Table S5. Gene Ontology (GO) enrichment analysis results for GNG12-associated genes in neuroblast

ONTOLOGY,"ID","Description","setSize","enrichmentScore","NES","pvalue","p.adjust","qvalue","ranl

BP,"GO:0006261","DNA-templated DNA replication",153,-0.691501406208541,-3.49030897084349,

CC,"GO:0098687","chromosomal region",320,-0.609049981563165,-3.37110482980753,1e-10,1.332

BP,"GO:0022613","ribonucleoprotein complex biogenesis",387,-0.586559000785767,-3.3064660333

CC,"GO:0062023","collagen-containing extracellular matrix",419,0.622223756928907,3.2852337349

CC,"GO:0000228","nuclear chromosome",193,-0.620804259952194,-3.2328406938282,1e-10,1.332

BP,"GO:0000819","sister chromatid segregation",186,-0.620613430366867,-3.21258246534091,1e-

CC,"GO:0000775","chromosome, centromeric region",196,-0.615461686166419,-3.20814744482112

MF,"GO:0140097","catalytic activity, acting on DNA",209,-0.60819485437218,-3.20209493001014,1

BP,"GO:0042254","ribosome biogenesis",281,-0.582649186214344,-3.18733078293648,1e-10,1.332

BP,"GO:0000070","mitotic sister chromatid segregation",161,-0.627975897472868,-3.182453246709

BP,"GO:0007059","chromosome segregation",313,-0.577262930774516,-3.18213711204669,1e-10,

BP,"GO:0032392","DNA geometric change",90,-0.69516461147157,-3.17167046785703,1e-10,1.332

MF,"GO:0005201","extracellular matrix structural constituent",169,0.661805991649657,3.16809244

BP,"GO:0000725","recombinational repair",152,-0.628377259852032,-3.1657811922283,1e-10,1.33

BP,"GO:0000724","double-strand break repair via homologous recombination",151,-0.626475482733

MF,"GO:0008094","ATP-dependent activity, acting on DNA",98,-0.679898426519146,-3.1552067841

BP,"GO:0006302","double-strand break repair",245,-0.584831621614919,-3.150734541599,1e-10,1.

BP,"GO:0006281","DNA repair",476,-0.542718273187727,-3.14399576928656,1e-10,1.3324590163

MF,"GO:0003678","DNA helicase activity",67,-0.720803991891163,-3.14387037068994,1e-10,1.332

BP,"GO:0006260","DNA replication",269,-0.578312928538059,-3.14288823919698,1e-10,1.3324590

BP,"GO:0098813","nuclear chromosome segregation",256,-0.582179073551851,-3.1401434907599,

BP,"GO:0071103","DNA conformation change",97,-0.67276664646157,-3.12047309863073,1e-10,1.

BP,"GO:0006397","mRNA processing",448,-0.542097240557803,-3.11639334192502,1e-10,1.332459

CC,"GO:0000793","condensed chromosome",232,-0.584606093834235,-3.10790364836942,1e-10,1

BP,"GO:0001525","angiogenesis",493,0.580117261017655,3.10094753501878,1e-10,1.3324590163

CC,"GO:0034708","methyltransferase complex",90,-0.678733732587434,-3.0967050100967,1e-10,1

CC,"GO:0000779","condensed chromosome, centromeric region",144,-0.620803819448571,-3.09234

BP,"GO:0033044","regulation of chromosome organization",199,-0.593446547950816,-3.088763552

BP,"GO:0032508","DNA duplex unwinding",84,-0.68465019804809,-3.07389911721772,1e-10,1.332

CC,"GO:0005681","spliceosomal complex",186,-0.593748577574394,-3.0735175486116,1e-10,1.332

CC,"GO:0000776","kinetochore",136,-0.625596838923358,-3.0721079181063,1e-10,1.3324590163

BP,"GO:0006364","rRNA processing",208,-0.584215091402252,-3.07172889709081,1e-10,1.332459

BP,"GO:0000377","RNA splicing, via transesterification reactions with bulged adenosine as nucleoph

BP,"GO:0000398","mRNA splicing, via spliceosome",281,-0.558911575392856,-3.05747628476704,1

BP,"GO:0000375","RNA splicing, via transesterification reactions",285,-0.557423614464953,-3.0433

BP,"GO:0044786","cell cycle DNA replication",42,-0.76700159166507,-3.03069519305819,1e-10,1.3

BP,"GO:0022618","ribonucleoprotein complex assembly",157,-0.599094471056734,-3.02511529600

CC,"GO:0000781","chromosome, telomeric region",136,-0.614828603276403,-3.01922852368356,1

BP,"GO:0016072","rRNA metabolic process",240,-0.562401349384295,-3.01639403629297,1e-10,1.

BP,"GO:0030198","extracellular matrix organization",296,0.590116098428915,3.00357960762237,1

BP,"GO:0034470","ncRNA processing",382,-0.530064291242845,-3.00177978563963,1e-10,1.33245

BP,"GO:0042060","wound healing",409,0.569170546667928,2.99387379380921,1e-10,1.332459016

BP,"GO:0043062","extracellular structure organization",297,0.587490540068228,2.9917751478439

BP,"GO:0045229","external encapsulating structure organization",299,0.587662236254983,2.99129

BP,"GO:0010594","regulation of endothelial cell migration",158,0.628398944008783,2.9816788107

BP,"GO:0043542","endothelial cell migration",205,0.609020613003933,2.98113909629659,1e-10,1.

MF,"GO:0003697","single-stranded DNA binding",110,-0.631053783909635,-2.98026798108356,1e-

BP,"GO:0006310","DNA recombination",286,-0.543880059107178,-2.97605959935723,1e-10,1.3324

BP,"GO:0071826","ribonucleoprotein complex subunit organization",165,-0.588455286482246,-2.97
BP,"GO:0033260","nuclear DNA replication",37,-0.773700048949352,-2.97359998536619,1e-10,1.3
BP,"GO:0060326","cell chemotaxis",292,0.586162066630755,2.97108350362329,1e-10,1.33245901
BP,"GO:0006270","DNA replication initiation",38,-0.771848563181422,-2.96339300215048,1e-10,1.
BP,"GO:0090329","regulation of DNA-templated DNA replication",51,-0.716705706241336,-2.96068
MF,"GO:0017116","single-stranded DNA helicase activity",23,-0.878904557756957,-2.95281073836
BP,"GO:1901342","regulation of vasculature development",266,0.583618057037819,2.9466656677
BP,"GO:0030335","positive regulation of cell migration",494,0.549117991027069,2.9372782785681
BP,"GO:0043534","blood vessel endothelial cell migration",111,0.657514897936861,2.93720941981
BP,"GO:0008380","RNA splicing",393,-0.519717855775575,-2.93584118184895,1e-10,1.332459016
BP,"GO:0045765","regulation of angiogenesis",262,0.582001862786258,2.92587448773592,1e-10,1
CC,"GO:0097525","spliceosomal snRNP complex",56,-0.70160985641532,-2.92343269195202,1e-10,
CC,"GO:0005788","endoplasmic reticulum lumen",302,0.571153775124633,2.92280418575661,1e-1
BP,"GO:0031589","cell-substrate adhesion",350,0.566793153499196,2.92262720390974,1e-10,1.33
MF,"GO:0004386","helicase activity",149,-0.582816522706993,-2.91668670920626,1e-10,1.332459
BP,"GO:0050900","leukocyte migration",365,0.563363124905596,2.91484043016874,1e-10,1.33245
CC,"GO:0035097","histone methyltransferase complex",62,-0.676941193500159,-2.8978488808230
BP,"GO:0034660","ncRNA metabolic process",493,-0.498009457898501,-2.8977697485395,1e-10,1.
BP,"GO:0043535","regulation of blood vessel endothelial cell migration",86,0.666248035804885,2.8
BP,"GO:0030595","leukocyte chemotaxis",222,0.586864113299345,2.88943089303962,1e-10,1.332
BP,"GO:0006399","tRNA metabolic process",157,-0.572162335987916,-2.88912203002505,1e-10,1.3
CC,"GO:0120114","Sm-like protein family complex",76,-0.650442689245156,-2.88420663789999,1e
BP,"GO:0010632","regulation of epithelial cell migration",217,0.585607159695483,2.877774799714
BP,"GO:0010631","epithelial cell migration",283,0.568592793503487,2.87577356424413,1e-10,1.33
CC,"GO:0044815","DNA packaging complex",95,-0.622031211224019,-2.87529083044308,1e-10,1.3
BP,"GO:0003158","endothelium development",121,0.635396417193443,2.87386215547539,1e-10,1
BP,"GO:0002685","regulation of leukocyte migration",208,0.584261198761452,2.87034511859676,1
MF,"GO:0019838","growth factor binding",131,0.624806648999997,2.86719631034891,1e-10,1.332
BP,"GO:0090130","tissue migration",291,0.566096884360221,2.86410417769686,1e-10,1.33245901
CC,"GO:0071013","catalytic step 2 spliceosome",85,-0.632974458094261,-2.85704042440175,1e-10
BP,"GO:2001251","negative regulation of chromosome organization",83,-0.638366540768413,-2.856
BP,"GO:0090132","epithelium migration",286,0.56408089167167,2.85617546290375,1e-10,1.33245
BP,"GO:0019882","antigen processing and presentation",101,0.64366685445766,2.8554193224083
MF,"GO:0005178","integrin binding",147,0.609512895723304,2.85460883898147,1e-10,1.3324590
BP,"GO:0002274","myeloid leukocyte activation",218,0.581290524494296,2.85032082712733,1e-10
CC,"GO:0030684","preribosome",71,-0.646651840434013,-2.84624733182232,1e-10,1.3324590163
BP,"GO:0000387","spliceosomal snRNP assembly",37,-0.738560912521466,-2.83854798981619,1e-1
BP,"GO:0071674","mononuclear cell migration",188,0.585025114663071,2.83295948678802,1e-10,
MF,"GO:0003735","structural constituent of ribosome",158,-0.560943830156022,-2.8269124261862
MF,"GO:0005518","collagen binding",66,0.686934508872684,2.8268771683447,1e-10,1.332459016
BP,"GO:0140014","mitotic nuclear division",278,-0.51705287806196,-2.82516561460702,1e-10,1.33
CC,"GO:0071005","U2-type precatalytic spliceosome",49,-0.687557742341283,-2.82168799683346,
CC,"GO:0030532","small nuclear ribonucleoprotein complex",65,-0.660315784107205,-2.820938304
CC,"GO:0043202","lysosomal lumen",94,0.645456716910524,2.81946861248461,1e-10,1.33245901
CC,"GO:0009897","external side of plasma membrane",365,0.543572241711648,2.81244241380561
BP,"GO:0048002","antigen processing and presentation of peptide antigen",58,0.700351814659555,
BP,"GO:0045766","positive regulation of angiogenesis",155,0.593923960607785,2.81125066589341
BP,"GO:1904018","positive regulation of vasculature development",155,0.593923960607785,2.8112
BP,"GO:0010595","positive regulation of endothelial cell migration",98,0.635067014067222,2.8039
MF,"GO:0140375","immune receptor activity",141,0.603742717958548,2.80390760515957,1e-10,1.

CC,"GO:0046540","U4/U6 x U5 tri-snRNP complex",30,-0.768076410964274,-2.80213247248776,1e-
CC,"GO:0097526","spliceosomal tri-snRNP complex",30,-0.768076410964274,-2.80213247248776,1
BP,"GO:0097529","myeloid leukocyte migration",214,0.570670154618005,2.80025860846535,1e-1
BP,"GO:0034341","response to interferon-gamma",136,0.605744148688211,2.79972483440991,1e-
BP,"GO:0045785","positive regulation of cell adhesion",427,0.529814109298378,2.79792223003987
BP,"GO:0007160","cell-matrix adhesion",227,0.567025330292988,2.79509988602908,1e-10,1.3324
CC,"GO:0005604","basement membrane",93,0.638316716597229,2.78898202392875,1e-10,1.3324
BP,"GO:0045446","endothelial cell differentiation",105,0.627093580196762,2.78868591224203,1e-
BP,"GO:0007596","blood coagulation",218,0.568302779316056,2.78663625113813,1e-10,1.332459
BP,"GO:0051306","mitotic sister chromatid separation",66,-0.644274196554109,-2.78612777149685
BP,"GO:0050921","positive regulation of chemotaxis",138,0.601864876966745,2.78561543261846,1
CC,"GO:0005684","U2-type spliceosomal complex",88,-0.616583020376941,-2.78435806180899,1e-
BP,"GO:0007599","hemostasis",224,0.563987851701172,2.78121099270702,1e-10,1.3324590163
BP,"GO:0008033","tRNA processing",114,-0.58206035174279,-2.77768141126673,1e-10,1.332459
BP,"GO:0050817","coagulation",222,0.563986865699807,2.77679455276283,1e-10,1.3324590163
CC,"GO:0030055","cell-substrate junction",419,0.525611413561173,2.77513728477415,1e-10,1.332
BP,"GO:0002687","positive regulation of leukocyte migration",137,0.600861410827739,2.7737354
BP,"GO:0048660","regulation of smooth muscle cell proliferation",132,0.603879259679859,2.77195
BP,"GO:0034502","protein localization to chromosome",82,-0.620024862640498,-2.7713915609992
CC,"GO:0071011","precatalytic spliceosome",51,-0.670644933943197,-2.77040763802937,1e-10,1.3
BP,"GO:0000280","nuclear division",414,-0.485193356147563,-2.76486284976832,1e-10,1.332459
BP,"GO:0051983","regulation of chromosome segregation",85,-0.612479729073122,-2.764533896
CC,"GO:0005925","focal adhesion",411,0.524554289276975,2.76416483088788,1e-10,1.332459016
BP,"GO:0010634","positive regulation of epithelial cell migration",140,0.593223032289053,2.76129
BP,"GO:0010965","regulation of mitotic sister chromatid separation",63,-0.643248320942919,-2.760
BP,"GO:0007091","metaphase/anaphase transition of mitotic cell cycle",61,-0.649855360827302,-2.
MF,"GO:0140101","catalytic activity, acting on a tRNA",99,-0.597422573201786,-2.75911021781733
BP,"GO:0001667","ameboidal-type cell migration",402,0.523917343085588,2.75565350051815,1e-1
BP,"GO:0042116","macrophage activation",97,0.625964164474863,2.75489050310319,1e-10,1.332
BP,"GO:0051304","chromosome separation",92,-0.599487085237169,-2.7525952243452,1e-10,1.33
CC,"GO:0044391","ribosomal subunit",173,-0.538178108562095,-2.75203352725915,1e-10,1.33245
MF,"GO:0050840","extracellular matrix binding",52,0.695950222951884,2.75084306007488,1e-10,1
BP,"GO:0070374","positive regulation of ERK1 and ERK2 cascade",200,0.558492781652149,2.74481
MF,"GO:0140098","catalytic activity, acting on RNA",339,-0.489367699282352,-2.74347534049363,
CC,"GO:0030667","secretory granule membrane",298,0.53974207527617,2.74235990546479,1e-10,
BP,"GO:0002237","response to molecule of bacterial origin",326,0.534979937801157,2.7417923667
BP,"GO:0033045","regulation of sister chromatid segregation",70,-0.625495834417207,-2.74097527
BP,"GO:0034508","centromere complex assembly",27,-0.777274746776758,-2.73897067063749,1e-
BP,"GO:0007229","integrin-mediated signaling pathway",100,0.61845777941577,2.73615808075818
BP,"GO:0001570","vasculogenesis",79,0.643309199184977,2.73594936998272,1e-10,1.3324590163
BP,"GO:0033002","muscle cell proliferation",183,0.565286343699671,2.73574444609345,1e-10,1.3
BP,"GO:0048659","smooth muscle cell proliferation",134,0.594706471675714,2.73149417051725,1
BP,"GO:0035924","cellular response to vascular endothelial growth factor stimulus",58,0.680004985
BP,"GO:0071824","protein-DNA complex subunit organization",158,-0.541475107400884,-2.728798
BP,"GO:0035150","regulation of tube size",138,0.588749518689348,2.72491352788867,1e-10,1.332
BP,"GO:0044784","metaphase/anaphase transition of cell cycle",64,-0.634543924372151,-2.723880
CC,"GO:0000123","histone acetyltransferase complex",82,-0.608871549960295,-2.72153840429284
CC,"GO:0031248","protein acetyltransferase complex",92,-0.592669050138347,-2.72128964443431
CC,"GO:1902493","acetyltransferase complex",92,-0.592669050138347,-2.72128964443431,1e-10,1
BP,"GO:0042274","ribosomal small subunit biogenesis",68,-0.61972582778805,-2.72042144549467,

BP,"GO:0032835","glomerulus development",65,0.658768010241731,2.71847072394938,1e-10,1.33
BP,"GO:0030071","regulation of mitotic metaphase/anaphase transition",59,-0.64161291899745,-2.
BP,"GO:0000075","cell cycle checkpoint signaling",161,-0.535645339426032,-2.71454088668205,1e
CC,"GO:0032432","actin filament bundle",70,0.653763285806984,2.71394881990767,1e-10,1.33245
MF,"GO:0019955","cytokine binding",138,0.586157771100956,2.71291812434218,1e-10,1.3324590
BP,"GO:0032496","response to lipopolysaccharide",310,0.528979457277084,2.71154855339548,1e-
BP,"GO:0035296","regulation of tube diameter",137,0.586965925526616,2.70959024602829,1e-10,
BP,"GO:0097746","blood vessel diameter maintenance",137,0.586965925526616,2.7095902460282
BP,"GO:0032611","interleukin-1 beta production",92,0.620715686438738,2.70879809108028,1e-10
BP,"GO:0032651","regulation of interleukin-1 beta production",92,0.620715686438738,2.70879809
BP,"GO:0097530","granulocyte migration",147,0.578363698146568,2.70872386205377,1e-10,1.332
CC,"GO:0070821","tertiary granule membrane",71,0.650224946244048,2.70762043400106,1e-10,1.
CC,"GO:0070820","tertiary granule",160,0.570642742167194,2.70739347185884,1e-10,1.33245901
BP,"GO:0071621","granulocyte chemotaxis",122,0.596149314949319,2.70572109119284,1e-10,1.33
CC,"GO:0015934","large ribosomal subunit",109,-0.572083544562858,-2.70528840453415,1e-10,1.3
BP,"GO:0006325","chromatin organization",490,-0.465243000126374,-2.70424877228943,1e-10,1.3
CC,"GO:0005657","replication fork",55,-0.651871657472716,-2.70367565585665,1e-10,1.33245901
CC,"GO:0005775","vacuolar lumen",166,0.565546853655549,2.70336561100841,1e-10,1.3324590
BP,"GO:0035825","homologous recombination",54,-0.651499555406718,-2.70085836145006,1e-10,
BP,"GO:0032635","interleukin-6 production",140,0.580135807717907,2.7003756026486,1e-10,1.33
BP,"GO:0032675","regulation of interleukin-6 production",140,0.580135807717907,2.700375602648
BP,"GO:0006338","chromatin remodeling",191,-0.518799031933499,-2.69722390512035,1e-10,1.33
BP,"GO:0042273","ribosomal large subunit biogenesis",68,-0.614331287949996,-2.69674093839624
CC,"GO:0042581","specific granule",158,0.567726051373983,2.69379309724181,1e-10,1.33245901
BP,"GO:0031123","RNA 3'-end processing",103,-0.578694952064689,-2.69119268957071,1e-10,1.33
BP,"GO:0030199","collagen fibril organization",60,0.659543664210017,2.68482483304963,1e-10,1.3
BP,"GO:0030336","negative regulation of cell migration",255,0.533921624625551,2.6821325230458
CC,"GO:0001725","stress fiber",64,0.65450576251168,2.68206351103625,1e-10,1.33245901639344
CC,"GO:0097517","contractile actin filament bundle",64,0.65450576251168,2.68206351103625,1e-1
BP,"GO:0032677","regulation of interleukin-8 production",82,0.626925381788962,2.680745722050
BP,"GO:0006909","phagocytosis",225,0.543395438125035,2.67936355055071,1e-10,1.3324590163
BP,"GO:1902099","regulation of metaphase/anaphase transition of cell cycle",62,-0.6257767950807
CC,"GO:0045335","phagocytic vesicle",134,0.582878967070405,2.67717029576583,1e-10,1.332459
CC,"GO:0030139","endocytic vesicle",334,0.520386466272431,2.67649654325305,1e-10,1.3324590
BP,"GO:1905818","regulation of chromosome separation",70,-0.610596009948102,-2.67568298816
BP,"GO:0032637","interleukin-8 production",83,0.62326190975743,2.67451760630913,1e-10,1.332
BP,"GO:1903053","regulation of extracellular matrix organization",46,0.692325164328467,2.673310
BP,"GO:0010569","regulation of double-strand break repair via homologous recombination",62,-0.62
BP,"GO:0032612","interleukin-1 production",108,0.598470941449337,2.66231528854326,1e-10,1.3
BP,"GO:0032652","regulation of interleukin-1 production",108,0.598470941449337,2.662315288543
MF,"GO:0042393","histone binding",238,-0.497677945389846,-2.66160181138796,1e-10,1.332459
BP,"GO:0050729","positive regulation of inflammatory response",130,0.580782343295887,2.661104
BP,"GO:0071346","cellular response to interferon-gamma",115,0.590209822587589,2.65876248766
BP,"GO:0051568","histone H3-K4 methylation",67,-0.609478506524855,-2.65831132983677,1e-10,
BP,"GO:0030168","platelet activation",129,0.580360325030155,2.65614171079024,1e-10,1.332459
BP,"GO:1904705","regulation of vascular associated smooth muscle cell proliferation",52,0.6719318
BP,"GO:0071706","tumor necrosis factor superfamily cytokine production",160,0.559764247285077
BP,"GO:1903555","regulation of tumor necrosis factor superfamily cytokine production",160,0.5597
BP,"GO:0048662","negative regulation of smooth muscle cell proliferation",51,0.673972275635498,
BP,"GO:0060191","regulation of lipase activity",86,0.610798978324348,2.65321995331159,1e-10,1.

BP,"GO:0006403","RNA localization",189,-0.510270879807715,-2.65307423853719,1e-10,1.332459(
BP,"GO:1903510","mucopolysaccharide metabolic process",86,0.609559557797978,2.64783609481
BP,"GO:0031497","chromatin assembly",128,-0.542096429832244,-2.64717336940745,1e-10,1.332
BP,"GO:0043410","positive regulation of MAPK cascade",464,0.497149788311755,2.6470559456457
BP,"GO:0019221","cytokine-mediated signaling pathway",457,0.498974576720999,2.646887291172
CC,"GO:0070603","SWI/SNF superfamily-type complex",84,-0.58947867997514,-2.64660406023424
BP,"GO:0032103","positive regulation of response to external stimulus",411,0.502007939944184,2.6
BP,"GO:0002042","cell migration involved in sprouting angiogenesis",53,0.665350798351863,2.6449
BP,"GO:1990868","response to chemokine",95,0.60400462707322,2.64464065116254,1e-10,1.3324
BP,"GO:1990869","cellular response to chemokine",95,0.60400462707322,2.64464065116254,1e-10
BP,"GO:0048285","organelle fission",456,-0.456228134963436,-2.63935344398546,1e-10,1.332459(
BP,"GO:0002040","sprouting angiogenesis",119,0.583790402677263,2.63878626215535,1e-10,1.33
BP,"GO:0046718","viral entry into host cell",141,0.568171540458247,2.63870760828868,1e-10,1.33
CC,"GO:0035579","specific granule membrane",89,0.60481192748739,2.63603616993122,1e-10,1.3
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BP,"GO:0002683","negative regulation of immune system process",394,0.501847723464121,2.6322(
BP,"GO:0032963","collagen metabolic process",91,0.602708309212878,2.63001532796484,1e-10,1.
BP,"GO:0045667","regulation of osteoblast differentiation",117,0.582286663375758,2.6299276369
BP,"GO:0050673","epithelial cell proliferation",383,0.504117739703876,2.62983725663591,1e-10,1
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CC,"GO:1904949","ATPase complex",85,-0.582224168668064,-2.62797015689477,1e-10,1.3324590
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BP,"GO:0010810","regulation of cell-substrate adhesion",205,0.536465507893027,2.625983858750
BP,"GO:0044409","entry into host",148,0.559580240081833,2.62446986403751,1e-10,1.332459016
BP,"GO:0061448","connective tissue development",243,0.527149282086498,2.62136708844985,1e
BP,"GO:0071219","cellular response to molecule of bacterial origin",205,0.534990803805135,2.6187
CC,"GO:0098798","mitochondrial protein-containing complex",237,-0.488807136834818,-2.618400
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BP,"GO:0001704","formation of primary germ layer",117,0.579225452259318,2.61610151964474,1
BP,"GO:0007093","mitotic cell cycle checkpoint signaling",122,-0.540805547095729,-2.6121290889
MF,"GO:0003684","damaged DNA binding",66,-0.603655716128185,-2.61047542199112,1e-10,1.33
BP,"GO:0070371","ERK1 and ERK2 cascade",305,0.50944932728668,2.61030025226511,1e-10,1.332
BP,"GO:0061041","regulation of wound healing",121,0.576860270374403,2.60910646514015,1e-10
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BP,"GO:0070555","response to interleukin-1",131,0.568354953215174,2.60814321908262,1e-10,1.3
BP,"GO:0008015","blood circulation",486,0.488222960735241,2.60547573891349,1e-10,1.3324590
BP,"GO:0050818","regulation of coagulation",70,0.627284046016697,2.60402630951128,1e-10,1.33
BP,"GO:0006405","RNA export from nucleus",82,-0.582212377674348,-2.60237704553401,1e-10,1.3
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BP,"GO:0001818","negative regulation of cytokine production",263,0.516192644296474,2.6004428
BP,"GO:0071347","cellular response to interleukin-1",105,0.584331976043497,2.59852500667246,1
BP,"GO:0001510","RNA methylation",72,-0.587038906391122,-2.59658568393433,1e-10,1.332459(
BP,"GO:0006275","regulation of DNA replication",127,-0.53053804315995,-2.59595038348229,1e-1
BP,"GO:0000723","telomere maintenance",127,-0.530513332152698,-2.59582947123183,1e-10,1.3
BP,"GO:0071222","cellular response to lipopolysaccharide",197,0.530087440002473,2.5955159457
BP,"GO:0032200","telomere organization",131,-0.531309647367033,-2.59521972645291,1e-10,1.33
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BP,"GO:0071260","cellular response to mechanical stimulus",70,0.624805118088617,2.5937356069
BP,"GO:0032602","chemokine production",91,0.594287106410193,2.59326804555241,1e-10,1.3324
BP,"GO:0002688","regulation of leukocyte chemotaxis",119,0.573441029993102,2.5920061466624
BP,"GO:0071675","regulation of mononuclear cell migration",112,0.577161482683609,2.588224882
BP,"GO:0002761","regulation of myeloid leukocyte differentiation",113,0.57671157589969,2.58759
CC,"GO:0042641","actomyosin",71,0.621017492961877,2.58599683621579,1e-10,1.332459016393
BP,"GO:0048771","tissue remodeling",161,0.54370705691195,2.58505158530524,1e-10,1.3324590
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BP,"GO:0043414","macromolecule methylation",289,-0.470956651028373,-2.57990980952048,1e-1
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BP,"GO:0140053","mitochondrial gene expression",97,-0.555972965266072,-2.5787525151603,1e-1
BP,"GO:0051216","cartilage development",183,0.532531420185522,2.57722460728118,1e-10,1.332
BP,"GO:0002573","myeloid leukocyte differentiation",205,0.526113149652038,2.57530935080731,
BP,"GO:2001252","positive regulation of chromosome organization",83,-0.575403906597907,-2.575
BP,"GO:0003018","vascular process in circulatory system",258,0.512098796430216,2.57488445542
BP,"GO:0070372","regulation of ERK1 and ERK2 cascade",282,0.509746579797069,2.573191461628
CC,"GO:0005581","collagen trimer",85,0.596409795368607,2.5727812046674,1e-10,1.3324590163
MF,"GO:0016887","ATP hydrolysis activity",296,-0.469240372582027,-2.57254715559515,1e-10,1.3
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BP,"GO:0016571","histone methylation",146,-0.515326224655451,-2.57037913151756,1e-10,1.332
BP,"GO:0007200","phospholipase C-activating G protein-coupled receptor signaling pathway",102,0.
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BP,"GO:0050920","regulation of chemotaxis",213,0.523411174018387,2.5684513762274,1e-10,1.33
BP,"GO:1903557","positive regulation of tumor necrosis factor superfamily cytokine production",102
BP,"GO:0022407","regulation of cell-cell adhesion",431,0.485334569915999,2.56539090089926,1e-
BP,"GO:0061982","meiosis I cell cycle process",120,-0.532102071102312,-2.56485877348844,1e-10,
BP,"GO:1902749","regulation of cell cycle G2/M phase transition",98,-0.552346021194507,-2.56327
CC,"GO:0098857","membrane microdomain",316,0.500281765206294,2.56288875990636,1e-10,1.3
BP,"GO:0001503","ossification",386,0.489731180152714,2.56244705187865,1e-10,1.332459016393
BP,"GO:0010389","regulation of G2/M transition of mitotic cell cycle",89,-0.564987965764925,-2.56
BP,"GO:0022612","gland morphogenesis",122,0.564184847497483,2.56064513189138,1e-10,1.3324
BP,"GO:0002460","adaptive immune response based on somatic recombination of immune receptor
BP,"GO:0032755","positive regulation of interleukin-6 production",90,0.586744844185159,2.558393
BP,"GO:0018022","peptidyl-lysine methylation",122,-0.529558982584783,-2.5578073859652,1e-10,
BP,"GO:0090287","regulation of cellular response to growth factor stimulus",278,0.506410661763
CC,"GO:0031091","platelet alpha granule",88,0.587151768663063,2.55694530554241,1e-10,1.3324
BP,"GO:0006282","regulation of DNA repair",132,-0.521783139661105,-2.55599959895095,1e-10,1.
CC,"GO:0045121","membrane raft",315,0.497971728029493,2.55208851045165,1e-10,1.332459016
BP,"GO:0032760","positive regulation of tumor necrosis factor production",98,0.577297327246977,
BP,"GO:0071216","cellular response to biotic stimulus",230,0.516544303124153,2.54880405659688
BP,"GO:0002250","adaptive immune response",423,0.483547202988343,2.54870712561792,1e-10,
BP,"GO:2000779","regulation of double-strand break repair",94,-0.550738349244185,-2.544597590
BP,"GO:0008217","regulation of blood pressure",178,0.526308959514702,2.54236519889221,1e-10
BP,"GO:0050878","regulation of body fluid levels",375,0.488313605673744,2.54182520486691,1e-1
BP,"GO:0070098","chemokine-mediated signaling pathway",88,0.583101055447398,2.539305143844
BP,"GO:0022409","positive regulation of cell-cell adhesion",280,0.502430089342149,2.537393940833

BP,"GO:0002064","epithelial cell development",198,0.518059975602645,2.53610076849434,1e-10,
BP,"GO:0030193","regulation of blood coagulation",66,0.615969220235266,2.53484037065444,1e-
BP,"GO:0034446","substrate adhesion-dependent cell spreading",106,0.569017647278691,2.533238
BP,"GO:0001935","endothelial cell proliferation",143,0.542993819169802,2.53292486516435,1e-10
BP,"GO:0002062","chondrocyte differentiation",102,0.569158471218153,2.53261151848948,1e-10,
BP,"GO:0007162","negative regulation of cell adhesion",276,0.501133862005046,2.5318144212264
BP,"GO:0040013","negative regulation of locomotion",305,0.494101401577878,2.53166103889564,
BP,"GO:0030278","regulation of ossification",110,0.567171967783122,2.53138720302536,1e-10,1.3
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BP,"GO:0008213","protein alkylation",175,-0.49443690056962,-2.53127811747533,1e-10,1.332459
BP,"GO:1903034","regulation of response to wounding",150,0.540089303273841,2.5301227985577
BP,"GO:0034109","homotypic cell-cell adhesion",91,0.57935167477691,2.52809487052749,1e-10,1.
BP,"GO:0010811","positive regulation of cell-substrate adhesion",119,0.559251352909348,2.527867
CC,"GO:0000428","DNA-directed RNA polymerase complex",96,-0.545255154470754,-2.526517108
BP,"GO:0007127","meiosis I",114,-0.528989863032316,-2.52442088675832,1e-10,1.332459016393
BP,"GO:0050866","negative regulation of cell activation",202,0.515479298613246,2.523606195363
BP,"GO:0071560","cellular response to transforming growth factor beta stimulus",236,0.510134651
BP,"GO:0072676","lymphocyte migration",112,0.562275850639865,2.52147170399931,1e-10,1.332
BP,"GO:0051028","mRNA transport",120,-0.523045193946187,-2.5212024674978,1e-10,1.3324590
BP,"GO:0034728","nucleosome organization",84,-0.561412493021173,-2.52059427078639,1e-10,1.
BP,"GO:0009451","RNA modification",151,-0.499944351300643,-2.51982209963951,1e-10,1.33245
BP,"GO:0150115","cell-substrate junction organization",98,0.570675153937046,2.51961053980646,
BP,"GO:0007044","cell-substrate junction assembly",92,0.577247708787138,2.51910419827515,1e-
CC,"GO:0055029","nuclear DNA-directed RNA polymerase complex",95,-0.543754194351308,-2.513
BP,"GO:0070663","regulation of leukocyte proliferation",244,0.504280379495133,2.5133216467663
BP,"GO:0007051","spindle organization",178,-0.489946551352116,-2.51267770588863,1e-10,1.332
BP,"GO:0052126","movement in host environment",172,0.522968240973289,2.50982343505107,1e
BP,"GO:0044839","cell cycle G2/M phase transition",144,-0.503856963598793,-2.50980610001721,
BP,"GO:0045132","meiotic chromosome segregation",82,-0.559763689859641,-2.5020357403479,1
BP,"GO:0002224","toll-like receptor signaling pathway",115,0.55531826822477,2.50158742259716,
BP,"GO:0071559","response to transforming growth factor beta",242,0.503181946190616,2.501091
BP,"GO:0050764","regulation of phagocytosis",93,0.572333056923897,2.50068119157239,1e-10,1.3
BP,"GO:0070661","leukocyte proliferation",320,0.486462059091001,2.49861055782891,1e-10,1.33
BP,"GO:0000086","G2/M transition of mitotic cell cycle",131,-0.511379191765534,-2.497867999845
MF,"GO:0030246","carbohydrate binding",258,0.496567022730209,2.49678912900945,1e-10,1.332
BP,"GO:0034968","histone lysine methylation",115,-0.522787040992229,-2.49590430116649,1e-10,
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BP,"GO:0001655","urogenital system development",349,0.483374762526683,2.49402437882033,1e
CC,"GO:0005766","primary lysosome",146,0.532370413964597,2.49196648924162,1e-10,1.332459
CC,"GO:0042582","azurophil granule",146,0.532370413964597,2.49196648924162,1e-10,1.332459
MF,"GO:0005539","glycosaminoglycan binding",219,0.507851961784653,2.48690740036443,1e-10,
BP,"GO:0007159","leukocyte cell-cell adhesion",358,0.481527621387204,2.48679013695928,1e-10,
CC,"GO:0030880","RNA polymerase complex",98,-0.535831414904993,-2.48663454647739,1e-10,1.
BP,"GO:0007179","transforming growth factor beta receptor signaling pathway",186,0.513587541
BP,"GO:0001763","morphogenesis of a branching structure",196,0.507528981150939,2.483567184
BP,"GO:0001936","regulation of endothelial cell proliferation",127,0.546559290363455,2.4833737
BP,"GO:0072006","nephron development",144,0.53043990568167,2.47918303097188,1e-10,1.3324
BP,"GO:1903046","meiotic cell cycle process",187,-0.478183427987971,-2.47890452380316,1e-10,1
BP,"GO:0002443","leukocyte mediated immunity",368,0.477874555885385,2.47856101515321,1e-

CC,"GO:0005814","centriole",129,-0.507380613568031,-2.47772167828868,1e-10,1.332459016393
BP,"GO:0061138","morphogenesis of a branching epithelium",182,0.512101042785602,2.47590259
BP,"GO:0030203","glycosaminoglycan metabolic process",118,0.549949849390467,2.475874897009
BP,"GO:0050730","regulation of peptidyl-tyrosine phosphorylation",249,0.49382096251778,2.47306
BP,"GO:0016525","negative regulation of angiogenesis",90,0.566632567030182,2.47069791399251,
BP,"GO:0072001","renal system development",309,0.481803163261385,2.47009225101941,1e-10,1
BP,"GO:0032259","methylation",328,-0.442816183953608,-2.46926973495342,1e-10,1.3324590163
BP,"GO:0030316","osteoclast differentiation",94,0.565200168947923,2.46889387679976,1e-10,1.33
BP,"GO:0140013","meiotic nuclear division",168,-0.48517205300336,-2.46859809380248,1e-10,1.33
CC,"GO:0031983","vesicle lumen",319,0.480241837885212,2.46822143828081,1e-10,1.3324590163
BP,"GO:0002367","cytokine production involved in immune response",99,0.554727348265592,2.464
CC,"GO:0034774","secretory granule lumen",314,0.481084429740277,2.46358335423869,1e-10,1.3
BP,"GO:1905952","regulation of lipid localization",142,0.52826785847303,2.4621843286186,1e-10,
MF,"GO:0005126","cytokine receptor binding",260,0.489236529811517,2.46132124770306,1e-10,1
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BP,"GO:0022408","negative regulation of cell-cell adhesion",182,0.508221395890014,2.4571453048
BP,"GO:0031570","DNA integrity checkpoint signaling",115,-0.514661251643386,-2.4571099336786
BP,"GO:0002221","pattern recognition receptor signaling pathway",162,0.514723134076097,2.4546
CC,"GO:0000922","spindle pole",160,-0.484206239052919,-2.45217232889046,1e-10,1.3324590163
BP,"GO:0050731","positive regulation of peptidyl-tyrosine phosphorylation",185,0.50657212440529
BP,"GO:0007052","mitotic spindle organization",117,-0.51191075512849,-2.45140696410746,1e-10
BP,"GO:0007266","Rho protein signal transduction",130,0.534974113733405,2.45121448577486,1e
BP,"GO:0002698","negative regulation of immune effector process",112,0.546525732913055,2.4508
BP,"GO:0090101","negative regulation of transmembrane receptor protein serine/threonine kinase
CC,"GO:0060205","cytoplasmic vesicle lumen",317,0.47699282585972,2.44623357657679,1e-10,1.3
BP,"GO:0007204","positive regulation of cytosolic calcium ion concentration",310,0.4771179842513
BP,"GO:0043123","positive regulation of I-kappaB kinase/NF-kappaB signaling",186,0.504164264148
BP,"GO:0051321","meiotic cell cycle",244,-0.453624987285421,-2.43866195274768,1e-10,1.332459
BP,"GO:0001822","kidney development",300,0.477970705887817,2.43419525106139,1e-10,1.3324
BP,"GO:0002718","regulation of cytokine production involved in immune response",97,0.552756231
BP,"GO:0051053","negative regulation of DNA metabolic process",124,-0.500627534872807,-2.4301
MF,"GO:0005125","cytokine activity",230,0.492401130886015,2.42967349031728,1e-10,1.3324590
BP,"GO:0050657","nucleic acid transport",152,-0.481961435359472,-2.42813441052857,1e-10,1.33
BP,"GO:0050658","RNA transport",152,-0.481961435359472,-2.42813441052857,1e-10,1.33245901
BP,"GO:0043491","protein kinase B signaling",184,0.501770896079846,2.4262546256595,1e-10,1.3
BP,"GO:0006022","aminoglycan metabolic process",129,0.529983373509388,2.42558094289175,1e
BP,"GO:0071356","cellular response to tumor necrosis factor",212,0.494935529129747,2.425486058
MF,"GO:0008083","growth factor activity",154,0.512789599750072,2.42460891242672,1e-10,1.332
BP,"GO:0051052","regulation of DNA metabolic process",430,-0.423839556180474,-2.42274722138
BP,"GO:0030282","bone mineralization",114,0.538583915066234,2.41959915452281,1e-10,1.33245
MF,"GO:0008757","S-adenosylmethionine-dependent methyltransferase activity",131,-0.495117711
BP,"GO:0050867","positive regulation of cell activation",339,0.469334393552458,2.4124261640669
BP,"GO:0051236","establishment of RNA localization",155,-0.478072972482189,-2.41174713496748
BP,"GO:1903844","regulation of cellular response to transforming growth factor beta stimulus",123,
BP,"GO:0001649","osteoblast differentiation",215,0.489385637879979,2.4060254848822,1e-10,1.3
BP,"GO:0090257","regulation of muscle system process",219,0.491058633702059,2.4046719163493
BP,"GO:0046634","regulation of alpha-beta T cell activation",102,0.540139071400031,2.403482514
BP,"GO:0000956","nuclear-transcribed mRNA catabolic process",113,-0.50473700046443,-2.401522
BP,"GO:0034612","response to tumor necrosis factor",231,0.484893027775907,2.40089324421649,
BP,"GO:0016570","histone modification",466,-0.41516925128202,-2.40082364794369,1e-10,1.3324

BP,"GO:1903037","regulation of leukocyte cell-cell adhesion",322,0.467837585407432,2.400543562
BP,"GO:0001952","regulation of cell-matrix adhesion",117,0.531271899554244,2.39951683467488,
BP,"GO:0032944","regulation of mononuclear cell proliferation",225,0.486623646340529,2.399434
BP,"GO:0110148","biomineralization",162,0.503133829472662,2.39933807343365,1e-10,1.332459
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BP,"GO:0007178","transmembrane receptor protein serine/threonine kinase signaling pathway",344
BP,"GO:0031214","biomineral tissue development",160,0.505301958131541,2.39738652868381,1e-
BP,"GO:0051480","regulation of cytosolic calcium ion concentration",340,0.465191497295014,2.393
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BP,"GO:0007249","I-kappaB kinase/NF-kappaB signaling",278,0.473629758981599,2.392010973869
BP,"GO:0001909","leukocyte mediated cytotoxicity",124,0.526050467331053,2.39167082912739,1e
BP,"GO:0006937","regulation of muscle contraction",155,0.505133548905754,2.3909744679659,1e-
BP,"GO:0050777","negative regulation of immune response",170,0.499131052880424,2.389176481
BP,"GO:0003012","muscle system process",409,0.45406913327168,2.38843293392143,1e-10,1.332
CC,"GO:0072686","mitotic spindle",151,-0.473638639499236,-2.38723591525422,1e-10,1.3324590
BP,"GO:0007098","centrosome cycle",126,-0.490373139798553,-2.38709733431244,1e-10,1.33245
BP,"GO:0045834","positive regulation of lipid metabolic process",144,0.510659407513435,2.386732
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BP,"GO:0007369","gastrulation",180,0.492241336920805,2.38165588237298,1e-10,1.33245901639
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BP,"GO:0072507","divalent inorganic cation homeostasis",492,0.442705008572182,2.366688261095
BP,"GO:0072503","cellular divalent inorganic cation homeostasis",473,0.444825443585475,2.36663
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BP,"GO:0042129","regulation of T cell proliferation",167,0.482986393460825,2.30850014137735,1e
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MF,"GO:0016741","transferase activity, transferring one-carbon groups",191,-0.440355133486395,-
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BP,"GO:1903522","regulation of blood circulation",244,0.457750135094226,2.28141599420214,1e-
BP,"GO:0009612","response to mechanical stimulus",195,0.466949720856193,2.28112660485839,1
BP,"GO:0001837","epithelial to mesenchymal transition",149,0.486114479256688,2.279816187849
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BP,"GO:0030099","myeloid cell differentiation",365,0.440577266086868,2.27954647904656,1e-10,
BP,"GO:0010959","regulation of metal ion transport",372,0.438951032751005,2.27901831439557,1
BP,"GO:1903311","regulation of mRNA metabolic process",279,-0.416199667384807,-2.277424893
BP,"GO:0032970","regulation of actin filament-based process",367,0.43917967874036,2.274286535
BP,"GO:0018394","peptidyl-lysine acetylation",177,-0.443027469123821,-2.27291854470452,1e-10,
BP,"GO:0007015","actin filament organization",417,0.430920000669374,2.2727712081608,1e-10,1.
BP,"GO:0045216","cell-cell junction organization",192,0.466313629519472,2.27147871966638,1e-1
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BP,"GO:1901987","regulation of cell cycle phase transition",370,-0.399027810403801,-2.260140344

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BP,"GO:0010876","lipid localization",419,0.427593132645476,2.25761772766324,1e-10,1.33245901
BP,"GO:0051048","negative regulation of secretion",153,0.479419158275078,2.2564724490536,1e-
BP,"GO:0015850","organic hydroxy compound transport",238,0.45417987459568,2.2556567142352
BP,"GO:0042110","T cell activation",490,0.421867026193808,2.25318011646218,1e-10,1.33245901
BP,"GO:0002764","immune response-regulating signaling pathway",384,0.431094610302273,2.2515
BP,"GO:1901888","regulation of cell junction assembly",194,0.461362877458592,2.2500818768082
BP,"GO:0002699","positive regulation of immune effector process",230,0.452924122348603,2.2348
BP,"GO:0006869","lipid transport",368,0.430675927511717,2.23375894562487,1e-10,1.332459016
BP,"GO:0042445","hormone metabolic process",227,0.453108172714549,2.23355560016847,1e-10,
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BP,"GO:0042063","gliogenesis",278,0.440606298965591,2.22522990225064,1e-10,1.332459016393
BP,"GO:0051249","regulation of lymphocyte activation",437,0.41895113348893,2.21970348254273,
CC,"GO:0030016","myofibril",225,0.450026703476612,2.21898282810441,1e-10,1.3324590163944
BP,"GO:0060485","mesenchyme development",286,0.438149868080866,2.2185344704346,1e-10,1.
BP,"GO:1902105","regulation of leukocyte differentiation",273,0.440003941610834,2.21550452753
CC,"GO:0043292","contractile fiber",235,0.447954674612073,2.21470977627506,1e-10,1.33245901
BP,"GO:0048705","skeletal system morphogenesis",222,0.449251090780176,2.21189190310887,1e
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BP,"GO:0051054","positive regulation of DNA metabolic process",246,-0.410792457770101,-2.2096
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CC,"GO:0019866","organelle inner membrane",483,-0.38050724773901,-2.20600189917482,1e-10,
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BP,"GO:0070302","regulation of stress-activated protein kinase signaling cascade",192,0.451001764
BP,"GO:0032956","regulation of actin cytoskeleton organization",327,0.427872357827748,2.196449
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BP,"GO:0006473","protein acetylation",206,-0.416396470983263,-2.18677957091015,1e-10,1.3324
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BP,"GO:1903706","regulation of hemopoiesis",343,0.42467059368797,2.18392730501653,1e-10,1.3
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BP,"GO:0006816","calcium ion transport",394,0.415445827890261,2.17902065566909,1e-10,1.3324
MF,"GO:0051015","actin filament binding",209,0.443647877605042,2.1788029600255,1e-10,1.3324
CC,"GO:0005765","lysosomal membrane",379,0.417252016501061,2.17650030743867,1e-10,1.3324
CC,"GO:0098852","lytic vacuole membrane",379,0.417252016501061,2.17650030743867,1e-10,1.33
BP,"GO:0019216","regulation of lipid metabolic process",310,0.424388795638188,2.175416850426
CC,"GO:0061695","transferase complex, transferring phosphorus-containing groups",246,-0.404235
BP,"GO:0031098","stress-activated protein kinase signaling cascade",241,0.436807244624956,2.170
BP,"GO:0050863","regulation of T cell activation",327,0.422810007940419,2.17046247185427,1e-1
BP,"GO:0110053","regulation of actin filament organization",247,0.432601219382502,2.166122856
BP,"GO:0060537","muscle tissue development",375,0.416015571561775,2.16549130134075,1e-10,
MF,"GO:0061134","peptidase regulator activity",219,0.441978466617804,2.16433055721782,1e-10
BP,"GO:0060249","anatomical structure homeostasis",302,0.422885891980828,2.16406283038095,
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BP,"GO:0002703","regulation of leukocyte mediated immunity",224,0.437828725159969,2.1590785

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CC,"GO:0005774","vacuolar membrane",421,0.408414828890034,2.15514623906867,1e-10,1.3324
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BP,"GO:0043068","positive regulation of programmed cell death",488,0.402303606123243,2.14637
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CC,"GO:0045178","basal part of cell",265,0.423874229576235,2.14039558368227,1e-10,1.3324590
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MF,"GO:0042277","peptide binding",296,0.419459283153443,2.13496861458583,1e-10,1.3324590
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BP,"GO:0003015","heart process",240,0.428463529685317,2.13032617309559,1e-10,1.332459016
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BP,"GO:0006401","RNA catabolic process",261,-0.392568376709041,-2.12920322858174,1e-10,1.33
BP,"GO:1901652","response to peptide",464,0.399646030957284,2.12790073991907,1e-10,1.3324
BP,"GO:0043065","positive regulation of apoptotic process",476,0.399311174621041,2.125934343
BP,"GO:0052548","regulation of endopeptidase activity",396,0.40400573358037,2.12208645269667
BP,"GO:0090066","regulation of anatomical structure size",466,0.397982908368766,2.11696500921
BP,"GO:0043270","positive regulation of ion transport",261,0.420491704367458,2.11447110965619
BP,"GO:0043543","protein acylation",247,-0.392911956228112,-2.11248461507576,1e-10,1.332459
BP,"GO:0042692","muscle cell differentiation",338,0.408927584475578,2.1039456480998,1e-10,1.3
BP,"GO:0043087","regulation of GTPase activity",353,0.407180765094997,2.10113165828303,1e-10
CC,"GO:0045177","apical part of cell",408,0.398851012432057,2.10015913074406,1e-10,1.3324590
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BP,"GO:0002831","regulation of response to biotic stimulus",327,0.402524832586747,2.066330093
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BP,"GO:0051047","positive regulation of secretion",294,0.401943208535085,2.04472695161304,1e-
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MF,"GO:0005543","phospholipid binding",445,0.379990488196828,2.01829270386271,1e-10,1.332
BP,"GO:0051090","regulation of DNA-binding transcription factor activity",419,0.382248845724267,
BP,"GO:0048608","reproductive structure development",426,0.379856434811658,2.004467794149
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BP,"GO:0045860","positive regulation of protein kinase activity",382,0.380417967640301,1.985475
BP,"GO:0007265","Ras protein signal transduction",335,0.386326038340023,1.98498954066018,1e-
BP,"GO:0008544","epidermis development",344,0.385390073906879,1.98369765606418,1e-10,1.3
CC,"GO:0005759","mitochondrial matrix",444,-0.347339225721544,-1.98334888528899,1e-10,1.33
BP,"GO:0030098","lymphocyte differentiation",374,0.379606665227931,1.97376978155857,1e-10,1
CC,"GO:0030133","transport vesicle",383,0.376483603972767,1.96400667991119,1e-10,1.3324590
BP,"GO:0045936","negative regulation of phosphate metabolic process",395,0.369735539328308,1.
BP,"GO:0048732","gland development",426,0.366354707561158,1.93322040971982,1e-10,1.33245
BP,"GO:0048568","embryonic organ development",445,0.363633743323646,1.93141500596784,1e-
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BP,"GO:0034765","regulation of ion transmembrane transport",442,0.355322929336881,1.8863403
BP,"GO:2001020","regulation of response to DNA damage stimulus",221,-0.398576936421909,-2.10
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BP,"GO:0007131","reciprocal meiotic recombination",52,-0.645455764795497,-2.6643546022087,1.
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BP,"GO:0002690","positive regulation of leukocyte chemotaxis",92,0.569133178152314,2.48369245
BP,"GO:0001906","cell killing",171,0.467577121404514,2.24640090551894,1.25699613730278e-10,
BP,"GO:0060541","respiratory system development",198,0.451570779032097,2.21061084365894,1
BP,"GO:0051250","negative regulation of lymphocyte activation",152,0.477014922876485,2.243283
BP,"GO:1900046","regulation of hemostasis",68,0.606513642736216,2.5058694003788,1.45882236
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BP,"GO:0060562","epithelial tube morphogenesis",325,0.390498349682356,2.00068002353117,1.5
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BP,"GO:0010608","post-transcriptional regulation of gene expression",479,-0.315120553954891,-1.8
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BP,"GO:0043536","positive regulation of blood vessel endothelial cell migration",49,0.669272075341
BP,"GO:0070167","regulation of biomineral tissue development",93,0.556022319085049,2.4294150
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BP,"GO:0051346","negative regulation of hydrolase activity",345,0.382474598443359,1.969840883
BP,"GO:0110149","regulation of biomineralization",95,0.551432258603307,2.41445197950097,1.93
CC,"GO:0042383","sarcolemma",128,0.505546118177919,2.30521458365461,1.92944794968458e-
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BP,"GO:0014065","phosphatidylinositol 3-kinase signaling",137,0.500970156488915,2.31261098905
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BP,"GO:0030324","lung development",174,0.462522493175716,2.22496208939856,2.25388950458
CC,"GO:0030670","phagocytic vesicle membrane",73,0.595689132131608,2.49419547668433,2.275
BP,"GO:0030323","respiratory tube development",178,0.465870962899835,2.25041603765033,2.28
BP,"GO:1902107","positive regulation of leukocyte differentiation",153,0.472614546460767,2.2244
BP,"GO:1903708","positive regulation of hemopoiesis",153,0.472614546460767,2.22444532034906
BP,"GO:0050728","negative regulation of inflammatory response",140,0.489211598629472,2.27714
BP,"GO:0032946","positive regulation of mononuclear cell proliferation",140,0.489135245096699,2.
BP,"GO:0002526","acute inflammatory response",110,0.52615610600893,2.34832627351209,2.781
BP,"GO:0006066","alcohol metabolic process",344,0.377958889396317,1.94544751862306,3.00899
BP,"GO:0050684","regulation of mRNA processing",137,-0.463565552477148,-2.28869249792314,3
BP,"GO:1900262","regulation of DNA-directed DNA polymerase activity",13,-0.907661425225828,-2.

