Wt/Ko	WT 5mth	KO 5mth		Descriptions
1154.982	31415.5	27.2	31442.7	gb:NM_008295.1 /DB_XREF=gi:6680292 /GEN=Hsd3b5 /FEA=FLmRNA /CNT=13 /TID=Mm.17910.1 /TIER=FL /STK=4 /UG=Mm.17910 /LL=15496 /DEF=Mus musculus hydroxysteroid dehydrogenase-5, delta<5>-3-beta (Hsd3b5), mRNA. /PROD=hydroxysteroid dehydrogenase-5, delta<5>-3-beta /FL=gb:L41519.1 gb:BC012715.1 gb:NM_008295.1
300.6563	1924.2	6.4	1930.6	gb:BB222737 /DB_XREF=gi:8891349 /DB_XREF=BB222737 /CLONE=A530079E05 /FEA=EST /CNT=10 /TID=Mm.164639.1 /TIER=Stack /STK=10 /UG=Mm.164639 /UG_TITLE=ESTs, Moderately similar to A39567 thyroxine-binding globulin precursor - rat (R.norvegicus)
117.0655	4647.5	39.7	4687.2	gb:NM_010702.1 /DB_XREF=gi:6754529 /GEN=Lect2 /FEA=FLmRNA /CNT=19 /TID=Mm.16973.1 /TIER=FL+Stack /STK=9 /UG=Mm.16973 /LL=16841 /DEF=Mus musculus leukocyte cell-derived chemotaxin 2 (Lect2), mRNA. /PROD=leukocyte cell-derived chemotaxin 2 /FL=gb:NM_010702.
116.1818	255.6	2.2	257.8	gb:BC024323.1 /DB_XREF=gi:19353046 /FEA=FLmRNA /CNT=46 /TID=Mm.38181.1 /TIER=FL+Stack /STK=16 /UG=Mm.38181 /DEF=Mus musculus, clone MGC:19361 IMAGE:4242170, mRNA, complete cds. /PROD=Unknown (protein for MGC:19361) /FL=gb:BC024323.1
84.08983	6365.6	75.7	6441.3	gb:AA267743 /DB_XREF=gi:1904477 /DB_XREF=mz68a06.r1 /CLONE=IMAGE:718546 /FEA=EST /CNT=3 /TID=Mm.90033.2 /TIER=ConsEnd /STK=0 /UG=Mm.90033 /UG_TITLE=ESTs, Weakly similar to COTR MOUSE CONTRAPSIN PRECURSOR (M.musculus)
78.95238	497.4	6.3	503.7	gb:AV360029 /DB_XREF=gi:6407241 /DB_XREF=AV360029 /CLONE=7530404L21 /FEA=EST /CNT=2 /TID=Mm.95361.1 /TIER=ConsEnd /STK=2 /UG=Mm.95361 /UG_TITLE=ESTs
64.68627	329.9	5.1	335	gb:BM195021 /DB_XREF=gi:17746312 /DB_XREF=C0301A09-3 /CLONE=C0301A09 /FEA=EST /CNT=8 /TID=Mm.74762.1 /TIER=ConsEnd /STK=4 /UG=Mm.74762 /UG_TITLE=ESTs
62.82666	29321.2	466.7	29787.9	gb:AK011413.1 /DB_XREF=gi:12847521 /FEA=mRNA /CNT=3 /TID=Mm.196051.1 /TIER=ConsEnd /STK=1 /UG=Mm.196051 /LL=76914 /UG_GENE=2610016E04Rik /UG_TITLE=RIKEN cDNA 2610016E04 gene /DEF=Mus musculus 10 days embryo whole body cDNA, RIKEN full-length enriched libr
54.5798	649.5	11.9	661.4	gb:NM_133232.1 /DB_XREF=gi:18875393 /GEN=Pfkfb3 /FEA=FLmRNA /CNT=148 /TID=Mm.19669.1 /TIER=FL+Stack /STK=90 /UG=Mm.19669 /LL=170768 /DEF=Mus musculus inducible 6-phosphofructo-2-kinase (Pfkfb3), mRNA. /PROD=inducible 6-phosphofructo-2-kinase /FL=gb:AF294617.1 gb:NM_133232.1
42.68627	217.7	5.1	222.8	gb:BB108799 /DB_XREF=gi:8761367 /DB_XREF=BB108799 /CLONE=9530014F13 /FEA=EST /CNT=2 /TID=Mm.180263.1 /TIER=ConsEnd /STK=2 /UG=Mm.180263 /UG_TITLE=ESTs
