

Primary Sequence Name	Sequence Code	Sequence Description
ANKRD34	AL359622	Hemochromatosis type 2 (juvenile)
LOC388494	AL365406	hypothetical gene supported by AL365406; BC034005
ABCA8	NM_007168	ATP-binding cassette, sub-family A (ABC1), member 8
TGFB2	AK021874	Transforming growth factor, beta 2
PAPLN	AL110280	Homo sapiens mRNA; cDNA DKFZp434F053 (from clone DKFZp434F053).
POLE4	NM_019896	Polymerase (DNA-directed), epsilon 4 (p12 subunit)
LGI1	NM_005097	Leucine-rich, glioma inactivated 1
NAT5	NM_016100	N-acetyltransferase 5 (ARD1 homolog, <i>S. cerevisiae</i> )
OGN	NM_014057	Osteoglycin (osteoinductive factor, mimecan)
VAT1	NM_006373	Vesicle amine transport protein 1 homolog (T californica)
	AL049974	Homo sapiens genomic DNA; cDNA DKFZp564B222 (from clone DKFZp564B222).
TMEM66	NM_016127	Transmembrane protein 66
UHMK1	AL137257	Homo sapiens, Similar to likely ortholog of yeast ARV1, clone IMAGE:3460560, mRNA
DDX10	NM_004398	DEAD (Asp-Glu-Ala-Asp) box polypeptide 10
BASP1	NM_006317	Brain abundant, membrane attached signal protein 1
CALU	NM_001219	Calumenin
PTGFRN	AB037857	Prostaglandin F2 receptor negative regulator
PIAS1	NM_016166	Protein inhibitor of activated STAT, 1
EMP2	NM_001424	Epithelial membrane protein 2
FOXJ1	NM_001454	Forkhead box J1
AEBP1	NM_001129	AE binding protein 1
PDZRN3	AB029018	PDZ domain containing RING finger 3
ARHGEF9	NM_015185	Cdc42 guanine nucleotide exchange factor (GEF) 9
MASK	NM_016542	Mst3 and SOK1-related kinase
HRASLS	NM_020386	HRAS-like suppressor
PPP1CB	NM_002709	Protein phosphatase 1, catalytic subunit, beta isoform
K-ALPHA-1	NM_006082	alpha tubulin
TAX1BP1	NM_006024	Tax1 (human T-cell leukemia virus type I) binding protein 1
C1QL1	NM_006688	Complement component 1, q subcomponent-like 1
TMEPAI	NM_020182	Transmembrane, prostate androgen induced RNA
C11orf2	NM_013265	Chromosome 11 open reading frame2
STAMBP	NM_006463	STAM binding protein
RBP4	NM_006744	Retinol binding protein 4, plasma
ATP6V1G1	NM_004888	ATPase, H <sup>+</sup> transporting, lysosomal 13kDa, V1 subunit G1
PICALM	NM_007166	Phosphatidylinositol binding clathrin assembly protein
MTCH1	NM_014341	Mitochondrial carrier homolog 1 ( <i>C. elegans</i> )
SLC17A7	NM_020309	Solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 7
SNX2	NM_003100	Sorting nexin 2

FCN1	NM_002003	Ficolin (collagen/fibrinogen domain containing) 1
RIMS2	NM_014677	Regulating synaptic membrane exocytosis 2
USP12	AF022789	Ubiquitin specific peptidase 12
ASS	NM_000050	Argininosuccinate synthetase
SCG2	NM_003469	Secretogranin II (chromogranin C)
RUVBL2	NM_006666	RuvB-like 2 (E. coli)
MTHFD1	NM_005956	Methylenetetrahydrofolate dehydrogenase (NADP+ dependent) 1, methenyltetrahydrofolate cyclohydrolase, formyltetrahydrofolate synthetase
SUB1	NM_006713	SUB1 homolog (S. cerevisiae)
EIF5A	NM_001970	Eukaryotic translation initiation factor 5A
	AK024054	Homo sapiens cDNA FLJ13992 fis, clone Y79AA1002139, weakly similar to DNAJ PROTEIN HOMOLOG 1.
RPL13	NM_000977	Ribosomal protein L13
COX6B1	NM_001863	Cytochrome c oxidase subunit Vib polypeptide 1 (ubiquitous)
KLK11	NM_006853	Kallikrein 11
GPC3	NM_004484	Glypican 3
PGAM1	NM_002629	Phosphoglycerate mutase 1 (brain)
NUDT3	NM_006703	Ribosomal protein S10
CIB1	NM_006384	Calcium and integrin binding 1 (calmyrin)
H1FX	NM_006026	H1 histone family, member X
TPI1	NM_000365	Triosephosphate isomerase 1
PRO0132	NM_014116	PRO0132 protein
C14orf132	AK001590	Chromosome 14 open reading frame 132
NOSIP	NM_015953	Nitric oxide synthase interacting protein
	AK001020	CDNA FLJ10158 fis, clone HEMBA1003463
LOC441207	AK026723	CDNA: FLJ23070 fis, clone LNG05629
FOXC1	AK021858	Forkhead box C1
NF2	NM_016418	Neurofibromin 2 (bilateral acoustic neuroma)
KCNG1	NM_002237	Potassium voltage-gated channel, subfamily G, member 1
LHFP	NM_005780	Lipoma HMGIC fusion partner
LOC440841	AK026368	Homo sapiens, clone IMAGE:4720764, mRNA
LOC339229	AY007126	hypothetical protein LOC339229
TFF3	NM_003226	Trefoil factor 3 (intestinal)
CCL26	NM_006072	Chemokine (C-C motif) ligand 26
TGFBR3	NM_003243	Transforming growth factor, beta receptor III (betaglycan, 300kDa)
PGF	NM_002632	Placental growth factor, vascular endothelial growth factor-related protein
ECGF1	NM_001953	Endothelial cell growth factor 1 (platelet-derived)
GRIK3	NM_000831	Glutamate receptor, ionotropic, kainate 3
GPATC1	NM_018025	G patch domain containing 1
TUBGCP3	NM_006322	Tubulin, gamma complex associated protein 3
FGF13	NM_004114	Fibroblast growth factor 13
TXNRD3	AF171055	Thioredoxin reductase 3
ANKRD16	AL137614	Ankyrin repeat domain 16

NKIRAS2	NM_017595	NFKB inhibitor interacting Ras-like 2
DBC1	NM_014618	Deleted in bladder cancer 1
FABP4	NM_001442	Fatty acid binding protein 4, adipocyte
LMNB1	NM_005573	Lamin B1
COMP	NM_000095	Cartilage oligomeric matrix protein
FER1L3	NM_013451	Fer-1-like 3, myoferlin ( <i>C. elegans</i> )
SULF1	AB029000	Sulfatase 1
SAR1B	NM_016103	SAR1 gene homolog B ( <i>S. cerevisiae</i> )
DARC	NM_002036	Duffy blood group, chemokine receptor
BCL3	NM_005178	B-cell CLL/lymphoma 3
PARVA	NM_018222	Parvin, alpha
SDCBP	NM_005625	Syndecan binding protein (syntenin)
RAB11A	NM_004663	RAB11A, member RAS oncogene family
PSD3	NM_015310	Pleckstrin and Sec7 domain containing 3
SLC4A4	NM_003759	Solute carrier family 4, sodium bicarbonate cotransporter, member 4
COX17	NM_005694	COX17 homolog, cytochrome c oxidase assembly protein (yeast)
MSN	NM_002444	Moesin
UBA52	NM_003333	Ubiquitin A-52 residue ribosomal protein fusion product 1
FKBP3	NM_002013	FK506 binding protein 3, 25kDa
EHD2	NM_014601	Glioma tumor suppressor candidate region gene 2
CPE	NM_001873	Carboxypeptidase E
RPS27A	NM_002954	Ribosomal protein S27a
SCGB1D1	NM_006552	Secretoglobin, family 1D, member 1
HSD17B2	NM_002153	Hydroxysteroid (17-beta) dehydrogenase 2
SEC11L1	NM_014300	SEC11-like 1 ( <i>S. cerevisiae</i> )
DYNLRB1	NM_014183	Dynein, light chain, roadblock-type 1
PIR	NM_003662	Pirin (iron-binding nuclear protein)
LOC51255	NM_016494	hypothetical protein LOC51255
NFE2L2	NM_006164	Nuclear factor (erythroid-derived 2)-like 2
SPEN	NM_015001	Spen homolog, transcriptional regulator ( <i>Drosophila</i> )
LSM8	NM_016200	LSM8 homolog, U6 small nuclear RNA associated ( <i>S. cerevisiae</i> )
EEF1G	NM_001404	Eukaryotic translation elongation factor 1 gamma
AARS	NM_001605	Alanyl-tRNA synthetase
NDUFB3	NM_002491	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3, 12kDa
POLRMT	NM_005035	Polymerase (RNA) mitochondrial (DNA directed)
LRRC23	NM_006992	Leucine rich repeat containing 23
UST	NM_005715	Uronyl-2-sulfotransferase
RNF10	NM_014868	Ring finger protein 10
ETFA	NM_000126	Electron-transfer-flavoprotein, alpha polypeptide (glutaric aciduria II)
SOD3	NM_003102	Superoxide dismutase 3, extracellular
CYP1B1	NM_000104	Cytochrome P450, family 1, subfamily B, polypeptide 1
COL12A1	AK022029	Collagen, type XII, alpha 1