BP,"GO:1900264","positive regulation of DNA-directed DNA polymerase activity",13,-0.9076614252
MF,"GO:0140030","modification-dependent protein binding",155,-0.442835714975167,-2.2339848
BP,"GO:0045786","negative regulation of cell cycle",344,-0.342251835903085,-1.92106239420132,3
CC,"GO:0030136","clathrin-coated vesicle",185,0.448913818306996,2.17240142962445,3.3442372
BP,"GO:0050819","negative regulation of coagulation",51,0.650154716430787,2.55975100354598,3
BP,"GO:0045598","regulation of fat cell differentiation",127,0.509115271534661,2.3132412783324
BP,"GO:0150063","visual system development",376,0.37046289973949,1.92735930711936,3.67017
BP,"GO:0070192","chromosome organization involved in meiotic cell cycle",62,-0.607875223330635
BP,"GO:0070527","platelet aggregation",69,0.598474285130029,2.48090478140996,3.7019331062
BP,"GO:0002822","regulation of adaptive immune response based on somatic recombination of imm
BP,"GO:2001236","regulation of extrinsic apoptotic signaling pathway",149,0.476665781502372,2.2
BP,"GO:0050671","positive regulation of lymphocyte proliferation",138,0.484798030851245,2.2437
BP,"GO:0061045","negative regulation of wound healing",70,0.597961841182638,2.4823018796845
MF,"GO:0030020","extracellular matrix structural constituent conferring tensile strength",41,0.6999
BP,"GO:0045841","negative regulation of mitotic metaphase/anaphase transition",40,-0.680277138
CC,"GO:0031093","platelet alpha granule lumen",65,0.615481378619718,2.53984419841478,4.405
BP,"GO:0016064","immunoglobulin mediated immune response",134,0.489820231249761,2.24975
BP,"GO:0048880","sensory system development",382,0.362119243256096,1.8899709945713,4.740
BP,"GO:0045787","positive regulation of cell cycle",304,-0.360926800827219,-1.97745682027953,4.
BP,"GO:0000245","spliceosomal complex assembly",56,-0.611496986300377,-2.54795491316822,4.
BP,"GO:0003007","heart morphogenesis",241,0.416585839651357,2.07022181317427,4.86202953
BP,"GO:0032872","regulation of stress-activated MAPK cascade",189,0.441641464085031,2.140104
BP,"GO:0001894","tissue homeostasis",259,0.413033886575817,2.07536049187309,5.330954011
BP,"GO:0035265","organ growth",155,0.472711146032013,2.23750784982274,5.44257900099539e
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BP,"GO:0031032","actomyosin structure organization",190,0.438336970946657,2.12794890714602
BP,"GO:0051985","negative regulation of chromosome segregation",44,-0.658617946743911,-2.620
BP,"GO:1905819","negative regulation of chromosome separation",44,-0.658617946743911,-2.6201
BP,"GO:0051225","spindle assembly",114,-0.483711352891275,-2.30834488094201,5.63890539881
BP,"GO:0060349","bone morphogenesis",97,0.53909922993411,2.37259484338318,5.75691937234
BP,"GO:0001885","endothelial cell development",63,0.620141185571185,2.52239391818213,5.806
BP,"GO:0071897","DNA biosynthetic process",178,-0.429280486292751,-2.20155342353992,5.8664
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BP,"GO:0006956","complement activation",56,0.625742429568888,2.51824331303546,6.4257635
BP,"GO:0014812","muscle cell migration",78,0.572204717615744,2.42617061601209,6.548316603
MF,"GO:0070851","growth factor receptor binding",137,0.49442015030062,2.28237442487393,6.64
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MF,"GO:0004866","endopeptidase inhibitor activity",173,0.456913798704498,2.199140949981,6.68
BP,"GO:0006417","regulation of translation",363,-0.329914214873299,-1.86387659918396,6.75957
BP,"GO:0002286","T cell activation involved in immune response",115,0.507859932622328,2.28779
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BP,"GO:1903531","negative regulation of secretion by cell",130,0.497096745958661,2.2776629994
BP,"GO:0043030","regulation of macrophage activation",55,0.628800500191165,2.52738581142267
BP,"GO:0070482","response to oxygen levels",299,0.392713149324366,1.99897326806135,7.15591
BP,"GO:0030490","maturation of SSU-rRNA",46,-0.646688058741686,-2.61687926680689,7.708651
BP,"GO:0072009","nephron epithelium development",111,0.526024604168621,2.34982420514834,
BP,"GO:0060047","heart contraction",231,0.412395894656734,2.04193185034112,7.725522790498
BP,"GO:0033047","regulation of mitotic sister chromatid segregation",45,-0.654871321600887,-2.63
BP,"GO:0006271","DNA strand elongation involved in DNA replication",15,-0.885284179864511,-2.6

CC,"GO:0101002","ficolin-1-rich granule",181,0.440049675489393,2.13125663199556,8.076285636
BP,"GO:0001933","negative regulation of protein phosphorylation",305,0.386870289609698,1.9822
BP,"GO:0032757","positive regulation of interleukin-8 production",62,0.620311447161395,2.529127
BP,"GO:0150076","neuroinflammatory response",65,0.610287281116824,2.51841024628074,8.336
CC,"GO:0030027","lamellipodium",194,0.437124024818003,2.13186819793224,8.34935939493927
BP,"GO:0001666","response to hypoxia",265,0.404681770396845,2.04348132940303,8.517513174
BP,"GO:0090501","RNA phosphodiester bond hydrolysis",139,-0.455645888933917,-2.25644372196
BP,"GO:0051784","negative regulation of nuclear division",55,-0.608471559120954,-2.52367122088
BP,"GO:0002444","myeloid leukocyte mediated immunity",101,0.531417023934196,2.35745934079
BP,"GO:0030850","prostate gland development",49,0.652746949066946,2.5492086926894,9.02535
BP,"GO:0045669","positive regulation of osteoblast differentiation",62,0.619000680294854,2.52378
MF,"GO:0004518","nuclease activity",187,-0.40972318599133,-2.12400639548408,9.209806506367
BP,"GO:0000018","regulation of DNA recombination",109,-0.489474910635314,-2.31464584611315
BP,"GO:0051146","striated muscle cell differentiation",250,0.40644430998907,2.02980212617852,9
BP,"GO:0008016","regulation of heart contraction",196,0.436683570874,2.13688878255731,9.3177
BP,"GO:0032722","positive regulation of chemokine production",68,0.591695572374419,2.4446471
CC,"GO:0098636","protein complex involved in cell adhesion",43,0.679843945040247,2.591345013
BP,"GO:0010543","regulation of platelet activation",47,0.669035656583565,2.59363028712556,9.98
BP,"GO:0032409","regulation of transporter activity",261,0.401442230907211,2.01867953786657,1
BP,"GO:0036293","response to decreased oxygen levels",277,0.395324516905558,1.996080570763
BP,"GO:0046635","positive regulation of alpha-beta T cell activation",67,0.595952519640124,2.4561
BP,"GO:0031532","actin cytoskeleton reorganization",108,0.51786135917478,2.30372123087203,1.
BP,"GO:0031341","regulation of cell killing",97,0.53297531430811,2.34564327337753,1.065550959
CC,"GO:0045334","clathrin-coated endocytic vesicle",87,0.553518584814313,2.39816510881825,1.0
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BP,"GO:0033048","negative regulation of mitotic sister chromatid segregation",42,-0.669250477796
BP,"GO:2000816","negative regulation of mitotic sister chromatid separation",42,-0.6692504777963
BP,"GO:1902100","negative regulation of metaphase/anaphase transition of cell cycle",42,-0.668975
CC,"GO:0098800","inner mitochondrial membrane protein complex",114,-0.47904635767954,-2.286
BP,"GO:0042326","negative regulation of phosphorylation",342,0.376859718544938,1.9372282227
BP,"GO:0030500","regulation of bone mineralization",74,0.582729343838823,2.44314918476465,1.
BP,"GO:0070588","calcium ion transmembrane transport",290,0.390675335597974,1.980818567041
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BP,"GO:0010001","glial cell differentiation",211,0.426491881081518,2.09123729948314,1.2715725
BP,"GO:0014068","positive regulation of phosphatidylinositol 3-kinase signaling",77,0.587013252084
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BP,"GO:0022616","DNA strand elongation",36,-0.696534740594061,-2.67074088506687,1.3060841
BP,"GO:0019884","antigen processing and presentation of exogenous antigen",45,0.6725097665475
MF,"GO:0048306","calcium-dependent protein binding",79,0.570096210204435,2.42457960979642
BP,"GO:0010466","negative regulation of peptidase activity",244,0.41147686632759,2.0507911022
CC,"GO:0001726","ruffle",172,0.448965869740744,2.15467206826228,1.45993399418662e-09,1.60
BP,"GO:0032732","positive regulation of interleukin-1 production",68,0.587254781332074,2.426299
BP,"GO:0045600","positive regulation of fat cell differentiation",63,0.612100841551326,2.48969021
BP,"GO:0006809","nitric oxide biosynthetic process",66,0.57924633187263,2.38371811179694,1.54
MF,"GO:0061135","endopeptidase regulator activity",186,0.437920819076398,2.11943461692274,1
BP,"GO:0019724","B cell mediated immunity",137,0.488467057171312,2.25489336954282,1.58988
BP,"GO:0030509","BMP signaling pathway",152,0.462958425510593,2.17717910778433,1.6099012
CC,"GO:0044665","MLL1/2 complex",32,-0.718415922045802,-2.65606348488441,1.618577794531
MF,"GO:0042562","hormone binding",84,0.562110915234994,2.42231359011027,1.670108896111
BP,"GO:0002478","antigen processing and presentation of exogenous peptide antigen",36,0.708087

BP,"GO:0003014","renal system process",108,0.513784162301062,2.28558370268225,1.834413241
BP,"GO:0051168","nuclear export",151,-0.430629883516326,-2.17046296138086,1.8373682595416
BP,"GO:0048247","lymphocyte chemotaxis",60,0.614761118876111,2.50252713795517,1.98719754
BP,"GO:0044706","multi-multicellular organism process",207,0.429182104685669,2.1107684230409
BP,"GO:0014706","striated muscle tissue development",224,0.415869495504374,2.0507902944000
BP,"GO:0000077","DNA damage checkpoint signaling",107,-0.484495900195567,-2.27825568064074
CC,"GO:0005770","late endosome",265,0.400073076737495,2.02020926692148,2.12552615806109
BP,"GO:0097553","calcium ion transmembrane import into cytosol",143,0.473619207042195,2.2093
BP,"GO:0031099","regeneration",174,0.446963740578652,2.15011678955355,2.14504324679489e-
BP,"GO:0009100","glycoprotein metabolic process",354,0.370493862096341,1.91387223804445,2.1
BP,"GO:0002504","antigen processing and presentation of peptide or polysaccharide antigen via MH
MF,"GO:0004857","enzyme inhibitor activity",365,0.364331839826264,1.88505269474518,2.18154
CC,"GO:0005796","Golgi lumen",97,0.526895801216464,2.3188871205014,2.2204574743275e-09,2
BP,"GO:0022604","regulation of cell morphogenesis",293,0.385064366515169,1.95623426396461,2
BP,"GO:0001654","eye development",372,0.362406826591096,1.88160348977148,2.368355939778
BP,"GO:2000379","positive regulation of reactive oxygen species metabolic process",64,0.602022130
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BP,"GO:0062013","positive regulation of small molecule metabolic process",134,0.477619477312640
BP,"GO:0051303","establishment of chromosome localization",76,-0.548907704529015,-2.43397807
BP,"GO:0060193","positive regulation of lipase activity",65,0.598800950623803,2.47101077835663,
BP,"GO:0071692","protein localization to extracellular region",339,0.366036585176932,1.881465000
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BP,"GO:0051017","actin filament bundle assembly",149,0.463613168797124,2.17428785240029,2.7
BP,"GO:0061572","actin filament bundle organization",149,0.463613168797124,2.17428785240029
BP,"GO:0033108","mitochondrial respiratory chain complex assembly",82,-0.529206326126616,-2.3
CC,"GO:0016607","nuclear speck",390,-0.319671968162625,-1.80319773274028,2.84295241499516
CC,"GO:0071339","MLL1 complex",31,-0.715567525824952,-2.65650079189066,3.01775891465576
BP,"GO:0061437","renal system vasculature development",28,0.753285730278956,2.579397374984
BP,"GO:0061440","kidney vasculature development",28,0.753285730278956,2.5793973749843,3.01
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BP,"GO:0002456","T cell mediated immunity",104,0.518176006932101,2.30234673874633,3.143431
BP,"GO:1903038","negative regulation of leukocyte cell-cell adhesion",128,0.484938096364519,2.21
CC,"GO:0032993","protein-DNA complex",123,-0.460973782271205,-2.23420931209943,3.3227744
BP,"GO:0072593","reactive oxygen species metabolic process",207,0.42596144870949,2.094928855
BP,"GO:0016055","Wnt signaling pathway",424,0.344257491697739,1.81395267060761,3.35889294
BP,"GO:0019229","regulation of vasoconstriction",63,0.603631545455715,2.45524176511188,3.421
BP,"GO:0048844","artery morphogenesis",69,0.577005142363138,2.39190697437539,3.547179805
BP,"GO:0046209","nitric oxide metabolic process",71,0.578254458776951,2.40792605340775,3.589
BP,"GO:0043010","camera-type eye development",325,0.374460061092688,1.9185094237858,3.68
BP,"GO:1903532","positive regulation of secretion by cell",269,0.393305039288328,1.984354825061
BP,"GO:0042476","odontogenesis",126,0.494171333250072,2.25361065367754,3.76991245762821
BP,"GO:0030856","regulation of epithelial cell differentiation",139,0.474766389144248,2.205168285
BP,"GO:0010517","regulation of phospholipase activity",64,0.597979183458682,2.45042632193046
BP,"GO:0048041","focal adhesion assembly",83,0.543070659256842,2.33040398733434,3.941983640
MF,"GO:0004527","exonuclease activity",74,-0.547981141330206,-2.42848044074674,3.981049761
BP,"GO:0072012","glomerulus vasculature development",26,0.767377898743779,2.5819609906564
BP,"GO:0007080","mitotic metaphase plate congression",51,-0.611896871101488,-2.527721719179
BP,"GO:0030100","regulation of endocytosis",200,0.420356172431227,2.06591390536274,4.256571
BP,"GO:0010951","negative regulation of endopeptidase activity",235,0.410091342570367,2.027513

BP,"GO:0001706","endoderm formation",54,0.624121317376201,2.49204777255563,4.3053422125
BP,"GO:0007272","ensheathment of neurons",136,0.469794505448347,2.17137110910843,4.32988
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BP,"GO:0010717","regulation of epithelial to mesenchymal transition",91,0.536017410269207,2.338
BP,"GO:0035023","regulation of Rho protein signal transduction",82,0.549476661564684,2.3495734
BP,"GO:0031663","lipopolysaccharide-mediated signaling pathway",60,0.606165726656738,2.46753
MF,"GO:1901681","sulfur compound binding",248,0.395771383205701,1.98389434497398,4.54259
BP,"GO:0006406","mRNA export from nucleus",60,-0.586818822401716,-2.47613411249772,4.8601
MF,"GO:0016779","nucleotidyltransferase activity",109,-0.47450571113202,-2.24385897900904,5.0
BP,"GO:0006023","aminoglycan biosynthetic process",80,0.551502532861362,2.34141656498983,5
BP,"GO:0044703","multi-organism reproductive process",199,0.421948066722328,2.069529028920
BP,"GO:0008608","attachment of spindle microtubules to kinetochore",33,-0.700189489381406,-2.6
BP,"GO:0007094","mitotic spindle assembly checkpoint signaling",38,-0.670574863513508,-2.57456
BP,"GO:0071173","spindle assembly checkpoint signaling",38,-0.670574863513508,-2.57456831915
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BP,"GO:0003073","regulation of systemic arterial blood pressure",94,0.531239564837366,2.320547
BP,"GO:1900047","negative regulation of hemostasis",49,0.634337517022078,2.47731332149948,5
BP,"GO:0198738","cell-cell signaling by wnt",426,0.343998030702437,1.81524626306695,5.929412
BP,"GO:1901222","regulation of NIK/NF-kappaB signaling",92,0.538510227118903,2.350054150582
BP,"GO:0030111","regulation of Wnt signaling pathway",311,0.372366566756184,1.9046879827503
BP,"GO:0014909","smooth muscle cell migration",65,0.59036833493486,2.43621276369849,6.1389
BP,"GO:0007043","cell-cell junction assembly",141,0.470911858155431,2.18701327762017,6.14367
BP,"GO:0034764","positive regulation of transmembrane transport",200,0.417415693915747,2.0514
BP,"GO:0071711","basement membrane organization",32,0.707593150986426,2.54017492751583,6
BP,"GO:0010633","negative regulation of epithelial cell migration",66,0.565240117555509,2.326079
BP,"GO:0062197","cellular response to chemical stress",310,0.37496969010454,1.92209452897059,
CC,"GO:0009925","basal plasma membrane",247,0.400450434476888,2.00513729576998,7.008279
BP,"GO:0050766","positive regulation of phagocytosis",68,0.573750385898769,2.37050485623545,
BP,"GO:0072080","nephron tubule development",91,0.531490320121559,2.31924409738131,7.496
BP,"GO:0043406","positive regulation of MAP kinase activity",115,0.488086800968458,2.19872435
BP,"GO:0006024","glycosaminoglycan biosynthetic process",76,0.559270676478353,2.35606977994
BP,"GO:0098742","cell-cell adhesion via plasma-membrane adhesion molecules",269,0.389181258
BP,"GO:0050000","chromosome localization",78,-0.525113399710669,-2.34000732066124,8.18280
MF,"GO:0008173","RNA methyltransferase activity",58,-0.586962278698696,-2.48630474628571,8.
BP,"GO:0043903","regulation of biological process involved in symbiotic interaction",55,0.60671986
BP,"GO:2000027","regulation of animal organ morphogenesis",128,0.478392695525958,2.1813990
BP,"GO:0007492","endoderm development",77,0.56696413195601,2.40256417959261,8.7767179
BP,"GO:0032368","regulation of lipid transport",107,0.506935507869848,2.2562435453777,8.95766
BP,"GO:0090049","regulation of cell migration involved in sprouting angiogenesis",37,0.689638317
BP,"GO:1902903","regulation of supramolecular fiber organization",346,0.361587532481776,1.8626
BP,"GO:0006939","smooth muscle contraction",107,0.506120352236799,2.25261549090727,9.4814
BP,"GO:0048738","cardiac muscle tissue development",211,0.415147155731263,2.03561018474256
BP,"GO:0031507","heterochromatin assembly",64,-0.566335676941373,-2.43108551019475,9.9903
BP,"GO:0002285","lymphocyte activation involved in immune response",192,0.421953528799012,2.
BP,"GO:0071900","regulation of protein serine/threonine kinase activity",355,0.358672666970559,1
MF,"GO:0008009","chemokine activity",48,0.636337670543906,2.4732300627941,1.083350845176
BP,"GO:0070304","positive regulation of stress-activated protein kinase signaling cascade",129,0.475
BP,"GO:0061028","establishment of endothelial barrier",47,0.645253144895207,2.50143334364152
BP,"GO:0001910","regulation of leukocyte mediated cytotoxicity",82,0.541661856203809,2.316157
BP,"GO:0030195","negative regulation of blood coagulation",48,0.635217130398501,2.4688749009

BP,"GO:0071375","cellular response to peptide hormone stimulus",279,0.380537783781174,1.9200
BP,"GO:0007088","regulation of mitotic nuclear division",111,-0.481032651098492,-2.27749042645
BP,"GO:0045861","negative regulation of proteolysis",327,0.36572319152789,1.87741171540555,1.
BP,"GO:0034501","protein localization to kinetochore",17,-0.815265431068131,-2.53068530280745
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MF,"GO:0019199","transmembrane receptor protein kinase activity",78,0.545632713960143,2.3135
BP,"GO:0014066","regulation of phosphatidylinositol 3-kinase signaling",104,0.507226946400774,2.
BP,"GO:0038084","vascular endothelial growth factor signaling pathway",39,0.673589937216138,2.5
BP,"GO:0001774","microglial cell activation",43,0.657233154649634,2.50516000068742,1.2748623
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MF,"GO:0035064","methylated histone binding",74,-0.537253417663055,-2.3809385362276,1.3229
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BP,"GO:0061326","renal tubule development",94,0.524432296653069,2.29081263088245,1.325233
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BP,"GO:0016485","protein processing",234,0.394113428059087,1.94990081235311,1.39073815582
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BP,"GO:0036297","interstrand cross-link repair",27,-0.719379000595481,-2.53495690150016,1.4208
BP,"GO:0030512","negative regulation of transforming growth factor beta receptor signaling pathwa
BP,"GO:0042255","ribosome assembly",57,-0.589991191962621,-2.48385987354009,1.4683445837
BP,"GO:0060317","cardiac epithelial to mesenchymal transition",32,0.695451014664802,2.4965861
BP,"GO:0050680","negative regulation of epithelial cell proliferation",121,0.491932980743304,2.224
BP,"GO:0010596","negative regulation of endothelial cell migration",51,0.613259517052593,2.4144
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BP,"GO:0071773","cellular response to BMP stimulus",163,0.447119244538865,2.12963805135746,
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BP,"GO:2001057","reactive nitrogen species metabolic process",72,0.56488626829812,2.353479999
MF,"GO:0042379","chemokine receptor binding",63,0.589847078911953,2.39917412215515,1.6215
BP,"GO:0060402","calcium ion transport into cytosol",160,0.444363048217447,2.10826411514531,
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CC,"GO:0071162","CMG complex",11,-0.90928858162355,-2.44732260705789,1.693098246977996
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BP,"GO:0072678","T cell migration",62,0.590025334086886,2.40564571640457,1.78667719193582
BP,"GO:0071887","leukocyte apoptotic process",107,0.500030111009383,2.22550934575502,1.800
BP,"GO:0030217","T cell differentiation",258,0.384384230316124,1.93272272157588,1.799886242
BP,"GO:0032231","regulation of actin filament bundle assembly",97,0.509990716362712,2.2444872
BP,"GO:0043628","ncRNA 3'-end processing",41,-0.638318102289515,-2.49797249188483,1.832120
BP,"GO:0032964","collagen biosynthetic process",40,0.659724266480076,2.47885336290641,1.845
BP,"GO:0046890","regulation of lipid biosynthetic process",160,0.443274122350893,2.10309774647
BP,"GO:0022898","regulation of transmembrane transporter activity",244,0.395275479390413,1.97
CC,"GO:0022626","cytosolic ribosome",99,-0.483100732430352,-2.23113124089037,1.941057322
CC,"GO:0000151","ubiquitin ligase complex",290,-0.337554131489134,-1.8495919887199,1.950245
BP,"GO:0060070","canonical Wnt signaling pathway",291,0.374584204637902,1.89516709072716,1
BP,"GO:1901654","response to ketone",184,0.424212349436909,2.05122932223515,1.985318349
BP,"GO:0052372","modulation by symbiont of entry into host",49,0.622760255518038,2.432100003
BP,"GO:0016125","sterol metabolic process",145,0.454505333720859,2.12667009546478,1.991437
MF,"GO:0031491","nucleosome binding",49,-0.612658479057513,-2.51430675571789,2.013546573

CC,"GO:0000794","condensed nuclear chromosome",69,-0.546363755744101,-2.39483072795515,2
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BP,"GO:0070997","neuron death",339,0.355943103436523,1.8295834855612,2.24142139386936e-
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BP,"GO:0006352","DNA-templated transcription, initiation",111,-0.473680768867832,-2.242682308
BP,"GO:0045670","regulation of osteoclast differentiation",63,0.585845048183376,2.382896058054
BP,"GO:0051783","regulation of nuclear division",138,-0.432064785430931,-2.13684012425255,2.4
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BP,"GO:0071459","protein localization to chromosome, centromeric region",23,-0.75468930772944,
CC,"GO:1990391","DNA repair complex",37,-0.657419565865775,-2.52669340539991,2.780586660
CC,"GO:0031261","DNA replication preinitiation complex",12,-0.891026375083373,-2.48933983215
BP,"GO:0008154","actin polymerization or depolymerization",185,0.418863797686406,2.026982186
BP,"GO:0062012","regulation of small molecule metabolic process",306,0.36836497401913,1.88695
BP,"GO:0085029","extracellular matrix assembly",45,0.641557333424307,2.4588965561674,2.9549
BP,"GO:0046849","bone remodeling",88,0.533756407632318,2.3244176610525,2.98371631699103
BP,"GO:0003002","regionalization",349,0.356684316618512,1.84035131776317,2.98167881458073
BP,"GO:0010038","response to metal ion",342,0.359945811856516,1.85028314539295,3.01522794
BP,"GO:0045814","negative regulation of gene expression, epigenetic",71,-0.536597035121684,-2.3
BP,"GO:0090288","negative regulation of cellular response to growth factor stimulus",96,0.5152313
CC,"GO:0042611","MHC protein complex",22,0.770153531901343,2.51940013217026,3.072337315
BP,"GO:0014074","response to purine-containing compound",132,0.45604121474585,2.093345113
BP,"GO:0050868","negative regulation of T cell activation",117,0.478793660820461,2.16249617274
BP,"GO:0031124","mRNA 3'-end processing",60,-0.56736717706566,-2.39405616829735,3.1763891
BP,"GO:0002700","regulation of production of molecular mediator of immune response",156,0.4402
CC,"GO:0072562","blood microparticle",112,0.47951358893568,2.15033234098273,3.28483935930
BP,"GO:0048008","platelet-derived growth factor receptor signaling pathway",52,0.610233047867
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BP,"GO:0014910","regulation of smooth muscle cell migration",58,0.592834864609162,2.380194519
BP,"GO:0032609","interferon-gamma production",113,0.486444128189512,2.18257881418028,3.50
BP,"GO:0032649","regulation of interferon-gamma production",113,0.486444128189512,2.1825788
BP,"GO:0046578","regulation of Ras protein signal transduction",180,0.418748946355075,2.026070
BP,"GO:0042113","B cell activation",251,0.384288496058359,1.92021778481816,3.6165727304711
MF,"GO:0017056","structural constituent of nuclear pore",26,-0.738516654197811,-2.58365759469
BP,"GO:0003044","regulation of systemic arterial blood pressure mediated by a chemical signal",48,
BP,"GO:0030510","regulation of BMP signaling pathway",97,0.50357243382268,2.21624015286321,
MF,"GO:0004879","nuclear receptor activity",52,0.608714559985798,2.40603159203131,3.8489194
MF,"GO:0098531","ligand-activated transcription factor activity",52,0.608714559985798,2.4060315
CC,"GO:0008305","integrin complex",31,0.700899278817369,2.4650106193422,3.86144054618236
MF,"GO:0045296","cadherin binding",309,0.359697600966496,1.8440855615047,3.8619797105103
CC,"GO:0016323","basolateral plasma membrane",222,0.391723192091892,1.92865276151768,3.8
BP,"GO:0051383","kinetochore organization",21,-0.774546875330817,-2.55368613477758,3.89787
BP,"GO:0097696","receptor signaling pathway via STAT",169,0.426260284471778,2.0405254758794

CC,"GO:0000315","organellar large ribosomal subunit",53,-0.580108348750222,-2.40519821235255
CC,"GO:0005762","mitochondrial large ribosomal subunit",53,-0.580108348750222,-2.40519821235
BP,"GO:2000106","regulation of leukocyte apoptotic process",83,0.524309629282121,2.249897374
MF,"GO:0016893","endonuclease activity, active with either ribo- or deoxyribonucleic acids and proc
BP,"GO:0000462","maturation of SSU-rRNA from tricistronic rRNA transcript (SSU-rRNA, 5.8S rRNA, L
MF,"GO:0000217","DNA secondary structure binding",35,-0.664840015310277,-2.54685736075128,
BP,"GO:0035592","establishment of protein localization to extracellular region",332,0.35440204101
CC,"GO:0043596","nuclear replication fork",27,-0.706748264290591,-2.49044855174209,4.4832642
BP,"GO:0001656","metanephros development",89,0.513890545831212,2.23976083246972,4.48040
BP,"GO:0007009","plasma membrane organization",144,0.456479344470037,2.13350434738679,4.
MF,"GO:0003899","DNA-directed 5'-3' RNA polymerase activity",36,-0.65561722664797,-2.5138498
BP,"GO:0031297","replication fork processing",37,-0.650821660357837,-2.50133534609968,4.5106
BP,"GO:0001913","T cell mediated cytotoxicity",45,0.633968428779749,2.42981056412365,4.57908
BP,"GO:0046328","regulation of JNK cascade",138,0.450631665419265,2.08566169859389,4.78207
BP,"GO:0046632","alpha-beta T cell differentiation",110,0.488562802540499,2.18054081738069,4.9
BP,"GO:0060401","cytosolic calcium ion transport",176,0.421760495220555,2.03485388104359,4.9
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BP,"GO:0060759","regulation of response to cytokine stimulus",154,0.434168863310534,2.0528686
BP,"GO:0035987","endodermal cell differentiation",45,0.631990542360043,2.42222991956933,5.39
CC,"GO:0034399","nuclear periphery",126,-0.437761380280655,-2.13098748508568,5.3965218045
BP,"GO:0006949","syncytium formation",56,0.585552131594872,2.3565011259319,5.42576102339
BP,"GO:0034655","nucleobase-containing compound catabolic process",380,-0.311818053292925,-1
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BP,"GO:0045428","regulation of nitric oxide biosynthetic process",54,0.595346581865583,2.377153
BP,"GO:0051282","regulation of sequestering of calcium ion",117,0.473160456650537,2.137053516
BP,"GO:0003205","cardiac chamber development",162,0.432142566352698,2.06079585959881,6.4
MF,"GO:0004867","serine-type endopeptidase inhibitor activity",94,0.510249243940746,2.2288585
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BP,"GO:0060538","skeletal muscle organ development",159,0.42382589500006,2.01282279122667,
BP,"GO:1905314","semi-lunar valve development",41,0.648030098527809,2.44837585307986,6.758
CC,"GO:0022625","cytosolic large ribosomal subunit",56,-0.563395466241767,-2.34752791661718,7
BP,"GO:0009306","protein secretion",331,0.354816685162664,1.8242361191357,7.3206756582124
CC,"GO:0005902","microvillus",89,0.509420176118476,2.22027699671818,7.38192403984655e-08,
BP,"GO:0032874","positive regulation of stress-activated MAPK cascade",127,0.465627145655585,2
MF,"GO:0001653","peptide receptor activity",150,0.447631122980215,2.09698970657453,7.419714
BP,"GO:1901989","positive regulation of cell cycle phase transition",112,-0.456275303565993,-2.165
BP,"GO:0006979","response to oxidative stress",401,0.339384443497433,1.78322986704262,7.498(
BP,"GO:1902751","positive regulation of cell cycle G2/M phase transition",31,-0.675575858040857,-
BP,"GO:0070200","establishment of protein localization to telomere",17,-0.794221118810725,-2.46
MF,"GO:0001664","G protein-coupled receptor binding",270,0.372076275939309,1.8792984794180
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BP,"GO:0071604","transforming growth factor beta production",42,0.639239204889086,2.4183436
BP,"GO:0051209","release of sequestered calcium ion into cytosol",115,0.467653938267077,2.1066
CC,"GO:0005876","spindle microtubule",81,-0.489984014778521,-2.20124582159817,8.573834383

BP,"GO:0060740","prostate gland epithelium morphogenesis",26,0.733381774409476,2.467575800
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BP,"GO:0000727","double-strand break repair via break-induced replication",12,-0.88084750487872
BP,"GO:2000781","positive regulation of double-strand break repair",55,-0.563683530328126,-2.337
MF,"GO:0004536","deoxyribonuclease activity",55,-0.563658853928364,-2.33780791679528,8.8749
CC,"GO:0101003","ficolin-1-rich granule membrane",57,0.590054770195595,2.36918824565958,8.9
BP,"GO:0006284","base-excision repair",42,-0.62076631931026,-2.45286779113696,9.0857935577
BP,"GO:1903035","negative regulation of response to wounding",85,0.522752895892608,2.2550414
CC,"GO:0031674","I band",138,0.443931933960993,2.0546532844817,9.27707371444429e-08,7.48
BP,"GO:0010518","positive regulation of phospholipase activity",53,0.596361062254291,2.3707266
BP,"GO:0071248","cellular response to metal ion",186,0.409245276606014,1.98065168009194,9.33
CC,"GO:1902562","H4 histone acetyltransferase complex",39,-0.638168709287452,-2.450806581564
CC,"GO:0000792","heterochromatin",62,-0.559165609588609,-2.39367533177118,9.478467682239
MF,"GO:0090079","translation regulator activity, nucleic acid binding",101,-0.460054058210405,-2.1
CC,"GO:0035577","azurophil granule membrane",54,0.590781212912071,2.35892440254178,1.0035
BP,"GO:0044774","mitotic DNA integrity checkpoint signaling",78,-0.502348770403552,-2.23856371
BP,"GO:0043484","regulation of RNA splicing",151,-0.399607732611633,-2.01410495628563,1.0151
BP,"GO:1902106","negative regulation of leukocyte differentiation",98,0.493320222782866,2.17807
BP,"GO:0051298","centrosome duplication",71,-0.521486967769531,-2.29533235318396,1.027963
MF,"GO:0008528","G protein-coupled peptide receptor activity",144,0.449614012846031,2.101416
BP,"GO:0002706","regulation of lymphocyte mediated immunity",165,0.42622142040107,2.036169
CC,"GO:0005643","nuclear pore",91,-0.480918003230566,-2.19773552003351,1.04588022869643e-
BP,"GO:0002495","antigen processing and presentation of peptide antigen via MHC class II",30,0.71
BP,"GO:1903313","positive regulation of mRNA metabolic process",115,-0.442372951040044,-2.111
BP,"GO:0042246","tissue regeneration",63,0.570485376350945,2.32042134468794,1.13985448607
BP,"GO:0097028","dendritic cell differentiation",46,0.616505193680674,2.38054291183217,1.1775
BP,"GO:0001937","negative regulation of endothelial cell proliferation",38,0.657845108878878,2.44
CC,"GO:0044853","plasma membrane raft",106,0.485504675658569,2.16144264047702,1.1975015
CC,"GO:0030662","coated vesicle membrane",167,0.421870418106265,2.01638790041862,1.19945
BP,"GO:0048024","regulation of mRNA splicing, via spliceosome",102,-0.456021654682261,-2.11707
BP,"GO:0007568","aging",157,0.424066003200018,2.00581498565453,1.23834582919628e-07,9.77
BP,"GO:0070199","establishment of protein localization to chromosome",22,-0.750011878179115,-2
BP,"GO:0007143","female meiotic nuclear division",27,-0.695803318318165,-2.45188061146808,1.2
BP,"GO:0071677","positive regulation of mononuclear cell migration",65,0.558268412889665,2.303
CC,"GO:0042575","DNA polymerase complex",19,-0.782900387804754,-2.50730250686409,1.32920
BP,"GO:0071774","response to fibroblast growth factor",109,0.473597465723576,2.1060900925525
BP,"GO:1901991","negative regulation of mitotic cell cycle phase transition",158,-0.39040375234230
BP,"GO:0006790","sulfur compound metabolic process",318,0.353511139582832,1.8149184245937
BP,"GO:2000401","regulation of lymphocyte migration",61,0.565990901324545,2.30781349831305,
BP,"GO:0051283","negative regulation of sequestering of calcium ion",116,0.473652883148535,2.13
BP,"GO:1903707","negative regulation of hemopoiesis",100,0.489491767071673,2.1655914090093,
BP,"GO:0015918","sterol transport",88,0.519391360680938,2.26186034397268,1.46174633155251
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BP,"GO:0001823","mesonephros development",99,0.491827189687853,2.18519354106367,1.51777
BP,"GO:0032868","response to insulin",245,0.379435152715788,1.89228172421417,1.52033735078
BP,"GO:0043299","leukocyte degranulation",74,0.535564809208212,2.2454073075949,1.54111820
BP,"GO:0031062","positive regulation of histone methylation",48,-0.59531946906703,-2.437266874
BP,"GO:0051057","positive regulation of small GTPase mediated signal transduction",66,0.53733382
BP,"GO:0009991","response to extracellular stimulus",462,0.323420339731232,1.72061061914108,

BP,"GO:1901224","positive regulation of NIK/NF-kappaB signaling",64,0.561355716745228,2.30034
BP,"GO:0000768","syncytium formation by plasma membrane fusion",53,0.586843901124225,2.332
BP,"GO:0140253","cell-cell fusion",53,0.586843901124225,2.33289292854887,1.6503348260904e-0
BP,"GO:0045778","positive regulation of ossification",48,0.610130698242588,2.37137239390106,1.
BP,"GO:0043967","histone H4 acetylation",72,-0.519388426960353,-2.29735463725445,1.6861538
BP,"GO:0002429","immune response-activating cell surface receptor signaling pathway",216,0.3905
BP,"GO:0002757","immune response-activating signal transduction",216,0.390586323936154,1.918
MF,"GO:0001968","fibronectin binding",30,0.703772875951423,2.46828826990118,1.77188316888
MF,"GO:0000400","four-way junction DNA binding",17,-0.782648313159053,-2.42943771181823,1.
BP,"GO:0055123","digestive system development",138,0.439314864914087,2.03328407141956,1.7
BP,"GO:0090303","positive regulation of wound healing",52,0.590845843564884,2.3354029278857
MF,"GO:0045182","translation regulator activity",135,-0.417703455187501,-2.05552695387041,1.8
BP,"GO:0072132","mesenchyme morphogenesis",53,0.585961205242511,2.32938392900646,1.830
BP,"GO:0030261","chromosome condensation",33,-0.654446468605455,-2.43565931767678,1.8434
BP,"GO:0003170","heart valve development",64,0.559705608759376,2.29358712506227,1.8626386
BP,"GO:0016578","histone deubiquitination",38,-0.628247727335849,-2.41205983610337,1.875347
BP,"GO:2001233","regulation of apoptotic signaling pathway",347,0.349045105538958,1.79899151
MF,"GO:0005496","steroid binding",100,0.48588294263959,2.14962538115658,2.02667905826183
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BP,"GO:0010718","positive regulation of epithelial to mesenchymal transition",49,0.5976862720513
BP,"GO:0030038","contractile actin filament bundle assembly",100,0.485011368310107,2.1457693
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BP,"GO:1904064","positive regulation of cation transmembrane transport",139,0.440846122163863
BP,"GO:0007389","pattern specification process",449,0.320087996452962,1.69851508627589,2.221
BP,"GO:0034113","heterotypic cell-cell adhesion",60,0.571380630559602,2.32593683981094,2.244
CC,"GO:0098802","plasma membrane signaling receptor complex",179,0.403542834043496,1.95171
MF,"GO:0001618","virus receptor activity",73,0.534570790035923,2.23828835302738,2.271843238
BP,"GO:0002440","production of molecular mediator of immune response",203,0.39652939656884,
BP,"GO:0051145","smooth muscle cell differentiation",57,0.580765322074478,2.33188924833104,2
MF,"GO:0140272","exogenous protein binding",74,0.532343572585315,2.23190196122377,2.28327
CC,"GO:0005938","cell cortex",289,0.359370416909726,1.81958694578233,2.28793622749272e-07
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BP,"GO:0010971","positive regulation of G2/M transition of mitotic cell cycle",28,-0.6968036986151
BP,"GO:0006334","nucleosome assembly",52,-0.569564660191821,-2.35108632753217,2.34806001
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BP,"GO:0051928","positive regulation of calcium ion transport",115,0.458687301996493,2.0662860
BP,"GO:0003231","cardiac ventricle development",122,0.450815393458082,2.04609933740783,2.45
CC,"GO:0000118","histone deacetylase complex",70,-0.516532991303125,-2.2634909418616,2.463
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BP,"GO:0010863","positive regulation of phospholipase C activity",39,0.640168706137055,2.392547
CC,"GO:0030665","clathrin-coated vesicle membrane",106,0.478745831718417,2.13135260381,2.5
BP,"GO:0046638","positive regulation of alpha-beta T cell differentiation",49,0.595414167758978,2.
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BP,"GO:0051208","sequestering of calcium ion",121,0.468362140466846,2.11837554339771,2.5338
BP,"GO:0045638","negative regulation of myeloid cell differentiation",76,0.527228243769783,2.221
BP,"GO:0072659","protein localization to plasma membrane",269,0.368819594158215,1.860817605
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BP,"GO:0051960","regulation of nervous system development",423,0.325033060425135,1.7132020
BP,"GO:0009595","detection of biotic stimulus",37,0.650271172769014,2.40233891105275,2.61788

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BP,"GO:0051569","regulation of histone H3-K4 methylation",38,-0.622209593012655,-2.388877354(
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BP,"GO:0060512","prostate gland morphogenesis",28,0.704368124606729,2.4118939448918,3.3424
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BP,"GO:0009410","response to xenobiotic stimulus",395,0.327945944765938,1.72130368604677,3.
BP,"GO:1903036","positive regulation of response to wounding",63,0.557371351925239,2.2670806
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BP,"GO:0043537","negative regulation of blood vessel endothelial cell migration",35,0.671405625222
BP,"GO:0010812","negative regulation of cell-substrate adhesion",58,0.562111837885543,2.256843
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BP,"GO:0030048","actin filament-based movement",127,0.448998146757983,2.04009014273822,6.
BP,"GO:0006958","complement activation, classical pathway",36,0.641185288532478,2.342957589
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BP,"GO:0090100","positive regulation of transmembrane receptor protein serine/threonine kinase s
MF,"GO:0042162","telomeric DNA binding",36,-0.620328723151549,-2.37854221268155,6.3421921
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BP,"GO:1903076","regulation of protein localization to plasma membrane",99,0.477760196134803,
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BP,"GO:0042088","T-helper 1 type immune response",45,0.606085795729576,2.32294480667372,6
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BP,"GO:0010712","regulation of collagen metabolic process",35,0.666702364565148,2.413512413
BP,"GO:0034405","response to fluid shear stress",32,0.647599084507671,2.32480339197086,6.789
BP,"GO:0014902","myotube differentiation",105,0.466256018457053,2.07344107997609,6.794284
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BP,"GO:1903556","negative regulation of tumor necrosis factor superfamily cytokine production",59
BP,"GO:0060350","endochondral bone morphogenesis",56,0.556897828563817,2.24118449789477,
BP,"GO:0003151","outflow tract morphogenesis",77,0.524978687247518,2.22464688317937,7.729
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CC,"GO:0098643","banded collagen fibril",12,0.845521781358393,2.35693835861634,8.506973581
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MF,"GO:0043394","proteoglycan binding",35,0.656783216195085,2.37760435456555,1.092871333
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BP,"GO:0060055","angiogenesis involved in wound healing",25,0.700878261792479,2.36193263897
MF,"GO:0005319","lipid transporter activity",154,0.407591247651118,1.92720244748273,1.135045
BP,"GO:0045661","regulation of myoblast differentiation",52,0.569050326882832,2.2492530225788
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BP,"GO:0110020","regulation of actomyosin structure organization",93,0.475608287753608,2.07806
BP,"GO:0001914","regulation of T cell mediated cytotoxicity",37,0.626900525665309,2.3159992157
BP,"GO:0035633","maintenance of blood-brain barrier",35,0.655563209813075,2.37318784023488,
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BP,"GO:0048525","negative regulation of viral process",86,0.486242286785166,2.11216420332187,
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BP,"GO:0071402","cellular response to lipoprotein particle stimulus",32,0.63815087543684,2.29088
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BP,"GO:0051223","regulation of protein transport",470,0.307231222889167,1.63378257587907,1.3
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MF,"GO:0098631","cell adhesion mediator activity",64,0.534824586084565,2.1916285375973,1.794
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BP,"GO:0003180","aortic valve morphogenesis",32,0.633389054015753,2.27379107914041,1.91523
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CC,"GO:0016591","RNA polymerase II, holoenzyme",74,-0.482583700210351,-2.13865950594425,1.
BP,"GO:0023061","signal release",440,0.307050955616668,1.62627071862036,2.002621142998 04e
BP,"GO:0098801","regulation of renal system process",33,0.638242877158262,2.30493995584986,2
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BP,"GO:0046486","glycerolipid metabolic process",370,0.321213349293027,1.66625202017741,2.02
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BP,"GO:0048144","fibroblast proliferation",78,0.489264930188685,2.07450263957127,2.109372373
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BP,"GO:0050779","RNA destabilization",86,-0.445067541755563,-1.99958317442211,2.2083782141
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BP,"GO:1905521","regulation of macrophage migration",39,0.6092789728572,2.27710088041765,2.
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BP,"GO:0000959","mitochondrial RNA metabolic process",45,-0.563455916942612,-2.26638866624
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MF,"GO:0004175","endopeptidase activity",406,0.312926358506458,1.646133957235,2.646611166
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BP,"GO:0030728","ovulation",23,0.708201960823872,2.32220006780839,2.7401364232329e-06,1.6
BP,"GO:0002027","regulation of heart rate",101,0.460127826770501,2.04120792959778,2.7480132
BP,"GO:0006941","striated muscle contraction",164,0.400953775315358,1.91367675924884,2.7808
CC,"GO:0005844","polysome",64,-0.504639594208194,-2.16624531227811,2.83334479071536e-06
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BP,"GO:0002293","alpha-beta T cell differentiation involved in immune response",67,0.5167480465C
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BP,"GO:0002821","positive regulation of adaptive immune response",104,0.457371633961433,2.032
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BP,"GO:0045685","regulation of glial cell differentiation",72,0.511256029272142,2.13004087235002
BP,"GO:0003197","endocardial cushion development",47,0.577532537517517,2.23890291401654,3
BP,"GO:0006413","translational initiation",118,-0.40720700246065,-1.96061773092987,3.0526220
BP,"GO:0055088","lipid homeostasis",147,0.407653033301633,1.90921301299049,3.127433100689
CC,"GO:0016363","nuclear matrix",108,-0.420626632338682,-1.98296984653695,3.2283095959944
CC,"GO:0031256","leading edge membrane",169,0.39239806475267,1.87842564034753,3.2525002
BP,"GO:0003416","endochondral bone growth",28,0.671544448845239,2.2994992722021,3.373540
CC,"GO:0070461","SAGA-type complex",32,-0.620114705201839,-2.29263296425306,3.3834190146
BP,"GO:0000380","alternative mRNA splicing, via spliceosome",77,-0.477106012108256,-2.1112384
MF,"GO:0000993","RNA polymerase II complex binding",34,-0.60766209531893,-2.28951365081969
BP,"GO:0090077","foam cell differentiation",37,0.61312974470057,2.26512492772186,3.44922968
CC,"GO:0071007","U2-type catalytic step 2 spliceosome",30,-0.645647400777238,-2.355481201699
BP,"GO:0060411","cardiac septum morphogenesis",67,0.51497524331006,2.12243417470718,3.489
BP,"GO:0055001","muscle cell development",168,0.395731285745663,1.89034866784533,3.496264
BP,"GO:0050714","positive regulation of protein secretion",126,0.43069796512236,1.96414777104
BP,"GO:0098754","detoxification",126,0.430347894293634,1.96255131391252,3.5035800442685e-
CC,"GO:0005689","U12-type spliceosomal complex",28,-0.654667646753672,-2.33591481517394,3.
BP,"GO:0006942","regulation of striated muscle contraction",87,0.475937603462868,2.0620390821
BP,"GO:0042303","molting cycle",113,0.443681564615859,1.99071162967285,3.58749939052187e
BP,"GO:0042633","hair cycle",113,0.443681564615859,1.99071162967285,3.58749939052187e-06,
CC,"GO:0042788","polysomal ribosome",31,-0.620144668643742,-2.30224925514839,3.613019540
CC,"GO:0005903","brush border",101,0.456503687584087,2.02513061104655,3.6167135549398e-0
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BP,"GO:0051988","regulation of attachment of spindle microtubules to kinetochore",13,-0.8012811
BP,"GO:0110151","positive regulation of biomineralization",49,0.562809856330364,2.19797239977
BP,"GO:0060993","kidney morphogenesis",93,0.463995680120528,2.02732527190463,3.809514982
BP,"GO:0007431","salivary gland development",36,0.618391213097306,2.25966567177665,3.82895
BP,"GO:0034599","cellular response to oxidative stress",262,0.344630476091356,1.7325468905969
BP,"GO:0001707","mesoderm formation",68,0.502399694833468,2.07571261936209,3.867429906
BP,"GO:0048333","mesodermal cell differentiation",30,0.660664046598681,2.31709600112172,3.92
BP,"GO:0001974","blood vessel remodeling",41,0.599507295108622,2.2650478556841,3.97022941
CC,"GO:0005682","U5 snRNP",17,-0.741009063382612,-2.30018430131717,4.05691932273438e-06
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BP,"GO:0002832","negative regulation of response to biotic stimulus",104,0.453257390845922,2.01
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BP,"GO:2000516","positive regulation of CD4-positive, alpha-beta T cell activation",37,0.609598680
BP,"GO:0009636","response to toxic substance",221,0.359405462772929,1.7696652510769,4.30359
BP,"GO:0099024","plasma membrane invagination",63,0.527217725559939,2.14443229292268,4.3
MF,"GO:0008170","N-methyltransferase activity",78,-0.457982428549332,-2.0408586719176,4.336
BP,"GO:0008037","cell recognition",154,0.396396950362996,1.87427276055779,4.37062681366589
BP,"GO:0006638","neutral lipid metabolic process",123,0.425832248612273,1.93533217579247,4.4
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BP,"GO:1902905","positive regulation of supramolecular fiber organization",168,0.39368565951988
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BP,"GO:0010565","regulation of cellular ketone metabolic process",119,0.419931295426855,1.8981
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BP,"GO:0043631","RNA polyadenylation",42,-0.565502698561746,-2.23450163443849,4.799422792
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BP,"GO:0050708","regulation of protein secretion",239,0.355856095212804,1.7697914761901,4.83
CC,"GO:0034709","methylosome",12,-0.828192604035027,-2.31379552341972,4.91312983808209e
CC,"GO:0005901","caveola",75,0.488587700555113,2.05533069486441,4.9306026880709e-06,2.79
BP,"GO:0055092","sterol homeostasis",81,0.471567266409403,2.00464555279713,4.938779475281
BP,"GO:1903169","regulation of calcium ion transmembrane transport",141,0.414145746959583,1.9
BP,"GO:0072171","mesonephric tubule morphogenesis",64,0.522254207031831,2.14011706602851
BP,"GO:0008347","glial cell migration",47,0.570555147823423,2.21185387850161,4.999650510334
BP,"GO:0045920","negative regulation of exocytosis",31,0.635095747171972,2.2335844940795,5.0
BP,"GO:1901617","organic hydroxy compound biosynthetic process",233,0.361402735279168,1.789
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BP,"GO:0034135","regulation of toll-like receptor 2 signaling pathway",11,0.837929565372418,2.29
BP,"GO:0051493","regulation of cytoskeleton organization",495,0.301356306271299,1.611891531
BP,"GO:0032330","regulation of chondrocyte differentiation",45,0.580143187280808,2.223514580
CC,"GO:0035267","NuA4 histone acetyltransferase complex",25,-0.676430334185413,-2.317765608
CC,"GO:0043189","H4/H2A histone acetyltransferase complex",25,-0.676430334185413,-2.3177656
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MF,"GO:0005516","calmodulin binding",199,0.370173726602379,1.81559138047064,5.2253745171
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BP,"GO:1902750","negative regulation of cell cycle G2/M phase transition",55,-0.512830917915176,
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BP,"GO:0002825","regulation of T-helper 1 type immune response",29,0.648807433088238,2.24745
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BP,"GO:0007188","adenylate cyclase-modulating G protein-coupled receptor signaling pathway",220
BP,"GO:0060323","head morphogenesis",34,0.622697587957703,2.25724440645678,6.5457683441
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BP,"GO:0000184","nuclear-transcribed mRNA catabolic process, nonsense-mediated decay",39,-0.57
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BP,"GO:0104004","cellular response to environmental stimulus",309,0.327623314823043,1.679648
CC,"GO:0005802","trans-Golgi network",237,0.348774500381309,1.72802856679636,7.1247974812
BP,"GO:0010883","regulation of lipid storage",51,0.546443503052724,2.15142529919538,7.135657
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BP,"GO:0034123","positive regulation of toll-like receptor signaling pathway",27,0.66687360981651
BP,"GO:0043627","response to estrogen",68,0.492733985102998,2.03577781074466,7.6561698092
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BP,"GO:0032354","response to follicle-stimulating hormone",18,0.7392042488641,2.284484562940
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BP,"GO:0032615","interleukin-12 production",62,0.521463652771807,2.12610667725405,7.939768
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BP,"GO:0042093","T-helper cell differentiation",64,0.516374728465306,2.11602387108599,8.05022
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BP,"GO:0006312","mitotic recombination",25,-0.666254726834368,-2.28289923447168,8.10915497
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BP,"GO:0010972","negative regulation of G2/M transition of mitotic cell cycle",54,-0.514884573936
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BP,"GO:0072088","nephron epithelium morphogenesis",74,0.489902510629724,2.05396370049663
BP,"GO:0042475","odontogenesis of dentin-containing tooth",88,0.475108629472397,2.069016640
BP,"GO:1902547","regulation of cellular response to vascular endothelial growth factor stimulus",21
BP,"GO:0060395","SMAD protein signal transduction",81,0.463951577141554,1.97227104610475,9
BP,"GO:0001990","regulation of systemic arterial blood pressure by hormone",38,0.6003938552799
BP,"GO:0044818","mitotic G2/M transition checkpoint",42,-0.55163167070809,-2.17969228606727,
CC,"GO:0005635","nuclear envelope",450,-0.266511016490642,-1.53240005258987,9.6654710057
BP,"GO:0010508","positive regulation of autophagy",132,0.402854482258695,1.8492044898504,9.6
BP,"GO:0003094","glomerular filtration",20,0.72161620648943,2.29685258647549,9.74521908941
BP,"GO:0034394","protein localization to cell surface",65,0.504503724233336,2.08188403676205,9
BP,"GO:0050931","pigment cell differentiation",34,0.618029140517051,2.24032154201015,9.78740
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BP,"GO:0036336","dendritic cell migration",33,0.614742250627065,2.22007017505349,1.00367576
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BP,"GO:2000403","positive regulation of lymphocyte migration",36,0.602127654598601,2.20023694
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BP,"GO:0006968","cellular defense response",51,0.540912542810998,2.12964912704507,1.037362
BP,"GO:0048715","negative regulation of oligodendrocyte differentiation",12,0.808899762829313,2
BP,"GO:1901136","carbohydrate derivative catabolic process",159,0.378257186249475,1.79640907
BP,"GO:0097242","amyloid-beta clearance",37,0.597116791646519,2.20596723810939,1.04941061
BP,"GO:0045429","positive regulation of nitric oxide biosynthetic process",38,0.599384948201903,2

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BP,"GO:1904407","positive regulation of nitric oxide metabolic process",40,0.582838832158132,2.1
BP,"GO:0002824","positive regulation of adaptive immune response based on somatic recombinatio
BP,"GO:0070169","positive regulation of biomineral tissue development",48,0.558959743627707,2.
BP,"GO:0006641","triglyceride metabolic process",95,0.45634554887353,1.99811381475016,1.1055
BP,"GO:1905523","positive regulation of macrophage migration",25,0.665155713277963,2.2415490
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MF,"GO:1901981","phosphatidylinositol phosphate binding",164,0.387853539120454,1.851151802
BP,"GO:0030879","mammary gland development",131,0.395916574165699,1.81683492400524,1.14
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BP,"GO:0006644","phospholipid metabolic process",353,0.318738458690583,1.64475221738649,1.
BP,"GO:0002691","regulation of cellular extravasation",36,0.600051865370668,2.19265179185249,
BP,"GO:0032210","regulation of telomere maintenance via telomerase",51,-0.508049597980358,-2.
BP,"GO:0098868","bone growth",31,0.621851889444828,2.18700683174612,1.17053594027906e-0
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BP,"GO:0043114","regulation of vascular permeability",47,0.555350850100886,2.15291183755119,
BP,"GO:0014911","positive regulation of smooth muscle cell migration",32,0.599946252028683,2.15
BP,"GO:0001658","branching involved in ureteric bud morphogenesis",57,0.532570087337684,2.138
BP,"GO:0051382","kinetochore assembly",16,-0.734825913171372,-2.28715738362069,1.24440341
BP,"GO:0051496","positive regulation of stress fiber assembly",49,0.548540643614079,2.14224605
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BP,"GO:0060021","roof of mouth development",91,0.456692923810212,1.99285354363631,1.2528
BP,"GO:0003081","regulation of systemic arterial blood pressure by renin-angiotensin",24,0.674597
BP,"GO:0006631","fatty acid metabolic process",369,0.316738569460367,1.64234285701423,1.263
BP,"GO:0044060","regulation of endocrine process",35,0.622374913371308,2.25304372541159,1.2
BP,"GO:0014855","striated muscle cell proliferation",59,0.523140918233294,2.11391765956659,1.2
MF,"GO:0005520","insulin-like growth factor binding",23,0.684738960234359,2.24526469545138,1.
BP,"GO:0061157","mRNA destabilization",82,-0.437716216410688,-1.956507071518,1.2896729595
BP,"GO:0061014","positive regulation of mRNA catabolic process",86,-0.425140308772557,-1.91005
BP,"GO:0031445","regulation of heterochromatin assembly",25,-0.658153912697848,-2.255142069
BP,"GO:0120261","regulation of heterochromatin organization",25,-0.658153912697848,-2.255142
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BP,"GO:0120263","positive regulation of heterochromatin organization",13,-0.780577410412614,-2.
BP,"GO:1905269","positive regulation of chromatin organization",13,-0.780577410412614,-2.23700
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BP,"GO:0071383","cellular response to steroid hormone stimulus",191,0.365139612882449,1.77772
BP,"GO:0098789","pre-mRNA cleavage required for polyadenylation",13,-0.779720728850965,-2.23
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BP,"GO:0031058","positive regulation of histone modification",101,-0.407792711364672,-1.897101
CC,"GO:0030686","90S preribosome",25,-0.654647766936011,-2.24312837987316,1.475078347107
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BP,"GO:0009620","response to fungus",61,0.507922597737317,2.07104146800456,1.52081282767
BP,"GO:0006287","base-excision repair, gap-filling",13,-0.777888962451544,-2.22929619989392,1.5
MF,"GO:0004252","serine-type endopeptidase activity",164,0.385290927341667,1.8389209395104
BP,"GO:0045639","positive regulation of myeloid cell differentiation",95,0.45277951286515,1.9824
BP,"GO:0010171","body morphogenesis",45,0.565989419585244,2.16926744055298,1.5573945566
CC,"GO:0005685","U1 snRNP",18,-0.716560998379404,-2.27070589082343,1.56978015286094e-05
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BP,"GO:0051169","nuclear transport",300,-0.295741302881188,-1.62058001457858,1.5787814847
BP,"GO:0045824","negative regulation of innate immune response",69,0.487123830368203,2.01931
BP,"GO:0090330","regulation of platelet aggregation",28,0.648836303185429,2.22174214903984,1.
BP,"GO:0003071","renal system process involved in regulation of systemic arterial blood pressure",2
BP,"GO:0010720","positive regulation of cell development",287,0.333980208125416,1.689605768
BP,"GO:0001890","placenta development",145,0.396894810831485,1.85710543445209,1.6148941
MF,"GO:0030021","extracellular matrix structural constituent conferring compression resistance",22
BP,"GO:0060986","endocrine hormone secretion",47,0.551887603778509,2.13948597531932,1.632
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BP,"GO:0002711","positive regulation of T cell mediated immunity",55,0.516349073494679,2.07540
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BP,"GO:0032691","negative regulation of interleukin-1 beta production",30,0.639163791487871,2.2
BP,"GO:0050830","defense response to Gram-positive bacterium",87,0.458301548211437,1.985629
BP,"GO:0007202","activation of phospholipase C activity",29,0.631966264862268,2.1891213172529
BP,"GO:0044728","DNA methylation or demethylation",91,-0.425291718458079,-1.9435303102664
BP,"GO:0048706","embryonic skeletal system development",126,0.413961835527066,1.887824606
BP,"GO:0050890","cognition",294,0.323961607067138,1.64802642560443,1.70605443359079e-05,
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BP,"GO:0061314","Notch signaling involved in heart development",11,0.815275226292889,2.23775
BP,"GO:0030834","regulation of actin filament depolymerization",49,0.544701886532939,2.127254
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BP,"GO:0032967","positive regulation of collagen biosynthetic process",23,0.678904681814179,2.22
BP,"GO:0001916","positive regulation of T cell mediated cytotoxicity",27,0.653041320453531,2.224
BP,"GO:0035025","positive regulation of Rho protein signal transduction",31,0.614008327518232,2.
MF,"GO:0019865","immunoglobulin binding",21,0.700670231592759,2.25118484018142,1.816912
BP,"GO:0072577","endothelial cell apoptotic process",54,0.522689995836802,2.08704366218125,1.
BP,"GO:0032465","regulation of cytokinesis",87,-0.427436823855022,-1.93282359478048,1.872226
BP,"GO:0061298","retina vasculature development in camera-type eye",19,0.718291415324668,2.25
BP,"GO:0061564","axon development",465,0.293632068922356,1.5625730647839,1.906748866559
BP,"GO:0002828","regulation of type 2 immune response",32,0.592491037060908,2.126972081374
BP,"GO:0048145","regulation of fibroblast proliferation",76,0.478050517752203,2.01390923131337
BP,"GO:0002532","production of molecular mediator involved in inflammatory response",74,0.4821
BP,"GO:1902895","positive regulation of miRNA transcription",43,0.5592671209989,2.13174519774
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BP,"GO:0000027","ribosomal large subunit assembly",25,-0.64883111286501,-2.22319781189211,1.
BP,"GO:0110110","positive regulation of animal organ morphogenesis",34,0.605706687299354,2.19
BP,"GO:0035313","wound healing, spreading of epidermal cells",19,0.717809385018752,2.2555277

BP,"GO:0002931","response to ischemia",50,0.533738028865523,2.10610057229691,2.0123497673
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BP,"GO:0051402","neuron apoptotic process",231,0.343887531267244,1.70272040077957,2.02907
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CC,"GO:0098553","lumenal side of endoplasmic reticulum membrane",26,0.651467654637317,2.19
BP,"GO:0035929","steroid hormone secretion",21,0.697859428159226,2.24215400400059,2.05219
BP,"GO:0010464","regulation of mesenchymal cell proliferation",29,0.627834969603761,2.1748105
BP,"GO:0002823","negative regulation of adaptive immune response based on somatic recombinatic
BP,"GO:0031400","negative regulation of protein modification process",473,0.294039337548515,1.5
BP,"GO:2000105","positive regulation of DNA-templated DNA replication",14,-0.767508067562503,-
CC,"GO:1990023","mitotic spindle midzone",14,-0.766448462062598,-2.25820556628231,2.193622
BP,"GO:0042554","superoxide anion generation",40,0.574614173997743,2.15906000424651,2.2070
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BP,"GO:0072078","nephron tubule morphogenesis",72,0.48274562098039,2.01125824393726,2.235
BP,"GO:0072010","glomerular epithelium development",26,0.649641072615197,2.18581732698654
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BP,"GO:0043550","regulation of lipid kinase activity",66,0.474896864236591,1.95429853281359,2.2
BP,"GO:0090051","negative regulation of cell migration involved in sprouting angiogenesis",15,0.770
CC,"GO:0005680","anaphase-promoting complex",21,-0.680130215825845,-2.24239378831129,2.29
BP,"GO:0035909","aorta morphogenesis",32,0.587608185478577,2.10944322719198,2.3013841169
BP,"GO:0048251","elastic fiber assembly",11,0.810778911259511,2.22541607514957,2.3050938193
BP,"GO:0032206","positive regulation of telomere maintenance",51,-0.49552057842443,-2.0469758
BP,"GO:0030833","regulation of actin filament polymerization",128,0.403486090508526,1.8398361
MF,"GO:0016888","endodeoxyribonuclease activity, producing 5'-phosphomonoesters",15,-0.75314
BP,"GO:0010721","negative regulation of cell development",170,0.376036311517666,1.7999623676
BP,"GO:0018023","peptidyl-lysine trimethylation",46,-0.518976401299065,-2.10008297843653,2.36
BP,"GO:0030042","actin filament depolymerization",54,0.517706303093419,2.06714432521827,2.3
BP,"GO:0072109","glomerular mesangium development",15,0.769908799908988,2.2654018304557
BP,"GO:0061337","cardiac conduction",93,0.441772547641387,1.93022626856015,2.39533343696
BP,"GO:0090382","phagosome maturation",26,0.648567360834087,2.18220465852349,2.39922769
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MF,"GO:0015485","cholesterol binding",50,0.530877346742847,2.09481247976857,2.41719687070
MF,"GO:0019003","GDP binding",71,0.483422034636139,2.01303162357586,2.42764821965041e-0
BP,"GO:0010522","regulation of calcium ion transport into cytosol",96,0.439665458241754,1.92468
BP,"GO:0000466","maturation of 5.8S rRNA from tricistronic rRNA transcript (SSU-rRNA, 5.8S rRNA, I
BP,"GO:2001022","positive regulation of response to DNA damage stimulus",114,-0.3860636493428
BP,"GO:0002468","dendritic cell antigen processing and presentation",14,0.773349206713845,2.255
CC,"GO:0030863","cortical cytoskeleton",105,0.427517867893079,1.90117247740138,2.498237105
BP,"GO:0042509","regulation of tyrosine phosphorylation of STAT protein",76,0.474835210188263,2
BP,"GO:0032535","regulation of cellular component size",334,0.317938170840956,1.635247014242
BP,"GO:0061614","miRNA transcription",58,0.515651619623496,2.07030866818388,2.5172347050
BP,"GO:0072028","nephron morphogenesis",77,0.485914775757001,2.05910986033791,2.5349758
CC,"GO:0031304","intrinsic component of mitochondrial inner membrane",51,-0.49295090292815,-
BP,"GO:0007263","nitric oxide mediated signal transduction",25,0.649964785310422,2.1903561920
BP,"GO:0002686","negative regulation of leukocyte migration",45,0.555921066352115,2.13067846