42.64179	285.7	6.7	292.4	gb:AW546342 /DB_XREF=gi:7191751 /DB_XREF=L0006H07-3 /CLONE=L0006H07 /FEA=EST /CNT=2 /TID=Mm.182598.1 /TIER=ConsEnd /STK=2 /UG=Mm.182598 /UG_TITLE=ESTs
38.58434	44240.8	1146.6	45387.4	gb:NM_009060.1 /DB_XREF=gi:6677738 /GEN=Rgn /FEA=FLmRNA /CNT=97 /TID=Mm.2118.1 /TIER=FL+Stack /STK=35 /UG=Mm.2118 /LL=19733 /DEF=Mus musculus regucalcin (Rgn), mRNA. /PROD=regucalcin /FL=gb:D86217.1 gb:NM_009060.1 gb:U28937.1 gb:BC012710.1
37.86447	1033.7	27.3	1061	gb:AW488243 /DB_XREF=gi:7058513 /DB_XREF=UI-M-BH3-ary-c-10-0-UI.s1 /CLONE=UI-M-BH3-ary-c-10-0-UI /FEA=EST /CNT=52 /TID=Mm.26627.1 /TIER=Stack /STK=11 /UG=Mm.26627 /LL=98863 /UG_GENE=AI159670 /UG_TITLE=expressed sequence AI159670
36.70968	227.6	6.2	233.8	gb:AF359384.1 /DB_XREF=gi:13936905 /FEA=FLmRNA /CNT=1 /TID=Mm.179740.4 /TIER=FL /STK=1 /UG=Mm.179740 /LL=71132 /UG_GENE=4933421A18Rik /DEF=Mus musculus calcium binding protein CBP86-4 mRNA, complete cds. /PROD=calcium binding protein CBP86-4 /FL=gb:AF3593
36.54878	299.7	8.2	307.9	gb:AK017612.1 /DB_XREF=gi:12856942 /FEA=mRNA /CNT=1 /TID=Mm.206820.1 /TIER=ConsEnd /STK=0 /UG=Mm.206820 /LL=70545 /UG_GENE=5730433N10Rik /UG_TITLE=RIKEN cDNA 5730433N10 gene /DEF=Mus musculus 8 days embryo whole body cDNA, RIKEN full-length enriched libra
35.00367	5716.1	163.3	5879.4	gb:BC019856.1 /DB_XREF=gi:18044344 /FEA=FLmRNA /CNT=95 /TID=Mm.26207.1 /TIER=FL+Stack /STK=40 /UG=Mm.26207 /DEF=Mus musculus, clone MGC:29968 IMAGE:5123684, mRNA, complete cds. /PROD=Unknown (protein for MGC:29968) /FL=gb:BC019856.1
32.35448	6325.3	195.5	6520.8	gb:AB031813.1 /DB_XREF=gi:9971230 /GEN=oatp1 /FEA=FLmRNA /CNT=46 /TID=Mm.103665.1 /TIER=FL /STK=5 /UG=Mm.103665 /LL=28248 /DEF=Mus musculus mRNA for organic anion transporting polypeptide 1, complete cds. /PROD=organic anion transporting polypeptide 1 /FL
31.1875	249.5	8	257.5	gb:BB521778 /DB_XREF=gi:16443564 /DB_XREF=BB521778 /CLONE=D930003F11 /FEA=EST /CNT=7 /TID=Mm.208205.1 /TIER=ConsEnd /STK=5 /UG=Mm.208205 /UG_TITLE=ESTs
31.15625	598.2	19.2	617.4	gb:BB138760 /DB_XREF=gi:8793697 /DB_XREF=BB138760 /CLONE=9830162B01 /FEA=mRNA /CNT=70 /TID=Mm.31626.1 /TIER=Stack /STK=11 /UG=Mm.31626 /LL=78252 /UG_GENE=4432416J03Rik /UG_TITLE=RIKEN cDNA 4432416J03 gene
29.97872	281.8	9.4	291.2	gb:BG067949 /DB_XREF=gi:12550518 /DB_XREF=H3060A07-3 /CLONE=H3060A07 /FEA=EST /CNT=6 /TID=Mm.25527.1 /TIER=ConsEnd /STK=6 /UG=Mm.25527 /UG_TITLE=ESTs
