0.030979	9.3	300.2	309.5	gb:NM_134256.1 /DB_XREF=gi:19527421 /GEN=mOAT6-L /FEA=FLmRNA /CNT=20 /TID=Mm.187584.1 /TIER=FL /STK=6 /UG=Mm.187584 /LL=171405 /DEF=Mus musculus organic anion transporter 6 (mOAT6-L), mRNA. /PROD=organic anion transporter 6 /FL=gb:NM_134256.1 gb:AB056442.1
0.0309	8	258.9	266.9	gb:AV234027 /DB_XREF=gi:16386106 /DB_XREF=AV234027 /CLONE=4632428K04 /FEA=EST /CNT=12 /TID=Mm.100306.1 /TIER=ConsEnd /STK=7 /UG=Mm.100306 /UG_TITLE=ESTs
0.030696	7.1	231.3	238.4	gb:BC006817.1 /DB_XREF=gi:14919421 /FEA=mRNA /CNT=14 /TID=Mm.402.1 /TIER=ConsEnd /STK=6 /UG=Mm.402 /DEF=Mus musculus, clone IMAGE:3156149, mRNA, partial cds. /PROD=Unknown (protein for IMAGE:3156149)
0.030387	8.8	289.6	298.4	gb:BG072280 /DB_XREF=gi:12554849 /DB_XREF=H3108G08-3 /CLONE=H3108G08 /FEA=EST /CNT=3 /TID=Mm.17642.1 /TIER=ConsEnd /STK=2 /UG=Mm.17642 /UG_TITLE=ESTs
0.030303	7.1	234.3	241.4	gb:BG074597 /DB_XREF=gi:12557166 /DB_XREF=H3137A08-3 /CLONE=H3137A08 /FEA=EST /CNT=3 /TID=Mm.197271.1 /TIER=ConsEnd /STK=2 /UG=Mm.197271 /UG_TITLE=ESTs
0.030019	6.4	213.2	219.6	gb:AK015427.1 /DB_XREF=gi:12853764 /FEA=mRNA /CNT=1 /TID=Mm.159289.1 /TIER=ConsEnd /STK=0 /UG=Mm.159289 /LL=74672 /UG_GENE=4930449A18Rik /UG_TITLE=RIKEN cDNA 4930449A18 gene /DEF=Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clo
0.029555	10.9	368.8	379.7	gb:AV258795 /DB_XREF=gi:6246254 /DB_XREF=AV258795 /CLONE=4930401P20 /FEA=mRNA /CNT=4 /TID=Mm.159963.1 /TIER=ConsEnd /STK=0 /UG=Mm.159963 /LL=77633 /UG_GENE=4930594C11Rik /UG_TITLE=RIKEN cDNA 4930594C11 gene
0.029378	10.1	343.8	353.9	gb:BB180412 /DB_XREF=gi:15409393 /DB_XREF=BB180412 /CLONE=A230084J22 /FEA=EST /CNT=67 /TID=Mm.121498.1 /TIER=Stack /STK=65 /UG=Mm.121498 /LL=101167 /UG_GENE=AI841794 /UG_TITLE=expressed sequence AI841794
0.029272	13.6	464.6	478.2	gb:NM_016768.1 /DB_XREF=gi:7949104 /GEN=Pbx3 /FEA=FLmRNA /CNT=26 /TID=Mm.7331.1 /TIER=FL /STK=2 /UG=Mm.7331 /LL=18516 /DEF=Mus musculus pre B-cell leukemia transcription factor 3 (Pbx3), mRNA. /PROD=pre B-cell leukemia transcription factor 3 /FL=gb:NM_016768.1 gb:AF020199.1
0.029228	7.5	256.6	264.1	gb:AK015242.1 /DB_XREF=gi:12853501 /FEA=mRNA /CNT=1 /TID=Mm.159059.1 /TIER=ConsEnd /STK=0 /UG=Mm.159059 /LL=73786 /UG_GENE=4930429L21Rik /UG_TITLE=RIKEN cDNA 4930429L21 gene /DEF=Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clo
0.028947	8.8	304	312.8	gb:AV020412 /DB_XREF=gi:4797404 /DB_XREF=AV020412 /CLONE=1190017H02 /FEA=EST /CNT=10 /TID=Mm.35390.1 /TIER=Stack /STK=9 /UG=Mm.35390 /UG_TITLE=ESTs
0.028407	22.2	781.5	803.7	gb:AI647939 /DB_XREF=gi:4726617 /DB_XREF=uk38a05.x1 /CLONE=IMAGE:1971248 /FEA=EST /CNT=45 /TID=Mm.25773.1 /TIER=Stack /STK=31 /UG=Mm.25773 /UG_TITLE=ESTs, Weakly similar to MOT2 MOUSE MONOCARBOXYLATE TRANSPORTER 2 (M.musculus)
0.027852	12.5	448.8	461.3	gb:BC002065.1 /DB_XREF=gi:12805212 /FEA=FLmRNA /CNT=42 /TID=Mm.15085.1 /TIER=FL+Stack /STK=11 /UG=Mm.15085 /LL=20715 /UG_GENE=Spi2-1 /DEF=Mus musculus, clone MGC:6156 IMAGE:3484201, mRNA, complete cds. /PROD=Unknown (protein for MGC:6156) /FL=gb:BC002065.