CC,"GO:0030687","preribosome, large subunit precursor",23,-0.652448298343063,-2.191997213544
CC,"GO:0055037","recycling endosome",184,0.366879915163426,1.77400502536357,2.558499025
MF,"GO:0004713","protein tyrosine kinase activity",135,0.394706350613435,1.81897861280093,2.5
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BP,"GO:0060412","ventricular septum morphogenesis",39,0.576112563841786,2.15314574240406,
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BP,"GO:0071354","cellular response to interleukin-6",32,0.585499479458769,2.10187322435413,2.
BP,"GO:0042886","amide transport",278,0.323627346974508,1.63444156691449,2.6767993190921
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BP,"GO:0000470","maturation of LSU-rRNA",27,-0.612395430610328,-2.15796683248716,2.705149
BP,"GO:0009266","response to temperature stimulus",170,0.372709524823597,1.78403813193637,
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MF,"GO:0051539","4 iron, 4 sulfur cluster binding",40,-0.539345411470382,-2.10741523480577,2.7
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BP,"GO:0070202","regulation of establishment of protein localization to chromosome",12,-0.800953
BP,"GO:0031638","zymogen activation",62,0.505676202994614,2.06173823621374,2.83277697835
BP,"GO:0060343","trabecula formation",24,0.663882585590394,2.20213095624945,2.8362070249
BP,"GO:0009409","response to cold",48,0.542190015304263,2.10730985712526,2.85386173670163
BP,"GO:0030574","collagen catabolic process",40,0.569677765968744,2.14051190428248,2.899402
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BP,"GO:0045910","negative regulation of DNA recombination",29,-0.599200202836956,-2.16227346
BP,"GO:0006303","double-strand break repair via nonhomologous end joining",55,-0.490251551557
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BP,"GO:0032981","mitochondrial respiratory chain complex I assembly",49,-0.51216557023027,-2.1
BP,"GO:2001237","negative regulation of extrinsic apoptotic signaling pathway",94,0.439154858621
MF,"GO:0048407","platelet-derived growth factor binding",11,0.807795400338014,2.21722697073
BP,"GO:0140448","signaling receptor ligand precursor processing",35,0.604331196713691,2.187724
BP,"GO:0051004","regulation of lipoprotein lipase activity",18,0.710915833386689,2.19706021633
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BP,"GO:0043038","amino acid activation",26,-0.636608126424975,-2.22713653284662,3.068789070
CC,"GO:0070469","respirasome",84,-0.428353434090532,-1.92319413134177,3.14453565344222e-
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BP,"GO:1901201","regulation of extracellular matrix assembly",15,0.763239621533871,2.24577824
BP,"GO:1902930","regulation of alcohol biosynthetic process",47,0.538557203094477,2.087808414
CC,"GO:0005697","telomerase holoenzyme complex",21,-0.673477888489161,-2.2204610213933,3.
CC,"GO:0005747","mitochondrial respiratory chain complex I",41,-0.541052752829422,-2.11733756
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BP,"GO:0090685","RNA localization to nucleus",19,-0.702217998468059,-2.24891055790758,3.4185
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BP,"GO:0045580","regulation of T cell differentiation",146,0.383292734299666,1.79415050947606,
BP,"GO:0046636","negative regulation of alpha-beta T cell activation",41,0.569663791340955,2.152
BP,"GO:0050918","positive chemotaxis",58,0.510343759943035,2.04899794697644,3.47615626787
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BP,"GO:0097006","regulation of plasma lipoprotein particle levels",70,0.476931132301173,1.97987
BP,"GO:0035930","corticosteroid hormone secretion",16,0.729436534118962,2.18652792378393,3.
MF,"GO:0043178","alcohol binding",89,0.443684337145779,1.93377132228037,3.53980786827197
BP,"GO:0061082","myeloid leukocyte cytokine production",29,0.617429121150572,2.13876487742
BP,"GO:0045071","negative regulation of viral genome replication",53,0.517426160070847,2.05693
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BP,"GO:0007520","myoblast fusion",40,0.565374994809568,2.12434463668348,3.74194516742324
BP,"GO:0032570","response to progesterone",37,0.577840607929873,2.13475398410351,3.776749
BP,"GO:0061097","regulation of protein tyrosine kinase activity",86,0.444629756485217,1.9314055
MF,"GO:0005507","copper ion binding",57,0.518628864376567,2.08239890838173,3.82690555085
BP,"GO:0051315","attachment of mitotic spindle microtubules to kinetochore",15,-0.745289150407
BP,"GO:1904874","positive regulation of telomerase RNA localization to Cajal body",15,-0.74521400
BP,"GO:0045777","positive regulation of blood pressure",35,0.600354007056947,2.1733263979076
BP,"GO:0090308","regulation of DNA methylation-dependent heterochromatin assembly",16,-0.713
BP,"GO:1900026","positive regulation of substrate adhesion-dependent cell spreading",41,0.567880
BP,"GO:0050767","regulation of neurogenesis",347,0.315470242400664,1.62594541604353,3.9017
BP,"GO:0002820","negative regulation of adaptive immune response",59,0.506486761904841,2.046
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BP,"GO:0061037","negative regulation of cartilage development",25,0.641903386182349,2.1631895
BP,"GO:0017004","cytochrome complex assembly",34,-0.567012471433281,-2.13635637886902,3.9
BP,"GO:0055117","regulation of cardiac muscle contraction",70,0.475201265739297,1.9726894158
BP,"GO:0098543","detection of other organism",20,0.695055536195496,2.21231187949784,3.9895
BP,"GO:0001942","hair follicle development",89,0.442377538327246,1.92807571874493,3.9965512
BP,"GO:0045732","positive regulation of protein catabolic process",223,0.3369074443523,1.660863
CC,"GO:0098803","respiratory chain complex",76,-0.439575465927075,-1.94917476612148,4.05886
BP,"GO:0032784","regulation of DNA-templated transcription, elongation",55,-0.485629185489977,
BP,"GO:0002715","regulation of natural killer cell mediated immunity",49,0.532844757194504,2.080
BP,"GO:0014743","regulation of muscle hypertrophy",54,0.508222907339592,2.02927816906142,4.
BP,"GO:0009952","anterior/posterior pattern specification",211,0.35018024196588,1.71705492185
BP,"GO:0150077","regulation of neuroinflammatory response",33,0.590466054982739,2.13239951
BP,"GO:0043981","histone H4-K5 acetylation",16,-0.710889495990142,-2.21265490308458,4.21330
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CC,"GO:0042827","platelet dense granule",20,0.692858410812954,2.20531858711843,4.306565512
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BP,"GO:0046850","regulation of bone remodeling",49,0.531546406595736,2.07587752374572,4.39
BP,"GO:0045662","negative regulation of myoblast differentiation",25,0.639107031170675,2.15376
BP,"GO:0070201","regulation of establishment of protein localization",494,0.280488059404183,1.50
CC,"GO:0031527","filopodium membrane",18,0.701489632334051,2.16792887566968,4.579425525
BP,"GO:0014014","negative regulation of gliogenesis",37,0.574653409039533,2.12297930881085,4.
BP,"GO:0016486","peptide hormone processing",33,0.588768563254706,2.12626922251978,4.6588
BP,"GO:0046887","positive regulation of hormone secretion",119,0.394727047158446,1.784202522
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BP,"GO:0048713","regulation of oligodendrocyte differentiation",40,0.561494259663652,2.1097631
BP,"GO:0009303","rRNA transcription",28,-0.609534951792797,-2.17487717824379,4.81209450321
BP,"GO:0044272","sulfur compound biosynthetic process",141,0.390488802869038,1.81351176838
CC,"GO:0032592","integral component of mitochondrial membrane",81,-0.414151656283022,-1.860
MF,"GO:0005161","platelet-derived growth factor receptor binding",13,0.76984736491694,2.20732
BP,"GO:0033673","negative regulation of kinase activity",213,0.345874007344416,1.697251824688
BP,"GO:0007611","learning or memory",256,0.327894801551646,1.64820299616153,5.0075642678
BP,"GO:1903306","negative regulation of regulated secretory pathway",23,0.661882354596222,2.17
BP,"GO:0001892","embryonic placenta development",86,0.441608399951315,1.91828123471196,5
CC,"GO:0098862","cluster of actin-based cell projections",152,0.373641271552569,1.757142597271
BP,"GO:0000154","rRNA modification",30,-0.59782812897544,-2.18102468615795,5.141142186470
BP,"GO:0022404","molting cycle process",92,0.441613404890479,1.92719722458772,5.255486201
BP,"GO:0022405","hair cycle process",92,0.441613404890479,1.92719722458772,5.2554862010989
CC,"GO:0000177","cytoplasmic exosome (RNase complex)",13,-0.755920028143482,-2.16633700631
BP,"GO:0002702","positive regulation of production of molecular mediator of immune response",11
BP,"GO:0071241","cellular response to inorganic substance",210,0.344781653106646,1.692571059
BP,"GO:0048645","animal organ formation",64,0.490041693189137,2.00811516027799,5.33084942
BP,"GO:0032620","interleukin-17 production",35,0.59514918382304,2.15448454860254,5.3383923
BP,"GO:0032660","regulation of interleukin-17 production",35,0.59514918382304,2.154484548602
BP,"GO:0045109","intermediate filament organization",68,0.464361064940068,1.91855236447948,
BP,"GO:0043254","regulation of protein-containing complex assembly",373,0.302654995660922,1.5
MF,"GO:0000175","3'-5'-exoribonuclease activity",34,-0.560052962718391,-2.1101347495647,5.409
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BP,"GO:0030888","regulation of B cell proliferation",63,0.487263352784705,1.98191983731113,5.4
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BP,"GO:0061318","renal filtration cell differentiation",22,0.655574460045019,2.14457807809735,5.
BP,"GO:0072112","glomerular visceral epithelial cell differentiation",22,0.655574460045019,2.1445
BP,"GO:0038066","p38MAPK cascade",52,0.511223585735196,2.02068453545633,5.531669100148
BP,"GO:0070203","regulation of establishment of protein localization to telomere",11,-0.801936496
BP,"GO:0006626","protein targeting to mitochondrion",97,-0.398159783178005,-1.8467724268114
BP,"GO:0070168","negative regulation of biomineral tissue development",26,0.632445807871801,2

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BP,"GO:0055067","monovalent inorganic cation homeostasis",155,0.373996388936383,1.77025624
BP,"GO:0002577","regulation of antigen processing and presentation",20,0.688514392276481,2.191
BP,"GO:0034308","primary alcohol metabolic process",104,0.424793677269216,1.88743269548757
CC,"GO:1904724","tertiary granule lumen",55,0.499613382414509,2.00813417535042,5.637195624
MF,"GO:0008327","methyl-CpG binding",25,-0.628888222332869,-2.15486417357738,5.646757947
BP,"GO:1904872","regulation of telomerase RNA localization to Cajal body",18,-0.689840130470072
MF,"GO:0032934","sterol binding",60,0.501416595348403,2.04113207350274,5.69353126371246e
BP,"GO:0014897","striated muscle hypertrophy",78,0.444582286886697,1.8850464661291,5.71110
CC,"GO:0042613","MHC class II protein complex",15,0.751170331166502,2.21026522025695,5.7555
BP,"GO:0060760","positive regulation of response to cytokine stimulus",61,0.490202497173412,1.9
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BP,"GO:0032715","negative regulation of interleukin-6 production",46,0.527930274576868,2.03852
MF,"GO:0043175","RNA polymerase core enzyme binding",39,-0.542879482874487,-2.0848603045
CC,"GO:0000307","cyclin-dependent protein kinase holoenzyme complex",46,-0.50445789587012,-2
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BP,"GO:0006469","negative regulation of protein kinase activity",194,0.353756005024092,1.725279
BP,"GO:0016049","cell growth",457,0.286122908931317,1.5177829226098,5.95531106412999e-05
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BP,"GO:0050853","B cell receptor signaling pathway",56,0.492911065769994,1.98367560939036,6.
BP,"GO:0043281","regulation of cysteine-type endopeptidase activity involved in apoptotic process"
MF,"GO:0044548","S100 protein binding",14,0.758078079453687,2.21063422353229,6.131857962
BP,"GO:2000112","regulation of cellular macromolecule biosynthetic process",446,-0.262955513844
MF,"GO:0048020","CCR chemokine receptor binding",40,0.557610910494418,2.09517180267938,6.
MF,"GO:0008017","microtubule binding",257,-0.29928689944308,-1.61902986565307,6.304878420
BP,"GO:0070646","protein modification by small protein removal",155,-0.341264605749506,-1.7215
BP,"GO:0043300","regulation of leukocyte degranulation",47,0.530431463186833,2.056307604277
BP,"GO:0046605","regulation of centrosome cycle",50,-0.493630591317444,-2.02180277360092,6.4
BP,"GO:0007548","sex differentiation",274,0.324218117234646,1.63615982198027,6.46671827521
BP,"GO:0048872","homeostasis of number of cells",271,0.322297667409402,1.62782466285767,6.4
BP,"GO:0001991","regulation of systemic arterial blood pressure by circulatory renin-angiotensin",1
BP,"GO:0006997","nucleus organization",128,-0.361015162639862,-1.76291462533387,6.536815400
CC,"GO:0043034","costamere",16,0.713549713360004,2.13890626571662,6.63189893443733e-05,
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BP,"GO:0051767","nitric-oxide synthase biosynthetic process",19,0.693512376077666,2.179180777
BP,"GO:0051769","regulation of nitric-oxide synthase biosynthetic process",19,0.693512376077666,
BP,"GO:0038127","ERBB signaling pathway",114,0.405859914697145,1.82333389279765,6.7400583
CC,"GO:0010369","chromocenter",15,-0.736238488693447,-2.22269511561682,6.7538588626889e
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BP,"GO:0035521","monoubiquitinated histone deubiquitination",28,-0.602019567925127,-2.148061
BP,"GO:0035522","monoubiquitinated histone H2A deubiquitination",28,-0.602019567925127,-2.14
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BP,"GO:0120192","tight junction assembly",68,0.461955823363866,1.90861487776585,6.88042816
BP,"GO:0072698","protein localization to microtubule cytoskeleton",53,-0.482680007476595,-2.001
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BP,"GO:0021782","glial cell development",108,0.412097467251881,1.83322749936308,6.930976503
BP,"GO:0036303","lymph vessel morphogenesis",22,0.649582762113622,2.12497746090221,6.9467
BP,"GO:0051279","regulation of release of sequestered calcium ion into cytosol",76,0.461107710953
BP,"GO:0030835","negative regulation of actin filament depolymerization",40,0.556364447598093,2
BP,"GO:0031668","cellular response to extracellular stimulus",230,0.331580605175196,1.636130698

BP,"GO:0007584","response to nutrient",148,0.379119840546278,1.77809816198461,7.044926337
MF,"GO:0016278","lysine N-methyltransferase activity",44,-0.507147987256469,-2.0175643365615
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BP,"GO:0042130","negative regulation of T cell proliferation",64,0.485393854713446,1.9890690361
BP,"GO:0070613","regulation of protein processing",62,0.491747221261753,2.00494712352128,7.1
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BP,"GO:0046165","alcohol biosynthetic process",135,0.382725708468786,1.76376660064336,7.404
BP,"GO:0002675","positive regulation of acute inflammatory response",26,0.626299469458354,2.10
BP,"GO:0071539","protein localization to centrosome",32,-0.565830683546773,-2.09193890485622
MF,"GO:0016896","exoribonuclease activity, producing 5'-phosphomonoesters",37,-0.54347487706
CC,"GO:0030894","replisome",15,-0.733767391719523,-2.21523490366307,7.47106499016006e-05
BP,"GO:0045628","regulation of T-helper 2 cell differentiation",11,0.790614999257491,2.17007041
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BP,"GO:0061517","macrophage proliferation",11,0.790551662811137,2.1698965701058,7.6470274
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BP,"GO:1904851","positive regulation of establishment of protein localization to telomere",10,-0.81
BP,"GO:2001212","regulation of vasculogenesis",16,0.71092813231405,2.1310479258977,7.710234
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BP,"GO:0043407","negative regulation of MAP kinase activity",57,0.509302842626425,2.044953060
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CC,"GO:0019898","extrinsic component of membrane",300,0.318428833955606,1.62168506535561
CC,"GO:0005847","mRNA cleavage and polyadenylation specificity factor complex",15,-0.728962204
BP,"GO:1903312","negative regulation of mRNA metabolic process",85,-0.41556056277167,-1.8757
BP,"GO:0071901","negative regulation of protein serine/threonine kinase activity",111,0.418954989
CC,"GO:0005721","pericentric heterochromatin",21,-0.654134756102597,-2.15668658687939,8.307
BP,"GO:0051204","protein insertion into mitochondrial membrane",23,-0.627139169395009,-2.106
MF,"GO:0016410","N-acyltransferase activity",78,-0.423213888769904,-1.88592330432387,8.4202
BP,"GO:0034143","regulation of toll-like receptor 4 signaling pathway",22,0.646400706606758,2.11
BP,"GO:0045668","negative regulation of osteoblast differentiation",44,0.530449412710835,2.0305
CC,"GO:0034358","plasma lipoprotein particle",35,0.587196126458863,2.12569388624255,8.55171
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CC,"GO:0090734","site of DNA damage",83,-0.415583190371125,-1.85991283476113,8.7378873121
BP,"GO:0045124","regulation of bone resorption",41,0.551448614394296,2.08347339848042,8.753
BP,"GO:1903670","regulation of sprouting angiogenesis",35,0.586044094921247,2.12152344593152
BP,"GO:0098787","mRNA cleavage involved in mRNA processing",14,-0.742288869041612,-2.18702
BP,"GO:0006970","response to osmotic stress",79,0.451039919173442,1.91824146811991,8.89782
BP,"GO:0010934","macrophage cytokine production",17,0.72080598504761,2.20020793749608,8.9
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BP,"GO:0002399","MHC class II protein complex assembly",14,0.749017808920748,2.184213535931
BP,"GO:0002503","peptide antigen assembly with MHC class II protein complex",14,0.74901780892
CC,"GO:1905354","exoribonuclease complex",20,-0.66876898444421,-2.18602385073693,9.086551
BP,"GO:0032355","response to estradiol",121,0.40562280452564,1.8346090657435,9.13417249888
BP,"GO:0098883","synapse pruning",11,0.784508150651371,2.15330841157832,9.20762331322495
BP,"GO:0032411","positive regulation of transporter activity",110,0.411667277229545,1.837342704
BP,"GO:0097581","lamellipodium organization",84,0.442957355256775,1.90884324142544,9.27231
BP,"GO:0070252","actin-mediated cell contraction",97,0.419772448022922,1.84743344132628,9.41

BP,"GO:0055078","sodium ion homeostasis",56,0.48548012544391,1.95377046807108,9.44275940
CC,"GO:0031305","integral component of mitochondrial inner membrane",50,-0.487205287613354,
BP,"GO:0010469","regulation of signaling receptor activity",165,0.365500212801245,1.7460887240
BP,"GO:0060260","regulation of transcription initiation from RNA polymerase II promoter",48,-0.494
BP,"GO:0007219","Notch signaling pathway",170,0.358916263038979,1.71801431620699,9.511814
BP,"GO:0070542","response to fatty acid",61,0.482712586882926,1.96824829022338,9.519994532
BP,"GO:0071320","cellular response to cAMP",53,0.502336952034149,1.99695067275,9.701538049
BP,"GO:0055017","cardiac muscle tissue growth",67,0.471967505483204,1.94518081403329,9.741
CC,"GO:0044291","cell-cell contact zone",67,0.471694392512727,1.94405519817189,9.7410274746
MF,"GO:0016279","protein-lysine N-methyltransferase activity",43,-0.518313831078632,-2.0461517
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CC,"GO:0030864","cortical actin cytoskeleton",77,0.466013889309261,1.97477797014139,9.871285
BP,"GO:0043280","positive regulation of cysteine-type endopeptidase activity involved in apoptotic
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MF,"GO:0019843","rRNA binding",61,-0.452241747171995,-1.92018205091366,0.00010107229465
BP,"GO:0031953","negative regulation of protein autophosphorylation",12,0.771890225551358,2.15
BP,"GO:0090050","positive regulation of cell migration involved in sprouting angiogenesis",20,0.676
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BP,"GO:0031109","microtubule polymerization or depolymerization",119,-0.359213121307141,-1.72
CC,"GO:0030134","COPII-coated ER to Golgi transport vesicle",89,0.428918021478014,1.869413229
BP,"GO:0045104","intermediate filament cytoskeleton organization",87,0.431459599967793,1.8693
BP,"GO:0043552","positive regulation of phosphatidylinositol 3-kinase activity",33,0.5704422414262
BP,"GO:0070741","response to interleukin-6",36,0.564772630369551,2.06373780573797,0.0001048
CC,"GO:0000421","autophagosome membrane",48,0.519762716472438,2.0201425051583,0.000105
BP,"GO:0043524","negative regulation of neuron apoptotic process",139,0.379588852859537,1.763
MF,"GO:0008191","metalloendopeptidase inhibitor activity",16,0.702948938540481,2.1071298340
BP,"GO:0010975","regulation of neuron projection development",411,0.293341515560884,1.54577
BP,"GO:0060143","positive regulation of syncytium formation by plasma membrane fusion",22,0.64
BP,"GO:0006925","inflammatory cell apoptotic process",21,0.663537946653126,2.13188244489145
CC,"GO:0097346","INO80-type complex",26,-0.610703406844271,-2.13651037688577,0.000108089
BP,"GO:0045738","negative regulation of DNA repair",28,-0.591577947012479,-2.1108049215644,0
BP,"GO:0070830","bicellular tight junction assembly",63,0.477058767180636,1.94041318485171,0.
BP,"GO:0010463","mesenchymal cell proliferation",41,0.547402420827124,2.06818614153162,0.00
BP,"GO:0072594","establishment of protein localization to organelle",406,-0.259645624890834,-1.4
BP,"GO:0098739","import across plasma membrane",178,0.360233704861483,1.74012928746704,0
BP,"GO:1901361","organic cyclic compound catabolic process",466,-0.25261420825079,-1.4608070
BP,"GO:0010927","cellular component assembly involved in morphogenesis",108,0.407224083623
MF,"GO:0005085","guanyl-nucleotide exchange factor activity",218,0.334845987157672,1.6418958
BP,"GO:0000288","nuclear-transcribed mRNA catabolic process, deadenylation-dependent decay",5
BP,"GO:0031057","negative regulation of histone modification",46,-0.490516493943955,-1.9849174
BP,"GO:0033619","membrane protein proteolysis",57,0.504660538875755,2.02631327996269,0.00
BP,"GO:2000113","negative regulation of cellular macromolecule biosynthetic process",179,-0.3211
BP,"GO:0001911","negative regulation of leukocyte mediated cytotoxicity",25,0.615078886136864,
BP,"GO:0006383","transcription by RNA polymerase III",52,-0.480850673900344,-1.9848869215498
BP,"GO:0098781","ncRNA transcription",51,-0.469093402356311,-1.93780625046395,0.000116990
BP,"GO:0090322","regulation of superoxide metabolic process",29,0.596853837966226,2.06749241
BP,"GO:1900363","regulation of mRNA polyadenylation",17,-0.678184038499238,-2.105167609744
MF,"GO:1902936","phosphatidylinositol bisphosphate binding",100,0.420542913861719,1.8605504
BP,"GO:0007498","mesoderm development",126,0.390973346515303,1.78298828711088,0.000118

BP,"GO:0060142","regulation of syncytium formation by plasma membrane fusion",27,0.622033252
MF,"GO:0005546","phosphatidylinositol-4,5-bisphosphate binding",75,0.447968566141071,1.88445
MF,"GO:0043177","organic acid binding",136,0.382084288730687,1.76597805247272,0.000118815
BP,"GO:0010847","regulation of chromatin assembly",28,-0.589602144043127,-2.10375507352187,
BP,"GO:0003382","epithelial cell morphogenesis",33,0.568918260883553,2.05458216308139,0.000
BP,"GO:0060441","epithelial tube branching involved in lung morphogenesis",29,0.59675484455678
BP,"GO:1900746","regulation of vascular endothelial growth factor signaling pathway",19,0.6844849
BP,"GO:0071372","cellular response to follicle-stimulating hormone stimulus",12,0.76833636029376
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BP,"GO:0002708","positive regulation of lymphocyte mediated immunity",107,0.405807530306179,
BP,"GO:1904754","positive regulation of vascular associated smooth muscle cell migration",11,0.776
BP,"GO:0034614","cellular response to reactive oxygen species",132,0.37490990995277,1.72093179
BP,"GO:1905508","protein localization to microtubule organizing center",34,-0.544345312010929,-2
BP,"GO:0050851","antigen receptor-mediated signaling pathway",163,0.365112661541552,1.73903
BP,"GO:0032886","regulation of microtubule-based process",235,-0.300305445598557,-1.60516321
CC,"GO:0097431","mitotic spindle pole",36,-0.524974336994905,-2.01292246274373,0.000129439
BP,"GO:0060415","muscle tissue morphogenesis",70,0.460135900514382,1.91014899629413,0.000
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BP,"GO:0043551","regulation of phosphatidylinositol 3-kinase activity",54,0.490149288771023,1.95
BP,"GO:0031623","receptor internalization",110,0.406715168089218,1.81524058003346,0.0001308
BP,"GO:0046641","positive regulation of alpha-beta T cell proliferation",23,0.647532862723882,2.12
CC,"GO:0098799","outer mitochondrial membrane protein complex",19,-0.67202795740464,-2.1522
BP,"GO:0045103","intermediate filament-based process",88,0.43725282195012,1.90416108762862
BP,"GO:0071277","cellular response to calcium ion",82,0.434068593066024,1.85608615911782,0.0
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CC,"GO:0000786","nucleosome",34,-0.543316528741292,-2.04707619391051,0.0001351504912861
BP,"GO:0060261","positive regulation of transcription initiation from RNA polymerase II promoter",4
BP,"GO:1902622","regulation of neutrophil migration",42,0.532550502364827,2.01472329817393,0
CC,"GO:1902554","serine/threonine protein kinase complex",95,-0.391175298265041,-1.808177351
BP,"GO:0062125","regulation of mitochondrial gene expression",26,-0.605109437655815,-2.116940
BP,"GO:0015711","organic anion transport",352,0.298838055656933,1.54350859486623,0.0001387
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BP,"GO:2000249","regulation of actin cytoskeleton reorganization",39,0.547548148874644,2.04638
BP,"GO:0050829","defense response to Gram-negative bacterium",75,0.444680032644087,1.87062
BP,"GO:0032928","regulation of superoxide anion generation",22,0.635215991304771,2.077979501
BP,"GO:0000966","RNA 5'-end processing",22,-0.630915544757613,-2.08082239698122,0.0001398
BP,"GO:0016482","cytosolic transport",154,0.36218772981283,1.71252224714346,0.000139829888
BP,"GO:0015718","monocarboxylic acid transport",115,0.391981858067716,1.76579259523219,0.0
BP,"GO:0042100","B cell proliferation",99,0.416547743791576,1.85072614601827,0.000141377114
BP,"GO:0032930","positive regulation of superoxide anion generation",19,0.68157252893365,2.141
BP,"GO:0048730","epidermis morphogenesis",34,0.572149581711317,2.07401067219515,0.000143
MF,"GO:0140104","molecular carrier activity",71,-0.430676171810845,-1.89562733490922,0.00014
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BP,"GO:0071880","adenylate cyclase-activating adrenergic receptor signaling pathway",25,0.611748
MF,"GO:0050135","NAD(P)+ nucleosidase activity",16,0.697813336531164,2.09173557197422,0.00
MF,"GO:0061809","NAD+ nucleotidase, cyclic ADP-ribose generating",16,0.697813336531164,2.091
BP,"GO:0070278","extracellular matrix constituent secretion",13,0.748105814513713,2.144988759

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BP,"GO:1904888","cranial skeletal system development",70,0.459247533646201,1.90646114433626
BP,"GO:0048593","camera-type eye morphogenesis",120,0.382848071774652,1.73189510727733,0
BP,"GO:0043370","regulation of CD4-positive, alpha-beta T cell differentiation",49,0.513084579446
BP,"GO:0045063","T-helper 1 cell differentiation",21,0.657331781447351,2.11194264383105,0.000
MF,"GO:1990782","protein tyrosine kinase binding",108,0.403195454135999,1.79362663660468,0.0
BP,"GO:0030206","chondroitin sulfate biosynthetic process",18,0.678362186404342,2.0964543227
BP,"GO:0061299","retina vasculature morphogenesis in camera-type eye",11,0.773029395022367,2
BP,"GO:0032986","protein-DNA complex disassembly",19,-0.668827999383678,-2.14197635566098
BP,"GO:0044089","positive regulation of cellular component biogenesis",467,0.28437312028254,1.5
BP,"GO:2000780","negative regulation of double-strand break repair",26,-0.600695864850786,-2.10
BP,"GO:0046545","development of primary female sexual characteristics",109,0.398706100284343,
BP,"GO:1904357","negative regulation of telomere maintenance via telomere lengthening",25,-0.60
BP,"GO:0042698","ovulation cycle",70,0.458085442182942,1.90163698729991,0.000158466385879
BP,"GO:0051098","regulation of binding",350,0.299466569960817,1.54417733987371,0.000158718
BP,"GO:0045981","positive regulation of nucleotide metabolic process",39,0.544542420181208,2.03
BP,"GO:1900544","positive regulation of purine nucleotide metabolic process",39,0.5445424201812
BP,"GO:0033003","regulation of mast cell activation",42,0.529264527837143,2.00229193362028,0.0
BP,"GO:0002291","T cell activation via T cell receptor contact with antigen bound to MHC molecule
BP,"GO:1902041","regulation of extrinsic apoptotic signaling pathway via death domain receptors",4
BP,"GO:0033209","tumor necrosis factor-mediated signaling pathway",99,0.415069380356061,1.84
BP,"GO:0045137","development of primary sexual characteristics",226,0.331366955835579,1.6347
BP,"GO:0009896","positive regulation of catabolic process",487,0.276526079418194,1.476101853
BP,"GO:0014031","mesenchymal cell development",87,0.424049897010019,1.83723129684715,0.0
BP,"GO:0044070","regulation of anion transport",87,0.424013367438468,1.83707302945318,0.000
CC,"GO:0031234","extrinsic component of cytoplasmic side of plasma membrane",95,0.4199399656
BP,"GO:2000104","negative regulation of DNA-templated DNA replication",14,-0.728704771722382,
BP,"GO:0071300","cellular response to retinoic acid",66,0.443399786336798,1.82468156171307,0.0
BP,"GO:0043583","ear development",213,0.332326962767262,1.63077459413784,0.000166007741
BP,"GO:0045913","positive regulation of carbohydrate metabolic process",72,0.452038505265374,1
BP,"GO:1904814","regulation of protein localization to chromosome, telomeric region",13,-0.73083
BP,"GO:0042269","regulation of natural killer cell mediated cytotoxicity",45,0.522330423899747,2.0
BP,"GO:0086065","cell communication involved in cardiac conduction",56,0.476936938297781,1.91
BP,"GO:0048260","positive regulation of receptor-mediated endocytosis",52,0.492645508864539,1.
CC,"GO:0005687","U4 snRNP",10,-0.794204094401041,-2.0644954123394,0.000171726531721605,
BP,"GO:1903018","regulation of glycoprotein metabolic process",49,0.511795328154607,1.9987425
CC,"GO:0031902","late endosome membrane",139,0.375681775695562,1.74494563259192,0.0001
BP,"GO:0033631","cell-cell adhesion mediated by integrin",16,0.694246118918136,2.08104263221
CC,"GO:0071004","U2-type prespliceosome",16,-0.684474646855859,-2.13043826353527,0.000174
CC,"GO:0071010","prespliceosome",16,-0.684474646855859,-2.13043826353527,0.0001748958500
MF,"GO:0042608","T cell receptor binding",10,0.806369308668751,2.15204521283985,0.00017507
CC,"GO:0005669","transcription factor TFIID complex",32,-0.547264414662573,-2.02329734594778,
BP,"GO:0008535","respiratory chain complex IV assembly",24,-0.608475121315491,-2.06583074105
BP,"GO:0000910","cytokinesis",155,-0.330760904226955,-1.66859812004605,0.000179858581419
BP,"GO:0010818","T cell chemotaxis",26,0.602362849865088,2.02674247345194,0.0001802122410
BP,"GO:0030032","lamellipodium assembly",67,0.460355239852628,1.89732167955976,0.0001806
BP,"GO:2000846","regulation of corticosteroid hormone secretion",15,0.722971777791278,2.12729
BP,"GO:0045622","regulation of T-helper cell differentiation",37,0.551349150538973,2.0368848772
BP,"GO:0048546","digestive tract morphogenesis",47,0.508392166520822,1.970868529727,0.0001

BP,"GO:0008625","extrinsic apoptotic signaling pathway via death domain receptors",81,0.4237692
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MF,"GO:0002039","p53 binding",64,-0.43859411962243,-1.88273466158674,0.0001839221039129
BP,"GO:0090025","regulation of monocyte chemotaxis",26,0.601495258504737,2.02382332885327,
BP,"GO:0060571","morphogenesis of an epithelial fold",23,0.641492672740141,2.1034597622446,0
BP,"GO:0010955","negative regulation of protein processing",27,0.613814807189842,2.0908739301
BP,"GO:1903318","negative regulation of protein maturation",27,0.613814807189842,2.090873930
BP,"GO:0034249","negative regulation of cellular amide metabolic process",181,-0.31413815615269
BP,"GO:0045055","regulated exocytosis",228,0.326555979864241,1.6096708482817,0.0001871622
MF,"GO:0003713","transcription coactivator activity",261,-0.288802806843612,-1.56640194482768
BP,"GO:0060419","heart growth",76,0.449114729235983,1.89200987247123,0.0001891299129227
MF,"GO:0001216","DNA-binding transcription activator activity",457,0.27830936968644,1.4763348
BP,"GO:2001185","regulation of CD8-positive, alpha-beta T cell activation",18,0.674289519123544,2
CC,"GO:0045120","pronucleus",12,-0.758095088644764,-2.11795784445153,0.00019021804971072
BP,"GO:0030166","proteoglycan biosynthetic process",56,0.474060387474555,1.90781277458727,0
CC,"GO:0031258","lamellipodium membrane",22,0.627676794813737,2.05331655872548,0.000192
BP,"GO:2000273","positive regulation of signaling receptor activity",44,0.515999582727267,1.9752
BP,"GO:0035886","vascular associated smooth muscle cell differentiation",27,0.612705999402504,2
CC,"GO:0009295","nucleoid",42,-0.500147933462203,-1.97626178906783,0.000193836184070802,
CC,"GO:0042645","mitochondrial nucleoid",42,-0.500147933462203,-1.97626178906783,0.0001938
MF,"GO:0017147","Wnt-protein binding",30,0.595379196125096,2.08812748566372,0.0001947234
BP,"GO:0072015","glomerular visceral epithelial cell development",12,0.756378620243112,2.10844
BP,"GO:0007019","microtubule depolymerization",44,-0.491325403359544,-1.95461805306076,0.0
BP,"GO:0002283","neutrophil activation involved in immune response",18,0.673893129994566,2.08
CC,"GO:0043296","apical junction complex",136,0.375315858782262,1.73469464435791,0.0001957
BP,"GO:0010875","positive regulation of cholesterol efflux",26,0.600877631970489,2.021745229365
BP,"GO:0106106","cold-induced thermogenesis",145,0.368544060543358,1.72444980128679,0.000
BP,"GO:0120161","regulation of cold-induced thermogenesis",145,0.368544060543358,1.72444980
BP,"GO:1902932","positive regulation of alcohol biosynthetic process",25,0.604455025789208,2.036
BP,"GO:0034698","response to gonadotropin",30,0.594871805734376,2.08634795452169,0.000200
MF,"GO:0034212","peptide N-acetyltransferase activity",41,-0.508910089929244,-1.991551549182
BP,"GO:0045987","positive regulation of smooth muscle contraction",30,0.594815363288112,2.086
MF,"GO:0019104","DNA N-glycosylase activity",13,-0.729170993331182,-2.08967886545548,0.0002
BP,"GO:0045833","negative regulation of lipid metabolic process",93,0.413523863712595,1.806799
BP,"GO:0010453","regulation of cell fate commitment",25,0.604002494370614,2.03546504899185,
BP,"GO:1904036","negative regulation of epithelial cell apoptotic process",46,0.506020440215037,1
BP,"GO:0007007","inner mitochondrial membrane organization",33,-0.532809350075048,-1.982961
BP,"GO:0000291","nuclear-transcribed mRNA catabolic process, exonucleolytic",19,-0.661548495834
BP,"GO:1903317","regulation of protein maturation",65,0.460550743687975,1.90050775712297,0.0
BP,"GO:0060688","regulation of morphogenesis of a branching structure",51,0.496967802502935,1.
BP,"GO:0051044","positive regulation of membrane protein ectodomain proteolysis",15,0.7200834
BP,"GO:0030449","regulation of complement activation",18,0.6721457802706,2.07724273965357,0
BP,"GO:0071825","protein-lipid complex subunit organization",47,0.504583831035605,1.956104870
BP,"GO:0001523","retinoid metabolic process",83,0.423547891261243,1.81751246876963,0.000216
BP,"GO:0110150","negative regulation of biomineralization",27,0.609886520056067,2.077492771777
BP,"GO:1900744","regulation of p38MAPK cascade",42,0.521876245282452,1.97434088497732,0.0
BP,"GO:0016266","O-glycan processing",42,0.521771829283803,1.97394586263813,0.00021776527
BP,"GO:0090503","RNA phosphodiester bond hydrolysis, exonucleolytic",40,-0.502760710522251,-1
BP,"GO:0060038","cardiac muscle cell proliferation",45,0.518598320592413,1.9876315928502,0.00
BP,"GO:0007040","lysosome organization",68,0.442632393368477,1.82877826977571,0.000220437

BP,"GO:0080171","lytic vacuole organization",68,0.442632393368477,1.82877826977571,0.000220
BP,"GO:1903580","positive regulation of ATP metabolic process",32,0.549956749020366,1.9742790
BP,"GO:0007173","epidermal growth factor receptor signaling pathway",102,0.394064255466892,1.
CC,"GO:0000178","exosome (RNase complex)",19,-0.659940448144988,-2.11351324611631,0.0002
BP,"GO:0032494","response to peptidoglycan",12,0.750865475832688,2.09307871325035,0.00022
BP,"GO:0007409","axonogenesis",422,0.278359989427849,1.46907544298749,0.000227556645676
BP,"GO:1901623","regulation of lymphocyte chemotaxis",26,0.597513587595087,2.0104263845535
BP,"GO:0015849","organic acid transport",309,0.299302276973008,1.5344528459694,0.000228552
CC,"GO:0005732","sno(s)RNA-containing ribonucleoprotein complex",23,-0.604383699577308,-2.03
BP,"GO:0002719","negative regulation of cytokine production involved in immune response",27,0.6
BP,"GO:1990748","cellular detoxification",101,0.404480412340441,1.79434621642235,0.00023102
CC,"GO:0031965","nuclear membrane",279,-0.278479307805994,-1.52382560043198,0.000231132
BP,"GO:0010878","cholesterol storage",19,0.671624188459651,2.11040288772118,0.00023325957
MF,"GO:0001848","complement binding",19,0.67129248659723,2.10936060160293,0.0002332595
BP,"GO:0051961","negative regulation of nervous system development",138,0.365677295944701,1.
MF,"GO:0001228","DNA-binding transcription activator activity, RNA polymerase II-specific",453,0.2
BP,"GO:0002833","positive regulation of response to biotic stimulus",168,0.353579357637594,1.68
BP,"GO:0045604","regulation of epidermal cell differentiation",54,0.477958111563726,1.908434168
BP,"GO:0034219","carbohydrate transmembrane transport",119,0.374278214457003,1.691771920
MF,"GO:0016798","hydrolase activity, acting on glycosyl bonds",135,0.368159871440771,1.6966408
BP,"GO:0010212","response to ionizing radiation",132,-0.33187107627821,-1.62569901821945,0.00
BP,"GO:0006301","postreplication repair",32,-0.540161089724898,-1.99703556442364,0.00023832
BP,"GO:0008585","female gonad development",105,0.400723530112914,1.78201802477255,0.0002
BP,"GO:0033028","myeloid cell apoptotic process",31,0.571127742788255,2.00861378163984,0.00
BP,"GO:0010559","regulation of glycoprotein biosynthetic process",44,0.510766546998313,1.95517
BP,"GO:0030177","positive regulation of Wnt signaling pathway",137,0.377075706704115,1.740681
BP,"GO:0002886","regulation of myeloid leukocyte mediated immunity",57,0.490837406919626,1.9
BP,"GO:0002437","inflammatory response to antigenic stimulus",67,0.455795240450691,1.8785279
BP,"GO:1901655","cellular response to ketone",88,0.428982017016778,1.86814314988908,0.00024
BP,"GO:0048736","appendage development",172,0.345246664156544,1.65690399661871,0.000243
BP,"GO:0060173","limb development",172,0.345246664156544,1.65690399661871,0.00024304026
BP,"GO:0048009","insulin-like growth factor receptor signaling pathway",34,0.561278506698033,2.0
BP,"GO:0071827","plasma lipoprotein particle organization",44,0.510066023467777,1.95248927271
MF,"GO:0016757","glycosyltransferase activity",256,0.314108952450913,1.57890675332713,0.0002
BP,"GO:0014033","neural crest cell differentiation",94,0.411838699877452,1.79898397102266,0.00
BP,"GO:0015914","phospholipid transport",95,0.412829805180851,1.80757967051209,0.00024987
BP,"GO:1905207","regulation of cardiocyte differentiation",27,0.606819403621119,2.06704507041
BP,"GO:0016101","diterpenoid metabolic process",86,0.418812775983543,1.81926043326038,0.00
BP,"GO:0001776","leukocyte homeostasis",88,0.428554788059889,1.86628264101555,0.00025319
BP,"GO:0043393","regulation of protein binding",188,0.332701458419575,1.61109280486321,0.000
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BP,"GO:0021700","developmental maturation",281,0.308813367097762,1.55958526368793,0.0002
BP,"GO:0055002","striated muscle cell development",66,0.437139633376821,1.7989197412712,0.0
BP,"GO:0072311","glomerular epithelial cell differentiation",23,0.63078624549378,2.068353299042
BP,"GO:0034104","negative regulation of tissue remodeling",22,0.621307959422578,2.0324822132
BP,"GO:0070525","tRNA threonylcarbamoyladenosine metabolic process",17,-0.660445097967155,-
BP,"GO:0060976","coronary vasculature development",46,0.502378997402763,1.9398616160563,0.
BP,"GO:0043523","regulation of neuron apoptotic process",198,0.334381080375334,1.6369226631
MF,"GO:0008194","UDP-glycosyltransferase activity",136,0.371671479067394,1.71785046944375,0
BP,"GO:0048146","positive regulation of fibroblast proliferation",46,0.502214402571054,1.9392260

BP,"GO:0016073","snRNA metabolic process",54,-0.461810373596891,-1.91448236392868,0.00026
CC,"GO:0005614","interstitial matrix",11,0.764467274617455,2.09830045926655,0.0002626789449
BP,"GO:0072207","metanephric epithelium development",24,0.618617304189268,2.051983807370
BP,"GO:0001765","membrane raft assembly",10,0.795327273945215,2.1225761374282,0.0002644
MF,"GO:0008080","N-acetyltransferase activity",59,-0.450968359542501,-1.90814835137982,0.000
MF,"GO:0004812","aminoacyl-tRNA ligase activity",22,-0.613759179240466,-2.02423899225216,0.0
BP,"GO:0006418","tRNA aminoacylation for protein translation",22,-0.613759179240466,-2.024238
MF,"GO:0016875","ligase activity, forming carbon-oxygen bonds",22,-0.613759179240466,-2.02423
BP,"GO:2000406","positive regulation of T cell migration",30,0.588827739374091,2.0651500672353
BP,"GO:1904994","regulation of leukocyte adhesion to vascular endothelial cell",29,0.581192345381
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BP,"GO:0002028","regulation of sodium ion transport",86,0.416987537417207,1.81133186828956,0
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BP,"GO:0002281","macrophage activation involved in immune response",17,0.696241953161317,2.
CC,"GO:0000314","organellar small ribosomal subunit",26,-0.58675211965864,-2.05271819063239,
CC,"GO:0005763","mitochondrial small ribosomal subunit",26,-0.58675211965864,-2.052718190632
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BP,"GO:0034763","negative regulation of transmembrane transport",108,0.394133905486092,1.753
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BP,"GO:0051770","positive regulation of nitric-oxide synthase biosynthetic process",17,0.695913140
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MF,"GO:0003925","G protein activity",39,0.529391657000508,1.97853243250551,0.000312414148
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BP,"GO:0007422","peripheral nervous system development",73,0.434602132392558,1.81971201800
BP,"GO:0050901","leukocyte tethering or rolling",33,0.545451427839169,1.9698344235342,0.0003
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CC,"GO:1904813","ficolin-1-rich granule lumen",124,0.375430233773618,1.7068809824358,0.0003

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BP,"GO:0010639","negative regulation of organelle organization",328,-0.266469633557481,-1.4859
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BP,"GO:0097048","dendritic cell apoptotic process",11,0.757624885655322,2.07951955342737,0.0
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BP,"GO:0036363","transforming growth factor beta activation",11,0.757553074129563,2.0793224
BP,"GO:0071229","cellular response to acid chemical",75,0.428302047958254,1.80172842018276,0
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BP,"GO:0002701","negative regulation of production of molecular mediator of immune response",38
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BP,"GO:0035108","limb morphogenesis",138,0.359508882432575,1.66391748279893,0.000374074
BP,"GO:0032740","positive regulation of interleukin-17 production",23,0.618705620197717,2.02874
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BP,"GO:0001886","endothelial cell morphogenesis",11,0.757379969896246,2.0788473116365,0.000
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BP,"GO:0032376","positive regulation of cholesterol transport",14,0.718811197727375,2.096127928
BP,"GO:0007006","mitochondrial membrane organization",103,-0.362033022049895,-1.6836169364
BP,"GO:0060394","negative regulation of pathway-restricted SMAD protein phosphorylation",11,0.7
MF,"GO:0032451","demethylase activity",38,-0.498562076713969,-1.91415186831736,0.00038505
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CC,"GO:0016010","dystrophin-associated glycoprotein complex",17,0.686728134850262,2.0961877
BP,"GO:0032410","negative regulation of transporter activity",72,0.437526394591919,1.822861834
BP,"GO:0014002","astrocyte development",40,0.519938035097289,1.9536194320704,0.000400387
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MF,"GO:0022803","passive transmembrane transporter activity",462,0.273527878976943,1.455180
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BP,"GO:0098751","bone cell development",34,0.55214050962101,2.00147888962897,0.000410969
BP,"GO:0042832","defense response to protozoan",23,0.616668976923871,2.02206265902177,0.0
BP,"GO:0072538","T-helper 17 type immune response",38,0.532346251192819,1.98032725547889,
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BP,"GO:0051349","positive regulation of lyase activity",49,0.493771363635529,1.9283525632407,0.
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BP,"GO:2000047","regulation of cell-cell adhesion mediated by cadherin",19,0.655439496153291,2.
BP,"GO:0002053","positive regulation of mesenchymal cell proliferation",23,0.614792435810992,2.(
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BP,"GO:0000082","G1/S transition of mitotic cell cycle",198,-0.299746291961976,-1.5632795357080
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MF,"GO:0070325","lipoprotein particle receptor binding",28,0.581316526299361,1.9905412537368
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BP,"GO:0071637","regulation of monocyte chemotactic protein-1 production",18,0.6550319092548
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BP,"GO:0021545","cranial nerve development",54,0.463659490312536,1.85134134644717,0.00052
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MF,"GO:0019001","guanyl nucleotide binding",367,0.284630133518755,1.47395362685932,0.00054
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BP,"GO:1903846","positive regulation of cellular response to transforming growth factor beta stimul
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BP,"GO:1902369","negative regulation of RNA catabolic process",68,-0.384006023443644,-1.685678
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BP,"GO:0032413","negative regulation of ion transmembrane transporter activity",64,0.4413929353
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BP,"GO:0010824","regulation of centrosome duplication",45,-0.459475051609471,-1.848146444255
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BP,"GO:0048026","positive regulation of mRNA splicing, via spliceosome",22,-0.579389429609511,-
BP,"GO:0070227","lymphocyte apoptotic process",74,0.416263689833353,1.74522581575156,0.001
BP,"GO:0051894","positive regulation of focal adhesion assembly",26,0.56658386846118,1.9063585
BP,"GO:0031331","positive regulation of cellular catabolic process",425,0.271945791600462,1.4348
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BP,"GO:0001101","response to acid chemical",116,0.361771894513254,1.62935010766025,0.00174
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BP,"GO:0044380","protein localization to cytoskeleton",57,-0.415660186175729,-1.7499272388044
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BP,"GO:0060390","regulation of SMAD protein signal transduction",30,0.517014632823844,1.81328
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CC,"GO:1990531","phospholipid-translocating ATPase complex",13,0.649230247523099,1.86149012
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BP,"GO:0099625","ventricular cardiac muscle cell membrane repolarization",26,0.515995696357699

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BP,"GO:0032060","bleb assembly",10,0.702861857591329,1.87580365430116,0.005810102083905
BP,"GO:0099177","regulation of trans-synaptic signaling",406,0.256400062400604,1.348780113536
BP,"GO:0043117","positive regulation of vascular permeability",17,0.603718857747354,1.84280798
BP,"GO:1902882","regulation of response to oxidative stress",83,0.36845553982611,1.581102282977
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BP,"GO:0014841","skeletal muscle satellite cell proliferation",14,0.624436215542761,1.8209207018
BP,"GO:0047484","regulation of response to osmotic stress",14,0.624014810076431,1.81969183984
BP,"GO:1901741","positive regulation of myoblast fusion",14,0.623860546156451,1.819241990269
BP,"GO:1903203","regulation of oxidative stress-induced neuron death",26,0.515013598762706,1.7
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BP,"GO:0097186","amelogenesis",23,0.551554840760452,1.80855286975477,0.0058565637470571
BP,"GO:0006971","hypotonic response",11,0.678713854400968,1.86292551648175,0.00586389932
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BP,"GO:0060037","pharyngeal system development",28,0.509926321278379,1.74608725702765,0.0
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BP,"GO:0006285","base-excision repair, AP site formation",12,-0.655903928938027,-1.83245728973
BP,"GO:0051573","negative regulation of histone H3-K9 methylation",12,-0.6558972307447,-1.832
BP,"GO:1902548","negative regulation of cellular response to vascular endothelial growth factor stin
BP,"GO:0001771","immunological synapse formation",14,0.622311576964775,1.81472503561914,0
CC,"GO:0005801","cis-Golgi network",65,0.395555766812807,1.63229961845808,0.006253664489
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MF,"GO:0048185","activin binding",15,0.620981874550581,1.82719495536743,0.00636976336014(
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BP,"GO:0001893","maternal placenta development",33,0.480318816141029,1.73461556797826,0.0
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BP,"GO:0033687","osteoblast proliferation",29,0.50985417593679,1.76612693950824,0.006557592
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BP,"GO:2000341","regulation of chemokine (C-X-C motif) ligand 2 production",16,0.5965902462144
BP,"GO:0043301","negative regulation of leukocyte degranulation",14,0.618725249507599,1.80426
MF,"GO:0004112","cyclic-nucleotide phosphodiesterase activity",25,0.516269866458086,1.7398094
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BP,"GO:0097320","plasma membrane tubulation",19,0.579548851399506,1.82108028654255,0.006
BP,"GO:0051155","positive regulation of striated muscle cell differentiation",43,0.440769893916868
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MF,"GO:0070700","BMP receptor binding",13,0.644332622107131,1.84744752050658,0.006786357
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BP,"GO:2000241","regulation of reproductive process",167,0.315274745805732,1.50689916966423
MF,"GO:0051393","alpha-actinin binding",28,0.506280293731548,1.7336025470359 5,0.006816899
BP,"GO:1900272","negative regulation of long-term synaptic potentiation",11,0.672794228002445,1
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BP,"GO:0060420","regulation of heart growth",58,0.409611061256173,1.64456252716095,0.00683(
BP,"GO:0045762","positive regulation of adenylate cyclase activity",36,0.47051299520385,1.719303
BP,"GO:0099622","cardiac muscle cell membrane repolarization",36,0.469841211127314,1.7168485
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MF,"GO:0106310","protein serine kinase activity",350,-0.236249365465961,-1.32953093381357,0.0
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BP,"GO:0070232","regulation of T cell apoptotic process",34,0.473280665636018,1.7156163053318
BP,"GO:0070166","enamel mineralization",18,0.569939649736385,1.76137831138206,0.007049293
BP,"GO:0046058","cAMP metabolic process",22,0.532537938715045,1.74208920378779,0.0070540
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BP,"GO:0051099","positive regulation of binding",166,0.307564725780203,1.47018747864848,0.00
BP,"GO:0034976","response to endoplasmic reticulum stress",243,0.286055816899525,1.42247619
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MF,"GO:0030228","lipoprotein particle receptor activity",18,0.569236646920217,1.7592057060653
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BP,"GO:0050769","positive regulation of neurogenesis",214,0.285822887856045,1.4025229736275
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BP,"GO:0035994","response to muscle stretch",24,0.531266996069392,1.76223856969776,0.00728
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BP,"GO:0001508","action potential",138,0.316350936827592,1.46416926036861,0.0074080858496
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MF,"GO:0008237","metallopeptidase activity",179,0.298570526644446,1.44402499024727,0.00742
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BP,"GO:0070050","neuron cellular homeostasis",31,0.492154067578309,1.73086924126954,0.0074
BP,"GO:0009791","post-embryonic development",84,0.366681589254331,1.58014685860116,0.007
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BP,"GO:0015844","monoamine transport",81,0.360104739045451,1.53081525179891,0.007468338
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BP,"GO:1903319","positive regulation of protein maturation",26,0.504338539794252,1.6969246025
MF,"GO:0008238","exopeptidase activity",96,0.349809126628708,1.53133023979716,0.007518210
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BP,"GO:0016973","poly(A)+ mRNA export from nucleus",22,-0.50871080666505,-1.6777789814985
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BP,"GO:0009168","purine ribonucleoside monophosphate biosynthetic process",18,-0.55134243502
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BP,"GO:0044088","regulation of vacuole organization",49,0.428138430118196,1.67203264495038,0
MF,"GO:0008376","acetylgalactosaminyltransferase activity",33,0.474696719859979,1.7143120208
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MF,"GO:0061980","regulatory RNA binding",40,-0.419204276014992,-1.63798089124604,0.008159
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MF,"GO:0120015","sterol transfer activity",23,0.540281581635266,1.77158776014893,0.008232921
BP,"GO:0042391","regulation of membrane potential",403,0.254564444928655,1.33842743462819,
BP,"GO:0051963","regulation of synapse assembly",98,0.348174816637289,1.53724046270862,0.00
BP,"GO:0021795","cerebral cortex cell migration",42,0.432203140592315,1.6350932597539,0.0083
BP,"GO:0070471","uterine smooth muscle contraction",11,0.661612249110883,1.81598523868852,
BP,"GO:0032836","glomerular basement membrane development",11,0.660983433865076,1.81425
BP,"GO:0055008","cardiac muscle tissue morphogenesis",59,0.404135443399038,1.6330380990710
BP,"GO:0070193","synaptonemal complex organization",25,-0.499810738037694,-1.7125845495586
BP,"GO:0007163","establishment or maintenance of cell polarity",210,0.287492229190586,1.41133(
BP,"GO:0032272","negative regulation of protein polymerization",73,0.369899073085134,1.548795
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BP,"GO:0097152","mesenchymal cell apoptotic process",15,0.606328449270253,1.78407829472344
BP,"GO:0002921","negative regulation of humoral immune response",16,0.587195979290918,1.760
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BP,"GO:0002065","columnar/cuboidal epithelial cell differentiation",87,0.358598155136515,1.5536
BP,"GO:0050852","T cell receptor signaling pathway",121,0.336556337564518,1.52222533136759,0
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BP,"GO:0002566","somatic diversification of immune receptors via somatic mutation",13,-0.625405
BP,"GO:0090181","regulation of cholesterol metabolic process",32,0.455160701765203,1.63397257
MF,"GO:1904047","S-adenosyl-L-methionine binding",14,-0.616920211977402,-1.81764688116192,
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BP,"GO:0030004","cellular monovalent inorganic cation homeostasis",105,0.339095855598336,1.50
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BP,"GO:0035725","sodium ion transmembrane transport",173,0.293886123708434,1.41448345642
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BP,"GO:2000191","regulation of fatty acid transport",31,0.464057339852132,1.63205514014698,0.(
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,1.33245901639344e-09,6.07765314926661e-10,2907,"tags=31%, list=16%, signal=26%","11031/532
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75e-09,1.81722156728197e-08,8.28876704305728e-09,5177,"tags=60%, list=29%, signal=43%","269
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98,2.01380894326167e-09,2.17662753867431e-08,9.92810064133202e-09,3459,"tags=37%, list=19%
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6,2.19371215909077,2.49215077516965e-09,2.6444127285351e-08,1.20617768725402e-08,4202,"t
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43,3.01888503885707e-09,3.15797909856246e-08,1.44042716342947e-08,3045,"tags=82%, list=17%
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0952809e-09,3.45364583128444e-08,1.5752875851875e-08,3527,"tags=44%, list=20%, signal=36%",
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8,4.04501659603938e-09,4.13558426322114e-08,1.88633544538242e-08,3045,"tags=85%, list=17%
938,4.25490974772133e-09,4.33551593953312e-08,1.97752889803093e-08,3104,"tags=57%, list=17
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4143874,4.41188247613316e-09,4.46572612279083e-08,2.03692076829534e-08,4027,"tags=52%, lis
3760847797,4.53012388252869e-09,4.5797073280091e-08,2.08891022705761e-08,4437,"tags=72%
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.69e-08,1.22480146547353e-07,5.58660220860672e-08,4315,"tags=67%, list=24%, signal=51%","709
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3513016677,1.40200925486664e-08,1.31892722494862e-07,6.01593152490291e-08,3104,"tags=60%
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5168e-08,1.54575936521574e-07,7.05056527700195e-08,3571,"tags=57%, list=20%, signal=45%","68
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e-08,1.64277287511927e-07,7.49306628959035e-08,4671,"tags=63%, list=26%, signal=47%","6387/4
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06556e-08,1.65163474077894e-07,7.533487304475e-08,5180,"tags=50%, list=29%, signal=36%","677
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54184531e-08,1.77509471042763e-07,8.09661672467636e-08,5141,"tags=43%, list=29%, signal=31%
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3058e-08,1.80297961378312e-07,8.22380620563635e-08,3309,"tags=38%, list=18%, signal=31%","1
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2,2.65663711066644e-08,2.36507627990108e-07,1.07876588502533e-07,2827,"tags=49%, list=16%
,-2.53549110915921,2.68525946160772e-08,2.38794189321089e-07,1.08919541906972e-07,2576,"
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62,2.41203363314359,3.40396293761782e-08,2.97179492555936e-07,1.35550426437786e-07,3123
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3538e-08,7.34088767766135e-07,3.34834832168505e-07,4365,"tags=67%, list=24%, signal=51%","54
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7771066987,1.0189043022878e-07,8.14321943854006e-07,3.7143103582856e-07,3806,"tags=50%,
82797249e-07,8.20755402137561e-07,3.74365484657129e-07,3083,"tags=48%, list=17%, signal=40%
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3757e-07,1.18364961563232e-06,5.3988991222765e-07,4126,"tags=36%, list=23%, signal=28%","110
391397e-07,1.19868026425002e-06,5.46745738019138e-07,3800,"tags=50%, list=21%, signal=40%",
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07,1.27630080556259e-06,5.82150258649947e-07,4132,"tags=55%, list=23%, signal=42%","135228/
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4034149,2.42527123464653e-07,1.81850595896743e-06,8.29462545001463e-07,4740,"tags=50%, l
5524322660908e-07,1.83928266782291e-06,8.38939281505425e-07,3488,"tags=39%, list=19%, sign
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723911533,2.49205981048559e-07,1.86171527018629e-06,8.49171309261827e-07,4027,"tags=72%,
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39,1.17480448437492e-06,7.53163701811906e-06,3.43535349899183e-06,5688,"tags=65%, list=32%
4,1.17496502694082e-06,7.53163701811906e-06,3.43535349899183e-06,4654,"tags=40%, list=26%,
963,2.22765177882857,1.18571873988313e-06,7.58483254299904e-06,3.45961720581779e-06,451
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4,1.639805487421e-06,1.02604611252948e-05,4.6800331645369e-06,5027,"tags=46%, list=28%, sig
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1,1.77256972590177e-06,1.1040189066766e-05,5.03568507733497e-06,4764,"tags=63%, list=27%,
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06,2.0501870842199e-05,9.35137654193752e-06,4944,"tags=43%, list=28%, signal=31%","9429/493
54991608915737e-06,2.0758070483936e-05,9.46823511246667e-06,3685,"tags=64%, list=21%, sigr
7846,3.57227769219533e-06,2.08738124242729e-05,9.52102768316014e-06,4744,"tags=47%, list=
-06,2.09326597603458e-05,9.54786931153363e-06,3259,"tags=37%, list=18%, signal=31%","135228
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83,3.80214599586703e-06,2.20584173122107e-05,1.00613533171371e-05,5074,"tags=67%, list=28
260608e-06,2.20854049776193e-05,1.00736630142943e-05,4238,"tags=48%, list=24%, signal=37%",
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8e-06,2.42447721336637e-05,1.10586002194835e-05,4050,"tags=73%, list=23%, signal=57%","5764
3275611e-06,2.45670200385387e-05,1.12055849274416e-05,2807,"tags=37%, list=16%, signal=31%
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398e-06,2.98835279857959e-05,1.36305669247272e-05,3820,"tags=46%, list=21%, signal=36%","32
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7723e-06,3.31181947436495e-05,1.51059730997635e-05,4481,"tags=36%, list=25%, signal=28%","3
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19634616113e-06,3.46015646790493e-05,1.57825723683707e-05,4263,"tags=55%, list=24%, signal=
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95,7.16442269991764e-06,3.91346960382598e-05,1.78502382209323e-05,4078,"tags=52%, list=23%
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430558338e-06,4.00471319354673e-05,1.82664212956791e-05,2801,"tags=82%, list=16%, signal=7
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031e-06,4.18785131069476e-05,1.910175652231e-05,3800,"tags=45%, list=21%, signal=36%","6284
86,7.79786838336928e-06,4.23387269338848e-05,1.93116706720294e-05,922,"tags=44%, list=5%,
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911,-2.13450691586584,8.50245608910823e-06,4.56761157252291e-05,2.08339307377042e-05,29
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909983413e-06,5.03178013390903e-05,2.29511106916023e-05,3553,"tags=55%, list=20%, signal=44
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n of immune receptors built from immunoglobulin superfamily domains",99,0.450310610073181,2.0
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51,2.14881793444014,1.13724421530651e-05,5.93294029654128e-05,2.70615102108609e-05,3318
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050307,1.21798450338457e-05,6.30960997036954e-05,2.87795875409779e-05,3632,"tags=32%, lis
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384e-05,0.00026127748159278 8,0.00011917469050064 8,4936,"tags=63%, list=28%, signal=46%","4(
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,0.00010718941211865 2,0.00044046286233589 8,0.00020090527884726 4,4691,"tags=67%, list=26%
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8,-0.467214537683279,-1.97906707932065,0.00011119620316359 47,0.00045776025811719,0.00020
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9,0.00718838517391895,0.0168767171269824,0.00769786025192087,3948,"tags=50%, list=22%, sig
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=62%, list=17%, signal=52%","4208/5592/183/5155/9510/3488/2771/387/4217/7043/2697/1906/93
, list=20%, signal=44%","6678/1969/23166/90993/183/10769/7057/7058/493/9510/3339/9314/128
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al=47%","8379/8697/23212/8452/54984/55166/150465/79003/54908/91782/81929/51112/283489/
%, signal=37%","56034/677/80310/5154/1956/183/5155/80243/375033/284/5159/4734/1634/3082/
0%, list=22%, signal=39%","6788/558/2152/3688/5154/1956/2022/7057/3488/284/388121/10979/1
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=47%","11031/3920/10981/567/306/9341/3133/3107/3916/23682/83547/3105/55647/7096/83871
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, list=27%, signal=42%","558/677/301/7048/4323/861/1435/2737/4853/3707/146433/9580/11454
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%, list=25%, signal=40%","7132/10203/10769/4907/5580/9314/114548/7538/58189/3082/7010/284
gs=51%, list=25%, signal=39%","683/3572/4208/301/7048/1435/3280/3133/604/3105/23529/4773/
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1%","51474/1601/55304/5321/1910/3269/54884/3953/8871/3707/9023/5978/151056/10577/1582
gnal=44%","283375/79811/10772/6432/58517/9667/6829/84530/29896/904/23144/1655/10914/5
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','51429/1601/4643/11037/1956/8301/8724/5872/1889/8322/928/9341/1839/7851/1906/5045/85
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"tags=58%, list=22%, signal=45%","23353/254528/57820/4292/27030/317761/9134/10734/546/101
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0752087,3.79043002142257e-10,4.68216036688794e-09,2.13564142304436e-09,4897,"tags=55%, li
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%, list=25%, signal=39%","683/3572/4208/301/7048/3280/3133/604/3105/23529/4773/8767/3479/
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gnal=48%","51340/6433/6431/55015/6606/60680/22827/91746/55692/11168/10291/6637/8175/16
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%","7049/5327/3673/677/2308/1410/7048/4323/4000/5154/4780/6387/283/84962/5351/824/7057
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, list=26%, signal=51%","965/7098/2152/84632/10392/23586/11027/7096/929/2149/7097/5054/70
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%, signal=41%","255374/114034/79042/90806/8602/57523/55720/9924/10556/27340/11340/8429
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%, list=18%, signal=50%","10413/25937/3572/4208/182/90/5587/1052/2737/860/1192/10979/3082
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=78%, list=28%, signal=56%","10392/114548/4023/351/3428/11027/8767/7099/79792/1958/7474/4
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26%, signal=31%","831/5627/135228/302/2/1410/8301/334/183/7057/871/2719/7078/7079/7035/3
29/23189/3685/7094/8826/3688/93663/1969/4651/1956/4627/6282/4641/23022/3678/152273/10
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, signal=30%","10413/7049/4208/90/3688/7048/4000/861/57493/6444/183/2022/2735/4853/5159
signal=41%","144715/1027/7158/25988/1112/7919/1386/80254/26058/51720/6118/545/54465/10
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gnal=29%","2059/1793/389/6678/23189/54873/3074/301/1601/57381/6281/4739/93663/4651/11
13/57381/1410/182/1969/147912/7048/4124/2200/1956/10252/1290/25932/4094/2737/7070/104
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9%, signal=69%","64785/9400/84296/3159/4174/9401/51659/4172/641/5888/4175/8318/9837/41
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2%, list=24%, signal=39%","301/861/6387/2615/3077/2737/3958/6385/9314/604/1051/8654/10875
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ıl=25%","3913/57381/1410/182/1969/147912/7048/4124/2200/1956/10252/25932/4094/2737/707
5%, signal=32%","79971/93664/1601/5321/6387/283/3077/183/3624/8754/6385/3916/28984/7043
'1435/3399/2737/1278/4038/860/3624/5066/116039/358/10468/1896/3912/54829/387/947/7043/
t=21%, signal=34%","10413/25937/7414/135228/677/7132/182/1903/10252/3280/857/7538/16174
signal=49%","22925/395/301/1956/283/183/5159/1909/3356/6401/2099/2769/400/185/2260/532
'7414/90102/395/3673/6281/4323/84962/7205/7070/7057/9748/2042/1307/79834/6385/3987/109
14034/90806/254528/57455/9924/27301/22894/81875/7486/64789/54512/10111/83940/129563/2
6, signal=70%","80310/3280/5155/4853/284/5159/2303/1909/947/8837/7010/4851/5175/4854/59/
%, signal=47%","8379/8697/23212/8452/54984/79003/54908/91782/81929/283489/27183/91272/
2%","11031/558/302/3685/3673/1601/11037/5587/8301/3077/30844/2042/2719/567/257364/857/
6, list=26%, signal=30%","831/5627/135228/302/2/1410/8301/334/183/7057/871/7078/7079/7035/