29.34247	214.2	7.3	221.5	gb:NM_053255.1 /DB_XREF=gi:16716584 /GEN=Elac1 /FEA=FLmRNA /CNT=7 /TID=Mm.204642.1 /TIER=FL /STK=2 /UG=Mm.204642 /LL=114615 /DEF=Mus musculus elaC homolog 1 (E. coli) (Elac1), mRNA. /PROD=elaC homolog 1 (E. coli) /FL=gb:NM_053255.1 gb:AF308697.1 gb:BC019371.1
28.69512	235.3	8.2	243.5	gb:BB284249 /DB_XREF=gi:16401421 /DB_XREF=BB284249 /CLONE=A930105D20 /FEA=EST /CNT=3 /TID=Mm.186070.1 /TIER=ConsEnd /STK=3 /UG=Mm.186070 /UG_TITLE=ESTs
28.33	283.3	10	293.3	gb:AY079003.1 /DB_XREF=gi:19335751 /GEN=Foxp2 /FEA=FLmRNA /CNT=15 /TID=Mm.133208.1 /TIER=FL /STK=1 /UG=Mm.133208 /LL=114142 /DEF=Mus musculus forkheadwinged-helix transcription factor 2 (Foxp2) mRNA, complete cds. /PROD=forkheadwinged-helix transcription
28.32468	218.1	7.7	225.8	gb:AK017253.1 /DB_XREF=gi:12856398 /FEA=mRNA /CNT=8 /TID=Mm.22800.1 /TIER=ConsEnd /STK=4 /UG=Mm.22800 /LL=71392 /UG_GENE=5430401F13Rik /UG_TITLE=RIKEN cDNA 5430401F13 gene /DEF=Mus musculus 6 days neonate head cDNA, RIKEN full-length enriched library, clo
28.13793	244.8	8.7	253.5	gb:AK016576.1 /DB_XREF=gi:12855386 /GEN=Pcdh13 /FEA=mRNA /CNT=1 /TID=Mm.160292.2 /TIER=ConsEnd /STK=0 /UG=Mm.160292 /LL=93723 /UG_TITLE=protocadherin gamma subfamily A, 11 /DEF=Mus musculus adult male testis cDNA, RIKEN full-length enriched library, clone
28.0715	35844.5	1276.9	37121.4	gb:AK012685.1 /DB_XREF=gi:12849599 /FEA=mRNA /CNT=51 /TID=Mm.30070.1 /TIER=Stack /STK=15 /UG=Mm.30070 /LL=76971 /UG_GENE=2810007J24Rik /UG_TITLE=RIKEN cDNA 2810007J24 gene /DEF=Mus musculus 10, 11 days embryo whole body cDNA, RIKEN full-length enriched li
26.36145	218.8	8.3	227.1	gb:NM_009104.1 /DB_XREF=gi:7106398 /GEN=Rrm2 /FEA=FLmRNA /CNT=287 /TID=Mm.99.1 /TIER=FL+Stack /STK=124 /UG=Mm.99 /LL=20135 /DEF=Mus musculus ribonucleotide reductase M2 (Rrm2), mRNA. /PROD=ribonucleotide reductase M2 /FL=gb:M14223.1 gb:NM_009104.1
26.28866	255	9.7	264.7	gb:AV091347 /DB_XREF=gi:5238895 /DB_XREF=AV091347 /CLONE=2310061J04 /FEA=EST /CNT=2 /TID=Mm.178097.1 /TIER=ConsEnd /STK=2 /UG=Mm.178097 /UG_TITLE=ESTs
26.19018	426.9	16.3	443.2	gb:BM120546 /DB_XREF=gi:17088572 /DB_XREF=L0941A12-3 /CLONE=L0941A12 /FEA=EST /CNT=2 /TID=Mm.215558.1 /TIER=ConsEnd /STK=2 /UG=Mm.215558 /UG_TITLE=ESTs, Weakly similar to YO11 MOUSE HYPOTHETICAL PROTEIN ORF-1137 (M.musculus)
26.04282	22988	882.7	23870.7	gb:BC025936.1 /DB_XREF=gi:19683977 /FEA=FLmRNA /CNT=100 /TID=Mm.56950.1 /TIER=FL+Stack /STK=36 /UG=Mm.56950 /DEF=Mus musculus, clone MGC:37794 IMAGE:5097698, mRNA, complete cds. /PROD=Unknown (protein for MGC:37794) /FL=gb:BC026582.1 gb:BC025936.1 gb:BC01
