

Supplementary Table S2. Cardiac gene expression. A GeneChip Mouse Genome 430 2.0 Array (Affymetrix, Santa Clara, CA, United States) was used to analyze the basal mRNA expression in hearts from PP2A-TG (n=3) and WT (n=3) mice. Here, only genes are shown that were expressed in both PP2A-TG and WT and with a 1.5-fold or higher down-regulation in PP2A-TG. For analysis, the TM4 application MeV was used (Saeed et al., 2003) and analyzed by a t-Test (Welch approximation; alpha (overall threshold p-value): 0.05; P-values based on t-distribution). Data are sorted by p value.

Gene descriptions	WT mean	WT SD	PP2A-TG mean	PP2A-TG SD	Absolute t value	Degrees of freedom	Adj p value	False discovery rate	PP2A-TG / WT	WT / PP2A-TG
gb.BC019117.1/DB_XREF=gi:5172286/FEA=FlmRNA/ CNT=92/ TID=Mm.27932/1/ TIER=FL+Stack/ STK=39/ UG=Mm.27932/ UG_GENE=Ansf2-pending/DEF=Mus musculus, clone MGC-29432 IMAGE:3709664, mRNA, complete cds. /PROD=Unknown (protein for MGC-29432) /FL-gb:BC019117.1 gb.NM.031405.1	838.2334	46.69404	477.6	51.20811	9.013391	3.0	0.0028832927	1.0524018	0.57	1.76
gb.AK004894.1/DB_XREF=gi:12834243/FEA=mrRNA/ CNT=70/ TID=Mm.1183.1/ TIER=Stack/ STK=31/ UG=Mm.1183/ LL=74153/ UG_GENE=130004C08Rik/ UG_TITLE=RIKEN cDNA 130004C08 gene/DEF=Mus musculus adult male liver cDNA, RIKEN full-length enriched library, clone:130004C08-UBE1L (FRAGMENT), full insert sequence.	610.3333	7.245238	400.9	19.546862	17.401026	2.0	0.003286286	0.5997472	0.66	1.52
gb.BC025144.1/DB_XREF=gi:19264066/FEA=FlmRNA/ CNT=27/ TID=Mm.219658.1/ TIER=FL+Stack/ STK=17/ UG=Mm.219658/DEF=Mus musculus, Similar to GTP binding protein 5 (putative), clone MGC:36295 IMAGE:4483303, mRNA, complete cds. /PROD=Similar to GTP binding protein 5 (putative) /FL-gb:BC025144.1	627.4666	47.652325	374.16666	36.395786	7.316819	3.0	0.0052728374	0.64152855	0.6	1.68
gb.U11548.1/DB_XREF=gi:511226/FEA=FlmRNA/ CNT=8/ TID=Mm.28587.1/ TIER=FL/ STK=1/ UG=Mm.28587/ LL=26407/ UG_GENE=Map3k4/DEF=Mus musculus TfIIi-associated transactivator factor p17 mRNA, complete cds. /PROD=TfIIi-associated transactivator factor p17 /FL-gb:U85608.1 gb.U11548.1 gb.NM.011948.1	500.83334	49.23366	250.59999	36.55603	7.0680494	3.0	0.0058227624	0.53132707	0.5	2
gb.NM.007981.1/DB_XREF=gi:6897938/GEN=Ec52/FEA=FlmRNA/ CNT=147/ TID=Mm.28982.1/ TIER=FL+Stack/ STK=45/ UG=Mm.28982/ LL=14081/DEF=Mus musculus fatty acid Coenzyme A ligase, long chain 2 (FACL2), mRNA. /PROD=fatty acid Coenzyme A ligase, long chain 2 /FL-gb:NM.007981.1 gb.U15977.1	3965.5664	415.30035	2066.2332	226.85002	6.9518514	3.0	0.006105606	0.44570923	0.52	1.92
gb.BG073508/DB_XREF=gi:12556007/DB_XREF=H3123D10.3/ CLONE=H3123D10/FEA=mrRNA/ CNT=46/ TID=Mm.21795.1/ TIER=Stack/ STK=53/ UG=Mm.21795/ UG_TITLE=Mus musculus, Similar to BCL2adenovirus E1B 19kD interacting protein 1, clone MGC-41600 IMAGE:1264175, mRNA, complete cds	559.89996	13.956017	347.1	27.04736	12.110144	2.0	0.0067497385	0.4106091	0.62	1.61
gb.AK003586.1/DB_XREF=gi:12834343/FEA=mrRNA/ CNT=119/ TID=Mm.20488.1/ TIER=Stack/ STK=50/ UG=Mm.20488/ LL=76301/ UG_GENE=1110008K06Rik/ UG_TITLE=RIKEN cDNA 1110008K06 gene/DEF=Mus musculus 18 days embryo whole body cDNA, RIKEN full-length enriched library, clone:1110008K06 homolog to FLN2 (FLN2 Gene PRODUCT), full insert sequence.	1412.2666	172.08234	591.0333	137.58932	6.455995	3.0	0.007538527	0.39308035	0.42	2.39
gb.BC006988.1/DB_XREF=gi:1387037/FEA=FlmRNA/ CNT=135/ TID=Mm.20336.1/ TIER=FL+Stack/ STK=80/ UG=Mm.20336/ LL=12908/ UG_GENE=Cral/DEF=Mus musculus, Similar to carnitine acetyltransferase, clone MGC:6046 IMAGE:3499169, mRNA, complete cds. /PROD=Similar to carnitine acetyltransferase /FL-gb:BC006988.1 gb.NM.007760.1	4650.7334	358.1267	2468.2334	466.95862	6.4237013	3.0	0.007646559	0.34887424	0.53	1.88
gb.BC014685.1/DB_XREF=gi:15928403/FEA=FlmRNA/ CNT=42/ TID=Mm.170023.1/ TIER=FL+Stack/ STK=18/ UG=Mm.170023/DEF=Mus musculus, clone MGC:25977 IMAGE:4241375, mRNA, complete cds. /PROD=Unknown (protein for MGC:25977) /FL-gb:BC014685.1	660.4333	51.97042	433.63333	39.917454	5.9945416	3.0	0.009296497	0.37702462	0.66	1.52
gb.NM.007533.1/DB_XREF=gi:6671623/GEN=Bckha/FEA=FlmRNA/ CNT=211/ TID=Mm.25848.1/ TIER=FL+Stack/ STK=76/ UG=Mm.25848/ LL=12039/DEF=Mus musculus branched chain ketoadic dehydrogenase E1, alpha polypeptide (Bckha), mRNA. /PROD=branched chain ketoadic dehydrogenase E1, alpha polypeptide /FL-gb:BC003787.1 gb.NM.007533.1 gb.U47335.1	2700.0334	286.70358	1546.7666	192.90247	5.780555	3.0	0.010295443	0.37578368	0.57	1.75
gb.BC048804.1/DB_XREF=gi:1235992/FEA=FlmRNA/ CNT=277/ TID=Mm.100540.1/ TIER=FL+Stack/ STK=17/ UG=Mm.100540/ LL=10450/ UG_GENE=Napa/DEF=Mus musculus, RIKEN cDNA 1500039N14 gene, clone MGC:6399 IMAGE:3584869, mRNA, complete cds. /PROD=RIKEN cDNA 1500039N14 gene /FL-gb:NM.025898.1 gb.BC048804.1	3788.9336	236.00606	2382.4668	356.9891	5.692439	3.0	0.010747662	0.35662696	0.63	1.59
gb.AB056479.1/DB_XREF=gi:15341205/FEA=FlmRNA/ CNT=427/ TID=Mm.34822.1/ TIER=FL+Stack/ STK=32/ UG=Mm.34822/ LL=12974/ UG_GENE=Cg/DEF=Mus musculus mRNA for citrate synthase, complete cds. /PROD=citrate synthase /FL-gb:BC013554.1 gb.NM.02644.1 gb.AB056479.1	3668.7334	270.4928	2305.8333	327.08664	5.5616684	3.0	0.011467959	0.34881708	0.63	1.59
gb.BC018375.1/DB_XREF=gi:17380875/FEA=FlmRNA/ CNT=59/ TID=Mm.4451.1/ TIER=FL+Stack/ STK=30/ UG=Mm.4451/ LL=15205/ UG_GENE=Hes1/DEF=Mus musculus, hairy and enhancer of split 1, (Drosophila) /FL-gb:NM.008235.1 gb.BC018375.1	557.3333	50.158768	345.19998	46.835567	5.354062	3.0	0.012746805	0.35789105	0.62	1.61
gb.NM.008761.1/DB_XREF=gi:6879448/DEF=Phps/FEA=FlmRNA/ CNT=174/ TID=Mm.1458.1/ TIER=FL+Stack/ STK=115/ UG=Mm.1458/ LL=10602/DEF=Mus musculus putative phosphatase (Psp), mRNA. /PROD=putative phosphatase /FL-gb:AF483523.1 gb.AF483522.1 gb.NM.008916.1 gb.U96724.1	1440.2001	142.18462	787.8	161.60974	5.2495723	3.0	0.013461299	0.3509553	0.55	1.83
gb.AK004710.1/DB_XREF=gi:12630083/GEN=Scs32/FEA=mrRNA/ CNT=23/ TID=Mm.21587.2/ TIER=Stack/ STK=16/ UG=Mm.21587/ LL=65962/ UG_TITLE=solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulator 2 /DEF=Mus musculus adult male lung cDNA, RIKEN full-length enriched library, clone:1200011K07/solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulator 2, full insert sequence.	764.56665	76.69448	400.46667	93.46831	5.215931	3.0	0.013702431	0.33342582	0.52	1.91
gb.AK012604.1/DB_XREF=gi:12849126/FEA=mrRNA/ CNT=49/ TID=Mm.172946.1/ TIER=Stack/ STK=15/ UG=Mm.172946/ LL=72579/ UG_GENE=2700049M22Rik/ UG_TITLE=RIKEN cDNA 2700049M22 gene/DEF=Mus musculus 11 days embryo whole body cDNA, RIKEN full-length enriched library, clone:2700049M22-hypothetical protein, full insert sequence.	717.3	14.961293	416.63333	60.370216	8.372979	2.0	0.013965844	0.31859583	0.58	1.72
gb.AK049477.1/DB_XREF=gi:12630083/FEA=FlmRNA/ CNT=907/ TID=Mm.22574.1/ TIER=FL+Stack/ STK=90/ UG=Mm.22574/ LL=12978/ UG_GENE=Colony stimulating factor 1 receptor /DEF=Mus musculus adult male liver cDNA, RIKEN full-length enriched library, clone:1300008N20.colony stimulating factor 1 receptor, full insert sequence. /FL-gb:NM.007779.1	1033.4667	80.77104	583.6	131.73853	5.0423894	3.0	0.015040075	0.32291928	0.56	1.77
gb.NM.007905.1/DB_XREF=gi:6681272/GEN=Erf12/FEA=FlmRNA/ CNT=64/ TID=Mm.2645.1/ TIER=FL+Stack/ STK=31/ UG=Mm.2645/ LL=13628/DEF=Mus musculus Eukaryotic translation elongation factor 1 alpha 2 (Erf12), mRNA. /PROD=eukaryotic translation elongation factor 1 alpha 2 /FL-gb:U36479.1 gb.NM.007905.1 gb.BC018215.1	17134.666	2077.833	9926.967	1596.6765	4.7641034	3.0	0.017563034	0.3561393	0.58	1.73
gb.NM.029632.1/DB_XREF=gi:18390326/GEN=Ppp1r11/FEA=FlmRNA/ CNT=146/ TID=Mm.46176.1/ TIER=FL+Stack/ STK=58/ UG=Mm.46176/ LL=76497/DEF=Mus musculus protein phosphatase 1, regulatory (inhibitor) subunit 11 (Ppp1r11), mRNA. /PROD=protein phosphatase 1, regulatory (inhibitor) subunit 11 /FL-gb:NM.029632.1	3319.6667	170.52585	1640.5333	382.8957	6.9386435	2.0	0.020145172	0.38699934	0.49	2.02
gb.BC008103.1/DB_XREF=gi:14788073/FEA=FlmRNA/ CNT=64/ TID=Mm.196749.1/ TIER=FL+Stack/ STK=24/ UG=Mm.196749/ LL=11840/DEF=Mus musculus, Similar to hypothetical protein FL10504, clone MGC:11722 IMAGE:5968515, mRNA, complete cds. /PROD=Similar to hypothetical protein FL10504 /FL-gb:BC008103.1	444.4	49.412437	264.86667	48.947556	4.470925	3.0	0.020850305	0.38051805	0.6	1.68
gb.AK020444.1/DB_XREF=gi:12861029/FEA=mrRNA/ CNT=70/ TID=Mm.36699.1/ TIER=Stack/ STK=42/ UG=Mm.36699/ LL=77254/ UG_GENE=9430028K06Rik/ UG_TITLE=RIKEN cDNA 9430028K10 gene/DEF=Mus musculus 12 days embryo whole body cDNA, RIKEN full-length enriched library, clone:9430028K10 homolog to SATMP, full insert sequence.	703.2334	78.743706	433.4	69.91444	4.438314	3.0	0.021263769	0.36958456	0.62	1.62
gb.NM.053014.1/DB_XREF=gi:16519540/GEN=Appa3/FEA=FlmRNA/ CNT=6/ TID=Mm.141230.1/ TIER=FL+Stack/ STK=1/ UG=Mm.141230/ LL=28169/DEF=Mus musculus 1-acetylglucosyl-3-phosphate O-acetyltransferase 3 (Appa3), mRNA. /PROD=1-acetylglucosyl-3-phosphate O-acetyltransferase 3 /FL-gb:NM.053014.1	1276.3666	167.27182	751.9666	118.466934	4.431236	3.0	0.021354912	0.3542974	0.59	1.7
gb.AK002437.1/DB_XREF=gi:1284241/FEA=FlmRNA/ CNT=204/ TID=Mm.180189.1/ TIER=FL+Stack/ STK=57/ UG=Mm.180189/ LL=67800/ UG_GENE=Dga2/ UG_TITLE=diacylglycerol O-acetyltransferase 2 /DEF=Mus musculus adult male kidney cDNA, RIKEN full-length enriched library, clone:010010B06.homolog to WUGSCH.DJ047478.5 PROTEIN (FRAGMENT), full insert sequence. /FL-gb:AF384160.1 gb.NM.026384.1	2753.4001	267.295	1639.7334	105.6523	6.7112303	2.0	0.021489117	0.34102297	0.6	1.68
gb.NM.007476.1/DB_XREF=gi:6680715/GEN=Atf1/FEA=FlmRNA/ CNT=64/ TID=Mm.6836.1/ TIER=FL+Stack/ STK=2/ UG=Mm.6836/ LL=11840/DEF=Mus musculus ADF-ribosylation factor 1 (Atf1), mRNA. /PROD=ADP-ribosylation factor 1 /FL-gb:DB7698.1 gb.NM.007476.1	1804.6666	201.61327	1130.6666	170.99333	4.415943	3.0	0.021553561	0.32779375	0.63	1.6
gb.BC027445.1/DB_XREF=gi:20071662/FEA=FlmRNA/ CNT=79/ TID=Mm.153891.1/ TIER=FL+Stack/ STK=28/ UG=Mm.153891/ LL=19245/ UG_GENE=Ptp4a3/DEF=Mus musculus, protein tyrosine phosphatase 4a3, clone MGC:36146 IMAGE:4482106, mRNA, complete cds. /PROD=protein tyrosine phosphatase 4a3 /FL-gb:NM.008975.1 gb.AF356645.1 gb.BC027445.1	3634.5999	385.0914	2027.2666	170.24298	6.6120944	3.0	0.022116944	0.3229074	0.56	1.79
gb.NM.028152.1/DB_XREF=gi:15100755/GEN=Mms19/FEA=FlmRNA/ CNT=164/ TID=Mm.28412.1/ TIER=FL+Stack/ STK=32/ UG=Mm.28412/ LL=72199/DEF=Mus musculus MMS19 (MEI18 S. cerevisiae)-like (Mms19), mRNA. /PROD=MMS19 (MEI18 S. cerevisiae)-like /FL-gb:AF319949.1 gb.NM.028152.1	386.7667	35.257385	238.33333	17.895628	6.5022836	3.0	0.022844655	0.3207038	0.62	1.62
gb.B147262/DB_XREF=gi:15173875/DB_XREF=602988681F1/ CLONE=IMAGE:5144372/FEA=mrRNA/ CNT=159/ TID=Mm.35345.1/ TIER=Stack/ STK=43/ UG=Mm.35345/ LL=11840/DEF=Mus musculus, clone IMAGE:5237602, mRNA, partial cds	512.1667	84.50942	246.5	68.90783	4.2199216	3.0	0.024325186	0.32884046	0.48	2.08
gb.BC017689.1/DB_XREF=gi:1284241/FEA=FlmRNA/ CNT=148/ TID=Mm.19258.1/ TIER=FL+Stack/ STK=1/ UG=Mm.19258/ LL=66951/ UG_GENE=2310028D20Rik/DEF=Mus musculus, Similar to tyrothroph embryonic factor, clone MGC:19233 IMAGE:4242534, mRNA, complete cds. /PROD=Similar to tyrothroph embryonic factor /FL-gb:BC017689.1	2227.3333	188.9985	1486.5	238.61961	4.2153683	3.0	0.024394894	0.31800488	0.67	1.5
gb.AK033605.1/DB_XREF=gi:12833890/FEA=mrRNA/ CNT=531/ TID=Mm.46367.1/ TIER=Stack/ STK=12/ UG=Mm.46367/ LL=68453/ UG_GENE=1110002J10Rik/ UG_TITLE=RIKEN cDNA 1110002J10 gene/DEF=Mus musculus 18 days embryo whole body cDNA, RIKEN full-length enriched library, clone:1110002J10.unclassified, full insert sequence.	5818.1997	927.7161	3272.6667	519.24005	4.1471405	3.0	0.025470959	0.32058278	0.56	1.78
