

## **Read**

HETEROM009E08.B1  
HETEROM009D02.B1  
HETEROM010G03.B1  
HETEROM010D12.B1  
HETEROM010B10.B1  
HETEROM010B10.G1  
HETEROM009C05.B1  
HETEROM009G02.B1  
HETEROM010G03.G1  
HETEROM010B03.B1  
HETEROM010D12.G1  
HETEROM009H04.G1  
HETEROM010E09.B1  
HETEROM010E09.G1  
HETEROM010H10.B1  
HETEROM009F02.G1  
HETEROM009F02.B1  
HETEROM009F11.B1  
HETEROM009D05.B1  
HETEROM009B09.B1  
HETEROM010H10.G1  
HETEROM009H08.B1  
HETEROM009F04.B1  
HETEROM009A12.B1  
HETEROM010H09.G1  
HETEROM010C09.B1  
HETEROM010A03.G1  
HETEROM010H09.B1  
HETEROM010H06.B1  
HETEROM010A03.B1  
HETEROM009F03.B1  
HETEROM009C11.B1  
HETEROM010H02.G1  
HETEROM010F12.G1  
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HETEROM009H08.G1  
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HETEROM009A12.G1  
HETEROM010H07.B1

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HETEROM010C06.B1  
HETEROM010B01.B1  
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HETEROM009D12.B1  
HETEROM010H01.B1  
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HETEROM009C02.G1  
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HETEROM010H07.G1  
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HETEROM009A09.B1  
HETEROM010C02.B1  
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HETEROM010C02.G1  
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HETEROM009B02.G1  
HETEROM009B02.B1  
HETEROM010G01.G1  
HETEROM010F10.B1  
HETEROM010E05.G1  
HETEROM010D08.B1  
HETEROM010D07.G1  
HETEROM010B04.G1  
HETEROM009H12.B1  
HETEROM009E09.B1  
HETEROM009D03.G1  
HETEROM009C08.B1  
HETEROM009C06.B1  
HETEROM010F09.G1  
HETEROM010A05.G1  
HETEROM009G09.B1  
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HETEROM009C08.G1  
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HETEROM010E04.G1  
HETEROM010D11.G1

HETEROM009H02.B1  
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HETEROM009A03.B1  
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HETEROM009G01.B1  
HETEROM009D02.G1  
HETEROM010B08.B1  
HETEROM009E01.B1  
HETEROM009A02.B1  
HETEROM009A01.G1

HETEROM009A01.B1  
HETEROM010D05.B1  
HETEROM009D12.G1  
HETEROM009D11.B1  
HETEROM009A02.G1  
HETEROM010G10.G1  
HETEROM010C07.B1  
HETEROM009G02.G1  
HETEROM009H03.B1  
HETEROM009E05.B1  
HETEROM009E05.G1  
HETEROM010B05.G1  
HETEROM010G09.B1  
HETEROM009D10.B1