1303/3685/4323/1290/3693/64321/83595/1289/3624/3678/1847/1295/1284/3912/1843/2335/809
%","3074/5376/443/928/58494/7851/3082/5803/3691/84668/79628/50846/5653/5354/7097/2313
074/5376/443/928/58494/7851/3082/5803/3691/84668/79628/50846/5653/5354/7097/23136/856
45%, list=15%, signal=38%","25937/90102/1601/182/90/7048/10252/2022/2042/10979/28984/7043
st=22%, signal=41%","2059/10161/143872/90627/23189/25959/3688/57493/2151/9754/5159/1281
, list=25%, signal=54%","23328/3656/4846/7040/654/23643/4067/11027/8767/929/7097/7099/479
%","7049/1842/11167/83716/3673/3075/2200/334/283/7057/7123/6649/7058/1289/9510/10468/1
9%","55308/23049/6428/56943/55677/26993/91746/56001/728343/4927/8888/84271/7175/5574
nal=40%","5429/9477/51082/10623/548644/64895/10728/5442/11201/5434/5423/10914/27434/64
nal=43%","9469/56548/5155/284/5159/10090/7040/6484/10678/11227/337876/4790/84752/2994
%, signal=32%","5125/3490/3673/367/85360/7048/7026/183/2013/824/7005/8528/1307/5069/348
=70%, list=9%, signal=63%","8379/56155/79003/81929/283489/55920/9184/91272/3192/4751/1061
%, list=17%, signal=55%","8379/25906/56155/7175/51451/54908/7517/84861/9184/55055/55755/7
7%, signal=55%","8379/25906/56155/7175/51451/54908/7517/84861/9184/55055/55755/7272/104
%, signal=55%","8379/25906/56155/7175/51451/54908/7517/84861/9184/55055/55755/7272/1040
list=20%, signal=40%","143872/5125/367/5547/7026/1910/183/2013/5155/1889/387/4846/1906/9
gnal=49%","5627/5327/5592/5154/7057/5155/7035/7056/928/5580/947/4846/10019/1906/710/53
%","79971/10413/25937/6788/26524/2239/23189/3487/1601/80114/2308/57664/23500/1956/299
ignal=37%","23328/7098/1956/55075/7205/4306/329/10392/1896/9111/3987/114548/387/351/19
%, signal=32%","79971/10413/25937/6788/26524/23189/3487/1601/80114/2308/57664/23500/195
47%","5327/3673/4208/5592/80310/5154/4780/183/5155/9510/6282/3488/5159/387/80005/9364
9%","7414/389/7094/1969/57493/5872/183/6591/4641/857/928/9948/387/28513/1009/2697/909
8%, list=26%, signal=35%","3688/4780/183/7070/2719/1909/154043/6548/1906/3779/5243/30845/
ignal=52%","90102/3913/858/3688/4811/83872/857/1282/3912/22795/1605/3914/23187/8985/15
list=17%, signal=40%","90627/4208/7026/6672/7057/493/79812/9314/28984/1634/4070/387/1617
signal=29%","558/22925/493869/677/301/2308/8531/80310/4780/1956/5587/4638/80303/6649/1
","9843/302/301/27202/2683/3680/1956/4478/57477/56606/493/5155/3953/4641/10351/358/103
gnal=46%","11031/3673/4240/10461/5322/54209/3601/6347/64581/5873/4153/2150/5806/2621/2
=42%","10413/25937/4208/182/26508/3280/183/2737/2719/4853/3217/30812/2246/4070/4851/2
t=21%, signal=38%","56034/23328/2690/80310/5154/1956/84962/7057/5155/146433/8322/8754/7
6%, signal=44%","9469/56548/5155/284/5159/10090/7040/6484/10678/11227/337876/4790/2994
3738,"tags=34%, list=21%, signal=27%","3556/8763/2239/90/3688/7048/84962/83872/10205/23022
0%","8379/8697/23212/8452/54984/55166/150465/79003/54908/91782/81929/51112/283489/271
gnal=53%","56257/60487/91893/29081/23070/29063/25917/1787/4234/115708/64216/64863/1529
,3676,"tags=56%, list=20%, signal=45%","3685/117854/2934/3956/8724/10581/10410/857/9023/10
3%, signal=38%","2239/1601/367/57664/5154/1435/10252/183/6591/2719/3217/860/30812/2246/
1303/3685/677/3915/4323/1290/3693/64321/83595/1289/55614/3624/3678/1847/1295/1284/391
7%","22925/302/3685/1601/5321/183/7057/5580/6342/9388/1906/5045/654/5468/185/114897/5
2626,"tags=57%, list=15%, signal=49%","90627/301/57381/10769/7057/79812/10519/27286/2303/9
9%, list=24%, signal=30%","51474/2059/90102/51429/389/23189/395/1410/6281/93663/11346/111
34%","143872/3673/5592/10203/1910/4638/183/493/857/477/624/387/1909/4629/3356/151/800
, signal=29%","10413/7049/4208/90/3688/7048/4000/861/57493/6444/183/2022/2735/4853/5159/
45%","26013/8085/22955/23347/353274/1017/55729/51548/51535/7175/57343/55183/8312/803
gs=45%, list=25%, signal=34%","301/3956/4853/3958/6778/3916/114548/604/7040/23529/8767/10
ags=34%, list=18%, signal=28%","6788/56034/26524/23328/2690/1601/8826/80310/7026/5154/574
7/58191/6358/2920/6360/6347/10344/6376/6368/2919/6356/2921/6357/6355/3576/6364/56477/
174e-08,3464,"tags=41%, list=19%, signal=33%","6788/23328/7098/4055/344558/7164/10392/4616
signal=47%","7414/7132/1903/1910/4478/5906/1909/91663/5175/1003/5962/9076/54538/7082/5
list=32%, signal=49%","100507436/3140/567/3133/3916/3105/3106/135250/3122/7305/8417/3123
25%, signal=50%","5627/5327/5592/5154/7057/5155/7035/7056/928/5580/947/4846/10019/1906/

6, list=23%, signal=28%","11031/5327/23189/858/2690/677/2308/2200/4780/81565/5872/183/5166
7%, signal=41%","8379/8697/25906/2253/56155/8452/23113/9820/5300/7175/51451/54908/7517/
gnal=24%","831/5627/135228/5327/302/2/64215/1410/8301/334/3077/183/29761/7057/871/2719
signal=76%","55166/79003/54908/51112/283489/27085/55920/9184/55055/7272/983/10403/921
3%, list=14%, signal=76%","55166/79003/54908/51112/283489/27085/55920/9184/55055/7272/983
, list=22%, signal=39%","558/7049/90/4052/1969/7048/1956/2022/2050/10461/2042/5159/51232/
=49%, list=21%, signal=39%","56034/80310/5154/1956/183/5155/284/5159/4734/1634/3082/7010
=54%, list=17%, signal=45%","5587/4641/5159/1634/967/64094/7424/2549/3791/3315/23554/5291
","7098/351/3459/120892/8877/7097/199/728/54209/7305/3569/6376/3460/712/1191/3684/1033
98/351/3459/120892/8877/7097/199/728/54209/7305/3569/6376/3460/712/1191/3684/10333/107
44%","5929/26108/9682/26013/727866/51147/8535/23512/29057/22823/26147/6879/79831/546/
25%, signal=44%","5929/26108/9682/26013/727866/51147/8535/23512/29057/22823/26147/6879/
%","10413/25937/4208/182/26508/3280/183/2737/2719/4853/3217/1282/30812/2246/4070/4851
gnal=48%","54623/55818/9682/3720/55011/22823/55677/26147/22938/546/4302/64324/10445/9
1/840/5327/5125/302/2152/6281/4323/2934/861/4627/824/2737/7057/7123/1889/25825/837/120
%, list=17%, signal=50%","8379/25906/56155/7175/51451/54908/7517/5885/84861/27085/1111/91
=65%","375748/545/7507/2072/9025/5892/64789/729475/165918/9937/64421/22909/7517/64710
86374979703e-08,2492,"tags=42%, list=14%, signal=36%","135228/7049/858/5654/4052/2200/1025
52/54505/6125/6135/27341/51388/4869/3692/29777/6208/26609/3921/29093/51335/6134/83939
t=18%, signal=56%","7049/182/90/7048/26508/10252/2022/6591/4851/23493/7042/23462/4487/6
43%, list=17%, signal=36%","135228/7049/6678/858/4208/1601/687/367/7026/6591/7057/79812/2
6, list=17%, signal=46%","90627/4208/7026/6672/7057/493/79812/9314/28984/1634/387/161742/7
=53%","8379/25906/56155/7175/51451/54908/7517/84861/9184/55055/55755/7272/10403/34865
49/11167/90/5654/2200/26508/3280/3077/2022/2719/4853/860/10272/10468/81792/7043/51232/
signal=35%","7049/11167/90/5654/2200/26508/3280/3077/2022/2719/4853/860/10272/10468/817
5","23328/7098/1956/55075/7205/4306/329/10392/1896/9111/3987/114548/387/351/1906/8797/5
:=24%, signal=44%","183/493/857/9314/947/4846/1728/10365/1906/5743/7097/6648/7099/80149/
=45%","6387/58191/10763/3716/6358/2920/6360/6347/10344/6376/6368/2919/9034/6356/57402/
signal=32%","5587/7070/493/857/3270/477/845/3356/5350/80024/4067/51393/59341/22821/2149
17%","2059/9936/389/3685/3688/4651/4643/2683/5154/25932/7402/4478/5205/10788/4641/284/
l=30%","51429/54843/302/221749/117583/7094/8826/57664/4651/5321/4643/2934/8301/8724/53
83/3688/80310/6387/3554/7070/58494/6484/6401/5175/7412/6403/1991/3676/5734/6347/10894
4174/51659/4172/4175/8318/9837/4173/4171/4176"
%","114034/90806/254528/9924/27301/22894/81875/7486/64789/54512/10111/83940/129563/29
58%","10199/92856/65083/8602/55813/27340/9724/84294/9277/55127/6201/25879/134430/5705
4478/58191/1901/1893/387/351/347/23143/1880/199/7474/3676/6347/10344/5339/3383/29108/
6%","558/4208/301/6387/2737/10461/355/3707/3958/604/4254/1514/4067/80301/652/7474/3586
7/301/7048/861/3140/2737/567/860/3953/3707/9580/6778/639/114548/387/604/1514/8546/8767
st=22%, signal=40%","90102/389/395/6281/11346/84952/1901/10174/6385/10979/28984/4070/387
0%","51574/114034/11340/118460/54931/26024/26173/54512/55520/90459/3028/345630/79650/
=48%","9697/3673/861/90993/2022/871/1289/7431/5159/28984/7043/7040/57333/1277/2149/65
3%, signal=31%","3490/301/1601/83643/5154/824/5166/6591/5155/5978/5580/2246/27122/15750
38%, list=24%, signal=30%","79026/2040/4208/3688/7402/23704/857/3270/477/4734/1909/15404
3521/6122/6232/6129/6159/6188/6231/6137/6146/6158/6152/25873/54505/6125/6204/6229/613
%","26260/57610/79139/1027/8454/8816/90864/7322/51433/26094/26133/342897/59349/25898/
gnal=29%","79971/10413/25937/6788/26524/23189/3487/1601/80114/2308/57664/23500/1956/1
.0413/7049/5327/2/3490/687/367/196/387758/2908/4478/7057/6446/857/114907/358/978/8837/
:=20%, signal=47%","3685/117854/2934/3956/8724/10581/10410/857/9023/10475/2519/81603/67
%","51474/3953/9023/10577/158219/2246/51700/351/10826/8542/3990/19/6646/8578/51302/94
"26013/26155/64431/6603/8289/3066/4302/84656/57459/6601/23523/8467/165918/6599/3151/5

ignal=39%","57820/4292/27030/317761/7273/10734/23212/10111/11144/6002/23476/84365/1012
76/443/928/58494/7851/3082/5803/3691/84668/79628/50846/5653/5354/7097/23136/8560/1039
ags=89%, list=16%, signal=74%","254048/546/23523/23421/79711/5928/29855/401541/387103/46
35,"tags=49%, list=25%, signal=37%","3556/7098/3656/367/117854/5587/183/857/10519/10392/18
5%","7049/4208/182/90/3688/4000/1956/6444/10252/3280/183/64321/5978/5159/1909/4629/227
=88%, list=16%, signal=74%","254048/546/23523/23421/79711/5928/29855/401541/387103/4678/
3572/2690/4208/2632/1435/8301/824/355/284/7068/5978/2730/10763/387/4217/947/6548/1728
%, signal=40%","22925/302/1601/5321/10226/5580/6342/4023/4481/9388/1906/654/114897/5170
0%","8260/8658/6421/10734/546/23383/79892/54921/5885/8313/9126/9985/55869/91272/11331
=26%, signal=38%","2957/9443/9014/64216/6881/29079/64425/54700/55832/9282/9441/2958/170
0%, signal=40%","7098/2200/1435/4853/10272/1051/7130/10766/4286/7107/2353/7099/84959/10
nal=36%","8379/8697/25906/2253/56155/8452/11105/23113/9820/5300/7175/51451/54908/7517/
ags=52%, list=22%, signal=41%","677/8826/10252/90993/7057/55614/860/2246/2730/7538/64094/
ags=39%, list=24%, signal=30%","79026/2040/4208/3688/7402/23704/857/3270/477/4734/1909/15
, signal=41%","90102/389/182/6239/84962/1289/4853/83478/3678/10979/1839/387/3214/4851/7
'tags=78%, list=14%, signal=67%","55166/79003/54908/51112/283489/27085/55920/9184/91272/5
5378/7507/2072/4292/27030/5395/23186/1161/79840/2547/64421/7520/51317/2071/23028/7515
, signal=83%","64785/84296/4174/51659/4172/4175/8318/9837/4173/4171/4176"
=27%, signal=34%","51474/2059/51429/23189/395/93663/11135/2934/23500/6387/148170/283/25
, list=20%, signal=28%","3487/301/1601/2308/5166/6591/3269/493/3953/857/4837/5978/158219/2
56%","90102/3913/183/1278/10516/79875/1281/28984/3912/4629/7040/84168/4851/30008/7148
166/4853/3953/1901/1513/7043/7040/2697/4286/7107/84959/6422/1436/1499/3690/3569/1200/
5125/4208/90/367/7026/2114/26508/10252/3280/64321/2737/2735/4038/2719/4853/3217/4188/
79901/140576/4208/8826/387758/311/4780/1956/25932/3077/7057/6649/567/5066/857/3707/14
=55%, list=23%, signal=43%","26013/8085/22955/23347/353274/1017/55729/10322/5987/51548/5
07,2587,"tags=42%, list=14%, signal=36%","5654/2200/10252/90993/183/7057/493/79812/10272/8
3140/567/3133/3107/3105/3106/3122/3115/3123/972/3108/3113/3127/3134/3111/3117/3119/31
=26%, signal=38%","5327/677/196/5906/3488/329/358/7056/845/351/1393/5028/5743/1277/4327/
4%, signal=39%","301/861/2615/3077/2737/3958/6385/604/1051/8654/10875/11326/80149/3566/
","6829/64895/642843/904/23144/10914/54623/255374/170506/3297/91746/23248/23283/56254
7e-07,5661,"tags=60%, list=32%, signal=42%","7098/117854/3554/3077/567/284/10417/3133/6778/
08/5627/2040/2/3075/2934/4478/183/2022/1192/590/2162/7040/2335/715/8542/716/710/6386/3
3,"tags=52%, list=17%, signal=43%","56034/5327/84159/2152/8826/80310/4643/5154/5155/64651/
301/306/5580/6814/11027/57580/1511/6810/7305/2150/3684/5169/6688/2207/2213/3576/6850/
5/3490/3673/367/85360/7048/7026/183/2013/824/7005/1307/5069/3488/5066/3678/7056/639/5
20%, signal=47%","3673/4208/5592/80310/5154/4780/183/5155/9510/6282/3488/5159/387/80005
=40%","558/7098/3554/3624/3105/10865/8767/929/7099/80149/602/7474/11118/7805/3586/9636
=27%, signal=40%","558/7098/3554/3624/3105/10865/8767/929/7099/80149/602/7474/11118/780
list=22%, signal=30%","2059/10161/143872/90627/23189/25959/3688/57493/1435/10252/2151/4
4208/677/196/3688/4323/3956/4853/3624/8935/5580/1316/6778/604/3087/7040/4067/23529/47
5%, signal=65%","4927/53371/7175/55746/8021/55706/129401/54830/9883/100101267/10204/96
22714e-07,3564,"tags=58%, list=20%, signal=47%","5125/1910/183/1889/387/4846/1906/1359/202
nal=37%","11167/5654/2200/3280/2022/2719/4853/10272/10468/51232/4091/4851/7130/4681/5
%","367/196/7026/2908/4306/7068/4929/2099/5468/2494/5241/8013/3164/5467/7376/6256/5916
ist=18%, signal=43%","367/196/7026/2908/4306/7068/4929/2099/5468/2494/5241/8013/3164/546
3673/3688/3680/3655/3693/3678/3691/4067/5354/22801/3676/3690/9358/3684/3672/11117/367
79026/51474/7414/90102/800/51429/22837/302/301/90/7094/8826/3688/93663/1969/11135/19
l=29%","9843/302/301/2683/1956/4478/56606/493/5155/3953/10351/358/10392/6814/6548/701
7%","79003/10051/51112/55839/387103/64105/9787/64946/10592/1062/83540/10403/1063/9735
signal=31%","3572/2690/7132/3280/183/857/54741/6778/9111/3716/7040/10019/3454/4851/546

signal=51%","28998/64979/51069/9553/57129/79590/54148/55168/29074/65080/64976/51642/2
4%, signal=51%","28998/64979/51069/9553/57129/79590/54148/55168/29074/65080/64976/5164
21%, signal=35%","558/4208/301/6387/10461/3707/3958/604/4254/4067/652/7474/3586/972/529
3,3.51201094345768e-07,1.60190926010904e-07,2899,"tags=59%, list=16%, signal=50%","10775/55
9e-07,1.63960756907001e-07,4700,"tags=79%, list=26%, signal=59%","29777/6208/10799/57647/86
gnal=61%","5892/7486/3146/7528/5932/7517/132660/56852/9464/3184/57697/63967/3159/4436
4,"tags=38%, list=27%, signal=28%","79971/5125/301/64215/64285/1910/283/5872/5066/8754/597
6","6118/6117/54465/10714/83637/8467/84083/56916/6119/9031/5558/54962/253714/5425/4172
9%","10413/25937/3913/5154/2200/10252/3280/3399/64321/2737/5155/2719/116039/5159/2303
nal=35%","51429/57088/858/26509/367/2934/284612/5205/4627/30846/1289/257364/857/6711/5
ist=29%, signal=56%","246721/9477/51082/10623/548644/5442/5434/64425/55703/5436/5437/17
58%","545/54465/1325/51750/7486/546/197342/9984/80198/84083/64710/201973/55218/57697/
2%","100507436/2013/3140/567/3133/3105/3106/3122/8417/3123/5873/5027/1075/3592/2213/3
%","6788/7098/4055/344558/1956/29969/7164/10392/10114/4616/4217/351/6386/1647/8767/59
nal=41%","301/7048/861/2737/3707/6778/639/114548/387/604/1514/8546/8767/80149/3566/602
nal=30%","5587/7070/493/857/3270/477/845/3356/5350/80024/4067/51393/59341/22821/2149/1
al=53%","9477/51082/10623/548644/5442/5434/64425/55703/5436/5437/171568/9533/5439/543
3%","9477/51082/10623/548644/5442/5434/64425/55703/5436/5437/171568/9533/5439/5438/54
list=24%, signal=43%","2615/6385/1051/8654/4067/90865/11326/652/3586/51043/4332/7305/312
8%, list=19%, signal=39%","858/2152/5587/6387/283/8742/5155/1893/7010/7424/654/185/116844
37e-07,5669,"tags=51%, list=32%, signal=36%","4208/27202/1997/123920/7006/7070/3958/5359/5
gs=50%, list=26%, signal=37%","3688/183/7070/1909/154043/6548/1906/3779/5243/30845/5348/2
st=24%, signal=35%","558/7132/3656/117854/83643/1435/3554/837/857/284/9314/23586/1893/81
al=46%","1303/3685/4323/1290/3693/64321/1289/3624/3678/1295/1284/3912/2335/8091/4313/3
152/6002/78994/7528/7175/5687/9984/10296/55791/10409/862/53335/5885/30827/4670/28564
5228/26509/85461/3688/4627/30846/824/8754/928/170689/58494/8837/4773/3566/54209/975/7
gs=38%, list=22%, signal=30%","51594/23049/196513/6499/54542/4292/9924/30833/65110/11340/
st=24%, signal=46%","183/493/857/9314/947/10365/1906/5743/6648/7099/80149/199/3586/6458
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%, signal=33%","5587/7070/3270/477/309/845/3356/5350/80024/4067/22821/2149/1756/3710/11
nal=31%","7798/7049/4208/182/90/7048/57493/26508/3280/2022/4853/9510/2303/23129/639/190
ist=29%, signal=35%","135228/302/2/334/183/871/7035/1293/7980/58189/51232/351/80760/5045
"tags=40%, list=9%, signal=37%","7015/25942/57504/57521/27085/5356/55920/595/163786/994/6
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l=49%","7132/182/26508/6591/4853/4846/7040/4091/4851/23493/7075/7042/652/23462/2702/66
ignal=51%","23521/6122/6129/6159/6137/6146/6158/6152/25873/6125/6135/6160/4736/6130/11
71/5125/301/64215/64285/1910/283/5872/5066/8754/5978/6814/4053/7043/5208/7040/7424/551
/4643/5154/25932/4478/10788/4641/284/8639/7040/7010/5962/120892/4642/7412/5420/1499/3
gs=40%, list=19%, signal=33%","6788/23328/7098/4055/344558/7164/10392/4616/4217/351/6386,
1%","10203/1910/2151/2840/624/1909/6751/57085/185/2149/2984/57007/4883/2357/55366/2150
42%, list=10%, signal=38%","23476/9984/8563/252983/5514/7015/25942/57504/57521/79447/2703
=25%","253827/5445/558/22925/11030/493869/301/2308/1410/80310/4323/4780/1956/5587/824,
tags=58%, list=9%, signal=53%","25942/57504/55920/595/994/1019/8317/80174/79915/983/26271,
2%, list=15%, signal=70%","10694/10575/7015/152485/10576/7203/22948/908/65057/55135/7903
%, signal=28%","6387/1910/183/58191/2771/1901/4734/10763/624/3716/6358/5028/3691/2769/19
=26%, signal=41%","3572/301/7048/3280/3133/3105/8767/3479/7412/199/975/3569/3115/942/874
=21%, signal=51%","3685/4052/2615/7057/1281/947/2335/4060/5045/5743/7042/7305/7481/6376
%, list=24%, signal=33%","5587/7070/3270/477/845/3356/5350/80024/4067/22821/2149/1756/371
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=15%, signal=52%","367/2099/4851/9420/652/7474/4313/6422/23213/3400/5916/8626/4824/3371
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),"tags=100%, list=12%, signal=88%","157777/84296/8317/4174/51659/4172/4175/8318/4173/4171
45%, list=14%, signal=39%","7336/196528/5932/55693/55183/84501/2187/8295/57634/8089/1090
:48%","254528/2072/1677/27301/64789/10111/83940/9937/5932/80198/64421/22909/284131/75
al=40%","965/3920/5547/22918/3958/3916/387/5269/6515/2357/10288/1832/10159/1604/2207/6
423/6118/7507/6117/27301/4595/7486/51548/6996/10075/3980/8505/23583/6119/4913/55170/1
list=25%, signal=42%","5627/90102/135228/5327/5592/5154/7057/5155/7035/7056/928/5580/947
.0/10398/11346/493/9124/23022/114907/3270/9531/10979/91624/2274/103910/845/1756/81493,
t=22%, signal=51%","395/1956/283/183/5159/1909/3356/6401/2099/2769/400/185/2260/5322/72
al=35%","4208/8826/4780/1956/25932/3077/567/3707/140609/358/4502/4499/4493/1728/351/13
5%, signal=53%","8473/1386/4300/339287/55167/26122/51105/10445/55683/4298/54934/8295/57
593/7317/7703/222229/4999/23378/56916/53615/11335/23310/22880/57673/3159/3070/157570/
=67%, list=37%, signal=43%","1974/9617/85476/10605/8663/26986/9669/8661/3646/56853/8894/1
42%","2040/3920/5547/2683/84061/11010/3916/967/54918/5660/684/4257/2357/5908/55754/719
st=20%, signal=39%","1017/79184/9656/84446/29086/60561/51499/197342/80198/65975/5514/58
%","2926/6733/58155/56853/57703/3167/4116/4928/2332/10521/6231/6625/283375/10772/5569
list=21%, signal=40%","7098/301/2200/861/2737/3624/10272/604/55365/7130/10766/3204/4067/
%","9851/84446/221908/116840/2648/9662/25914/55125/7514/7517/284001/9738/79959/7840/1
list=28%, signal=32%","10203/1910/2151/2840/624/1909/6751/57085/185/2149/57007/4883/2357
list=32%, signal=42%","196/100507436/3554/3077/3140/567/3133/6778/3916/114548/604/7040/3
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-72e-07,4028,"tags=80%, list=22%, signal=62%","567/1514/5641/54209/3122/3115/3123/972/3108/
49%, list=22%, signal=38%","196513/54542/4292/9924/11340/170506/6434/3297/22894/246744/27
0413/10404/301/7048/861/7006/688/928/8837/6442/347/4851/2697/3479/5653/84525/8324/4815
5%","558/4055/7048/4853/1051/55365/4067/28959/3106/54209/5971/2150/10288/55509/2621/6
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","858/26509/7048/30846/493/4038/857/7035/477/4846/3356/161742/6401/5743/120892/1495/5
2%","966/1601/11015/1956/51226/567/8322/928/9341/3133/3107/1839/3105/8546/89866/3106/
%, list=25%, signal=39%","1655/57060/8087/51574/55234/6433/6428/6431/7072/55225/6434/6068
30/1910/183/3488/23268/5159/1284/2730/1909/1728/7043/3356/5028/347/1906/710/3779/5743,
gs=73%, list=15%, signal=62%","10694/10575/7015/222229/152485/10576/7203/22948/908/65057/
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%, list=25%, signal=43%","80310/6387/3958/387/58494/351/245937/5641/5054/199/7474/3676/26
"5558/11232/5426/54107/5425/4172/5427/5557/23649/5424/5422"
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5,"tags=32%, list=9%, signal=29%","5885/835/144455/29117/80279/91603/84861/8883/29980/593
, signal=30%","686/8801/2690/9469/56548/4780/1910/2687/5166/9645/284/84836/1192/23743/2
signal=51%","6387/4478/1893/387/58494/351/347/80024/23143/199/7474/3676/6347/29108/876
44%, list=24%, signal=33%","5587/7070/3270/477/309/845/3356/5350/80024/4067/22821/2149/17
gnal=37%","7098/301/2200/861/2737/3624/10272/604/55365/7130/10766/3204/4067/28959/108
4/302/10351/857/10577/6342/4481/9388/3990/54762/5045/5360/55207/19/26119/5244/4790/106
ignal=38%","23347/9025/7334/26122/1054/79184/29086/3146/7336/196528/165918/5932/55693/
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=19%, signal=51%","22823/26147/22938/10445/94104/56254/55683/7936/57343/23476/55183/80
07,4009,"tags=55%, list=22%, signal=43%","10161/3685/1435/2151/4853/3707/5159/1281/2840/425
gnal=27%","558/2690/2/3673/25959/2308/3920/7048/4780/5587/3077/5166/6591/355/2771/2730

list=17%, signal=39%","23328/7098/1956/7205/4306/10392/1896/9111/3987/387/351/1906/929/8
%, list=23%, signal=42%","135228/26509/85461/3688/4627/30846/824/8754/928/170689/58494/8
26509/85461/3688/4627/30846/824/8754/928/170689/58494/8837/4773/3566/54209/975/7305/1
gnal=47%","4208/90/116039/7043/7040/10766/654/7020/7042/652/7474/2201/9636/5027/9365/5
","339287/55167/51147/26122/55011/7994/55677/6879/10445/2648/55683/11143/54934/8295/5
27e-07,5669,"tags=52%, list=32%, signal=36%","4208/27202/1997/123920/7006/7070/3958/5359/55
2%, list=32%, signal=36%","4208/27202/1997/123920/7006/7070/3958/5359/5580/3105/4067/110
35/3688/7057/3488/10272/6385/1513/7130/1514/151887/4313/3676/3690/59339/3489/1520/367
gnal=71%","5892/7486/3146/7528/7517/57697/4436/348654/5889/4221/7516/3148/641/2956/314
al=30%","79971/10413/56034/5125/3572/7048/1956/3280/1910/64321/2737/2735/2300/688/128
signal=33%","23189/301/2152/3688/4780/6239/4638/7057/7056/2303/10979/1839/64094/6484/6
al=41%","9617/85476/10605/646799/8663/26986/9669/1153/8661/3646/56853/8894/1984/10767
46%","90/7048/26508/2022/6591/2303/4846/4851/23493/59/7042/7474/51176/653/23462/4487/7
=56%","5619/9131/1677/10595/7273/26993/8467/79648/10051/22985/9031/23310/8815/10592/29
%","4208/7132/182/90/7048/26508/6591/4853/639/4846/7040/4091/4851/23493/7075/56999/704
%","6881/51720/57602/56943/171023/79184/9960/6878/55578/56970/6879/6873/27005/2648/117
=21%, signal=29%","10413/64114/6788/3685/8531/90/259230/367/4055/4000/8406/4780/6672/63
08/4306/857/477/10577/6342/309/5375/347/2099/54762/5241/19/6646/1880/975/706/950/9365
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5,"tags=53%, list=13%, signal=46%","25937/1601/182/90/7048/2022/10979/28984/7043/7040/7041
st=22%, signal=38%","90102/389/395/6281/3693/84952/9124/1901/10174/6385/10979/28984/407
'90102/389/395/6281/3693/84952/9124/1901/10174/6385/10979/28984/4070/387/6548/7043/22
tags=50%, list=26%, signal=37%","183/7070/1909/154043/6548/1906/3779/30845/5348/2149/1756
l=22%","79971/5125/4208/80114/90/367/7048/7026/2114/23500/26508/10252/3280/2022/64321/
=45%","965/3685/3688/7070/3678/9314/91663/7412/3586/3676/3690/1824/10288/1832/4155/36
%, list=30%, signal=37%","3597/2059/3572/3685/3673/90/3688/7048/3680/9180/10203/3655/3693
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tags=37%, list=17%, signal=31%","7098/117854/3554/3077/567/284/10417/3133/6778/23586/1145
gnal=47%","4208/90/3280/1910/2022/5155/1909/9096/4851/79652/93649/255743/30008/23493/4
%","558/3685/3673/3688/1969/1956/3693/3678/5359/3916/3356/4360/9076/1605/975/3690/942
02/800/51429/9936/389/117583/10398/8826/4739/4651/9322/4643/2934/5587/7402/83872/5747
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6%, signal=35%","54984/27000/10694/100134934/10575/11143/5609/7520/3183/54921/64785/67
ist=26%, signal=37%","6387/4638/7070/5155/857/3958/5159/6548/11027/51393/30845/2149/535
al=31%","7798/7049/4208/182/90/7048/57493/26508/3280/2022/2303/639/1909/4846/7040/409
=43%","8841/55818/64431/57649/25855/10428/10847/3066/23186/81611/57459/84312/64426/30
23%, signal=36%","90806/51594/196513/54542/4292/9924/11340/66037/170506/207/22894/60493
, list=22%, signal=56%","395/1956/283/183/5159/1909/3356/6401/2099/2769/2260/728/64805/94
l=34%","1601/1956/8322/928/9341/1839/8546/26119/7474/3122/348/3115/5868/3123/10618/97
gs=76%, list=28%, signal=54%","301/7048/861/2737/3707/114548/387/8546/8767/3566/51043/312
159/83716/6237/64651/7043/444/1277/25976/4313/51176/4487/7122/22943/5083/6497/83690"
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st=20%, signal=42%","831/10413/22925/677/2934/4780/7057/284/28984/7538/8837/7010/641700
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=20%, signal=25%","10413/3556/3572/23500/4124/26508/81565/3280/6387/2737/7070/7058/4038
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%, list=18%, signal=44%","7049/11030/1601/90/7048/3077/2022/56937/3624/7043/7040/4091/638
59/83716/6237/64651/1909/7043/444/1277/220/25976/5604/7474/4313/51176/4487/5916/7122/
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=15%, signal=52%","10445/55683/7936/54934/4089/23168/51230/4602/84148/55170/54880/2840
46%","23166/3426/58191/4481/51435/5167/8578/9332/57007/84466/950/84171/4035/4017/8443
tags=42%, list=22%, signal=33%","683/2308/1410/5547/7048/4780/183/7057/10516/5155/329/277
18%, signal=61%","135228/7049/3685/90/4052/7048/2615/2022/7057/7043/8425/94/57045/3753
ist=22%, signal=28%","51594/23049/196513/6499/54542/4292/9924/30833/65110/11340/23708/5
=13%, signal=36%","56034/80310/5154/2022/5155/857/9961/7424/5167/11261/54206/5912/1045
3%, list=24%, signal=41%","23381/545/6117/2072/170506/8658/9894/51750/54984/10111/10694/1
7/3133/3107/7040/3105/11027/6892/3106/6890/958/6891/3676/3122/3115/3123/3113/55080/11
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nal=30%","11015/2934/5587/6672/7070/857/3270/477/309/845/3356/5350/80024/4067/23345/22
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39%","558/117854/2934/3956/8724/10581/10410/9023/5359/10475/64135/4851/56829/8091/816
%","558/3572/2690/4208/2632/1435/8301/824/284/5978/2730/10763/387/4217/947/1728/7043/4
, signal=43%","23381/545/23049/2072/170506/8658/51750/546/54984/10111/51548/10694/10013
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=35%","196513/54542/4292/9924/11340/66037/170506/207/22894/60493/27327/340719/10642/4
%, signal=55%","9469/56548/1634/633/337876/29940/133584/22856/83539/79586/55454/2131/50
=49%","367/2099/4851/9420/652/7474/4313/6422/23213/3400/5916/8626/4824/3371/6662/3169
50%, list=17%, signal=42%","10413/25937/3913/4208/182/8826/4643/3280/1910/2737/5155/4853/
8%","5587/5872/7057/2246/64094/4851/2260/5641/3791/3164/3315/1960/9037/8823/8829/7114
l=51%","311/567/3122/975/3115/3123/972/3108/3113/10288/55080/3127/3111/3117/3119/920/3
%, list=15%, signal=60%","22938/10445/55683/7936/54934/4089/23168/51230/4602/84148/284058
list=24%, signal=51%","3685/1435/183/4023/4481/7040/5468/185/19/5583/6646/5322/4792/4790
0%, signal=34%","1601/5321/1910/54884/2052/11343/5978/151056/27122/6342/216/157506/5742
%","5327/5125/2152/5654/5547/4323/11098/3426/1513/715/57552/716/5045/5328/4327/94031/
signal=27%","9445/9469/56548/2683/4124/81031/84061/4121/285203/2300/25825/117248/1839/
%","947/3082/7097/7099/602/9636/54209/7305/3569/85480/3123/64581/2150/28514/29108/729
9%, signal=41%","947/3082/7097/7099/602/9636/54209/7305/3569/85480/3123/64581/2150/2851
signal=36%","1601/2308/10252/1848/183/7078/857/1847/5580/10114/154043/1843/161742/100
,"tags=53%, list=26%, signal=40%","83860/2957/9443/6881/29079/55832/9282/9441/2958/170506/
5%","51474/3953/9023/10577/158219/2246/351/8542/3990/19/6646/8578/51302/9420/26119/23
ist=31%, signal=45%","301/861/6778/114548/604/1514/8767/80149/3566/602/1880/51176/51043/
/4055/344558/1956/29969/148170/9459/7164/10392/10114/4616/4217/351/6386/1647/51135/12
list=29%, signal=48%","301/7048/861/2737/3707/639/114548/387/604/8546/8767/80149/3566/51
%, list=21%, signal=42%","7098/857/329/2099/4067/11027/7096/338382/7097/3430/3659/54209/7
01/306/5580/6814/11027/57580/1511/7305/2150/3684/6688/2207/2213/3576/6850/10758/5724/5
gnal=26%","7098/3673/4208/196/7048/5154/4780/1848/2022/7057/3624/7079/5066/4837/2052/5
ist=10%, signal=32%","23189/301/2152/3688/4780/6239/81565/4638/7057/7056/2303/10979/1839
57%, list=27%, signal=42%","301/6778/114548/604/8767/10875/80149/3566/602/1880/51176/5104
26%, signal=39%","301/7048/861/6778/114548/604/1514/8767/80149/3566/602/1880/51176/5104
74/857/10226/158219/4023/4481/5468/8605/5167/19/6646/80149/4792/4790/2629/54209/1066/3

gs=56%, list=25%, signal=43%","3140/567/3133/3916/3105/3106/135250/3122/7305/8417/3123/21
=33%, list=13%, signal=29%","64114/2239/5525/1601/367/3688/1969/4323/1956/8301/5906/493/2(
, signal=29%","858/4208/8531/90/7026/26508/6840/688/857/30812/79784/387/10791/8837/7040
al=71%","11143/5001/4999/5000/4998/1660/4174/1663/8318/55388/990/4171/5422"
%","10252/3280/183/2737/2719/3217/6385/2303/30812/2246/4070/4091/255743/2297/2668/652
6,3260,"tags=55%, list=18%, signal=45%","11030/1601/90/7048/3077/2022/56937/3624/7043/7040
57%, list=29%, signal=48%","196/100507436/3077/3133/604/3105/3106/684/56940/84868/1604/66
, signal=47%","117854/2934/3956/8724/10581/10410/9023/10475/81603/6737/10346/85363/851
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,1167,"tags=55%, list=7%, signal=51%","4208/5592/4780/183/5155/9510/3488/387/80005/93649/8(
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3%, signal=33%","2/117854/3075/100507436/4780/329/5359/3133/9111/3916/81844/3105/51752,
18%, signal=46%","57647/55308/6499/170506/8449/51428/22907/55794/11218/9879/79009/8886
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gnal=33%","6788/7098/4055/344558/7164/10392/4616/4217/351/6386/1647/8767/59341/153769
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72,"tags=76%, list=19%, signal=61%","3077/567/3133/3107/3105/51752/6892/3106/6890/6891/312
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%, signal=27%","11031/64114/2239/5525/1601/7132/367/6281/3688/1969/4323/196383/1956/830
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899,"tags=51%, list=11%, signal=46%","90627/4208/7057/493/79812/9314/28984/387/161742/704(
, list=11%, signal=37%","90102/23189/395/182/4323/7057/8322/4070/387/604/347/4851/81848/32
/3688/4651/2683/7402/4478/10788/10519/351/84168/4254/5962/2195/7408/59/59341/7412/175
%, list=17%, signal=46%","114548/3479/80149/338557/3586/54209/1992/54206/114769/6376/7128
t=28%, signal=46%","51497/6829/7919/904/284695/54623/64425/51574/8621/9025/56943/51755/
t=23%, signal=37%","26013/8085/22955/23347/3720/353274/1017/23512/64783/55729/10336/917
2e-06,4021,"tags=82%, list=22%, signal=64%","3077/3140/567/3133/3107/3105/51752/6892/3106/6
=12%, signal=28%","80198/51451/65975/54908/7517/5514/5883/5885/835/144455/29117/7976/8(
9/84962/1289/83478/3678/10979/1839/387/2316/94/4488/10630/977/5728/23671"
1%","4208/182/90/7048/26508/6591/4853/4846/7040/4091/4851/23493/7075/56999/7042/652/2
2%","3956/4853/112616/9452/602/1880/3586/3569/28514/6688/3955/2213/4242/5336/6482/339
7414/389/1969/5872/6591/4641/9948/387/90952/5175/1003/9076/1495/5583/79778/137075/48(
=25%, signal=31%","3572/2690/7132/3280/183/857/54741/6778/9111/3716/10019/3454/4851/164
38/4811/64129/3956/7057/3958/4059/8754/1893/1605/59339/1520/3679/3675/27076/59277/284
=20%, signal=48%","4208/90/116039/7043/7040/654/7020/652/2201/9636/5027/9365/56975/1304
23%, signal=33%","2152/2934/8301/55075/83595/355/857/5978/10392/114548/387/4217/1611/88
3400,"tags=53%, list=19%, signal=43%","558/5587/79812/3678/8322/2303/4734/161198/3087/7424
0%","5784/1969/7026/57493/2303/161198/79987/7424/79652/7075/3791/94/2296/54345/10630/
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al=34%","4643/122786/2934/6444/2013/23704/4627/4641/857/55930/477/79784/6548/6324/5350
21%, signal=50%","3426/715/716/710/54209/729/735/4153/730/712/713/1191/718/721/51279/160
l=34%","683/4208/677/196/4323/3624/6778/604/7040/4067/23529/4773/7099/1880/6422/7805/9
3061271e-06,5521,"tags=65%, list=31%, signal=45%","301/6778/114548/604/8767/80149/3566/602/
gs=54%, list=24%, signal=41%","2615/6385/1051/8654/4067/11326/652/3586/51043/4332/7305/312
0%, signal=34%","996/79176/26260/1027/8454/8816/90864/51433/26094/26133/342897/59349/54
517,"tags=53%, list=25%, signal=40%","7098/857/329/55601/2099/4067/11027/7096/338382/51296
4%","54885/1793/143872/90627/395/8826/57514/93663/57480/85360/141/9459/23348/7070/590

, list=20%, signal=24%","54885/1793/143872/90627/395/8826/57514/93663/57480/85360/141/945
%","55214/5836/5351/2022/5066/1203/4360/80216/8974/6403/6515/282969/5033/283208/80201/
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","23381/6118/6117/23049/65110/9894/7486/10111/100134934/4691/2547/9887/7520/7015/3178
27%","2690/2/3673/25959/2308/3920/7048/4780/5587/3077/5166/6591/355/2771/2730/4217/753
gs=35%, list=13%, signal=31%","64114/5525/1601/367/3688/1969/4323/1956/8301/493/2042/6711
gnal=26%","79971/11031/51429/5125/7045/9469/259230/10981/56548/2683/4124/5587/5205/587
5%, signal=47%","51535/57343/80312/23378/23133/9869/53615/22880/23028/57673/3070/90378
%, list=28%, signal=39%","301/1601/83643/824/5580/2246/157506/23682/3356/654/5743/11261/34
gnal=46%","301/5321/3554/6778/10865/8767/7099/3566/602/51176/3123/5971/9655/7292/84868
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5%, signal=51%","3673/3688/861/90993/2022/7431/5159/28984/7043/7040/1471/2191/2149/652/
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%","858/26509/4208/85461/3688/4323/4627/30846/824/688/928/170689/58494/79365/8837/4851
t=13%, signal=43%","51429/152503/4627/7078/25825/257364/7079/8754/51752/5045/7077/7076/
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signal=42%","22925/302/1601/5321/5580/6342/9388/1906/654/114897/51703/26119/5244/348/7
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/183/196740/624/4846/10365/3779/6648/4883/348/5467/1373/2150/2702/3672/8985/2131/153/
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signal=40%","3269/857/387/3356/6915/5743/2149/146/54795/6916/2702/552/10800/6869/147"
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38%","7049/4208/182/7048/26508/3280/2022/64321/2303/23129/1909/4091/10512/4851/1906/23
gnal=24%","10413/4208/90/1969/8463/7048/4323/2200/3280/2022/64321/2737/2735/4853/3217/8
54%, list=29%, signal=46%","558/301/7048/4323/861/2737/3707/9580/114548/387/604/8546/8767
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=33%","5327/5125/2152/5654/5547/4323/11098/3426/1513/715/716/5045/5328/4327/94031/7092
=29%, signal=47%","301/7048/861/2737/3707/9580/114548/387/604/8546/8767/8808/3566/51176
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%","7132/182/26508/6591/4846/7040/4091/4851/23493/7075/652/23462/6615/1942/7133/11117/
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4%","56034/2308/3910/1910/63976/3953/857/6778/2303/216/3356/1051/8878/4851/2697/5743/
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, signal=33%","22925/3280/23704/857/7779/477/4734/4846/845/28954/3356/5350/2669/5743/40
al=41%","3140/567/3133/3916/3105/3106/135250/3122/7305/8417/3123/64581/2150/5027/3684/
%, signal=43%","196/3554/3077/3140/567/3133/114548/3105/10865/3106/54795/3122/975/3569/
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:ags=48%, list=20%, signal=39%","677/861/1435/4853/146433/7040/4254/2353/51176/54209/3398,
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ıl=55%","7132/182/26508/4853/4846/4091/4851/23493/7042/652/23462/2702/7133"
l=38%","831/10413/6788/22925/677/7048/2934/4780/7057/284/28984/1909/4217/7538/8837/701
,3260,"tags=36%, list=18%, signal=30%","6788/56034/23328/2690/8826/80310/5154/1956/84962/7
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%","2690/90/10203/1910/1889/3953/3624/1393/2984/4883/4881/552/1240/2587/5618/10148/30
4323/1435/1910/7057/79148/3958/146433/928/59341/5919/728/1436/64805/54209/975/6347/63
gs=54%, list=22%, signal=42%","389/6281/11346/10174/6385/10979/28984/387/7043/22899/7984/
3673/4811/3075/7057/1289/10468/50509/1513/4023/2335/1514/6386/1508/348/10457/4035/337
9142e-06,3440,"tags=65%, list=19%, signal=52%","183/5155/9510/3488/2771/4217/2697/1906/420
/56257/1984/6625/139804/55696/27288/7919/904/27258/51574/114034/55285/10594/905/7906
2/7132/2308/29995/2934/4000/183/90523/493/3488/114907/4846/6548/8837/6442/4851/1906/8
%, list=26%, signal=35%","64114/2040/2239/5525/1601/367/3688/1969/4323/2934/1956/8301/590
%, signal=57%","5547/2683/4162/2303/947/64094/5054/3791/1605/3690/6376/7128/5291/240/28
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signal=30%","6907/4831/10987/9443/5585/58508/55758/9500/64216/440498/51132/9139/2023/
4,"tags=58%, list=25%, signal=44%","1793/23189/1601/6281/4739/6239/10519/55679/3987/10979/
=20%, signal=38%","1017/79184/9656/84446/29086/60561/51499/197342/80198/65975/5514/588
%, list=23%, signal=30%","79971/10413/25937/6788/26524/23189/3487/1601/80114/2308/57664/2
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%, list=22%, signal=38%","90102/389/395/6281/84952/1901/10174/6385/10979/28984/4070/387/6
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signal=34%","51474/2059/51429/23189/93663/11135/2934/23500/6387/148170/25802/6840/6711
tags=54%, list=20%, signal=44%","7098/2200/3624/10272/7130/10766/3204/4067/7107/7099/8495
al=37%","5327/677/196/5906/3488/329/358/7056/351/1393/5028/5743/1277/4327/3309/2353/51
l=39%","64114/861/183/5166/3953/1901/7040/2697/1471/5468/7107/84959/6422/1436/3690/356
signal=42%","117854/2934/8724/10581/10410/9023/5359/7538/64135/56829/8091/81603/3428/6
8%","2040/558/117854/2934/3956/29969/8724/10581/10410/9023/5359/10475/7538/64135/4851
%, list=25%, signal=56%","3688/928/4023/5468/19/7099/54209/1066/975/348/9655/10333/2207/39
ıal=31%","3572/7132/2308/29995/4000/183/90523/493/3488/114907/4846/6548/6442/4851/1906
=26%","79971/2040/5784/64215/161291/64285/5587/29969/6672/81565/283/8724/2737/4641/50
'302/3074/5611/5580/10577/2720/8804/2519/6386/10487/175/1511/5476/1991/4332/1727/29108
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0312/6996/200424/54496/54784/56154/23583/53615/4913/9760/51010/7874/10419/56915/1789
869/85459/80254/56890/26005/4591/1017/9851/221908/116840/2648/9662/25914/55125/28400
ignal=54%","64328/3842/26953/64901/23039/30000/55374/3843/8021/7514/79711/55705/56478/
=25%, signal=45%","6281/1435/7402/2013/6237/7070/54453/1307/10519/27185/6385/3987/1097
'tags=45%, list=9%, signal=41%","3685/1435/183/4023/4481/5468/185/19/5583/5322/4792/4790/3
'tags=61%, list=13%, signal=53%","51429/152503/7078/257364/7079/8754/5045/7077/7076/3586/
=44%, list=26%, signal=33%","831/1410/2152/2934/55075/7057/83595/355/329/5978/358/10392/9
list=20%, signal=54%","7098/2200/10272/7130/10766/7107/7099/84959/10234/6422/1499/9935/3
:13%, signal=34%","26073/23476/6198/1977/5514/7015/25942/57504/284001/23326/440/57521/2
9147/79693/6741/51095/60487/55152/348180/129450/51067/4234/9255/6165/54938/124454/839
4034/90806/10208/6606/1017/2483/96764/55341/9129/9879/54433/345630/50628/51602/7514/

gnal=33%","1601/1910/54884/1889/5978/151056/27122/216/157506/2099/654/6121/220/25976/9
","55214/7026/5836/5351/8309/5066/6342/4846/5375/6401/5660/8974/5468/2350/2173/6403/3
","57647/55308/6499/170506/8449/51428/22907/55794/11218/9879/79009/8886/64794/3508/971
%, list=6%, signal=38%","143872/1910/183/387/1909/3356/151/80005/1906/59"
3%, signal=36%","143872/3673/5592/10203/857/477/387/151/80005/1906/93649/255743/6915/57
gs=46%, list=27%, signal=34%","51474/2059/51429/23189/93663/11135/2934/23500/6387/148170/
signal=47%","683/27202/1435/7057/146433/7130/1906/59341/5919/728/1436/23604/972/57402/
9%, signal=38%","558/677/301/7048/4323/861/2737/3624/3707/9580/639/114548/387/604/8546/8
8281291911209e-06,3026,"tags=88%, list=17%, signal=73%","3077/567/3133/3107/3105/51752/689
l=37%","302/3673/301/3688/11135/6282/9124/23022/4059/91624/1266/90952/27020/22801/5453
nal=29%","51429/302/1601/2152/6281/4323/2934/344558/152503/4780/8301/55075/4627/28567
l=41%","79971/5154/1956/6591/1896/23129/7043/3082/7040/10512/7042/7052/1605/2252/2296/
,3823,"tags=67%, list=21%, signal=53%","10556/201164/10775/55520/131870/27161/79897/51163/
4%, list=24%, signal=34%","558/7132/3656/117854/83643/1435/3554/837/857/284/1893/81844/10
4%","182/26508/6591/4846/7040/4091/4851/23493/7075/652/23462/6615/1942/11117/54567/66
47%","5154/1956/6591/1896/23129/7043/3082/7040/10512/7042/7052/1605/2252/57045/4824/8
301/8826/57477/25802/9124/4641/23022/8522/7010/10634/1730/7171/7456/199/7169/5756/27
%, signal=31%","558/301/1410/80310/4780/1956/824/6649/10516/8754/358/5159/5580/6778/388/
gnal=39%","83860/548644/2957/5434/6881/54623/728340/2958/4331/2068/6878/5436/5437/6879
5125/93664/4208/301/1601/5321/6387/1910/3077/183/5906/3624/8322/5978/7779/6814/3356/5
ignal=49%","1910/183/2013/5155/2771/1906/185/2149/2150/2702/4881/2621/552/10391/6869/1
=51%","7048/3133/3105/8767/3659/2000/51043/975/7292/57045/29126/8764/1604/3965/1794/1
64272e-06,2937,"tags=68%, list=16%, signal=57%","4208/5592/387/7043/93649/30008/5468/6648/
al=27%","5321/5154/283/84962/5337/824/126969/5155/8871/857/3707/11343/5359/5159/15105
","7049/57493/2022/9510/1901/387/2274/7010/4851/1277/23493/7042/4313/6422/2201/653/234
%, signal=47%","22925/3685/7057/158219/5468/19/4792/4790/54209/1066/3690/3569/5467/7376/
gnal=46%","10111/5932/80198/64421/22909/284131/7517/84083/1773/7515/55345/348654/1775
gs=62%, list=16%, signal=52%","546/54984/10694/10575/7520/7015/152485/10576/7203/22948/90
336/10291/6634/6627/51634/10992/6637/8175/6629/6633/6635/84844/6628/6632/23450"
%","11031/1793/3673/4240/2934/4627/7057/4481/51454/19/391/199/54209/64581/2150/94134/8
"56034/80310/5154/1956/183/5155/2355/358/5159/1281/2335/2099/6277/26471/3479/8877/664
%, list=15%, signal=44%","947/3082/7097/7099/602/9636/54209/3569/85480/64581/2150/29108/7
%, signal=39%","23353/254528/57820/4292/27030/317761/9134/10734/11144/51547/256006/443
%, signal=51%","311/567/3122/975/3115/3123/972/3108/3113/3127/3111/3117/3119/920/3109"
al=73%","567/6892/3122/2923/3115/3123/3108/3113/55080/3127/3111/3117/3119/3109/811/312
gs=94%, list=22%, signal=73%","567/6892/3122/2923/3115/3123/3108/3113/55080/3127/3111/3117
","51147/56943/6885/55011/7994/6878/56970/27154/6879/6873/8888/8861/2648/6871/11143/13
96513/54542/4292/9924/11340/170506/22894/27327/340719/10642/4904/5436/79066/27257/129
gs=48%, list=24%, signal=37%","3554/3140/567/3133/6778/3916/23586/114548/7040/3105/10865/
2239/3685/3688/2022/23643/10082/4919/57007/3676/3690/4804/942/1464/8829/920/10268/843
2%, list=7%, signal=49%","4208/5592/4780/183/5155/9510/3488/387/80005/93649/8013"
%","51750/9361/79840/3146/55226/65057/11232/79035/2177/55215/81620/5111/5888/6597"
57%, list=26%, signal=42%","7098/3554/3105/10865/8767/929/7099/602/7474/11118/9636/3115/6
ags=57%, list=18%, signal=47%","3956/4853/9452/602/1880/3586/3569/28514/6688/3955/2213/42
677/3688/4323/3956/4853/3624/1316/604/3087/23529/7412/112616/9452/602/1880/6422/3586/
","11031/51429/1793/3673/4240/2934/8724/4627/7057/257364/387/4481/51454/19/391/199/542
st=22%, signal=50%","7414/7132/182/1903/4851/654/1003/652/3397/1499/81575/694/84913/94/2
signal=46%","56853/28998/79727/23521/5914/54464/7248/6188/2332/55571/79068/26065/13140
al=52%","4323/1435/7057/79148/146433/928/59341/5919/728/1436/64805/54209/975/6376/719/
, list=24%, signal=41%","2615/6385/1051/8654/4067/11326/652/3586/51043/4332/7305/3123/942

ignal=28%","3673/183/23704/493/857/3270/1901/477/79784/845/6548/9096/6324/5350/2697/30
0%, list=23%, signal=39%","7414/90102/395/6281/4323/7070/7057/9748/2042/1307/79834/6385/3
tags=45%, list=18%, signal=37%","558/1969/1956/2050/10461/2042/5159/51232/7010/7075/2260/
9%, signal=36%","3490/1601/6591/5978/2246/27122/654/50846/27120/1958/4790/653/1066/8644
al=35%","90806/196513/54542/4292/9924/11340/66037/170506/207/22894/60493/27327/340719,
%, signal=49%","5442/10922/64216/54938/124454/64863/80298/60493/54931/130916/81892/790
gnal=47%","3913/3280/5155/5159/30812/947/2297/2668/1958/652/1499/6943/55366/6299/13049
340/1519/5327/5125/2152/5654/4323/11098/81565/3426/824/29761/5069/79148/1889/9510/258
al=41%","26013/8085/22955/23347/353274/1017/55729/3066/51548/51535/7175/57343/55183/8
0/10398/11346/493/9124/23022/114907/3270/9531/91624/2274/103910/845/1756/81493/9457/2
624/4846/7130/5743/4327/5241/3566/4313/8204/3625/4609/2069/3248"
","1910/183/23704/857/3270/1909/845/6324/5350/1906/23493/6330/146/1756/2034/3752/54795,
2%","2934/6444/23704/857/3270/477/79784/845/6548/6324/5350/2697/8654/59341/30845/6330
727/6188/2332/6231/1996/6152/25873/6135/6160/26523/4736/6130/6235/689/6228/6191/6170/6
%","7175/26073/440145/79598/55125/91782/285643/56647/9126/11200/3192/4751/8086/9493/9
5, signal=31%","79026/2040/4208/3688/23704/857/3270/1909/845/28954/351/1393/5350/2669/19
","6234/64968/6233/6232/6188/6231/28973/54505/6204/6229/55168/6183/63931/60488/6235/62
nal=55%","25917/1787/4234/115708/152992/131965/6894/54931/54482/158234/80745/55039/22
list=22%, signal=50%","1601/183/2246/654/185/26119/1066/348/6720/1081/1581/5562/9572/335,
%, signal=36%","3490/1601/183/6591/5978/158219/2246/27122/654/50846/185/27120/26119/195
"tags=64%, list=31%, signal=45%","301/6778/114548/604/8767/80149/3566/602/1880/51176/5104
521,"tags=64%, list=31%, signal=45%","301/6778/114548/604/8767/80149/3566/602/1880/51176/
=25%, signal=50%","3572/5592/928/5580/3716/10019/5962/4067/114897/84959/1511/3569/10630
%, list=17%, signal=35%","3572/5321/3554/3140/567/3133/6778/114548/7040/3105/10865/23529/
5","10413/3490/8463/5154/2042/8322/358/5159/9314/3199/654/4067/1277/19/7075/8324/8273/7
signal=37%","3572/23500/3280/146433/443/9948/7040/5803/4851/23529/7097/1600/54209/1499
gnal=45%","182/90/7048/26508/2022/6591/4846/4851/23493/2065/7042/652/10370/653/23462/4
"6188/2332/1982/8664/1979/54505/728689/51441/4869/27335/3692/1964/10987/5499/8666/621
474/8309/857/10577/158219/727910/4023/7851/63874/9388/3990/29116/57333/5468/2494/19/6
2/6002/7528/5687/9984/10296/55791/10409/862/53335/5885/30827/4670/285643/4602/51588/8
,"2059/23189/3685/7094/3688/1969/1956/7070/4641/6711/3678/152273/10519/23129/10979/23
%","7048/85301/1893/309/2121/5916/54796/4488/2131/5915/6662/2261/6781/4921/25987/1026
56943/6885/6878/55578/56970/6879/6873/51616/2648/6871/117143/9913/8295/23326/10629/68
ist=27%, signal=43%","84991/84530/11171/1655/8087/55234/8621/6433/6431/7072/55225/6434/5
, signal=44%","23248/57466/84337/27161/23476/23160/58506/55250/123169/23168/55197/6827/
,"3685/1435/183/4023/4481/7040/5468/185/19/5583/6646/5322/4792/4790/3690/7376/8435/338
%, signal=49%","10594/22938/6634/6627/9716/6637/6629/6633/51645/5356/1659/6635/56949/27
37%","7049/182/90/7048/26508/3280/2022/4853/4846/4091/10512/4851/23493/7042/8321/9249
%","1465/858/26509/10398/3688/4000/6444/3280/284612/183/2022/25802/5159/9948/1909/4629
, signal=32%","79971/283/8754/7043/5208/7040/7424/55107/654/5468/3479/6934/7042/7097/27
3869/4780/6649/10516/7779/216/4502/4846/4499/4493/1728/5742/5743/4258/5243/2878/84203,
nal=51%","51639/55285/4904/6634/10992/6637/6633/10081/11066/6635/51690/84844/10262/662
26%, signal=35%","857/3270/477/845/6548/5350/8654/59341/30845/1756/80149/1193/54795/316
3/5154/1956/4038/3624/3488/10468/1896/9079/4851/5743/7042/26018/7474/5187/60485/1499/8
54/1956/4038/3624/3488/10468/1896/9079/4851/5743/7042/26018/7474/5187/60485/1499/8323
6231/1996/6152/25873/6135/6160/4736/6130/6235/689/6228/6170/6156/6141/6142/8667/6132/
059/79901/9429/4643/2683/4627/4641/1192/358/151056/79784/103910/2028/4642/3710/2317/1
"tags=47%, list=24%, signal=35%","25937/6788/26524/3487/1601/80114/2308/64321/2737/6591/2
57159/126731/10844/5605/3796/23332/25876/374654/23093/51361/3925/79848/51181/1453/497
signal=33%","135228/966/683/2239/4907/7070/2719/50863/7035/10392/1368/2350/10082/5129