gb.AF136571.1/DB_XREF=gi:1507284/FEA=FlmRNA/ CNT=14/ TID=Mm.10211.2/ TIER=FL+Stack/ STK=1/ UG=Mm.10211/ LL=12499/ UG_GENE=Entp5/DEF=Mus musculus PcpH proto-oncogene protein mRNA, complete cds. /PROD=PcpH proto-oncogene protein /FL-gb:AF136571.1	602.6667	72.00232	401.03336	45.249798	4.106742	3.0	0.026137074	0.31800106	0.67	1.5
gb.BC027445.1/DB_XREF=gi:12847833/FEA=mrRNA/ CNT=96/ TID=Mm.21693.1/ TIER=Stack/ STK=4/ UG=Mm.21693/ LL=76201/ UG_GENE=2610028P08Rik/ UG_TITLE=RIKEN cDNA 2610028P08 gene/DEF=Mus musculus 10 days embryo whole body cDNA, RIKEN full-length enriched library, clone:2610028P08.homolog to CDNA FLJ20128 FIS. CLONE COL06181, full insert sequence.	428.7333	51.981567	236.03333	62.753677	4.095951	3.0	0.02631879	0.30988252	0.55	1.82
gb.NM.011776.1/DB_XREF=gi:7242190/GEN=Pasp/FEA=FlmRNA/ CNT=332/ TID=Mm.3383.1/ TIER=FL+Stack/ STK=1/ UG=Mm.3383/ LL=19156/DEF=Mus musculus prosaposin (Pasp), mRNA. /PROD=prosaposin /FL-gb:NM.011776.1	5425.933	829.29877	2215.7332	1085.7566	4.06974	3.0	0.026767017	0.3053113	0.41	2.45
gb.AK00460.1/DB_XREF=gi:12844269/FEA=mrRNA/ CNT=27/ TID=Mm.218627.2/ TIER=Stack/ STK=20/ UG=Mm.218627/ LL=66053/ UG_GENE=Ppil2/UG_TITLE=peptidylglyoxyl isomerase (cyclophilin-like) 2 /DEF=Mus musculus adult male tongue cDNA, RIKEN full-length enriched library, clone:2310022D1.homolog to CYCLOPHILIN-LIKE PROTEIN CYP-60, full insert sequence.	726.69995	100.58585	387.8667	104.56621	4.044865	3.0	0.027201526	0.30086538	0.53	1.87
gb.BC024138.1/DB_XREF=gi:12838268/FEA=FlmRNA/ CNT=226/ TID=Mm.22626.1/ TIER=FL+Stack/ STK=1/ UG=Mm.22626/ LL=66951/ UG_GENE=2310078J06Rik/DEF=Mus musculus, Similar to tyrothroph embryonic factor, clone MGC:37973 IMAGE:5136320, mRNA, complete cds. /PROD=RIKEN cDNA 2310078J06 gene /FL-gb:BC024138.1	1782.4	251.44804	1036.3667	201.14285	4.012932	3.0	0.027772743	0.29814857	0.58	1.72
gb.BC018323.1/DB_XREF=gi:17380875/FEA=FlmRNA/ CNT=100/ TID=Mm.3459.1/ TIER=FL+Stack/ STK=28/ UG=Mm.3459/ LL=13170/ UG_GENE=Ddp/DEF=Mus musculus, D site albumin promoter binding protein, clone MGC:18803 IMAGE:4195116, mRNA, complete cds. /PROD=D site albumin promoter binding protein /FL-gb:NM.016974.1 gb.BC018323.1	2181.5	125.08818	1068.4	314.64484	5.693912	2.0	0.029487085	0.30750817	0.49	2.04
gb.NM.008398.1/DB_XREF=gi:6680479/GEN=Ilgav/FEA=FlmRNA/ CNT=47/ TID=Mm.179747/ TIER=FL+Stack/ STK=24/ UG=Mm.179747/ LL=16404/DEF=Mus musculus integrin alpha 7 (Ilgav), mRNA. /PROD=integrin alpha 7 /FL-gb:NM.008398.1 gb.U123423.1	982.36664	117.43058	634.99994	100.058136	3.8998327	3.0	0.029924314	0.3033993	0.65	1.55
gb.AF206720.1/DB_XREF=gi:6636320/FEA=FlmRNA/ CNT=130/ TID=Mm.38469.1/ TIER=FL+Stack/ STK=83/ UG=Mm.38469/ LL=11785/ UG_GENE=Alpo1/DEF=Mus musculus p165 mRNA, complete cds. /PROD=P65 /FL-gb:AF206720.1	1659.7334	267.31207	867.3	232.41673	3.8747864	3.0	0.030429503	0.30018294	0.52	1.91
gb.B408679/DB_XREF=gi:15169602/DB_XREF=602994304F1/ CLONE=IMAGE:5118956/FEA=mrRNA/ CNT=61/ TID=Mm.29389.1/ TIER=Stack/ STK=30/ UG=Mm.29389/ UG_TITLE=Mus musculus, Similar to KIAA1075 protein, clone IMAGE:5099327, mRNA, partial cds	1488.9	179.23257	810.2	246.92395	3.8527741	3.0	0.030882569	0.29663518	0.54	1.84
gb.NM.010116.1/DB_XREF=gi:7106302/GEN=Edh1/FEA=FlmRNA/ CNT=383/ TID=Mm.30169.1/ TIER=FL+Stack/ STK=145/ UG=Mm.30169/ LL=13660/DEF=Mus musculus E1 domain containing 1 (Edh1), mRNA. /PROD=E1 domain containing 1 /FL-gb:NM.010116.1 gb.AF099186.1 gb.AF173156.1	517.63336	69.581696	323.16666	53.06084	3.8492374	3.0	0.03095617	0.289718	0.62	1.6
gb.NM.009983.1/DB_XREF=gi:6753555/GEN=Ctsd/FEA=FlmRNA/ CNT=522/ TID=Mm.2147.1/ TIER=FL+Stack/ STK=193/ UG=Mm.2147/ LL=13033/DEF=Mus musculus cathepsin D (Ctsd), mRNA. /PROD=cathepsin D /FL-gb:NM.009983.1	5454.1333	777.9049	3241.0332	625.04315	3.8412483	3.0	0.031123256	0.2839997	0.59	1.68
gb.U082071.1/DB_XREF=gi:2680307/GEN=Col2A1/FEA=FlmRNA/ CNT=189/ TID=Mm.18917.1/ TIER=FL+Stack/ STK=1/ UG=Mm.18916/ LL=14719/DEF=Mus musculus aspartate aminotransferase precursor (Col2A1) mRNA, Col2A1 allele, complete cds; nuclear gene for mitochondrial product. /PROD=aspartate aminotransferase precursor /FL-gb:U82470.1 gb.U082071.1 gb.NM.010325.1	4684.267	741.73334	2668.0999	544.53687	3.795118					

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gb BC024394.1 DB_XREF=gi:19354424 FEA=FLmRNA CNT=2007 TID=Mm.34570.2 TIER=FL+Slack STK=106 UG=Mm.34570.1 UG_GENE=Irfp3 DEF=Mus musculus, RNA polymerase 1, 116 kDa subunit, clone MGC 36184 IMAGE:4987975, mRNA, complete cds. /PROD=RNA polymerase 1, 116 kDa subunit FL-gb BC024394.1	1394.3667	266.72238	763.7	316.98685	2.6367848	3.0	0.07787033	0.28141257	0.55	1.83																																																																																																																																																																																												
gb BC013486.1 DB_XREF=gi:15488676 FEA=FLmRNA CNT=1989 TID=Mm.41389.1 TIER=FL+Slack STK=141 UG=Mm.41389.1 LL=66277 UG_GENE=K15 DEF=Mus musculus, Kruppel-like factor 15, clone MGC:19125 IMAGE:4211067, mRNA, complete cds.	164.1.1333	293.1515	993.6667	309.8631	2.6290479	3.0	0.07838921	0.28051043	0.61	1.65																																																																																																																																																																																												
/PROD=Kruppel-like factor 15 FL-gb NM_023184.1 gb AF1317225.1 gb BC013486.1																																																																																																																																																																																																						
gb NM_009777.1 DB_XREF=gi:10671071 FEA=FLmRNA CNT=1487 TID=Mm.2570.1 TIER=FL+Slack STK=58	2006.4667	304.08218	1285.4332	365.01932	2.6287265	3.0	0.078410864	0.27786374	0.64	1.56																																																																																																																																																																																												
UG=Mm.2570.1 LL=12260 DEF=Mus musculus complement component 1, q subcomponent, beta polypeptide (C1qb), mRNA. /PROD=complement component 1, q subcomponent, beta polypeptide (C1qb) FL-gb NM_009777.1																																																																																																																																																																																																						
gb AF041858.1 DB_XREF=gi:3241986 FEA=mRNA CNT=17 TID=Mm.30717.5 TIER=ConsEnd STK=0 UG=Mm.30717.1 LL=20975 UG_GENE=Synp DEF=Mus musculus synaptobrevin 2 isoform delta mRNA, partial cds. /PROD=synaptobrevin 2 isoform delta	608.3667	48.258926	377.56668	109.53727	3.3397484	2.0	0.07915579	0.27780637	0.62	1.61																																																																																																																																																																																												
gb NM_023371.1 DB_XREF=gi:12863822 GEN=PinP1 FEA=FLmRNA CNT=1737 TID=Mm.7906.1 TIER=FL+Slack STK=110 UG=Mm.7906.1 LL=23988 DEF=Mus musculus protein (peptidyl-prolyl cistrans isomerase) NIMA-interacting 1 (Pin1), mRNA. /PROD=protein (peptidyl-prolyl cistrans isomerase)NIMA-interacting 1 FL-gb NM_023371.1	3336.7666	745.0641	1999.5333	480.16516	2.613035	3.0	0.07947676	0.27627635	0.6	1.67																																																																																																																																																																																												
gb NM_008155.1 DB_XREF=gi:6680086 GEN=Sp11 FEA=FLmRNA CNT=26 TID=Mm.588.1 TIER=FL STK=2 UG=Mm.588.1 LL=14751 DEF=Mus musculus glucose phosphate isomerase 1 complex (Sp1), mRNA. /PROD=glucose phosphate isomerase 1 complex FL-gb MI4220.1 gb NM_008155.1	6017.467	1636.3217	3145.5999	981.9437	2.6065686	3.0	0.07992121	0.27520037	0.52	1.91																																																																																																																																																																																												
gb U48800.1 DB_XREF=gi:1234824 GEN=VEGF-B FEA=FLmRNA CNT=77 TID=Mm.15807.2 TIER=FL STK=1 UG=Mm.15807.1 LL=22340 DEF=Mus musculus vascular endothelial growth factor B precursor (VEGF-B) mRNA, complete cds. /PROD=vascular endothelial growth factor B precursor FL-gb NM_043837.1 gb U48800.1	3644.2334	921.23413	1713.7333	417.65958	3.3057415	2.0	0.08060011	0.2749443	0.47	2.13																																																																																																																																																																																												
gb NM_130811.1 DB_XREF=gi:18700010 GEN=MGC6685 FEA=FLmRNA CNT=43 TID=Mm.12267.1 TIER=FL STK=2 UG=Mm.12267.1 LL=177075 DEF=Mus musculus similar to poly(A)-binding protein, cytoplasmic 4 (inducible form) (MGC6685), mRNA. /PROD=similar to poly(A)-binding protein, cytoplasmic (inducible form) FL-gb NM_130811.1 gb BC003283.1	2363.5667	555.67523	1319.0	424.0053	2.5884504	3.0	0.081182934	0.27436823	0.56	1.79																																																																																																																																																																																												
gb NM_130864.1 DB_XREF=gi:18700003 GEN=PCAB2 FEA=FLmRNA CNT=1237 TID=Mm.205266.1 TIER=FL+Slack STK=602 UG=Mm.205266.1 LL=113866 DEF=Mus musculus acetyl-Coenzyme A acyltransferase (peroxisomal 3-oxoacyl-Coenzyme A thioesterase) (PCAB2), mRNA. /PROD=acetyl-Coenzyme A acyltransferase (peroxisomal 3-oxoacyl-Coenzyme A thioesterase) FL-gb NM_130864.1	796.2	190.25824	459.66666	122.50037	2.5759327	3.0	0.08206902	0.27481827	0.58	1.73																																																																																																																																																																																												
gb BC012400.1 gb BC026669.1																																																																																																																																																																																																						
gb AK003239.1 DB_XREF=gi:12833782 FEA=FLmRNA CNT=199 TID=Mm.34244.1 TIER=Slack STK=14 UG=Mm.34244.1 LL=67458 UG_GENE=1200007D18Rk UG_TITLE=RIKEN cDNA 1200007D18 gene (DEF=Mus musculus 18 days embryo whole body cDNA, RIKEN full-length enriched library, clone:1110001K09 homolog to KIAA1181 PROTEIN (FRAGMENT)), full insert sequence. /FL-gb NM_026170.1 gb BC005516.1	510.6	123.16634	300.93335	69.6006	2.5669823	3.0	0.08270989	0.27444646	0.59	1.7																																																																																																																																																																																												
gb NM_053082.1 DB_XREF=gi:16718364 GEN=Tm4d7 FEA=FLmRNA CNT=147 TID=Mm.195408.1 TIER=FL+Slack STK=105 UG=Mm.195498 LL=20524 DEF=Mus musculus transmembrane 4 superfamily member 7 (Tm4d7), mRNA. /PROD=transmembrane 4 superfamily member 7 FL-gb NM_053082.1 gb BC003482.1	1432.9668	371.01614	723.1333	83.90889	3.2321563	2.0	0.083857775	0.27574852	0.5	1.98																																																																																																																																																																																												
gb NM_015816.1 DB_XREF=gi:7454381 FEA=FLmRNA CNT=133 TID=Mm.14585.1 TIER=FL+Slack STK=52 UG=Mm.14585.1 LL=50783 DEF=Mus musculus U6 snRNA-associated SM-like protein 4 (Lsm4-pending), mRNA. /PROD=U6 snRNA-associated SM-like protein 4 FL-gb NM_015816.1 gb BC026747.1	2082.0667	419.71677	1322.8334	300.10544	2.5486553	3.0	0.08404149	0.27388522	0.64	1.57																																																																																																																																																																																												