0.027391	6.3	230	236.3	gb:BM223267 /DB_XREF=gi:17784175 /DB_XREF=k0133G01-3 /CLONE=k0133G01 /FEA=EST /CNT=3 /TID=Mm.217939.1 /TIER=ConsEnd /STK=2 /UG=Mm.217939 /UG_TITLE=ESTs
0.027214	10.6	389.5	400.1	gb:AI327383 /DB_XREF=gi:4061812 /DB_XREF=mp96c10.x1 /CLONE=IMAGE:577074 /FEA=EST /CNT=16 /TID=Mm.28494.4 /TIER=Stack /STK=12 /UG=Mm.28494 /LL=66419 /UG_GENE=Mrpl11 /UG_TITLE=mitochondrial ribosomal protein L11
0.027203	13.4	492.6	506	gb:BC026925.1 /DB_XREF=gi:20070837 /FEA=FLmRNA /CNT=234 /TID=Mm.16228.1 /TIER=FL+Stack /STK=24 /UG=Mm.16228 /LL=11739 /UG_GENE=Slc25a4 /DEF=Mus musculus, clone MGC:25083 IMAGE:4500784, mRNA, complete cds. /PROD=Unknown (protein for MGC:25083) /FL=gb:BC026
0.027087	8.5	313.8	322.3	gb:BF451327 /DB_XREF=gi:11517496 /DB_XREF=uz68h03.x1 /CLONE=IMAGE:3674261 /FEA=EST /CNT=2 /TID=Mm.54213.1 /TIER=ConsEnd /STK=2 /UG=Mm.54213 /UG_TITLE=ESTs
0.026945	17.8	660.6	678.4	gb:NM_023135.1 /DB_XREF=gi:12963514 /GEN=Ste /FEA=FLmRNA /CNT=8 /TID=Mm.89655.1 /TIER=FL /STK=5 /UG=Mm.89655 /LL=20860 /DEF=Mus musculus sulfotransferase, estrogen preferring (Ste), mRNA. /PROD=sulfotransferase, estrogen preferring /FL=gb:NM_023135.1
0.026543	8	301.4	309.4	gb:AW488042 /DB_XREF=gi:7058312 /DB_XREF=UI-M-BH3-arr-a-08-0-UI.s1 /CLONE=UI-M-BH3-arr-a-08-0-UI /FEA=EST /CNT=4 /TID=Mm.101379.1 /TIER=ConsEnd /STK=3 /UG=Mm.101379 /UG_TITLE=ESTs
0.026512	8.9	335.7	344.6	gb:NM_007734.1 /DB_XREF=gi:6680967 /GEN=Col4a3 /FEA=FLmRNA /CNT=15 /TID=Mm.8069.1 /TIER=FL /STK=2 /UG=Mm.8069 /LL=12828 /DEF=Mus musculus procollagen, type IV, alpha 3 (Col4a3), mRNA. /PROD=procollagen, type IV, alpha 3 /FL=gb:NM_007734.1 gb:AF169387.1
0.026277	5.3	201.7	207	gb:AV260432 /DB_XREF=gi:6247891 /DB_XREF=AV260432 /CLONE=4930413D15 /FEA=EST /CNT=9 /TID=Mm.87305.1 /TIER=ConsEnd /STK=7 /UG=Mm.87305 /UG_TITLE=ESTs
0.026199	11.2	427.5	438.7	gb:NM_033652.1 /DB_XREF=gi:17298669 /GEN=Lmx1a /FEA=FLmRNA /CNT=4 /TID=Mm.103799.1 /TIER=FL /STK=2 /UG=Mm.103799 /LL=110648 /DEF=Mus musculus LIM homeobox transcription factor 1 alpha (Lmx1a), mRNA. /PROD=LIM homeobox transcription factor 1 alpha /FL=gb:NM_033652.1 gb:AF226662.1
0.026142	9.9	378.7	388.6	gb:NM_009917.1 /DB_XREF=gi:6753467 /GEN=Cmkbr5 /FEA=FLmRNA /CNT=2 /TID=Mm.14302.1 /TIER=FL /STK=1 /UG=Mm.14302 /LL=12774 /DEF=Mus musculus chemokine (C-C) receptor 5 (Cmkbr5), mRNA. /PROD=chemokine (C-C) receptor 5 /FL=gb:NM_009917.1