413/7048/10252/157506/8546/5604/3202/652/1499/1605/6943/2252/1522/8985/10253/2131/666
)5,1131,"tags=69%, list=6%, signal=65%","55920/3192/4751/10615/1894/891/9212/90417/29127"
%, signal=48%","4208/90/116039/7043/7040/1051/654/7020/652/2201/9636/5027/9365/56975/13
"25937/3280/183/2737/2719/3217/5159/30812/2246/4070/255743/120892/2297/2668/51301/652
.6%","5154/1956/6591/1896/23129/7043/3082/7040/10512/7042/7052/1605/2252/57045/4824/88
, signal=27%","558/22925/493869/301/2308/80310/4780/1956/5587/6649/10516/5978/358/5580/
79971/3673/90/3688/1969/3624/2303/9314/3691/2697/8091/6886/652/7474/51176/3690/7481/2
l=42%","3673/3688/3624/2303/9314/3691/2697/8091/6886/652/3690/2139/22943/3675/2294"
,"64114/558/4208/182/183/2303/1909/4846/7040/10512/1471/84870/3200/2034/2316/94/2296/2
)4/6634/6637/10421/6633/25980/6635/10907/23020/6628/24148/9416/6632"
tags=44%, list=17%, signal=37%","558/27202/947/3479/7099/80149/602/3586/54209/54206/11213
:44%, list=20%, signal=36%","2/196/5654/100507436/3133/10475/9111/3105/710/5468/3428/1102
.7/8602/55720/55813/26747/55127/131870/54433/51602/9790/7520/10885/9221/10607/10528/5
","135228/5154/1956/4038/3624/3488/10468/1896/9079/4851/8546/7042/7474/60485/1499/832
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%","5445/9429/493869/196/4780/6649/10516/2052/5159/7779/5446/216/4502/4846/4499/4493/1
al=45%","11031/51429/1793/3673/4240/2934/4627/7057/257364/387/4481/51454/19/391/199/54
=50%","60487/80335/91893/9070/29081/51108/29063/10196/6419/29104/92342/10400/58508/64
/2683/4478/2042/50863/23022/375033/3958/928/10417/91624/351/5175/2350/30815/27020/532
al=34%","5321/283/857/11343/151056/4023/9388/3990/7148/150763/348/7376/1373/847/718/33
","140576/4208/8826/311/25932/7057/857/3707/8754/4734/845/1393/1906/3779/3726/5641/330
tags=47%, list=27%, signal=35%","51429/389/6281/11346/11135/2934/148170/25802/9948/10174/
al=34%","5321/283/857/11343/151056/4023/9388/3990/7148/150763/348/7376/1373/847/718/33
643/4162/2013/5155/1909/947/2149/23213/2150/2702/2621/10391/3675/1535"
3/10627/4651/4643/4627/84952/4641/55930/79784/4629/103910/4642/92922/4635/4634/4644/4
t=26%, signal=51%","7048/3133/3105/8767/3659/51043/975/7292/57045/29126/8764/1604/3965/
e-05,4517,"tags=65%, list=25%, signal=49%","7098/857/55601/7096/51296/7097/3430/7099/7100/
%, list=20%, signal=29%","301/1601/5166/493/857/5978/27122/157506/654/5468/5743/8013/271
,"6829/64895/904/23144/10914/54623/3297/91746/56254/7936/11052/29894/55149/11051/5533
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","51474/857/10577/158219/4023/7851/9388/3990/29116/2494/19/6646/51302/9420/26119/106
22%, signal=29%","10413/25959/1601/367/196/5547/7026/2908/26508/56937/6591/4306/329/706
al=29%","79971/64215/64285/283/5066/8754/5978/7043/5208/7040/7424/55107/654/5468/80024
5637/6633/6635/79084/10419/2079/6628/6636/6632/3276"
48/30846/493/857/7035/477/4846/3356/161742/6401/5743/120892/1495/5348/2149/1013/1499/1
474/857/10577/158219/4023/7851/9388/3990/29116/2494/19/6646/51302/9420/26119/1066/348
=38%, list=23%, signal=30%","79026/5587/7070/3270/477/845/6548/28954/5350/2669/401152/117
signal=41%","3280/183/2737/2719/3217/30812/2246/4070/255743/2297/2668/652/1499/6943/74
74/4323/1435/2737/928/27185/3912/5028/7042/1880/1600/64805/54209/1499/6347/6376/706/56
al=58%","301/5906/2771/5978/6814/4851/5908/6688/3162/3965/7879/2213/3134/2294/3071/113
40%, list=28%, signal=29%","1601/55304/6591/3269/9839/3707/9023/5978/2246/27122/6342/5170
908/348235/91272/10403/1063/891/220134/221150/701/5347"
82%, list=13%, signal=71%","4067/7096/54209/2150/7128/10333/64127/9867/1535"
4%, signal=28%","51474/2059/90102/51429/683/389/23189/395/6281/8826/93663/11346/85360/1
%, signal=48%","4094/2737/6591/860/81792/5744/654/652/7227/1499/2202/5916/94/4017/7704/
%, signal=47%","4298/8295/57634/8089/10902/86/51412/55689/6944/10856/54799/55257/8607"
11%, signal=47%","4298/8295/57634/8089/10902/86/51412/55689/6944/10856/54799/55257/8607
signal=65%","4067/7096/8767/7097/54209/2150/7128/10333/64127/9867/1535/118788"
00/6641/5136/8826/4651/4643/4638/4627/493/10788/355/6004/4641/6711/3707/55930/79784/46
7%, list=10%, signal=60%","683/5587/81031/2013/55679/3987/967/7076/1495/2316"

","3572/23500/3280/146433/443/30812/7040/5803/4851/5324/4067/23529/5583/7097/1600/542
, signal=28%","11031/23189/858/2690/677/2308/81565/5872/183/5166/4641/23371/2771/5580/2
t=16%, signal=46%","3673/861/90993/2022/7431/5159/28984/7043/7040/2149/652/54206/3569/5
280/5205/2737/1889/1909/3198/1906/4803/7412/7020/2065/3200/26018/10370/57007/6507/367
8%, signal=59%","5587/7057/2246/64094/4851/2260/5641/3791/3315/9037/8823/7114/8817/2247
39%","4208/3280/183/2737/2719/3217/1282/30812/2246/4070/255743/2297/2668/652/1499/694
signal=43%","4094/2737/6591/860/81792/1513/5744/654/652/7474/7227/1499/2202/7481/5916/9
, list=22%, signal=29%","117854/1956/5587/493/857/284/3270/8412/624/154043/3082/7040/351/
","10413/2/4208/182/90/117854/8463/4780/3280/1910/64321/6591/860/5978/2303/30812/4070/
22%, signal=27%","51594/23049/196513/6499/54542/4292/9924/30833/65110/11340/23708/5101
%, signal=45%","4300/64863/353274/1017/55729/51535/7175/57343/80312/54784/25942/23133/9
'tags=44%, list=16%, signal=37%","84446/29086/60561/54984/5885/80279/91603/8883/29980/111
6%, signal=37%","54984/10111/10694/100134934/10575/54433/5609/7520/3183/7015/6714/3178
st=16%, signal=56%","1969/9314/347/3586/1991/3569/2150/4938/9655/3965/140885/4940/59307
ist=26%, signal=43%","5654/2200/10272/51232/4091/4851/7130/4681/5468/94031/8321/7474/642
843/6856/302/10981/3688/311/4323/3916/967/4629/309/23682/6386/51393/8934/813/3309/936
/54843/6856/302/10981/3688/311/4323/3916/967/4629/309/23682/6386/51393/8934/813/3309/9
059/7049/2/3490/687/196/7048/6446/3624/329/387/978/1728/10365/7040/1393/2720/54762/377
st=22%, signal=41%","23353/254528/57820/4292/27030/317761/9134/10734/11144/51547/256006
list=19%, signal=55%","5321/7057/120892/90865/7099/3566/7474/3586/54209/84868/10333/1075
"64785/675/4172/5888/4175/8318/9837/3978/4173/4171/5422"
.82/1969/4124/2200/1290/2737/7070/1289/4853/1295/7068/30812/8412/639/2295/5797/6121/21
l=31%","10413/135228/677/301/182/1969/3280/25932/2735/23022/2355/8754/9314/7538/4851/3
, list=21%, signal=43%","56943/171023/6878/55578/56970/6879/6873/7398/2648/117143/9913/82
01/1601/3077/183/3624/8322/5978/6814/5208/1393/5028/55107/2697/1906/654/5468/80024/40
%, list=26%, signal=53%","301/5321/3554/10865/8767/3566/9655/7292/84868/3142/9173/6556/35
=42%","9148/6045/54623/10016/84108/6015/4591/84759/84541/10336/56970/1643/8452/55120/
=41%","6239/63976/9580/1051/284207/5743/59341/116844/338557/64641/6517/51043/3914/101
, list=26%, signal=35%","2152/2934/8301/55075/83595/355/5978/10392/114548/387/4217/1611/8
73066551e-05,5835,"tags=46%, list=33%, signal=32%","1903/10203/493/2771/1901/1909/5744/502
84159/83716/6237/64651/7043/444/1277/25976/4313/51176/4487/10159/7122/22943/5083/6497
ist=17%, signal=43%","23500/3280/4851/1600/54209/1499/3400/1745/1746/3949/11221/9283/957
78527e-05,4237,"tags=56%, list=24%, signal=43%","23381/51594/23049/196513/6499/65110/23708
6","301/6778/114548/1893/604/3566/602/3586/975/3569/942/255877/972/9655/7292/3142/6412
6, signal=27%","10413/7098/3673/677/7132/8531/1410/4055/4638/80303/183/2022/79073/6591/3
list=20%, signal=27%","10413/7098/3673/677/7132/8531/1410/4055/4638/80303/183/2022/79073
79971/11031/51429/5125/7045/9469/10981/56548/5587/5205/5872/25825/5066/58475/9341/236
%","3685/10226/158219/4023/4481/5468/19/80149/4792/4790/54209/1066/3690/3569/5467/737
%, signal=41%","7414/90102/395/6281/4323/7070/7057/9748/2042/1307/79834/6385/3987/10979
list=23%, signal=41%","7414/90102/395/6281/4323/7070/7057/9748/2042/1307/79834/6385/3987
42%","683/4780/183/6649/10516/2771/5580/5446/4846/1728/7040/1906/6648/57580/7305/6458
24,"tags=50%, list=26%, signal=37%","2152/2934/55075/83595/355/5978/10392/114548/387/4217/
0%","6606/11218/6634/50628/6637/79760/6633/25929/6635/8487/6628/6636/6632/79833"
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6, list=18%, signal=44%","3688/6387/2113/9314/387/58494/6484/6401/7412/6403/1991/3676/3569
,"tags=74%, list=25%, signal=56%","7098/857/7096/7097/3430/7100/2150/57590/9867/91543/5714
57/7048/4323/2737/857/5159/7043/1393/4091/2099/10865/1495/50846/10095/3566/4313/6422/
./5359/4067/8013/8773/10397/3566/284021/85480/2242/3937/5169/5294/3162/3965/8673/64109
signal=42%","7049/5327/5069/3624/2730/1909/4851/1906"
signal=50%","10392/3105/23643/7096/3106/7097/7099/54209/3123/64581/10333/721/64127/912

%","7098/7057/8767/3106/7097/7099/3659/7805/958/4790/3586/11213/64581/23236/7292/1240
7%, signal=41%","7098/7057/8767/3106/7097/7099/3659/7805/958/4790/3586/11213/64581/2323
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%","301/6778/114548/604/8767/80149/3566/602/1880/51176/51043/3122/5734/64332/3569/312
%","79971/3673/90/3688/1969/3624/2303/9314/3691/2697/8091/6886/652/7474/51176/3690/748
","5893/2072/4292/79677/5892/2068/10111/11144/7517/25788/51002/126549/7515/7014/4436/6
gnal=59%","3685/158219/5468/19/4792/54209/1066/3690/3569/5467/7376/51099/9619/1401/546
%, signal=53%","5592/5154/5155/7056/928/5580/4846/10019/114897/84959/348"
al=60%","3688/928/4023/5468/19/7099/54209/1066/975/348/9655/10333/2207/3949/6850/4615/
=43%, list=22%, signal=34%","6788/4055/8406/51330/183/8682/8742/7057/7078/355/138151/3624
40%","3280/183/2737/2719/3217/30812/2246/4070/255743/2297/2668/652/1499/6943/7481/553
31/840/5327/5125/302/2152/6281/4323/2934/861/4627/824/2737/7057/7123/871/1889/25825/8
nal=58%","23283/11052/29894/11051/55339/10978/8189/81608/10898/1477/79869/51692/1478"
ags=65%, list=29%, signal=47%","196/100507436/3077/3133/604/3105/3106/56940/84868/1604/64
35,"tags=44%, list=16%, signal=37%","84446/29086/60561/54984/5885/80279/91603/8883/29980/1
gnal=45%","100507436/3133/3916/3105/55207/3106/2214/135250/5873/84868/3965/353091/3592
%, signal=43%","1974/9617/85476/8663/9669/8661/3646/8894/1984/10767/1982/8664/54505/728
7048/1896/5604/3202/652/51176/1499/5916/6662/2294/7477/5594/340419/6469/5595"
5%","54885/1793/143872/90627/395/8826/57514/93663/85360/141/9459/7070/54453/80243/107
3%","2934/6444/23704/857/3270/477/845/6548/6324/5350/2697/8654/30845/6330/1756/80149/1
1/3688/81031/3077/183/7057/493/7779/477/6548/285195/2350/51393/59341/51296/6560/2668/
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7%","23405/26523/10799/79042/10556/80746/54931/10775/27161/3028/79897/6895/29102/5448
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1%","1969/5872/6591/4641/79987/7043/90952/5175/1003/9076/5583/79778/137075/4867/1366/7
5%, signal=55%","29777/10799/8602/55720/27340/11340/84294/23212/90459/9790/112479/6227/
35/2/3688/7538/5355/57007/3676/3690/9034/1524/10630/1238/1230/2247/1232/1234/3579/2532
4%","1435/3624/146433/7040/351/7097/652/1436/2629/3398/3123/3059/714/6688/375387/841/1
signal=40%","3280/183/2737/2719/3217/30812/2246/4070/255743/2297/2668/652/1499/6943/74
=29%, signal=40%","5654/1435/2737/4038/860/10468/1896/1495/5054/652/51176/1499/4487/480
0851049e-05,2785,"tags=57%, list=16%, signal=48%","493/4641/81792/1634/967/64094/23554/529
gnal=41%","25937/11030/1601/81031/3624/7431/6711/7043/7040/57552/4091/654/5468/7042/23
4,"tags=55%, list=20%, signal=44%","5125/1910/183/1889/387/4846/1906/1359/2028/185/2149/15
gnal=37%","80279/91603/8883/29980/1111/63967/79035/983/58493/4998/51514/11073/641/672/
27866/274/57647/23353/23588/10456/10527/2869/3756/10434/56943/7341/27101/163087/6490
l=32%","2308/117854/5587/10769/79443/9531/10392/10475/1634/26353/81603/5899/55062/673
5/4643/4162/2013/5155/1909/947/2149/23213/2150/2702/2621/10391/1535"
gnal=34%","7414/8301/3077/2013/10516/257364/284/54741/30851/8546/6810/1499/7305/10076
%","4208/10981/1910/2737/9839/967/23682/4254/4286/5167/56999/6648/5873"
5, signal=53%","10392/23643/929/7097/7099/85363/64581/51284/64127/81035/5724/51311/8685
'2059/7984/1880/54795/2621/5294/6688/91662/3965/3592/2131/7226/81704/998/1230/240/811
gs=72%, list=20%, signal=58%","90102/861/183/28984/8837/7040/30008/7148/1605/3569/23187/1
t=25%, signal=46%","6387/387/58494/351/199/7474/3676/29108/8764/6364/81704/284114/6348/
al=28%","4704/551/4713/57602/2932/6341/9131/51537/10989/64863/80025/23408/10456/9026/9
5,4237,"tags=54%, list=24%, signal=42%","23381/545/2072/170506/8658/51750/54984/10694/1001
%","3688/23643/90865/8808/728/8061/4332/7305/10288/23601/10871/4688/8915/1524/4046/10
gs=67%, list=15%, signal=57%","23500/3280/4851/1499/3400/1745/1746/11221"
t=28%, signal=35%","3074/4121/4907/1486/1806/5580/81792/216/978/4758/7040/27115/2720/25
","8301/4038/4481/3459/93649/728/54209/348/5868/1191/3684/718/4035/7274/4043/3949/4311
gs=61%, list=24%, signal=46%","183/9314/10365/1906/5743/6648/7099/199/64581/1191/5806/103

=57%, list=22%, signal=44%","10413/182/1903/10252/3280/857/161742/4851/3204/3397/1499/285
60%, list=24%, signal=46%","183/9314/10365/1906/5743/6648/7099/199/64581/1191/5806/10333/
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=67%, list=28%, signal=48%","4208/90/116039/7043/7040/1051/654/7020/652/2201/9636/5027/93
:34%","857/11343/151056/4023/9388/3990/7148/150763/348/7376/1373/847/718/338328/5294/9
list=25%, signal=60%","4323/1435/7057/146433/59341/5919/728/1436/64805/54209/6376/719/15
20%, signal=56%","23107/54931/26995/92170/79072/22868/6472/708/8846/51204/26164/10128/5
0640/9960/23265/84440/55763/9150/56155/27309/11113/9727/129138/5300/26073/9817/57448/
%, list=29%, signal=47%","1969/6591/8754/10519/2303/9780/5328/4067/7042/5420/5054/23075/6
","tags=51%, list=18%, signal=42%","6788/4055/8406/51330/183/8682/8742/7057/7078/3624/857/1
.601/5321/183/3624/8322/1906/654/185/114897/5027/4824/3553/2587/1588/3784/6696/5443/56
=29%, signal=33%","54843/302/8826/57664/4651/5321/2934/8301/8724/6840/55614/10788/10979
l=32%","3673/367/1969/7048/1435/6239/2737/3488/5066/857/6778/7043/1051/2099/2697/5241/
","8295/57634/8089/10902/86/6944/79829/10856/54799/55257/8607"
gnal=24%","3074/259230/5321/5154/84962/166929/5337/824/126969/5155/8871/10351/3707/535
signal=48%","683/80310/6387/3554/7070/6484/6401/6403/1991/3676/5734/10894/3383/23236/25
=45%, list=16%, signal=38%","54984/10694/100134934/10575/5609/7520/3183/6714/3178/10576/
53/85301/1893/309/2121/5916/54796/4488/2131/5915/6662/2261/6781/4921/25987/10269"
e-05,4550,"tags=61%, list=25%, signal=46%","3426/715/716/710/602/54209/975/729/735/4153/730
8%, list=25%, signal=29%","163255/23410/26058/51335/9470/8087/9148/6134/8872/83939/7349/2
t=20%, signal=26%","64114/26524/5525/1601/64285/29969/6672/8301/8724/257364/284/9531/59
gnal=40%","3269/284/10979/624/7040/7010/654/1003/59341/348/7082/5339/4881/7122/9590/94
=56%, list=16%, signal=47%","3673/80310/183/5155/9510/6282/3488/5159/80005/3479/8013/7099
53%, list=24%, signal=40%","183/2737/2719/3217/30812/2246/4070/255743/2297/2668/652/1499/
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ags=54%, list=24%, signal=41%","301/7048/861/114548/604/8767/80149/3566/51043/3122/975/643
34%","7049/84159/7048/2615/2737/6591/3624/64651/116039/7043/444/2295/5396/25976/7020/7
e-05,3404,"tags=67%, list=19%, signal=54%","5125/1910/183/387/1359/2028/185/2149/1511/2150/
=29%","301/7132/5321/56898/2687/5166/8309/5066/857/2052/11343/6342/80221/4023/27306/93
=44%","1601/183/3624/1393/2697/654/185/2149/114897/2150/4824/2587/51762/3625/668/5443/
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%, list=22%, signal=38%","196513/54542/4292/9924/11340/170506/22894/27327/340719/10642/49
=21%, signal=48%","353274/1017/55729/51535/7175/57343/80312/25942/23133/9869/22880/230
list=21%, signal=48%","353274/1017/55729/51535/7175/57343/80312/25942/23133/9869/22880/2
985/675/641/672/3619/5888"
77%, list=20%, signal=62%","1017/55729/51535/7175/57343/25942/9869/22880/10155/1786"
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, list=20%, signal=62%","1017/55729/51535/7175/57343/25942/9869/22880/10155/1786"
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85%, list=15%, signal=72%","23283/11052/29894/11051/4686/81608/10898/64852/79869/1478/22
, signal=47%","1974/8663/9669/8661/3646/8894/1982/8664/54505/728689/27335/3692/1964/109
l=33%","7798/7049/90/7048/57493/26508/3280/639/4846/4091/4851/23493/7042/8321/652/747
","tags=58%, list=16%, signal=49%","10413/25937/3913/5155/947/2668/652/1499/6943/55366/629
list=19%, signal=36%","22823/26993/26147/22938/84656/10445/94104/2648/6871/56254/55683/7
1/10199/92856/65083/8602/26156/26155/55813/27340/55127/134430/9790/65095/10885/10607/
6/5375/377677/4642/5354/3710/6810/5339/4359/7122/2752/4155/4099/400746/153562/51090/8

2%","3490/1601/6591/9023/5978/2246/27122/6342/51700/10826/654/50846/27120/51302/9420/
signal=51%","6118/54465/84101/8379/25906/7175/9984/8563/51451/7517/5514/79621/11200/55
3/22919/9585/55920/3192/9928/1062/201161/55143/9055/81930/9212/3619/990/146909/6790/2
417/5322/7099/5919/1511/1991/64581/5806/6688/8915/1524/5336/6850/6283/4057/6279/50856
al=56%","3980/8505/5426/328/5111/3978/5424/2237"
, signal=35%","5327/5125/2152/5654/4323/11098/3426/1513/715/716/5045/5328/4327/94031/70
%, list=20%, signal=36%","677/182/861/1435/4853/3624/146433/2113/7040/10826/4254/338382/2
90102/84159/83716/6237/2719/64651/7043/444/1277/25976/4313/51176/4487/1605/23187/1015
015/55692/6634/6637/25766/6629/6633/51747/6635/6631/6628/6636/6632/6626"
:=20%, signal=26%","55970/51429/54843/858/301/6281/8826/57664/2791/4240/59345/8301/2771
al=30%","23381/2010/8841/65083/2932/51366/54535/55308/23049/10527/6428/26953/6059/651
3381/2010/8841/65083/2932/51366/54535/55308/23049/10527/6428/26953/6059/65110/56943/7
%, list=20%, signal=38%","2/100507436/3133/9111/3105/710/5468/3428/54625/6737/3106/11326/
nal=45%","3572/5592/928/5580/10019/4067/114897/84959/1511/3569/10630/22915/6850/3717/8
20965864e-05,3782,"tags=64%, list=21%, signal=51%","5125/1910/183/2013/5155/387/185/2149/2
23%, signal=27%","1793/3572/1601/6281/4739/4124/6239/81565/3280/6387/2737/146433/10519/
","6788/677/5654/7026/1956/3280/283/5155/4853/329/4188/639/1051/3726/5468/5743/84870/2
77273137e-05,2576,"tags=50%, list=14%, signal=43%","127435/3339/1634/54829/633/4969/4060/2
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=30%, signal=49%","10772/6432/6829/904/57060/6433/6431/708/56254/7936/3190/79882/9810/1
73/3655/1278/3691/1277/2629/54206/8626/1041"
s=53%, list=17%, signal=44%","114548/3479/80149/338557/54209/1992/54206/114769/6376/7128/
, list=28%, signal=40%","1969/283/3140/10392/3133/2634/3105/8767/7097/6039/5919/79792/728,
%, signal=54%","395/1956/283/183/1909/3356/6401/2769/728/64805/942/3357/552/115727/5031/
signal=28%","80312/6996/200424/54496/54784/56154/53615/9760/7874/10419/1789/121642/921
=21%, signal=33%","56243/5125/7048/4323/81031/2737/3217/860/116039/4188/2303/1909/15750
1/3688/1956/8301/84061/57477/183/10769/3269/6446/567/3678/477/3356/25978/590/351/1393
1%, list=12%, signal=36%","5609/7520/54921/64785/6714/3178/10576/7203/22948/908/4931/8429
7263e-05,3511,"tags=54%, list=20%, signal=44%","11030/1601/90/7048/3077/2022/3624/7043/704
list=14%, signal=71%","182/26508/6591/4853/4851/23493/23462/6615/54567"
ist=23%, signal=40%","51474/2059/2934/25802/6840/6711/130271/9948/5962/11034/23075/5756/
8%, list=8%, signal=44%","3673/861/90993/2022/7431/5159/28984/7043/7040/2149/652"
=48%, list=8%, signal=44%","3673/861/90993/2022/7431/5159/28984/7043/7040/2149/652"
4%, list=26%, signal=55%","3140/567/3133/3105/3106/3122/8417/3123/5027/3592/3134/5817/313
gs=65%, list=22%, signal=50%","10161/2151/5159/1281/2840/11214/2149/2150/8477/2846/9002/5
6","2217/3958/2212/11027/2205/2214/2206/2207/2209/2213/7450/2204"
nal=42%","4780/7057/284/28984/4217/7010/641700/1003/5054/958/3586/3676/3791/6347/8644/
5","84440/79848/56155/8452/11113/9727/5300/26073/57448/144455/57026/9738/9585/80179/66
%, list=18%, signal=60%","57381/25932/8322/5159/1282/196740/22899/94/4693/1545/8829/5156/
14/3913/3688/1969/81565/8301/6387/334/2737/7070/2050/4038/2042/3908/1889/4853/91653/2
%, signal=41%","301/6778/114548/1893/604/3566/975/3569/942/972/9655/7292/3142/64127/915
signal=39%","56034/80310/5154/1956/183/5155/2355/358/5159/2335/2099/6277/26471/3479/88
2e-05,5018,"tags=59%, list=28%, signal=43%","57326/347/4067/8773/5054/7099/80149/3566/5187/
1%, signal=48%","2908/5155/688/2113/9314/7040/4091/5468/7042/2353/1958/3586/8061/4804/7
9/23189/3685/7094/3688/1969/1956/3678/152273/10519/23586/387/81848/11170/2191/59341/19
nal=59%","6152/6125/6135/6134/6147/6123/23195/51154/55299/23212/22907/55794/22868/612
=59%, list=21%, signal=47%","367/1435/10252/30812/2246/947/1906/2668/652/7474/1499/4804/74
ist=15%, signal=49%","90102/6239/84962/1289/83478/3678/10979/1839/94/4488/5728"

1435/857/10519/5978/4217/1728/8878/1958/1436/54209/5606/6376/5291/5027/65018/1524/270
57,"tags=83%, list=25%, signal=63%","3688/928/4023/5468/19/1066/975/9655/2207/3949/6850/36
%","558/840/3572/4208/2632/355/284/7068/2730/10763/387/1728/7043/4929/351/1051/5803/26
855e-05,3649,"tags=73%, list=20%, signal=58%","3133/3107/3105/6892/3106/3122/3115/3123/972
3%, list=20%, signal=58%","3133/3107/3105/6892/3106/3122/3115/3123/972/3113/84888/3127/31
%","1601/183/3624/8322/654/185/114897/4824/2587/3784/5443/56729/1408"
st=20%, signal=44%","7048/4000/5154/5396/652/7474/3791/1499/7481/8646/6662/4609/2294/64
07667394339487,2.15324201951868e-05,0.000105685695257535,4.82056851814441e-05,4897,"tag
=32%, list=22%, signal=26%","25937/26524/135228/5611/7026/57493/10252/1848/183/2022/5693
tags=93%, list=20%, signal=75%","1017/6742/546/30968/144455/8317/80174/10926/25792/79733/
","9585/55920/3192/1062/9055/81930/9212/146909"
=46%","183/2771/5580/5446/7040/1906/6648/7305/64581/2150/285590/3684/4688/4689/1535/6
41%","4714/9167/4700/1345/1340/7388/4696/4695/10975/126328/1349/6834/4720/1350/51079/4
9%","3280/183/2737/2719/3217/30812/2246/4070/255743/2297/2668/652/1499/6943/7481/5536
, signal=47%","3913/182/8826/4643/1910/4853/2303/1909/947/5175/5420/652/2296/2131"
gnal=45%","2908/5155/688/5978/2113/9314/7040/4091/5468/7042/2353/1958/652/3586/8061/54
, list=8%, signal=65%","91603/8883/63967/4998/11073/90381/990"
%, signal=34%","29995/3269/493/10519/2274/6548/1906/4773/3479/2065/3710/3708/3709/1126
t=20%, signal=25%","79971/10413/25937/2040/3688/1969/161291/1956/5587/4478/283/2737/493
gnal=35%","5525/5587/5155/5159/388121/7040/7010/113791/4067/9021/64805/975/5305/9655/2
4336e-05,2434,"tags=73%, list=14%, signal=63%","90627/7057/79812/9314/387/161742/4851/6915/
nal=44%","8697/25906/29882/9820/64682/51529/51434/991/701/27338/11065"
182/3280/4638/2022/5159/1281/4851/30008/23493/56999/7042/23462/94/4035/54567"
10516/79875/1281/4629/30008/7148/4015/11117/4239/4054"
list=20%, signal=42%","170506/8658/51750/546/10111/51548/10694/10575/5609/7520/5514/3178
list=27%, signal=33%","2059/51429/23189/93663/11135/2934/23500/148170/25802/6840/6711/55
3e-05,2048,"tags=67%, list=11%, signal=59%","22909/284131/1773/55345/348654/1775/1763/328/9
7%, signal=26%","23189/1903/23500/2200/3280/1910/7070/4038/567/5978/4070/5744/56920/105
al=40%","9682/8085/79609/55011/23067/4800/55677/10322/546/11105/11137/57343/23476/556
nal=37%","51474/2059/2934/25802/6840/9645/6711/130271/9948/5962/11034/23075/5756/2150/
signal=66%","80310/5155/5159/2303/947/8837/4851/59/1958/652/3570"
3/23704/493/857/3270/477/845/6548/9096/6324/5350/2697/30845/6330/3752/54795/10060/270
","11031/10981/8406/23682/55647/83871/338382/55207/5322/5027/339122/7879/10312"
14034/11340/118460/26173/54512/55174/79650/80789/55756/51010/25896/56915/57508/64852/
-05,4730,"tags=74%, list=26%, signal=55%","301/114548/8767/3566/51043/3122/64332/3123/942/
=45%","6387/1893/387/351/347/23143/199/7474/3676/29108/8764/6364/81704/326/284114/110
57/10577/6342/309/5375/347/54762/19/6646/975/706/950/80765/6820/8435/335/114876/54847
5004/6237/51285/5906/9077/388/387/2669/5899/2633/55207/8934/9367/64121/5912/5868/5908
%, list=24%, signal=32%","5587/7070/857/3270/477/845/5350/4067/2149/1756/1193/84959/6376/
5298,5.40427763322163e-05,4700,"tags=75%, list=26%, signal=55%","29777/10799/8602/27340/113
12,"tags=34%, list=14%, signal=30%","7336/9521/196528/165918/5932/55693/55183/83933/84501/
79%, list=19%, signal=64%","7057/10392/10875/3122/3123/972/64127/6556/2213/50856/968"
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tags=42%, list=24%, signal=32%","3572/2690/7132/3280/857/10019/4067/54625/83666/23529/3479
7%, signal=27%","51474/2059/51429/23189/93663/11135/2934/23500/81565/8301/6387/148170/4
2908/5155/688/5978/2113/9314/7040/4091/5468/7042/2353/1958/652/3586/8061/5467/4804/737
6","3280/183/2737/2719/3217/5159/30812/2246/4070/255743/2297/2668/652/1499/6943/7481/5
tags=73%, list=27%, signal=53%","9389/100287932/9997/345778/6687/10440/10939/100131801/60
=13%, signal=38%","1956/183/7057/493/51655/4846/8654/348/2357/2977/23576"
t=25%, signal=45%","27202/6387/79148/79890/1843/347/7130/4681/3204/23143/653/5734/6347/

ignal=60%","10412/90441/51388/3692/23195/26155/84549/51154/23212/84261/81887/117246/54
3688/141/3077/5872/30844/30846/567/9341/3133/3107/54918/351/23678/3105/285195/79628/8
gnal=30%","558/1969/1956/7006/2050/10461/2042/79834/5159/5580/10114/3716/51232/7010/40
51/857/10577/158219/5468/5360/19/6646/4792/54209/1066/348/7376/6256/4035/1950/344/843
l=31%","6788/1601/1410/7048/80014/85004/183/64321/2719/3624/3488/978/604/4502/4499/449
1435/3280/3953/146433/4851/5324/4067/5583/1436/2629/54209/3569/6376/706/1191/5179/649
signal=44%","7049/90/7048/26508/3280/4846/4851/23493/7042/8321/652/7474/10370/60485/23
/8406/141/5587/79443/3916/25978/8878/23710/55062/6737/120892/81631/92421/10133/10241,
st=16%, signal=39%","5327/677/301/687/90/2908/114907/5978/358/7538/8837/1906/11170/678/5
gnal=45%","10413/3572/3716/6997/4790/55022/3569/3570/6774/6688/22904/1050/361/8737/371
9/301/3077/5978/6814/5208/1393/5660/55107/2697/1906/2350/80024/5243/6892/5360/19/5934
9/3958/5734/6347/10344/23604/6376/6356/6357/6355/6348/11314/6354/7857/6351/6352/38837
=68%, list=28%, signal=49%","9469/56548/1634/633/337876/29940/133584/22856/83539/79586/5
, signal=42%","6419/29104/58508/5929/8085/23512/23067/10322/64324/11105/9757/345630/104
","79707/6129/10412/9875/4736/6130/11224/3692/65008/26156/84549/8886/23560/285855/8436
ignal=28%","3490/2308/5611/100507436/6387/11080/7057/9531/2246/4023/947/4846/6548/3356
%, signal=33%","51474/3953/9023/10577/158219/2246/351/8542/3990/19/6646/8578/51302/942(
st=16%, signal=37%","5327/677/301/687/90/2908/6446/114907/5978/358/7538/8837/1906/11170,
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st=18%, signal=69%","10594/11218/9129/57819/10421/25980/135295/26121/27339/9733/24148"
e-05,2652,"tags=75%, list=15%, signal=64%","10694/10575/10576/7203/22948/908/55135/10574/1
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9/9510/2274/7010/1277/23493/4313/6422/2201/23462/1101/4488/133522/30061"
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2%","3688/4323/79148/9902/1513/1514/1471/5045/4327/2191/5653/4313/1508/1520/9509/9508/
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22%, signal=48%","23347/4292/51750/56893/9984/55183/84083/2187/126549/9400/7014/4436/55
38,"tags=67%, list=30%, signal=47%","25913/2074/50485/7158/6419/10213/24144/5423/27434/233
=31%, signal=41%","4696/4695/126328/4720/28976/51079/4710/29078/80224/4704/4713/55967/4
s=59%, list=31%, signal=41%","4696/4695/126328/4720/28976/51079/4710/29078/80224/4704/471
32,"tags=31%, list=10%, signal=28%","10413/64114/3685/90/367/4000/3655/8682/6591/7057/854
%, signal=76%","5154/1278/1289/5155/5159/1282/1281/1277/1291/5156"
, list=20%, signal=41%","5125/1889/25825/590/1514/5045/1359/2028/1511/834/1522/10159/1363
9%, signal=50%","5125/5045/7376/51129/338328/344/51099/6272/350/335/341"
,"tags=59%, list=15%, signal=51%","55729/51535/57343/80312/9869/53615/22880/23028/3070/903
=34%","302/6342/9388/5045/26119/5244/4790/7376/706/4035/154664/344/335/341"
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08/2934/6444/79073/7057/3270/51655/845/444/5350/813/3710/3708/6517/3709/202333/5339/6
2%","6603/8289/6601/196528/6599/6598/6418/6605/3159/6602/11198/6749/6597"
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"302/3074/4124/8406/141/9531/1203/25978/23120/26471/55647/23710/8546/5899/55062/83871
%, signal=63%","7098/857/23180/2150/7128/4938/57590/84868/81622/57162/81035/79155/38832
st=14%, signal=69%","90102/183/28984/7040/84168/4851/7075/1605/23187/3037/11117/6662"
=27%, signal=45%","1601/3269/5978/2246/27122/5028/654/4790/653/348/6720/9563/5031/1581/
gnal=57%","6634/54433/9887/3183/7015/55651/6635/4931/55226/65057/3192/55135/6628/1736
t=39%, signal=45%","4719/4709/4708/4698/4714/4700/4696/4695/126328/4720/51079/4710/4704
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45%","4719/4709/4708/4698/4714/4700/4696/4695/126328/4720/51079/4710/4704/4713/55967/4
list=29%, signal=34%","3556/6281/1969/183/7070/7058/1307/857/27286/6385/3987/10979/7010/
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%","55728/255374/79042/2072/57523/10556/1677/5395/10595/201164/80746/27301/22894/5405
ıal=54%","10694/10575/10576/7203/22948/55651/908/55135/10856/10574/1736/8607"
st=15%, signal=54%","10694/10575/10576/7203/22948/55651/908/55135/10856/10574/1736/8607
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1%","3133/3107/3105/6892/3106/5476/3122/3115/3123/972/3113/84888/3127/3134/3117/3119/
3%","558/84159/2/4323/7026/283/10461/9510/3624/8322/10468/4846/1051/2099/4254/7130/264
ignal=39%","301/7048/861/2737/3707/9580/639/114548/387/604/8546/8767/10875/8808/80149/3
51%, list=24%, signal=46%","301/861/3077/2737/604/80149/3566/51043/9655/7292/57045/29126/
673/2152/6387/5155/3958/284/1901/3082/7424/6360/2668/652/7474/3791/6376/2252/2150/104
7%, signal=42%","9150/6879/23248/57466/84337/27161/23476/6872/23160/51602/2962/6908/585
9,"tags=50%, list=17%, signal=42%","9150/6879/23248/57466/84337/27161/23476/6872/23160/516
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gnal=46%","1601/183/654/185/114897/2587/3784/5443/1408"
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6%, signal=52%","10417/23586/604/7043/7040/8013/7042/7099/7474/7805/11213/972/8915/3162
%, list=22%, signal=47%","117854/10581/10410/5359/64135/56829/8091/3428/3434/684/8519/963
ignal=33%","9843/79901/56898/8301/3077/567/7779/391059/4502/4499/4493/351/654/8546/664
/85461/3688/4627/30846/824/928/170689/58494/8837/4773/3566/975/11096/9518/8038/5738/9
0%","10413/7057/857/7043/7040/7042/7097/2353/8061/706/2354/10628/4155/6720/2706/1545/6
st=24%, signal=37%","4739/183/84632/7070/5155/857/146433/9961/1839/351/9467/2065/9052/14
343/6284/285362/283/6649/5066/8639/26002/4015/144811/2157/84171/5621/4017/6276/6283/6
104,"tags=80%, list=17%, signal=66%","8379/79003/81929/283489/91272/3192/1062/83540/11004/
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%, signal=41%","7026/183/387/151/6915/146/3398/552/6869/1535/7168/153/147/5144/1268/8076
127e-05,3442,"tags=62%, list=19%, signal=51%","55729/51535/57343/80312/9869/22880/23028/903
05,4514,"tags=61%, list=25%, signal=46%","1793/1601/6281/4739/6239/10519/55679/3987/10979/
24%","10413/3572/23500/4124/26508/81565/3280/6387/2737/7070/4038/567/146433/443/5978/
56%, list=27%, signal=41%","196/3077/604/80149/3566/51043/64092/56940/7292/84868/8764/160
, signal=28%","558/3572/2690/4208/2632/1435/284/5978/2730/10763/387/947/7043/4929/1051/5
ist=23%, signal=50%","6591/81792/1513/5744/652/1499/2202/7481/5916/7483/5915/150356/6662
nal=47%","51287/116228/6834/100131801/1352/6341/7386/79072/285521/51241/493753/790955
18%, signal=32%","857/3270/477/845/6548/5350/8654/30845/1756/80149/1193/54795/5142/2702
52%","10392/3105/7096/3106/7097/7099/3123/64581/10333/64127/912/284297/1755"
%","135228/5154/1956/4038/3624/3488/10468/1896/9079/4851/7042/7474/60485/1499/8323/480
, list=20%, signal=28%","51429/1601/2308/344558/152503/4780/4478/10769/285671/8742/64321/
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3%, signal=25%","79971/5125/7026/2114/26508/3280/64321/2737/2719/3217/8553/2303/3087/79
st=25%, signal=48%","26471/5743/120892/3479/90865/8877/54209/3569/6376/7133/1075/3553/39
54%","10445/55683/11143/54934/51230/84148/84289/284058/79960/9767/3054/11091"
54%","10445/55683/11143/54934/51230/84148/84289/284058/79960/9767/3054/11091"
%, list=13%, signal=44%","5154/824/5155/23682/3356/11261/3357/2170"
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%","2040/4774/117854/10581/10410/5359/10475/64135/4851/56829/8091/3428/2494/8934/3434
24%, signal=28%","79971/2040/161291/5587/283/2737/4641/8754/10519/9531/54741/5580/7043,
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=21%, signal=35%","7048/4323/2737/3217/860/116039/4188/2303/157506/7043/3201/3216/3198/
726/10930/23091/23070/115708/142940/83480/54931/26995/79066/54517/54888/56339/80324/
al=41%","5166/3953/1901/2697/7107/84959/6422/1436/3690/3569/7128/5027/3635/338773/1335
, list=32%, signal=52%","3399/30812/7040/4851/652/5467/28514/144165/9573/6662/116173/1649
=33%, list=24%, signal=25%","79971/2040/5784/64215/161291/64285/5587/29969/6672/81565/28
,"3685/4651/7402/84168/7408/1756/57348/3690/10630/3675/3135"
gnal=38%","23500/3280/4851/1600/54209/1499/3400/706/1745/6497/1030/1746/3949/11221/29
=39%","5125/1889/25825/590/1514/5045/1359/2028/1511/1522/10159/1363/5126/1215/1803/51
t=30%, signal=33%","1601/3077/3624/5208/5028/55107/1906/654/5468/114897/6934/2746/33855
ignal=29%","301/1601/3077/183/3624/5978/5208/1393/5028/55107/2697/1906/654/5468/80024/
=34%","7026/5836/8309/6342/5375/5660/5468/2173/338557/8644/5467/6256/241/80765/2170/5
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)%, signal=33%","683/29995/493/10519/2274/6548/120892/2633/3479/2065/3708/11261/54209/6
list=19%, signal=38%","23500/3280/146433/443/9948/5803/4851/7097/1499/1605/3400/7133/174
.0360/51547/4691/11137/84365/2975/9329/2972/9330/84172/6598/6949/9328/25926/1663/2976,
%, signal=33%","9469/56548/4780/2687/5166/284/23743/2730/10090/9060/6484/10678/4258/51
3%, list=23%, signal=33%","6341/10989/55333/55750/9804/55735/119559/8834/23761/285521/91
%, list=17%, signal=70%","56034/80310/5154/3554/5155/3678/5159/4067/3690/5156/5795"
, signal=29%","25937/26524/5525/7026/57493/10252/84962/183/7070/5787/857/5580/9961/1011
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s=70%, list=20%, signal=56%","5906/2771/5978/6814/4851/5908/6688/3162/3965/2213/3134/2294
signal=32%","6788/677/7026/1956/3280/5155/4188/639/1051/3726/84870/23493/9021/5604/2034
7%, signal=29%","51474/2059/79901/389/9429/4643/2683/5205/4627/4641/1192/358/151056/797
4216/115939/51504/79922/79066/54433/345630/10436/4839/2091/117246/55651/55226/27079/5
"135228/5154/1956/4038/3624/3488/10468/1896/9079/4851/7042/26018/7474/60485/1499/8323
5228/5154/1956/4038/3624/3488/10468/1896/9079/4851/7042/26018/7474/60485/1499/8323/48
=21%, signal=67%","11340/118460/22894/54512/6832/87178/9567/51010/56915/5393/23404"
11307484067,3097,"tags=39%, list=17%, signal=32%","117854/3554/567/10417/3133/6778/23586/
0%, signal=33%","4208/2308/8826/4780/1956/25932/3077/567/3707/140609/358/4502/4499/449
'4208/367/7048/10252/3280/64321/2737/2246/157506/4851/5604/3200/2668/652/7474/10370/1
","7040/10865/8877/7099/51043/3569/23180/7292/5621/64127/5008/3592/3071/4615/1649/946
%, signal=51%","7040/10865/8877/7099/51043/3569/23180/7292/5621/64127/5008/3592/3071/46
signal=36%","7431/4534/5339/1832/1674/25984/3861/144501/631/342574/3885/3856/7062/4744,
gs=38%, list=27%, signal=28%","2059/51429/389/23189/1410/93663/11135/2934/23500/3280/1481
l=45%","114034/90806/9924/22894/81875/54512/129563/29883/90459/23019/79650/112479/228
,4246,"tags=44%, list=24%, signal=34%","3572/2690/7132/3280/4067/54625/83666/23529/3479/14
al=36%","683/4208/196/604/4067/23529/4773/7099/1880/958/3586/4332/975/7305/972/5074/36
7098/84159/2/8531/367/4323/7026/5154/283/5872/9510/3624/8322/10468/10046/5159/30812/1
%","114034/11340/118460/26173/54512/79650/80789/51010/25896/56915/57508/64852/5393/23
45,"tags=53%, list=31%, signal=37%","1997/123920/7070/3958/4067/2633/8934/5583/7805/975/56
"57470/55152/51067/54938/23395/124454/10056/55278/25973/23438/79731/5188/57505/92935,
nal=42%","3913/182/8826/4643/1910/4853/2303/1909/5420/652"
%, list=8%, signal=42%","3913/182/8826/4643/1910/4853/2303/1909/5420/652"
328/677/4616/4217/7538/1843/3082/1647/9261/8877/80149/652/5187/54209/5606/768211/6688
95,2652,"tags=82%, list=15%, signal=70%","10694/10575/10576/7203/22948/908/55135/10574/173
%, signal=35%","57602/9131/10456/9049/9530/55750/283514/9520/27429/9804/158234/140823/1
ags=65%, list=18%, signal=54%","54829/1893/4846/7040/4851/5167/23493/5552/54795/23462/262

signal=34%","302/301/11135/6282/9124/23022/91624/1266/90952/27020/54538/1824/29766/183
st=26%, signal=29%","1910/25932/183/5166/2771/1203/477/1909/23682/6548/285195/1906/5564
=80%, list=22%, signal=62%","3077/7057/10392/10875/54209/972/10288/55080/29108/64127/6556
, signal=34%","1601/1910/54884/5978/151056/27122/216/157506/654/6121/3479/220/9249/653/
","567/978/51646/7130/1265/1471/7077/116844/79792/25840/427/25797/5806/2919/1509/1520/
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","tags=61%, list=15%, signal=52%","10694/10575/10576/7203/22948/55651/908/10856/10574/173
0577/6342/309/5375/347/54762/19/6646/1880/975/706/950/80765/6820/3638/8435/335/114876
2%","3572/7132/2308/29995/4000/183/90523/493/3488/6548/4851/1906/11214/8654/5468/2349
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s=38%, list=10%, signal=34%","558/117854/1435/3554/837/23586/81844/64135/1906/54625/8366
3,"tags=43%, list=23%, signal=33%","57602/9131/10456/9049/9530/55750/283514/9520/27429/483
%, list=17%, signal=36%","27202/3082/10365/7099/80149/3586/2629/11213/6376/7128/2621/965
.6%, signal=39%","23248/57466/84337/27161/23476/23160/58506/55250/123169/23168/55197/68
:ags=59%, list=25%, signal=44%","904/645121/8621/85417/51755/1018/1017/1020/4331/9134/905,
3572/7132/2308/29995/4000/183/90523/493/3488/6548/4851/1906/11214/8654/5468/23493/347
list=22%, signal=29%","25937/26524/7026/57493/10252/183/7070/5787/857/5580/9961/10114/46
.14/3913/3487/3490/1601/1410/8826/3688/7048/4323/1956/81565/6387/85004/183/64321/58192
023/4481/8542/3990/5360/348/1191/51599/3949/3931/344/350/338/335/51449/341/5444/3250/3
gnal=42%","4208/123920/7006/4067/4773/5583/4332/975/23180/29760/2213/5336/6850/3071/60
)120772345008281,4142,"tags=38%, list=23%, signal=29%","1410/2152/2934/55075/7057/83595/35
"79026/302/8826/311/6282/2246/6277/6285/6271/7430"
405,"tags=37%, list=25%, signal=28%","163255/10585/26058/51335/9470/8087/9148/6134/8872/83
ignal=43%","10763/3716/6358/6360/6347/10344/6376/6368/9034/6356/6357/6355/6364/6348/63
'246176/285643/2597/81624/9585/6683/55920/29911/348235/7840/79929/80817/10013/166614/
46%, list=31%, signal=32%","51138/389856/8239/377630/401447/9098/27351/9736/10869/57559/1
)%, signal=41%","3916/4067/3566/6810/2242/2150/3684/6688/3162/3965/2213/8673/3134/6850/2
gnal=36%","221908/2648/79648/7514/7517/79959/7840/163786/55226/6491/55755/55835/34865
58/7098/84159/2/8531/367/4323/7026/5154/283/5872/10461/9510/3624/8322/10468/10046/515
gnal=24%","558/7049/11167/4208/677/301/51705/1435/10461/567/355/3624/3707/2113/80177/7
8723335224294,3404,"tags=69%, list=19%, signal=56%","5125/1910/1359/2028/2149/1511/2150/15
","4926/124491/90780/55968/9972/26108/2010/55818/57602/274/23353/54700/9025/10208/734
14/6840/6385/1756/1605/5239/5339/23336"
%, signal=45%","367/10252/2246/2668/652/7474/10370/1499/23213/7481/22943/2247/650/6495/
=22%, signal=58%","1906/120892/90865/7097/7099/3791/6347/5606/9590/10135/64127/3717/677
ags=74%, list=22%, signal=58%","1906/120892/90865/7097/7099/3791/6347/5606/9590/10135/641
%","8826/1956/64285/183/84632/55614/9961/8412/1839/7040/79628/5641/9052/2549/54206/22
55143/23468/9212/11339/3619/10951"
82395140517,5022,"tags=87%, list=28%, signal=63%","1969/7057/79812/185/652/4804/94/3162/90
%, list=21%, signal=45%","56943/171023/6878/55578/56970/6879/6873/2648/117143/9913/8295/2
=57%, list=21%, signal=45%","56943/171023/6878/55578/56970/6879/6873/2648/117143/9913/82
nal=30%","3074/5321/56898/8309/11343/5580/151056/6342/4023/4758/79887/9388/3990/8605/4
1969/5872/6591/4641/90952/5175/1003/9076/5583/79778/137075/4867/1366/7082/7481/6615/5
49%, list=23%, signal=38%","2932/26005/51104/5108/9851/1453/9419/137886/50855/11190/7964
%, list=23%, signal=39%","367/5154/183/6591/79812/3217/30812/81792/2246/4070/3082/2099/269
","3913/7431/443/54587/351/3459/3691/4067/79628/4919/5354/7097/7099/10397/728/54209/16
=53%","5784/1969/2303/161198/79987/7424/7075/94/2296/54345/10630/2324/659/85329/3964"
38,"tags=41%, list=24%, signal=31%","5587/7070/3270/477/845/5350/4067/2149/1756/1193/84959
tags=57%, list=27%, signal=42%","51474/2059/2934/25802/6840/6711/130271/5962/23075/5756/2
st=21%, signal=29%","558/25959/2308/3920/4780/5587/5166/6591/355/4217/4023/4330/1611/49

."2/3673/25959/7048/6591/2771/2730/4023/4330/1728/590/9388/5468/2350/6915/5743/1277/74
%, signal=43%","6419/29104/58508/5929/8085/79609/23067/151194/10322/64324/11105/9757/54
:ags=78%, list=25%, signal=59%","139322/10989/9804/55735/10651/79135/401505/387990/54927/
=24%, signal=39%","2615/6385/1051/8654/11326/652/3586/51043/3123/942/10288/10457/84868/
;nal=30%","831/302/6281/4323/2934/861/4627/7057/7123/120892/9528/5054/10418/114769/835
,"2/1910/5359/2335/1051/5743/9332/23780/3569/3570/4153/2157/445/3553/5265/59307/12/126
=29%","1601/55304/3269/3707/5978/2246/27122/5028/654/80271/26002/8877/4790/2629/653/1(
=65%, list=27%, signal=48%","3572/9180/3133/5743/3569/241/718/5294/3553/5008/1268/695/388
5, signal=45%","55968/2932/26005/5108/9851/1453/137886/50855/11190/79648/55125/22919/647
4155,"tags=57%, list=23%, signal=44%","114034/90806/9924/22894/81875/54512/129563/29883/9
30/5558/5425/4172/5111/5557/23649/5424/5422"
st=11%, signal=65%","301/114548/604/3566/942/9655/7292/3142"
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","1435/146433/90865/1436/2629/54209/6376/1191"
s=55%, list=6%, signal=51%","3280/2022/1909/4851/30008/23493"
4408597,2652,"tags=90%, list=15%, signal=77%","10694/10575/10576/7203/22948/908/55135/105
6%","2013/6237/5906/54453/947/23493/3791/23462/51666/133/55273"
=43%, list=23%, signal=34%","6341/10989/55333/55750/9804/55735/119559/8834/23761/285521/9
18%, signal=38%","3572/7132/2308/29995/4000/183/90523/493/6548/4851/1906/8654/5468/3479
t=19%, signal=37%","10252/183/857/5580/10114/1843/10019/81848/5468/4067/652/6422/2629/3
%, list=22%, signal=54%","79008/54464/548593/54552/25913/167227/2072/8658/51750/54984/10
5, signal=24%","55970/51429/54843/683/858/301/6281/8826/57664/2791/4240/59345/8301/30846
93,2713,"tags=73%, list=15%, signal=62%","23283/11052/29894/11051/55339/8189/81608/10898/7
5%, list=23%, signal=35%","6433/90806/51594/6431/66037/10642/4904/79066/708/56254/7936/81
5,3758,"tags=41%, list=21%, signal=32%","26524/7026/57493/10252/183/857/5580/10114/3087/18
al=56%","546/114991/55693/222229/11335/23310/3070/157570/23468/1786/2146/3619/10951/10
70%, list=28%, signal=50%","51079/581/55750/4836/158234/140823/9141/285521/26520/3308/265
%","79612/570/4836/1387/8260/10847/7994/253782/204219/91012/339983/9862/8888/79969/92
73%, list=25%, signal=55%","4067/11027/338382/3430/54209/2150/7128/57590/57162/4814/4057/
%, list=20%, signal=42%","3399/3488/51232/4091/7424/117581/4851/7130/3199/5468/6422/54795
5%","4023/4481/8542/3990/5360/348/1191/51599/3949/3931/344/350/338/335/51449/341/5444/
23/4481/8542/3990/5360/348/1191/51599/3949/3931/344/350/338/335/51449/341/5444/3250/3
3%, signal=33%","57602/9131/10456/9049/9530/55750/283514/9520/23788/27429/4836/207/980
'tags=100%, list=21%, signal=79%","7132/7040/84168/4851/1471/2191/7133/11117/150356/1803/8
/11052/29894/5438/11051/4686/81608/10898/1479/64852/79869/51692/1478/22916"
"79840/56893/10111/51547/221443/165918/11143/5932/1642/51444/55693/55183/7520/56916/5
=42%","5166/1901/7107/84959/1436/3690/3569/7128/5027/3635/338773/133522/6850/6696/578
%, signal=46%","57381/90993/7057/3678/9314/2246/3716/10365/56999/3586/7082/5291/28514/6
%, list=15%, signal=67%","23283/11052/29894/11051/4686/81608/10898/64852/79869/1478/2291
31%","7098/3673/677/8531/4638/80303/358/624/5743/9076/5243/59341/5269/6517/54206/92241
;nal=64%","10417/7043/7040/7042/7099/7474/7805/11213/972/51704/5336/3135/10859"
, list=16%, signal=64%","10417/7043/7040/7042/7099/7474/7805/11213/972/51704/5336/3135/10
2%, signal=72%","567/3122/3115/3123/3108/3113/3127/3111/3117/3119/3109/3120/3112"
028,"tags=93%, list=22%, signal=72%","567/3122/3115/3123/3108/3113/3127/3111/3117/3119/310
5%","11340/51013/118460/22894/54512/129563/6832/87178/81887/9567/51010/56915/5393/234
3673/301/1410/7026/5154/1956/183/5066/5159/8837/1728/7040/1393/2099/6751/5743/1277/508
09/6376/712/713/3684/718/22943/714/1524"
st=24%, signal=33%","6446/5580/1909/154043/6548/1906/5243/30845/1756/11261/54209/6347/7
42%","7414/23189/3688/1969/6239/84962/9459/1901/29984/123720/10163/64805/5756/221178/
=33%","122786/2934/6444/2013/23704/857/477/6548/6324/5350/2697/59/6330/3752/54795/514