gb NM_009045.1 DB_XREF=gi:6677708 GEN=Raga FEA=FLmRNA CNT=10 TID=Mm.28170.1 TIER=FL+Slack STK=9 UG=Mm.28170.1 LL=19687 DEF=Mus musculus v-rel reticulendotheliosis viral oncogene homolog A, (avian) (Rela), mRNA. /PROD=avian reticulendotheliosis viral oncogene homolog A FL-gb NM_009045.1 gb BC01909.1	1960.0332	383.5544	1213.4333	334.97433	2.5393877	3.0	0.08472489	0.2736689	0.62	1.62																																																																																																																																																																																												
gb NM_011125.1 DB_XREF=gi:6755111 GEN=Plp FEA=FLmRNA CNT=67 TID=Mm.8105.1 TIER=FL+Slack STK=32 UG=Mm.8105.1 LL=18830 DEF=Mus musculus phospholipid transfer protein (Plp), mRNA. /PROD=phospholipid transfer protein (Plp) FL-gb BC003782.1	2012.7001	384.84665	1234.1333	370.92523	2.5229409	3.0	0.08594584	0.27520546	0.61	1.63																																																																																																																																																																																												
gb NM_008155.1 DB_XREF=gi:6680086 GEN=Sp11 FEA=FLmRNA CNT=26 TID=Mm.588.1 TIER=FL STK=2 UG=Mm.588.1 LL=14751 DEF=Mus musculus glucose phosphate isomerase 1 complex (Sp1), mRNA. /PROD=glucose phosphate isomerase 1 complex FL-gb MI4220.1 gb NM_008155.1	8020.0	2118.5437	4423.6665	1272.875	2.5203187	3.0	0.08615266	0.27344105	0.55	1.81																																																																																																																																																																																												
gb NM_025529.1 DB_XREF=gi:13384949 GEN=Z1310039H17Rk FEA=FLmRNA CNT=18 TID=Mm.44092.1 TIER=FL STK=2 UG=Mm.44092.1 LL=6837 DEF=Mus musculus RIKEN cDNA Z1310039H17 gene (Z1310039H17Rk), mRNA. /PROD=transmembrane cDNA Z1310039H17 FL-gb NM_025529.1	3700.1335	1125.2148	1606.6998	191.09187	3.1769488	2.0	0.08642728	0.27194792	0.43	2.3																																																																																																																																																																																												
gb AF072307.1 DB_XREF=gi:6882347 GEN=Unc51.1 FEA=FLmRNA CNT=148 TID=Mm.34216.1 TIER=FL+Slack STK=85 UG=Mm.34216.1 LL=22241 DEF=Mus musculus UNC51.1 serine/threonine kinase (Unc51.1), mRNA, complete cds. /PROD=UNC51.1 serine/threonine kinase FL-gb AF072307.1 gb AF053756.1 gb NM_009469.1	983.8667	138.9416	596.06665	228.08847	2.514983	3.0	0.08655745	0.27002966	0.61	1.65																																																																																																																																																																																												
gb AB054000.1 DB_XREF=gi:14017772 GEN=Cg10671-like FEA=FLmRNA CNT=35 TID=Mm.36758.1 TIER=FL+Slack STK=15 UG=Mm.36758.1 LL=66880 DEF=Mus musculus Cg10671-like mRNA, complete cds. /PROD=Cg10671-like FL-gb AB054000.1	1886.9332	390.19482	1093.9	189.40694	3.1668425	2.0	0.086909845	0.2688313	0.58	1.72																																																																																																																																																																																												
gb BC024621.1 DB_XREF=gi:19354257 FEA=FLmRNA CNT=208 TID=Mm.30064.1 TIER=FL+Slack STK=62 UG=Mm.30064.1 LL=56486 UG_GENE=Gabarp DEF=Mus musculus, GABA(A) receptor-associated protein, clone MGC 25654 IMAGE:4459651, mRNA, complete cds. /PROD=GABA(A) receptor-associated protein FL-gb NM_019749.1 gb BC002126.1 gb BC024621.1 gb AF161587.1	10245.833	2222.5093	5753.9663	1048.8822	3.165773	2.0	0.086961135	0.26672953	0.56	1.78																																																																																																																																																																																												
gb NM_023041.1 DB_XREF=gi:12746415 GEN=Pp1 FEA=FLmRNA CNT=210 TID=Mm.144235.1 TIER=FL STK=1 UG=Mm.144235.1 LL=19298 DEF=Mus musculus peroxisomal fatty acyltransferase protein (Pp1), mRNA. /PROD=peroxisomal fatty acyltransferase protein FL-gb NM_023041.1 gb BC019767.1	883.8	254.4808	419.06665	37.01004	3.130145	2.0	0.0886949	0.26755074	0.47	2.11	Mus musculus REF=MGC2599 DEF=Mus musculus glyceraldehyde-3-phosphate dehydrogenase mRNA, complete cds LEN=1228 C_5_..._3 represent transcript regions 5 prime, Middle, and 3 prime respectively	38164.9	9613.984	22698.932	5409.2744	2.4283547	3.0	0.093467414	0.2796361	0.59	1.68	gb NM_017393.1 DB_XREF=gi:8393155 GEN=Cp FEA=FLmRNA CNT=134 TID=Mm.15243.1 TIER=FL+Slack STK=78 UG=Mm.15243.1 LL=53895 DEF=Mus musculus caseinolytic protease, ATP-dependent, proteolytic subunit homolog (E. coli) (Cp), mRNA. /PROD=caseinolytic protease, ATP-dependent, proteolytic subunit homolog (E. coli) FL-gb NM_017393.1 gb BC001998.1 gb BC019420.1 DB_XREF=gi:18043845 FEA=FLmRNA CNT=96 TID=Mm.46654.1 TIER=FL+Slack STK=59 UG=Mm.46654.1 LL=19420 DEF=Mus musculus, Similar to chromosome 6 open reading frame 35, clone MGC:30293 IMAGE:5053733, mRNA, complete cds. /PROD=similar to chromosome 6 open reading frame 35 FL-gb BC019420.1	1815.1333	453.52466	927.93335	445.7683	2.416461	3.0	0.09446801	0.27807117	0.51	1.96	gb NM_031888.1 DB_XREF=gi:1394194 GEN=Pp1G FEA=FLmRNA CNT=386 TID=Mm.1970.1 TIER=FL+Slack STK=206 UG=Mm.1970.1 LL=10045 DEF=Mus musculus protein phosphatase 1, catalytic subunit, alpha isoform (Pp1c), mRNA. /PROD=protein phosphatase 1, catalytic subunit, alpha isoform FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	537.10004	157.27226	241.5	70.8365	2.968276	2.0	0.0972279	0.28165224	0.45	2.22	gb BC002073.1 DB_XREF=gi:12805224 FEA=FLmRNA CNT=1111 TID=Mm.137.1 TIER=FL+Slack STK=50 UG=Mm.137.1 LL=20005 UG_GENE=Soya6 DEF=Mus musculus, clone MGC 8215 IMAGE:3492808, mRNA, complete cds. /PROD=Unknown (protein for MGC 8215) FL-gb NM_009139.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	1573.6334	216.04079	988.26666	366.27304	2.384262	3.0	0.09724311	0.27947822	0.63	1.59	gb NM_009519.1 DB_XREF=gi:6678588 GEN=Wr11 FEA=FLmRNA CNT=24 TID=Mm.22182.1 TIER=FL+Slack STK=10 UG=Mm.22182.1 LL=11141 DEF=Mus musculus wingless-related MMTV integration site 11 (Wr11), mRNA. /PROD=wingless-related MMTV integration site 11 FL-gb NM_009519.1	505.4	71.000206	316.5	117.53268	2.3827572	3.0	0.09737522	0.27767152	0.63	1.6	gb BC026669.1 DB_XREF=gi:1234824 FEA=FLmRNA CNT=77 TID=Mm.15807.2 TIER=FL STK=1 UG=Mm.15807.1 LL=22340 DEF=Mus musculus vascular endothelial growth factor B precursor (VEGF-B) mRNA, complete cds. /PROD=vascular endothelial growth factor B precursor FL-gb NM_043837.1 gb U48800.1	3644.2334	921.23413	1713.7333	417.65958	3.3057415	2.0	0.08060011	0.2749443	0.47	2.13	gb AF151792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	2269.7334	651.43805	1033.5333	312.80685	2.9629393	2.0	0.097528845	0.2759537	0.46	2.2	gb AF302138.1 DB_XREF=gi:15420074 FEA=FLmRNA CNT=158 TID=Mm.22029.1 TIER=FL+Slack STK=86 UG=Mm.22029.1 LL=10222 UG_GENE=Hs7 DEF=Mus musculus Hs7-interacting protein kinase mRNAs, complete cds. /PROD=Hs7-interacting protein kinase FL-gb BC18463.1 gb AF302138.1 gb BC004756.1	1104.3334	297.56613	577.73334	241.51155	2.3799613	3.0	0.097621255	0.27409044	0.52	1.91	gb AK008049.1 DB_XREF=gi:12840262 GEN=Mg1 FEA=mRNA CNT=5 TID=Mm.194795.3 TIER=ConsEnd STK=0 UG=Mm.194795.3 LL=23945 UG_TITLE=monoglyceride lipase DEF=Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone:1700474K monoglyceride lipase, full insert sequence.	1528.3666	498.10037	664.73334	387.59006	2.3701098	3.0	0.0984942	0.27444304	0.43	2.3	gb BC021405.1 DB_XREF=gi:13007473 FEA=FLmRNA CNT=557 TID=Mm.1034.1 TIER=FL+Slack STK=330 UG=Mm.1034.1 LL=51792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	1429.6333	272.46213	914.4666	261.58444	2.362404	3.0	0.099183634	0.27425778	0.64	1.56	gb BM196939 DB_XREF=gi:17748959 DB_XREF=0304006.3 CLONE=0304006.3 FEA=mRNA CNT=57 TID=Mm.220848.1 TIER=Slack STK=16 UG=Mm.220848 UG_TITLE=Slack DEF=Mus musculus, similar to putative, clone MGC:5345087, mRNA, partial cds. /PROD=Slack FL-gb NM_0207247.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	366.83334	48.94613	229.76666	88.11454	2.355309	3.0	0.09982363	0.27395207	0.63	1.6	gb NM_023631.1 DB_XREF=gi:12072471 FEA=FLmRNA CNT=119 TID=Mm.21897.1 TIER=FL+Slack STK=30 UG=Mm.21897.1 LL=11923 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	687.3	287.4756	202.13335	12.589808	2.9203472	2.0	0.09997883	0.2723304	0.29	3.4	gb BC003744.1 DB_XREF=gi:13277677 FEA=FLmRNA CNT=135 TID=Mm.3234.1 TIER=FL+Slack STK=52 UG=Mm.3234.1 LL=19060 UG_GENE=Pp2b DEF=Mus musculus, protein phosphatase 5, catalytic subunit, clone MGC:5847 IMAGE:3599302, mRNA, complete cds. /PROD=protein phosphatase 5, catalytic subunit FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	1420.2334	425.87183	649.23334	171.36757	2.9095585	2.0	0.10061329	0.27202854	0.46	2.19	gb NM_130880.1 DB_XREF=gi:18698997 GEN=Cdk9 FEA=FLmRNA CNT=119 TID=Mm.27557.1 TIER=FL+Slack STK=50 UG=Mm.27557.1 LL=107951 DEF=Mus musculus cyclin-dependent kinase 9 (CDC-related kinase) (Cdk9), mRNA. /PROD=cyclin-dependent kinase 9 FL-gb AF327431.1 gb BC003901.1 gb NM_130860.1	513.26666	108.13493	311.26666	103.22113	2.340426	3.0	0.1011825	0.271556	0.61	1.65	gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus
Mus musculus REF=MGC2599 DEF=Mus musculus glyceraldehyde-3-phosphate dehydrogenase mRNA, complete cds LEN=1228 C_5_..._3 represent transcript regions 5 prime, Middle, and 3 prime respectively	38164.9	9613.984	22698.932	5409.2744	2.4283547	3.0	0.093467414	0.2796361	0.59	1.68	gb NM_017393.1 DB_XREF=gi:8393155 GEN=Cp FEA=FLmRNA CNT=134 TID=Mm.15243.1 TIER=FL+Slack STK=78 UG=Mm.15243.1 LL=53895 DEF=Mus musculus caseinolytic protease, ATP-dependent, proteolytic subunit homolog (E. coli) (Cp), mRNA. /PROD=caseinolytic protease, ATP-dependent, proteolytic subunit homolog (E. coli) FL-gb NM_017393.1 gb BC001998.1 gb BC019420.1 DB_XREF=gi:18043845 FEA=FLmRNA CNT=96 TID=Mm.46654.1 TIER=FL+Slack STK=59 UG=Mm.46654.1 LL=19420 DEF=Mus musculus, Similar to chromosome 6 open reading frame 35, clone MGC:30293 IMAGE:5053733, mRNA, complete cds. /PROD=similar to chromosome 6 open reading frame 35 FL-gb BC019420.1	1815.1333	453.52466	927.93335	445.7683	2.416461	3.0	0.09446801	0.27807117	0.51	1.96	gb NM_031888.1 DB_XREF=gi:1394194 GEN=Pp1G FEA=FLmRNA CNT=386 TID=Mm.1970.1 TIER=FL+Slack STK=206 UG=Mm.1970.1 LL=10045 DEF=Mus musculus protein phosphatase 1, catalytic subunit, alpha isoform (Pp1c), mRNA. /PROD=protein phosphatase 1, catalytic subunit, alpha isoform FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	537.10004	157.27226	241.5	70.8365	2.968276	2.0	0.0972279	0.28165224	0.45	2.22	gb BC002073.1 DB_XREF=gi:12805224 FEA=FLmRNA CNT=1111 TID=Mm.137.1 TIER=FL+Slack STK=50 UG=Mm.137.1 LL=20005 UG_GENE=Soya6 DEF=Mus musculus, clone MGC 8215 IMAGE:3492808, mRNA, complete cds. /PROD=Unknown (protein for MGC 8215) FL-gb NM_009139.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	1573.6334	216.04079	988.26666	366.27304	2.384262	3.0	0.09724311	0.27947822	0.63	1.59	gb NM_009519.1 DB_XREF=gi:6678588 GEN=Wr11 FEA=FLmRNA CNT=24 TID=Mm.22182.1 TIER=FL+Slack STK=10 UG=Mm.22182.1 LL=11141 DEF=Mus musculus wingless-related MMTV integration site 11 (Wr11), mRNA. /PROD=wingless-related MMTV integration site 11 FL-gb NM_009519.1	505.4	71.000206	316.5	117.53268	2.3827572	3.0	0.09737522	0.27767152	0.63	1.6	gb BC026669.1 DB_XREF=gi:1234824 FEA=FLmRNA CNT=77 TID=Mm.15807.2 TIER=FL STK=1 UG=Mm.15807.1 LL=22340 DEF=Mus musculus vascular endothelial growth factor B precursor (VEGF-B) mRNA, complete cds. /PROD=vascular endothelial growth factor B precursor FL-gb NM_043837.1 gb U48800.1	3644.2334	921.23413	1713.7333	417.65958	3.3057415	2.0	0.08060011	0.2749443	0.47	2.13	gb AF151792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	2269.7334	651.43805	1033.5333	312.80685	2.9629393	2.0	0.097528845	0.2759537	0.46	2.2	gb AF302138.1 DB_XREF=gi:15420074 FEA=FLmRNA CNT=158 TID=Mm.22029.1 TIER=FL+Slack STK=86 UG=Mm.22029.1 LL=10222 UG_GENE=Hs7 DEF=Mus musculus Hs7-interacting protein kinase mRNAs, complete cds. /PROD=Hs7-interacting protein kinase FL-gb BC18463.1 gb AF302138.1 gb BC004756.1	1104.3334	297.56613	577.73334	241.51155	2.3799613	3.0	0.097621255	0.27409044	0.52	1.91	gb AK008049.1 DB_XREF=gi:12840262 GEN=Mg1 FEA=mRNA CNT=5 TID=Mm.194795.3 TIER=ConsEnd STK=0 UG=Mm.194795.3 LL=23945 UG_TITLE=monoglyceride lipase DEF=Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone:1700474K monoglyceride lipase, full insert sequence.	