

201291_s_at Consensus includes gb:AU159942
/FEA=EST /DB_XREF=gi:11021463
/DB_XREF=est:AU159942
/CLONE=Y79AA1000724 /UG=Hs.156346
topoisomerase (DNA) II alpha (170kD)
/FL=gb:J04088.1 gb:NM_001067.1
201664_at "gb:AL136877.1 /DEF=Homo sapiens mRNA; cDNA
DKFZp434F205 (from clone DKFZp434F205); complete cds. /FEA=mRNA
/GEN=DKFZp434F205 /PROD=hypothetical protein /DB_XREF=gi:6807670
/UG=Hs.50758 SMC4 (structural maintenance of chromosomes 4, yeast)-like
1 /FL=gb:AB019987.1 gb:NM_005496.1 gb:AL136877.1"
201666_at "gb:NM_003254.1 /DEF=Homo sapiens tissue inhibitor of metalloproteinase 1
(erythroid potentiating activity, collagenase inhibitor) (TIMP1), mRNA. /FEA=mRNA /GEN=TIMP1
/PROD=tissue inhibitor of metalloproteinase 1precursor /DB_XREF=gi:4507508 /UG=Hs.5831 tissue
inhibitor of metalloproteinase 1 (erythroid potentiating activity, collagenase inhibitor)
/FL=gb:BC000866.1 gb:M12670.1 gb:M59906.1 gb:NM_003254.1"
201752_s_at Consensus includes
gb:AI763123 /FEA=EST
/DB_XREF=gi:5178790
/DB_XREF=est:wi06f09.x1
/CLONE=IMAGE:2389481 /UG=Hs.324470
adducin 3 (gamma) /FL=gb:D67031.1
gb:NM_019903.1
202912_at "gb:NM_001124.1 /DEF=Homo
sapiens adrenomedullin (ADM), mRNA.
/FEA=mRNA /GEN=ADM /PROD=adrenomedullin
/DB_XREF=gi:4501944 /UG=Hs.394
adrenomedullin /FL=gb:NM_001124.1
gb:D14874.1"
203554_x_at "gb:NM_004219.2 /DEF=Homo sapiens pituitary tumor-
transforming 1 (PTTG1), mRNA. /FEA=mRNA /GEN=PTTG1
/PROD=pituitary tumor-transforming protein 1 /DB_XREF=gi:11038651
/UG=Hs.252587 pituitary tumor-transforming 1 /FL=gb:NM_004219.2
gb:AF095287.1 gb:AF062649.1 gb:AF075242.1"
204713_s_at "Consensus includes gb:AA910306
/FEA=EST /DB_XREF=gi:3049596
/DB_XREF=est:ok83a04.s1 /CLONE=IMAGE:1520526
/UG=Hs.30054 coagulation factor V (proaccelerin, labile
factor) /FL=gb:NM_000130.2 gb:M16967.1 gb:M14335.1"
205137_x_at "gb:NM_005709.1 /DEF=Homo sapiens

PDZ-73 protein (PDZ-73NY-CO-38), mRNA. /FEA=mRNA
/GEN=PDZ-73NY-CO-38 /PROD=PDZ-73 protein
/DB_XREF=gi:5031978 /UG=Hs.132945 PDZ-73 protein
/FL=gb:AF039700.1 gb:NM_005709.1 gb:AB018687.1"
209155_s_at "gb:BC001595.1 /DEF=Homo sapiens, 5-nucleotidase
(purine), cytosolic type B, clone MGC:1109, mRNA, complete cds.
/FEA=mRNA /PROD=5-nucleotidase (purine), cytosolic type B
/DB_XREF=gi:12804388 /UG=Hs.138593 5-nucleotidase (purine),
cytosolic type B /FL=gb:BC001595.1"
210512_s_at "gb:AF022375.1 /DEF=Homo sapiens vascular
endothelial growth factor mRNA, complete cds. /FEA=mRNA
/PROD=vascular endothelial growth factor /DB_XREF=gi:3719220
/UG=Hs.73793 vascular endothelial growth factor /FL=gb:M32977.1
gb:AF022375.1 gb:NM_003376.1 gb:AB021221.1 gb:AF091352.1"
211184_s_at "gb:AB006955.1 /DEF=Homo
sapiens mRNA for AIE-75, complete cds.
/FEA=mRNA /GEN=aie-75 /PROD=AIE-75
/DB_XREF=gi:5152287 /UG=Hs.132945 PDZ-
73 protein /FL=gb:AB006955.1"
211527_x_at "gb:M27281.1 /DEF=Human
vascular permeability factor mRNA, complete
cds. /FEA=mRNA /DB_XREF=gi:340300
/UG=Hs.73793 vascular endothelial growth
factor /FL=gb:M27281.1"
211964_at "Consensus includes
gb:X05610.1 /DEF=Human mRNA for type IV
collagen alpha (2) chain. /FEA=mRNA
/PROD=alpha (2) chain /DB_XREF=gi:29550
/UG=Hs.75617 collagen, type IV, alpha 2"
211966_at "Consensus includes
gb:AA909035 /FEA=EST
/DB_XREF=gi:3048440
/DB_XREF=est:ol11h01.s1
/CLONE=IMAGE:1523185 /UG=Hs.75617
collagen, type IV, alpha 2"
211980_at "Consensus includes
gb:AI922605 /FEA=EST
/DB_XREF=gi:5658569
/DB_XREF=est:wm90c05.x1
/CLONE=IMAGE:2443208 /UG=Hs.119129