LOH3CR2A	NM_013343	Loss of heterozygosity, 3, chromosomal region 2, gene A
IFT20	AF070643	Intraflagellar transport 20 homolog (Chlamydomonas)
C11orf43	AK025719	Chromosome 11 open reading frame 43
POLR2I	NM_006233	Polymerase (RNA) II (DNA directed) polypeptide I, 14.5kDa
PITX2	NM_000325	Paired-like homeodomain transcription factor 2
DSCR1	NM_004414	Down syndrome critical region gene 1
HSPB2	NM_001541	Heat shock 27kDa protein 2
NDUFA8	NM_014222	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 8, 19kDa
FHL1	NM_001449	Four and a half LIM domains 1
CHGA	NM_001275	Chromogranin A (parathyroid secretory protein 1)
ERBB2IP	NM_018695	ErbB2 interacting protein
GNG11	NM_004126	Guanine nucleotide binding protein (G protein), gamma 11
EMP1	NM_001423	Epithelial membrane protein 1
RASGRP1	NM_005739	RAS guanyl releasing protein 1 (calcium and DAG-regulated)
ARHGAP29	NM_004815	Rho GTPase activating protein 29
KIAA1600	AB046820	KIAA1600
DKK3	NM_013253	Dickkopf homolog 3 (Xenopus laevis)
RGC32	NM_014059	response gene to complement 32
PALM	NM_002579	Paralemmin
LMTK2	NM_014916	Lemur tyrosine kinase 2
CDH1	NM_004360	Cadherin 1, type 1, E-cadherin (epithelial)
RND3	NM_005168	Rho family GTPase 3
KIT	NM_000222	V-kit Hardy-Zuckerman 4 feline sarcoma viral oncogene homolog
DLG4	NM_001365	Discs, large homolog 4 (Drosophila)
CHN1	NM_001822	Chimerin (chimaerin) 1
TP53	NM_000546	Tumor protein p53 (Li-Fraumeni syndrome)
TBCA	NM_004607	Tubulin-specific chaperone a
GABARAPL2	NM_007285	GABA(A) receptor-associated protein-like 2
BTG2	NM_006763	BTG family, member 2
SPTBN5	NM_016642	Spectrin, beta, non-erythrocytic 5
SPOCK3	NM_016950	Sparc/osteonectin, cwcv and kazal-like domains proteoglycan (testican) 3
FCGBP	NM_003890	Fc fragment of IgG binding protein
CASC4	AK022565	Cancer susceptibility candidate 4
15-sep	NM_004261	15 kDa selenoprotein
CD99	NM_002414	CD99 molecule
CAT	NM_001752	Catalase
SLC29A3	NM_018344	Solute carrier family 29 (nucleoside transporters), member 3
CHRNA1	NM_000079	Cholinergic receptor, nicotinic, alpha 1 (muscle)
RABL3	AK025772	RAB, member of RAS oncogene family-like 3
CTH	NM_001902	Cystathionase (cystathionine gamma-lyase)
RBM14	NM_006328	RNA binding motif protein 4

RPS10	NM_001014	Ribosomal protein S10
RPL39	NM_001000	Ribosomal protein L39
COL4A3BP	NM_005713	Collagen, type IV, alpha 3 (Goodpasture antigen) binding protein
RPL15	NM_002948	Ribosomal protein L15
QPCT	NM_012413	GlutaminyI-peptide cyclotransferase (glutaminyI cyclase)
GAPDH	NM_002046	Glyceraldehyde-3-phosphate dehydrogenase
YPEL5	NM_016061	Yippee-like 5 (Drosophila)
LSM4	NM_012321	LSM4 homolog, U6 small nuclear RNA associated (S. cerevisiae)
SCAND1	NM_016558	SCAN domain containing 1
ST8SIA5	NM_013305	ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 5
GMDS	NM_001500	GDP-mannose 4,6-dehydratase
MAN1C1	NM_020379	Mannosidase, alpha, class 1C, member 1
AK1	NM_000476	Adenylate kinase 1
GUK1	NM_000858	Guanylate kinase 1
HIRIP3	NM_003609	HIRA interacting protein 3
HERC5	NM_016323	Hect domain and RLD 5
ABCC5	AL359600	ATP-binding cassette, sub-family C (CFTR/MRP), member 5
C10orf56	AL049949	Chromosome 10 open reading frame 56
TUBB4	NM_006087	Tubulin, beta 4
SH3PXD2A	AK025783	SH3 and PX domains 2A
PRO1777	NM_018605	Homo sapiens hypothetical protein PRO1777 (PRO1777), mRNA.
HNRPH2	NM_019597	Ribosomal protein L36a
SPP1	NM_000582	Secreted phosphoprotein 1 (osteopontin, bone sialoprotein I, early T-lymphocyte activation 1)
PDE6C	NM_006204	Phosphodiesterase 6C, cGMP-specific, cone, alpha prime
LOC493869	AK022110	similar to RIKEN cDNA 2310016C16
NECAP1	NM_015509	NECAP endocytosis associated 1
BM88	NM_016564	BM88 antigen
EDN3	NM_000114	Endothelin 3
CP	NM_000096	Ceruloplasmin (ferroxidase)
AOC3	NM_003734	Amine oxidase, copper containing 3 (vascular adhesion protein 1)
ChGn	NM_018371	chondroitin beta1,4 N-acetylgalactosaminyltransferase
CDC25C	NM_001790	Cell division cycle 25C
CUL2	NM_003591	Cullin 2
MAD1L1	NM_003550	MAD1 mitotic arrest deficient-like 1 (yeast)
EDNRB	NM_000115	Endothelin receptor type B
TREM2	NM_018965	Triggering receptor expressed on myeloid cells 2
IL11RA	NM_004512	Interleukin 11 receptor, alpha
MN1	NM_002430	Meningioma (disrupted in balanced translocation) 1
CFDP1	NM_006324	Craniofacial development protein 1

FLJ10633	NM_018162	Homo sapiens hypothetical protein FLJ10633 (FLJ10633), mRNA.
COBL	AB014533	Cordon-bleu homolog (mouse)
SLC2A5	NM_003039	Solute carrier family 2 (facilitated glucose/fructose transporter), member 5
SERHL2	NM_014509	Serine hydrolase-like 2
C16orf5	NM_013399	Chromosome 16 open reading frame 5
CXCL14	NM_004887	Chemokine (C-X-C motif) ligand 14
PROCR	NM_006404	Protein C receptor, endothelial (EPCR)
UBQLN1	NM_013438	Ubiquilin 1
ADAMTS4	NM_005099	ADAM metallopeptidase with thrombospondin type 1 motif, 4
LGMN	NM_005606	Legumain
HBA1	NM_000558	Hemoglobin, alpha 1
ARFGAP1	NM_018209	ADP-ribosylation factor GTPase activating protein 1
SPCS2	NM_014752	Signal peptidase complex subunit 2 homolog (S. cerevisiae)
COQ6	NM_015940	Homo sapiens coenzyme Q6 homolog (yeast) (COQ6), transcript variant 1, mRNA.
KIAA1199	AB033025	KIAA1199
RPS15A	NM_001019	Ribosomal protein S15a
TINP1	NM_014886	TGF beta-inducible nuclear protein 1
DNAJB2	NM_006736	DnaJ (Hsp40) homolog, subfamily B, member 2
DNAJB1	NM_006145	DnaJ (Hsp40) homolog, subfamily B, member 1
ERGIC3	NM_015966	ERGIC and golgi 3
P4HA1	NM_000917	Procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide I
UCHL5	NM_015984	Ubiquitin carboxyl-terminal hydrolase L5
TPP1	NM_000391	Tripeptidyl peptidase I
PLTP	NM_006227	Phospholipid transfer protein
CHD1L	NM_004284	Chromodomain helicase DNA binding protein 1-like
ZHX2	NM_014943	Zinc fingers and homeoboxes 2
SAP30BP	NM_013260	SAP30 binding protein
PTOV1	NM_017432	Prostate tumor overexpressed gene 1
ARNT	NM_001668	Aryl hydrocarbon receptor nuclear translocator
MLLT10	NM_004641	Myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila)
NDUFS8	NM_002496	NADH dehydrogenase (ubiquinone) Fe-S protein 8, 23kDa (NADH-coenzyme Q reductase)
FLJ20701	NM_017933	hypothetical protein FLJ20701
OTUB1	NM_017670	OTU domain, ubiquitin aldehyde binding 1
TMEM51	NM_018022	Transmembrane protein 51
BTD	NM_000060	Biotinidase
CYB5R3	NM_007326	Cytochrome b5 reductase 3
NCKAP1	NM_013436	NCK-associated protein 1
CNGA3	NM_001298	Cyclic nucleotide gated channel alpha 3
WFDC2	NM_006103	WAP four-disulfide core domain 2
PDGFRA	NM_006206	Platelet-derived growth factor receptor, alpha polypeptide