1528.3666	498.10037	664.73334	387.59006	2.3701098	3.0	0.0984942	0.27444304	0.43	2.3	gb BC021405.1 DB_XREF=gi:13007473 FEA=FLmRNA CNT=557 TID=Mm.1034.1 TIER=FL+Slack STK=330 UG=Mm.1034.1 LL=51792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	1429.6333	272.46213	914.4666	261.58444	2.362404	3.0	0.099183634	0.27425778	0.64	1.56	gb BM196939 DB_XREF=gi:17748959 DB_XREF=0304006.3 CLONE=0304006.3 FEA=mRNA CNT=57 TID=Mm.220848.1 TIER=Slack STK=16 UG=Mm.220848 UG_TITLE=Slack DEF=Mus musculus, similar to putative, clone MGC:5345087, mRNA, partial cds. /PROD=Slack FL-gb NM_0207247.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	366.83334	48.94613	229.76666	88.11454	2.355309	3.0	0.09982363	0.27395207	0.63	1.6	gb NM_023631.1 DB_XREF=gi:12072471 FEA=FLmRNA CNT=119 TID=Mm.21897.1 TIER=FL+Slack STK=30 UG=Mm.21897.1 LL=11923 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	687.3	287.4756	202.13335	12.589808	2.9203472	2.0	0.09997883	0.2723304	0.29	3.4	gb BC003744.1 DB_XREF=gi:13277677 FEA=FLmRNA CNT=135 TID=Mm.3234.1 TIER=FL+Slack STK=52 UG=Mm.3234.1 LL=19060 UG_GENE=Pp2b DEF=Mus musculus, protein phosphatase 5, catalytic subunit, clone MGC:5847 IMAGE:3599302, mRNA, complete cds. /PROD=protein phosphatase 5, catalytic subunit FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	1420.2334	425.87183	649.23334	171.36757	2.9095585	2.0	0.10061329	0.27202854	0.46	2.19	gb NM_130880.1 DB_XREF=gi:18698997 GEN=Cdk9 FEA=FLmRNA CNT=119 TID=Mm.27557.1 TIER=FL+Slack STK=50 UG=Mm.27557.1 LL=107951 DEF=Mus musculus cyclin-dependent kinase 9 (CDC-related kinase) (Cdk9), mRNA. /PROD=cyclin-dependent kinase 9 FL-gb AF327431.1 gb BC003901.1 gb NM_130860.1	513.26666	108.13493	311.26666	103.22113	2.340426	3.0	0.1011825	0.271556	0.61	1.65	gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus											
gb NM_017393.1 DB_XREF=gi:8393155 GEN=Cp FEA=FLmRNA CNT=134 TID=Mm.15243.1 TIER=FL+Slack STK=78 UG=Mm.15243.1 LL=53895 DEF=Mus musculus caseinolytic protease, ATP-dependent, proteolytic subunit homolog (E. coli) (Cp), mRNA. /PROD=caseinolytic protease, ATP-dependent, proteolytic subunit homolog (E. coli) FL-gb NM_017393.1 gb BC001998.1 gb BC019420.1 DB_XREF=gi:18043845 FEA=FLmRNA CNT=96 TID=Mm.46654.1 TIER=FL+Slack STK=59 UG=Mm.46654.1 LL=19420 DEF=Mus musculus, Similar to chromosome 6 open reading frame 35, clone MGC:30293 IMAGE:5053733, mRNA, complete cds. /PROD=similar to chromosome 6 open reading frame 35 FL-gb BC019420.1	1815.1333	453.52466	927.93335	445.7683	2.416461	3.0	0.09446801	0.27807117	0.51	1.96	gb NM_031888.1 DB_XREF=gi:1394194 GEN=Pp1G FEA=FLmRNA CNT=386 TID=Mm.1970.1 TIER=FL+Slack STK=206 UG=Mm.1970.1 LL=10045 DEF=Mus musculus protein phosphatase 1, catalytic subunit, alpha isoform (Pp1c), mRNA. /PROD=protein phosphatase 1, catalytic subunit, alpha isoform FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	537.10004	157.27226	241.5	70.8365	2.968276	2.0	0.0972279	0.28165224	0.45	2.22	gb BC002073.1 DB_XREF=gi:12805224 FEA=FLmRNA CNT=1111 TID=Mm.137.1 TIER=FL+Slack STK=50 UG=Mm.137.1 LL=20005 UG_GENE=Soya6 DEF=Mus musculus, clone MGC 8215 IMAGE:3492808, mRNA, complete cds. /PROD=Unknown (protein for MGC 8215) FL-gb NM_009139.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	1573.6334	216.04079	988.26666	366.27304	2.384262	3.0	0.09724311	0.27947822	0.63	1.59	gb NM_009519.1 DB_XREF=gi:6678588 GEN=Wr11 FEA=FLmRNA CNT=24 TID=Mm.22182.1 TIER=FL+Slack STK=10 UG=Mm.22182.1 LL=11141 DEF=Mus musculus wingless-related MMTV integration site 11 (Wr11), mRNA. /PROD=wingless-related MMTV integration site 11 FL-gb NM_009519.1	505.4	71.000206	316.5	117.53268	2.3827572	3.0	0.09737522	0.27767152	0.63	1.6	gb BC026669.1 DB_XREF=gi:1234824 FEA=FLmRNA CNT=77 TID=Mm.15807.2 TIER=FL STK=1 UG=Mm.15807.1 LL=22340 DEF=Mus musculus vascular endothelial growth factor B precursor (VEGF-B) mRNA, complete cds. /PROD=vascular endothelial growth factor B precursor FL-gb NM_043837.1 gb U48800.1	3644.2334	921.23413	1713.7333	417.65958	3.3057415	2.0	0.08060011	0.2749443	0.47	2.13	gb AF151792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	2269.7334	651.43805	1033.5333	312.80685	2.9629393	2.0	0.097528845	0.2759537	0.46	2.2	gb AF302138.1 DB_XREF=gi:15420074 FEA=FLmRNA CNT=158 TID=Mm.22029.1 TIER=FL+Slack STK=86 UG=Mm.22029.1 LL=10222 UG_GENE=Hs7 DEF=Mus musculus Hs7-interacting protein kinase mRNAs, complete cds. /PROD=Hs7-interacting protein kinase FL-gb BC18463.1 gb AF302138.1 gb BC004756.1	1104.3334	297.56613	577.73334	241.51155	2.3799613	3.0	0.097621255	0.27409044	0.52	1.91	gb AK008049.1 DB_XREF=gi:12840262 GEN=Mg1 FEA=mRNA CNT=5 TID=Mm.194795.3 TIER=ConsEnd STK=0 UG=Mm.194795.3 LL=23945 UG_TITLE=monoglyceride lipase DEF=Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone:1700474K monoglyceride lipase, full insert sequence.	1528.3666	498.10037	664.73334	387.59006	2.3701098	3.0	0.0984942	0.27444304	0.43	2.3	gb BC021405.1 DB_XREF=gi:13007473 FEA=FLmRNA CNT=557 TID=Mm.1034.1 TIER=FL+Slack STK=330 UG=Mm.1034.1 LL=51792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	1429.6333	272.46213	914.4666	261.58444	2.362404	3.0	0.099183634	0.27425778	0.64	1.56	gb BM196939 DB_XREF=gi:17748959 DB_XREF=0304006.3 CLONE=0304006.3 FEA=mRNA CNT=57 TID=Mm.220848.1 TIER=Slack STK=16 UG=Mm.220848 UG_TITLE=Slack DEF=Mus musculus, similar to putative, clone MGC:5345087, mRNA, partial cds. /PROD=Slack FL-gb NM_0207247.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	366.83334	48.94613	229.76666	88.11454	2.355309	3.0	0.09982363	0.27395207	0.63	1.6	gb NM_023631.1 DB_XREF=gi:12072471 FEA=FLmRNA CNT=119 TID=Mm.21897.1 TIER=FL+Slack STK=30 UG=Mm.21897.1 LL=11923 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	687.3	287.4756	202.13335	12.589808	2.9203472	2.0	0.09997883	0.2723304	0.29	3.4	gb BC003744.1 DB_XREF=gi:13277677 FEA=FLmRNA CNT=135 TID=Mm.3234.1 TIER=FL+Slack STK=52 UG=Mm.3234.1 LL=19060 UG_GENE=Pp2b DEF=Mus musculus, protein phosphatase 5, catalytic subunit, clone MGC:5847 IMAGE:3599302, mRNA, complete cds. /PROD=protein phosphatase 5, catalytic subunit FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	1420.2334	425.87183	649.23334	171.36757	2.9095585	2.0	0.10061329	0.27202854	0.46	2.19	gb NM_130880.1 DB_XREF=gi:18698997 GEN=Cdk9 FEA=FLmRNA CNT=119 TID=Mm.27557.1 TIER=FL+Slack STK=50 UG=Mm.27557.1 LL=107951 DEF=Mus musculus cyclin-dependent kinase 9 (CDC-related kinase) (Cdk9), mRNA. /PROD=cyclin-dependent kinase 9 FL-gb AF327431.1 gb BC003901.1 gb NM_130860.1	513.26666	108.13493	311.26666	103.22113	2.340426	3.0	0.1011825	0.271556	0.61	1.65	gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																						
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AF128189.1 gb AF128188.1	1573.6334	216.04079	988.26666	366.27304	2.384262	3.0	0.09724311	0.27947822	0.63	1.59	gb NM_009519.1 DB_XREF=gi:6678588 GEN=Wr11 FEA=FLmRNA CNT=24 TID=Mm.22182.1 TIER=FL+Slack STK=10 UG=Mm.22182.1 LL=11141 DEF=Mus musculus wingless-related MMTV integration site 11 (Wr11), mRNA. /PROD=wingless-related MMTV integration site 11 FL-gb NM_009519.1	505.4	71.000206	316.5	117.53268	2.3827572	3.0	0.09737522	0.27767152	0.63	1.6	gb BC026669.1 DB_XREF=gi:1234824 FEA=FLmRNA CNT=77 TID=Mm.15807.2 TIER=FL STK=1 UG=Mm.15807.1 LL=22340 DEF=Mus musculus vascular endothelial growth factor B precursor (VEGF-B) mRNA, complete cds. /PROD=vascular endothelial growth factor B precursor FL-gb NM_043837.1 gb U48800.1	3644.2334	921.23413	1713.7333	417.65958	3.3057415	2.0	0.08060011	0.2749443	0.47	2.13	gb AF151792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	2269.7334	651.43805	1033.5333	312.80685	2.9629393	2.0	0.097528845	0.2759537	0.46	2.2	gb AF302138.1 DB_XREF=gi:15420074 FEA=FLmRNA CNT=158 TID=Mm.22029.1 TIER=FL+Slack STK=86 UG=Mm.22029.1 LL=10222 UG_GENE=Hs7 DEF=Mus musculus Hs7-interacting protein kinase mRNAs, complete cds. /PROD=Hs7-interacting protein kinase FL-gb BC18463.1 gb AF302138.1 gb BC004756.1	1104.3334	297.56613	577.73334	241.51155	2.3799613	3.0	0.097621255	0.27409044	0.52	1.91	gb AK008049.1 DB_XREF=gi:12840262 GEN=Mg1 FEA=mRNA CNT=5 TID=Mm.194795.3 TIER=ConsEnd STK=0 UG=Mm.194795.3 LL=23945 UG_TITLE=monoglyceride lipase DEF=Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone:1700474K monoglyceride lipase, full insert sequence.	1528.3666	498.10037	664.73334	387.59006	2.3701098	3.0	0.0984942	0.27444304	0.43	2.3	gb BC021405.1 DB_XREF=gi:13007473 FEA=FLmRNA CNT=557 TID=Mm.1034.1 TIER=FL+Slack STK=330 UG=Mm.1034.1 LL=51792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	1429.6333	272.46213	914.4666	261.58444	2.362404	3.0	0.099183634	0.27425778	0.64	1.56	gb BM196939 DB_XREF=gi:17748959 DB_XREF=0304006.3 CLONE=0304006.3 FEA=mRNA CNT=57 TID=Mm.220848.1 TIER=Slack STK=16 UG=Mm.220848 UG_TITLE=Slack DEF=Mus musculus, similar to putative, clone MGC:5345087, mRNA, partial cds. /PROD=Slack FL-gb NM_0207247.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	366.83334	48.94613	229.76666	88.11454	2.355309	3.0	0.09982363	0.27395207	0.63	1.6	gb NM_023631.1 DB_XREF=gi:12072471 FEA=FLmRNA CNT=119 TID=Mm.21897.1 TIER=FL+Slack STK=30 UG=Mm.21897.1 LL=11923 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	687.3	287.4756	202.13335	12.589808	2.9203472	2.0	0.09997883	0.2723304	0.29	3.4	gb BC003744.1 DB_XREF=gi:13277677 FEA=FLmRNA CNT=135 TID=Mm.3234.1 TIER=FL+Slack STK=52 UG=Mm.3234.1 LL=19060 UG_GENE=Pp2b DEF=Mus musculus, protein phosphatase 5, catalytic subunit, clone MGC:5847 IMAGE:3599302, mRNA, complete cds. /PROD=protein phosphatase 5, catalytic subunit FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	1420.2334	425.87183	649.23334	171.36757	2.9095585	2.0	0.10061329	0.27202854	0.46	2.19	gb NM_130880.1 DB_XREF=gi:18698997 GEN=Cdk9 FEA=FLmRNA CNT=119 TID=Mm.27557.1 TIER=FL+Slack STK=50 UG=Mm.27557.1 LL=107951 DEF=Mus musculus cyclin-dependent kinase 9 (CDC-related kinase) (Cdk9), mRNA. /PROD=cyclin-dependent kinase 9 FL-gb AF327431.1 gb BC003901.1 gb NM_130860.1	513.26666	108.13493	311.26666	103.22113	2.340426	3.0	0.1011825	0.271556	0.61	1.65	gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																	