26	226.2	8.7	234.9	gb:BC026654.1 /DB_XREF=gi:20071860 /FEA=FLmRNA /CNT=59 /TID=Mm.31552.1 /TIER=FL+Stack /STK=36 /UG=Mm.31552 /DEF=Mus musculus, clone MGC:37581 IMAGE:4988382, mRNA, complete cds. /PROD=Unknown (protein for MGC:37581) /FL=gb:BC018222.1 gb:BC026654.1
25.7375	205.9	8	213.9	gb:NM_007540.1 /DB_XREF=gi:6680777 /GEN=Bdnf /FEA=FLmRNA /CNT=5 /TID=Mm.1442.1 /TIER=FL /STK=1 /UG=Mm.1442 /LL=12064 /DEF=Mus musculus brain derived neurotrophic factor (Bdnf), mRNA. /PROD=brain derived neurotrophic factor /FL=gb:AY057913.1 gb:NM_007540.1
24.9571	110390	4423.2	114813.6	gb:NM_021282.1 /DB_XREF=gi:11276064 /GEN=Cyp2e1 /FEA=FLmRNA /CNT=381 /TID=Mm.21758.1 /TIER=FL+Stack /STK=151 /UG=Mm.21758 /LL=13106 /DEF=Mus musculus cytochrome P450, 2e1, ethanol inducible (Cyp2e1), mRNA. /PROD=cytochrome P450, 2e1, ethanol inducible /FL=gb:NM_021282.1 gb:L11650.1 gb:BC013451.1
24.70103	239.6	9.7	249.3	gb:BG072064 /DB_XREF=gi:12554633 /DB_XREF=H3106C12-3 /CLONE=H3106C12 /FEA=EST /CNT=3 /TID=Mm.17896.1 /TIER=ConsEnd /STK=2 /UG=Mm.17896 /UG_TITLE=ESTs
24.34409	226.4	9.3	235.7	gb:AF263536.1 /DB_XREF=gi:9964289 /GEN=Fgf23 /FEA=FLmRNA /CNT=5 /TID=Mm.154766.1 /TIER=FL /STK=1 /UG=Mm.154766 /LL=64654 /DEF=Mus musculus FGF23 (Fgf23) mRNA, complete cds. /PROD=FGF23 /FL=gb:AB037889.1 gb:NM_022657.1 gb:AF263536.1
24.31	243.1	10	253.1	gb:AI551101 /DB_XREF=gi:4483464 /DB_XREF=vx37b06.x1 /CLONE=IMAGE:1277363 /FEA=EST /CNT=3 /TID=Mm.23537.2 /TIER=ConsEnd /STK=1 /UG=Mm.23537 /UG_TITLE=ESTs
24.17886	297.4	12.3	309.7	gb:AK008865.1 /DB_XREF=gi:12843315 /FEA=mRNA /CNT=1 /TID=Mm.41036.1 /TIER=ConsEnd /STK=0 /UG=Mm.41036 /LL=70144 /UG_GENE=2210409B11Rik /UG_TITLE=RIKEN cDNA 2210409B11 gene /DEF=Mus musculus adult male stomach cDNA, RIKEN full-length enriched library, clon
23.97647	203.8	8.5	212.3	gb:AK013818.1 /DB_XREF=gi:12851323 /FEA=mRNA /CNT=10 /TID=Mm.63579.1 /TIER=ConsEnd /STK=7 /UG=Mm.63579 /LL=73075 /UG_GENE=2900084F20Rik /UG_TITLE=RIKEN cDNA 2900084F20 gene /DEF=Mus musculus adult male hippocampus cDNA, RIKEN full-length enriched library,
23.91608	342	14.3	356.3	gb:AW910654 /DB_XREF=gi:8075897 /DB_XREF=ur86d09.x1 /CLONE=IMAGE:3157169 /FEA=mRNA /CNT=56 /TID=Mm.200834.1 /TIER=Stack /STK=8 /UG=Mm.200834 /LL=116848 /UG_GENE=Baz2a /UG_TITLE=bromodomain adjacent to zinc finger domain, 2A