RGS20	NM_003702	Regulator of G-protein signalling 20
GPS2	NM_004489	G protein pathway suppressor 2
FJX1	NM_014344	Four jointed box 1 (Drosophila)
HBEGF	NM_001945	Heparin-binding EGF-like growth factor
GNRH2	NM_001501	Gonadotropin-releasing hormone 2
PSMD10	NM_002814	Proteasome (prosome, macropain) 26S subunit, non-ATPase, 10
KATNB1	NM_005886	Katanin p80 (WD repeat containing) subunit B 1
SNX14	AK026479	Homo sapiens cDNA: FLJ22826 fis, clone KAIA4022, highly similar to AF121863 Homo sapiens sorting nexin 14 (SNX14) mRNA.
HUS1	NM_004507	HUS1 checkpoint homolog (S. pombe)
CD302	NM_014880	CD302 molecule
CHCHD2	NM_016139	Coiled-coil-helix-coiled-coil-helix domain containing 2
SNX3	NM_003795	Sorting nexin 3
RNUT1	NM_005701	RNA, U transporter 1
TLR6	NM_006068	Toll-like receptor 1
NEF3	NM_005382	Neurofilament 3 (150kDa medium)
RPS16	NM_001020	Ribosomal protein S16
SLC22A18	NM_002555	Solute carrier family 22 (organic cation transporter), member 18
FDFT1	NM_004462	Farnesyl-diphosphate farnesyltransferase 1
CCL19	NM_006274	Chemokine (C-C motif) ligand 19
CNN3	NM_001839	Calponin 3, acidic
DUSP6	NM_001946	Dual specificity phosphatase 6
UBE2L3	NM_003347	Ubiquitin-conjugating enzyme E2L 3
CTSB	NM_001908	Cathepsin B
MTHFD1L	AL117452	Methylenetetrahydrofolate dehydrogenase (NADP+ dependent) 1-like
BZW2	NM_014038	Basic leucine zipper and W2 domains 2
MLANA	NM_005511	Melan-A
PECI	NM_006117	Peroxisomal D3,D2-enoyl-CoA isomerase
CDR2L	L02867	Cerebellar degeneration-related protein 2-like
RPS9	NM_001013	Ribosomal protein S9
SRBD1	NM_018079	S1 RNA binding domain 1
ZMPSTE24	NM_005857	Zinc metalloproteinase (STE24 homolog, yeast)
RPL17	NM_000985	Ribosomal protein L17
NHSL1	AB037778	NHS-like 1
DYNC1LI2	NM_006141	Dynein, cytoplasmic 1, light intermediate chain 2
LDHB	NM_002300	Lactate dehydrogenase B
ELL	NM_006532	Elongation factor RNA polymerase II
C14orf94	NM_017815	Chromosome 14 open reading frame 94
GFPT2	NM_005110	Glutamine-fructose-6-phosphate transaminase 2
PRO0529	NM_014074	Homo sapiens PRO0529 protein (PRO0529), mRNA.
OPN1MW	NM_000513	Opsin 1 (cone pigments), medium-wave-sensitive (color blindness, deutan)
NEFL	NM_006158	Neurofilament, light polypeptide 68kDa
TIMP2	AL110197	TIMP metalloproteinase inhibitor 2

GDF15	NM_004864	Growth differentiation factor 15
TEGT	NM_003217	Testis enhanced gene transcript (BAX inhibitor 1)
COL9A3	NM_001853	Collagen, type IX, alpha 3
BTG1	NM_001731	B-cell translocation gene 1, anti-proliferative
BCL2	NM_000633	B-cell CLL/lymphoma 2
PLSCR4	NM_020353	Phospholipid scramblase 4
EI24	NM_004879	Etoposide induced 2.4 mRNA
LTBP1	NM_000627	Latent transforming growth factor beta binding protein 1
EGF	NM_001963	Epidermal growth factor (beta-urogastrone)
ITGB4	NM_000213	Integrin, beta 4
LIMS2	NM_017980	LIM and senescent cell antigen-like domains 2
CENPB	X05299	Centromere protein B, 80kDa
PXMP3	NM_000318	Peroxisomal membrane protein 3, 35kDa (Zellweger syndrome)
PLEK2	NM_016445	Pleckstrin 2
NBL1	NM_005380	Neuroblastoma, suppression of tumorigenicity 1
KRT19	NM_002276	Keratin 17
ARMCX3	NM_016607	Armadillo repeat containing, X-linked 3
MYCBP2	NM_015057	MYC binding protein 2
PTGS2	NM_000963	Prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase)
TNNT1	NM_003283	Troponin T type 1 (skeletal, slow)
PTX3	NM_002852	Pentraxin-related gene, rapidly induced by IL-1 beta
KBTBD10	NM_006063	Kelch repeat and BTB (POZ) domain containing 10
CXCL2	NM_002089	Chemokine (C-X-C motif) ligand 2
UBE2N	NM_003348	Ubiquitin-conjugating enzyme E2N (UBC13 homolog, yeast)
HPGD	NM_000860	Hydroxyprostaglandin dehydrogenase 15-(NAD)
PLA1A	NM_015900	Phospholipase A1 member A
TRPC3	NM_003305	Transient receptor potential cation channel, subfamily C, member 3
SF3B14	NM_016047	splicing factor 3B, 14 kDa subunit
TUFM	NM_003321	Tu translation elongation factor, mitochondrial
RALY	NM_016732	RNA binding protein, autoantigenic (hnRNP-associated with lethal yellow homolog (mouse))
PCSK7	NM_004716	Proprotein convertase subtilisin/kexin type 7
ACOT7	NM_007274	Acyl-CoA thioesterase 7
OAZ1	NM_004152	Ornithine decarboxylase antizyme 1
SETBP1	NM_015559	SET binding protein 1
LSM3	NM_014463	LSM3 homolog, U6 small nuclear RNA associated (S. cerevisiae)
NACA	NM_005594	Nascent-polypeptide-associated complex alpha polypeptide
RPL7	NM_000971	Ribosomal protein L7
FBN1	NM_000138	Fibrillin 1 (Marfan syndrome)
SDHD	NM_003002	Succinate dehydrogenase complex, subunit D, integral membrane protein
GPD1L	D42047	Glycerol-3-phosphate dehydrogenase 1-like



EIF3S6	NM_001568	Eukaryotic translation initiation factor 3, subunit 6 48kDa
BHLHB2	NM_003670	Basic helix-loop-helix domain containing, class B, 2
PTMA	NM_002823	Prothymosin, alpha (gene sequence 28)
FOXC1	NM_001453	Forkhead box C1
ZIC1	NM_003412	Zic family member 1 (odd-paired homolog, Drosophila)
TPST1	NM_003596	Tyrosylprotein sulfotransferase 1
GLO1	NM_006708	Glyoxalase I
NDUFB7	NM_004146	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 7, 18kDa
RICS	NM_014715	Rho GTPase-activating protein
LAX1	NM_017773	Lymphocyte transmembrane adaptor 1
C17orf59	NM_017622	Chromosome 17 open reading frame 59
NS5ATP13TP2	AK021720	NS5ATP13TP2 protein
SYN1	NM_006950	Synapsin I
C20orf24	NM_018840	Chromosome 20 open reading frame 24
MAP1B	NM_005909	Microtubule-associated protein 1B
FRZB	NM_001463	Frizzled-related protein
CAPZB	NM_004930	Capping protein (actin filament) muscle Z-line, beta
GOLGA7	NM_016099	Golgi autoantigen, golgin subfamily a, 7
STMN2	NM_007029	Stathmin-like 2
PROS1	NM_000313	Protein S (alpha)
AKAP12	NM_005100	A kinase (PRKA) anchor protein (gravin) 12
NMUR1	NM_006056	Neuromedin U receptor 1
PMP22	NM_000304	Peripheral myelin protein 22
FLJ10292	NM_018048	mago-nashi homolog
COL9A1	NM_001851	Collagen, type IX, alpha 1
RUFY3	NM_014961	RUN and FYVE domain containing 3
OLFML3	NM_020190	Olfactomedin-like 3
ARHGAP21	AB037845	Rho GTPase activating protein 21
CHN2	NM_004067	Chimerin (chimaerin) 2
CLEC2D	NM_013269	C-type lectin domain family 2, member D
FLJ10357	NM_018071	hypothetical protein FLJ10357
GEM	NM_005261	GTP binding protein overexpressed in skeletal muscle
TUBB3	NM_006086	Tubulin, beta 3
THBS4	NM_003248	Thrombospondin 4
EPB41L3	NM_012307	Erythrocyte membrane protein band 4.1-like 3
ARF4	NM_001660	ADP-ribosylation factor 4
GHR	NM_000163	Growth hormone receptor
WDR45	NM_007075	WD repeat domain 45
TUBB2A	NM_001069	Tubulin, beta 2B
SERPINF1	NM_002615	Serpin peptidase inhibitor, clade F (alpha-2 antiplasmin, pigment epithelium derived factor), member 1
M6PRBP1	NM_005817	Mannose-6-phosphate receptor binding protein 1
OSBPL7	NM_017731	Oxysterol binding protein-like 7
KIAA1458	AB040891	KIAA1458 protein