gb BC002073.1 DB_XREF=gi:12805224 FEA=FLmRNA CNT=1111 TID=Mm.137.1 TIER=FL+Slack STK=50 UG=Mm.137.1 LL=20005 UG_GENE=Soya6 DEF=Mus musculus, clone MGC 8215 IMAGE:3492808, mRNA, complete cds. /PROD=Unknown (protein for MGC 8215) FL-gb NM_009139.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	1573.6334	216.04079	988.26666	366.27304	2.384262	3.0	0.09724311	0.27947822	0.63	1.59	gb NM_009519.1 DB_XREF=gi:6678588 GEN=Wr11 FEA=FLmRNA CNT=24 TID=Mm.22182.1 TIER=FL+Slack STK=10 UG=Mm.22182.1 LL=11141 DEF=Mus musculus wingless-related MMTV integration site 11 (Wr11), mRNA. /PROD=wingless-related MMTV integration site 11 FL-gb NM_009519.1	505.4	71.000206	316.5	117.53268	2.3827572	3.0	0.09737522	0.27767152	0.63	1.6	gb BC026669.1 DB_XREF=gi:1234824 FEA=FLmRNA CNT=77 TID=Mm.15807.2 TIER=FL STK=1 UG=Mm.15807.1 LL=22340 DEF=Mus musculus vascular endothelial growth factor B precursor (VEGF-B) mRNA, complete cds. /PROD=vascular endothelial growth factor B precursor FL-gb NM_043837.1 gb U48800.1	3644.2334	921.23413	1713.7333	417.65958	3.3057415	2.0	0.08060011	0.2749443	0.47	2.13	gb AF151792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	2269.7334	651.43805	1033.5333	312.80685	2.9629393	2.0	0.097528845	0.2759537	0.46	2.2	gb AF302138.1 DB_XREF=gi:15420074 FEA=FLmRNA CNT=158 TID=Mm.22029.1 TIER=FL+Slack STK=86 UG=Mm.22029.1 LL=10222 UG_GENE=Hs7 DEF=Mus musculus Hs7-interacting protein kinase mRNAs, complete cds. /PROD=Hs7-interacting protein kinase FL-gb BC18463.1 gb AF302138.1 gb BC004756.1	1104.3334	297.56613	577.73334	241.51155	2.3799613	3.0	0.097621255	0.27409044	0.52	1.91	gb AK008049.1 DB_XREF=gi:12840262 GEN=Mg1 FEA=mRNA CNT=5 TID=Mm.194795.3 TIER=ConsEnd STK=0 UG=Mm.194795.3 LL=23945 UG_TITLE=monoglyceride lipase DEF=Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone:1700474K monoglyceride lipase, full insert sequence.	1528.3666	498.10037	664.73334	387.59006	2.3701098	3.0	0.0984942	0.27444304	0.43	2.3	gb BC021405.1 DB_XREF=gi:13007473 FEA=FLmRNA CNT=557 TID=Mm.1034.1 TIER=FL+Slack STK=330 UG=Mm.1034.1 LL=51792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	1429.6333	272.46213	914.4666	261.58444	2.362404	3.0	0.099183634	0.27425778	0.64	1.56	gb BM196939 DB_XREF=gi:17748959 DB_XREF=0304006.3 CLONE=0304006.3 FEA=mRNA CNT=57 TID=Mm.220848.1 TIER=Slack STK=16 UG=Mm.220848 UG_TITLE=Slack DEF=Mus musculus, similar to putative, clone MGC:5345087, mRNA, partial cds. /PROD=Slack FL-gb NM_0207247.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	366.83334	48.94613	229.76666	88.11454	2.355309	3.0	0.09982363	0.27395207	0.63	1.6	gb NM_023631.1 DB_XREF=gi:12072471 FEA=FLmRNA CNT=119 TID=Mm.21897.1 TIER=FL+Slack STK=30 UG=Mm.21897.1 LL=11923 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	687.3	287.4756	202.13335	12.589808	2.9203472	2.0	0.09997883	0.2723304	0.29	3.4	gb BC003744.1 DB_XREF=gi:13277677 FEA=FLmRNA CNT=135 TID=Mm.3234.1 TIER=FL+Slack STK=52 UG=Mm.3234.1 LL=19060 UG_GENE=Pp2b DEF=Mus musculus, protein phosphatase 5, catalytic subunit, clone MGC:5847 IMAGE:3599302, mRNA, complete cds. /PROD=protein phosphatase 5, catalytic subunit FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	1420.2334	425.87183	649.23334	171.36757	2.9095585	2.0	0.10061329	0.27202854	0.46	2.19	gb NM_130880.1 DB_XREF=gi:18698997 GEN=Cdk9 FEA=FLmRNA CNT=119 TID=Mm.27557.1 TIER=FL+Slack STK=50 UG=Mm.27557.1 LL=107951 DEF=Mus musculus cyclin-dependent kinase 9 (CDC-related kinase) (Cdk9), mRNA. /PROD=cyclin-dependent kinase 9 FL-gb AF327431.1 gb BC003901.1 gb NM_130860.1	513.26666	108.13493	311.26666	103.22113	2.340426	3.0	0.1011825	0.271556	0.61	1.65	gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																												
gb NM_009519.1 DB_XREF=gi:6678588 GEN=Wr11 FEA=FLmRNA CNT=24 TID=Mm.22182.1 TIER=FL+Slack STK=10 UG=Mm.22182.1 LL=11141 DEF=Mus musculus wingless-related MMTV integration site 11 (Wr11), mRNA. /PROD=wingless-related MMTV integration site 11 FL-gb NM_009519.1	505.4	71.000206	316.5	117.53268	2.3827572	3.0	0.09737522	0.27767152	0.63	1.6	gb BC026669.1 DB_XREF=gi:1234824 FEA=FLmRNA CNT=77 TID=Mm.15807.2 TIER=FL STK=1 UG=Mm.15807.1 LL=22340 DEF=Mus musculus vascular endothelial growth factor B precursor (VEGF-B) mRNA, complete cds. /PROD=vascular endothelial growth factor B precursor FL-gb NM_043837.1 gb U48800.1	3644.2334	921.23413	1713.7333	417.65958	3.3057415	2.0	0.08060011	0.2749443	0.47	2.13	gb AF151792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	2269.7334	651.43805	1033.5333	312.80685	2.9629393	2.0	0.097528845	0.2759537	0.46	2.2	gb AF302138.1 DB_XREF=gi:15420074 FEA=FLmRNA CNT=158 TID=Mm.22029.1 TIER=FL+Slack STK=86 UG=Mm.22029.1 LL=10222 UG_GENE=Hs7 DEF=Mus musculus Hs7-interacting protein kinase mRNAs, complete cds. /PROD=Hs7-interacting protein kinase FL-gb BC18463.1 gb AF302138.1 gb BC004756.1	1104.3334	297.56613	577.73334	241.51155	2.3799613	3.0	0.097621255	0.27409044	0.52	1.91	gb AK008049.1 DB_XREF=gi:12840262 GEN=Mg1 FEA=mRNA CNT=5 TID=Mm.194795.3 TIER=ConsEnd STK=0 UG=Mm.194795.3 LL=23945 UG_TITLE=monoglyceride lipase DEF=Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone:1700474K monoglyceride lipase, full insert sequence.	1528.3666	498.10037	664.73334	387.59006	2.3701098	3.0	0.0984942	0.27444304	0.43	2.3	gb BC021405.1 DB_XREF=gi:13007473 FEA=FLmRNA CNT=557 TID=Mm.1034.1 TIER=FL+Slack STK=330 UG=Mm.1034.1 LL=51792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	1429.6333	272.46213	914.4666	261.58444	2.362404	3.0	0.099183634	0.27425778	0.64	1.56	gb BM196939 DB_XREF=gi:17748959 DB_XREF=0304006.3 CLONE=0304006.3 FEA=mRNA CNT=57 TID=Mm.220848.1 TIER=Slack STK=16 UG=Mm.220848 UG_TITLE=Slack DEF=Mus musculus, similar to putative, clone MGC:5345087, mRNA, partial cds. /PROD=Slack FL-gb NM_0207247.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	366.83334	48.94613	229.76666	88.11454	2.355309	3.0	0.09982363	0.27395207	0.63	1.6	gb NM_023631.1 DB_XREF=gi:12072471 FEA=FLmRNA CNT=119 TID=Mm.21897.1 TIER=FL+Slack STK=30 UG=Mm.21897.1 LL=11923 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	687.3	287.4756	202.13335	12.589808	2.9203472	2.0	0.09997883	0.2723304	0.29	3.4	gb BC003744.1 DB_XREF=gi:13277677 FEA=FLmRNA CNT=135 TID=Mm.3234.1 TIER=FL+Slack STK=52 UG=Mm.3234.1 LL=19060 UG_GENE=Pp2b DEF=Mus musculus, protein phosphatase 5, catalytic subunit, clone MGC:5847 IMAGE:3599302, mRNA, complete cds. /PROD=protein phosphatase 5, catalytic subunit FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	1420.2334	425.87183	649.23334	171.36757	2.9095585	2.0	0.10061329	0.27202854	0.46	2.19	gb NM_130880.1 DB_XREF=gi:18698997 GEN=Cdk9 FEA=FLmRNA CNT=119 TID=Mm.27557.1 TIER=FL+Slack STK=50 UG=Mm.27557.1 LL=107951 DEF=Mus musculus cyclin-dependent kinase 9 (CDC-related kinase) (Cdk9), mRNA. /PROD=cyclin-dependent kinase 9 FL-gb AF327431.1 gb BC003901.1 gb NM_130860.1	513.26666	108.13493	311.26666	103.22113	2.340426	3.0	0.1011825	0.271556	0.61	1.65	gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																																							
gb BC026669.1 DB_XREF=gi:1234824 FEA=FLmRNA CNT=77 TID=Mm.15807.2 TIER=FL STK=1 UG=Mm.15807.1 LL=22340 DEF=Mus musculus vascular endothelial growth factor B precursor (VEGF-B) mRNA, complete cds. /PROD=vascular endothelial growth factor B precursor FL-gb NM_043837.1 gb U48800.1	3644.2334	921.23413	1713.7333	417.65958	3.3057415	2.0	0.08060011	0.2749443	0.47	2.13	gb AF151792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	2269.7334	651.43805	1033.5333	312.80685	2.9629393	2.0	0.097528845	0.2759537	0.46	2.2	gb AF302138.1 DB_XREF=gi:15420074 FEA=FLmRNA CNT=158 TID=Mm.22029.1 TIER=FL+Slack STK=86 UG=Mm.22029.1 LL=10222 UG_GENE=Hs7 DEF=Mus musculus Hs7-interacting protein kinase mRNAs, complete cds. /PROD=Hs7-interacting protein kinase FL-gb BC18463.1 gb AF302138.1 gb BC004756.1	1104.3334	297.56613	577.73334	241.51155	2.3799613	3.0	0.097621255	0.27409044	0.52	1.91	gb AK008049.1 DB_XREF=gi:12840262 GEN=Mg1 FEA=mRNA CNT=5 TID=Mm.194795.3 TIER=ConsEnd STK=0 UG=Mm.194795.3 LL=23945 UG_TITLE=monoglyceride lipase DEF=Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone:1700474K monoglyceride lipase, full insert sequence.	1528.3666	498.10037	664.73334	387.59006	2.3701098	3.0	0.0984942	0.27444304	0.43	2.3	gb BC021405.1 DB_XREF=gi:13007473 FEA=FLmRNA CNT=557 TID=Mm.1034.1 TIER=FL+Slack STK=330 UG=Mm.1034.1 LL=51792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	1429.6333	272.46213	914.4666	261.58444	2.362404	3.0	0.099183634	0.27425778	0.64	1.56	gb BM196939 DB_XREF=gi:17748959 DB_XREF=0304006.3 CLONE=0304006.3 FEA=mRNA CNT=57 TID=Mm.220848.1 TIER=Slack STK=16 UG=Mm.220848 UG_TITLE=Slack DEF=Mus musculus, similar to putative, clone MGC:5345087, mRNA, partial cds. /PROD=Slack FL-gb NM_0207247.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	366.83334	48.94613	229.76666	88.11454	2.355309	3.0	0.09982363	0.27395207	0.63	1.6	gb NM_023631.1 DB_XREF=gi:12072471 FEA=FLmRNA CNT=119 TID=Mm.21897.1 TIER=FL+Slack STK=30 UG=Mm.21897.1 LL=11923 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	687.3	287.4756	202.13335	12.589808	2.9203472	2.0	0.09997883	0.2723304	0.29	3.4	gb BC003744.1 DB_XREF=gi:13277677 FEA=FLmRNA CNT=135 TID=Mm.3234.1 TIER=FL+Slack STK=52 UG=Mm.3234.1 LL=19060 UG_GENE=Pp2b DEF=Mus musculus, protein phosphatase 5, catalytic subunit, clone MGC:5847 IMAGE:3599302, mRNA, complete cds. /PROD=protein phosphatase 5, catalytic subunit FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	1420.2334	425.87183	649.23334	171.36757	2.9095585	2.0	0.10061329	0.27202854	0.46	2.19	gb NM_130880.1 DB_XREF=gi:18698997 GEN=Cdk9 FEA=FLmRNA CNT=119 TID=Mm.27557.1 TIER=FL+Slack STK=50 UG=Mm.27557.1 LL=107951 DEF=Mus musculus cyclin-dependent kinase 9 (CDC-related kinase) (Cdk9), mRNA. /PROD=cyclin-dependent kinase 9 FL-gb AF327431.1 gb BC003901.1 gb NM_130860.1	513.26666	108.13493	311.26666	103.22113	2.340426	3.0	0.1011825	0.271556	0.61	1.65	gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																																																		
gb AF151792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	2269.7334	651.43805	1033.5333	312.80685	2.9629393	2.0	0.097528845	0.2759537	0.46	2.2	gb AF302138.1 DB_XREF=gi:15420074 FEA=FLmRNA CNT=158 TID=Mm.22029.1 TIER=FL+Slack STK=86 UG=Mm.22029.1 LL=10222 UG_GENE=Hs7 DEF=Mus musculus Hs7-interacting protein kinase mRNAs, complete cds. /PROD=Hs7-interacting protein kinase FL-gb BC18463.1 gb AF302138.1 gb BC004756.1	1104.3334	297.56613	577.73334	241.51155	2.3799613	3.0	0.097621255	0.27409044	0.52	1.91	gb AK008049.1 DB_XREF=gi:12840262 GEN=Mg1 FEA=mRNA CNT=5 TID=Mm.194795.3 TIER=ConsEnd STK=0 UG=Mm.194795.3 LL=23945 UG_TITLE=monoglyceride lipase DEF=Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone:1700474K monoglyceride lipase, full insert sequence.	1528.3666	498.10037	664.73334	387.59006	2.3701098	3.0	0.0984942	0.27444304	0.43	2.3	gb BC021405.1 DB_XREF=gi:13007473 FEA=FLmRNA CNT=557 TID=Mm.1034.1 TIER=FL+Slack STK=330 UG=Mm.1034.1 LL=51792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	1429.6333	272.46213	914.4666	261.58444	2.362404	3.0	0.099183634	0.27425778	0.64	1.56	gb BM196939 DB_XREF=gi:17748959 DB_XREF=0304006.3 CLONE=0304006.3 FEA=mRNA CNT=57 TID=Mm.220848.1 TIER=Slack STK=16 UG=Mm.220848 UG_TITLE=Slack DEF=Mus musculus, similar to putative, clone MGC:5345087, mRNA, partial cds. /PROD=Slack FL-gb NM_0207247.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	366.83334	48.94613	229.76666	88.11454	2.355309	3.0	0.09982363	0.27395207	0.63	1.6	gb NM_023631.1 DB_XREF=gi:12072471 FEA=FLmRNA CNT=119 TID=Mm.21897.1 TIER=FL+Slack STK=30 UG=Mm.21897.1 LL=11923 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	687.3	287.4756	202.13335	12.589808	2.9203472	2.0	0.09997883	0.2723304	0.29	3.4	gb BC003744.1 DB_XREF=gi:13277677 FEA=FLmRNA CNT=135 TID=Mm.3234.1 TIER=FL+Slack STK=52 UG=Mm.3234.1 LL=19060 UG_GENE=Pp2b DEF=Mus musculus, protein phosphatase 5, catalytic subunit, clone MGC:5847 IMAGE:3599302, mRNA, complete cds. /PROD=protein phosphatase 5, catalytic subunit FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	1420.2334	425.87183	649.23334	171.36757	2.9095585	2.0	0.10061329	0.27202854	0.46	2.19	gb NM_130880.1 DB_XREF=gi:18698997 GEN=Cdk9 FEA=FLmRNA CNT=119 TID=Mm.27557.1 TIER=FL+Slack STK=50 UG=Mm.27557.1 LL=107951 DEF=Mus musculus cyclin-dependent kinase 9 (CDC-related kinase) (Cdk9), mRNA. /PROD=cyclin-dependent kinase 9 FL-gb AF327431.1 gb BC003901.1 gb NM_130860.1	513.26666	108.13493	311.26666	103.22113	2.340426	3.0	0.1011825	0.271556	0.61	1.65	gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																																																													