1910/183/2771/477/1909/6548/1906/185/6332/6337/6507/4881/552/6869/2131/482/66002/6558
"tags=72%, list=27%, signal=53%","9389/100287932/9997/345778/6687/10440/10939/100131801/6
3%, signal=34%","4208/3688/3077/5580/1839/27122/351/1393/5028/151/1906/5468/5328/27120/9
98930151,4834,"tags=56%, list=27%, signal=41%","83860/2957/6881/29079/55832/9282/2958/1705
7%","10413/307/3572/84441/182/1903/7048/4323/26508/3280/6591/4853/7079/2303/604/4846/3
"5327/2308/3280/3399/5166/4023/1906/5743/9076/7097/80149/55022/8644/1373/6916/847/232
'677/196/5906/3488/358/351/1393/3309/3709/1373/2230/5908/5294/7430/10411/5179/445/1545
l=31%","10413/7049/4208/7048/861/57493/183/2735/1901/2303/4851/1906/11214/23493/3479/7
'79026/7414/3688/477/58494/10791/6548/6324/2697/1495/5348/6330/1499/5239/5912/2702/182
, list=27%, signal=44%","6419/29104/58508/5929/8085/79609/23067/151194/10322/64324/11105/
35%","8314/100463289/100287932/5663/55288/3099/10105/255027/8034/55072/56947/6687/552
0%","800/51429/8826/4643/2934/7402/57477/4627/824/6711/130271/9948/2037/91663/11214/5
48,0.000186621588773012,4142,"tags=44%, list=23%, signal=34%","2152/2934/55075/83595/355/5
gnal=30%","558/301/1410/80310/4780/824/8754/358/5159/5580/6778/388/4217/1728/3082/1036
=58%, list=27%, signal=43%","558/6387/10461/3707/604/4254/652/972/1831/2621/3059/1942/891
/64968/1153/6122/6188/2971/6125/6135/55168/29074/6191/6167/92856/6222/6208/51023/5133
gs=67%, list=13%, signal=58%","2022/857/9961/5167/11261/54206/91662/3725"
470129,3072,"tags=55%, list=17%, signal=46%","301/57381/10769/10519/27286/2303/5743/3791/3
=34%","196/2052/2329/5446/1728/57552/2948/1429/2327/9/27233/2326/316/1545/29785/874/62
=40%, list=23%, signal=31%","7756/81551/57704/2258/403/50861/57159/126731/10844/3796/233
st=31%, signal=41%","966/567/3133/3107/351/3105/89866/3106/3122/3115/3123/50999/972/3113
57%, list=38%, signal=36%","81565/7431/10763/286077/4534/5339/1832/1674/25984/3861/144501
52,"tags=52%, list=20%, signal=41%","5587/5155/5159/388121/7040/7010/4067/64805/64127/2321
%","10413/3572/3716/6997/4790/55022/3569/3570/6774/6688/22904/1050/361/8737/3717/5781
35%","3920/5587/3916/25978/23710/55062/81631/92421/10241/10159/55283/81671/7879/12886
=39%, list=22%, signal=30%","558/3572/4208/2632/284/2730/10763/387/7043/4929/1051/5803/23
18%, signal=56%","7078/7079/7077/56925/7076/4803/684/50859/8434/9806/5918"
%, list=19%, signal=22%","64764/23189/1601/4780/5587/81565/3280/6387/8724/3655/183/7070/59
73303874,4773,"tags=64%, list=27%, signal=47%","30846/824/8754/4773/3566/54209/7305/10411,
%, signal=49%","4208/301/3707/1514/80301/3569/5291/64127/3665/10269/3661/1234/356/23657
","29844/54617/51377/64431/10428/10847/125476/81611/79913/10445/7528/8295/57634/10902,
signal=47%","7158/24144/51366/23347/3297/56893/78990/55183/2187/55611/84309/9400/7014/
ignal=29%","1969/5872/6591/4641/90952/5175/1003/9076/5583/79778/137075/4867/1366/7082/
l=41%","7048/4000/5154/2719/5396/652/7474/3791/1499/4487/7481/2252/130497/8646/6662/46
s=26%, list=16%, signal=23%","9141/285521/26520/55374/4927/728489/3843/27092/10694/89781/
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)8795020344067,3933,"tags=53%, list=22%, signal=42%","196513/54542/4292/9924/11340/170506/
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'tags=88%, list=29%, signal=63%","100507436/3133/3105/3106/84868/1524/3965/2213/3134/5272/
%, signal=49%","84232/2475/9188/25824/6618/6617/7702/55814/10302/2971/2074/55290/10623/
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7%, signal=56%","6829/904/23144/3297/91746/56254/7936/11052/11051/79882/9810/79869/6626
, list=20%, signal=30%","54843/302/57664/2934/8301/8724/6840/55614/6548/55803/6386/5333/5
5%","79971/10413/3673/677/90/3688/1969/8463/2114/3624/2303/9314/3691/2697/8091/7075/6

2,4773,"tags=59%, list=27%, signal=44%","30846/824/8754/8837/4773/3566/54209/7305/10411/95
%, list=18%, signal=31%","54843/302/57664/2934/8301/6840/6548/55803/6386/5333/341359/8324
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al=39%","367/1306/122786/196383/57493/25932/80781/23022/26084/2195/2317/8985/4855/6781
2,"tags=55%, list=23%, signal=43%","10413/10252/157506/3202/652/1499/1605/1522/10253/2131
416,2785,"tags=53%, list=16%, signal=45%","4641/1634/967/64094/23554/5291/28514/3592/10253
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%, list=19%, signal=42%","4780/7057/284/28984/7010/641700/1003/5054/958/3676/3791/6347/86
"tags=45%, list=24%, signal=34%","3554/3140/567/3133/6778/3916/114548/7040/3105/10865/235
67193607,1167,"tags=55%, list=7%, signal=51%","183/5155/9510/3488/80005/8013"
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ags=59%, list=26%, signal=43%","4926/55968/2932/26005/5108/9851/1453/137886/50855/11190/7
)%, list=31%, signal=35%","4208/1997/123920/7006/7070/3958/3105/4067/10384/4773/8767/2633
list=24%, signal=27%","5563/56890/27124/2932/81551/57704/2258/4591/339829/1946/403/50861
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al=28%","7049/57493/4638/2022/1901/2303/1281/1909/7040/4851/30008/23493/7042/8321/7474
62%","57461/58155/56853/10772/51631/10946/6433/6431/55015/60680/22827/91746/55692/111
s=44%, list=21%, signal=35%","5587/5155/5159/388121/7040/7010/113791/4067/9021/64805/9655
2%","11031/302/3688/8301/10203/857/284/928/4734/79890/967/6401/6386/55788/26119/57007/
gs=74%, list=26%, signal=55%","7048/3133/3105/8767/975/7292/1604/3965/10148/6850/3717/649
74%, list=25%, signal=56%","139322/10989/9804/55735/10651/79135/401505/387990/54927/2581
6, signal=36%","81565/7431/10763/286077/81493/4534/5339/1832/1674/25984/3861/144501/667
gnal=35%","4208/8826/25932/3707/1393/1906/3726/5641/3309/2353/341359/7474/8644/144402/
3%","9645/5066/9023/2329/4846/10826/26002/51302/9420/2328/8644/6916/2327/3162/2326/154
signal=26%","23476/6872/5932/637/6198/59343/1874/1977/252983/55626/7015/245711/835/144
221656/23522/56848/5619/26122/7994/84656/8295/57634/8089/10902/86/6944/10856/54799/55
247431279847945,4834,"tags=57%, list=27%, signal=42%","83860/2957/6881/29079/55832/9282/2
ignal=47%","683/27202/3554/7130/1906/391/728/23604/972/2526/719/64127/3576/83700/3071/
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","22925/9429/301/3688/5321/81031/26503/5205/183/56606/477/624/51700/5660/2697/1906/89
nal=45%","306/9948/6814/51135/1511/1991/3569/2150/3684/6688/54210/6850/5724/57126/4615
3%, list=20%, signal=45%","10788/4853/7010/90952/1436/7082/2242/51225/3059/9535/397/10411
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st=21%, signal=50%","183/2771/5580/5446/7040/7305/64581/2150/3684/1535/6850/57126/3689/
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601/10749/4478/8724/2013/55614/7813/1203/9341/3916/83547/2060/400/5962/55062/120892/3
nal=30%","22925/9429/301/5321/5205/624/1906/6578/5322/11309/345274/5244/8273/123264/10
33/4208/196/5580/604/4067/23529/4773/7099/1880/958/51176/3586/4332/975/7305/972/5074/3
=74%, list=24%, signal=56%","183/2771/5580/7040/7305/64581/2150/3684/1535/6850/57126/3689
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%, signal=34%","7048/4323/2737/860/2303/1909/157506/7043/3198/5396/3199/7020/7042/652/1
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30,"tags=61%, list=26%, signal=45%","301/861/114548/604/8767/80149/3566/51043/3122/64332/3
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=28%, signal=52%","9469/56548/337876/29940/22856/83539/79586/55454/166012/50515/54480/(
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s=36%, list=27%, signal=27%","3556/2059/90102/51429/389/6281/1969/11346/11135/81565/1481
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2/9510/3624/7079/5066/8322/4846/7043/2099/7130/4327/5241/7042/3309/1958/4313/2662/303
","6788/7049/302/3673/1601/6281/117854/3433/9236/6672/3280/57326/3077/5155/567/857/284
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%, list=15%, signal=22%","7098/84159/2/8531/367/4323/7026/5154/283/5872/9510/3624/8322/10
=20%, signal=24%","51429/677/1601/2308/117854/344558/152503/4780/5587/4478/10769/28567
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5,3418,"tags=33%, list=19%, signal=27%","55970/51429/858/8826/57664/2791/59345/2771/10979/3
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ags=35%, list=15%, signal=30%","302/1601/3077/567/284/967/6401/5054/26119/5538/1191/6424/
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, list=12%, signal=32%","9445/81031/25825/1839/1471/6646/3479/9528/90865/6934/9452/123264
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28%, signal=63%","3688/3678/8754/255743/9780/5420/23075/3676/1803/83706/6352/10563/100/
8%","6625/51631/55660/55015/55692/10291/6637/8175/25766/51747/51690/6631/11338/6628"
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3925/79848/56155/8452/11113/116840/9727/9820/5300/26073/57448/91782/84501/54784/1444
191/1880/7474/10344/5339/5294/284114/6348/6558/102/6352/3627/8740/10563/729230/6373/6
%","7414/3688/1969/84962/9459/1901/29984/123720/10163/64805/5756/221178/51225/64083/1
%, list=19%, signal=43%","1601/183/654/185/114897/2587/5443/1408"
:24%, signal=43%","301/114548/604/8767/80149/3566/51043/3122/64332/3123/942/9655/7292/3
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3860/25898/3329/4194/2932/23368/1020/1387/55602/51499/56254/23476/6872/9100/8312/1128
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signal=39%","7798/367/1956/157506/652/7474/653/1499/23213/8626"
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st=13%, signal=39%","831/7057/120892/9528/5054/114769/83596/1522/7850/5621/2619/2734"
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list=16%, signal=22%","64764/2004/4774/169792/2120/4208/2308/367/147912/2908/861/26508/4
tags=83%, list=28%, signal=60%","861/3077/3133/3105/3659/51043/29126/10859/3071/865/56253
5183/200424/890/1063/2146/10951/6790"
signal=36%","9469/56548/81031/2300/10090/3479/6934/337876/1499/29940/22856/83539/7958
5%","3685/1969/23129/10979/84168/9780/26499/7408/2191/3690/1464/10630/81704/3737"
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ags=59%, list=19%, signal=48%","3280/2022/5155/1909/4851/93649/30008/23493/4773/6648/2627
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%","5442/116447/64216/80298/60493/54931/9361/22907/55794/6742/22868/6472/292/26073/30
79971/8322/4919/8321/8324/388630/6422/8323/6424/11211/2535/164284/8434/8325/2487/1159
%, list=8%, signal=54%","3913/182/8826/4643/4853/5420/652"
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=78%, list=19%, signal=63%","306/6814/11027/7305/3684/6688/2207/6850/5724/57126/4615/3689
%","117583/8531/11346/122786/54532/57477/5872/84952/5376/23586/387/3801/58494/54587/9
21%, signal=49%","10351/857/5468/5360/19/4792/54209/1066/348/7376/6256/4035/335/80820/5
al=33%","56034/3910/63976/3953/857/6778/2303/216/1051/4851/2697/51393/7099/3566/338557
list=31%, signal=33%","56034/3910/63976/3953/857/6778/2303/216/1051/4851/2697/51393/7099,
60%, list=22%, signal=47%","1601/3269/2246/5028/654/6720/5031/1581/5562/5745/5724/553/662
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,3923,"tags=74%, list=22%, signal=58%","6499/11340/118460/22894/5436/54512/129563/29883/57
ignal=27%","831/302/6281/4323/2934/861/4627/7057/7123/120892/9528/5054/10418/114769/83
ags=35%, list=10%, signal=32%","367/5154/183/6591/3217/30812/4070/3082/2099/120892/2668/6
733,2295,"tags=60%, list=13%, signal=52%","51429/152503/257364/8754/5045/348/7133/29763/35
ignal=47%","966/2/3075/710/11326/54209/718/1604/3553/725"
ist=26%, signal=39%","51429/183/4023/9388/3990/5360/185/19/6646/5322/348/7376/3931/344/8
%","54884/151056/216/157506/6121/220/9249/5919/8644/5467/57016/4035/1545/29785/125/33
=18%, signal=52%","54829/1893/4846/7040/4851/5167/23493/5552/54795/23462/2621/6662/1230
gnal=46%","23328/4616/4217/1843/3082/1647/8877/80149/652/5187/54209/768211/6688/7430/9
117248/727936/6484/10678/11227/51301/29071/93010/55568/2590/10610/6482/51146/2650/10
tags=55%, list=23%, signal=42%","114034/90806/9924/11340/22894/81875/54512/129563/29883/9
nal=31%","10413/7049/4208/7048/861/2735/2303/4851/23493/7042/10370/60485/2296/2627/968
","3074/8406/1203/25978/23120/55647/8546/83871/120892/338382/55207/19/5322/3257/7805/2

1%","3074/8406/1203/25978/23120/55647/8546/83871/120892/338382/55207/19/5322/3257/780
, list=23%, signal=43%","3356/351/26471/3479/8671/54209/55022/5027/65018/5562/4609/7114/9
ags=44%, list=26%, signal=33%","8826/1956/64285/183/84632/55614/9961/8412/1839/7040/9052/
=54%","11340/51013/118460/22894/54512/129563/6832/87178/9567/51010/56915/5393/23404"
55%","8767/245937/728/54209/3569/11213/64127/4615/820/3663"
/3913/3688/1969/81565/8301/6387/334/2737/7070/2050/4038/2042/3908/1889/4853/91653/230
%, signal=50%","80024/7474/6347/284114/6348/2185/6354/102/6351/6352/3627/11240/8740/105
","22925/9429/301/3688/5321/26503/5205/183/7057/477/624/5375/2697/1906/89874/5468/2350
=96%, list=38%, signal=59%","54913/51367/55505/92345/138716/9136/9410/10199/10799/10556/1
57782716,3012,"tags=52%, list=17%, signal=43%","3077/284/604/7043/7040/7042/684/3586/11213
,"9429/493869/4780/6649/10516/216/4846/1728/5742/5743/4258/2878/84203/6648/1193/348/42
,"57647/23353/23588/10456/2869/3756/10434/7341/163087/6434/79041/22932/1861/64783/865
.58219/4023/4481/5468/19/6646/54209/1066/5467/7376/338/9619/5465/948"
966/22918/11326/84466/5806/3684/721/629/720/3689/811/1401/731"
ags=30%, list=17%, signal=26%","23500/3280/7070/4038/567/5978/56920/10512/4851/58190/1495
701424269,2888,"tags=25%, list=16%, signal=22%","64764/2004/4774/169792/2120/4208/2308/36
42%, list=24%, signal=32%","23328/117854/3140/5359/3133/55601/9111/3916/81844/654/3428/23
ist=15%, signal=31%","135228/677/3280/7538/4851/3204/23493/5583/652/54206/28514/6424/229
t=22%, signal=28%","4780/81031/26503/56606/8682/2719/358/6814/1909/1906/5167/3479/8013/
list=26%, signal=32%","3556/683/7098/3074/2632/4124/141/3554/4121/1486/1203/4758/2720/57
nal=28%","85453/165918/2547/1196/64421/55693/2002/64782/7520/5810/5883/64710/25788/59
","11201/25898/7335/7334/51455/10714/83932/7336/26073/9100/55339/201973/56852/4436/54
=28%","84159/2/4323/7026/283/9510/3624/8322/10468/4846/1051/2099/4254/7130/26471/4327/
nal=38%","4208/301/3707/4254/1514/80301/3569/5291/23601/64127/5728/3665/350/3661/1234"
6%, list=12%, signal=32%","9445/81031/25825/1839/6646/3479/9528/90865/6934/9452/123264/11
%, list=13%, signal=27%","79971/10413/23189/1601/57664/23500/1956/57326/2719/857/9839/189
gs=49%, list=20%, signal=39%","3133/23586/4067/3566/6810/7305/2242/2150/3684/718/6688/316
st=21%, signal=37%","2/10769/4853/3133/4851/5045/4067/3586/1991/2214/54209/975/3123/796
=36%","5327/687/367/196/2908/4478/6446/114907/358/8837/10365/6422/5734/8644/54206/2230
1%","5125/7026/3655/4154/2737/4038/2719/1889/4853/860/5066/116039/10979/1281/157506/44
125/7026/3655/4154/2737/4038/2719/1889/4853/860/5066/116039/10979/1281/157506/444/369
ags=53%, list=20%, signal=42%","2690/3487/367/3488/51232/3479/653/23236/4824/3489/9451/348
26%, signal=40%","183/4023/9388/3990/5360/185/19/6646/5322/348/7376/3931/344/8435/338/33
l=27%","3074/2632/5836/2683/2992/83468/285203/117248/727936/53947/6484/10678/54625/83
nal=34%","4208/182/90/3280/1910/6591/2303/30812/1909/157506/309/56920/2335/10512/4254/
","57088/161291/5205/5359/57194/388121/5580/10577/6342/23120/5243/5360/19/5244/348/55
23%, signal=49%","4208/1956/64321/7040/93649/23493/7042/8324/652/28514/144165/22943/262
nal=33%","54884/151056/216/157506/6121/220/9249/5919/8644/5467/57016/4035/1545/29785/1
","558/4208/301/10461/355/3707/10019/4254/4067/1880/3569/972/7128/5142/5291/1831/5027/
al=29%","6788/7049/302/1601/3433/3077/5155/567/857/284/5580/27185/23129/10763/351/1516
/6601/8110/6599/5977/26039/6598/51412/6605/6602/6597/8193"
=27%","4774/558/10981/8301/3280/1910/283/860/1901/1203/27185/30812/81792/387/1909/967
nal=31%","1465/10398/3688/25802/5159/9948/4629/8837/1906/11214/5239/5339/29766/7140/71
3%, signal=40%","3913/182/8826/4643/1910/4853/2303/1909/5420/652"
3%, signal=53%","183/1471/5468/7107/84959/6422/3569/7128/5027/3635/338773/57817/4982/79
"tags=65%, list=20%, signal=52%","55644/54931/64172/3028/112858/8270/80324/51654/51002/55
gnal=35%","7049/6444/2719/5159/23129/639/161742/4091/4851/93649/23493/652/144165/1117
=22%, signal=29%","558/3572/4208/2632/284/2730/10763/387/1728/7043/4929/1051/5803/26471
ignal=28%","3074/2683/2992/285203/117248/727936/53947/10678/11227/374378/51301/337876/
list=17%, signal=45%","56034/80310/5154/1956/183/5155/2355/358/5159/2335/2099/6277/3479/8

46%","26512/23016/55505/6618/6617/7702/4654/55656/10302/56257/92105/79871/23379/5559
42/7373/64856/151887/81621/54507/133584/3371/5212"
0%, signal=55%","10413/25937/3913/3280/30812/55366/130497/7484/6662/3976/7477/6772/749
","302/858/6281/2013/857/11252/23180"
al=40%","1386/6881/9027/79612/1387/8260/10847/7994/339983/9862/8888/79969/2648/6871/11
gnal=52%","57470/55152/51067/54938/23395/124454/10056/25973/23438/79731/57505/92935/5
%, list=33%, signal=52%","57470/55152/51067/54938/23395/124454/10056/25973/23438/79731/57
%, list=33%, signal=52%","57470/55152/51067/54938/23395/124454/10056/25973/23438/79731/5
%, signal=42%","6387/387/351/199/7474/3676/29108/8764/6364/81704/284114/102/6352/3627/8
22,"tags=55%, list=18%, signal=45%","6387/2113/9314/387/6484/6401/6403/1991/3676/3569/252
"7132/355/8797/4804/7133/8764/55504"
gnal=38%","3270/1906/23493/146/54795/552/7168/153/147/3784/5144/133"
tags=36%, list=22%, signal=28%","64764/4780/5587/81565/8724/3655/183/5906/2042/27185/8837,
881934150699,1923,"tags=65%, list=11%, signal=58%","90/7048/26508/2022/6591/4851/23493/70
list=34%, signal=54%","196/3077/56940/64127/2213/4092/3134/3135/10859/3071/3824/5788/35
signal=34%","2040/7402/493/477/4734/154043/4846/6548/6324/5348/6330/1756/5349/5187/1126
list=14%, signal=29%","7049/57493/2022/2303/1909/7040/4851/23493/56999/84525/7042/3398/1
ignal=27%","5321/5154/283/84962/5337/824/126969/5155/8871/3707/5359/4023/23167/23682/6
31%","3572/7132/2308/29995/4000/183/90523/493/6548/4851/1906/11214/8654/5468/23493/34
/25937/6788/26524/8463/80014/84962/7005/79817/60485/5606/126374/9867/154810/9686/700
7%, signal=35%","5321/824/126969/151056/154141/23682/63874/3990/8605/57153/5322/56994/7
196/100507436/3140/4853/9314/3105/4851/7805/3123/51225/84868/29126/6688/10628/3592/58
20%, signal=61%","8260/8658/6421/546/5885/8313/55869/10664/699/9700/1663/113130/2237"
9,"tags=46%, list=23%, signal=36%","51366/23347/2072/3297/22954/51750/4282/51616/56893/51
7040/729/735/730/4277/4814/732/3002/731/10670/4069/3001/3929/2999/5320"
474/2059/23189/81565/83478/857/26084/4070/5962/391/199/1436/64805/54869/3383/51225/14
=82%, list=24%, signal=63%","9111/8291/3430/54209/7305/84868/1524/22904/5336/6850/2896/1
signal=51%","28973/55168/6183/63931/60488/51023/64965/23107/51650/54460/64963/64951/51
1%, signal=51%","28973/55168/6183/63931/60488/51023/64965/23107/51650/54460/64963/6495
19%, signal=45%","1601/183/654/185/114897/4824/2587/5443/56729/1408"
38%, list=24%, signal=29%","23704/8682/7057/857/7779/477/4734/6814/845/28954/5350/2669/40
l=30%","23500/3280/2737/146433/443/9948/9580/30812/5803/4851/4067/5354/7097/1499/1605
24%","966/6856/26503/284612/567/5066/3133/3107/3105/120892/127262/89866/3106/79772/34
%, signal=56%","5893/55100/7334/317761/7336/5932/51444/56916/55218/84464/1763/641/8399
80%, list=23%, signal=61%","3572/3716/1511/3569/10630/5329/22915/3717/84034/50848/6352/2
40,"tags=46%, list=26%, signal=34%","7070/6548/30845/2149/5354/6376/3306/9104/81671/6786/9
7,3955,"tags=76%, list=22%, signal=60%","120892/90865/7097/7099/3791/6347/5606/9590/10135/
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3411,"tags=73%, list=19%, signal=59%","7057/10392/10875/972/64127/6556/2213/968"
signal=41%","64425/84128/54700/55011/55127/2068/51547/4691/11137/6872/9533/23160/5691
004/51285/5906/79785/387/5899/55207/387496/5912/5868/5908/5873/6016/7879/998/5861"
63%, list=18%, signal=52%","6591/81792/5744/652/1499/2202/5916/7483/5915/150356/6662/405
10445/55683/54934/51230/84148/284058/3054/11091"
ags=62%, list=19%, signal=50%","3656/7096/51135/7097/7099/3659/7100/11213/10318/10333/366
,2881,"tags=31%, list=16%, signal=27%","307/3656/26508/6672/3280/3399/4188/8553/9314/11454
3%, signal=29%","3913/861/1910/5376/30812/3691/9076/79628/4803/2065/2668/10397/10370/16
45%","3688/6387/58494/6484/6401/7412/6403/1991/3676/2526/1524/64083/2131/9435/56477/5
4,4212,"tags=55%, list=23%, signal=42%","1601/4478/8724/2013/55614/83547/5962/4642/5604/58
238,"tags=41%, list=24%, signal=31%","79026/3270/477/845/6548/28954/5350/2669/401152/30845
33%","7414/26578/83716/5836/2934/2992/9961/978/51646/2720/7130/1471/7077/10875/116844

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066/57459/5928/5931/57504/53615/9219/9112/1108/1457"
signal=23%","301/5321/2687/5166/493/1806/11343/23743/6342/216/157506/23498/4023/27306/9
.0252/2246/2668/652/7474/1499/7481/22943/2247/650/6495/3227"
40%, list=23%, signal=31%","5166/4846/3356/351/26471/3479/8671/5957/54209/55022/144811/50
8801/389/57381/10981/85004/5872/25923/6237/30844/30846/51285/5906/2771/84617/51655/26
nal=35%","5321/283/4023/150763/7376/718/9104/6720/23659/3949/9663/51196/64900/7069/392
nal=35%","5321/283/4023/150763/7376/718/9104/6720/23659/3949/9663/51196/64900/7069/392
50630298716384,3442,"tags=70%, list=19%, signal=57%","55729/51535/57343/9869/22880/10155/1
13%, signal=44%","23560/146059/84083/3980/91603/7270/29980/54962/675/51053/26271/81620/
558/301/10461/3707/10019/3569/5142/5291/83700"
9%","4124/5154/3488/7043/93649/5241/84525/3202/652/3791/6943/54206/2627/2324/9394/316
st=19%, signal=49%","3678/9314/2246/3716/3586/7082/28514/6271/221074/168667/2247/63923"
%","4240/375033/10417/5322/54209/64581/4153/84466/5806/721/81035/725/8547/399687/2219
nal=34%","3913/81565/7070/9314/3082/347/5396/1495/8578/5604/4313/1605/706/2316/4099/37
8082/6641/6444/7402/6442/1756/1605/5239/8910/6645"
gnal=45%","2908/4306/2099/5241/8013/5467/6256/6258/9572/85315/81285/11057/5465/2852"
2%","64848/904/6881/284695/54623/274/54700/905/23248/57466/84337/27161/23476/23160/58
5%","29777/10799/8602/10556/27340/54974/54931/10775/3028/79897/11102/10607/10940/1614
=28%","4208/10981/1910/2737/9839/967/23682/4254/4286/5167/56999/6648/5873/10159"
/7169/29766/7140/8736/7168/7170/56203/7139/7134/7136/29767/343263/4607/442721"
nal=56%","100507436/3133/3105/3106/84868/1524/3965/2213/3134/5272/3135/10859/3071/387
, signal=55%","3673/1282/1284/1287/1277/22801/84959/6850/3717/4921/1288/126014"
gs=85%, list=25%, signal=64%","24144/2072/51750/9937/64421/55183/64858/65057/7515/7014/32
","302/2200/4853/604/10365/2313/351/10019/10163/7097/6886/10370/7305/10006/29766/5305/
%, list=20%, signal=26%","207/57159/8260/84101/8658/51750/23332/25876/1496/4331/8195/392
=9%, signal=24%","308/307/302/301/5321/311/306/309/310/8605/8291/57488/341359/144402"
2763,3222,"tags=59%, list=18%, signal=49%","2113/387/6484/6401/6403/1991/3676/3569/2526/94
477/81857/23097/9443/29079/55090/9282/9441/10001/9968/9862/80306/1024/6837/9969/21954
40/7334/26747/84280/66037/55127/27301/51447/51386/5987/79840/55269/118487/7150/9618/9
=18%, signal=36%","493/857/477/4846/6548/5350/4635/817/4634/153/133"
tags=51%, list=23%, signal=39%","255374/79042/8602/57523/55720/10556/27340/84294/10595/20
=46%","558/6387/4067/2621/3965/10859"
t=16%, signal=46%","558/6387/4067/2621/3965/10859"
19%, signal=67%","3685/4052/2615/5045/2627/375387/3696/4054/3694"
gnal=36%","56034/80310/1956/1290/824/1278/1307/1282/1281/2730/10365/1051/1277/4313/64
gs=49%, list=21%, signal=39%","4739/183/84632/5155/146433/1839/2065/1436/10006/5649/2621/1
230306181966,2652,"tags=75%, list=15%, signal=64%","10694/10575/10576/7203/22948/908/55135
368771194679,6039,"tags=74%, list=34%, signal=49%","3077/284/604/7043/7040/7042/684/3586/1
25%","5125/4154/2737/4038/2719/1889/4853/860/116039/1281/157506/444/10512/5396/4851/2
5125/4154/2737/4038/2719/1889/4853/860/116039/1281/157506/444/10512/5396/4851/2697/70
%, list=36%, signal=56%","7040/10865/8877/51043/3569/23180/64127/5008/4615/149233/3558/99
ignal=46%","4208/4038/1893/4091/4851/5167/5552/6422/54795/22856/22943/4092/6662/1230/6
=61%","1306/57493/25932/80781/26084/8985/4855/6781"
signal=33%","677/8531/4638/80303/358/624/5743/59341/5269/6517/54206/92241/5971/706"
nal=47%","994/7272/10403/9133/993/995"
gnal=52%","6387/2920/6376/2919/2921/3576/83737/3627/5197/6372/10563/6373"
0610773348030012,2802,"tags=77%, list=16%, signal=65%","7158/23347/56893/55183/2187/9400/
7907980778,1141,"tags=36%, list=6%, signal=34%","3487/367/3488/3479"
843/93664/6284/301/5872/4627/5906/2771/306/5978/9341/6385/3916/6814/3356/1393/5028/48

nal=51%","23244/27127/23126/8260/8658/546/23383/54921/5885/9126/9985/91272/991/79075/
%, signal=43%","302/6342/9388/26119/5244/7376/154664"
st=14%, signal=43%","302/6342/9388/26119/5244/7376/154664"
33%, signal=36%","3099/10105/345778/255027/6687/293/28978/10449/10939/664/7417/572/8426
L8239115,4078,"tags=91%, list=23%, signal=70%","56937/4091/5468/753/22943/11117/4035/4092/
'10765/55818/9682/3720/25974/79831/65094/8242/8846/23210/55693/51780/54784/84678/2303
al=43%","55214/8974/10491/5033/283208/64175/54583"
ist=39%, signal=44%","4719/4709/4708/4698/4714/4700/4696/4695/4720/51079/4710/4704/4713/
,"5872/1909/1906/5962/120892/9465/133558/8736/9590/7430/9867/154/10268/5141/2641/799"
%, signal=41%","302/4481/9388/3990/57552/29116/6646/26119/54209/348/338328/3162/4043/398
signal=22%","301/5321/2687/5166/493/1806/11343/23743/6342/216/157506/23498/4023/27306/9
5/64431/10847/81611/23421/401541/387103/4678/6944/55355/11339/54069"
37%","117854/6672/10581/10410/9023/81844/3716/3454/3434/7474/8519/9636/11274/4938/493
2%, signal=24%","5321/5154/84962/5337/824/126969/5155/8871/3707/5359/5159/151056/15414
145/84159/8826/3688/7057/9124/6385/7040/59/5339/79899/9590/7763/5371/8692/4921/2247/2
ist=15%, signal=50%","8082/6641/6444/7402/6442/1756/1605/5239/8910/6645"
=24%, signal=31%","23704/857/477/4734/845/28954/5350/2669/401152/1193/817/7881/130497/6
13/7431/351/3459/4919/5354/7099/728/54209/3569/10319/712/28514/4035/3553/3949/6279/95
, signal=42%","90/7048/26508/2022/6591/4846/4851/23493/7042/653/23462/4487/6615/94/4488/
'55214/6444/79073/7057/3270/51655/845/444/5350/813/3710/3708/6517/3709/202333/6271/81
31/3556/4774/2239/3913/4208/85461/11346/7058/4038/8404/5978/27286/27185/1282/4734/231
, list=26%, signal=25%","302/3685/25932/23704/79073/1192/358/309/6324/8542/55107/2697/377
=31%","7049/84159/57493/4638/2022/1901/2303/1281/1909/7040/4851/30008/23493/7042/8321
al=28%","64764/5611/23471/4780/90993/11080/7057/871/9531/26353/9709/8988/467/10133/330
signal=49%","3280/5159/30812/2668/652/1499/55366/6299/7484/6662/3976/6772/7490/26585/5
28%, signal=50%","861/3077/3133/3105/3659/51043/944/29126/10859/3071/50856/865/56253/8
, list=17%, signal=51%","307/301/9314/602/3586/1991/10333/671/945/7114/284297"
'302/2200/4853/2313/10019/10163/6886/7305/10006/5305/79705/56975/7066/4057/10859"
al=49%","2634/3566/602/958/3586/64581/115361/6556/163351/3394/4615/1130/388646/116071
gnal=44%","10865/4851/80149/51043/64332/3569/64581/3570/6774/4092/10758/9466/3594/237
%, list=27%, signal=42%","7098/8767/7097/7099/3659/7805/958/64581/23236/7292/64127/3965/5
al=55%","60493/54931/79072/22868/3028/6832/79675/80324/51654/87178/9238/55006/57570"
%, signal=39%","5587/5155/5159/388121/7040/7010/4067/64805/975/64127/2321/2261/998/515
gnal=39%","1909/4846/140/10487/5957/84876/55366/6786/23479/153/154/49/10486/796/1816/2
=40%","308/307/302/301/306"
, list=17%, signal=40%","8531/857/329/8837/26471/427/841/57162/8737/11035/330"
, signal=36%","117854/6672/10581/10410/81844/3716/3454/3434/7474/8519/9636/11274/4938/4
signal=65%","8899/1207/55696/84991/6433/55660/55015/51729/58509/11168/10291/25766/1352
","10413/3487/3490/1601/1410/4323/1956/81565/6387/85004/183/64321/58191/10516/6446/36
%","302/3685/309/117531/51393/59341/22821/3710/3708/80036/3709/93589/84876/54795/1408
gnal=47%","3673/2152/6387/1901/7424/3791/2150/4908/3576/7857/4088/3603/1012/5228/1233/
=19%, signal=47%","9804/140823/131118/728489/387990/51025/10469/29090/92609/10452/1024
501/90/7048/1290/90993/56937/1278/10979/1281/28984/7041/4091/8091/93649/5468/2353/103
signal=60%","23395/1968/27000/6193/57505/80222/57176/51654/80755/283459/92675"
/54496/56154/53615/10419/1789/9219/79813/8493/57673/3070/10664/10155/1786/1788/142/21
24/54496/56154/53615/10419/1789/9219/79813/8493/57673/3070/10664/10155/1786/1788/142/
nal=24%","3074/259230/55304/285362/5587/166929/10351/5580/4758/53947/2720/5660/30815/
signal=53%","23198/23140/11176/4297/138474/54454/23613/4300/55023/80764/23476/6872/104
39%, signal=53%","23198/23140/11176/4297/138474/54454/23613/4300/55023/80764/23476/687
"301/861/6711/7538/947/8767/2633/11326/7805/942/64581/7128/5142/84868/9454/7430/5621/

signal=40%","301/861/6711/7538/947/8767/2633/11326/7805/942/64581/7128/5142/84868/9454
st=21%, signal=33%","6284/4067/8013/8773/3566/284021/2242/5169/5294/3162/3965/8673/6850,
al=58%","158219/4023/4481/5468/19/54209/1066/5467/7376/338/9619/5465/948"
1%","3673/6387/1910/11343/6548/3356/5028/55107/1906/56925/5743/2149/7432/3586/6347/32
%, signal=40%","183/4023/9388/3990/5360/185/5322/348/3931/344/338/335/341/4353/9619/336
%, signal=58%","3269/3707/5028/23236/5336/5031/5724/2185/2852"
=74%, list=30%, signal=51%","28984/4851/654/7474/7082/10076/4855/9647/3169/201627/10211/
gs=61%, list=23%, signal=47%","7048/5154/5396/7474/3791/1499/8646/6662/4609/2294/6495/677
","1435/1910/7057/79148/3958/146433/59341/5919/728/1436/6347/719/1240/1588/6348/2185/
%, signal=25%","5932/637/6198/59343/1874/1977/55626/7015/245711/835/144455/29117/25799/
gnal=37%","3572/3280/146433/443/30812/7040/5803/4851/5324/4067/23529/5583/7097/1605/33
tags=42%, list=27%, signal=31%","966/4121/51226/871/2335/123720/79056/9780/4924/6892/3309
%, signal=42%","1601/8301/3339/26119/348/5649/1191/22943/4035/4043/338/335/1401/336/7184
gnal=38%","2737/2735/4038/3219/1499/22856/28514/2736/8626/3206/3226/1745/130497/1746/
/3685/25932/23704/79073/1192/358/309/6324/8542/55107/2697/3779/9424/9780/117531/51393
790,"tags=56%, list=16%, signal=47%","54984/100134934/3183/6714/3178/55226/65057/3192/701
list=15%, signal=48%","347/59341/64581/4938/9655/64127/3965/3553/6850/4940"
1,"tags=56%, list=15%, signal=48%","347/59341/64581/4938/9655/64127/3965/3553/6850/4940"
","7538/57007/9034/1524/1238/1230/1232/1234/3579/2532/729230/10663/1235/51554/1233"
tags=87%, list=26%, signal=64%","29777/10799/8602/55720/27340/84294/23212/9790/6227/65095
205,0.000729233242041282,4700,"tags=87%, list=26%, signal=64%","29777/10799/8602/55720/2734
15%, signal=33%","11105/196528/5932/55183/2187/8295/57634/8089/10902/51010/86/2521/799
","255374/114034/79042/90806/57523/9924/10556/10595/201164/80746/22894/81875/54059/27
al=64%","6742/30968/9093/201973/3980/11232/92667/1763/2021/6240"
9%, signal=55%","2308/4000/90523/493/5468/54206"
1560,"tags=60%, list=9%, signal=55%","2308/4000/90523/493/5468/54206"
79642/340075/412/153642/23213/411/2588/55959/410/22901"
nal=41%","1465/10398/91624/4629/444/7171/1756/7169/1605/4604/5339/8736/4634/7168/2333
6","56159/10609/27127/171482/57820/4292/27030/317761/10734/5885/9126/9985/675/7329/256
%","56159/10609/27127/171482/57820/4292/27030/317761/10734/5885/9126/9985/675/7329/25
gs=75%, list=16%, signal=63%","10417/7099/7474/7805/972/51704/5336/3135/10859"
35%, signal=48%","1153/56853/3167/4928/6625/283375/29896/54623/170506/6434/3297/246744
100%, list=29%, signal=71%","56159/164045/254528/1325/2072/4292/197342/80198/57697/84464,
3/8531/1969/11346/54532/5872/84952/5376/23586/58494/54587/2697/90952/1003/9076/7408/7
%, signal=57%","4780/9314/7043/10365/4091/19/445/4092"
11,"tags=61%, list=28%, signal=44%","2074/7428/51497/6829/904/54623/51574/8621/9025/56943/
4276,4975,"tags=63%, list=28%, signal=46%","51079/581/55750/4836/158234/140823/9141/285521
'10392/3105/7096/3106/7097/3123/10333/64127/912/284297/1755"
30%","5321/11343/151056/4023/9388/3990/8605/5322/1373/23236/338328/5294/23659/3949/34
4238,"tags=43%, list=24%, signal=33%","1909/154043/6548/1906/5243/30845/1756/11261/54209/6
8658/51750/10111/100134934/9937/5609/64421/7874/64858/65057/3184/4751/7515/5591/7014,
%","7048/871/860/85301/1893/5744/23213/22856/5916/4488/9451/2131/4092/6662/8817/25987,
332,3664,"tags=52%, list=20%, signal=42%","3487/367/3488/3479/653/4824/3489/3484/8483/650/
ist=30%, signal=53%","51287/116228/6834/100131801/6341/79072/285521/51241/493753/51204/
ignal=36%","301/7048/688/928/8837/6442/2697/3479/84525/8324/1605/975/5467"
signal=24%","10413/4208/90/8463/7048/3280/2022/64321/1889/4853/9645/2303/1909/4851/269
%, signal=40%","2239/1601/57664/23500/2719/8322/387/84870/10082/4919/8321/8324/166336/7
9%, signal=39%","51455/7486/10714/83932/26073/9100/56897/201973/5426/5983/5425/51514/51
%","3280/5205/2737/3198/7020/2065/3200/10370/57007/6507/1499/1959/104/6299/9935/10312
%, signal=44%","3572/3280/4851/23529/1600/54209/3398/3569/3400/4099/3949/3976/9283/9572

),"tags=39%, list=24%, signal=30%","9697/6238/259230/23471/166929/285671/567/3133/3107/310
)81/1910/2737/6591/9839/1896/967/23682/4254/4286/8546/5167/63971/8934/56999/6648/3257
list=26%, signal=37%","1601/10252/1848/7078/154043/161742/81848/4067/2633/467/57447/709
=20%, signal=44%","1435/3624/146433/7040/3398/3059/714/841/3976/8737/10221/5925"
35%","7779/4502/4499/4493/3309/4489/4501"
50%, list=8%, signal=46%","5592/928/5580/10019/5962/114897/84959"
ignal=23%","3074/259230/5321/5154/84962/166929/5337/824/126969/5155/8871/10351/3707/53
/7122/4155/4099/400746/153562/51090/83700/5728"
, list=31%, signal=37%","1997/7070/3958/2633/8934/7805/975/56940/5074/8915/7430/5621/1038
%","4208/1910/2737/9839/4254/4286/5167/56999/5873"
25%","11031/8801/389/5592/57381/10981/85004/5872/25923/6237/30844/30846/51285/5906/27
ıal=25%","11031/8801/389/5592/57381/10981/85004/5872/25923/6237/30844/30846/51285/5906
ıal=32%","677/3280/3726/84870/23493/9021/5604/51176/3586/653/23462/841/9394/113026/676
%, signal=51%","4208/4780/9314/7980/10365/5743/4313/5339/3037/9655/445"
573/183/23371/1284/2730/284207/1906/654/6648/4313/3586/3569/5467/847/9365/1535/825/21
337/116236/2151/11343/151056/624/1909/4023/63874/79887/9388/3990/57552/5333/8605/196
,"51574/118460/54512/7737/345630/2091/51010/56915/23404/1736"
27%, signal=62%","4869/3692/84946/6059/6209/23212/55341/55153/4927/7514/5901/161424"
,"4869/3692/84946/6059/6209/23212/55341/55153/4927/7514/5901/161424"
4/79831/65094/8242/8846/23210/55693/80312/51780/6996/200424/54784/84678/23030/23133/9
%","54901/80224/317671/7386/6059/55316/11019/4595/441250/51750/64789/2068/3658/374291
54901/80224/317671/7386/6059/55316/11019/4595/441250/51750/64789/2068/3658/374291/472
list=26%, signal=30%","10769/3908/351/8878/11170/5743/5641/27020/3710/6507/6810/6347/730
059/2934/25802/6840/6711/5962/5756/29766/6708/11344/832/85477/56203/829/2314/830/2976
98/8531/5836/355/857/329/4217/197259/8837/26471/114908/427/706/841/57162/4814/8737/11
58%, list=22%, signal=46%","7049/7048/2022/10979/7043/7040/4091/116844/7042/4092/25805/4
=32%","2934/6444/23704/857/477/6548/6324/5350/2697/6330/3752/54795/5142/2702/2316/182
25,1430,"tags=44%, list=8%, signal=41%","1435/183/4023/4481/185/5583/5322/4790"
","1655/57060/8087/90806/26156/7072/11340/66037/84294/170506/6201/7073/10642/54512/56
)%, signal=42%","3624/604/4067/6422/7805/3586/64092/4332/3398/7305/7128/5074/3635/2213/1
32%","54514/10146/9271/167227/26523/80336/4848/6191/23112/11201/9898/10922/64848/9118
gnal=69%","2013/5155/2149/2150/2702/2621/10391/1535"
%, signal=64%","10694/10575/10576/7203/22948/908/55135/10574/1736"
signal=64%","10694/10575/10576/7203/22948/908/55135/10574/1736"
,"tags=61%, list=7%, signal=57%","29922/83549/54963/4833/790/7371/1503/56474/4830/1841/729
ıal=29%","5872/30844/30846/567/9341/3133/3107/54918/3105/30845/3106/10133/8324/10397/2
list=27%, signal=43%","6419/29104/58508/5929/8085/23067/10322/64324/11105/9757/54093/84
/967/1513/351/1514/4803/5641/1508/1520/4043/3592/5618/4012/3717/8572/338"
signal=29%","4208/90/3688/4000/6444/183/5159/4629/2274/6548/9096/7040/1906/93649/11214
ignal=44%","966/3075/5873/718/1604"
gnal=32%","29995/493/10519/2274/6548/4773/3479/2065/11261/64581/9454/5621/5533/1827/5
44%, list=23%, signal=34%","301/5468/5743/8013/5467/7376/972/552/9104/3553/344/335/8660/5
%, signal=31%","8872/1981/51247/57409/66037/1975/1965/6199/5436/4528/8667/8892/27161/88
)2014/125965/6834/664/4720/1350/4710/3421/9650/1352/29796/23410/1347/4704/27089/2271/5
, signal=24%","10413/6788/26524/7049/2690/4208/8531/367/7048/861/1435/81565/6387/1848/8
79%, list=28%, signal=57%","2737/860/654/94/4017/7704/4092/6662/8817/55553/4088/392255/91
ist=23%, signal=55%","55818/9682/3720/55011/22823/4302/55693/55170/54880/6598/1789/2302
55252/9898/6045/84108/6015/3720/8535/171023/23512/84759/10336/3066/1911/7528/5928/770
1%","114034/90806/9924/22894/81875/54512/129563/29883/90459/23019/79650/112479/22803/
=57%","7123/7078/1893/5919/3699/3959/350/10447/55664"

34%","2690/3488/54741/6778/3716/4067/1373/6774/5829/445/6776/9518/4609/3717/5295/3158
%","3488/10468/4851/7042/1499/4804/2252/6615/2736/8626/2131/94234/342574"
385,"tags=32%, list=22%, signal=25%","64285/29969/6672/8724/284/9531/5978/347/120892/2149/
"26578/1793/57088/395/57514/6451/84632/134265/8871/8754/5359/9467/2060/4067/10163/740
36%","558/54843/302/7094/4240/7057/309/6121/341359/79792/54209/2621/950/115548/57158/
%, signal=49%","3913/5155/5159/947/1958/6943/55366/130497/5156/7490"
9%","10413/6788/26524/7049/4208/8531/7048/861/1848/80014/2735/2303/4851/1906/23493/34
st=28%, signal=43%","55290/83860/9014/6881/64425/54700/55832/2958/55011/55729/6878/6879
%, signal=52%","51366/55100/7334/56254/78990/9810/55611/23028"
list=21%, signal=63%","7057/1281/4060/5045/7305/6376/10159/2627/375387/23607/25987/8710
s=48%, list=13%, signal=41%","4208/5592/4780/3488/387/93649/5054/199/5467/4035/7168"
54%","302/858/6281/2934/2013/857/91663/11252/23180/5538/1794/4118/4864/5788/10211/173
gs=48%, list=19%, signal=39%","161291/89866/123264/975/55754/27248/9695/10956/3638/55741/
3189/81565/83478/857/26084/4070/5962/391/199/64805/54869/3383/51225/1464"
ags=94%, list=34%, signal=62%","5321/3554/10865/8767/6556/3553/8809/9466/3594/149233/7292
ignal=56%","10413/10252/2246/157506/2099/652/7474/10253/6662/6495/374/51804/6469"
43%","10226/4023/5468/5167/54209/3553/51099/123/5465/57104"
7%, signal=51%","6829/904/23144/10914/54623/170506/3297/91746/56254/7936/11052/11051/1
t=26%, signal=35%","966/2239/7070/2719/10392/2350/10082/51299/929/1942/338328/5621/6412
9%, signal=42%","966/567/3133/3107/3105/89866/3106/3122/3115/3123/972/3113/9871/6720/312
367484,0.00095166341060567 7,2796,"tags=40%, list=16%, signal=34%","3270/477/845/5350/1756/
301/9322/375033/7099/199/94134/84466/353376/26253/51411/9938/3936/10347/57192/1785/71
=31%","54884/151056/216/157506/6121/220/9249/5919/8644/5467/57016/4035/1545/29785/125,
2190,"tags=46%, list=12%, signal=40%","64114/5525/1601/8301/7040/2633/9528/55022/116372/51
21,5244,"tags=83%, list=29%, signal=59%","7428/51497/6829/51574/9025/165918/7936/8312/1091
ags=32%, list=22%, signal=25%","117854/1956/5587/493/857/284/8412/7040/351/5743/8767/4803/
%, signal=39%","117854/10581/10410/5359/10475/64135/4851/56829/8091/3428/2494/3434/684/
3,"tags=41%, list=17%, signal=34%","558/117854/1435/3554/837/81844/1906/54625/83666/8767/7
","6591/7043/54209/6376/712/713/3684/718/22943/714/1524"
7%","387758/8754/351/5743/120892/6648/3309/706/9451"
18%, signal=42%","1027/8816/26094/26133/25853/285429/80067/25879/1161/80344/1643/79269/
","4162/2303/947/64094/5054/3791/7128/240/6558/79625"
t=38%, signal=40%","4709/1329/54949/4708/1738/4698/4714/9167/4700/7388/4696/4695/4720/1
,"tags=64%, list=38%, signal=40%","4709/1329/54949/4708/1738/4698/4714/9167/4700/7388/4696
","3554/3077/567/114548/3105/10865/54795/975/3569/7292/7133/1604/91543/3553/4092/3134/
5, signal=37%","3554/3077/567/114548/3105/10865/54795/975/3569/7292/7133/1604/91543/3553
l=37%","7049/57493/2022/9510/1901/2274/7010/4851/23493/7042/653/23462"
521,2250,"tags=50%, list=13%, signal=44%","7057/947/4060/5045/5743/7481/6376/10159/2627/396
02/858/301/6281/10981/8406/8301/8724/5872/257364/857/9839/9341/6385/947/23682/25978/8
gnal=35%","117854/6672/10581/10410/81844/3716/3454/7474/8519/9636/11274/4938/4939/342
","7048/5604/3202/652/1499/7477/5594/340419/6469/5595"
643/9493/9055/9212/24137/29127"
9/81031/493/3953/477/5243/11309/6515/6517/8671/6507/387700/348/114134/53919/10060/652
signal=30%","9429/81031/493/3953/477/5243/11309/6515/6517/8671/6507/387700/348/114134/5
2%, signal=33%","4853/9096/4851/2697/23493/84525/652/23462/4487/3398/2702/9242/4488/262
0/6387/493/5155/3958/5159/7779/5350/51393/59341/84876/1499/6347/7223/5621/1950/7226/1
%, signal=52%","64863/9361/6742/30968/9093/7156/201973/3980/11232/92667/1763/2021/142/6
al=44%","183/4023/9388/3990/5360/185/5322/348/3931/344/338/335/341/4353/9619/336/5320/
, signal=44%","183/4023/9388/3990/5360/185/5322/348/3931/344/338/335/341/4353/9619/336/5
3/185/6646/348/3931/8435/335/341/9619/336/38/55004/134429/337"

/185/6646/348/3931/8435/335/341/9619/336/38/55004/134429/337"
"183/185/6646/348/3931/8435/335/341/9619/336/38/55004/134429/337"
214/5351/9023/23498/10826/8974/51302/9420/23474/5033/283208/2230/26234/5352/6916/6417
list=20%, signal=37%","2690/54741/6778/3716/4067/6774/5829/6776/9518/4609/3717/5295"
signal=39%","3459/93649/54209/348/1191/4035/7274/4043"
al=32%","10769/351/8878/11170/5641/27020/3710/6810/7305/348/5649/10288/4915/80863/2338
3%, list=14%, signal=46%","301/604/80149/3566/51043/9655/7292/3142/4092"
57%, list=17%, signal=48%","8531/857/329/26471/841/57162/8737/330"
signal=70%","23643/7096/7097/7099/54209/10333/721/64127/284297/1755/3929"
=31%","29995/493/10519/2274/6548/2494/4773/3479/2065/11261/64581/9454/5621/5533/1827/
%, signal=26%","10413/26524/2690/25959/1601/367/5547/2908/26508/56937/4306/7068/4734/38
t=24%, signal=33%","3280/7779/477/4846/845/28954/5350/2669/5743/401152/1193/10288/29763
9%","687/196/6446/3624/10365/54762/3779/1495/6648/6422/3709/1499/1066/1605/5734/8644/2
51%, list=21%, signal=41%","558/3624/7099/80149/7805/3586/3123/2621/7292/84868/29126/8038
"4627/3958/27020/3122/975/8417/3123/3383/84868/8915/7430/5795/3002/963/84766/56253/10
4208/182/3280/64321/4853/9314/5396/3199/4170/64641/652/1499/28514/4855/2627"
al=29%","7049/4208/182/7048/57493/26508/2022/2303/1909/7040/10512/4851/23493/7042/234
signal=33%","9429/493869/4780/6649/10516/216/4846/1728/5742/5743/4258/2878/84203/6648/
%","2022/23493/23462/4804/94/4824/54567"
"tags=36%, list=23%, signal=28%","51474/2059/90102/23189/395/1410/2934/84952/25802/6840/6
52%","3140/11027/10288/11024/11025/10990/10859/353514/1512/79168/51348/23547"
4722182813963,5244,"tags=82%, list=29%, signal=58%","7428/51497/51574/9025/165918/7936/83
%","7414/3693/3801/1495/5271/1832/1545/29119/7060"
291,3182,"tags=41%, list=18%, signal=33%","1997/7070/3958/2633/7805/56940/5074/7430/5621/22
=22%, signal=45%","80298/6606/57466/22803/7978/7270/10419/8458/1660/4150/51728"
%","23166/7057/4023/4481/3990/5360/8578/23780/54209/348/338328/3949/81035/335/1401/33
47%","23166/7057/4023/4481/3990/5360/8578/23780/54209/348/338328/3949/81035/335/1401/3
%","142940/83480/115939/26995/54517/54433/83448/80324/55651/27079/113000/1736"
=23%, signal=43%","1601/4478/8724/2013/55614/83547/5962/5604/5868/2316/7430/7879/64400/
.7,3954,"tags=69%, list=22%, signal=54%","3685/5468/19/4792/3690/7376/9619/1401/5465"
=22%, signal=40%","90102/395/182/4323/7057/604/347/3204/11170/5054/94/56940/5728/10395/
/6117/10714/6119/5558/5425/4172/5557/23649/5424/5422"
22/4153/5806/721/725/8547/399687/2219/1401/3929/722/2220"
8/27030/994/10403/26271/9319/9133/6790"
al=35%","677/84870/9021/51176/3586/653/6768/7477/133"
2/2934/3270/477/5028/5350/8654/5142/817/5144/3759/5025/6588/134/390594/2946/3778/481/
5%, list=25%, signal=27%","2690/3280/183/857/54741/7040/10019/4851/5468/1647/54625/9021/2
24%, signal=76%","64901/23039/8021/7514/56478/5901/23214/1434/57510"
,"5429/10728/11201/5423/27434/51455/10714/7015/201973/11232/7014/5426/10721/54107/542
list=14%, signal=33%","29995/493/10519/2274/6548/3479/2065/11261/64581/9454/5621/1827/53
069/6387/2737/2050/2042/3908/1889/4853/91653/23022/23129/91624/1909/56920/351/10678/63
=82%, list=34%, signal=55%","4297/80335/9070/6419/58508/5929/8085/23067/11105/9757/54093/
2%, signal=48%","143872/5592/10203/477/80005/8654/5743/80149/5294/154/150/134/796"
24/23022/5468/79778/4790/27295/1605/9260/9499/8572/64236/4892/129446"
039,"tags=47%, list=17%, signal=39%","301/861/604/80149/3566/51043/9655/7292/57045/29126/3
ignal=44%","4638/2022/1281/30008/25976/652/7763/130497/6662/6495/6469/5311/1739"
nal=59%","10413/25937/3280/30812/55366/130497/7484/6662/3976/7477/6772/5311/38/793/54
%, list=23%, signal=62%","80310/5155/2719/5159/8837/1958/652/3570/130497/4609/650/6772/20
32/5205/1192/8542/28962/55107/5243/5348/6560/5349/57348/2564/1193/6507/94015/5172/706/
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1%, signal=40%","3673/2152/6387/1901/7424/3791/2150/4908/3576/7857/4088/3603"
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l=24%","5592/23704/493/6446/857/7779/4734/28954/23678/6324/2669/3779/5348/6330/5349/37
/6434/60680/55285/10594/90459/6895/7884/3178/3191/11189/6631/7536/3192/3181/11338/10
al=60%","54840/254528/27301/64789/83940/9937/64421/5810/5883/55218/64858/92667/5426/3
533,4014,"tags=77%, list=22%, signal=60%","54840/254528/27301/64789/83940/9937/64421/5810/
"558/3685/301/4240/7057/10461/375033/391/7052/54209/6347/3690/7305/84466/2621/718/721
=47%","4638/183/1909/3356/1906"
gnal=46%","4208/351/1393/6347/3460/5649/65018/5923/6536/5924/2185/22871/3458"
ags=38%, list=24%, signal=30%","23704/857/7779/477/4734/845/28954/5350/2669/401152/1193/8
49%","25802/7171/7169/29766/7140/7168/7170/56203/7139/7134/7136/29767/442721"
8,"tags=53%, list=19%, signal=43%","29922/83549/54963/4833/790/7371/1503/56474/4830"
/9076/59341/6337/2629/7357/64834/5339/3037/7466/8626/51599/8692/2131/361/2312/388697/
signal=50%","3913/182/8826/4643/4853/5420/652"
16346857085188,1396,"tags=35%, list=8%, signal=33%","7049/1601/81031/7057/9839/7041/93649
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2/9510/3624/5066/8322/4846/7043/2099/7130/4327/5241/7042/3309/4313/5806/8204/2069/5150
list=23%, signal=28%","302/3685/1601/8301/3077/567/284/79890/967/6401/6386/5054/26119/36
, signal=36%","1956/64285/183/84632/9961/8412/1839/79628/5641/9052/54206/79651/2537/965
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'tags=74%, list=30%, signal=52%","6281/7070/1307/6385/3987/10979/7010/6422/3791/9647/8829/
=23%, signal=31%","5166/978/4846/3356/351/26471/3479/8671/5957/54209/55022/144811/5027/
, signal=25%","23500/3280/7070/4038/567/5978/56920/10512/4851/58190/1495/7474/3397/1600/
22918/11326/64581/84466/5806/3684/720/3689/811/1401"
8%","51082/10623/5434/55703/5437/171568/9533/10621/661/11128/55718/51728"
=53%","9510/4846/4327/5241/4313/8204"
2345,"tags=33%, list=13%, signal=29%","3688/1969/1956/493/2042/6711/3958/10519/8878/5583/3
9,"tags=54%, list=10%, signal=49%","7049/90/4052/7048/2022/8425/94"
).00119277858345228,4517,"tags=75%, list=25%, signal=56%","329/51296/7099/7128/330/57864/33
, signal=63%","80310/5155/5159/8837/1958/652/3570"
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%, list=15%, signal=35%","2239/1601/57664/2719/387/84870/10082/4919/8321/8324/166336/747
","54797/9439/10025/9477/81857/9443/29079/55090/9282/9441/10001/9862/80306/6837/21954
%","11343/151056/4023/9388/3990/1373/338328/5294/344/145447/51099/338/335/200879/13/1
%","11343/151056/4023/9388/3990/1373/338328/5294/344/145447/51099/338/335/200879/13/1
, signal=43%","55214/8974/5033/283208/64175/54583"
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.%","493869/4780/6649/10516/4846/1728/5742/5743/4258/2878/84203/6648/1193/348/4257/260
49%","558/3433/10581/10410/3454/3437/684/8519/2621/3455/4600"
34%","182/90/3280/1910/6591/2303/30812/1909/157506/309/56920/2335/10512/4254/1906/266
7011,2696,"tags=60%, list=15%, signal=51%","367/5660/4851/1499/8626/3169"
%, signal=64%","158219/5468/19/54209/1066/5467/7376"
st=20%, signal=57%","10524/9575/2033/138474/114799/23522/1386/6881/1387/10847/7994/9862
9%","54840/5378/4292/27030/5395/4595/4439/6996/4436/5111/2956"
=52%, list=23%, signal=40%","301/5743/7376/972/552/9104/3553/344/335/51085/60481"
t=19%, signal=35%","367/7026/5066/639/2099/3726/5743/5241/10370/23780/1508/5467/84868/4
5%","3280/5159/30812/2668/652/1499/55366/6299/7484/6662/3976/7477/6772/80144/7490/265