CCS	NM_005125	Copper chaperone for superoxide dismutase
SLC22A17	NM_016609	Solute carrier family 22 (organic cation transporter), member 17
ATP6AP1	NM_001183	ATPase, H <sup>+</sup> transporting, lysosomal accessory protein 1
EMILIN1	NM_007046	Elastin microfibril interfacier 1
SNAP25	NM_003081	Synaptosomal-associated protein, 25kDa
ACSL3	NM_004457	Acyl-CoA synthetase long-chain family member 3
LTBP2	NM_000428	Latent transforming growth factor beta binding protein 2
MCCC1	NM_020166	Methylcrotonoyl-Coenzyme A carboxylase 1 (alpha)
PSMB5	NM_002797	Proteasome (prosome, macropain) subunit, beta type, 5
PSMD3	NM_002809	Proteasome (prosome, macropain) 26S subunit, non-ATPase, 3
CYP51A1	NM_000786	Cytochrome P450, family 51, subfamily A, polypeptide 1
PRDX6	NM_004905	Peroxiredoxin 6
SLC17A6	NM_020346	Solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 6
LSM1	NM_014462	LSM1 homolog, U6 small nuclear RNA associated (S. cerevisiae)
RPS5	NM_001009	Ribosomal protein S5
RPL11	NM_000975	Ribosomal protein L11
SIX6	NM_007374	Sine oculis homeobox homolog 6 (Drosophila)
TMLHE	NM_018196	Trimethyllysine hydroxylase, epsilon
CEBPA	NM_004364	CCAAT/enhancer binding protein (C/EBP), alpha
NOX4	NM_016931	NADPH oxidase 4
CRAT	NM_004003	Carnitine acetyltransferase
COX7A2L	NM_004718	Cytochrome c oxidase subunit VIIa polypeptide 2 like
VGLL4	D50911	Vestigial like 4 (Drosophila)
SHOX	NM_000451	Short stature homeobox
H3F3A	NM_002107	H3 histone, family 3A
OSTM1	NM_014028	Osteopetrosis associated transmembrane protein 1
	AL109783	MRNA full length insert cDNA clone EUROIMAGE 163507
KIAA1345	AB037766	KIAA1345 protein
SLIT1	NM_003061	Slit homolog 1 (Drosophila)
ACTG1	NM_001614	Actin, gamma 1
GYPC	NM_002101	Glycophorin C (Gerbich blood group)
TNNI3	NM_000363	Troponin T type 1 (skeletal, slow)
SFRP4	NM_003014	Secreted frizzled-related protein 4
CDH13	NM_001257	Cadherin 13, H-cadherin (heart)
ECEL1	NM_004826	Endothelin converting enzyme-like 1
PTPLA	NM_014241	Protein tyrosine phosphatase-like (proline instead of catalytic arginine), member A
RARRES1	NM_002888	Retinoic acid receptor responder (tazarotene induced) 1
CEP250	NM_007186	Centrosomal protein 250kDa

IER3IP1	NM_016097	Immediate early response 3 interacting protein 1
COLEC10	NM_006438	Collectin sub-family member 10 (C-type lectin)
PACSIN2	NM_007229	Protein kinase C and casein kinase substrate in neurons 2
MAL	NM_002371	Mal, T-cell differentiation protein
ADM	NM_001124	Adrenomedullin
FLJ20674	NM_019086	hypothetical protein FLJ20674
PLS3	NM_005032	Plastin 3 (T isoform)
KRT14	NM_000526	Keratin 17
FHIT	NM_002012	Fragile histidine triad gene
AVPR2	NM_000054	Arginine vasopressin receptor 2 (nephrogenic diabetes insipidus)
LAPTM4A	NM_014713	Lysosomal-associated protein transmembrane 4 alpha
CENTB1	NM_014716	Centaurin, beta 1
ANP32B	NM_006401	Acidic (leucine-rich) nuclear phosphoprotein 32 family, member B
EVI1	AK025934	Ecotropic viral integration site 1
XLKD1	NM_006691	Extracellular link domain containing 1
C11orf58	NM_014267	Chromosome 11 open reading frame 58
SERPINB5	NM_002639	Serpin peptidase inhibitor, clade B (ovalbumin), member 5
CPAMD8	AB033109	C3 and PZP-like, alpha-2-macroglobulin domain containing 8
CD8B	NM_004931	CD8b molecule
LCN2	NM_005564	Lipocalin 2 (oncogene 24p3)
IDI1	NM_004508	Isopentenyl-diphosphate delta isomerase 1
OCA2	NM_000275	Oculocutaneous albinism II (pink-eye dilution homolog, mouse)
ATP2C1	NM_014382	ATPase, Ca <sup>++</sup> transporting, type 2C, member 1
APOD	NM_001647	Apolipoprotein D
ATP13A1	NM_020410	ATPase type 13A1
NPTX1	NM_002522	Neuronal pentraxin I
EEF1E1	NM_004280	Thioredoxin domain containing 5
PRKCZ	NM_002744	Protein kinase C, zeta
GIMAP5	NM_018384	GTPase, IMAP family member 5
CKB	NM_001823	Creatine kinase, brain
DIO3	NM_001362	Deiodinase, iodothyronine, type III
CPNE3	NM_003909	Copine III
AGPS	NM_003659	Alkylglycerone phosphate synthase
EIF2S2	NM_003908	Eukaryotic translation initiation factor 2, subunit 2 beta, 38kDa
EEF1A1	NM_001402	Eukaryotic translation elongation factor 1 alpha 1
GMFB	NM_004124	Glia maturation factor, beta
ADH1A	NM_000667	Alcohol dehydrogenase 1C (class I), gamma polypeptide
NDUFA4	NM_002489	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 4, 9kDa
RRM1	NM_001033	Ribonucleotide reductase M1 polypeptide
SFRS2	NM_003016	Splicing factor, arginine/serine-rich 2

SOD1	NM_000454	Superoxide dismutase 1, soluble (amyotrophic lateral sclerosis 1 (adult))
PTS	NM_000317	6-pyruvoyltetrahydropterin synthase
ALDH2	NM_000690	Aldehyde dehydrogenase 2 family (mitochondrial)
PGD	NM_002631	Phosphogluconate dehydrogenase
KIAA1155	AB032981	KIAA1155 protein
C14orf2	NM_004894	Chromosome 14 open reading frame 2
SERINC5	AK026295	Serine incorporator 5
AURKAIP1	NM_017900	Aurora kinase A interacting protein 1
KIAA1432	AB037853	KIAA1432
RNF133	AK022391	Ring finger protein 133
FST	NM_006350	Follistatin
RANGNRF	NM_014185	RAN guanine nucleotide release factor
PDE6H	NM_006205	Phosphodiesterase 6H, cGMP-specific, cone, gamma
ACN9	NM_020186	ACN9 homolog (S. cerevisiae)
HEBP1	NM_015987	Heme binding protein 1
DKFZP564O0823	AK025205	DKFZP564O0823 protein
RGS2	NM_002923	Regulator of G-protein signalling 2, 24kDa
AOC2	NM_009590	Amine oxidase, copper containing 2 (retina-specific)
MORF4L1	NM_006791	Mortality factor 4 like 1
IL1R1	NM_000877	Interleukin 1 receptor, type I
FARP1	NM_005766	FERM, RhoGEF (ARHGEF) and pleckstrin domain protein 1 (chondrocyte-derived)
MLF2	NM_005439	Myeloid leukemia factor 2
KAL1	NM_000216	Kallmann syndrome 1 sequence
GAS1	NM_002048	Growth arrest-specific 1
ALDOA	NM_000034	Aldolase A, fructose-bisphosphate
TUBB4Q	NM_020040	Tubulin, beta polypeptide 4, member Q
SYMPK	NM_004819	Symplekin
FOLR1	NM_016730	Folate receptor 1 (adult)
C20orf3	AB033767	Chromosome 20 open reading frame 3
SLC1A2	NM_004171	Solute carrier family 1 (glial high affinity glutamate transporter), member 2
SEC24D	NM_014822	SEC24 related gene family, member D (S. cerevisiae)
OFD1	NM_003611	Oral-facial-digital syndrome 1
PHLDB1	AB014538	Pleckstrin homology-like domain, family B, member 1
DACT1	NM_016651	Dapper, antagonist of beta-catenin, homolog 1 (Xenopus laevis)
TMEFF2	NM_016192	Transmembrane protein with EGF-like and two follistatin-like domains 2
PAFAH1B3	NM_002573	Platelet-activating factor acetylhydrolase, isoform Ib, gamma subunit 29kDa
RARRES2	NM_002889	Retinoic acid receptor responder (tazarotene induced) 2
PCCA	NM_000282	Propionyl Coenzyme A carboxylase, alpha polypeptide
TFPI2	NM_006528	Tissue factor pathway inhibitor 2
NT5C2	NM_012229	5'-nucleotidase, cytosolic II