gb AF302138.1 DB_XREF=gi:15420074 FEA=FLmRNA CNT=158 TID=Mm.22029.1 TIER=FL+Slack STK=86 UG=Mm.22029.1 LL=10222 UG_GENE=Hs7 DEF=Mus musculus Hs7-interacting protein kinase mRNAs, complete cds. /PROD=Hs7-interacting protein kinase FL-gb BC18463.1 gb AF302138.1 gb BC004756.1	1104.3334	297.56613	577.73334	241.51155	2.3799613	3.0	0.097621255	0.27409044	0.52	1.91	gb AK008049.1 DB_XREF=gi:12840262 GEN=Mg1 FEA=mRNA CNT=5 TID=Mm.194795.3 TIER=ConsEnd STK=0 UG=Mm.194795.3 LL=23945 UG_TITLE=monoglyceride lipase DEF=Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone:1700474K monoglyceride lipase, full insert sequence.	1528.3666	498.10037	664.73334	387.59006	2.3701098	3.0	0.0984942	0.27444304	0.43	2.3	gb BC021405.1 DB_XREF=gi:13007473 FEA=FLmRNA CNT=557 TID=Mm.1034.1 TIER=FL+Slack STK=330 UG=Mm.1034.1 LL=51792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	1429.6333	272.46213	914.4666	261.58444	2.362404	3.0	0.099183634	0.27425778	0.64	1.56	gb BM196939 DB_XREF=gi:17748959 DB_XREF=0304006.3 CLONE=0304006.3 FEA=mRNA CNT=57 TID=Mm.220848.1 TIER=Slack STK=16 UG=Mm.220848 UG_TITLE=Slack DEF=Mus musculus, similar to putative, clone MGC:5345087, mRNA, partial cds. /PROD=Slack FL-gb NM_0207247.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	366.83334	48.94613	229.76666	88.11454	2.355309	3.0	0.09982363	0.27395207	0.63	1.6	gb NM_023631.1 DB_XREF=gi:12072471 FEA=FLmRNA CNT=119 TID=Mm.21897.1 TIER=FL+Slack STK=30 UG=Mm.21897.1 LL=11923 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	687.3	287.4756	202.13335	12.589808	2.9203472	2.0	0.09997883	0.2723304	0.29	3.4	gb BC003744.1 DB_XREF=gi:13277677 FEA=FLmRNA CNT=135 TID=Mm.3234.1 TIER=FL+Slack STK=52 UG=Mm.3234.1 LL=19060 UG_GENE=Pp2b DEF=Mus musculus, protein phosphatase 5, catalytic subunit, clone MGC:5847 IMAGE:3599302, mRNA, complete cds. /PROD=protein phosphatase 5, catalytic subunit FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	1420.2334	425.87183	649.23334	171.36757	2.9095585	2.0	0.10061329	0.27202854	0.46	2.19	gb NM_130880.1 DB_XREF=gi:18698997 GEN=Cdk9 FEA=FLmRNA CNT=119 TID=Mm.27557.1 TIER=FL+Slack STK=50 UG=Mm.27557.1 LL=107951 DEF=Mus musculus cyclin-dependent kinase 9 (CDC-related kinase) (Cdk9), mRNA. /PROD=cyclin-dependent kinase 9 FL-gb AF327431.1 gb BC003901.1 gb NM_130860.1	513.26666	108.13493	311.26666	103.22113	2.340426	3.0	0.1011825	0.271556	0.61	1.65	gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																																																																								
gb AK008049.1 DB_XREF=gi:12840262 GEN=Mg1 FEA=mRNA CNT=5 TID=Mm.194795.3 TIER=ConsEnd STK=0 UG=Mm.194795.3 LL=23945 UG_TITLE=monoglyceride lipase DEF=Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone:1700474K monoglyceride lipase, full insert sequence.	1528.3666	498.10037	664.73334	387.59006	2.3701098	3.0	0.0984942	0.27444304	0.43	2.3	gb BC021405.1 DB_XREF=gi:13007473 FEA=FLmRNA CNT=557 TID=Mm.1034.1 TIER=FL+Slack STK=330 UG=Mm.1034.1 LL=51792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	1429.6333	272.46213	914.4666	261.58444	2.362404	3.0	0.099183634	0.27425778	0.64	1.56	gb BM196939 DB_XREF=gi:17748959 DB_XREF=0304006.3 CLONE=0304006.3 FEA=mRNA CNT=57 TID=Mm.220848.1 TIER=Slack STK=16 UG=Mm.220848 UG_TITLE=Slack DEF=Mus musculus, similar to putative, clone MGC:5345087, mRNA, partial cds. /PROD=Slack FL-gb NM_0207247.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	366.83334	48.94613	229.76666	88.11454	2.355309	3.0	0.09982363	0.27395207	0.63	1.6	gb NM_023631.1 DB_XREF=gi:12072471 FEA=FLmRNA CNT=119 TID=Mm.21897.1 TIER=FL+Slack STK=30 UG=Mm.21897.1 LL=11923 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	687.3	287.4756	202.13335	12.589808	2.9203472	2.0	0.09997883	0.2723304	0.29	3.4	gb BC003744.1 DB_XREF=gi:13277677 FEA=FLmRNA CNT=135 TID=Mm.3234.1 TIER=FL+Slack STK=52 UG=Mm.3234.1 LL=19060 UG_GENE=Pp2b DEF=Mus musculus, protein phosphatase 5, catalytic subunit, clone MGC:5847 IMAGE:3599302, mRNA, complete cds. /PROD=protein phosphatase 5, catalytic subunit FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	1420.2334	425.87183	649.23334	171.36757	2.9095585	2.0	0.10061329	0.27202854	0.46	2.19	gb NM_130880.1 DB_XREF=gi:18698997 GEN=Cdk9 FEA=FLmRNA CNT=119 TID=Mm.27557.1 TIER=FL+Slack STK=50 UG=Mm.27557.1 LL=107951 DEF=Mus musculus cyclin-dependent kinase 9 (CDC-related kinase) (Cdk9), mRNA. /PROD=cyclin-dependent kinase 9 FL-gb AF327431.1 gb BC003901.1 gb NM_130860.1	513.26666	108.13493	311.26666	103.22113	2.340426	3.0	0.1011825	0.271556	0.61	1.65	gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																																																																																			
gb BC021405.1 DB_XREF=gi:13007473 FEA=FLmRNA CNT=557 TID=Mm.1034.1 TIER=FL+Slack STK=330 UG=Mm.1034.1 LL=51792 UG_GENE=Pp2b1A DEF=Mus musculus, protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform, clone MGC:7086 IMAGE:3157293, mRNA, complete cds. /PROD=protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform FL-gb NM_016891.1 gb BC006606.1 gb AB021743.1	1429.6333	272.46213	914.4666	261.58444	2.362404	3.0	0.099183634	0.27425778	0.64	1.56	gb BM196939 DB_XREF=gi:17748959 DB_XREF=0304006.3 CLONE=0304006.3 FEA=mRNA CNT=57 TID=Mm.220848.1 TIER=Slack STK=16 UG=Mm.220848 UG_TITLE=Slack DEF=Mus musculus, similar to putative, clone MGC:5345087, mRNA, partial cds. /PROD=Slack FL-gb NM_0207247.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	366.83334	48.94613	229.76666	88.11454	2.355309	3.0	0.09982363	0.27395207	0.63	1.6	gb NM_023631.1 DB_XREF=gi:12072471 FEA=FLmRNA CNT=119 TID=Mm.21897.1 TIER=FL+Slack STK=30 UG=Mm.21897.1 LL=11923 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	687.3	287.4756	202.13335	12.589808	2.9203472	2.0	0.09997883	0.2723304	0.29	3.4	gb BC003744.1 DB_XREF=gi:13277677 FEA=FLmRNA CNT=135 TID=Mm.3234.1 TIER=FL+Slack STK=52 UG=Mm.3234.1 LL=19060 UG_GENE=Pp2b DEF=Mus musculus, protein phosphatase 5, catalytic subunit, clone MGC:5847 IMAGE:3599302, mRNA, complete cds. /PROD=protein phosphatase 5, catalytic subunit FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	1420.2334	425.87183	649.23334	171.36757	2.9095585	2.0	0.10061329	0.27202854	0.46	2.19	gb NM_130880.1 DB_XREF=gi:18698997 GEN=Cdk9 FEA=FLmRNA CNT=119 TID=Mm.27557.1 TIER=FL+Slack STK=50 UG=Mm.27557.1 LL=107951 DEF=Mus musculus cyclin-dependent kinase 9 (CDC-related kinase) (Cdk9), mRNA. /PROD=cyclin-dependent kinase 9 FL-gb AF327431.1 gb BC003901.1 gb NM_130860.1	513.26666	108.13493	311.26666	103.22113	2.340426	3.0	0.1011825	0.271556	0.61	1.65	gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																																																																																														
gb BM196939 DB_XREF=gi:17748959 DB_XREF=0304006.3 CLONE=0304006.3 FEA=mRNA CNT=57 TID=Mm.220848.1 TIER=Slack STK=16 UG=Mm.220848 UG_TITLE=Slack DEF=Mus musculus, similar to putative, clone MGC:5345087, mRNA, partial cds. /PROD=Slack FL-gb NM_0207247.1 gb AF128192.1 gb AF128191.1 gb MS8004.1 gb BC002073.1 gb AF128190.1 gb AF128189.1 gb AF128188.1	366.83334	48.94613	229.76666	88.11454	2.355309	3.0	0.09982363	0.27395207	0.63	1.6	gb NM_023631.1 DB_XREF=gi:12072471 FEA=FLmRNA CNT=119 TID=Mm.21897.1 TIER=FL+Slack STK=30 UG=Mm.21897.1 LL=11923 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	687.3	287.4756	202.13335	12.589808	2.9203472	2.0	0.09997883	0.2723304	0.29	3.4	gb BC003744.1 DB_XREF=gi:13277677 FEA=FLmRNA CNT=135 TID=Mm.3234.1 TIER=FL+Slack STK=52 UG=Mm.3234.1 LL=19060 UG_GENE=Pp2b DEF=Mus musculus, protein phosphatase 5, catalytic subunit, clone MGC:5847 IMAGE:3599302, mRNA, complete cds. /PROD=protein phosphatase 5, catalytic subunit FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	1420.2334	425.87183	649.23334	171.36757	2.9095585	2.0	0.10061329	0.27202854	0.46	2.19	gb NM_130880.1 DB_XREF=gi:18698997 GEN=Cdk9 FEA=FLmRNA CNT=119 TID=Mm.27557.1 TIER=FL+Slack STK=50 UG=Mm.27557.1 LL=107951 DEF=Mus musculus cyclin-dependent kinase 9 (CDC-related kinase) (Cdk9), mRNA. /PROD=cyclin-dependent kinase 9 FL-gb AF327431.1 gb BC003901.1 gb NM_130860.1	513.26666	108.13493	311.26666	103.22113	2.340426	3.0	0.1011825	0.271556	0.61	1.65	gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																																																																																																									
gb NM_023631.1 DB_XREF=gi:12072471 FEA=FLmRNA CNT=119 TID=Mm.21897.1 TIER=FL+Slack STK=30 UG=Mm.21897.1 LL=11923 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	687.3	287.4756	202.13335	12.589808	2.9203472	2.0	0.09997883	0.2723304	0.29	3.4	gb BC003744.1 DB_XREF=gi:13277677 FEA=FLmRNA CNT=135 TID=Mm.3234.1 TIER=FL+Slack STK=52 UG=Mm.3234.1 LL=19060 UG_GENE=Pp2b DEF=Mus musculus, protein phosphatase 5, catalytic subunit, clone MGC:5847 IMAGE:3599302, mRNA, complete cds. /PROD=protein phosphatase 5, catalytic subunit FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	1420.2334	425.87183	649.23334	171.36757	2.9095585	2.0	0.10061329	0.27202854	0.46	2.19	gb NM_130880.1 DB_XREF=gi:18698997 GEN=Cdk9 FEA=FLmRNA CNT=119 TID=Mm.27557.1 TIER=FL+Slack STK=50 UG=Mm.27557.1 LL=107951 DEF=Mus musculus cyclin-dependent kinase 9 (CDC-related kinase) (Cdk9), mRNA. /PROD=cyclin-dependent kinase 9 FL-gb AF327431.1 gb BC003901.1 gb NM_130860.1	513.26666	108.13493	311.26666	103.22113	2.340426	3.0	0.1011825	0.271556	0.61	1.65	gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																																																																																																																				
gb BC003744.1 DB_XREF=gi:13277677 FEA=FLmRNA CNT=135 TID=Mm.3234.1 TIER=FL+Slack STK=52 UG=Mm.3234.1 LL=19060 UG_GENE=Pp2b DEF=Mus musculus, protein phosphatase 5, catalytic subunit, clone MGC:5847 IMAGE:3599302, mRNA, complete cds. /PROD=protein phosphatase 5, catalytic subunit FL-gb U25809.1 gb BC001468.1 gb BC001864.1 DB_XREF=gi:1284269 FEA=FLmRNA CNT=29 TID=Mm.23836.1 TIER=Slack STK=23 UG=Mm.23836.1 LL=70231 UG_GENE=Cn2-pending UG_TITLE=RIKEN cDNA Z1310039H17 gene (DEF=Mus musculus ES cells cDNA, RIKEN full-length enriched library, clone:2410043M02.gtg) assembly stacking protein 2, full insert sequence.	1420.2334	425.87183	649.23334	171.36757	2.9095585	2.0	0.10061329	0.27202854	0.46	2.19	gb NM_130880.1 DB_XREF=gi:18698997 GEN=Cdk9 FEA=FLmRNA CNT=119 TID=Mm.27557.1 TIER=FL+Slack STK=50 UG=Mm.27557.1 LL=107951 DEF=Mus musculus cyclin-dependent kinase 9 (CDC-related kinase) (Cdk9), mRNA. /PROD=cyclin-dependent kinase 9 FL-gb AF327431.1 gb BC003901.1 gb NM_130860.1	513.26666	108.13493	311.26666	103.22113	2.340426	3.0	0.1011825	0.271556	0.61	1.65	gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																																																																																																																															
gb NM_130880.1 DB_XREF=gi:18698997 GEN=Cdk9 FEA=FLmRNA CNT=119 TID=Mm.27557.1 TIER=FL+Slack STK=50 UG=Mm.27557.1 LL=107951 DEF=Mus musculus cyclin-dependent kinase 9 (CDC-related kinase) (Cdk9), mRNA. /PROD=cyclin-dependent kinase 9 FL-gb AF327431.1 gb BC003901.1 gb NM_130860.1	513.26666	108.13493	311.26666	103.22113	2.340426	3.0	0.1011825	0.271556	0.61	1.65	gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																																																																																																																																										