0.026104	10.7	409.9	420.6	gb:NM_013650.1 /DB_XREF=gi:7305452 /GEN=S100a8 /FEA=FLmRNA /CNT=18 /TID=Mm.21567.1 /TIER=FL+Stack /STK=10 /UG=Mm.21567 /LL=20201 /DEF=Mus musculus S100 calcium binding protein A8 (calgranulin A) (S100a8), mRNA. /PROD=S100 calcium binding protein A8 (calgranulin A) /FL=gb:M83218.1 gb:NM_013650.1
0.025948	5.2	200.4	205.6	gb:BB306677 /DB_XREF=gi:16402231 /DB_XREF=BB306677 /CLONE=B230209B12 /FEA=EST /CNT=3 /TID=Mm.131623.1 /TIER=ConsEnd /STK=3 /UG=Mm.131623 /UG_TITLE=ESTs
0.025614	7.4	288.9	296.3	gb:BB028429 /DB_XREF=gi:15403532 /DB_XREF=BB028429 /CLONE=5430403O20 /FEA=EST /CNT=6 /TID=Mm.32742.1 /TIER=ConsEnd /STK=3 /UG=Mm.32742 /LL=103745 /UG_GENE=AI464234 /UG_TITLE=expressed sequence AI464234
0.02548	175.8	6899.5	7075.3	gb:NM_010000.1 /DB_XREF=gi:6753577 /GEN=Cyp2b9 /FEA=FLmRNA /CNT=28 /TID=Mm.876.1 /TIER=FL+Stack /STK=8 /UG=Mm.876 /LL=13094 /DEF=Mus musculus cytochrome P450, 2b9, phenobarbitol inducible, type a (Cyp2b9), mRNA. /PROD=cytochrome P450, 2b9, phenobarbitol inducible,type a /FL=gb:NM_010000.1 gb:M21855.1
0.025433	7.2	283.1	290.3	gb:NM_030556.1 /DB_XREF=gi:13385307 /GEN=Slc19a3 /FEA=FLmRNA /CNT=2 /TID=Mm.160173.1 /TIER=FL /STK=2 /UG=Mm.160173 /LL=80721 /DEF=Mus musculus solute carrier family 19 (sodiumhydrogen exchanger), member 3 (Slc19a3), mRNA. /PROD=solute carrier family 19 (s
0.024869	7.1	285.5	292.6	gb:R74675 /DB_XREF=gi:849878 /DB_XREF=MDB0826 /FEA=EST /CNT=2 /TID=Mm.5279.1 /TIER=ConsEnd /STK=2 /UG=Mm.5279 /LL=52819 /UG_GENE=D7Bwg0826e /UG_TITLE=DNA segment, Chr 7, Brigham & Womens Genetics 0826 expressed
0.024827	6.8	273.9	280.7	gb:NM_011720.1 /DB_XREF=gi:6756004 /GEN=Wnt8b /FEA=FLmRNA /CNT=11 /TID=Mm.88365.1 /TIER=FL /STK=2 /UG=Mm.88365 /LL=22423 /DEF=Mus musculus wingless related MMTV integration site 8b (Wnt8b), mRNA. /PROD=wingless related MMTV integration site 8b /FL=gb:AF130349.1 gb:NM_011720.1
0.024534	7.1	289.4	296.5	gb:BE634580 /DB_XREF=gi:9917268 /DB_XREF=uv63e06.x1 /CLONE=IMAGE:3413794 /FEA=EST /CNT=4 /TID=Mm.32834.1 /TIER=ConsEnd /STK=2 /UG=Mm.32834 /UG_TITLE=ESTs
0.024528	5.2	212	217.2	gb:AV124256 /DB_XREF=gi:5310490 /DB_XREF=AV124256 /CLONE=2700017F10 /FEA=EST /CNT=11 /TID=Mm.181847.4 /TIER=Stack /STK=11 /UG=Mm.181847 /LL=56612 /UG_GENE=Pfdn5 /UG_TITLE=prefoldin 5