RPN1	NM_002950	Ribophorin I
LSM7	NM_016199	LSM7 homolog, U6 small nuclear RNA associated (S. cerevisiae)
DNAJB6	NM_005494	DnaJ (Hsp40) homolog, subfamily B, member 6
C15orf15	NM_016304	Chromosome 15 open reading frame 15
JUNB	NM_002229	Jun B proto-oncogene
RPP14	NM_007042	Ribonuclease P 14kDa subunit
RPL28	NM_000991	Ribosomal protein L28
IDH3A	NM_005530	Isocitrate dehydrogenase 3 (NAD+) alpha
B3GAT3	NM_012200	Beta-1,3-glucuronyltransferase 3 (glucuronosyltransferase I)
ERCC5	NM_000123	Excision repair cross-complementing rodent repair deficiency, complementation group 5 (xeroderma pigmentosum, complementation group G (Cockayne syndrome))
ELF2	NM_006874	E74-like factor 2 (ets domain transcription factor)
ZNF133	NM_003434	Zinc finger protein 133 (clone pHZ-13)
ZNF532	NM_018181	Zinc finger protein 532
LOC130074	AL137406	hypothetical protein LOC130074
IDH3G	NM_004135	Isocitrate dehydrogenase 3 (NAD+) gamma
PRO1598	NM_018503	hypothetical protein PRO1598
LOC283824	AL050125	hypothetical protein LOC283824
MTBP	AK022122	Homo sapiens cDNA FLJ12060 fis, clone HEMBB1002142.
ANG	NM_001145	Ribonuclease, RNase A family, 4
ENY2	NM_020189	Enhancer of yellow 2 homolog (Drosophila)
RAFTLIN	D42043	raft-linking protein
TAOK1	AK025902	TAO kinase 1
RELN	NM_005045	Reelin
TMCC1	AB018322	Transmembrane and coiled-coil domain family 1
FEZ2	NM_005102	Fasciculation and elongation protein zeta 2 (zygin II)
ACTG2	NM_001615	Actin, gamma 2, smooth muscle, enteric
PRELP	NM_002725	Proline/arginine-rich end leucine-rich repeat protein
PPAP2B	NM_003713	Phosphatidic acid phosphatase type 2B
APXL	NM_001649	Apical protein-like (Xenopus laevis)
RAB3GAP1	D31886	RAB3 GTPase activating protein subunit 1 (catalytic)
ECT2	NM_018098	Epithelial cell transforming sequence 2 oncogene
GNB3	NM_002075	Triosephosphate isomerase 1
SCG5	NM_003020	Secretogranin V (7B2 protein)
YWHAB	NM_003404	Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, beta polypeptide
ADCY7	NM_001114	Adenylate cyclase 7
ISLR	NM_005545	Immunoglobulin superfamily containing leucine-rich repeat
ADAM15	NM_003815	ADAM metalloproteinase domain 15 (metargidin)
CRABP1	NM_004378	Cellular retinoic acid binding protein 1
ITPR3	NM_002224	Inositol 1,4,5-triphosphate receptor, type 3
YY1	NM_003403	YY1 transcription factor
PCGF1	AL049296	Polycomb group ring finger 1

MMD	NM_012329	Monocyte to macrophage differentiation-associated
LAMB1	NM_002291	Laminin, beta 1
NGRN	NM_016645	Neugrin, neurite outgrowth associated
RAB28	NM_004249	RAB28, member RAS oncogene family
CMKOR1	U67784	Chemokine orphan receptor 1
COPE	NM_007263	Coatomer protein complex, subunit epsilon
OPN1SW	NM_001708	Opsin 1 (cone pigments), short-wave-sensitive (color blindness, tritan)
ABCD3	NM_002858	ATP-binding cassette, sub-family D (ALD), member 3
TERF2IP	NM_018975	Telomeric repeat binding factor 2, interacting protein
EIF3S6IP	NM_016091	Eukaryotic translation initiation factor 3, subunit 6 interacting protein
GPR143	NM_000273	G protein-coupled receptor 143
SLC16A1	NM_003051	Solute carrier family 16 (monocarboxylic acid transporters), member 1
RAB10	NM_016131	RAB10, member RAS oncogene family
ATP5L	NM_006476	ATP synthase, H <sup>+</sup> transporting, mitochondrial F0 complex, subunit G
CTSK	NM_000396	Aryl hydrocarbon receptor nuclear translocator
AKR1C1	NM_001353	Aldo-keto reductase family 1, member C2 (dihydrodiol dehydrogenase 2; bile acid binding protein; 3-alpha hydroxysteroid dehydrogenase, type III)
USP9X	NM_004652	Ubiquitin specific peptidase 9, X-linked (fat facets-like, Drosophila)
PGCP	NM_006102	Homo sapiens plasma glutamate carboxypeptidase (PGCP), mRNA.
TYR	NM_000372	Tyrosinase (oculocutaneous albinism IA)
HSF1	NM_005526	Heat shock transcription factor 1
PARP1	NM_001618	Poly (ADP-ribose) polymerase family, member 1
PRPF8	NM_006445	PRP8 pre-mRNA processing factor 8 homolog (yeast)
TMEM59	NM_004872	Transmembrane protein 59
EEF1A2	NM_001958	Eukaryotic translation elongation factor 1 alpha 2
CTBP2	NM_001329	C-terminal binding protein 2
RPS3A	NM_001006	Ribosomal protein S3A
F13A1	NM_000129	Coagulation factor XIII, A1 polypeptide
NDUFB2	NM_004546	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2, 8kDa
KIAA0963	NM_014963	KIAA0963
POLR2J	NM_006234	DNA directed RNA polymerase II polypeptide J-related gene
KIAA0265	D87454	KIAA0265 protein
TENC1	AB028998	Tensin like C1 domain containing phosphatase (tensin 2)
MOSC2	NM_017898	MOCO sulphurase C-terminal domain containing 2
ALDH1A3	NM_000693	Aldehyde dehydrogenase 1 family, member A3
GTF3C1	NM_001520	General transcription factor IIIC, polypeptide 1, alpha 220kDa

SAMD14	AF035306	Sterile alpha motif domain containing 14
TUBB2C	NM_006088	Tubulin, beta 2C
PFN2	NM_002628	Profilin 2
RDH8	NM_015725	Retinol dehydrogenase 8 (all-trans)
CLEC1A	NM_016511	C-type lectin domain family 1, member A
CACNA1E	NM_000721	Calcium channel, voltage-dependent, alpha 1E subunit
CD248	NM_020404	CD248 molecule, endosialin
CST3	NM_000099	Cystatin C (amyloid angiopathy and cerebral hemorrhage)
STAT4	NM_003151	Signal transducer and activator of transcription 4
MYL4	NM_002476	Myosin, light polypeptide 4, alkali; atrial, embryonic
HK2	NM_000189	Hexokinase 2
WDR68	NM_005828	WD repeat domain 68
PDZRN4	NM_013377	PDZ domain containing RING finger 4
S100A1	NM_006271	S100 calcium binding protein A1
SH3GL2	NM_003026	SH3-domain GRB2-like 2
LRP1	NM_002332	Low density lipoprotein-related protein 1 (alpha-2-macroglobulin receptor)
DCTN3	NM_007234	Dynactin 3 (p22)
HSPC152	NM_016404	hypothetical protein HSPC152
RHEB	NM_005614	Ras homolog enriched in brain
TGFB1	NM_000358	Transforming growth factor, beta-induced, 68kDa
LEPROTL1	NM_015344	Leptin receptor overlapping transcript-like 1
LAMA2	NM_000426	Laminin, alpha 2 (merosin, congenital muscular dystrophy)
SFT2D1	AK021489	MRNA; cDNA DKFZp667D087 (from clone DKFZp667D087)
WWC1	AB020676	WW, C2 and coiled-coil domain containing 1
GSTT1	NM_000853	Glutathione S-transferase theta 1
HSPA8	NM_006597	Heat shock 70kDa protein 8
BCL2L13	NM_015367	BCL2-like 13 (apoptosis facilitator)
USP18	NM_017414	Ubiquitin specific peptidase 18
SLC7A6	NM_003983	Lysophospholipase 3 (lysosomal phospholipase A2)
KIAA0888	AB020695	KIAA0888 protein
KCNK1	NM_002245	Potassium channel, subfamily K, member 1
TMED2	NM_006815	Transmembrane emp24 domain trafficking protein 2
EIF2S1	NM_004094	Eukaryotic translation initiation factor 2, subunit 1 alpha, 35kDa
PSMD14	NM_005805	Proteasome (prosome, macropain) 26S subunit, non-ATPase, 14
UCHL1	NM_004181	Ubiquitin carboxyl-terminal esterase L1 (ubiquitin thiolesterase)
TIMM17B	NM_005834	Translocase of inner mitochondrial membrane 17 homolog B (yeast)
P8	NM_012385	p8 protein (candidate of metastasis 1)
BACE2	NM_012105	Beta-site APP-cleaving enzyme 2
SOHLH2	NM_017826	Spermatogenesis and oogenesis specific basic helix-loop-helix 2
RPS11	NM_001015	Ribosomal protein S11

SLC25A4	NM_001151	Solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 4
ENO2	NM_001975	Enolase 2 (gamma, neuronal)
MCM6	NM_005915	MCM6 minichromosome maintenance deficient 6 (MIS5 homolog, <i>S. pombe</i> ) ( <i>S. cerevisiae</i> )
TOMM7	NM_019059	Translocase of outer mitochondrial membrane 7 homolog (yeast)
RAD51L3	NM_002878	RAD51-like 3 ( <i>S. cerevisiae</i> )
HMX1	NM_018942	Homeobox (H6 family) 1
SEC14L5	AB007880	SEC14-like 5 ( <i>S. cerevisiae</i> )
LOC144871	AK024941	hypothetical protein LOC144871
INHBB	NM_002193	Inhibin, beta B (activin AB beta polypeptide)
LYNX1	AL353948	Ly6/neurotoxin 1
HEPH	NM_014799	Hephaestin
PDLIM5	AL049969	PDZ and LIM domain 5
CDH11	NM_001797	Cadherin 11, type 2, OB-cadherin (osteoblast)
TARBP1	NM_005646	Tar (HIV-1) RNA binding protein 1
MAOB	NM_000898	Monoamine oxidase B
PRKAR2B	NM_002736	Protein kinase, cAMP-dependent, regulatory, type II, beta
PVALB	NM_002854	Parvalbumin
PRL	NM_000948	Prolactin
SPARC	NM_003118	Secreted protein, acidic, cysteine-rich (osteonectin)
SH3YL1	NM_015677	SH3 domain containing, Ysc84-like 1 ( <i>S. cerevisiae</i> )
GPR133	AL162032	G protein-coupled receptor 133
MYOC	NM_000261	Myocilin, trabecular meshwork inducible glucocorticoid response
CD164	NM_006016	CD164 molecule, sialomucin
FLRT2	NM_013231	Fibronectin leucine rich transmembrane protein 2
IGF2	NM_000612	Insulin-like growth factor 2 (somatomedin A)
NLE1	NM_018096	Notchless homolog 1 ( <i>Drosophila</i> )
ADAM17	NM_003183	ADAM metallopeptidase domain 17 (tumor necrosis factor, alpha, converting enzyme)
COL2A1	NM_001844	Collagen, type II, alpha 1 (primary osteoarthritis, spondyloepiphyseal dysplasia, congenital)
TESC	NM_017899	Tescalcin
RHOBTB3	NM_014899	Rho-related BTB domain containing 3
LMOD1	NM_012134	Leiomodin 1 (smooth muscle)
DDEFL1	NM_017707	Development and differentiation enhancing factor-like 1
ARPC1A	NM_006409	Actin related protein 2/3 complex, subunit 1A, 41kDa
C1RL	NM_016546	Complement component 1, r subcomponent
KIAA0251	D87438	KIAA0251 protein
SUMO1	NM_003352	SMT3 suppressor of mif two 3 homolog 1 (yeast)
GATM	NM_001482	Glycine amidinotransferase (L-arginine:glycine amidinotransferase)
BAMBI	NM_012342	BMP and activin membrane-bound inhibitor homolog ( <i>Xenopus laevis</i> )
CH25H	NM_003956	Cholesterol 25-hydroxylase