gb NM_010742.1 DB_XREF=gi:7940177 GEN=Cdc37 FEA=FLmRNA CNT=193 TID=Mm.4117.1 TIER=FL+Slack STK=68 UG=Mm.4117.1 LL=12536 UG_GENE=Cdc37 DEF=Mus musculus cyclin division cycle 37 homolog (S. cerevisiae) (Cdc37), mRNA. /PROD=cyclin division cycle 37 homolog (S. cerevisiae) FL-gb U43076.1 gb NM_016742.1	5240.6997	1181.1212	3308.3665	816.72766	2.3307126	3.0	0.10208148	0.2719689	0.63	1.58	gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																																																																																																																																																					
gb BC002148.1 DB_XREF=gi:12805356 FEA=FLmRNA CNT=59 TID=Mm.582.2 TIER=FL+Slack STK=49 UG=Mm.582.1 LL=11770 UG_GENE=fbp4 DEF=Mus musculus, clone MGC:7265 IMAGE:3484864, mRNA, complete cds. /PROD=Unknown (protein for MGC:7265) FL-gb BC002148.1	610.1333	54.303703	384.66666	124.90342	2.8673048	2.0	0.103153974	0.27283478	0.63	1.59	gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																																																																																																																																																																
gb NM_031260.1 DB_XREF=gi:13994126 FEA=Mov101 FEA=FLmRNA CNT=18 TID=Mm.102985.1 TIER=FL+Slack STK=11 UG=Mm.102985.1 LL=83456 DEF=Mus musculus Moloney leukemia virus 10-like 1 (Mov10), mRNA. /PROD=Moloney leukemia virus 10-like 1 FL-gb AF285587.1 gb AF340211.1 gb NM_031260.1	3885.3	658.83813	2512.8667	787.5243	2.3151417	3.0	0.10354287	0.27189314	0.65	1.55	gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																																																																																																																																																																											
gb NM_019652.1 DB_XREF=gi:12025541 GEN=Anaf1 FEA=FLmRNA CNT=69 TID=Mm.41475.1 TIER=FL+Slack STK=26 UG=Mm.41475.1 LL=56495 DEF=Mus musculus ara (bacterial) anisole transporter, ATP-binding, homolog 1 (Anaf1), mRNA. /PROD=ara (bacterial) anisole transporter ATP-binding, homolog 1 FL-gb AF039405.2 gb NM_019652.1 gb BC016453.1 gb BC018430.1 gb NM_133777.1 DB_XREF=gi:19527003 GEN=BT2046SF12Rk FEA=FLmRNA CNT=180 TID=Mm.22491.1 TIER=FL+Slack STK=71 UG=Mm.22491.1 LL=77891 DEF=Mus mus																																																																																																																																																																																																						

gb:AK019396.1 /DB_XREF=gi:12859591 /FEA=mRNA /CNT=69 /TID=Mm.44226.2 /TIER=Stack /STK=39 /UG=Mm.44226 /LL=68066 /UG_GENE=3010027G13R/K /UG_TITLE=RIKEN cDNA 3010027G13 gene /DEF=Mus musculus 12 days embryo head cDNA, RIKEN full-length enriched library, clone 3010027G13.homolog to HYPOTHETICAL 38.5 KDA PROTEIN, full insert sequence.	1165.7667	342.06116	664.5	142.46161	2.3431087	2.0	0.14385577	0.2599374	0.57	1.75
gb:NM_019722.1 /DB_XREF=gi:9878878 /GEN=Ariz /FEA=FLmRNA /CNT=74 /TID=Mm.21071.1 /TIER=FL+Stack /STK=39 /UG=Mm.21071 /LL=56327 /DEF=Mus musculus, Ariz-iboyastion-like 2 (Ariz), mRNA. /PROD=Ariz-iboyastion-like 2 (Ariz), cDNA, 019722.1 gb:AF143680	1832.2001	534.95233	1139.4667	293.3553	1.9666193	3.0	0.14392015	0.2587727	0.62	1.61
gb:NM_021443.1 /DB_XREF=gi:10846817 /GEN=ScybA /FEA=FLmRNA /CNT=15 /TID=Mm.42029.1 /TIER=FL+Stack /STK=7 /UG=Mm.42029 /LL=20307 /DEF=Mus musculus small inducible cytokine A8 (ScybA), mRNA. /PROD=small inducible cytokine A8 /FL=gb:AB023418.1 gb:NM_021443.1	517.3	171.56436	271.66666	59.66963	2.3422065	2.0	0.14394383	0.25754657	0.53	1.9
gb:NM_013602.1 /DB_XREF=gi:7305284 /GEN=MT /FEA=FLmRNA /CNT=145 /TID=Mm.219629.1 /TIER=FL+Stack /STK=36 /UG=Mm.219629 /LL=17448 /DEF=Mus musculus melanin-concentrating hormone 1 (M1), mRNA. /PROD=melanin-concentrating hormone 1 (M1), cDNA, 013602.1 gb:BCU26842.1 /DB_XREF=gi:20071854 /FEA=FLmRNA /CNT=89 /TID=Mm.20499.1 /TIER=FL+Stack /STK=49 /UG=Mm.20499 /DEF=Mus musculus, Similar to hypothetical protein FLJ20234, clone MGC:37525 IMAGE:4986113, mRNA, complete cds. /PROD=Similar to hypothetical protein FLJ20234 /FL=gb:BC026642.1	11612.833	3314.8948	7316.1665	1936.7217	1.9384388	3.0	0.14793773	0.26340133	0.63	1.59
gb:U43884.1 /DB_XREF=gi:1247492 /GEN=IG1B /FEA=FLmRNA /CNT=3 /TID=Mm.444.2 /TIER=FL /STK=1 /UG=Mm.444 /LL=15901 /DEF=Mus musculus transcription factor IG1B (IG1B) mRNA, complete cds. /PROD=IG1B /FL=gb:U43884.1	1394.6	364.95178	889.7	273.0372	1.9186989	3.0	0.15083033	0.26595685	0.64	1.57
gb:BC026751.1 /DB_XREF=gi:20071192 /FEA=FLmRNA /CNT=36 /TID=Mm.41486.1 /TIER=FL /STK=1 /UG=Mm.41486 /LL=70103 /UG_GENE=2700001K05R/K /DEF=Mus musculus, Similar to putative cyclin G1 interacting protein (FL=gb:BC026751.1) mRNA, complete cds. /PROD=Similar to putative cyclin G1 interacting protein (FL=gb:BC026751.1)	1355.3667	312.11258	905.0999	262.02222	1.9137493	3.0	0.15156595	0.2659691	0.67	1.5
gb:BC026751.1 /DB_XREF=gi:1933035 /FEA=FLmRNA /CNT=39 /TID=Mm.46628.2 /TIER=FL+Stack /STK=21 /UG=Mm.46628 /LL=22344 /UG_GENE=Vesf /DEF=Mus musculus, vascular endothelial zinc finger 1, clone MGC:29455 IMAGE:4159675, mRNA, complete cds. /PROD=vascular endothelial zinc finger 1 /FL=gb:BC024610.1	1260.9	340.7898	789.93335	122.7772	2.251978	2.0	0.15314065	0.26744658	0.63	1.6
gb:AK01380.1 /DB_XREF=gi:12847432 /FEA=mRNA /CNT=64 /TID=Mm.142187.1 /TIER=Stack /STK=24 /UG=Mm.142187 /LL=89902 /UG_GENE=2610009E16R/K /UG_TITLE=RIKEN cDNA 2610009E16 gene /DEF=Mus musculus 10 days embryo whole body cDNA, RIKEN full-length enriched library, clone 2610009E16.homolog to PARATHYMOSIN (ZINC-BINDING 11.5 KDA PROTEIN), full insert sequence.	3908.8	1215.2148	2384.5	673.47577	1.9002765	3.0	0.15358968	0.2669535	0.61	1.64
gb:NM_009088.1 /DB_XREF=gi:6671792 /GEN=Rpo1.4 /FEA=FLmRNA /CNT=139 /TID=Mm.7387.1 /TIER=FL+Stack /STK=36 /UG=Mm.7387 /LL=20019 /DEF=Mus musculus RNA polymerase 1-4 (194 kDa subunit) (Rpo1.4), mRNA. /PROD=RNA polymerase 1-4 (194 kDa subunit) /FL=gb:NM_009088.1 gb:AF000938.1	442.7667	120.27112	271.9	99.23125	1.8980509	3.0	0.15392701	0.26627183	0.61	1.63
gb:AK072897.1 /DB_XREF=gi:12840800 /GEN=Gapg1 /FEA=mRNA /CNT=2 /TID=Mm.2945.3 /TIER=ConsEnd /STK=1 /UG=Mm.2945 /LL=12345 /UG_TITLE=capping protein beta 1 /DEF=Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone 1700120C01.capping protein beta 1, full insert sequence.	3347.5	1118.8156	1890.4668	725.1011	1.892879	3.0	0.15471424	0.26637122	0.56	1.77
gb:NM_017476.1 /DB_XREF=gi:8567377 /GEN=Nakap95-pending /FEA=FLmRNA /CNT=87 /TID=Mm.9590.1 /TIER=FL+Stack /STK=37 /UG=Mm.9590 /LL=54194 /DEF=Mus musculus neighbor of A-kinase anchoring protein 95 (Nakap95-pending), mRNA. /PROD=neighbor of A-kinase anchoring protein 95 (Nakap95-pending) /FL=gb:AB028921.1 /FL=gb:AB028921.1	538.3333	148.90977	345.46664	20.051022	2.2232726	2.0	0.15623604	0.26772842	0.64	1.56
gb:BC007140.1 /DB_XREF=gi:13938048 /FEA=FLmRNA /CNT=129 /TID=Mm.219663.2 /TIER=FL+Stack /STK=107 /UG=Mm.219663 /LL=14114 /UG_GENE=Fbln1 /DEF=Mus musculus, Similar to fibulin 1, clone MGC:6128 IMAGE:3495754, mRNA, complete cds. /PROD=Similar to fibulin 1 /FL=gb:BC007140.1	2261.4333	914.7857	1040.7667	275.78848	2.2128296	2.0	0.15738344	0.26843438	0.46	2.17
gb:BE84258 /DB_XREF=gi:10856305 /DB_XREF=U01Mm.C0501bgd-c-01-U-31 /CLONE=U01Mm.C0501bgd-c-01-U /FEA=FLmRNA /CNT=16 /TID=Mm.22543.1 /TIER=Stack /STK=15 /UG=Mm.22543 /UG_TITLE=Mus musculus brain and reproductive organ-expressed protein (BrE) mRNA, complete cds /FL=gb:AF40752.1	915.3333	239.33093	607.63336	154.42604	1.8711394	3.0	0.15807502	0.2683559	0.66	1.51
gb:NM_010312.1 /DB_XREF=gi:13937390 /GEN=Gnb2 /FEA=FLmRNA /CNT=367 /TID=Mm.30141.1 /TIER=FL+Stack /STK=141 /UG=Mm.30141 /LL=14693 /DEF=Mus musculus guanine nucleotide binding protein, beta 2 (Gnb2), mRNA. /PROD=guanine nucleotide-binding protein, beta-2subunit /FL=gb:U54560.1 gb:NM_010312.1	1156.7001	482.16302	477.7333	227.77388	2.2053285	2.0	0.15821478	0.26735365	0.41	2.42
gb:NM_011707.1 /DB_XREF=gi:13933353 /FEA=FLmRNA /CNT=47 /TID=Mm.34556.1 /TIER=FL+Stack /STK=16 /UG=Mm.34556 /LL=23831 /DEF=Mus musculus carbonic anhydrase 14 (Car14), mRNA. /PROD=carbonic anhydrase 14 /FL=gb:NM_011707.1 gb:AB005450.1	1273.2001	392.50107	750.2666	284.58118	1.8682411	3.0	0.15852948	0.26665094	0.59	1.7
gb:AK040893.1 /DB_XREF=gi:12835132 /FEA=mRNA /CNT=542 /TID=Mm.29458.1 /TIER=Stack /STK=17 /UG=Mm.29458 /LL=68746 /UG_GENE=1110034G11R/K /UG_TITLE=RIKEN cDNA 1110034G11 gene /DEF=Mus musculus 18 days embryo whole body cDNA, RIKEN full-length enriched library, clone 1110034G11 related to CG12379 PROTEIN, full insert sequence.	2279.8	659.10315	1438.1333	63.327145	2.2016687	2.0	0.15862256	0.26558363	0.63	1.59
gb:U18321.1 /DB_XREF=gi:1185384 /FEA=FLmRNA /CNT=80 /TID=Mm.4269.2 /TIER=FL+Stack /STK=55 /UG=Mm.4269 /LL=21413 /UG_GENE=Tcf4 /DEF=Mus musculus basic transcription factor MITF-2A mRNA, complete cds. /PROD=MITF-2A protein	1709.9332	404.62988	1114.2999	333.23102	1.8674502	3.0	0.15865374	0.2644229	0.65	1.53
gb:AK00892.1 /DB_XREF=gi:12842855 /FEA=mRNA /CNT=284 /TID=Mm.35719.1 /TIER=Stack /STK=41 /UG=Mm.35719 /LL=11552 /UG_GENE=1300004C11R/K /UG_TITLE=RIKEN cDNA 1300004C11 gene /DEF=Mus musculus adult male small intestine cDNA, RIKEN full-length enriched library, clone 2010319A12.homolog to THFIC2 SUBUNIT, full insert sequence.	595.0	171.37785	374.63333	27.457663	2.1991158	2.0	0.15890788	0.2636426	0.63	1.59
gb:AF252873.1 /DB_XREF=gi:7644395 /FEA=FLmRNA /CNT=73 /TID=Mm.30211.1 /TIER=FL /STK=5 /UG=Mm.30211 /LL=57286 /UG_GENE=Scyb14 /DEF=Mus musculus CXC chemokine MIP-2gamm precursor, mRNA, complete cds. /PROD=CXC chemokine MIP-2gamm precursor /FL=gb:AF192557.1 gb:NM_010568.1 gb:AF144754.1 gb:AF152377.1 gb:AF252873.1	540.5	160.72278	317.30002	72.58843	2.1921403	2.0	0.15969104	0.2637431	0.59	1.7
gb:AF332085.1 /DB_XREF=gi:14193736 /GEN=Rad21 /FEA=FLmRNA /CNT=282 /TID=Mm.182628.1 /TIER=FL+Stack /STK=2 /UG=Mm.182628 /LL=19357 /DEF=Mus musculus strain ISS Rad21 (Rad21) mRNA, complete cds. /PROD=Rad21 /FL=gb:AF332086.1 gb:AF332085.1	823.76666	335.08176	380.09998	241.96138	1.8592664	3.0	0.15994632	0.2629748	0.46	2.17
gb:NM_009009.1 gb:D49429.1										
gb:BC004826.1 /DB_XREF=gi:13435955 /FEA=FLmRNA /CNT=206 /TID=Mm.2079.1 /TIER=FL+Stack /STK=57 /UG=Mm.2079 /LL=30273 /UG_GENE=Syny2 /DEF=Mus musculus synaptogyrin 2, clone MGC:6483 IMAGE:2946657, mRNA, complete cds. /PROD=synaptogyrin 2 /FL=gb:AF394049.1 gb:BC004829.1 gb:AF151985.1 gb:NM_009304.1	1586.0668	384.5148	1058.5	310.58392	1.8486891	3.0	0.1616351	0.2645597	0.67	1.5
gb:AK007572.1 /DB_XREF=gi:12841200 /FEA=mRNA /CNT=171 /TID=Mm.161667.1 /TIER=ConsEnd /STK=0 /UG=Mm.161667 /LL=89115 /UG_GENE=1810021D01R/K /UG_TITLE=RIKEN cDNA 1810021D01 gene /DEF=Mus musculus 10 day old day hatched cDNA, RIKEN full-length enriched library, clone 1810021D01.homolog to CDNA FLJ10479 FIS, CLONE NT2RP200120 (DC1) HYPOTHETICAL 39.8 KDA PROTEIN), full insert sequence.	941.5667	57.336903	622.8334	247.26883	2.1749341	2.0	0.16164558	0.2633957	0.66	1.51
gb:NM_026465.1 /DB_XREF=gi:13385959 /GEN=Atpgs2 /FEA=FLmRNA /CNT=334 /TID=Mm.10314.1 /TIER=FL+Stack /STK=154 /UG=Mm.10314 /LL=67942 /DEF=Mus musculus ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2 (Atpgs2), mRNA. /PROD=ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2 /FL=gb:NM_026468.1 gb:BC068913.1	13885.966	3905.741	8379.966	1993.4896	2.1748064	3.0	0.16166022	0.2622488	0.6	1.66
gb:BC019442.1 /DB_XREF=gi:18043191 /FEA=FLmRNA /CNT=127 /TID=Mm.185518.1 /TIER=FL+Stack /STK=87 /UG=Mm.185518 /DEF=Mus musculus, clone MGC:25803 IMAGE:4036849, mRNA, complete cds. /PROD=Unknown (protein for MGC:25803) /FL=gb:BC014795.1 gb:BC019442.1	772.5	196.32268	505.13333	85.66692	2.1619692	2.0	0.16314003	0.26347837	0.65	1.53
gb:NM_006713.1 /DB_XREF=gi:6079069 /GEN=Nos3 /FEA=FLmRNA /CNT=49 /TID=Mm.12637.1 /TIER=FL+Stack /STK=25 /UG=Mm.12637 /UG_GENE=181017 /DEF=Mus musculus nitric oxide synthase 3, endothelial cell (Nos3), mRNA. /PROD=nitric oxide synthase 3, endothelial cell /FL=gb:U53142.1 gb:NM_006713.1	1535.2333	382.03305	1020.69995	303.48578	1.8265747	3.0	0.16523309	0.26568317	0.66	1.5