## Top match

gi|66809617|ref|XP\_638531.1| hypothetical protein DDB0218610 [Di...  
gi|7270417|emb|CAB80184.1| Putative S-phase-specific ribosomal p...  
gi|68351589|gb|EAN32350.1| 40S ribosomal protein S15a, putative ...  
gi|67587050|ref|XP\_665229.1| senescence-associated protein [Cryp...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|2995405|emb|CAA73042.1| polyprotein [Ananas comosus] >gi|7489...  
gi|31431781|gb|AAP53507.1| putative polyprotein [Oryza sativa (j...  
gi|68351589|gb|EAN32350.1| 40S ribosomal protein S15a, putative ...  
gi|7297878|gb|AAF53124.1| CG14941-PA [Drosophila melanogaster] >...  
gi|13359451|dbj|BAB33421.1| putative senescence-associated prote...  
gi|20799715|gb|AAM28620.1| adenosine monophosphate binding prote...  
gi|53804933|ref|YP\_113254.1| hypothetical protein MCA0751 [Methy...  
gi|53804933|ref|YP\_113254.1| hypothetical protein MCA0751 [Methy...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|23014786|ref|ZP\_00054586.1| COG1024: Enoyl-CoA hydratase/carn...  
gi|23014786|ref|ZP\_00054586.1| COG1024: Enoyl-CoA hydratase/carn...  
gi|7489212|pir||T02995 unspecific monooxygenase (EC 1.14.14.1) -...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|29881613|gb|AAH51156.1| Unknown (protein for MGC:55807) [Dani...  
gi|2133292|pir||S60179 pol polyprotein homolog - fungus (Fusariu...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|67542872|ref|ZP\_00420807.1| hypothetical protein Bcep1808DRAF...  
gi|441217|dbj|BAA03709.1| S-adenosyl-L-homocystein hydrolase [Ni...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|67542872|ref|ZP\_00420807.1| hypothetical protein Bcep1808DRAF...  
gi|38108640|gb|EAA54628.1| hypothetical protein (T18348 probable...  
gi|67542872|ref|ZP\_00420807.1| hypothetical protein Bcep1808DRAF...  
gi|67542872|ref|ZP\_00420807.1| hypothetical protein Bcep1808DRAF...  
gi|15593252|gb|AAL02222.1| cysteine protease CP14 precursor [Fra...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68386573|ref|XP\_687778.1| PREDICTED: similar to Zinc finger p...  
gi|68386573|ref|XP\_687778.1| PREDICTED: similar to Zinc finger p...  
gi|6321025|ref|NP\_011104.1| 14-3-3 protein, major isoform; binds...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...

gi|67542872|ref|ZP\_00420807.1| hypothetical protein Bcep1808DRAF...  
gi|38108640|gb|EAA54628.1| hypothetical protein (T18348 probable...  
gi|38108640|gb|EAA54628.1| hypothetical protein (T18348 probable...  
gi|32420589|ref|XP\_330738.1| hypothetical protein [Neurospora cr...  
gi|40741157|gb|EAA60347.1| RS27\_XENLA 40S ribosomal protein S27 ...  
gi|7489212|pir||T02995 unspecific monooxygenase (EC 1.14.14.1) -...  
gi|46431455|gb|EAK91016.1| hypothetical protein CaO19.6835 [Cand...  
gi|32420589|ref|XP\_330738.1| hypothetical protein [Neurospora cr...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|50725962|dbj|BAD33489.1| zinc finger (C3HC4-type RING finger)...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|2133292|pir||S60179 pol polyprotein homolog - fungus (Fusariu...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|71005880|ref|XP\_757606.1| hypothetical protein UM01459\_1 [Ust...  
gi|5917803|gb|AAD56048.1| S-adenosyl-L-homocysteinase [Lupinus l...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|12652527|gb|AAH00001.1| MGC3234 protein [Homo sapiens]  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|12652527|gb|AAH00001.1| MGC3234 protein [Homo sapiens]  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...

gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|68558534|ref|ZP\_00597872.1| hypothetical protein RmetDRAFT\_17...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|67867504|gb|AAH98085.1| Unknown (protein for MGC:107782) [Xen...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|18767126|gb|AAL79278.1| unknown [Saccharomyces cerevisiae]  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642984|emb|CAH00946.1| unnamed protein product [Kluyveromyc...  
gi|27361021|gb|AAO09931.1| Unknown [Vibrio vulnificus CMCP6] >gi...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642984|emb|CAH00946.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|23479823|gb|EAA16552.1| hypothetical protein [Plasmodium yoel...  
gi|15217223|gb|AAK92567.1| Putative polycomb protein [Oryza sativa]  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|37955146|gb|AAP75564.1| malate synthase [Chlamydomonas reinha...  
gi|37955146|gb|AAP75564.1| malate synthase [Chlamydomonas reinha...  
gi|37955146|gb|AAP75564.1| malate synthase [Chlamydomonas reinha...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...

gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|15217223|gb|AAK92567.1| Putative polycomb protein [Oryza sativa]  
gi|37955146|gb|AAP75564.1| malate synthase [Chlamydomonas reinha...  
gi|34419534|ref|NP\_899547.1| hypothetical protein KVP40.0300 [Ba...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642984|emb|CAH00946.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642984|emb|CAH00946.1| unnamed protein product [Kluyveromyc...  
gi|49642984|emb|CAH00946.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|62718561|ref|XP\_580193.1| PREDICTED: hypothetical protein XP\_...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|38110542|gb|EAA56245.1| hypothetical protein MG01897.4 [Magna...  
gi|49642984|emb|CAH00946.1| unnamed protein product [Kluyveromyc...  
gi|9759046|dbj|BAB09568.1| unnamed protein product [Arabidopsis ...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642984|emb|CAH00946.1| unnamed protein product [Kluyveromyc...  
gi|66857206|ref|ZP\_00401261.1| hypothetical protein AdehDRAFT\_10...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|45504831|gb|AAS66950.1| polyprotein [Melampsora lini]  
gi|49642984|emb|CAH00946.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|34894214|ref|NP\_908432.1| P0439B06.28 [Oryza sativa (japonica...  
gi|9759046|dbj|BAB09568.1| unnamed protein product [Arabidopsis ...

gi|62718743|ref|XP\_578945.1| PREDICTED: similar to Ac1147 [Rattu...  
gi|9759046|dbj|BAB09568.1| unnamed protein product [Arabidopsis ...  
gi|49642978|emb|CAH00940.1| unnamed protein product [Kluyveromyc...  
gi|66798217|ref|ZP\_00396973.1| hypothetical protein DgeoDRAFT\_06...  
gi|39588015|emb|CAE57246.1| Hypothetical protein CBG00124 [Caeno...  
gi|9759046|dbj|BAB09568.1| unnamed protein product [Arabidopsis ...  
gi|3874243|emb|CAA94113.1| Hypothetical protein C11G6.3 [Caenorh...  
gi|66802564|ref|XP\_635154.1| hypothetical protein DDB0230207 [Di...  
gi|3874243|emb|CAA94113.1| Hypothetical protein C11G6.3 [Caenorh...  
gi|66802564|ref|XP\_635154.1| hypothetical protein DDB0230207 [Di...  
gi|3874243|emb|CAA94113.1| Hypothetical protein C11G6.3 [Caenorh...  
gi|67989224|gb|EAM76999.1| Death-on-curing protein [Kineococcus ...  
gi|23270399|ref|NP\_690845.1| Mitochondrial protein of unknown fu...  
gi|66798217|ref|ZP\_00396973.1| hypothetical protein DgeoDRAFT\_06...  
gi|51535965|dbj|BAD38046.1| hypothetical protein [Oryza sativa (...  
gi|11357142|pir||T44430 protein PV100 [imported] - winter squash...  
gi|50751282|ref|XP\_430156.1| PREDICTED: hypothetical protein XP\_...  
gi|9759046|dbj|BAB09568.1| unnamed protein product [Arabidopsis ...  
gi|49648048|emb|CAG82501.1| unnamed protein product [Yarrowia li...  
gi|61658795|gb|AAX49605.1| RPGR [Gorilla gorilla]  
gi|11357142|pir||T44430 protein PV100 [imported] - winter squash...  
gi|60602612|gb|AAX27763.1| unknown [Schistosoma japonicum]  
gi|67521327|gb|EAM25267.1| Transthyretin [Chromohalobacter salex...  
gi|1066466|gb|AAC48486.1| zonadhesin >gi|7522633|pir||T34022 zon...  
gi|3191990|emb|CAA19389.1| putative signal peptidase I [Streptom...  
gi|3191990|emb|CAA19389.1| putative signal peptidase I [Streptom...  
gi|26335533|dbj|BAC31467.1| unnamed protein product [Mus musculus]  
gi|68192204|gb|EAN06858.1| Transthyretin [Mesorhizobium sp. BNC1...  
gi|62666723|ref|XP\_346291.2| PREDICTED: similar to 60S ribosomal...  
gi|26352113|dbj|BAC39693.1| unnamed protein product [Mus musculus]  
gi|34912742|ref|NP\_917718.1| B1040D09.19 [Oryza sativa (japonica...  
gi|62664649|ref|XP\_225708.3| PREDICTED: similar to LRRG00122 [Ra...  
gi|68004178|ref|ZP\_00536677.1| Polynucleotide adenylyltransferas...  
gi|9635644|ref|NP\_061549.1| Orf53 [Pseudomonas phage D3] >gi|889...  
gi|9294219|dbj|BAB02121.1| unnamed protein product [Arabidopsis ...  
gi|71021915|ref|XP\_761188.1| hypothetical protein UM05041\_1 [Ust...  
gi|23488487|gb|EAA21331.1| hypothetical protein [Plasmodium yoel...  
gi|30526291|gb|AAP32203.1| latency associated nuclear antigen [S...  
gi|34101408|gb|AAQ57777.1| conserved hypothetical protein [Chrom...  
gi|11282893|pir||F81737 hypothetical protein TC0129 [imported] -...