ATP5D	NM_001687	ATP synthase, H <sup>+</sup> transporting, mitochondrial F1 complex, delta subunit
VAX2	NM_012476	Ventral anterior homeobox 2
RPL26	NM_000987	Ribosomal protein L26
RPL23A	NM_000984	Ribosomal protein L23a
ERP29	NM_006817	Endoplasmic reticulum protein 29
TCEB2	NM_007108	Transcription elongation factor B (SIII), polypeptide 2 (18kDa, elongin B)
GZMB	NM_004131	Granzyme B (granzyme 2, cytotoxic T-lymphocyte-associated serine esterase 1)
KIAA0284	AB006622	KIAA0284
SMG5	AB029012	Smg-5 homolog, nonsense mediated mRNA decay factor (C. elegans)
LOC56902	NM_020143	putative 28 kDa protein
MIER2	AB033019	KIAA1193
FOXF2	NM_001452	Forkhead box F2
ZNFN1A2	NM_016260	Zinc finger protein, subfamily 1A, 2 (Helios)
ZNF529	AK025110	Zinc finger protein 529
TGFB3	NM_003239	Transforming growth factor, beta 3
IDH2	NM_002168	Isocitrate dehydrogenase 2 (NADP <sup>+</sup> ), mitochondrial
KMO	NM_003679	Kynurenine 3-monooxygenase (kynurenine 3-hydroxylase)
NDUFB5	NM_002492	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 5, 16kDa
NDUFS4	NM_002495	NADH dehydrogenase (ubiquinone) Fe-S protein 4, 18kDa (NADH-coenzyme Q reductase)
EGFR	BF512905	Epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian)
KIAA0241	AA205569	KIAA0241
FKBP9	AW965093	FK506 binding protein 9, 63 kDa
AI422523	AI422523	tf86a06.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:2106130 3', mRNA sequence.
FLJ25076	AK057805	similar to CG4502-PA
FLJ27365	BE877828	FLJ27365 protein
MLL3	BE551763	Myeloid/lymphoid or mixed-lineage leukemia 3
EPPK1	NM_031308	Epiplakin 1
CA3	BG212781	Transcribed locus
BF693375	BF693375	602081004F1 NIH_MGC_81 Homo sapiens cDNA clone IMAGE:4245528 5', mRNA sequence.
AW305170	AW305170	Transcribed locus
DKFZP566N034	AK054738	hypothetical protein DKFZp566N034
AI421736	AI421736	tf39a05.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:2098544 3', mRNA sequence.
TRAK2	AI360167	Trafficking protein, kinesin binding 2
DNAJA3	AF244136	DnaJ (Hsp40) homolog, subfamily A, member 3
EFNA5	BG929325	Nuclear RNA-binding protein, putative

BG054770	BG054770	nac89c10.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:3441426 3' similar to contains Alu repetitive element;contains element MER19 repetitive element ; mRNA sequence.
MYO1E	NM_004998	Myosin IE
AL037421	AL037421	Transcribed locus
C12orf22	AI417295	Transcribed locus
CACNG4	AI953087	Calcium channel, voltage-dependent, gamma subunit 4
COL1A2	NM_000089	Collagen, type I, alpha 2
TBC1D5	AW090566	TBC1 domain family, member 5
CXXC5	BC006428	CXXC finger 5
QKI	AB067801	Quaking homolog, KH domain RNA binding (mouse)
AW090131	AW090131	xc91g04.x1 NCI_CGAP_Brn35 Homo sapiens cDNA clone IMAGE:2591670 3' similar to gb:X14723 CLUSTERIN PRECURSOR (HUMAN);contains PTR5.b3 MSR1 repetitive element ; mRNA sequence.
HLCS	AI089425	Holocarboxylase synthetase (biotin-(propionyl-Coenzyme A-carboxylase (ATP-hydrolysing)) ligase)
CD79B	NM_000626	CD79b molecule, immunoglobulin-associated beta
AL513921	AL513921	AL513921 Homo sapiens PLACENTA Homo sapiens cDNA clone CL0BA006ZG09 3-PRIME, mRNA sequence.
KALRN	H12097	Kalirin, RhoGEF kinase
MS4A6A	NM_022349	Membrane-spanning 4-domains, subfamily A, member 6A
AI983521	AI983521	Transcribed locus
DOCK1	BE223068	Dedicator of cytokinesis 1
R45557	R45557	Transcribed locus
NAP1L5	AK054689	Nucleosome assembly protein 1-like 5
BF594228	BF594228	Transcribed locus
FLJ35390	BC014556	hypothetical protein FLJ35390
RAB38	NM_022337	RAB38, member RAS oncogene family
BE261437	BE261437	601146805F1 NIH_MGC_19 Homo sapiens cDNA clone IMAGE:3162312 5', mRNA sequence.
HIST3H2A	NM_033445	Histone 3, H2a
DKFZP686E2158	BC014242	hypothetical protein LOC643155
NUSAP1	BC016329	Nucleolar and spindle associated protein 1
RSAFD1	AA757250	Radical S-adenosyl methionine and flavodoxin domains 1
AA179146	AA179146	zp12a01.s1 Stratagene fetal retina 937202 Homo sapiens cDNA clone IMAGE:609192 3', mRNA sequence.
TRIM63	AF361946	Homo sapiens striated muscle RING zinc finger protein SMRZ mRNA, complete cds.
SLC25A37	NM_016612	Solute carrier family 25, member 37
C4A	AK054612	Complement component 4A (Rodgers blood group)
IGL@	AK057065	Immunoglobulin lambda locus
RNF20	BG105573	Ring finger protein 20

IGFBP5	L27560	Insulin-like growth factor binding protein 5
TSEN34	AA176280	TRNA splicing endonuclease 34 homolog (S. cerevisiae)
SYNC1	NM_030786	Syncoilin, intermediate filament 1
SESN3	AL598437	MRNA; cDNA DKFZp313B1017 (from clone DKFZp313B1017)
SCN1A	AW157289	Sodium channel, voltage-gated, type I, alpha
LRIG1	AB050468	Leucine-rich repeats and immunoglobulin-like domains 1
ELAVL1	BC003376	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R)
SLC24A6	NM_024959	Solute carrier family 24 (sodium/potassium/calcium exchanger), member 6
IGSF9	AK056650	Immunoglobulin superfamily, member 9
SMARCAL1	BG204606	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a-like 1
SNX10	BG208557	Sorting nexin 10
UCHL1	AK055249	Ubiquitin carboxyl-terminal esterase L1 (ubiquitin thiolesterase)
TANC1	AB051515	Tetratricopeptide repeat, ankyrin repeat and coiled-coil containing 1
FLJ30672	AK055382	hypothetical protein FLJ30672
BTBD6	AF353674	BTB (POZ) domain containing 6
DSCAML1	R59091	Transcribed locus
DOK6	AK057795	Docking protein 6
ABCA10	NM_080282	ATP-binding cassette, sub-family A (ABC1), member 10
AASDH	BG681142	2-aminoadipic 6-semialdehyde dehydrogenase
KIAA1754	AB051541	KIAA1754
TPSAB1	NM_003294	Tryptase beta 2
HLA-DOA	NM_002119	Major histocompatibility complex, class II, DP alpha 1
COL9A2	NM_001852	Collagen, type IX, alpha 2
F33254	F33254	HSPD26568 HM3 Homo sapiens cDNA clone s3000057H05, mRNA sequence.
TRIM39	BC007661	Tripartite motif-containing 39
TCTEX1D1	AI421187	Tctex1 domain containing 1
COTL1	BC010039	Coactosin-like 1 (Dictyostelium)
MORN1	NM_024848	MORN repeat containing 1
INT1	BE048245	integrator complex subunit 1
RAB18	NM_021252	RAB18, member RAS oncogene family
PCP4L1	W22165	Purkinje cell protein 4 like 1
BI522966	BI522966	603175461R1 NIH_MGC_121 Homo sapiens cDNA clone IMAGE:5239791 3', mRNA sequence.
MIZF	NM_015517	MBD2-interacting zinc finger
DIXDC1	BI914728	DIX domain containing 1
CDH23	NM_022124	Cadherin-like 23
MAP4	H11737	Microtubule-associated protein 4