gb:AC040695.1 /DB_XREF=gi:12836055 /FEA=mRNA /CNT=184 /TID=Mm.34763.1 /TIER=Stack /STK=111 /UG=Mm.34763 /LL=71732 /UG_GENE=1200011A11R/K /UG_TITLE=RIKEN cDNA 1200011A11 gene /DEF=Mus musculus adult male lung cDNA, RIKEN full-length enriched library, clone 1200011A11.homolog to VACUOLAR PROTEIN SORTING PROTEIN 11, full insert sequence.	578.6333	199.9345	324.93332	53.362926	2.123492	2.0	0.1676879	0.26844773	0.56	1.78
gb:BC019836.1 /DB_XREF=gi:18043722 /FEA=FLmRNA /CNT=420 /TID=Mm.22248.2 /TIER=FL+Stack /STK=125 /UG=Mm.22248 /LL=16010 /UG_GENE=Igf2bp4 /DEF=Mus musculus, insulin-like growth factor binding protein 4, clone MGC:29917 IMAGE:5123738, mRNA, complete cds. /PROD=insulin-like growth factor binding protein 4 /FL=gb:BC019836.1	5639.267	1676.5708	3555.6335	1113.3978	1.7931857	3.0	0.17084232	0.27230328	0.63	1.59
gb:BC016699.1 /DB_XREF=gi:12836055 /FEA=mRNA /CNT=207 /TID=Mm.28492.1 /TIER=FL+Stack /STK=110 /UG=Mm.28492 /DEF=Mus musculus, clone MGC:6816 IMAGE:2648707, mRNA, complete cds. /PROD=Unknown (protein for MGC:6816) /FL=gb:BC016699.1	860.1333	288.7083	484.16666	115.139244	2.0950768	2.0	0.17115806	0.27162042	0.56	1.78
gb:BC020255.1 /DB_XREF=gi:12805192 /FEA=FLmRNA /CNT=194 /TID=Mm.1506.1 /TIER=FL+Stack /STK=73 /UG=Mm.1508 /LL=12257 /UG_GENE=Grpr /DEF=Mus musculus, Similar to benzodiazepine receptor, peripheral, clone MGC:6086 IMAGE:3493196, mRNA, complete cds. /PROD=Similar to benzodiazepine receptor, peripheral /FL=gb:NM_009775.1 gb:L17306.1 gb:BC020255.1 gb:D21207.1	1181.9333	402.33136	650.36664	178.55083	2.091686	2.0	0.17157866	0.27110913	0.55	1.82
gb:NM_008281.2 /DB_XREF=gi:7949140 /GEN=Zfp143 /FEA=FLmRNA /CNT=51 /TID=Mm.10815.1 /TIER=FL+Stack /STK=29 /UG=Mm.10815 /LL=20841 /DEF=Mus musculus zinc finger protein 143 (Zfp143), mRNA. /PROD=zinc finger protein 143	352.66666	124.71	200.73334	18.59068	2.087083	2.0	0.17215186	0.27084237	0.57	1.76
gb:AF011758.2 gb:NM_009281.2										
gb:NM_009612.1 /DB_XREF=gi:1012066 /GEN=Fxyd5 /FEA=FLmRNA /CNT=58 /TID=Mm.1870.1 /TIER=FL+Stack /STK=26 /UG=Mm.1870 /LL=18301 /DEF=Mus musculus FXYD domain-containing ion transport regulator 5 (Fxyd5), mRNA. /PROD=related to ion channel /FL=gb:U72680.1 gb:NM_008761.2	1579.3999	339.63223	955.8667	502.16968	1.7814642	3.0	0.17286329	0.27079442	0.61	1.65
gb:NM_008940.1 /DB_XREF=gi:6679486 /GEN=Klk6 /FEA=FLmRNA /CNT=11 /TID=Mm.5193.2 /TIER=FL /STK=7 /UG=Mm.5193 /LL=16624 /DEF=Mus musculus kallikrein 6 (Klk6), mRNA. /PROD=protease, serine, 19 (neuropilin) /FL=gb:NM_008940.1	514.4666	147.93588	316.46667	73.88783	2.0739162	2.0	0.173806	0.27110764	0.62	1.63
gb:NM_008592.1 /DB_XREF=gi:6679834 /GEN=Foxc1 /FEA=FLmRNA /CNT=61 /TID=Mm.12948.1 /TIER=FL+Stack /STK=9 /UG=Mm.12949 /LL=17300 /DEF=Mus musculus forkhead box C1 (Foxc1), mRNA. /PROD=forkhead box C1 /FL=gb:AF045017.1	284.06668	94.30697	169.09999	61.060375	1.7724156	3.0	0.17444223	0.27094218	0.6	1.68
gb:NM_008592.1										
gb:NM_022602.1 /DB_XREF=gi:12863570 /GEN=Ndub2 /FEA=FLmRNA /CNT=13 /TID=Mm.29513.1 /TIER=FL /STK=0 /UG=Mm.29513 /LL=66416 /DEF=Mus musculus NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (14.5KD, B14.5E) (Ndub2), mRNA. /PROD=NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (14.5KD, B14.5E) /FL=gb:NM_023202.1	12473.101	4055.233	7557.9	2600.8333	1.7671416	3.0	0.17537016	0.27122927	0.61	1.65
gb:NM_007472.1 /DB_XREF=gi:6680709 /GEN=Aqp1 /FEA=FLmRNA /CNT=282 /TID=Mm.18825.1 /TIER=FL+Stack /STK=113 /UG=Mm.18825 /LL=11826 /DEF=Mus musculus aquaporin 1 (Aqp1), mRNA. /PROD=aquaporin 1 /FL=gb:NM_007472.1 gb:BC007125.1 gb:U02914.1	9189.899	2563.1602	5969.3003	1845.5371	1.7661334	3.0	0.17554818	0.270359	0.65	1.54
gb:NM_007632.1 /DB_XREF=gi:8724312 /GEN=Ccn3 /FEA=FLmRNA /CNT=286 /TID=Mm.7417.1 /TIER=FL+Stack /STK=172 /UG=Mm.7417 /LL=12445 /DEF=Mus musculus cyclin D3 (Ccn3), mRNA. /PROD=cyclin D3 /FL=gb:BC004076.1 gb:NM_007632.1 gb:M86183.1 gb:BC005605.1	1193.2001	406.31543	681.76666	146.32567	2.051192	2.0	0.17671187	0.27100772	0.57	1.75
gb:AK074262.1 /DB_XREF=gi:12851922 /FEA=mRNA /CNT=54 /TID=Mm.45020.1 /TIER=Stack /STK=47 /UG=Mm.45020 /LL=73224 /UG_GENE=3110080A02R/K /UG_TITLE=RIKEN cDNA 3110080A02 gene /DEF=Mus musculus 13 days embryo head cDNA, RIKEN full-length enriched library, clone 3110080A02.unclassified transcript, full insert sequence.	483.83334	61.59841	314.56668	129.55139	2.0437663	2.0	0.17767572	0.27134576	0.65	1.54
gb:NM_080837.1 /DB_XREF=gi:18250287 /GEN=Ii25 /FEA=FLmRNA /CNT=150 /TID=Mm.29925.1 /TIER=FL+Stack /STK=73 /UG=Mm.29925 /LL=28106 /DEF=Mus musculus interleukin 25 (Ii25), mRNA. /PROD=interleukin 25 /FL=gb:AY038184.1 gb:NM_080837.1	531.9666	172.19682	323.43335	40.16644	2.0427072	2.0	0.17781377	0.2704251	0.61	1.64
gb:NM_025691.1 /DB_XREF=gi:13359595 /GEN=Mgsl3 /FEA=FLmRNA /CNT=210 /TID=Mm.28823.1 /TIER=FL+Stack /STK=96 /UG=Mm.28823 /LL=66447 /DEF=Mus musculus microsomal glutathione S-transferase 3 (Mgsl3), mRNA. /PROD=microsomal glutathione S-transferase 3 /FL=gb:NM_025691.1	5083.5	1483.3075	3071.2334	1323.4027	1.7533147	3.0	0.17782989	0.26932743	0.6	1.66
gb:NM_007932.1 /DB_XREF=gi:6679648 /GEN=Eng /FEA=FLmRNA /CNT=113 /TID=Mm.4851.1 /TIER=FL+Stack /STK=50 /UG=Mm.4851 /LL=13605 /DEF=Mus musculus endothelial growth factor (Eng), mRNA. /PROD=englerin /FL=gb:NM_007932.1	2786.6335	1294.6776	1264.8334	141.4751	2.0238533	2.0	0.18029575	0.27193367	0.45	2.2
gb:NM_008898.1 /DB_XREF=gi:6679420 /GEN=Por /FEA=FLmRNA /CNT=175 /TID=Mm.3883.1 /TIER=FL+Stack /STK=63 /UG=Mm.3883 /LL=18984 /DEF=Mus musculus P450 (cytochrome) oxidoreductase (Por), mRNA. /PROD=P450 (cytochrome) oxidoreductase /FL=gb:D15751.1 gb:NM_008898.1	721.4001	186.83733	446.43335	200.67395	1.7369798	3.0	0.18078685	0.27155226	0.62	1.62
gb:NM_008322.1 /DB_XREF=gi:6680342 /GEN=Idh2 /FEA=FLmRNA /CNT=70 /TID=Mm.2966.1 /TIER=FL /STK=2 /UG=Mm.2966 /LL=18227 /DEF=Mus musculus isocitrate dehydrogenase 2 (NADP+), mitochondrial (Idh2), mRNA. /PROD=isocitrate dehydrogenase 2 (NADP+) /FL=gb:NM_008322.1 gb:U15167.1	11169.633	2583.2095	7453.2993	2658.36	1.736538	3.0	0.1808676	0.27056015	0.67	1.5
gb:NM_1										

[illegible]

[illegible]

gb:LA2114.1 /DB_XREF=gi:18654373 /GEN=GDF10 /FEA=FLmRNA /CNT=85 /TID=Mm.40323.1 /TIER=FL+Stack /STK=85 /UG=Mm.40323 /DEF=Mus musculus growth/differentiation factor 10 (GDF10) mRNA, complete cds. /PROD=growth/differentiation factor 10 /FL=gb:LA2114.1	339.76666	192.45802	215.83333	47.5599	1.0827825	2.0	0.39208058	0.40771914	0.64	1.57
gb:NM_007657.1 /DB_XREF=gi:6680893 /GEN=Cd9 /FEA=FLmRNA /CNT=324 /TID=Mm.2956.1 /TIER=FL+Stack /STK=135 /UG=Mm.2956 /LL=12527 /DEF=Mus musculus CD9 antigen (Cd9), mRNA. /PROD=CD9 antigen /FL=gb:L08115.1 gb:NM_007657.1	2590.3333	1327.7048	1686.2667	684.79297	1.0481877	2.0	0.40454426	0.41948482	0.65	1.54
gb:AK004342.1 /DB_XREF=gi:12835491 /FEA=FLmRNA /CNT=37 /TID=Mm.178550.2 /TIER=Stack /STK=1919 /UG=Mm.178550 /LL=66478 /UG_GENE=2410020K10Rik /UG_TITLE=RIKEN cDNA 2410020K10 gene /DEF=Mus musculus 18 days embryo whole body cDNA, RIKEN full-length enriched library, clone:110061M03:homolog to T-CELL SURFACE GLYCOPROTEIN E2 PRECURSOR (E2 ANTIGEN) (CD99) (MC2 PROTEIN) (I2E7), full insert sequence.	849.5334	656.2247	453.93335	76.852875	1.0370659	2.0	0.40864527	0.4225369	0.53	1.87
gb:BC010564.1 /DB_XREF=gi:14714826 /FEA=FLmRNA /CNT=83 /TID=Mm.219659.1 /TIER=FL+Stack /STK=24 /UG=Mm.219659 /LL=15267 /UG_GENE=Hist2 /DEF=Mus musculus, H2A histone family, member O, clone MGC:5966 IMAGE:3582122, mRNA, complete cds. /PROD=H2A histone family, member O /FL=gb:NM_013549.1 gb:BC010564.1	1537.0668	912.65375	992.8333	134.19629	1.0218685	2.0	0.41432413	0.42719862	0.65	1.55
gb:BC019459.1 /DB_XREF=gi:15930030 /FEA=FLmRNA /CNT=83 /TID=Mm.18814.1 /TIER=FL+Stack /STK=10 /UG=Mm.18814 /LL=66898 /UG_GENE=1300006M19Rik /DEF=Mus musculus, RIKEN cDNA 1300006M19 gene, clone MGC:8269 IMAGE:3592632, mRNA, complete cds. /PROD=RIKEN cDNA 1300006M19 gene /FL=gb:BC015459.1	992.26666	1056.4161	371.96667	48.13329	1.015961	2.0	0.41655517	0.42828912	0.37	2.67
gb:NM_010890.1 /DB_XREF=gi:13964148 /GEN=Nedd4a /FEA=FLmRNA /CNT=23 /TID=Mm.18553.1 /TIER=FL /STK=1 /UG=Mm.18553 /LL=17599 /DEF=Mus musculus neural precursor cell expressed, developmentally down-regulated gene 4a (Nedd4a), mRNA. /PROD=neural precursor cell expressed, developmentally down-regulated gene 4a /FL=gb:NM_010890.1	1768.7333	992.5988	1140.6	479.7904	0.98683304	2.0	0.42775115	0.4385651	0.64	1.55
gb:BC019775.1 /DB_XREF=gi:18043802 /FEA=FLmRNA /CNT=156 /TID=Mm.214514.1 /TIER=FL+Stack /STK=98 /UG=Mm.214514 /LL=116847 /UG_GENE=Prep /DEF=Mus musculus, proline arginine-rich end leucine-rich repeat, clone MGC:25825 IMAGE:4165772, mRNA, complete cds. /PROD=proline arginine-rich end leucine-rich repeat /FL=gb:BC019775.1 gb:NM_054077.1	3675.5667	1965.7422	2444.2334	1320.7708	0.90055424	3.0	0.4341985	0.44392845	0.66	1.5
gb:AK000203.1 /DB_XREF=gi:12842246 /FEA=FLmRNA /CNT=83 /TID=Mm.18814.1 /TIER=Stack /STK=25 /UG=Mm.18814 /LL=66898 /UG_GENE=1300006M19Rik /UG_TITLE=RIKEN cDNA 1300006M19 gene /DEF=Mus musculus adult male small intestine cDNA, RIKEN full-length enriched library, clone:2010012D17:homolog to INSULIN RECEPTOR TYROSINE KINASE SUBSTRATE, full insert sequence. /FL=gb:BC015459.1	2052.7	2641.6304	592.10004	181.3516	0.95542985	2.0	0.4401908	0.44879788	0.29	3.47
gb:NM_013550.1 /DB_XREF=gi:7305140 /GEN=Hist4 /FEA=FLmRNA /CNT=9 /TID=Mm.193557.1 /TIER=FL /STK=1 /UG=Mm.193557 /LL=15269 /DEF=Mus musculus histone 4 protein (Hist4), mRNA. /PROD=histone 4 protein /FL=gb:NM_013550.1	2056.1667	1616.3438	1194.2001	14.616449	0.9236334	2.0	0.45318398	0.4607581	0.58	1.72
gb:AB010742.1 /DB_XREF=gi:3062840 /FEA=FLmRNA /CNT=27 /TID=Mm.6254.1 /TIER=FL /STK=3 /UG=Mm.6254 /LL=22229 /UG_GENE=Ucp3 /DEF=Mus musculus mRNA for UCP3, complete cds. /PROD=UCP3 /FL=gb:AB010742.1 gb:AF030164.1	529.93335	380.70724	337.53336	105.21071	0.84371	2.0	0.4876572	0.4944302	0.64	1.57
gb:NM_009464.1 gb:AB008216.1 gb:AF053352.1 gb:AF032902.1										
gb:NM_017370.1 /DB_XREF=gi:3850218 /GEN=Hp /FEA=FLmRNA /CNT=154 /TID=Mm.26730.1 /TIER=FL+Stack /STK=33 /UG=Mm.26730 /LL=15439 /DEF=Mus musculus haptoglobin (Hp), mRNA. /PROD=haptoglobin /FL=gb:NM_017370.1 gb:M66827.1	705.19995	506.99698	440.69998	253.31256	0.8083316	2.0	0.5037641	0.50934595	0.62	1.6
gb:NM_010180.1 /DB_XREF=gi:6753821 /GEN=Fbinn1 /FEA=FLmRNA /CNT=91 /TID=Mm.219663.1 /TIER=FL+Stack /STK=41 /UG=Mm.219663 /LL=14114 /DEF=Mus musculus fibulin 1 (Fbinn1), mRNA. /PROD=fibulin 1 /FL=gb:NM_010180.1	835.6333	728.61206	505.8666	168.10336	0.76385224	2.0	0.52476585	0.5291147	0.61	1.65
gb:NM_009052.1 /DB_XREF=gi:6677720 /GEN=Reox3 /FEA=FLmRNA /CNT=77 /TID=Mm.14768.1 /TIER=FL+Stack /STK=56 /UG=Mm.14768 /LL=19716 /DEF=Mus musculus reduced expression 3 (Reo3), mRNA. /PROD=reduced expression 3 /FL=gb:AF051347.1	1297.2999	1452.1323	657.06665	84.7658	0.76234937	2.0	0.5254902	0.52838546	0.51	1.97
gb:NM_009052.1 gb:AF097438.1										
gb:AK004564.1 /DB_XREF=gi:12835818 /FEA=mRNA /CNT=74 /TID=Mm.46346.1 /TIER=Stack /STK=30 /UG=Mm.46346 /LL=66866 /UG_GENE=1200003G01Rik /UG_TITLE=RIKEN cDNA 1200003G01 gene /DEF=Mus musculus adult male lung cDNA, RIKEN full-length enriched library, clone:1200003G01:NHL repeat containing protein, full insert sequence.	495.19998	386.70343	330.69998	64.495186	0.72675955	2.0	0.5429257	0.54441726	0.67	1.5
gb:NM_050037.1 /DB_XREF=gi:16850086 /GEN=Scgb3a1 /FEA=FLmRNA /CNT=13 /TID=Mm.22802.1 /TIER=FL /STK=7 /UG=Mm.22802 /LL=68662 /DEF=Mus musculus secretoglobin, family 3A, member 1 (Scgb3a1), mRNA. /PROD=UGRP type A /FL=gb:NM_054037.1 gb:AF313456.1	6362.733	6998.1064	3526.5332	5637.334	0.5466609	3.0	0.62264764	0.62264764	0.55	1.8