collagen, type IV, alpha 1
/FL=gb:NM_001845.1"
212171_x_at Consensus includes
gb:H95344 /FEA=EST
/DB_XREF=gi:1102977
/DB_XREF=est:yu21b08.s1
/CLONE=IMAGE:234423 /UG=Hs.73793
vascular endothelial growth factor
/FL=gb:AF214570.1
213217_at Consensus includes
gb:AU149572 /FEA=EST
/DB_XREF=gi:11011093
/DB_XREF=est:AU149572
/CLONE=NT2RM4002598 /UG=Hs.2352
adenylate cyclase 2 (brain)
213388_at Consensus includes gb:H15535
/FEA=EST /DB_XREF=gi:880355
/DB_XREF=est:ym27c01.s1
/CLONE=IMAGE:49385 /UG=Hs.52792 Homo
sapiens mRNA; cDNA DKFZp586l1823 (from
clone DKFZp586l1823)
213433_at Consensus includes
gb:AF038193.1 /DEF=Homo sapiens clone
23608 mRNA sequence. /FEA=mRNA
/DB_XREF=gi:2795913 /UG=Hs.6220 Homo
sapiens clone 23608 mRNA sequence
213689_x_at Consensus includes
gb:AL137958 /FEA=EST
/DB_XREF=gi:6854638
/DB_XREF=est:DKFZp761C1715_r1
/CLONE=DKFZp761C1715
/UG=Hs.180946 ribosomal protein L5
213900_at Consensus includes
gb:AA524029 /FEA=EST
/DB_XREF=gi:2264957
/DB_XREF=est:ng32f02.s1
/CLONE=IMAGE:936507 /UG=Hs.77889
Friedreich ataxia region gene X123
214930_at Consensus includes
gb:AW449813 /FEA=EST