3%, list=13%, signal=29%","4208/301/4067/7474/3586/5291/5027/29126/3965"
3%, signal=48%","3656/3554/51135/8808/1958/3569/11213/23236/7444/7850/3553/4615/79155/3
al=30%","283/10417/3133/3105/2920/6039/5919/602/1511/1991/2919/6590/2627/2921/6357/641
,"tags=69%, list=20%, signal=55%","996/51433/1017/8697/8452/64682/51434/991/27338/11065/53
gnal=32%","79848/8452/11113/26073/9585/6683/994/9928/1894/9493/201161/9212/990/23636/2
gnal=40%","3673/6591/2335/7424/6386/22801/64805/1464/23607/8829/4690"
signal=47%","683/27202/7130/1906/728/23604/972/719/64127/3576/83700/3071/1803/7060/9055
5240/5584/11021/51335/7507/64780/9657/14/9094/252839/92400/56155/9727/11103/79650/102
6%","301/7132/5321/2687/2052/11343/27306/5742/1906/8605/5743/4258/9261/7097/5322/8644
2308/355/857/5159/1277/6648/4313/847/10533/10630"
0%, signal=44%","1601/5978/27122/654/27120/1958/653/8644/9563/1588/133/650/3293"
al=28%","558/3280/1910/283/860/1901/1203/30812/81792/387/1909/10365/4929/351/2697/190
38%","11343/151056/4023/9388/3990/1373/338328/5294/344/51099/338/335/200879/13"
2%","687/3280/4005/7068/2730/10634/1514/5962/2629/1508/1520/3625/717/1512/3249"
79875/339366/30008/8425/2201/4239/8076"
%, signal=45%","10361/1017/27154/6742/546/8467/11143/30968/144455/8317/80174/10926/799
%","80335/9070/259282/5929/23067/84661/143241/30827/91272/54554/84332/9739/3054/1109
,"1435/860/947/1906/652/4487/4804/5083/650/4982/340419/6469/80320/147495"
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ignal=43%","1386/6881/1387/10847/7994/9862/8888/2648/6871/11143/6872/9913/23326/9329/8
50%, list=17%, signal=42%","5154/6591/23129/3082/10512/7052/1605/2252/8829/3911/59277"
3%, signal=38%","4208/8531/90/3488/7040/2697/4015/2065/652/51176/4534/1499/84466/28514/
st=23%, signal=34%","90806/51594/66037/170506/55051/81892/10642/4904/79066/8148/23367/2
nal=35%","57088/161291/5205/5359/57194/388121/23120/5243/5360/19/5244/55754/51228/816
, signal=30%","135228/677/3280/7538/4851/3204/23493/5583/652/54206/5467/28514/6424/2296
75771789,4042,"tags=38%, list=23%, signal=29%","3270/477/845/5350/1756/1193/817/9446/6348/
238,"tags=41%, list=24%, signal=31%","23704/857/477/4734/845/28954/5350/2669/401152/1193/8
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, signal=34%","221908/2648/7514/7517/79959/7840/163786/55226/6491/55755/55835/348654/10
%, list=19%, signal=45%","7057/4790/3586/11213/1240/64127/4210/10859/51311/353514"
03/8289/6601/196528/6599/53335/65980/86/6598/51412/6605/605/6602/6597"
=24%","10413/7049/2690/4208/8531/7048/4323/1435/1956/81565/6387/58191/2735/10519/2718
gs=55%, list=28%, signal=40%","347/7099/80149/5187/54209/3569/64581/29108/6774/10333/7850
782,5018,"tags=55%, list=28%, signal=40%","347/7099/80149/5187/54209/3569/64581/29108/6774
033741,3606,"tags=51%, list=20%, signal=41%","170506/8658/51750/10694/10575/5609/7520/3178
ags=51%, list=21%, signal=41%","10769/3133/5045/4067/3586/54209/975/3123/79651/718/64127/
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284/4067/8013/8773/3566/284021/2242/5294/3162/3965/8673/6850/2294/5729/2268/11314/681
402/10174/1495/1756/1605/10391/5829/23336/4892"
%","64789/9937/22803/64421/22909/1797/64858/51692/9156/2237"
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signal=31%","6281/23704/857/10519/3270/6324/9709/1003/4067/120892/5348/6330/27020/3752/
8/367/10461/2295/2099/654/7474/653/1499/55366/8626/1588/7484/5156/3293/7490/6469/4920
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'tags=43%, list=21%, signal=34%","2040/7402/477/4734/154043/6548/6324/5348/6330/1756/5349/
nal=65%","857/4851/93649/4855/3553/3576/64083/81035/361/335/2650"

%","4651/4643/4627/4641/55930/79784/4642/4644/4648/4542/4637"
, list=27%, signal=31%","51429/389/6281/11346/11135/2934/148170/25802/9948/10174/6385/10
29,3039,"tags=52%, list=17%, signal=44%","301/861/604/80149/3566/51043/9655/7292/3142/4092
'23704/857/9948/477/845/6324/2697/6330/3752/2702/2316/3762/482/123041/3784/3759/10269,
=24%","10413/6788/10161/4071/677/8531/90/367/3688/4643/7048/7026/4124/57493/1956/1239
77%, list=35%, signal=50%","1153/56853/3167/4928/6625/283375/29896/6434/246744/22938/469
5%","2737/355/3958/604/4067/652/7474/3586/972/1831/155038/5027/1942/29126/8915/3965/81
t=22%, signal=48%","6281/7070/1307/6385/3987/10979/7010/6422/3791/9647/8829/5795/4088/4
list=20%, signal=24%","51429/677/1601/2308/117854/344558/152503/4780/5587/4478/10769/285
3%","1277/4313/6422/2201/1101/4488/22904/133522"
tags=52%, list=25%, signal=39%","1655/8087/55234/6431/7072/55225/6434/60680/64783/22827/9
t=20%, signal=47%","2934/8724/10581/10410/9023/81603/85363/8519/5806/4261/8547/11043/22
0%, signal=33%","10440/92014/29074/6576/10166/10939/11194/57380/3329/9131/55186/7352/9
=27%, signal=37%","3958/604/4067/652/7474/3586/972/1831/155038/5027/1942/29126/8915/396
nal=25%","677/2308/4000/4780/84962/355/857/358/28984/132864/947/6548/8837/4851/1906/37
nal=37%","7779/4502/4499/4493/4489/847/4501"
gnal=41%","5322/5919/64581/6688/8915/1524/5336/50856/26253/55223/245932"
, signal=31%","558/301/80310/4780/358/5580/6778/388/4217/1728/3082/10365/1906/120892/80
17%, signal=31%","10413/7049/4208/7048/861/2735/4851/23493/10370/60485/2627/9686/5728/2
signal=41%","1601/1910/5978/27122/654/1958/653/8644/706/1583/10170/6718/1586/133/650/3
signal=54%","120892/8877/54209/3569/6376/1075/3949/9572/6348/2896/8530/5788"
list=37%, signal=44%","6618/6617/2963/387332/10302/138474/2968/9013/83860/2957/6881/5470
, signal=58%","8301/7078/84168/7077/7076/6774/4035/4155/8434"
688/477/10791/6548/6324/2697/1495/5348/6330/1499/5239/2702/1824/1832/1674/29109/4892/
al=57%","84101/8379/25906/7175/51451/7517/55755/10403/348654/891/4085/81620"
9,"tags=71%, list=20%, signal=57%","84101/8379/25906/7175/51451/7517/55755/10403/348654/8
0%, signal=57%","84101/8379/25906/7175/51451/7517/55755/10403/348654/891/4085/81620"
st=25%, signal=43%","2151/624/1909/5333/23236/5336/113026/1230/51196/6352/5330/1234/230
5,4836,"tags=38%, list=27%, signal=28%","10413/2059/23189/1969/4651/11135/23500/4134/81565
6%","2059/23189/81565/83478/857/4070/5962/391/64805/54869/3383/51225"
78/58494/8013/3037/6793/3553/6279/960/6280"
al=60%","10694/10575/10576/7203/22948/908/10574"
list=19%, signal=29%","83643/5154/6591/5155/5978/27122/27120/8877/4790/653/8644/348/661
","25932/5205/1192/1051/8542/28962/55107/5167/5243/5348/6560/5349/57348/2564/1193/867
list=18%, signal=23%","302/3685/493/309/80024/117531/51393/59341/22821/3710/3708/80036/3
nal=69%","545/4292/5395/4595/7486/4436/84515/5111/2956"
=60%, list=24%, signal=46%","301/861/2737/604/80149/3566/51043/9655/7292/3142/4092/865/8
signal=26%","1410/1969/147912/7048/10252/4094/7431/8412/1414/131544/161742/2697/2195/6
8%, list=26%, signal=35%","2040/7402/493/477/4734/154043/6548/6324/5348/6330/1756/5349/11
45%","6419/55818/8085/55011/23067/9757/54093/6598/79813/9739/10919/79723/6839"
%","10413/7068/10365/57333/5604/3202/652/60485/54206/2252/2627/9719/6662/998/3169/747
0%, signal=58%","10413/25937/3280/30812/55366/130497/6662/3976/7477/6772/5311/38/793/54
nal=29%","4780/81031/56606/8682/2719/6814/1909/1906/5167/3479/8013/6515/338557/6517/55
915/3912/3914/3909/3911/59277"
gs=48%, list=25%, signal=36%","2690/3280/183/7040/4851/2149/3586/3588/3569/85480/3570/154
=39%","182/7431/10392/4851/8767/4792/7128/64127"
50%, list=27%, signal=37%","83860/2957/6881/29079/54700/55832/9282/2958/170506/55729/687
0%, signal=25%","3913/57381/4124/25932/7070/10461/493/8322/7068/5159/1282/30812/196740/
=90%, list=22%, signal=70%","7048/2022/10979/7043/7040/4091/4092/2280/58533"
nal=51%","90627/81792/388/2246/387/652/1499/94/54567/3687/3627"

l=51%","90627/81792/388/2246/387/652/1499/94/54567/3687/3627"
","23643/7096/7097/10333/81622/6850/6279/4615/6280/948"
st=24%, signal=57%","11326/80149/6376/671/145581/2213/3949/9572/2896/133396/8530/5788"
=54%, list=18%, signal=44%","857/624/154043/3082/1647/1756/54541/22943/9647/9770/4092/572
'1387/23512/3297/6603/3066/57459/6601/6599/55693/3151/164684/51780/5928/3183/53615/55
,"2308/117854/5587/10769/79443/3953/9531/10392/10475/4734/1634/1611/26353/3082/5660/2
2%, signal=60%","7414/7132/1903/1003/23236/7122/3553/50848/5624/3551/118/6093/7422/7124
, list=32%, signal=60%","7414/7132/1903/1003/23236/7122/3553/50848/5624/3551/118/6093/742
tags=31%, list=13%, signal=27%","3688/1969/1956/493/2042/6711/3958/10519/2037/8878/5583/3
s=29%, list=10%, signal=27%","84630/245711/6714/51654/56647/80279/10983/595/124790/1164/5
t=25%, signal=54%","1435/7057/146433/59341/5919/728/1436/719/1240/6352/5594/5595/2660"
"7984/1880/54795/2621/5294/6688/3965/3592/7226/1230/811/196527/6352/1234/3579"
/8531/5836/355/857/329/197259/8837/841/57162/8737/11035/330/171389"
list=17%, signal=68%","857/4851/93649/4855/3553/3576/81035/361/335"
al=54%","2690/183/3570/5618/3717/11314/2185/6352/8651/149233/51561"
3%","51574/118460/54512/345630/2091/51010/56915/23404/1736"
=55%","6237/5906/54453/947/3791/51666/133/55273"
113/1969/8463/3399/2735/7474/7481/9573/1942/6662"
signal=42%","3685/30812/387/4330/6422/4883/338773/4057/650/63923/4088/115908/26585"
3,2095,"tags=34%, list=12%, signal=30%","2114/64321/2303/9314/8324/652/1499/2139/79577/229
al=46%","4240/311/5360/57488/54209"
%","10351/857/158219/5468/5360/19/4792/54209/1066/348/7376/6256/4035/1950/335/80820/5
=33%, signal=38%","358/9076/59341/6337/2629/7357/64834/5339/3037/7466/8626/51599/8692/2
%, list=19%, signal=43%","1435/7097/7099/1436/54209/5153/81501"
1,3481,"tags=54%, list=19%, signal=43%","1435/7097/7099/1436/54209/5153/81501"
","677/4000/4780/84962/358/28984/132864/947/6548/8837/4851/1906/3779/5468/5743/10397/
%","3074/2632/5836/2683/2992/285203/117248/53947/10678/11227/374378/51301/337876/2629
gs=70%, list=30%, signal=49%","10772/6432/57060/6433/6431/708/3190/135295/3181/11338/1092
38523,3143,"tags=79%, list=18%, signal=65%","493/857/2771/1909/5350/1906/11214/5142/998/10
8%, signal=46%","5592/928/5580/10019/114897/84959"
=20%, signal=35%","2690/54741/6778/3716/4067/6774/5829/6776/9518/3717/5295"
nal=30%","56034/7098/2690/7048/283/183/2013/5155/284/5580/79817/2246/28984/151/8797/3
22%, signal=62%","301/8767/3566/9655/7292/3142/246778/941"
"3556/3133/114548/1051/51176/3123/942/7292/84868/2207/3965/9308/6850/80762/940/1649/1
signal=42%","3556/3133/114548/1051/51176/3123/942/7292/84868/2207/3965/9308/6850/80762
signal=44%","64863/55186/9361/4976/6742/30968/9093/7156/201973/3980/11232/92667/80119/
840/134265/23022/9948/7171/7412/7984/7082/285590/5782/85477/83706/10097/51411/9644/3
,"tags=62%, list=19%, signal=50%","161291/89866/123264/975/55754/9695/55741/56681"
ignal=32%","55758/440498/51132/9139/2023/54531/10765/3281/8841/8409/64207/84458/161882
14%, signal=32%","29995/493/10519/2274/6548/3479/2065/11261/64581/9454/5621/1827/5336"
522106916,5738,"tags=73%, list=32%, signal=50%","7099/3569/64581/6774/10333/64127/63940/30
l=35%","7098/8531/5836/355/857/329/4217/197259/8837/26471/427/841/57162/8737/11035/330
44%, list=24%, signal=33%","83643/857/9314/1893/10019/5468/54625/9636/54209/114769/3569/7
,"tags=40%, list=27%, signal=30%","51429/389/11135/148170/4478/84962/25802/3958/140609/109
%, signal=63%","9972/55051/55677/26993/7175/11052/54888/10657/1660/22916"
s=80%, list=21%, signal=63%","7132/183/28984/4851/7133/57589/650/84830"
.0%, signal=64%","3572/1511/3569/10630/22915/3717/84034/50848"
ıgs=71%, list=28%, signal=51%","8754/10519/2303/9780/7042/5420/64805/6850/3976/3071/6352/
","10413/135228/677/687/5587/6591/4853/10468/7538/5467/2252/3037/8626/5796/7421/2069/1
signal=29%","2239/1601/57664/4478/2719/9948/387/2295/84870/10082/2195/4919/8321/8324/

8%, list=19%, signal=23%","3556/3572/4124/81565/3280/6387/2737/7058/146433/443/27286/271
3%, list=21%, signal=34%","3910/2303/216/4851/7099/3397/4883/7376/55366/406/11096/3226/70
st=22%, signal=44%","6672/9111/3716/5468/54625/83666/3659/7376/84166/3055/3717/6772/865
50%, list=27%, signal=37%","2059/23189/2934/25802/6840/6711/5580/91663/5962/256949/5756/2
"tags=83%, list=21%, signal=66%","7057/4060/5045/7305/6376/10159/2627/23607/25987/8710"
14%, signal=41%","7132/1410/2303/4851/7042/51176/2296/7133/6688/5519/10018/7421"
796,"tags=38%, list=16%, signal=32%","3270/477/845/6548/5350/1756/1193/817/9446"
, signal=32%","5321/5066/2052/11343/6342/80221/5742/8605/2948/5743/5354/51703/8644/6483
ıal=38%","27202/7057/79148/79890/1843/56920/10512/4851/7130/4681/199/7474/1991/6347/10
%, list=15%, signal=36%","90/7048/2022/1282/2303/23129/1909/1906/51176/3791/1499/4855/545
ist=11%, signal=40%","1969/6591/4641/9076/5583/4867/7082/6615/7122"
'1910/6446/5446/1909/1906"
!%, signal=41%","57088/161291/5205/5359/57194/23120/5243/19/5244/55754/79895/81671/1482
=47%, list=10%, signal=42%","9445/81031/25825/1839/1471/9528/9452/5806"
857/130132/23586/64135/5899/8767/3434/4792/4790/11213/23180/5027/64127/57162/81035/14
","2968/5257/8408/901/92/64210/7046/5260/3480/269/5584/904/645121/8941/5563/8621/72834
gnal=36%","8301/824/5978/387/4217/1728/26471/4170/7020/7042/2353/7099/1958/7474/1499/7
6/7779/23498/4502/4499/4493/7412/6648/4489/1373/706/5027/4501/445/210/6279/57817/5025
57/7056/710/5328/2191/5054"
624801,1651,"tags=29%, list=9%, signal=27%","245711/6714/51654/56647/80279/10983/595/12479
","4323/57326/8091/51176/2131/4092/6662/2261/4057/6781/4921/659"
ist=15%, signal=62%","10694/10575/10576/7203/22948/908/10574/1736"
s=73%, list=15%, signal=62%","10694/10575/10576/7203/22948/908/10574/1736"
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382,4491,"tags=45%, list=25%, signal=34%","966/51226/123720/79056/9780/6892/6890/6891/6468
/3916/1393/8878/120892/5322/4688/4689/2512/8031/84557"
'57602/54700/6223/10436/9221/23378/23133/84172/57673/1663/6839"
s=48%, list=12%, signal=42%","81031/6646/3479/90865/6934/123264/11261/1499/23236/5074/233
327/302/6281/861/7057/7123/5328/5054/1522"
9%, list=18%, signal=32%","8826/10252/90993/7057/55614/860/2246/64094/81848/2260/27020/7
s=72%, list=24%, signal=55%","7057/28984/641700/958/3676/6347/8644/57124/3135/27250/19652
29%","54884/151056/10577/216/157506/6121/220/9249/5919/8644/5467/57016/4035/1545/2978
signal=38%","5125/2737/2295/7042/5919/6943/3398/6299/2736/3142/3576/348093/5915/2294/5
","1277/4313/6422/2201/1101/4488/133522"
,"tags=71%, list=24%, signal=54%","23328/654/23643/11027/9450/85363/114769/6376/7128/1604,
=44%","90102/3913/3912/1605/3914/23187/8985/59277"
"tags=40%, list=24%, signal=31%","9697/6238/259230/23471/166929/567/3133/3107/3105/6892/3
nal=53%","3685/3487/3655/3488/3691/3690/3489/3484"
gnal=45%","2934/8724/10581/10410/9023/81603/684/85363/8519/5806/4261/8547/11043/2219"
=33%","367/1969/7048/1435/2737/3488/857/6778/2099/5241/652/7474/1436/4488/6356/5371/74
gnal=45%","302/6281/861/7057/7123/5054/1522"
%","367/654/653/1499/55366/7484/5156/3293/7490/6469/4920/1718/6926/3209"
24%, list=13%, signal=21%","10413/64114/3685/8531/90/367/4000/4780/6387/3655/90993/8682/6
al=61%","55035/7486/23212/54984/10445/7015/4931/5901/25885/705"
","837/351/2697/3479/7412/5641/7099/4313/54209/3676/4804/3383/10333/4035/5621/2213/100
858/26509/301/5872/9023/928/9341/25978/1514/55647/2350/83871/8676/338382/55207/7075/8
ignal=36%","143872/5547/4846/7424/9424/6648/2702/9365/153/59342/154/9283/5724/1268/659
gnal=42%","367/157506/652/7474/1499/23213/8626"
088,"tags=57%, list=12%, signal=51%","5154/5155/5159/93649/4015/5157/5782/4035"
signal=49%","25913/29844/54617/51377/125476/79913/10445/7528/64710/86/10856/8607"

=56%","604/10875/3122/975/3123/7292/8764/6482/3594/149233/4179/5105/51561"
=22%, signal=40%","10413/25937/2308/182/122953/10979/7040/7041/5167/59341/6648/7474/67
6342/5375/5468/2173/338557/5467/241/2170/57834/116362/6279/5306/341/165140/6280/5730,
%, list=12%, signal=44%","27202/728/2357/2359/719/1240"
'858/3572/493/4038/7431/4846/5028/5660/2697/4067/5604/9332/3709/54209/28514/1832/841/
","5327/5328/5269/5054/1508/114769/834/1475/7448/5104"
/389/57381/10981/2791/85004/5872/25923/6237/51285/5906/6004/2771/84617/51655/2634/907
17583/8531/11346/54532/5872/84952/5376/23586/58494/54587/90952/1003/9076/7408/79778/
%","90/7048/26508/2022/6591/4851/23493/7042/653/4487/6615/4488/650/55273/187"
,"306/6814/3684/6688/6850/5724/57126/3689/6813"
%","1910/2771/1909/4846/140/10487/5957/84876/4883/55366/6786/2587/153/10886/154/53829/
93/29081/29063/64216/51504/79922/79066/345630/10436/4839/2091/117246/55006/55695"
634/3566/602/958/3586/64581/115361/6556/163351/3394/4615/1130/388646/116071/10538"
=67%, list=18%, signal=55%","2150/84868/11117/64127/1535/84034/5650/2896"
910/6446/1909/1906/8644/3037/2621/8692/3784"
signal=42%","1909/140/10487/55366/153/154/49/10486/796/1816/2696/59350/122042/1812/116/
ags=33%, list=13%, signal=29%","7070/857/5580/9961/10019/9467/54625/9021/9052/6422/64092/
gnal=53%","91893/29081/29063/64216/115939/79922/79066/345630/10436/4839/2091/117246/5
51599/344/350/338/335/341/336/55911"
=25%","4154/2737/4038/2719/1889/4853/860/116039/157506/5396/4851/2697/7020/7042/34164
signal=25%","4154/2737/4038/2719/1889/4853/860/116039/157506/5396/4851/2697/7020/7042,
=29%, signal=38%","6934/1499/29940/23213/9394/2131/3340/26035/9375/2132/55959/9653/266
22,"tags=67%, list=21%, signal=53%","1282/1284/1287/1277/84959/6850/4921/1288"
44%","4719/4709/4708/4698/4714/4700/4696/4695/4720/51079/4710/4704/4713/9131/55967/47
55214/5351/5066/8974/5033/283208/5352/64175/8985/54583"
signal=22%","51429/302/1601/2308/344558/152503/4780/1956/64285/4478/8724/3077/10769/28
"56034/80310/1956/1290/824/1278/493/1307/1282/1281/2730/387/1728/10365/1051/1906/4067
58494/3082/5803/1499/1605/1959/5916/7133/1075/4099/400746/6258/9451/163175/5728/1890,
4%","546/51548/57343/55693/55183/4089/4602/9869/1111/6598/1789/23028/79813/57673/2625,
%","5327/387758/2908/114907/358/8837/1906/9076/2629/54206/1373/54541/5176/445/2706/15
9%, signal=47%","4067/3586/4332/7305/5074/3635/2213/11314/27242/695"
),"tags=69%, list=38%, signal=43%","83752/134266/131474/100287932/9409/10440/5190/5830/332
3%, signal=31%","3673/387/151/1906/93649/255743/6915/5743/59341/2149/8877/6869/9446/378
30%","54840/7804/4923/8153/30819/6118/3516/10456/2072/6812/4331/10640/2068/6804/51164
54%","5155/7040/80760/7130/4790/3037/3699/1950/3553/8692/2247/3373/960/3697/3036/3473
84/81565/4627/4067/83871/55207/8013/8773/3566/284021/2242/5294/7942/3162/3965/8673/23
84/81565/4627/4067/83871/55207/8013/8773/3566/284021/2242/5294/7942/3162/3965/8673/232
gnal=59%","11340/51013/118460/22894/54512/51010/56915/5393/23404"
%, signal=50%","100134934/2547/55183/7520/7015/65057/7014"
98,3923,"tags=82%, list=22%, signal=64%","6499/11340/118460/54512/129563/51010/56915/5393/
=11%, signal=39%","10413/182/117854/4780/3280/5978/4851/678/144165/6774"
t=21%, signal=44%","3572/7040/5743/57447/728/3569/23213/3570/718/719/2324/3553/1545/945
=33%","7414/10979/3801/28513/1009/654/1003/5962/1499/7082/5339/1832/1004/4092/1016/998
21%, signal=35%","7048/4323/2737/860/2303/157506/7043/3198/5396/3199/7020/652/1746/7484
69/6649/1728/5742/5743/4258/2878/84203/6648/1193/348/4257/26034/847/241/373156/83667/
7,"tags=68%, list=20%, signal=55%","3133/23586/6810/7305/2150/3684/718/6688/5724/57126/368
ignal=34%","683/4208/4323/6778/604/7040/23529/4773/7099/1880/958/3586/975/3569/30011/9
nal=34%","301/5321/27306/5742/1906/5743/8644/6916/972/552/3553"
=34%","301/5321/27306/5742/1906/5743/8644/6916/972/552/3553"
ignal=31%","1956/183/493/857/2771/10392/8639/1906/120892/185/2668/4790/4883/348/2230/5

al=33%","302/3685/5587/8301/2042/257364/857/3958/79890/6386/79772/3690/11252/7376/297
528,"tags=69%, list=20%, signal=55%","59341/64581/4938/64127/3965/3553/6850/4940/8832"
nal=41%","90/7048/26508/4091/4851/23493/7042/23462/2702/1942/650/659/187"
%, signal=43%","51497/6829/23379/904/4300/79664/54623/8621/26747/51755/905/6919/84337/7
23%","56034/80310/5154/1956/5587/7006/2022/7070/2050/9748/5155/857/79834/5159/9961/70
39%, list=11%, signal=35%","677/1601/8826/1956/6591/7538/8837/1277/6997/5151/3397/678/542
=29%, signal=50%","2737/860/654/652/7474/94/649/4017/7704/4092/6662/8817/55553/650/659/
:=17%, signal=39%","90102/395/4323/7057/347/11170/7097/7474/94/56940/22943/3553/5728/10
ist=19%, signal=41%","79184/84446/29086/60561/80279/29980/1111/63967/983/51514/11073/64
3133/4067/3106/11213/7128/10288/84868/29126/3965/326/3135/83737/868/5678/11006/3559/1
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%, list=22%, signal=40%","2771/5028/9934/2857/140/1511/64805/5027/5029/5031/5724/286530/5
2%, signal=27%","301/5166/857/5468/5743/8013/5467/7376/972/552/9104/6720/2170/3553/37/3
5321/5337/2151/11343/151056/624/1909/4023/63874/79887/9388/3990/5333/8605/196463/5322
6/7919/27258/51574/10594/79066/51691/57819/9128/51690/9733/64852/25804"
5%, list=13%, signal=31%","56034/80310/5154/5155/7424/5912/10457/4683/9770"
24%, signal=37%","3673/4208/5159/255743/23645/3672/9104/945/133522/3717/11035/11314/650
t=14%, signal=50%","687/4005/7068/2730/10634/1514/5962/1508/1520/3625"
2,"tags=44%, list=15%, signal=38%","10694/10575/5609/7520/3178/10576/7203/22948/908/65057/
3%, list=27%, signal=31%","3077/493/7779/477/6548/285195/51393/59341/6560/6337/54795/1006
list=27%, signal=31%","3077/493/7779/477/6548/285195/51393/59341/6560/6337/54795/10060/6
st=15%, signal=34%","2239/1601/57664/2719/387/84870/10082/4919/8321/8324/166336/7474/642
%","3913/1969/6387/2737/2050/2042/3908/1889/4853/91653/23022/23129/91624/1909/56920/3
1906/3708/3037/6563/8692/2131/364/361/2132/1080/360/282679/65010/51763/1437/7380"
4,0.00260548551798272,2509,"tags=60%, list=14%, signal=52%","8886/23560/285855/84365/11724
gnal=37%","2932/26005/51104/5108/84687/9851/1453/9419/137886/50855/11190/79648/55125/2
0%, signal=30%","301/7132/5321/2052/11343/6342/27306/5742/1906/8605/5743/8560/8644/6483
31%","84159/301/3280/2735/27122/1909/1906/654/51092/5604/3200/3202/652/7474/10370/653
28/10847/81611/8295/57634/10902/10856/8607"
, signal=32%","51474/2059/2934/25802/6840/6711/10519/130271/9948/10763/444/5962/59341/1
=41%, list=24%, signal=32%","2040/3688/1969/1956/493/4038/2042/4641/6711/3958/10519/8878
3%","27154/2648/6871/11143/84289/55689/23774/2625/11091"
signal=30%","558/2934/5159/113791/222584/118788/10870/3717/5295/5783/8660"
3%","55567/90990/55605/146754/127602/3796/374654/8632/196385/10513/10126/285643/1768/
=58%, list=30%, signal=40%","2840/57007/9034/1524/1240/1238/1230/1232/1234/3579/729230/10
0%","2840/57007/9034/1524/1240/1238/1230/1232/1234/3579/729230/10663/1235/51554/1233"
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3%","301/114548/604/3566/602/942/9655/7292/3142"
32%","2239/1601/57664/2719/9948/387/2295/84870/10082/4919/8321/8324/166336/7474/4867/
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077/5978/6814/5208/1393/55107/2697/1906/80024/6892/19/59341/51296/51092/6934/2746/743
'10765/55818/9682/3720/79831/65094/8242/23210/55693/51780/54784/84678/23030/23133/230
0765/55818/9682/3720/79831/65094/8242/23210/55693/51780/54784/84678/23030/23133/23028
7098/84159/8531/367/4323/5154/5872/3624/10046/5159/30812/2099/4254/2697/26471/654/149
list=23%, signal=45%","2022/5155/93649/30008/23493/4773/6648/652/348093/93166/2069/2725
17%, signal=58%","8379/7175/7517/10403/57805/348654/4085"
t=12%, signal=37%","3280/2022/2719/4853/4851/1003/2297/652/3791/4487/23213/94/57045/448
%, list=20%, signal=54%","1435/146433/7040/3398/3059/841/3976/8737/10221/5925"
4%","8473/339287/55167/55011/10445/55683/54934/51230/84148/284058/26053/3054/672/1109
078839,4380,"tags=39%, list=24%, signal=30%","3270/477/845/5350/1756/1193/817/9446/6348/514

=48%, list=16%, signal=41%","81031/6646/3479/90865/6934/123264/11261/1499/23236/5074/2338
=43%","4055/7048/4853/3586/5971/55509/29108/84868/6688/1794/3696/3694/81501/116071/10
)7/6778/387/1514/3569/972/3570/6774/1794/326/6850/920/912/9474/940/3594/6469/149233/57
23%, signal=26%","301/5321/5166/11343/6342/157506/4023/27306/9388/3990/5742/1906/2948/5
nal=68%","2475/7702/4654/56257/64768/51574/55011/546/51547/4691/11137/6598/25926/1663
%, signal=49%","3454/26471/3588/65018/23479/4609/23596/3458/7415"
nal=25%","301/1601/1910/80219/5166/493/857/5978/27122/157506/654/5468/5743/8013/27120/
t=20%, signal=44%","3916/3566/6810/2150/3684/8673/3134/6850/5724/2268/57126/3689/3822"
3,0.00281909681375661,4903,"tags=81%, list=27%, signal=59%","7057/79812/185/4804/3162/3592/
,"tags=35%, list=24%, signal=27%","79971/2040/161291/5587/283/2737/4641/8754/10519/9531/54
17%, signal=45%","8531/857/329/841/57162/8737/330"
=27%","301/5321/5166/11343/4023/27306/9388/3990/5742/1906/2948/5743/54988/5354/8560/8
'857/10226/1896/157506/63874/8605/9249/84883/25840/1066/5908/65084/1727/5010/54947/91
","4208/4323/860/654/1277/4209/652/2296/338773/2131/2261/8817/3636/162466"
"4208/4323/860/654/1277/4209/652/2296/338773/2131/2261/8817/3636/162466"
2%","50834/50836/50838/50835/50837/5726/259296/50839/54429/259295/50831/50840/50832/2
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.0028734134321984,4785,"tags=50%, list=27%, signal=37%","55107/54795/6810/5467/3123/25874/
457,"tags=46%, list=25%, signal=35%","1956/64285/183/84632/9961/8412/1839/9052/54206/79651
signal=44%","1435/7057/79148/146433/59341/5919/728/1436/719/1240/1588/2185/6352/5594/5
49%","5926/55904/80349/63925/4297/80335/8473/8085/23067/4800/11105/200424/23168/4801/
27,0.00289351179762293,3517,"tags=59%, list=20%, signal=48%","10417/23586/8013/7099/7474/78
s=56%, list=28%, signal=41%","64114/5525/1601/8301/7040/2633/9528/55022/116372/51225/2343
%, signal=25%","4208/4323/2022/5155/79365/7040/4851/1906/93649/30008/9780/23493/4773/34
signal=45%","493/2771/5350/151/11214/146/5142/153/147/3784/154/5144/150/1813/64784/181
,"7098/117854/10581/824/10410/5359/3428/684/3659/8519/4332/4938/3455/54739"
=41%","8542/3990/5360/348/1191/3931/344/350/338/335/341/5444/3250/336"
%, list=18%, signal=46%","9656/197342/80198/5883/11200/54962/4436/2177/146956"
ignal=26%","2040/5784/302/1601/1410/161291/5587/29969/6672/81565/4478/8724/30846/2737/
6%, signal=63%","1051/55365/28959/3106/10288/3965/2213/3135/10859"
s=44%, list=25%, signal=33%","493869/5742/5743/4258/2878/1193/4257/26034/847/241/373156/8
gs=42%, list=13%, signal=36%","966/2152/7070/8754/2350/929/2159/338328/5621/8692"
%, signal=34%","56034/80310/1956/1290/824/1278/1307/1282/1281/2730/1051/1277/4313/64121
6, list=29%, signal=43%","4208/8531/90/7040/4534/1499/84466/28514/5562/56203/3481/6469/102
49/25855/3066/84312/64426/5928/25942/79595/1457"
5, list=8%, signal=46%","56852/55920/3192/891/9212/81620/990"
1,3271,"tags=64%, list=18%, signal=53%","5028/9934/2857/64805/5029/5031/5724/286530/53829"
ags=64%, list=18%, signal=53%","5028/9934/2857/64805/5029/5031/5724/286530/53829"
%","4502/4499/4493/4489/4501"
2%","4502/4499/4493/4489/4501"
tags=62%, list=30%, signal=44%","387/2099/10133/80149/5187/54209/7128/10318/29108/91662/8
"440145/10426/80097/653784/7283/55835/27229/672"
=28%","55970/51429/858/8826/57664/2791/59345/57477/4627/329/2771/10979/387/2669/6277/
al=52%","91893/29081/29063/64216/79922/79066/345630/10436/4839/2091/117246/55006/5569
46/80312/6996/200424/54784/9760/7874/121642/10155/2625/328"
:=17%, signal=26%","1410/29969/6672/8724/284/9531/347/5350/120892/11261/392/27248/51715
%","30812/2668/652/1499/6299/7484/6662/3976/6772/26585/342184/54361/7849/5076/6608"
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%, signal=28%","7098/4208/196/4780/2052/5978/2329/5446/1728/1393/57552/1906/2948/3309/51
al=29%","4780/81031/56606/8682/2719/6814/1909/1906/5167/3479/8013/6515/338557/6517/55

1%","9388/3990/19/1066/348/1191/3931/344/335/9619/336"
5/3490/85360/7048/2013/3691/2697/5743/7076/4313/5467/397/81671/3553/5618/51673/3976/6
4%, list=17%, signal=53%","26471/120892/90865/54209/3569/1075/3553/5336/6348"
gs=79%, list=25%, signal=59%","9014/64425/55011/51547/4691/11137/6598/6949/25926/1663/659
"9469/56548/10090/8459/83539/27233/9394/8460/9957/9435/6820/3340/27284/6783/166012/41
%, signal=41%","1601/5978/27122/157506/654/6934/1958/4790/653/8644/53919/9563/3091/133/
s=62%, list=13%, signal=53%","493/857/947/80149/3586/706/9104/140885"
62%, list=13%, signal=53%","493/857/947/80149/3586/706/9104/140885"
"tags=68%, list=32%, signal=46%","3133/3916/135250/353091/3592/3134/5817/3135/7409/3822/5
%, list=22%, signal=45%","1601/2246/654/6720/1081/1581/5562/9572/133/3552/9619/3458"
3664/301/5872/5906/2771/5978/6385/3916/6814/3356/5028/4851/6386/4067/120892/341359/35
35%","5321/2687/4258/9261/7097/5322/241/22949/57834/6850/66002/240/64174"
:ags=42%, list=12%, signal=37%","64114/5525/1601/8301/7040/2633/9528/55022/116372/51225/2
al=43%","837/351/2697/3479/7412/5641/7099/54209/3676/4804/3383/10333/4035/5621/2213/10
22%, signal=26%","25937/26524/5525/7026/57493/10252/84962/183/7070/5787/857/5580/9961/1
%, signal=30%","9843/79901/3077/7779/391059/4502/4499/4493/351/654/8546/4489/79689/48/6
ist=13%, signal=46%","114548/3566/975/3569/942/972/7292/64127/91543"
ıal=39%","183/3217/30812/4070/2297/2668/652/55366/130497/6662/4609/6495/6772/51804/749
43/79901/56898/8301/3077/567/391059/654/6648/79689/2034/48/26234/64116/3162/6556/2347
"tags=58%, list=14%, signal=50%","7132/2303/4851/7042/2296/7133/7421"
ags=58%, list=14%, signal=50%","7132/2303/4851/7042/2296/7133/7421"
ıal=40%","57493/2022/10512/4851/94/2294/659/187"
%, signal=44%","558/100507436/3133/3916/639/2214/3601/7305/2621/84868/3592/3134/3717/22
5%, signal=32%","4208/301/3707/4254/5291/23601/64127/5728/3665/350"
list=18%, signal=32%","6591/5978/27122/27120/4790/653/348/6615/3638/650"
30812/2668/652/1499/6299/130497/7484/6662/3976/6772/26585/342184/54361/7849/5076/6608
"tags=32%, list=8%, signal=29%","857/54741/10019/5468/54625/9021/602/1600"
=22%, signal=28%","4780/81031/56606/8682/2719/6814/1909/1906/5167/3479/8013/6515/33855
ignal=59%","10694/10575/10576/7203/22948/908/55135/10574/1736"
al=51%","142940/83480/26995/54517/83448/80324/27079/113000/1736"
/50863/23022/91624/351/27020/54538/9037/57142/2131/152330/8829/4978/27242/1268/51294
6","7098/10475/9111/23586/81844/64135/8767/84282/7097/7099/3659/9636/5971/79132/4938/2
%, signal=40%","7098/10475/9111/23586/81844/64135/8767/84282/7097/7099/3659/9636/5971/7
6","389856/8239/377630/401447/9098/9736/10869/57559/10213/9097/123228/10987/51377/576
nal=45%","2475/1974/8661/8894/23521/8664/54505/728689/51441/1981/57409/10480/1975/196
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39%, signal=43%","4719/4709/4708/4698/4714/4700/4696/4695/4720/51079/4710/4704/4713/559
, signal=52%","7107/84959/6422/3569/7128/5027/3635/338773/57817/4982/796/26585"
,"546/11143/1663/641/5888/8914"
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51867,3603,"tags=73%, list=20%, signal=58%","55011/51547/4691/11137/6598/25926/1663/6597"
list=16%, signal=61%","79026/3915/2200/10516/2199/2201/11117/84034"
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63848,4639,"tags=65%, list=26%, signal=48%","329/8767/51296/7099/4792/144811/7128/64127/33
910/183/2771/1906/185/4881/552/6869/6781/80763/134/1813"
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al=52%","135228/6678/334/928/9749/5175/6403/3690/10184/51706/6622"
ıal=51%","221908/2648/79959/7840/163786/6491/55755/55835/10432/10733/672/23636"
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1%","10765/55818/9682/3720/79831/65094/8242/23210/55693/51780/84678/23030/23133/2302
7%","301/7132/5321/27306/5742/1906/5743/8644/6916/972/22949/552/3553/29785"
l=27%","301/7132/5321/27306/5742/1906/5743/8644/6916/972/22949/552/3553/29785"
gnal=39%","7048/85301/309/5916/5915/6662/6781/25987/10269"
=29%","283/10417/3133/3105/6039/1511/1991/6590/64127/4057/6037/5650/820/7018/1755/846
nal=33%","23189/858/2308/81565/183/5166/23371/2771/5580/2303/8412/5791/5167/6121/9021/
4780/6649/5066/358/4502/4499/4493/351/4489/4501/5621/4017/3931/3552/84557/6622"
26%, signal=38%","3572/9180/1910/3133/114548/5743/3569/241/718/5294/3553/5008/2213/1268
934/4627/6840/11034/9535/85477/23380/2314"
14/23189/2308/367/83439/64321/2737/30851/9314/4929/2099/2697/1003/9351/1495/6934/5117
nal=51%","3720/55011/23383/51535/91782/222229/5885/55869/4436/157570/10856/2146/84515,
%","302/10577/6342/54762/55207/19/26119/1066/5305/3949/4864/9619"
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","558/7049/677/301/10461/567/3624/3707/2113/7538/604/10365/3216/10019/4254/4067/1777/
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63%","1435/1880/64805/54209/6376/1524/482/29956/6348/5025"
ignal=43%","1910/183/2771/1906/185/4881/552/6869/80763/134/1813"
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091/23070/115708/54931/79066/54888/56339/79872/63935/57721/51605/55006"
signal=41%","10226/4023/5468/54209/51099/123/5465/57104"
28%, signal=27%","5906/2771/5978/3916/6814/3356/5028/4851/4067/120892/341359/3566/6810/
gnal=34%","443/5354/7097/3400/1191/4099/23385/5728/4118"
st=17%, signal=34%","443/5354/7097/3400/1191/4099/23385/5728/4118"
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903,5389,"tags=54%, list=30%, signal=38%","6125/6135/143941/65992/9097/8473/51377/6201/28
"2151/624/1909/5333/2923/23236/5336/113026/1230/51196/6352/5330/1234/23007/81544/8986
nal=31%","3074/285362/5580/4758/53947/2720/30815/338596/2629/7357/3073/51228/81849/87
","23381/170506/54984/6634/54433/9887/3183/7015/55651/3192/55135/6628/1736"
, list=28%, signal=42%","1969/10519/10019/4534/1605/5291/23612/8692/10385/5728/3135/5795/
gnal=26%","5321/4907/1806/8871/5580/151056/27115/79887/9388/3990/8654/8605/5167/196463
%","10577/54762/5360/19/348/706/80765/338/335/9619/336"
signal=27%","11113/116840/9820/26073/91782/84501/54784/285643/10617/9585/6683/54443/27
, signal=27%","302/6281/4323/2934/4627/7123/10418"
st=17%, signal=26%","677/4000/4780/84962/358/28984/132864/947/6548/8837/4851/1906/3779/5
1%, signal=53%","10498/54496/55471/55170/10419/3276"
2/4015/388630/84171/4017/335/4016/2876/336/129293"
558/1956/81565/6387/2737/27185/3912/387/120892/26468/1600/64805/1499/5649/285025/5714
040/1910/7402/493/6446/477/4734/5446/1909/154043/4846/6548/6324/285195/1906/80024/534
5883271731028,5627,"tags=63%, list=31%, signal=43%","4780/90993/9531/4734/10365/1051/4851/
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4780/5066/400/2327/2326/1545"
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2%, list=16%, signal=43%","22925/2934/7057/28984/641700/958/3676/6347/8644/3569/6424/5712
%","5592/23704/493/6446/857/7779/4734/28954/23678/6324/2669/3779/5348/6330/5349/3708/1
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35%","1601/3077/5978/27122/654/1958/4790/653/2230/706/83539/9563/1588/1583/50700/1586,
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55,6970,"tags=64%, list=39%, signal=40%","4719/4709/1329/4708/4698/4714/9167/4700/1340/738
32%","301/5321/2687/27306/5742/1906/5743/4258/5322/8644/6916/972/241/552/3553/6850/24
%","4121/81792/1514/1471/56999/9695/26270/4126/26232/55741"
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gnal=32%","2152/4780/7057/7056/6484/6915/2149/5054"
2%","2152/4780/7057/7056/6484/6915/2149/5054"
822,3259,"tags=45%, list=18%, signal=37%","2690/7048/871/85301/309/1277/59341/5916/2131/59
nal=27%","259230/55304/5587/166929/10351/5580/53947/30815/375298/8877/338596/8560/262
%, signal=42%","3842/26953/30000/55374/3843/79711/55705/23534/3838/3837"
s=43%, list=7%, signal=40%","9445/81031/25825/1839/9528/9452"
0%, signal=46%","284021/5169/3162/3965/2294/11314/1268/8832"
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, list=14%, signal=28%","7049/57493/2022/2303/1909/7040/4851/23493/7042/2296/1832/4634/71
, list=17%, signal=60%","6484/6401/6403/1991/3676/2526/9435/5724"
5%, signal=46%","7078/8797/467/6688/841/5728"
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","285643/9493/9055/9212/24137/3619/29127"
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302,4166,"tags=73%, list=23%, signal=56%","182/3280/1909/10512/1906/652/10370/998/5594/530
gs=51%, list=26%, signal=38%","329/55601/23586/80216/64135/8767/51296/84282/7099/4792/549
gnal=25%","4208/182/26508/3280/6387/1910/2737/567/5978/30812/387/604/351/151/4851/4681
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4/4121/57134/23324/9695/4126/55741/4125"
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%, signal=46%","5205/57194/23120/5243/5360/19/5244/5306/9619/57605"
14/5351/444/8974/10491/5033/283208/5352/64175/8985/54583"
1%","4780/358/4502/4499/4493/351/4489/4501/5621"
4617/51377/125476/79913/10445/7528/86/10856/8607"
24%, signal=32%","1909/1906/30845/54209/6347/3460/5649/65018/1520/6786/5336/57158/5031,
33%","3356/351/26471/3479/8671/5027/54541/6774/51341/51422/5562/4609/957/3091/51085/5
, signal=50%","302/29116/5045/151636/7805/348"
6%, signal=47%","10413/3572/3716/3569/3570/6774/6688/1050/8737/3717"
signal=32%","23704/857/9948/845/6324/2697/6330/2702/2316/3784/3759/10269/23630"
%","302/55614/1889/9341/4734/7040/29116/5045/151636/8934/30845/5641/10133/7805/348/394

nal=44%","6234/6233/6232/6188/6231/54505/6204/6229/6235/6228/6191/6222/6208/6218/3921/
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%","7388/51292/4696/4695/10975/126328/4720/51079/4710/3421/29796/2653/4704/27089/5575
%, list=31%, signal=59%","143872/5592/10203/80005/5743/154/150/134/796/3778/5997/136"
)42/8467/5928/5931/2186/9031/54107"
35%","9429/23704/477/6548/2697/5243/19/5348/3752/8671/3737/482/123041/3784/30061/8645,
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%","158/2618/471/5198/5471/10606"
signal=28%","5784/5525/5136/1848/5337/4907/2151/8871/5787/23371/1847/64759/477/5446/624
signal=44%","5429/11201/5423/27434/51455/10714/201973/11232/5426/10721/54107/5427/542
gnal=26%","3399/857/4188/7779/4734/6324/5350/4091/3708/51176/3397/3398/3400/29763/4824
ıal=53%","6881/6878/6879/6873/51616/2648/8295/6877/6874"
2308/3920/4780/5587/3077/5166/355/4217/7538/1611/10634/6751/3428/23710/5899/55062/120
1910/183/2771/1906/185/4881/552/6869/80763/134/1813"
=62%, list=17%, signal=51%","567/3133/8546/6891/3134/3135/563/912"
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%, signal=48%","1806/978/7378/27350/151531/60489/200315/140564/81602/164668"
01/3077/567/654/7020/48/3162/210/30061/3091/57817/84557/9474/6622/7018/1813"
%","79228/7919/9984/57187/8563/26097/84321/10189"
=29%","8531/857/329/8837/26471/427/706/841/57162/8737/11035/330"
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=28%","493/6446/4734/23678/6324/5348/6330/5349/817/53827/6327/7533/2257"
gnal=33%","4208/367/10252/2737/2246/2668/652/7474/1499/7481/22943/2131/2247/650/6495"
37886/116840/64793/1111/4751/3832/5902/23636/6790"
2%, signal=57%","2072/79677/5892/10111/7517/51002/7515/7014/675/5889/5888"
0989/55735/10651/79135/54927/25813/4580/84303"
)989/55735/10651/79135/54927/25813/4580/84303"
al=46%","3487/3488/26471/51092/8013/6422/3164/1992/54206"
gnal=26%","3074/259230/55304/285362/5587/166929/10351/5580/4758/53947/6484/2720/5660/
=19%, signal=36%","3572/3280/146433/443/7040/5803/4851/23529/7097/1605/3398/1959/7133/4
ignal=25%","10413/3572/182/4323/3280/604/4846/23555/4851/23493/51162/7042/4792/23462/2
2%, signal=35%","3678/5359/1839/2202/222584/9655/1950/685/2069/7039/374/6643/10451/378
,4238,"tags=48%, list=24%, signal=37%","3270/845/5350/1756/1193/817/57158/9446/5144/2280/2
signal=60%","8663/8661/3646/8664/728689/27335/8666/8669/10480/51386/1965/8667/8665/866
%, list=36%, signal=60%","8663/8661/3646/8664/728689/27335/10987/8666/8669/10480/51386/8
","4739/2934/83872/123720/57091/54857/90550"
al=50%","8724/5359/1265/8291/7097/2213/140885/11314/64422/9474/5657/26060"
13/116840/9820/26073/91782/285643/10617/9585/6683/54443/27183/55165/26586/1894/9493/
21%, signal=40%","90102/395/84952/1901/4070/10163/7082/10391/23671/10395/5295/50848/745
"10417/23643/929/6403/7097/7099/54209/5027/671/4814/4057/6037/5724/820/84659/3929/245
%","9023/51302/9420/1545/29785/1588/1593/1581/1583/1586/113612/1559/10858/1589/1548/1
18%, signal=41%","1601/5978/27122/654/1958/653/706/1583/133/650"
t=8%, signal=37%","2022/6591/5159/1909/10512/4851/1906/652"
28%, signal=50%","6387/245937/5641/5054/199/57402/1524/1230/196527/6352/3627/2358/72923
.00443775157332758,3143,"tags=32%, list=18%, signal=27%","3270/477/845/5350/1756/1193/817/
4027,"tags=40%, list=22%, signal=31%","143872/23189/3688/57493/10252/84952/83478/604/81848
%, list=21%, signal=55%","11340/22894/29883/57472/115024/51010/56915/5393/23404"
28/3691/7097/23136/4099/400746"

%, list=32%, signal=55%","1843/347/4681/23143/653/6347/23236/84830/9466/11240/26585/1314
t=19%, signal=32%","23704/857/6324/6330/3752/54795/1824/1832/3762/817/3784/6327/29119/3
.6%","4208/1514/80301/64127/3665/3661/1234"
ist=26%, signal=49%","4023/3990/348/3931/344/335/336/9971"
list=26%, signal=49%","4023/3990/348/3931/344/335/336/9971"
,3124,"tags=48%, list=17%, signal=40%","10252/90993/7057/860/64094/81848/27020/7474/1499/4
8,"tags=86%, list=19%, signal=70%","10201/347736/4831/29922/83549/54963/4833/790/7371/150
3297/10694/10575/55172/10576/7203/22948/10963/908/23234/10856/10574/5536/8607"
=19%, signal=59%","3956/10475/6737/3123/972/3965/920/7113"
, list=19%, signal=59%","3956/10475/6737/3123/972/3965/920/7113"
048/3202/652/9719/6662/7477"
signal=50%","3082/1958/728/1191/22943/5925/3458"
ıgs=71%, list=23%, signal=54%","1601/4478/8724/5962/5604/7430/23011/55823/29934/51652/559
%, signal=52%","6387/6484/6401/6403/1991/3676/2526/9435/56477/5724"
al=33%","4323/2737/7043/652/4488/51715/3638"
7%","861/7040/5468/7076/2149/3091/3552"
ignal=34%","1910/2771/1909/140/10487/84876/4883/55366/6786/2587/153/10886/154/53829/49
%","4643/10749/81565/79443/4641/55930/8546/63971/4642/81/1176/66008/4644/4648/4542/586
=35%, list=24%, signal=27%","5563/27124/2932/81551/57704/2258/1946/403/50861/57159/12673
l=29%","57088/161291/5205/5359/57194/388121/5580/10577/6342/23120/5243/5360/19/5244/34
=53%","8663/8661/3646/8664/54505/728689/27335/8666/8669/10480/51386/1965/8667/8665/86
%","8531/857/329/8837/841/57162/8737/11035/330"
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302/2200/4853/7305/79705/56975/4057/10859"
st=10%, signal=35%","302/301/11135/6282/9124/1266/29766"
%, list=10%, signal=57%","4851/654/652/1499/694/84913/94"
%","7283/259266/10403/26271/9133/9700/6790"
list=21%, signal=21%","64764/2059/3673/11135/4780/6239/5587/81565/6387/148170/8724/3655
=31%","302/6281/8301/8724/83547/8546/89866/92421/6517/9871/5027/54732/7879/64083/2628
98/8531/80303/358/9076/59341/54206/92241/23308/361/6558/553/10367"
48%, list=21%, signal=38%","84836/9060/5167/8459/27233/8460/6820/27284/6783/6817"
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919/246176/10615/55755/122060/11004/9793/90417/146909"
913/81565/9314/347/4067/1495/8578/5604/6648/199/4313/64805/54209/1605/7305/706/2316/4
59%, list=25%, signal=51%","3269/5028/5031/5745/5724/2185/553/6622/5341/2852/51744"
5155/3958/3082/7424/6360/2668/652/7474/6376/2252/10457/4908/6348/2247/7857/6352/3627
gnal=49%","4055/7048/4853/5971/55509/6688/3696/3694/81501/116071/10538/3662/7189/1437"
list=18%, signal=31%","4851/654/5583/5054/2668/652/1499/694/84913/94/6424/2296/1030/7421
, list=20%, signal=64%","90102/28984/7040/1605/23187/11117/6662/4088"
5%, signal=21%","10413/4774/307/4208/6867/367/117854/196/2114/4780/29969/3280/57326/850
5%, signal=58%","604/10875/3122/3123/7292/6482/3594/149233/4179/5105/51561"
3454381,4517,"tags=71%, list=25%, signal=53%","8767/51296/7099/4792/144811/7128/64127/3304
2%, list=15%, signal=52%","824/5580/23682/3356/11261/3357/2170/344"
gnal=33%","6788/302/3077/567/857/23129/351/151636/5899/6737/120892/8767/6934/652/7474
/10392/8767/114769/834/29775/8915/64127"
signal=45%","3356/351/3479/8671/5027/5562/4609/957/3091/51085/3458"
%, signal=35%","367/1969/1435/6239/6778/1051/2099/5241/3202/7474/1436/84876/4487/3398/4
!4%, signal=47%","183/3990/185/5322/348/338/4353/9619/336/5320"
st=34%, signal=44%","1435/4853/2353/54209/7305/133522/1230/3458/149233/51561/3605/7071/
395/7430/6404/3384/388325/10211/9051"

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gnal=32%","677/1601/8826/1956/6591/7538/8837/1277/6997/5151/3397/678/54206/56940/2296/
737/355/3958/652/7474/1831/155038/1942/29126/8915/3965/81704/10018/8737/6482/11035/30
%, signal=39%","8531/857/329/26471/841/57162/8737/330"
/307/302/301/306"
%","3688/6387/183/477/3356/5028/151/140/341359/7432/2668/10550/6529/5027/65018/552/37
1910/284/1909/2131/3340/26035/2132/84068"
'8,"tags=52%, list=23%, signal=41%","183/3217/30812/4070/2668/652/55366/6662/6495/51804/26
st=18%, signal=60%","1956/23493/8324/28514/144165/22943/55553/650"
gnal=51%","677/1435/146433/3204/3123/2242/972/3635/920/4609/3665/81501"
signal=54%","7919/6118/54465/9984/8563/5514/79621/11200/57805/79968/55159"
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64321/30812/9096/652/130497/6662/2294/6495/84445/27031/6469/2018"
17%, signal=41%","4907/1806/7378/1807/272/1890/151531/9615/4860"
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36,"tags=32%, list=8%, signal=29%","857/54741/10019/54625/9021/602/1600"
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%, signal=42%","23189/1969/6239/10163/5756/51225/10630/11344/998/832/10096/23191/5295/5
1340/90459/112479/6227/51010/5393/23404"
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, list=25%, signal=53%","55968/2932/137886/50855/11190/55722/23636"
signal=48%","22925/301/283/185/5538/4035"
tags=82%, list=25%, signal=62%","3430/2150/57590/57162/4814/4057/353376/3929/26191"
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5","3920/4734/112770/8546/55788/7107/79589/7805/975/1191/950/7879/57142/64747/6272/640
ignal=33%","10413/135228/677/687/5587/6591/4853/7538/2252/3037/8626/5796/7421"
signal=39%","3269/3707/5028/80271/5336/5031/5745/2247/5724/2185/553/6622/5341/2852"
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","7098/84159/8531/367/4323/5154/5872/3624/10046/5159/30812/2099/4254/2697/26471/1495/
%","1956/81565/6387/2737/27185/3912/387/120892/26468/1600/64805/1499/5649/285025/5714
55%","2737/972/1794/326/9474/940/6469/5788"
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signal=50%","23166/7057/4481/3990/5360/8578/54209/3949/81035/1401/948/1012"
, signal=38%","3691/79628/10397/1605/5339/4915/400746/6497/163175/4653/3611/57716/5453"
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0%","2308/3920/4780/5587/5166/355/4217/1611/10634/3428/23710/5899/55062/120892/81631/
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5, signal=48%","55818/9682/3720/55011/546/57343/55693/55183/4602/6598/1789/23028/1786/10
6","1601/1910/5978/27122/654/653"
ags=55%, list=21%, signal=44%","3572/7040/5743/57447/728/3569/23213/718/719/2324/3553/154
.0201/347736/4831/29922/83549/54963/4833/7371/1503/56474/4830"
73%, list=28%, signal=53%","51079/55750/158234/140823/26520/26517/26519/1678"
","79441/54801/7283/23354/3192/115106/93323"
32%","54984/100134934/10575/5609/7520/6714/10576/4931/65057/3184/4751/3181/55135/8550
4,"tags=50%, list=27%, signal=37%","83860/2957/6881/29079/55832/9282/2958/170506/6878/9968
4780/81031/26503/56606/8682/2719/358/6814/1909/1906/5167/3479/8013/6515/338557/6517/5