AW971942	AW971942	Transcribed locus, weakly similar to NP_872601.1 tetratricopeptide repeat protein isoform 1; ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome; ubiquitously transcribed TPR gene on Y chromosome [Homo sapiens]
ERCC5	AI368638	Excision repair cross-complementing rodent repair deficiency, complementation group 5 (xeroderma pigmentosum, complementation group G (Cockayne syndrome))
BRD3	AW138687	Bromodomain containing 3
NF2	AI937250	Neurofibromin 2 (bilateral acoustic neuroma)
CRTC1	NM_025021	CREB regulated transcription coactivator 1
PRKRIR	AF081567	Protein-kinase, interferon-inducible double stranded RNA dependent inhibitor, repressor of (P58 repressor)
AL534569	AL534569	Transcribed locus
LEPR	U50748	Leptin receptor
KITLG	AK055903	KIT ligand
C1orf54	NM_024579	Chromosome 1 open reading frame 54
AI114492	AI114492	Transcribed locus
AK055112	AK055112	3'UTR of hypothetical protein (ORF1)
AI792841	AI792841	qv38c04.y5 NCI_CGAP_Ov31 Homo sapiens cDNA clone IMAGE:1983846 5', mRNA sequence.
ATP2B2	AW090039	ATPase, Ca <sup>++</sup> transporting, plasma membrane 2
TBN	AK057383	Taube nuss homolog (mouse)
UCK2	AL563776	Uridine-cytidine kinase 2
SPACA1	NM_030960	Sperm acrosome associated 1
T26536	T26536	AB300G7F Infant brain, LLNL array of Dr. M. Soares 1NIB Homo sapiens cDNA clone LLAB300G7 3', mRNA sequence.
COX7CP1	AF042165	Homo sapiens cytochrome c oxidase subunit VIIc (COX7CP1) pseudogene, complete sequence.
LOC440065	N63497	Transcribed locus, strongly similar to XP_498526.1 PREDICTED: hypothetical protein XP_498526 [Homo sapiens]
AW088087	AW088087	xc97d07.x1 NCI_CGAP_Brn35 Homo sapiens cDNA clone IMAGE:2592205 3' similar to contains PTR5.b2 MER22 repetitive element ; mRNA sequence.
AI870750	AI870750	wl76d11.x1 NCI_CGAP_Brn25 Homo sapiens cDNA clone IMAGE:2430837 3', mRNA sequence.
KCNC2	AF268897	Homo sapiens voltage gated potassium channel Kv3.2a mRNA, complete cds; alternatively spliced.
PLAGL1	AI953762	Pleiomorphic adenoma gene-like 1
DCHS1	NM_003737	Dachsous 1 (Drosophila)
C19orf23	BE393028	Chromosome 19 open reading frame 23
OPA3	NM_025136	Optic atrophy 3 (autosomal recessive, with chorea and spastic paraplegia)
SHC4	AL035883	Transcribed locus
IGFBP6	NM_002178	Insulin-like growth factor binding protein 6
PCM1	BC000453	Pericentriolar material 1

AL120181	AL120181	Full length insert cDNA clone YY86C01
AW264180	AW264180	xq97a02.x1 NCI_CGAP_Brn53 Homo sapiens cDNA clone IMAGE:2758538 3', mRNA sequence.
AW969067	AW969067	Transcribed locus, weakly similar to XP_342865.1 PREDICTED: similar to LRRG00116 [Rattus norvegicus]
BE260094	BE260094	601146938F1 NIH_MGC_19 Homo sapiens cDNA clone IMAGE:3162369 5', mRNA sequence.
AK001536	AK001536	CDNA FLJ10674 fis, clone NT2RP2006436
GIMAP5	AI587192	GTPase, IMAP family member 5
BG819298	BG819298	602781479F1 NCI_CGAP_Brn67 Homo sapiens cDNA clone IMAGE:4932306 5', mRNA sequence.
AP1G1	AI939422	Adaptor-related protein complex 1, gamma 1 subunit
BG190000	BG190000	Transcribed locus
SMOC2	AJ420521	SPARC related modular calcium binding 2
TRIM37	NM_015294	Tripartite motif-containing 37
R43203	R43203	yg18h01.s1 Soares infant brain 1NIB Homo sapiens cDNA clone IMAGE:32865 3', mRNA sequence.
AW148831	AW148831	Transcribed locus
SCGB1D2	NM_006551	Secretoglobin, family 1D, member 2
CHPF	NM_024536	chondroitin polymerizing factor
ENSA	AF067170	Endosulfine alpha
ABCA6	AY028898	ATP-binding cassette, sub-family A (ABC1), member 6
UNQ9217	BG208711	AASA9217
ACBD6	NM_032360	Acyl-Coenzyme A binding domain containing 6
GNGT2	BG396167	Guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 2
ABI3BP	NM_015429	ABI gene family, member 3 (NESH) binding protein
INPP4B	AI953893	Inositol polyphosphate-4-phosphatase, type II, 105kDa
YIPF6	BC012469	Yip1 domain family, member 6
CRR9	NM_030782	cisplatin resistance related protein CRR9p
FMNL2	AB067489	Formin-like 2
LOC388170	AI557230	Transcribed locus, strongly similar to XP_373646.2 PREDICTED: hypothetical protein XP_373646 [Homo sapiens]
MOV10L1	NM_018995	Mov10l1, Moloney leukemia virus 10-like 1, homolog (mouse)
CRSP3	BG149642	Cofactor required for Sp1 transcriptional activation, subunit 3, 130kDa
AKAP12	BF035258	A kinase (PRKA) anchor protein (gravin) 12
TPSB2	NM_024164	Tryptase beta 2
TEAD2	BC007556	TEA domain family member 2
DDR1	NM_001954	Discoidin domain receptor family, member 1
TAF3	AW005174	TAF3 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 140kDa
R13901	R13901	yf62b07.r1 Soares infant brain 1NIB Homo sapiens cDNA clone IMAGE:26699 5', mRNA sequence.

ATP2B4	AW888226	ATPase, Ca <sup>++</sup> transporting, plasma membrane 4
AI860987	AI860987	Transcribed locus, strongly similar to XP_542173.2 PREDICTED: similar to CG1676-PA [Canis familiaris]
CBR3	NM_001236	Carbonyl reductase 3
ITGB1	NM_002211	Integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)
AI146787	AI146787	Transcribed locus
T02973	T02973	Transcribed locus
VIT	AK056772	Vitrin
AK055040	AK055040	CDNA FLJ30478 fis, clone BRAWH1000167
AQP9	NM_020980	Aquaporin 9
PLEKHA2	AK057205	Pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 2
LAMC1	NM_002293	Laminin, gamma 1 (formerly LAMB2)
BG212324	BG212324	Transcribed locus
ZNF713	AA774885	Zinc finger protein 713
AV723968	AV723968	Transcribed locus
BG254392	BG254392	602368978F1 NIH_MGC_91 Homo sapiens cDNA clone IMAGE:4477195 5', mRNA sequence.
RBM35B	NM_024939	RNA binding motif protein 35B
FAM33A	AI792525	Family with sequence similarity 33, member A
TMTC1	BC017984	Transmembrane and tetratricopeptide repeat containing 1
MGC4562	BC014124	hypothetical protein MGC4562
TMED10	AK056156	Transmembrane emp24-like trafficking protein 10 (yeast)
N47786	N47786	Transcribed locus
SKP1A	NM_006930	S-phase kinase-associated protein 1A (p19A)
KRTAP4-7	NM_033061	Keratin associated protein 4-7
AI361968	AI361968	qy37g09.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:2014240 3', mRNA sequence.
TRAK1	AK023962	Trafficking protein, kinesin binding 1
WDR22	AB058727	WD repeat domain 22
SHANK3	BF986692	SH3 and multiple ankyrin repeat domains 3
SUMF1	AK057983	Sulfatase modifying factor 1
AW662487	AW662487	hi28b03.x1 NCI_CGAP_Co14 Homo sapiens cDNA clone IMAGE:2973581 3' similar to contains Alu repetitive element;contains element MER26 repetitive element ; mRNA sequence.
MRLC2	NM_033546	myosin regulatory light chain MRLC2
LTBP3	NM_021070	Latent transforming growth factor beta binding protein 3
ZNF503	NM_032772	Zinc finger protein 503
EGFR	AK026818	Epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian)
PPP1R12B	R41870	Protein phosphatase 1, regulatory (inhibitor) subunit 12B
BIC	AF402776	BIC transcript