/DB_XREF=gi:6990589
/DB_XREF=est:UI-H-BI3-akm-b-07-0-
UI.s1 /CLONE=IMAGE:2734788
/UG=Hs.58009 KIAA0918 protein
216044_x_at "Consensus includes gb:AK027146.1
/DEF=Homo sapiens cDNA: FLJ23493 fis, clone
LNG01831, highly similar to HSU66589 Human ribosomal
protein L5 pseudogene mRNA. /FEA=mRNA
/DB_XREF=gi:10440199 /UG=Hs.180946 ribosomal protein
L5"
218891_at "gb:NM_024541.1 /DEF=Homo sapiens
hypothetical protein FLJ13114 (FLJ13114), mRNA.
/FEA=mRNA /GEN=FLJ13114 /PROD=hypothetical protein
FLJ13114 /DB_XREF=gi:13375700 /UG=Hs.9444
hypothetical protein FLJ13114 /FL=gb:NM_024541.1"
219410_at "gb:NM_018004.1 /DEF=Homo sapiens
hypothetical protein FLJ10134 (FLJ10134), mRNA.
/FEA=mRNA /GEN=FLJ10134 /PROD=hypothetical protein
FLJ10134 /DB_XREF=gi:8922242 /UG=Hs.104800
hypothetical protein FLJ10134 /FL=gb:NM_018004.1"
219978_s_at "gb:NM_018454.1 /DEF=Homo sapiens uncharacterized
bone marrow protein BM037 (BM037), mRNA. /FEA=mRNA
/GEN=BM037 /PROD=uncharacterized bone marrow protein BM037
/DB_XREF=gi:8922094 /UG=Hs.283649 uncharacterized bone marrow
protein BM037 /FL=gb:AF217513.1 gb:NM_018454.1"
220005_at "gb:NM_023914.1 /DEF=Homo sapiens G protein-
coupled receptor 86 (GPR86), mRNA. /FEA=mRNA /GEN=GPR86
/PROD=G protein-coupled receptor 86 /DB_XREF=gi:13194202
/UG=Hs.13040 G protein-coupled receptor 86 /FL=gb:AF295368.1
gb:NM_023914.1 gb:AF178982.1 gb:AF345565.1"
221011_s_at "gb:NM_030915.1 /DEF=Homo sapiens
hypothetical protein DKFZp566J091 (DKFZP566J091),
mRNA. /FEA=mRNA /GEN=DKFZP566J091
/PROD=hypothetical protein DKFZp566J091
/DB_XREF=gi:13569871 /FL=gb:NM_030915.1"
221362_at "gb:NM_024012.1 /DEF=Homo sapiens 5-
hydroxytryptamine (serotonin) receptor 5A (HTR5A), mRNA.
/FEA=CDS /GEN=HTR5A /PROD=5-hydroxytryptamine (serotonin)
receptor 5A /DB_XREF=gi:13236496 /UG=Hs.248137 5-
hydroxytryptamine (serotonin) receptor 5A /FL=gb:NM_024012.1"

221527_s_at "gb:AF196185.1 /DEF=Homo sapiens atypical PKC isotype-specific interacting protein long variant mRNA, complete cds. /FEA=mRNA /PROD=atypical PKC isotype-specific interactingprotein long variant /DB_XREF=gi:13491609 /UG=Hs.72249 three-PDZ containing protein similar to C. elegans PAR3 (partitioning defect) /FL=gb:AF196185.1"

221729_at "Consensus includes gb:AL575735 /FEA=EST /DB_XREF=gi:12937190 /DB_XREF=est:AL575735 /CLONE=CS0DI070YK23 (3 prime) /UG=Hs.82985 collagen, type V, alpha 2 /FL=gb:NM_000393.1"

203891_s_at "gb:NM_001348.1 /DEF=Homo sapiens death-associated protein kinase 3 (DAPK3), mRNA. /FEA=mRNA /GEN=DAPK3 /PROD=death-associated protein kinase 3 /DB_XREF=gi:4557510 /UG=Hs.25619 death-associated protein kinase 3 /FL=gb:AB007144.1 gb:NM_001348.1 gb:AB022341.1"

204563_at "gb:NM_000655.2 /DEF=Homo sapiens selectin L (lymphocyte adhesion molecule 1) (SELL), mRNA. /FEA=mRNA /GEN=SELL /PROD=selectin L /DB_XREF=gi:5713320 /UG=Hs.82848 selectin L (lymphocyte adhesion molecule 1) /FL=gb:M25280.1 gb:NM_000655.2"

206932_at "gb:NM_003956.1 /DEF=Homo sapiens cholesterol 25-hydroxylase (CH25H), mRNA. /FEA=mRNA /GEN=CH25H /PROD=cholesterol 25-hydroxylase /DB_XREF=gi:4502498 /UG=Hs.194687 cholesterol 25-hydroxylase /FL=gb:AF059214.1 gb:NM_003956.1"

212623_at Consensus includes gb:AU153138 /FEA=EST /DB_XREF=gi:11014659 /DB_XREF=est:AU153138 /CLONE=NT2RP3002507 /UG=Hs.174905 KIAA0033 protein

216150_at "Consensus includes gb:AK021609.1 /DEF=Homo sapiens cDNA FLJ11547 fis, clone HEMBA1002934. /FEA=mRNA /DB_XREF=gi:10432824 /UG=Hs.306605 Homo sapiens cDNA FLJ11547 fis, clone HEMBA1002934"

216984_x_at "Consensus includes gb:D84143.1 /DEF=Human

immunoglobulin (mAb59) light chain V region mRNA, partial
sequence. /FEA=mRNA /PROD=immunoglobulin light chain V-J region
/DB_XREF=gi:1255613 /UG=Hs.121508 Human immunoglobulin
(mAb59) light chain V region mRNA, partial sequence"