0.024482	20.1	821	841.1	gb:NM_021442.1 /DB_XREF=gi:10946815 /GEN=Mds1 /FEA=FLmRNA /CNT=2 /TID=Mm.42023.1 /TIER=FL /STK=1 /UG=Mm.42023 /LL=17251 /DEF=Mus musculus myelodysplasia syndrome 1 homolog (human) (Mds1), mRNA. /PROD=myelodysplasia syndrome 1 protein homolog /FL=gb:NM_021
0.024384	10.1	414.2	424.3	gb:BG073495 /DB_XREF=gi:12556064 /DB_XREF=H3123C05-3 /CLONE=H3123C05 /FEA=EST /CNT=2 /TID=Mm.182657.1 /TIER=ConsEnd /STK=2 /UG=Mm.182657 /UG_TITLE=ESTs
0.02391	5.1	213.3	218.4	gb:NM_008863.1 /DB_XREF=gi:6679360 /GEN=Pkib /FEA=FLmRNA /CNT=13 /TID=Mm.1499.1 /TIER=FL /STK=2 /UG=Mm.1499 /LL=18768 /DEF=Mus musculus protein kinase inhibitor beta, cAMP dependent, testis specific (Pkib), mRNA. /PROD=protein kinase inhibitor beta, cAMP dependent,testis specific /FL=gb:U02241.1 gb:NM_008863.1
0.023418	8.4	358.7	367.1	gb:BF018708 /DB_XREF=gi:10750040 /DB_XREF=uv97b10.x1 /CLONE=IMAGE:3656443 /FEA=EST /CNT=29 /TID=Mm.87397.1 /TIER=Stack /STK=25 /UG=Mm.87397 /UG_TITLE=ESTs, Weakly similar to Charcot-Marie-Tooth duplicated region transcript 1 (Homo sapiens) (H.sapiens)
0.022295	6.8	305	311.8	gb:BM239335 /DB_XREF=gi:17874605 /DB_XREF=k0532H06-3 /CLONE=K0532H06 /FEA=EST /CNT=3 /TID=Mm.218225.1 /TIER=ConsEnd /STK=3 /UG=Mm.218225 /UG_TITLE=ESTs
0.022273	4.9	220	224.9	gb:AK019817.1 /DB_XREF=gi:12860164 /FEA=mRNA /CNT=1 /TID=Mm.195817.1 /TIER=ConsEnd /STK=0 /UG=Mm.195817 /LL=78933 /UG_GENE=4930578N11Rik /UG_TITLE=RIKEN cDNA 4930578N11 gene /DEF=Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clo
0.021783	6.5	298.4	304.9	gb:BB529729 /DB_XREF=gi:16445661 /DB_XREF=BB529729 /CLONE=E030003N18 /FEA=EST /CNT=5 /TID=Mm.209868.1 /TIER=ConsEnd /STK=3 /UG=Mm.209868 /UG_TITLE=ESTs
0.021007	8.3	395.1	403.4	gb:BB488016 /DB_XREF=gi:16441592 /DB_XREF=BB488016 /CLONE=D430047D06 /FEA=EST /CNT=4 /TID=Mm.135160.1 /TIER=ConsEnd /STK=2 /UG=Mm.135160 /UG_TITLE=ESTs
0.020991	7.5	357.3	364.8	gb:NM_009876.1 /DB_XREF=gi:6753387 /GEN=Cdkn1c /FEA=FLmRNA /CNT=214 /TID=Mm.168789.1 /TIER=FL+Stack /STK=39 /UG=Mm.168789 /LL=12577 /DEF=Mus musculus cyclin-dependent kinase inhibitor 1C (P57) (Cdkn1c), mRNA. /PROD=cyclin-dependent kinase inhibitor 1C (P57) /FL=gb:U20553.1 gb:NM_009876.1 gb:U22399.1 gb:BC005412.1
0.020866	5.3	254	259.3	gb:BE952702 /DB_XREF=gi:10593432 /DB_XREF=UI-M-BH4-azj-b-03-0-UI.s1 /CLONE=UI-M-BH4-azj-b-03-0-UI /FEA=mRNA /CNT=6 /TID=Mm.53687.1 /TIER=ConsEnd /STK=0 /UG=Mm.53687 /LL=72708 /UG_GENE=2810047E18Rik /UG_TITLE=RIKEN cDNA 2810047E18 gene