, list=11%, signal=29%","301/861/2737/3624/604/10875/80149/3566/652/6422/3659/51043/3398/9
%, signal=26%","8801/1969/4907/5166/84836/387/80221/9060/5744/3356/5208/7040/351/27115/
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%","4709/1329/205327/54949/4708/1738/4698/4714/9167/4700/10105/7388/4696/4695/92014/1
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l=40%","4719/4709/1329/54949/4708/1738/4698/4714/9167/4700/7388/4696/4695/92014/4720/
/183/5166/8309/8871/154141/6342/6484/347/10903/8821/185/6646/8013/375298/8877/4534/26
","tags=59%, list=31%, signal=40%","4780/90993/9531/4734/10365/1051/4851/467/3309/1958/203
"4869/4926/2010/137886/9787/9212/6790/10101"
8%, signal=34%","4780/183/824/9531/4217/8837/26471/1906/93649/5468/23493/3479/6648/8014
signal=38%","857/3270/477/845/2697/2702/817/2706/482/3736/5144/288/57165/10804/481/476/
nal=63%","84868/81622/51284/91543/118788/285852/388325/26191/23547/1654"
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signal=32%","117854/5654/10475/23586/151636/83666/10875/3434/80149/54941/7128/79132/29
5%, list=14%, signal=22%","7098/84159/8531/367/4323/5154/5872/3624/10046/5159/30812/2099
29%, signal=51%","1601/5978/27122/654/1958/4790/653/9563/3091/133/650/2865/8614/54361/4
4321/3214/4851/7042/8324/678/7704/4674/2131/84750/2124"
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ignal=31%","1956/493/857/10392/2668/4790/4883/348/5291/6271/3357/3553/23576/7421/3091/1
.39,"tags=45%, list=29%, signal=32%","1639/859/1027/4204/4926/10142/27124/81551/57704/2258
gnal=28%","10413/2059/23189/1969/4651/11135/23500/4134/81565/148170/5205/5337/83478/6
"1386/8443/6881/9027/254531/79612/1387/8260/10847/7994/339983/9862/2194/8888/79969/26
signal=22%","840/1519/824/29761/837/1513/8837/1514/843/5641/826/6422/27283/1508/2923/1
=70%, list=23%, signal=54%","2022/93649/30008/6648/1491/2852/6469"
1%, list=24%, signal=31%","2040/858/5978/2335/8542/8091/80149/1436/1511/51176/1991/9367/3
27122/120892/27120/652/27123/22943/7484/650"
,"tags=45%, list=25%, signal=34%","2690/3280/183/4851/2149/3586/3588/3569/3570/1545/3587/5
=30%","4733/7175/79598/91782/3192/10615/79866/84722/56992/8480/10460/22974/79187/2363
"tags=89%, list=31%, signal=62%","100507436/3133/3105/3106/84868/3965/3134/5272/3135/1085
513/1514/5641/1508/51284/1520/3949/53373/51311/6609"
l=27%","4627/10519/4023/9351/55788/23345/23136"
=21%, signal=22%","10413/7049/2690/4208/8531/7048/1435/81565/6387/2735/27185/2335/4851/
%","3673/4208"
767/929/7097/7099/54209"
46%","8767/929/7097/7099/54209"
st=28%, signal=65%","4052/2200/4681/2201/375387/64388/26585/4179/9350"
29/11201/25898/51455/10714/83932/26073/9100/201973/5425/51514/5111/5427/5424"
3113,"tags=58%, list=17%, signal=48%","2335/5045/7305/2627/23607/10859/25987"
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80312/6996/200424/54784/9760/7874/121642/10155/2625/328"
%, list=36%, signal=39%","81031/26503/56606/358/6515/6517/55314/154091/200010/364/144195/
","3673/677/7048/10203/7057/2730/387/4023/1728/5208/55107/1292/5743/4067/51092/6934/66
%","138474/8900/91646/11077/29988/54514/56159/11022/254528/4292/27030/56155/11144/643
signal=36%","5525/3673/4208/5159/5580/7040/255743/1003/23645/2629/10288/65018/3672/91
=17%, signal=40%","81565/2737/27185/3912/26468/1600/64805/1499/5649/285025/57142/85458/
8,3044,"tags=67%, list=17%, signal=55%","2022/5155/93649/30008/23493/4773/6648/27250/4772/
st=17%, signal=49%","90102/395/4323/7057/347/11170/94/56940/5728/10395"
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, signal=22%","51727/3074/259230/5321/5836/5154/84962/166929/5337/824/5166/126969/3269/
/3688/64321/2737/9096/4851/50846/8324/7474/1499/7481/2139/28514/2736/22943/3226/54345
al=45%","367/64321/8324/7474/7481/22943/2247"
=44%, list=26%, signal=33%","7070/857/2149/6376/5027/9002/57158/5031/825/9446/10268/9001,
signal=37%","1910/11343/6548/1906/2149/7432/3586/6348/5724/55273/80763/3552/134"
37%","1910/11343/6548/1906/2149/7432/3586/6348/5724/55273/80763/3552/134"
477/6548/7043/6324/5962/5348/8291/6517/361/825/779/3759"
5%, signal=30%","4733/7175/79598/91782/3192/79866/84722/56992/8480/10460/22974/79187/23
l=39%","1601/1910/5978/27122/654/653"
nal=44%","8894/79727/2332/1979/6229/728689/26608/1981/22954/10480/5436/23367/1977/889(
gnal=53%","8663/8661/3646/8664/54505/728689/27335/8666/8669/10480/51386/8667/8665/866
48/3624/7040/54762/1066/1605/23209/3625/1581/6348"
53%","137886/64793/1111/4751/3832/5902/23636/6790"
gs=40%, list=12%, signal=35%","158/2618/471/8833/5198/5471/1716/10606"
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=40%","57007/9034/1524/1238/1230/1232/1234/3579/729230/10663/1235/51554/1233"
signal=65%","11027/10288/11024/11025/10990/10859/353514/79168/23547"
st=19%, signal=65%","677/1435/146433/3123/2242/972/920/81501"
s=27%, list=13%, signal=24%","79971/10413/23189/57664/23500/1956/2719/857/9839/1896/3087/
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=62%","100130958/644186/10388/93426/56159/10609/27030/10734/56154/5885/7917/9985/931
%, list=13%, signal=34%","5631/158/2618/471/54963/790/8833/7371/5198/5471/1854/1716/10606/
23%, signal=43%","27202/79148/79890/1843/7130/4681/6347/1588/1803/11240/26585"
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","90/57493/2022/4853/4851/23493/7042/7474/7481/2702/54567/650/659/27031"
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2%, signal=24%","5321/5154/84962/5337/824/126969/5155/8871/3707/5359/23167/23682/63874/
nal=24%","3556/2239/4208/85461/7058/4038/27286/27185/4734/387/26037/351/22899/7162/120
al=45%","2059/2934/6840/5962/5756/11344/832/85477/829/2314/830/120/118"
signal=43%","55809/10463/7251/23082/5585/29079/56943/9282/84660/83637/56970/55749/9968
6%","7049/7048/2615/7043/7042/8321/7474/51176/7481/2535/3696"
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0134934/2547/7520/7015/65057/7014"
7%","100134934/2547/7520/7015/65057/7014"
gnal=30%","5872/5045/6517/10618/9182/54732/51762/3482/83988/10972/6272"
","7098/857/130132/23586/64135/5899/8767/3434/4792/11213/23180/64127/81035/148022/366
:=18%, signal=44%","25937/11030/1601/81031/6711/7043/7040/654/5468/652/4035/79674/8483/
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3214,"tags=57%, list=18%, signal=46%","3624/4091/654/1003/6997/652/60485/653/29108/656/951
","2308/2934/4000/90523/493/3488/8837/5468/23493/54206/84676/2627"
%, signal=42%","161291/5205/57194/23120/55754/79895"
","6788/90/64321/4851/3791/54345"
5%, signal=50%","966/2/710/11326/1604/725/4179/722"
84,"tags=45%, list=17%, signal=37%","8528/8639/4015/7412/2746/84171/2747/4017/4016"
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13%, signal=37%","3554/567/114548/3105/10865/975/3569/7292/1604/91543/3553"
13/2120/677/861/64321/6591/10979/10763/947/7424/4254/2697/8091/5243/2122/7474/2629"
%, list=17%, signal=23%","10413/23189/1601/4134/3280/7070/5376/4038/567/83478/7431/10519/
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%, signal=45%","284/7040/654/59341/348/7082/9414/6869/5139"
, signal=30%","4780/10516/8309/5978/3082/120892/4170/79772/8013/6648/8321/7099/84883/51
","4124/4121/57134/23324/9695/55741/4125"
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signal=44%","8531/80303/624/5743/5243/59341/23209"
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38%","196/100507436/3140/3105/3123/84868/29126/3592/5817/8751/56253"
%, list=18%, signal=36%","6591/5978/27122/27120/4790/653/348/6615/3638/650"
7/56975/6786/10312/123041/3694/5465/9313/1080/64065/9622/84957/401138/6522/25818/1011
538/59341/706"
493/4846/64116/9056/445/23576/435/4942/2184"
"10577/54762/5360/19/348/706/80765/338/335/114876/114879/4864/9619/336/114884/114883'
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19%, signal=38%","81565/7070/9314/3082/5396/8578/5604/2316/5728/6696/29956/2896/5764"
=37%","90/3280/1889/1909/1906/3199/7042/652/653/4824/659/6495/51804"
st=19%, signal=49%","51750/7486/7520/7517/84464/7014/1763/9156/641"
/7486/7520/7517/84464/7014/1763/9156/641"
st=19%, signal=49%","51750/7486/7520/7517/84464/7014/1763/9156/641"
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/5742/5743/51302/9420/55640/79689/6916/847/2977/3162/1545/29785/57834/1535/55652/1588
%","302/3077/567/4015/3586/8829/7262/8751/64388"
30%","11015/4052/2934/2200/6672/857/4188/4681/23345/3309/5552/4792/2201/3586/5756/706/
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%, list=19%, signal=21%","10413/7414/3913/8826/3688/7048/1435/10252/81565/6387/1901/2718
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,"2934/3270/477/5350/8654/5142/817/5144/3759/5025/2946/481/476/5997"
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5,"tags=40%, list=27%, signal=29%","2059/23189/1410/2934/56937/25802/6840/6711/329/5580/40
%, list=17%, signal=51%","4780/3082/120892/79772/8013/5151/3586/65018/55074/135112/9283"
, signal=52%","8930/4595/6996/23583/4913/55247/7374"
, list=23%, signal=57%","55818/9682/55011/55693/6598/1789/23028/1786/672"
0683307935267462,2421,"tags=50%, list=13%, signal=43%","493/81792/1634/5291/3592/10253"
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4124/51226/8934/50999/643699/100128327/56927/4043/27333/10972/3135/101059918/65258/5
%","558/54843/302/7094/4240/7057/309/2350/2948/4258/6121/341359/79792/54209/1373/2621
2%, signal=44%","9388/3990/5360/348/3931/335/341/9619/336"
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25%","1601/3433/6672/3077/5155/567/857/4188/5580/27185/10763/5028/5350/8091/3428/4011
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t=22%, signal=34%","10413/3572/182/3280/4846/4851/7042/28514/6774/8626/64127/3955/54567
048/2022/10272/10468/94/4092"
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, signal=37%","7098/10475/9111/23586/81844/64135/6737/8767/929/84282/7097/7099/3659/358
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ıal=49%","5725/58155/56853/6231/10772/6432/57060/6433/6431/6207/708/3190/135295/3181/1
=39%, signal=42%","4719/4709/4708/4698/4714/4700/4696/4695/4720/51079/4710/4704/4713/55
646,"tags=66%, list=37%, signal=42%","6618/6617/2963/387332/10302/138474/2968/83860/2957/(
"90/1910/30812/1909/309/56920/2335/10512/4254/1906/2668/652/10501/3357/9037/8829/998/
%, signal=40%","824/126969/23682/57153/56994/2170/23446/3931/25979/335/336/80736/376497
5,"tags=53%, list=17%, signal=44%","83460/10328/8266/51234/7917/6449/51608/51016"
signal=40%","3270/477/1906/23493/146/7042/54795/552/7168/153/147/3784/5144/133/1908/10
%","7026/639/3726/5743/10370/1508/5467/4693/2706/7421/51673/3976/6696/6781/659/5764"
st=16%, signal=34%","55075/9111/114548/3430/9516/80149/6688/91662/9770/7114"
, list=20%, signal=35%","5359/5580/4067/8013/2207/2213/6850/3055/5795/2268/5338/7409/2606
5/116039/30812/387/4330/3726/6422/4883/338773/4057/650/63923/4088/115908/26585"
.8%, signal=46%","9314/4023/7099/972/2150/4938/4155/4940/4615"
gs=56%, list=18%, signal=46%","9314/4023/7099/972/2150/4938/4155/4940/4615"
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%","2152/3075/7056/5328/4153/5329/8547/2219"
%, signal=30%","135228/677/7538/4851/3204/5583/54206/2296/81537/4488/8626/7421"
/1003/652/653/29108/656/9518/651/650"
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24/23022/5468/27295/1605/9260/9499/8572/64236/4892/129446"
=73%, list=21%, signal=57%","351/11170/7305/348/1524/5621/5764/134"
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10413/7049/4208/7048/861/1848/2735/4851/1906/23493/3479/10370/60485/2627/32/9686/5728
34%, signal=39%","1909/140/10487/84876/55366/6786/153/154/49/10486/796/1816/2696/59350/1
30%, signal=39%","23704/9948/6324/2697/6330/3752/2702/2316/3762/3784/3759/10269/23630/3
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ıgs=33%, list=10%, signal=30%","161291/5205/57194/23120/5243/19/5244/55754/79895"
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28,"tags=41%, list=20%, signal=32%","4067/3566/2242/5169/3162/3965/8673/6850/2294/2268/113
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02,4990,"tags=45%, list=28%, signal=33%","7070/2149/6376/9002/57158/5031/825/9446/9001/187
37/183/477/3356/5028/151/140/341359/7432/2668/10550/6529/5027/65018/552/3737/6623/655
8%","6788/302/3673/6281/117854/3280/3077/567/857/1203/9314/23129/7040/351/93649/15163
6%, signal=26%","64764/5611/8406/4780/90993/4121/29761/7057/837/857/4217/1051/26471/573
","124491/9972/53371/7175/55746/55706/81929/129401/9688/57122/23165"
gnal=34%","26020/23166/58191/4973/4035/53353/3949"
7/972/1794/326/9474/940/6469/5788"
gnal=39%","22925/3685/7057/4790/3690/7376/344/341/1408/8660/336"
l=26%","7804/55567/26153/140735/90990/55605/11258/146754/127602/57159/85478/3796/1012
gnal=23%","3572/4124/81565/3280/6387/2737/146433/443/27185/30812/23129/7040/2335/5803/
61%, list=28%, signal=44%","9469/56548/337876/29940/22856/83539/79586/54360/55454/166012
4%, signal=48%","79792/64581/5027/6590/6850/4057/219972/6280/84659/3458/130120/1668/50
","2934/6548/1906/1756/2353/4792/4790/1499/3725"
%","80149/51043/64332/3569/3570/6774/4092/3594/149233/3558/84807/10538/6097/51561/366
0799/10556/54931/10775/3028/79897/11102/10940/10248"
,"10201/347736/4831/29922/83549/54963/4833/7371/1503/56474/4830"
l=30%","683/196883/2984/4881/2977/8635/4337"
=30%, signal=35%","23704/857/3752/54795/1824/1832/3762/3784/29119/23630/3764/6640/3753/
91231,6004,"tags=78%, list=33%, signal=52%","183/3217/30812/2668/55366/6662/6495/51804/265
357/477/79784/6548/9096/6324/2697/2769/6330/1756/6332/3752/54795/9630/2629/2702/2316/1
0791087431628186,2199,"tags=64%, list=12%, signal=56%","51594/22794/9704/9775/79048/3182/
","10404/254827/4323/5069/79148/1889/9510/8754/170689/81792/1368/51752/119587/1359/432
list=13%, signal=44%","7132/355/7133/8764/55504"
nal=70%","23555/975/10867/10630/81619/25939/102/388325/340348/51206"
4/7057/64856/2701/706/5649/6869/57121/5025/644139/7060"
6","23498/25978/5028/3569/6376/55277/1134/5562/4130/1268/134/10590/5023/794"
6","84159/4643/2200/57493/64651/9314/639/4929/10512/25976/6648/3708/652/48/3791/10848/
nal=35%","2308/355/857/5159/1277/6648/4313/847/10533/10630"
37/183/3356/5028/151/140/341359/7432/2668/3690/65018/3737/6850/6623/10859/6581/1268/65
%, signal=41%","8826/80310/5155/5159/5962/7099/54206/3037/4609/3373/3036"
, signal=25%","302/6281/4323/2934/4627/7123/10418"
404/254827/5547/4323/1368/51752/119587/1359/2028/2191/5476/1200/54504/290/1522/25797/
nal=35%","90/7048/26508/4091/4851/23493/7042/23462/1942"
=23%, signal=40%","493/51393/59341/5621/7226/162514/123041/10268/779/773/9187/799/5023/
signal=39%","3270/477/1906/23493/146/7042/54795/552/7168/153/147/3784/5144/133/1908/100
gs=73%, list=20%, signal=58%","51117/51004/51805/79133/10229/84274/51025/84273"
=45%","2303/7474/51176/7481/2296"
nal=39%","5580/4067/2214/2207/2213/6850/3055/5795/2268/5338/7409/26060/10451/26253/578
1936/1937/26046/1802/120526/1938/84164/6181/51611/8846/6726/9775/90850/60678/7284/19
list=38%, signal=50%","6224/28951/51343/7332/55294/550/6125/6135/6167/6015/6209/6201/577
3/4124/84061/4121/285203/117248/727936/6484/10678/57134/30815/9528/55808/11227/338596
%","2683/4124/84061/4121/285203/117248/727936/6484/10678/57134/30815/9528/55808/11227
, list=18%, signal=35%","3572/183/6548/1906/8654/3479/8013/5756/817/4205/57817"
%, signal=35%","3572/183/6548/1906/8654/3479/8013/5756/817/4205/57817"

signal=33%","28984/4851/654/1003/7474/7082/10076/4855/9647/3169"
nal=28%","2059/51429/23189/93663/11135/2934/23500/148170/25802/6840/6711/5580/387/916
%","3688/5166/6591/857/3678/4851/4170/4915/7088"
%, signal=44%","3075/11326/735/730/718/629/1675/732/731"
","4780/183/824/9531/4217/8837/26471/1906/93649/5468/23493/3479/6648/80149/817/3162/2
%, signal=49%","978/7378/27350/151531/60489/200315/140564/81602/164668"
%, signal=49%","978/7378/27350/151531/60489/200315/140564/81602/164668"
al=53%","29118/55795/9877/160419/55308/56943/55677/56001/728343/8888/84271/55746/5718
20%, signal=24%","2040/5784/161291/5587/29969/6672/81565/2737/4641/284/10519/9531/5474
al=29%","1346/387/3356/351/26471/3479/8671/54209/55022/706/5027/54541/65018/6774/5134
signal=63%","23522/7862/51147/7994/27154/11143/9678/84289/23774"
ags=37%, list=23%, signal=29%","1793/23189/1601/6281/4739/6239/10519/55679/3987/10979/407
list=16%, signal=32%","54984/27000/100134934/3183/6714/3178/1111/55226/65057/3192/7014/8
54913/51367/138716/6741/10799/10556/54974/54931/10775/3028/79897/11102/10940/10248"
',2590,"tags=60%, list=14%, signal=51%","367/4851/1499/5176/4824/6662"
/55704/84984/8481/8558/9231/57539/55764/2909/79989/124590/26005/51098/80184/11158/207
2/858/6281/8301/2013/4038/857/6711/727910/25978/221035/92421/6810/7357/11252/23180/57
17%, signal=45%","1601/4478/5962/7430/23011/55823"
7,"tags=39%, list=12%, signal=34%","158/2618/471/8833/5198/5471/10606"
tags=59%, list=31%, signal=41%","10252/161742/7042/6422/653/753/1942/114990/10253/4092/31
%","3280/3399/5166/4023/1906/9076/80149/55022/8644/1373/23236/445/6720/3949/5724/8128
%","22823/26147/57343/79813/57673/2625/2146/10919"
01/6599/86/6598/6605/6602/6597"
al=28%","302/141/26471/5899/55062/120892/4534/5868/5305/950/65018/23232/51375/128866/2
%, signal=35%","117248/11227/374378/337876/22856/79586/55568/2590/55454/26301/28/8693/5
070077,2664,"tags=62%, list=15%, signal=53%","2074/7428/51497/904/51574/8621/9025/51755/88
2332/131405/23405/9271/26523/4204/6659/29066/4904/23264/27161/6895/23369/3178/400629
al=34%","22925/3688/5321/183/7057/477/1906/51703/10550/6529/5027/552/2170/3553/6850/65
77/54762/5360/19/348/706/80765/338/335/9619/336"
nal=25%","4208/23704/857/1192/9948/477/79784/4734/2730/1634/845/6548/9096/351/6324/535
1%","3556/2239/4208/7058/27286/351/7162/10082/7097/8321/7474/347902/51225/22943/4915/
","1956/81565/2737/27185/3912/387/26468/1600/64805/1499/5649/285025/57142/85458/84750
nal=42%","183/151/6869/150/2852/6865"
st=22%, signal=57%","3913/4811/4643/23213/2131/55959/7490/1286"
gnal=29%","7049/57493/2022/1901/2303/1909/7040/4851/23493/7042/7474/2296/1832/4634/716
gnal=29%","56154/5885/7917/9985/9319/256126/10919/5347"
t=22%, signal=25%","389/23189/117583/3688/2934/4000/4134/10252/81565/3280/25932/4478/4
=27%, signal=32%","2059/23189/2934/25802/6840/6711/5580/91663/1003/5962/256949/5756/29
202333/84676/4635/8736/127294/4660/8850/4634/667"
"11340/10595/118460/22894/27044/54512/131870/129563/54888/29102/87178/51010/56915/10
st=6%, signal=38%","994/7272/993/995"
2059/51429/23189/395/117854/93663/11135/2934/23500/81565/148170/283/79649/25802/6840/
nal=56%","5978/27122/653/9563/1583/1586/650/1589/54361/476/1585/1609/1392"
=55%","147912/284/8837/5467/84466/3717/2247/2047/6495/6469/2660"
58,"tags=41%, list=14%, signal=35%","5787/5791/5803/5797/10076/5793/5796"
=41%, list=14%, signal=35%","5787/5791/5803/5797/10076/5793/5796"
%","79008/548593/25913/51750/7486/7517/84464/7014/9401/641"
=23%, signal=27%","3487/2308/5166/3269/5155/3953/4837/3356/5208/7040/351/5028/26471/516
%, signal=22%","7798/6788/367/8463/6385/30812/157506/9096/7040/7041/7424/7408/220/7020/
736,"tags=51%, list=26%, signal=38%","90993/29761/9709/120892/3309/23645/7376/1191/7466/70

'387758/5155/91663/7474/958/7481/3357/9590/2324/54986/5341/10211/1608/23171/7422/5590
/23332/84687/8697/9150/51547/91782/55844/6683/54443/11331/9735/27338/11065/113130"
signal=39%","9972/3842/26953/30000/3843/8021/79711/9221/9883/100101267/8315/3838/3837
30714/23347/6015/79677/7341/1017/23512/10336/55791/6599/7703/6598/3192/157570/1788/10
=35%","51398/55750/4836/158234/140823/83460/9141/285521/26520/285093/10328/3308/2651
tags=53%, list=12%, signal=47%","1601/57664/2719/84870/7474/6422/22943/22881"
nal=49%","2120/1909/4487/4488/6662/3091/6469/8200/3209/7849/5076"
, list=16%, signal=37%","966/2/710/11326/1604/2213/725"
","tags=83%, list=27%, signal=60%","6688/3162/1524/3965/2213/2294/11314/8832/729230/3598"
ist=18%, signal=27%","10413/3572/3280/688/639/1051/4851/654/51092/5054/3202/652/7474/604
ıl=29%","1997/7006/7070/3958/3105/10384/8767/2633/8934/80149/11118/7805/975/64332/1146
66/26271/3619/6790"
4365,"tags=69%, list=24%, signal=52%","5423/4292/5395/8888/4436/10721/9156/7374/2956"
%, signal=36%","158219/2246/26119/1066/348/6720/37/3949/1581/5562/338/9619/2330/9971/235
al=59%","29081/29063/875/29104/115939/27232/64745/56339/79918/63935/3276/79723"
3%, signal=24%","3556/2239/4208/85461/7058/4038/27286/27185/4734/387/26037/351/22899/71
ist=9%, signal=21%","4208/4323/2022/7040/1906/93649/30008/9780/3479/84525/6648/652/2216
, signal=29%","4208/1910/824/7057/64856/477/4929/2701/348/4862/706/5649/116372/5179/686
list=23%, signal=31%","55208/7334/6209/84101/6187/6201/9529/57761/23142/9349/5300/8312/8
ist=24%, signal=31%","117854/5359/3133/9111/3916/81844/3428/4067/11027/83666/5322/3430/
%, list=31%, signal=69%","2213/3135/10859/3071/3824/5788/3575/5532/3821/634"
360/5744/3199/23314/2736/4488/23601/5745/8994/63923/4088/6469/43"
43,"tags=33%, list=6%, signal=31%","857/10019/54625/9021"
signal=28%","2730/1634/6324/2697/120892/54795/3791/706/2702/2537/5027/7871/817/4092"
35%","22821/3710/3708/3709/55283/7226/53373"
t=26%, signal=41%","301/861/6711/8767/942/64581/5142/3553/9308/5336/920/4860/5144/3552/
0%, list=21%, signal=56%","2246/26119/1066/6720/1581/5562/9619"
gnal=41%","1601/5978/27122/654/653/9563/650"
, signal=34%","683/4208/604/23529/4773/7099/1880/958/975/972/57162/10673/1026/3071/1480
","25959/6591/4023/4330/2350/1277/6422/2242/6256/2621/3162/5179/1030/7421"
167/79660/3122/3123/64581/648791/5806/5507/8909"
5%","157506/220/8644/29785/125/339761/59342/10170/126/83875"
gnal=60%","5434/5202/8409/55011/10471/81572/10856/8607"
514/2060/684/53373/440400/30835/6614"
3%, signal=29%","10413/7049/4208/7048/861/2735/4851/1906/23493/3479/10370/60485/2627/96
943,"tags=38%, list=11%, signal=34%","183/114907/6442/1605/84676"
ist=29%, signal=38%","6934/1499/29940/9394/2131/3340/26035/9375/2132/9653/266722/64131/
0/8321/8324/7474/6422/7481/4693/7483/7484/7477/25805/54894/4653/115908/7475/4920/625
ıal=45%","4023/8542/348/51599/344/350/338/335/51449/341/336"
%, signal=45%","4023/8542/348/51599/344/350/338/335/51449/341/336"
24/7040/54762/1066/1605/23209/3625/1581/3638/6348"
364/857/9261/7376/5538/29108/1794/998"
6%, signal=35%","4643/2200/9314/3791/2202/10018/3717"
2308/5547/5836/5166/3953/25825/3488/2730/6814/55107/5468/80024/59341/51092/6934/6517/
5547/5836/5166/3953/25825/3488/2730/6814/55107/5468/80024/59341/51092/6934/6517/5479
%, signal=34%","1956/493/857/10392/4883/348/5291/6271/3357/3553/23576/3091/1268/2185/35
signal=33%","3685/8406/6591/355/3624/9531/4254/1495/4170/2260/2065/2668/6376/2139/2537
gs=45%, list=26%, signal=33%","3685/8406/6591/355/3624/9531/4254/1495/4170/2260/2065/2668
26%, signal=29%","8801/5166/84836/80221/9060/23597/5167/54988/51703/8459/10157/64834/5
list=26%, signal=29%","8801/5166/84836/80221/9060/23597/5167/54988/51703/8459/10157/648

)%, list=26%, signal=29%","8801/5166/84836/80221/9060/23597/5167/54988/51703/8459/10157/6
ignal=64%","10196/80204/10498/54496/55471/55170/10419/3276"
t=11%, signal=64%","10196/80204/10498/54496/55471/55170/10419/3276"
=64%","10196/80204/10498/54496/55471/55170/10419/3276"
308/5611/6548/4792/9451"
st=5%, signal=44%","54962/8317/80174/10926/58525/81620"
33%","677/4323/3624/6422/3586/678/3398/3635/6850/3071/150372/695"
signal=30%","3074/285203/10678/51301/93010/3073/4245/3955/2131/4242/2134/51146/2650/10
=24%, signal=36%","351/8878/11170/5641/27020/7305/348/5649/10288/23385/1524/5621/4311/5
, list=17%, signal=41%","4907/1806/7378/1807/272/9615/4860"
ignal=47%","3673/3075/10468/4023/348/4035/9037/129804/375790/60495/5788"
70,"tags=67%, list=29%, signal=47%","996/9587/51433/8697/8452/9787/51529/51434/9700/81620"
ags=67%, list=29%, signal=47%","996/9587/51433/8697/8452/9787/51529/51434/9700/81620"
9/64129/8301/8724/5872/30844/30846/257364/7813/1203/54741/9341/54918/25978/83547/8878
'57088/161291/5205/5359/57194/5580/23120/5243/19/5244/55754/79895/5027/81671/114786/1
%","6045/54623/84108/6015/4591/84759/10336/56970/1643/56254/165918/1642/7703/123169/8
","57728/79809/92104/64792/26160/57560/51668/150737/23059/57539/55764/112752/79989/51
4566,"tags=73%, list=25%, signal=54%","904/8941/4331/905/8851/1164/1163/891"
l=38%","2040/858/80149/1436/348/5806/6355/3429/3482/9218/4057/1401/5479/9189"
8,3726,"tags=54%, list=21%, signal=43%","3572/7040/5743/728/3569/23213/718/719/2324/3553/1
%","5429/50485/7158/6419/124817/1386/6118/23347/55100/9025/79677/79840/10111/51547/165
.16039/1909/5396/1906/3199/4487/130497/3638/6495"
25%, signal=53%","2934/5205/4627/57194/5580/23207/51100/79400/356/8976"
854,"tags=56%, list=21%, signal=44%","7049/90/4052/7048/2022/8425/94/659/130399/25928"
gnal=20%","10413/4208/90/8463/7048/3280/2022/64321/4853/9645/4851/2697/7474/10370/2851
","284/2131/3340/26035/2132/84068"
, signal=32%","857/477/5350/54795/5142/2702/1824/1832/817/6781/5144/29119/3759"
gnal=47%","81031/116039/652/1499/130497/7483/2131/3373/3227"
0%, list=37%, signal=50%","54795/2702/1824/1832/29119/51086/6910/3741/6336/5318/6331/1825
8%, list=21%, signal=38%","90102/395/84952/1901/4070/10163/7082/10391/23671/10395/5295/50
%","5978/27185/6548/1728/6121/929/6648/199/7832/4313/3164/5027/84676/9590/6869/5728"
l=28%","1601/1410/85004/183/64321/3624/978/604/58189/56920/7040/10512/2697/5167/7042/7
727/23405/26523/27161/6895/29102/8575/144233/389421"
5606/11218/50628/79760/25929/8487/79833"
,"4831/170506/9361/7486/10111/80119/1663/641"
%","7057/7412/3586/7128/1588/8876/6279/6352/6280/10563"
al=43%","1969/10163/5756/51225/11344/998/832/10096/23191/5295/51196/7454/10097/5879/1
%","5611/351/4642/3309/57348/7871/338/3759/821/3284/811/5479/10525/3283"
=34%","7026/3280/4853/3726/23493/9021/5604/8061/23462/9394/113026/3696"
./6403/4017/3965/85329/130120/55600/6402/142683/5068/78989/5968"
"9429/10351/23460/10350/5243/6892/19/6890/5244/10349/6891/340273/10060"
3913/3691/79628/10397/1605/5339/4915/400746/6497/163175/4653/3611/57716/5453"
5,1149,"tags=46%, list=6%, signal=43%","904/645121/8941/1031/85417/4331/9134/905/8851/10983
al=59%","3569/3570/6774/3594/149233/10538/51561/3662/4063/30009/3593/114836"
al=65%","57703/54883/55696/10594/22794/9775/27339/9939/55110/23020"
"57703/54883/55696/10594/22794/9775/27339/9939/55110/23020"
%, signal=28%","10413/7049/4208/2735/4851/1906/23493/3479"
t=24%, signal=57%","5423/4292/5395/8888/4436/10721/9156/7374/2956"
5, list=17%, signal=25%","29969/6672/284/9531/347/120892/11261/27248/51715/10956/3638/1114
, signal=42%","301/861/2737/604/10875/80149/3566/652/3659/51043/972/9655/7292/3142/4092

3/9388/3990/8605/23659/51365"
nal=49%","7107/84959/3569/7128/5027/3635/338773/57817/4982/796"
al=36%","4643/4641/55930/4642/81/4644/4648/4542"
3%","389/7094/5872/30846/9948/2037/388/387/29984/59341/23136/391/91010/5339/7430/2343
, signal=31%","2697/6892/51296/6890/6891"
63/1017/1111/9031/983/7443/9212/6790"
55%, list=22%, signal=43%","8826/80310/5155/5159/5962/7099/54206/3037/4609/3373/3036"
, signal=52%","5136/27115/5151/5137/5142/5153/5139/8622/5141/5144"
%, signal=48%","7048/975/57045/29126/1604/3965"
=55%, list=13%, signal=48%","7048/975/57045/29126/1604/3965"
l=40%","367/652/7474/23213/8626"
'1499/4488/3625"
=35%","22925/301/51129/6272/341/336/5997/440503"
gs=62%, list=32%, signal=42%","3133/3916/135250/3592/3134/5817/7409/3822/56253/3823/8302/
5%, list=26%, signal=34%","4780/90993/29761/857/26471/9709/120892/3309/23645/7376/1191/74
'5742/5743/4258/2878/1193/4257/26034/847/241/373156/5052/4353/2876/4259/3048/2946/953
,"56898/4780/5066/9839/5446/400/7474/8644/57016/126321/2327/27233/2326/1545"
1984/2332/6728/57003/26135/10196/27335/3692/56926/3921/23480/83939/79612/23107/25875/
%, list=6%, signal=26%","10413/7049/4208/2735/4851/23493"
gnal=54%","11031/3673/54209/2150/718/5336/3071/11314/196527/948/10347"
al=54%","11031/3673/54209/2150/718/5336/3071/11314/196527/948/10347"
35/4853/55679/2246/8837/1051/3569/7128/2267/7039/5764/3249"
s=57%, list=22%, signal=45%","2735/4853/55679/2246/8837/1051/3569/7128/2267/7039/5764/324
99/1832/1674/4892/29119/87"
329/387/1728/1393/3779/6915/9076/7412/929/761/6648/6517/3709/6916/847/7122/5179/841/6
4061/4121/285203/1203/117248/727936/6484/10678/57134/30815/9528/55808/11227/338596/3
5%, signal=32%","23704/857/845/28954/2669/1193/7881/6536/9446/6236/23630/2946/51185/181
5/27233/2131/8460/6820/3340/27284/6783/2132/6817"
034/80310/1956/1290/824/1278/1307/1282/1281/2730/387/1728/1051/1906/4067/1277/4313/64
gnal=32%","3280/4478/387/2195/7474/64083/55691/998/23150/2294/399687/9265"
54/387/56920/10512/7474/3690/59352/10501/9037/23768/335/1803/7869/91584/6586/6259/10
2%, signal=36%","7538/947/2633/11326/7805/7128/84868/9454/7430/5621/64127"
'6281/8301/2013/4038/857/6711/727910/25978/221035/92421/6810/7357/11252/23180"
6%","51727/8801/1969/4907/5166/1806/84836/387/80221/978/9060/5744/3356/5208/7040/351/
54%","58494/1075/9451/5728/27242/7124/9896/8898/11202"
941/27089/429/401/8851/23513"
3%, signal=29%","3270/477/845/5350/1756/1193/817/9446/6348/5144/10269/6588/1234/2946"
list=12%, signal=39%","6239/9580/284207/5743/338557/51043/7030"
2/2246/157506/2099/652/7474/6422/7481/10253/6662/6495/374/51804/6469"
'7098/9111/23586/64135/8767/7099/3586/84868/51284/3665/51311/6772/3661/29110/3663"
signal=43%","7098/9111/23586/64135/8767/7099/3586/84868/51284/3665/51311/6772/3661/291
, signal=35%","7132/2308/4000/90523/493/3488/4851/5468/54206/7133/84676"
9/728/54209/3569/712/4035/3553/3949/9572/2896/3458/2358"
ıal=56%","2/710/4153/4239/8547/2219/3848/2220/78989/5648"
%, list=15%, signal=42%","1903/1901/8877/5291/5294/7430/53637"
7506/6121/220/8644/1545/339761/83875/195814"
8/4023/3990/19/54209/4035/3931/821"
signal=49%","2908/4734/5187/406/3717/1408/811/7533"
,"604/10875/3122/3123/7292/3594/149233/4179/5105/51561"
%","10769/5906/5962/5912/5908/10411"

:0%, signal=53%","2151/10788/2149/2150/9002/3248/6622/5341"
43%","3280/4851/5324/4067/5583/3569/706/6497/3553/1030/4609/23376/29956"
signal=34%","5587/29969/2737/284/9531/5580/7040/347/5743/80149/652/11261/2316/3059/869
1%","127733/54778/55968/55666/11124/55728/55208/51366/3720/23062/9025/10208/8975/7334
nal=56%","23710/81631/10533/643246/440738/84557/9474"
56%","23710/81631/10533/643246/440738/84557/9474"
27910/9388/57333/19/2170/335/3694/9619"
.1%, signal=31%","3956/4853/6778/604/7040/23529/9452/7099/602/1880/23075/958/3586/3569/
897312,4078,"tags=67%, list=23%, signal=52%","2668/652/1499/6299/7484/3976/6772/26585"
%, list=19%, signal=54%","147912/284/8837/5467/3717/2247/2047/6495"
/6548/5142/5179/3784/6622"
1277/7020/4488/54757/56975/6786/10312/123041/3694/5465/9313/7475/1080/401138/6522"
0%, list=27%, signal=36%","117248/11227/374378/55568/2590/8693/50614/79623/79695"
6203723206817,4639,"tags=62%, list=26%, signal=46%","329/8767/51296/7099/4792/144811/7128
/114548/79792/834/29108/838/4210/171389/22861/22900"
al=27%","283/5155/2246/151/3479/7474/2621/65018/4908"
0%, signal=40%","123920/4067/5583/975/2213/11314/150372/26191/27086/9404/148823/257144"
list=27%, signal=29%","25932/183/1203/477/23682/6548/285195/55647/120892/8992/8671/6337/
signal=36%","3688/7057/7040/59/9590/4921"
304/6342/54988/3032/123876/30/6296/348158/10558/23464/38"
'5587/29969/2737/284/9531/5580/7040/347/5743/80149/7984/652/11261/2316/3059/8692/5171
57/10392/351/2335/834/29108/65018/1075/4824/91662/4118/1512/5720/22861/10010/7415/229
4/54840/1161/7515/7014/142/55775"
:ags=77%, list=21%, signal=61%","80310/5155/8837/1958/652/3570/4609/201163/7490/8710"
list=25%, signal=45%","143872/5547/9424/6648/9365/153/154/9283/659/134/796/585"
016/8559/54464/55795/11340/118460/54512/1797/51010/56915/5393/23404"
23016/8559/54464/55795/11340/118460/54512/1797/51010/56915/5393/23404"
:=13%, signal=44%","301/5743/972/552/3553"
signal=53%","2668/652/1499/6299/7484/3976/6772/26585/342184/7849/5076/6608"
5959/6591/4330/590/2350/5743/1277/6648/6422/5467/2242/6256/706/847/2621/9365/5179/894
signal=58%","147912/284/8837/5467/3717/2247/2047/6495/6469/2660"
ags=64%, list=25%, signal=48%","3269/5028/5031/5745/5724/553/6622/2852/51744"
.4%, signal=35%","4323/604/23555/23493/51162/4792/23462/28514/221336/1745/51341/3955/54
st=25%, signal=38%","4208/2697/2065/4534/1499/28514/5562/56203/3481/6469/10211"
9023/2246/51700/10826/1066/348/1727/6720/51422/1581/6309/5562/3638/338/335/51706/3158,
L/8661/9140/8664/8666/51386/28985/5094/8562/8665/8662/1660"
st=19%, signal=42%","5136/27115/8654/5151/5137/5142/5153/5139/8622/5141/5144/5149"
=31%","7049/6444/5159/161742/4851/23493"
9/56947/10939/664/581/7037/23788/54974/201164/4976/55486/30968/54332/57143/10075/5492
","26013/23512/8467/7528/23378/11335/22893/2146/6839"
:=29%, signal=28%","10404/285671/9839/1909/1513/4929/26002/7042/51301/7474/2034/1508/53
%, signal=30%","301/5166/5743/7376/972/552/9104/3553/37/1581/344/3638/335/341/51085/604
ist=20%, signal=40%","7132/2308/4000/90523/493/4851/5468/54206/7133/84676/221476/5025/4
32%","285671/29958/3176/590/23446/3067/5153/6581/121278"
0/183/857/5580/1281/5743/57085/185/4790/9365/1535/4609/1536"
6/11037/81565/8301/26503/284612/5872/27185/3916/351/120892/127262/4803/8676/3479/797
%","4323/5069/79148/1889/9510/8754/170689/81792/4327/7092/56999/388630/4313/80332/10
ignal=35%","84876/7225/93129/7223"
87/207/22954/51455/7486/2068/10714/1965/1161/84446/83932/1643/51499/79913/7398/7528/
70,"tags=62%, list=29%, signal=44%","996/9587/51433/8697/8452/9787/51529/51434/9700/81620

l=29%","79443/9531/10392/1634/26353/5899/55062/8767/10133/10241/3791/5305/65018/3162/
692,4261,"tags=42%, list=24%, signal=32%","27124/2258/57159/23332/25876/3925/10297/22919/7
/114548/7097/54209/64127/84659/130120/5068/3098/114770/5968/57115"
9%","2072/8658/51750/10111/5609/7874/3184/4751/7515/7014/8550/79035/9212"
=33%","5205/57194/23120/5243/5244"
gs=50%, list=25%, signal=38%","7412/6403/3791/5621/6364/7226/5795/4935/5788/5737"
392/5742/5743/55788/51302/9420/55640/79689/6916/847/2977/3162/1545/29785/57834/1535/
337/688/5420/7430/83715/53827"
29%, list=19%, signal=24%","23189/1601/3280/7070/5376/4038/567/7431/10519/387/56920/10512
,"115939/54433/55651/27079/113000/1736"
075/3383/2621/3625/4057/30835/131578"
list=17%, signal=34%","4780/183/824/9531/4217/8837/26471/93649/23493/817/2627/3717/4058"
%, signal=45%","80310/3554/7070/10894/3383/83483/83700/5724/7499"
","54885/143872/4739/1969/85360/141/81565/83478/7813/30851/26084/26037/22899/7984/8414
7/4067/1326/975/942/10288/7292/29126/8764/80380/10673/1947/1803/7409/940/941/8740"
074/5580/4758/2629/3073/4668/2581"
=29%, signal=58%","4067/10288/945/3135/10859/387837/5678/11006/3821"
al=74%","54913/51367/60528/10799/79042/10556/80746/10775/79897/11102/10940/10248"
460/54512/6832/1797/51010/56915/5393/23404"
nal=51%","6472/5631/2618/8833/5471/10606"
34%","2624/26155/55011/4800/55677/6879/84656/80764/2648/6871/11143/4089/1487/25942/29
/2984/4881/2977"
%","10056/155061/4144/11052/27153/144404/11051/84869/2193/6241/79869/6240"
","3624/54762/1066/1605/23209/3625/1581"
gnal=29%","4780/5978/5580/4217/3082/120892/4170/8013/6648/8321/7099/5151/3586/54209/14
35%","10226/4023/4481/80149/4790/7376/718/32/338/123"
al=38%","2200/4853/7305/4057/10859"
=17%, signal=26%","2308/2737/2735/2771/55679/27185/10763/387/5803/7424/4851/4067/120892
/10577/6342/54762/55207/19/26119/1066/5305/81537/2170/3949"
st=23%, signal=39%","493/2274/3708/11261/9454/5621/1827/123041/5341/58529"
=27%, signal=35%","23189/183/23371/2771/5580/5791/5167/9021/51043/55022/340706/5305/896
","5321/5580/151056/79887/9388/3990/8605/196463/5322/23236/23659/3949/5336/344/145447/
/1345/1340/7388/10975/125965/1349/1350/1352/29796/1347/27089/7386/440567/341947/7385/
124/23268/6484/3105/83871/54922/9528/337876/23213/22856/170384/643699/79586/25963/51
ist=23%, signal=77%","5093/54386/80169/25913/6118/6117/3297/10111/3178/3181/7014/79035"
ignal=33%","23328/4616/4217/1647/8877/80149/652/768211/6688/9867/3553"
5166/1203/23682/6548/285195/1906/55647/120892/8992/8671/11261/5172/144811/5538/10159/
=26%","5445/5321/116236/11343/151056/5446/4023/63874/590/9388/3990/57552/7130/8605/23
9%, signal=46%","4208/4000/2274/93649/23493/5915/4205/9474"
signal=39%","6241/1854/1716/1841/7298/6240"
%, signal=39%","6241/1854/1716/1841/7298/6240"
%, signal=39%","6241/1854/1716/1841/7298/6240"
55%","5547/5476/1200/54504/59342/1362/3249"
, signal=72%","5725/6741/8661/8664/28985/5094/8562/8665/8662"
ist=26%, signal=43%","29969/284/347/11261/51715/11142/948/51569"
al=43%","29969/284/347/11261/51715/11142/948/51569"
5468/5467/241/2170/6279/5306/165140/6280/2167"
t=18%, signal=44%","22938/9150/51193/6882/64710/6598/7023/23076/6597"
2300383344,4643,"tags=45%, list=26%, signal=33%","90993/837/4217/1051/9709/120892/84216/37
=23%, signal=42%","23213/3357/5724/80763/55959/6865"

signal=35%","5327/3624/2730/1909/4851/1906/2627/1545"
2983/4976/84446/2011/131870/5495/79178/129563/51251/83983/5636/5631/85865/5609/26164
0%","1723/6472/5631/2618/790/8833/1503/56474/5471/10606/7372"
7,2850,"tags=43%, list=16%, signal=36%","56034/284/6886/1950/5618/154"
, signal=25%","64764/5611/23471/4780/90993/11080/7057/871/9531/26353/9709/8988/467/1013
7/5552/1992/127544/3002/6813/811/2999"
340/10595/118460/22894/54512/29102/51010/56915/54555/5393/23404"
nal=56%","5580/5583/54209/6347/28514/2621/9590"
51364/54919/55172/10963/57418/10856/8607"
, signal=38%","57664/5321/8724/55614/55062/26499/285590/4689/23612/57158/90203/29934/28
st=19%, signal=42%","2/3133/1991/2214/718/2209/2213/2131/1268/133396/171389/695"
100%, list=25%, signal=75%","9013/9014/64425/54700/55011/6872/6598/7343/6597"
853/55679/2246/8837/1051/3569/7128/2267/7039/5764/3249"
%, list=21%, signal=20%","10413/25937/7414/2059/23189/2012/3688/1969/4651/11135/2934/235
ist=11%, signal=48%","3280/4851/3204/28514/9573/4488/8626"
1%, signal=48%","3280/4851/3204/28514/9573/4488/8626"
398/10627/2934/57477/387/103910/54558/1756/4604/79083/5873/29108/4635/4634/66008"
signal=53%","9580/51176/1960/64127/6850/10859/3071"
,"1601/5978/27122/654/653"
=32%","23704/857/477/9096/6324/2697/6330/1756/3752/54795/2702/2316/1824/7871/1832/376
%, signal=28%","9429/81031/25932/1192/6548/8542/55107/2697/285195/89874/5348/6578/6560/
al=34%","10413/182/117854/8463/4780/3280/5978/4070/3214/4851/7042/2668/678/221662/144
%, list=31%, signal=39%","4846/5957/54209/55022/65018/4609/7114/5465/7415/2978/136/2980"
ags=57%, list=31%, signal=39%","4846/5957/54209/55022/65018/4609/7114/5465/7415/2978/136/
357/2697/1495/2702/2705/2706/3553"
al=26%","51727/8801/1969/5836/4907/5166/1806/84836/387/80221/978/9060/5744/3356/5208/7
","3673/183/1901/79784/6324/6332/2629/5339/552/4915/4099/3737/83700/1134/3736/1890/103
=50%","477/6548/5350/5348/10060/482/3764/483/481/476/486/480/496/23439"
%, signal=56%","7388/10975/29796/27089/7386/440567/7385/1537/617"
","7388/10975/29796/27089/7386/440567/7385/1537/617"
27%, signal=51%","5978/27122/653/9563/650/54361/476"
9%","975/3383/2621/3625/30835/131578"
0/4067/6403/54209/1200/5538/3306/54749/51228/10026/4099"
nal=25%","1601/3433/3077/5155/567/857/5580/27185/10763/5899/120892/3434/3397/3586/3676
,3738,"tags=27%, list=21%, signal=22%","3688/83872/10205/23022/91624/28513/1009/54510/9095
10%, signal=29%","10413/3572/688/639/654/51092/3202/652/7474/653/5916/28514/406"
3%","6472/2618/471/441024/4522/1719/7298"
,"7431/10763/286077/4534/3861/23336"
=26%, signal=57%","11031/3673/54209/2150/718/5336/3071/196527/948/10347"
6%, signal=57%","11031/3673/54209/2150/718/5336/3071/196527/948/10347"
3%, list=19%, signal=31%","23704/9948/6324/2697/6330/2702/2316/3784/10269/23630"
9%, signal=46%","80218/51126/254042/2033/28989/5494/64146/122830/6659/60560/79612/4836/
2%","51474/2059/3915/3920/4323/2934/81565/57326/79443/6591/25802/6840/9645/6711/10519
70506/10642/4904/79066/8148/23367/23185/54888/3183/56478/8313/3184/3192/2521/1994/718
, signal=35%","7070/9314/91663/3586/4155/3553/335/3557/960/10211/2266"
2%, signal=48%","3688/183/477/5243/10550/5621/8737/5724/285966/8501/124935/1080/59272/8
ignal=23%","5154/5155/8871/3707/5159/23167/3356/10903/8395/5333/80271/84668/8821/9021/
5/4804/4915/6272/9001"
6121/5360/19/57488/950/335/5306/341/51652/57605/336/5104/5130"
24%","7414/2239/57493/84952/5906/10979/3716/5175/1003/10082/7474/1605/7082/3315/5649/

15%, signal=34%","3556/3133/114548/1051/942/7292/84868/2207/3965/6850"
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","10445/129563/27161/3508/23210/9987/9716/284131/1653/3183/124540/80324/79882/3178/2
4/56123/124590/51098/403/9657/80184/9742/126731/5108/261734/123016/6903/84747/1070/5
signal=27%","3685/9388/347/2629/3690"
553/153/154/1268/3552/9536/376497/8792/57561/7124"
%, list=15%, signal=39%","6591/5054/23075/1545/83700"
378/120892/2512/8031/84557"
6%, list=23%, signal=28%","51474/2059/2934/79443/25802/6840/6711/10519/130271/9948/10763,
=11%, signal=25%","845/9096/6324/2697/6330/54795/2702/2316/3762"
31/64764/11015/10749/5587/8724/5872/51226/55614/6711/9341/3916/10791/123720/2060/400/
gnal=30%","3077/5208/55107/6934/2746/338557/54795/6810/5467/3123/2752/7226/2182/51762/
4/2335/8291/945/8692/102/11006/27086"
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3678/3356/1051/123803/5743/6285/4803/5641/3710/6517/54209/348/5649/6529/23236/5176/15
35963419162,5484,"tags=69%, list=31%, signal=48%","3678/3690/3553/8817/3091/2887/7448/405/
al=25%","2040/7402/493/477/4734/154043/6548/6324/285195/80024/5348/6330/1756/5349/633
al=25%","8801/1969/4907/5166/84836/387/80221/9060/4846/5744/3356/5208/7040/351/27115/
7%","10413/8463/2303/7474/51176/7481/2296"
38%","1806/978/2519/2629/7378/27350/4668/151531/60489/9615/4860/200315/140564/81602/1
"10979/3801/2149/1436/7082/2150/57126/5657"
signal=32%","54514/10146/9271/167227/26523/80336/4848/6191/23112/11201/9898/9118/11022
0485/8087/1975/84083/8438/9401/641"
=32%","22821/3710/3708/3709/54795/55283/7226/53373/5025/2904/5311"
%, signal=27%","2040/302/1601/161291/5587/4478/30846/2737/4641/10519/9531/54741/5580/47
gnal=49%","1511/1991/2150/54210/4615/171389/6372"
","92292/23409/440/790/8833/5198/1503/56474/5471/89839"
=25%","1601/1910/5978/27122/216/157506/223/654/6121/220/653/8644/1545/339761/83875/65
=17%, signal=31%","4780/5978/3082/120892/4170/8013/6648/8321/7099/5151/3586/54209/1499,
11108/55352/10196/80204/65094/10498/54496/55471/55170/10419/3276"
8/7042/7474/5187/4804/406/4488/8717/162514/25987"
54%","6045/84108/6015/4591/84759/10336/1643/165918/1642/7703/84678/54880"
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01/3077/567/55640/79689/57348/392/64116/6556/23479/55652/2512/6536/30061/4057/286133/
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gnal=53%","140609/114548/7099/54209/10333/5336/4210/4615/22900/948/373509/80772/22933
, signal=28%","182/3280/2737/567/5978/30812/387/351/4851/3199/58190/4854/23462/3400/285
al=34%","84269/57486/4713/9131/10989/80025/23408/10456/55750/27429/4702/4976/26520/514
6","92104/64792/26160/57560/51668/150737/23059/79989/51098/28981/11020"
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57%, list=25%, signal=50%","51132/23347/3720/23512/64783/10336/91746/55791/56339/29890/31
1/27295/56940/9260/83715/23607/83700/8572/64236/23301/6525/254102/10787"
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49/182/4000/6444/10252/183/5159/4629/2274/9096/4851/1906/11214/23493/3479/652/5339/42

6045/51366/84108/6015/4591/9025/84759/10336/1643/165918/1642/78990/7703/84678/55611/5
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nal=32%","5208/55107/6934/2746/54795/6810/5467/3123/2752/7226/2182/25874/3717/3091/264
%, list=7%, signal=21%","143872/23189/3688/57493/10252/83478/604/81848/54922/22821/7042"
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gs=67%, list=26%, signal=49%","23097/8621/51755/1024/23476/1019/1021/983"
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45,3143,"tags=32%, list=18%, signal=26%","857/3270/477/2697/2702/817/482/5144"
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signal=44%","493/6548/5142/3784/6622/22933/5313"
%, signal=40%","8301/3459/4067/3479/10418/348/114134/1191/1942/4915/23385/5621/57142/10
t=23%, signal=36%","5155/79365/4851/93649/23493/4773/8324/4487/3398/28514/22943/338773/
48%, list=19%, signal=39%","4780/284/7010/1003/5054/3791/51129/3383/7128/2621/9037/79625/
42%, list=13%, signal=36%","301/5743/972/552/3553"
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"183/477/6548/6337/6507/482/6558/3552/483/481/476"
3304,5311,"tags=78%, list=30%, signal=55%","10252/90993/7057/81848/7474/4804/23213/10253/
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1%, list=22%, signal=34%","2258/57159/23332/25876/10297/79929/10013/26586/157922/10300/22
33%","90/7048/3624/10272/10468/6997/10370/94/6497/4092/3625/659/60436/130399/4088/228
nal=44%","5321/151056/3990/8605/23659/10908"
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08,0.0166517729009427,3230,"tags=22%, list=18%, signal=18%","301/10203/183/3269/3356/6751/
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33%, signal=60%","7376/338328/344/51099/350/335/5406/337/10062"
%, signal=62%","90/4681/1499/64853/64388/1649/25928/26585"
29%","10981/967/23682"
al=29%","10981/967/23682"
0981/967/23682"
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57%, list=21%, signal=45%","7098/7099/958/10333/3665/353376/148022/3661"
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29%","677/84870/9021/51176/3586/653/6768/7477/133"
1906/7412/4313/4790/3775/116372/3162/5179/6869/3784/1268/5594/2876/1140/5465/1813/255
gs=43%, list=27%, signal=32%","9469/56548/10090/55034/8459/4338/83539/27233/9394/8460/995
=30%, signal=44%","134266/131474/130617/100287932/9409/133022/10440/613/5190/5830/6729
gs=71%, list=19%, signal=58%","9972/55051/55677/26993/7175/11052/54888/10657/1660/22916"
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, signal=41%","3554/1906/728/23604/972/2526/719/3576/3071/3552/7060/7499/90550/5879/392
, signal=52%","1893/5167/5552/54795/6662/1230/6348/3091/2185/4054/26585/6779"
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)/358/1906/3708"
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9%","3420/79944/2806/3418/84706"
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78252839818056,2084,"tags=38%, list=12%, signal=34%","64114/8682/85450/4846/8837/3082/505
50%, list=11%, signal=45%","4780/3082/120892/8013/5151/3586/65018"
s=50%, list=11%, signal=45%","4780/3082/120892/8013/5151/3586/65018"
5671/493/29958/4837/8639/6303/1909/3176/23498/4929/26002/7412/7042/2034/11185/27163/4
l=41%","157506/9249/8644/125/10170/126/50700/195814/124/145226/5959"
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80/5583/54209/6347/28514/2621/9590/27242/5925/842"
30%","3673/4208/3958/2771/5159/255743/7042/145376/11261/23645/3672/9104/945/133522/18
1%, signal=36%","6209/84101/6201/9349/84930/991/26271/4085/27338/11065/5347"
L/2150/5594/87/58529/5595"
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358/4478/8724/5872/9341/6814/5899/8676/7412/57488/11261/6810/8417/3383/7430/81671/787
al=41%","367/1393/2099/10865/6422/5272/4609/8031"
7007,4668,"tags=69%, list=26%, signal=51%","1839/1950/685/2069/7039/374/150/134/653361"
signal=35%","84656/2648/6871/11143/4089/29117/3670/6598/26053/10856/22976/2625/672"
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gnal=46%","5028/9934/2857/64805/5027/5029/5031/5724/286530/5025/53829/5023/9127"
"5028/9934/2857/64805/5027/5029/5031/5724/286530/5025/53829/5023/9127"
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62%, list=39%, signal=38%","4719/4709/4708/1738/4698/4714/4700/4696/4695/4720/4710/4704/4
79727/10521/131405/23405/54542/3265/27044/129563/27161/29102/87178/144233/8570/38942
t=17%, signal=50%","7048/85301/309/5916/6662/25987"
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t=20%, signal=35%","2308/4000/90523/493/5468/23493/54206/2627/23125/4088"
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=24%","5321/2052/11343/5742/8605/5743/8644/6916/1545/29785/57834/66002/240/113612/155
list=22%, signal=27%","8531/259230/4780/6387/90993/6591/138151/857/624/26471/9709/5743/1
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signal=47%","2908/4734/5187/406/1408/811/7533"
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al=42%","2059/81565/64805/54869/23191/22871/23242/5216"
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830/6729/9972/3842/30819/26953/9804/30000/3843/55176/8021/6730/79711/9221/387990/9883
l=46%","6484/10678/93010/11046/9435/6482/10331/4166/10402/11041"
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%","8322/4919/8321/8324/8323/11211/2535/8325"
56/857/4067/1326/975/942/10288/7292/29126/8764/80380/10673/1947/1803/7409/940/941/874
61/2099/7474"
2380,"tags=23%, list=13%, signal=20%","307/4208/6867/367/196/2114/4780/29969/3280/85004/7
36%","8528/443/2746/2752/2747/122970/137872/10840"
signal=41%","9394/9957/3340/9653/9956/266722/9953/8509/64579/222537"
%, list=31%, signal=34%","79053/9526/91869/79796/144245/84920/10195/79087/57171"
574,3628,"tags=60%, list=20%, signal=48%","5169/3162/3965/2294/11314/8832"
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18%, signal=30%","10642/4904/79066/8148/23367/23185/9704/3183/56478/79048/8313/3184/31
%, signal=37%","3688/183/477/6507/10550/6583/6542/6536/23428/55244/6582/8501/124935/547
872/9341/25978/55647/83871/8676/338382/55207/8773/8291/8877/5322/6810/8417/5305/5480
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%","2902/3858/5310/10056/11052/11051/84869/2193/6241/79869/6240"
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al=46%","9780/117531/59341/9725/11322/63895/79905/147798/342125/441027"
22%, signal=28%","10413/25959/1601/367/2908/26508/56937/4306/4734/387/2099/5241/8013/94
, list=20%, signal=48%","9314/91663/3586/4155/335/3557"
0%, list=20%, signal=48%","9314/91663/3586/4155/335/3557"
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=43%","81605/90353/55140/27304/55687/84705"
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al=25%","7132/8531/259230/1969/4780/6387/90993/79073/6591/138151/837/857/5580/624/4217
,"9124/23022/27295/9260/8572/64236/4892"
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3368,"tags=79%, list=19%, signal=64%","134266/100287932/293/10440/140823/131118/26519/510
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3%, list=27%, signal=53%","1435/860/652/4804/650/4982/340419/147495"
32%","55764/56123/124590/51098/403/9657/80184/9742/261734/123016/6903/1070/57670/2332
/6778/1514/3569/972/3570/6774/1794/3594/6469/149233/5788/10538/51561/3662/4063/38735
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41%, list=19%, signal=33%","4067/11027/338382/54209/11213/55366/2150/7128/10333/56927/64
14%, signal=29%","57664/8724/55614/55062/26499/285590/23612/53373/57158"
23%, signal=49%","54209/55022/65018/4609/7114/5465/7415"
, list=10%, signal=16%","9429/493/10351/477/23460/8992/10350/5243/6892/19/6890/5244/10349
t=23%, signal=37%","84128/55011/55127/51547/4691/11137/23160/56919/23133/6598/25926/73
15%, signal=36%","10468/7042/7474/4488/8717"
%, list=10%, signal=31%","252983/7015/57521/79447/27085/5356/595/6909/163786/6491/6241/55
3%, signal=30%","3356/351/26471/79660/3479/8671/5027/54541/950/6774/51341/51422/5562/46
"51082/5434/64425/5437/9533/84172/25885"
, list=20%, signal=33%","2308/10114/1843/153769/2122/6422/5187/54209/2150/65018/7430/1408
9,3585,"tags=41%, list=20%, signal=33%","2308/10114/1843/153769/2122/6422/5187/54209/2150/
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3%","5321/2687/4258/5322/241/6850/240"
1%, signal=33%","196/3140/3123/84868/29126/3592/5817"
s=45%, list=18%, signal=37%","5327/2149/27020/4915/1268"
067/8773/8673/6581/695/6582"
29/7097/10333/912/3929/911"
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1/4487/8323/4488"
nal=57%","1910/8412/1909/1906/998/2185/6495"
l=54%","7040/5468/7076/2149/7099/3162/240/3091/2896/3552"
43/5360/81537/51228/154664/9218/65977"
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043/9025/221927/26155/346007/2063/1258/2792/170506/6010/4836/1020/1387/207/3297/22954
32%","2683/53947/10678/93010/8985/8705/23127/28/10331/8707/84002/10317"
%","81492/54839/4926/374407/54777/644890/9148/79989/7756/339829/146754/79582/85478/23
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'1775/1763/4521/25804/7186/1660/5393/10949/2021/5150/22916/80153/328/5433/10535/23404,

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/169714/9112/51530/7329/8480/1434/22929/7112/57122/1063/23468/4521/84823/5902/142/400

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