23.81818	209.6	8.8	218.4	gb:AI413247 /DB_XREF=gi:4256751 /DB_XREF=me93d10.x1 /CLONE=IMAGE:403123 /FEA=EST /CNT=4 /TID=Mm.151794.1 /TIER=ConsEnd /STK=2 /UG=Mm.151794 /UG_TITLE=ESTs, Weakly similar to FMN2_MOUSE Formin 2 (M.musculus)
23.75728	244.7	10.3	255	gb:AW553766 /DB_XREF=gi:7199189 /DB_XREF=L0231D06-3 /CLONE=L0231D06 /FEA=EST /CNT=7 /TID=Mm.77786.1 /TIER=ConsEnd /STK=5 /UG=Mm.77786 /UG_TITLE=ESTs
23.62637	215	9.1	224.1	gb:BM218963 /DB_XREF=gi:17778633 /DB_XREF=C0918C06-3 /CLONE=C0918C06 /FEA=mRNA /CNT=18 /TID=Mm.40497.1 /TIER=ConsEnd /STK=7 /UG=Mm.40497 /LL=73131 /UG_GENE=3110027P15Rik /UG_TITLE=RIKEN cDNA 3110027P15 gene
23.13793	671	29	700	gb:NM_010701.1 /DB_XREF=gi:6754527 /GEN=Lect1 /FEA=FLmRNA /CNT=18 /TID=Mm.46561.1 /TIER=FL+Stack /STK=10 /UG=Mm.46561 /LL=16840 /DEF=Mus musculus leukocyte cell derived chemotaxin 1 (Lect1), mRNA. /PROD=leukocyte cell derived chemotaxin 1 /FL=gb:NM_010701
22.98947	218.4	9.5	227.9	gb:BB360492 /DB_XREF=gi:16405866 /DB_XREF=BB360492 /CLONE=C030047H17 /FEA=EST /CNT=2 /TID=Mm.155868.1 /TIER=ConsEnd /STK=2 /UG=Mm.155868 /UG_TITLE=ESTs, Moderately similar to S12207 hypothetical protein (M.musculus)
22.87821	356.9	15.6	372.5	gb:U14648.1 /DB_XREF=gi:555923 /FEA=mRNA /CNT=1 /TID=Mm.218761.1 /TIER=ConsEnd /STK=0 /UG=Mm.218761 /LL=20381 /UG_GENE=Sfrs10 /UG_TITLE=splicing factor, arginineserine-rich 10 /DEF=Mus musculus putative myelin regulatory factor 1 mRNA, partial cds.
22.52239	301.8	13.4	315.2	gb:BB822150 /DB_XREF=gi:16994779 /DB_XREF=BB822150 /CLONE=G830017F04 /FEA=EST /CNT=9 /TID=Mm.141848.1 /TIER=Stack /STK=8 /UG=Mm.141848 /UG_TITLE=ESTs
22.5	238.5	10.6	249.1	gb:BB472451 /DB_XREF=gi:16428046 /DB_XREF=BB472451 /CLONE=D230045E19 /FEA=EST /CNT=2 /TID=Mm.216010.1 /TIER=ConsEnd /STK=2 /UG=Mm.216010 /UG_TITLE=ESTs
22.49138	260.9	11.6	272.5	gb:AK007714.1 /DB_XREF=gi:12841434 /FEA=mRNA /CNT=17 /TID=Mm.46042.1 /TIER=ConsEnd /STK=6 /UG=Mm.46042 /LL=69169 /UG_GENE=1810037B05Rik /UG_TITLE=RIKEN cDNA 1810037B05 gene /DEF=Mus musculus 10 day old male pancreas cDNA, RIKEN full-length enriched librar
22.42188	287	12.8	299.8	gb:BG092512 /DB_XREF=gi:12575075 /DB_XREF=mac12d03.x1 /CLONE=IMAGE:3999413 /FEA=EST /CNT=23 /TID=Mm.200270.1 /TIER=ConsEnd /STK=1 /UG=Mm.200270 /LL=100702 /UG_GENE=AI595338 /UG_TITLE=expressed sequence AI595338
22.34653	225.7	10.1	235.8	gb:Y11009.1 /DB_XREF=gi:1841447 /FEA=mRNA /CNT=3 /TID=Mm.204652.1 /TIER=ConsEnd /STK=1 /UG=Mm.204652 /UG_TITLE=M.musculus mRNA for L41 ribosomal like-protein /DEF=M.musculus mRNA for L41 ribosomal like-protein.