0.020507	7.2	351.1	358.3	gb:NM_020620.1 /DB_XREF=gi:10181169 /GEN=U48620 /FEA=FLmRNA /CNT=1 /TID=Mm.103774.1 /TIER=FL /STK=1 /UG=Mm.103774 /LL=57384 /DEF=Mus musculus cDNA sequence U48620 (U48620), mRNA. /PROD=cDNA sequence U48620 /FL=gb:NM_020620.1
0.019452	1479.7	76067.5	77547.2	gb:NM_013602.1 /DB_XREF=gi:7305284 /GEN=Mt1 /FEA=FLmRNA /CNT=145 /TID=Mm.219629.1 /TIER=FL+Stack /STK=36 /UG=Mm.219629 /LL=17748 /DEF=Mus musculus metallothionein 1 (Mt1), mRNA. /PROD=metallothionein 1 /FL=gb:NM_013602.1
0.018505	7.5	405.3	412.8	gb:BB037094 /DB_XREF=gi:15403721 /DB_XREF=BB037094 /CLONE=5930423H14 /FEA=EST /CNT=49 /TID=Mm.20258.1 /TIER=Stack /STK=43 /UG=Mm.20258 /UG_TITLE=ESTs, Weakly similar to YR86_CAEEL HYPOTHETICAL 93.7 KDA PROTEIN F48E8.6 IN CHROMOSOME III (C.elegans)
0.018146	4.6	253.5	258.1	gb:BB104606 /DB_XREF=gi:8757174 /DB_XREF=BB104606 /CLONE=9430092D12 /FEA=mRNA /CNT=2 /TID=Mm.159918.1 /TIER=ConsEnd /STK=1 /UG=Mm.159918 /LL=77427 /UG_GENE=9430092D12Rik /UG_TITLE=RIKEN cDNA 9430092D12 gene
0.016859	3.5	207.6	211.1	gb:BI652465 /DB_XREF=gi:15566701 /DB_XREF=602929555F1 /CLONE=IMAGE:5340012 /FEA=FLmRNA /CNT=60 /TID=Mm.24081.1 /TIER=ConsEnd /STK=0 /UG=Mm.24081 /LL=80281 /UG_GENE=BC003236 /UG_TITLE=hypothetical protein, MGC:7720 /FL=gb:BC003236.1 gb:NM_030249.1
0.015703	30.7	1955	1985.7	gb:NM_009998.1 /DB_XREF=gi:6753575 /GEN=Cyp2b10 /FEA=FLmRNA /CNT=2 /TID=Mm.218749.1 /TIER=FL /STK=1 /UG=Mm.218749 /LL=13088 /DEF=Mus musculus cytochrome P450, 2b10, phenobarbitol inducible, type b (Cyp2b10), mRNA. /PROD=cytochrome P450, 2b10, phenobarbitol
0.015209	3.2	210.4	213.6	gb:AV048291 /DB_XREF=gi:5134063 /DB_XREF=AV048291 /CLONE=1810004D23 /FEA=EST /CNT=8 /TID=Mm.29140.3 /TIER=Stack /STK=8 /UG=Mm.29140 /LL=67389 /UG_GENE=1110035L05Rik /UG_TITLE=RIKEN cDNA 1110035L05 gene
0.013192	704.5	53405.1	54109.6	gb:AA796766 /DB_XREF=gi:2859721 /DB_XREF=vp27c11.r1 /CLONE=IMAGE:1077908 /FEA=mRNA /CNT=34 /TID=Mm.142740.1 /TIER=Stack /STK=16 /UG=Mm.142740 /LL=17750 /UG_GENE=Mt2 /UG_TITLE=metallothionein 2
0.004551	95.3	20941.1	21036.4	gb:NM_008030.1 /DB_XREF=gi:6679814 /GEN=Fmo3 /FEA=FLmRNA /CNT=23 /TID=Mm.2900.1 /TIER=FL+Stack /STK=8 /UG=Mm.2900 /LL=14262 /DEF=Mus musculus flavin containing monooxygenase 3 (Fmo3), mRNA. /PROD=flavin containing monooxygenase 3 /FL=gb:U87147.1 gb:NM_008