gi|68560883|ref|ZP\_00600191.1| hypothetical protein Rxy|DRAFT\_21...  
gi|18378780|gb|AAL68699.1| unknown [Bradyrhizobium sp. ORS278]  
gi|50940129|ref|XP\_479592.1| unknown protein [Oryza sativa (japo...  
gi|46580215|ref|YP\_011023.1| GGDEF domain protein [Desulfovibrio...  
gi|68125435|emb|CAJ03295.1| hypothetical protein, conserved [Lei...  
gi|34912742|ref|NP\_917718.1| B1040D09.19 [Oryza sativa (japonica...  
gi|50253297|dbj|BAD29566.1| hypothetical protein [Oryza sativa (...  
gi|61191888|gb|AAX39410.1| lipase [Bombyx mandarina]  
gi|58391017|ref|XP\_318209.2| ENSANGP00000001363 [Anopheles gambi...  
gi|46437082|gb|EAK96434.1| hypothetical protein CaO19.10591 [Can...  
gi|46437082|gb|EAK96434.1| hypothetical protein CaO19.10591 [Can...  
gi|39579352|emb|CAE56520.1| Hypothetical protein CBG24243 [Caeno...  
gi| \*\*\*\*\* No hits found \*\*\*\*\*  
gi| \*\*\*\*\* No hits found \*\*\*\*\*

<b>score</b>	<b>e-value</b>
246	2.00E-64
199	6.00E-50
191	8.00E-48
174	1.00E-42
172	2.00E-42
158	4.00E-38
147	2.00E-34
145	4.00E-34
143	5.00E-34
136	5.00E-31
135	4.00E-43
135	1.00E-30
131	8.00E-30
127	8.00E-29
114	1.00E-24
111	8.00E-24
111	8.00E-24
110	1.00E-23
110	1.00E-23
110	1.00E-30
109	2.00E-23
109	4.00E-23
109	4.00E-23
108	4.00E-23
107	4.00E-25
107	2.00E-22
107	2.00E-22
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101	1.00E-20

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44 0.002  
44 0.001  
43 0.004  
43 0.004  
43 0.003  
43 0.004  
42 0.005  
41 0.01  
40 0.018  
40 0.023  
39 0.041  
39 0.041  
39 0.051  
38 0.086  
38 0.087  
38 0.11  
37 0.19  
37 0.25  
37 0.19  
37 0.19  
37 0.26  
37 0.19  
36 0.052  
36 0.43  
36 0.42  
36 0.37

36	0.38
35	0.95
35	0.74
35	0.57
35	0.56
34	1.3
32	0.3
26	1.00E-12
24	0.032
24	7.00E-06
23	0.003
22	4.00E-06