TSPAN16	NM_012466	Protein kinase C substrate 80K-H
AI870981	AI870981	Transcribed locus
PRR14	NM_024031	Hypothetical protein MGC3121
LOC285550	BC017971	hypothetical protein LOC285550
PLXNC1	AF035307	Plexin C1
VPS35	NM_018206	Vacuolar protein sorting 35 (yeast)
TRIP6	AF000974	Thyroid hormone receptor interactor 6
MUC20	AI457171	Mucin 20, cell surface associated
TTRAP	AW972292	TRAF and TNF receptor associated protein
RBJ	AL117490	Ras-associated protein Rap1
RABEP1	AI806781	Rabaptin, RAB GTPase binding effector protein 1
TRIOBP	NM_007032	TRIO and F-actin binding protein
GLUL	NM_002065	Glutamate-ammonia ligase (glutamine synthetase)
RNF187	BC012758	Ring finger protein 187
TCEAL2	NM_080390	Transcription elongation factor A (SII)-like 2
MRPL13	NM_014078	Mitochondrial ribosomal protein L13
MLR2	AB058698	ligand-dependent corepressor
HSPB6	AK056951	Heat shock protein, alpha-crystallin-related, B6
AA218641	AA218641	Transcribed locus
MESDC1	NM_022566	Mesoderm development candidate 1
FAM103A1	NM_031452	Family with sequence similarity 103, member A1
MIB2	BC016490	Mindbomb homolog 2 (Drosophila)
MT1E	BG191659	Metallothionein 1E (functional)
CALD1	AJ223812	Caldesmon 1
NEDD4L	AA019094	Neural precursor cell expressed, developmentally down-regulated 4-like
AI003379	AI003379	Transcribed locus
MGC10471	NM_030818	hypothetical protein MGC10471
1-sep	NM_052838	Septin 1
C4orf15	BF960556	Chromosome 4 open reading frame 15
BG192694	BG192694	Transcribed locus
PCDH18	NM_019035	Protocadherin 18
FLJ31306	AK055868	hypothetical protein FLJ31306
BE967835	BE967835	601649073R2 NIH_MGC_73 Homo sapiens cDNA clone IMAGE:3932826 3', mRNA sequence.
SLC35D2	AJ005866	Solute carrier family 35, member D2
SYT4	AB037763	Synaptotagmin IV
LOC442075	BG204924	weakly similar to serine/threonine protein kinase Kp78
AI884753	AI884753	wl84c10.x1 NCI_CGAP_Brn25 Homo sapiens cDNA clone IMAGE:2431602 3', mRNA sequence.
LOC284033	BF679019	hypothetical protein LOC284033
CCDC81	NM_021827	Coiled-coil domain containing 81
UGCG	AJ420423	UDP-glucose ceramide glucosyltransferase
MAN1B1	NM_007230	Homo sapiens mannosidase, alpha, class 1B, member 1 (MAN1B1), mRNA.
LOC441351	AA046461	CDNA clone IMAGE:4797645
TSSK1	NM_032028	Testis-specific serine kinase 1

RLTPR	BE312807	RGD, leucine-rich repeat, tropomodulin and proline-rich containing protein
ACOT8	AI885067	Acyl-CoA thioesterase 8
RMND5B	AI992288	Transcribed locus
TMEM68	AK056932	Transmembrane protein 68
SYDE1	NM_033025	Synapse defective 1, Rho GTPase, homolog 1 (C. elegans)
LOC201895	BC008502	hypothetical protein LOC201895
LOC149194	AK057535	hypothetical protein LOC149194
FLJ13195	NM_022906	Homo sapiens hypothetical protein FLJ13195 similar to stromal antigen 3 (FLJ13195), mRNA.
MPPE1	AI582762	Metallophosphoesterase 1
GNB4	NM_021629	Guanine nucleotide binding protein (G protein), beta polypeptide 4
CNTNAP2	AA174106	Contactin associated protein-like 2
PLAGL1	U72621	Pleiomorphic adenoma gene-like 1
COL18A1	NM_030582	Collagen, type XVIII, alpha 1
MPZL1	NM_024569	Myelin protein zero-like 1
BC008122	BC008122	Homo sapiens, clone MGC:18053 IMAGE:4148889, mRNA, complete cds.
PLEKHA4	NM_020904	Pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 4
BI545849	BI545849	Transcribed locus
RAB10	AA017090	RAB10, member RAS oncogene family
GLCCI1	AK055741	Glucocorticoid induced transcript 1
AW157250	AW157250	au93d09.x1 Schneider fetal brain 00004 Homo sapiens cDNA clone IMAGE:2783825 3' similar to gb:L05093 60S RIBOSOMAL PROTEIN L18A (HUMAN);contains PTR5.b3 MSR1 repetitive element ; mRNA sequence.
GPHN	NM_020806	Gephyrin
C5orf23	NM_024563	Hypothetical protein FLJ14054
ECHDC3	NM_024693	Enoyl Coenzyme A hydratase domain containing 3
IQCH	N59381	Transcribed locus
CSMD2	AI929247	CUB and Sushi multiple domains 2
KCNA2	BF513715	Transcribed locus
CDKL1	NM_004196	Cyclin-dependent kinase-like 1 (CDC2-related kinase)
BC012900	BC012900	Homo sapiens, clone IMAGE:3881549, mRNA
LOC442165	BI753564	Transcribed locus, strongly similar to XP_496797.1 PREDICTED: similar to Keratin, type I cytoskeletal 18 (Cytokeratin 18) (K18) (CK 18) [Homo sapiens]
AI739080	AI739080	Transcribed locus, moderately similar to XP_573212.1 PREDICTED: hypothetical protein XP_573212 [Rattus norvegicus]
C14orf78	BC011859	Chromosome 14 open reading frame 78
ARD1A	NM_003491	ARD1 homolog A, N-acetyltransferase (S. cerevisiae)
RSPO3	NM_032784	R-spondin 3 homolog (Xenopus laevis)
EPS8L2	NM_022772	EPS8-like 2



PLEKHB2	AK054998	Pleckstrin homology domain containing, family B (evectins) member 2
LOC348801	AK056320	hypothetical protein LOC348801
FAM38B	NM_022068	Family with sequence similarity 38, member B
BC007261	BC007261	Homo sapiens, clone MGC:15545 IMAGE:3050745, mRNA, complete cds.
TNNT1	BG217313	Troponin T type 1 (skeletal, slow)
SMC4L1	BG943520	SMC4 structural maintenance of chromosomes 4-like 1 (yeast)
AA020740	AA020740	Transcribed locus, moderately similar to NP_694957.1 RPE-spondin [Homo sapiens]
PDGFD	NM_025208	Platelet derived growth factor D
FLJ33534	Z40688	hypothetical protein FLJ33534
HTRA3	AY040094	HtrA serine peptidase 3
KCNMA1	N51068	Potassium large conductance calcium-activated channel, subfamily M, alpha member 1
KIF3C	AF035621	Kinesin family member 3C
FKRP	NM_024301	Fukutin related protein
COL6A2	NM_001849	Collagen, type VI, alpha 2
VPS37A	AK057204	Vacuolar protein sorting 37A (yeast)
EMX2	NM_004098	Empty spiracles homolog 2 (Drosophila)
GNGT2	NM_031498	Guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 2
VANGL1	NM_024062	Homo sapiens vang-like 1 (van gogh, Drosophila) (VANGL1), mRNA.
C1orf36	AV721413	Chromosome 1 open reading frame 36
STAG2	NM_006603	Stromal antigen 2
RAP1GDS1	NM_021159	RAP1, GTP-GDP dissociation stimulator 1
KRTHB6	NM_002284	Keratin, hair, basic, 6 (monilethrix)
C2orf32	BC011535	Chromosome 2 open reading frame 32
NEIL1	AK026216	Nei endonuclease VIII-like 1 (E. coli)
CREG2	BF966862	Cellular repressor of E1A-stimulated genes 2
FLJ22655	NM_024730	hypothetical protein FLJ22655
BAD	NM_004322	BCL2-antagonist of cell death
KIAA0773	BG026700	KIAA0773 gene product
CNDP1	NM_032649	Carnosine dipeptidase 1 (metallopeptidase M20 family)
BG335831	BG335831	602404536F1 NIH_MGC_21 Homo sapiens cDNA clone IMAGE:4542008 5', mRNA sequence.
MCF2L	AK057899	Homo sapiens cDNA FLJ25170 fis, clone CBR08752.
NPHP3	AI200954	Nephronophthisis 3 (adolescent)
MSI2	AI949570	Transcribed locus
TOLLIP	AL136835	Toll interacting protein
ANKS6	BF312789	Ankyrin repeat and sterile alpha motif domain containing 6
F11R	AF172398	F11 receptor
SOX4	AJ420500	SRY (sex determining region Y)-box 4
FLJ11895	NM_032161	Homo sapiens hypothetical protein FLJ11895 (FLJ11895), mRNA.

AI241800	AI241800	qu72a04.x1 NCI_CGAP_Brn35 Homo sapiens cDNA clone IMAGE:1977582 3', mRNA sequence.
C3orf58	BI552539	Chromosome 3 open reading frame 58
GPR39	AA167540	Transcribed locus
PIK3C2G	NM_004570	Phosphoinositide-3-kinase, class 2, gamma polypeptide
BRD3	NM_007371	Bromodomain containing 3
SS18	NM_005637	Synovial sarcoma translocation, chromosome 18
AW088056	AW088056	Transcribed locus
AA020917	AA020917	ze64a10.r1 Soares retina N2b4HR Homo sapiens cDNA clone IMAGE:363738 5', mRNA sequence.
CCDC12	AK055537	Coiled-coil domain containing 12
AK055418	AK055418	CDNA FLJ30856 fis, clone FEBRA2003258
DCT	NM_001922	Dopachrome tautomerase (dopachrome delta-isomerase, tyrosine-related protein 2)
LRRC29	AA167217	Leucine rich repeat containing 29
CCDC74A	BC016861	Coiled-coil domain containing 74A
YIPF5	NM_030799	Yip1 domain family, member 5
LOC146177	BI916787	hypothetical protein LOC146177
CYBRD1	AL136693	Cytochrome b reductase 1
HP	NM_005143	Haptoglobin
LOC441434	BI497434	Homo sapiens, clone IMAGE:5213378, mRNA
DIO2	NM_000793	Deiodinase, iodothyronine, type II
ADH1C	NM_000669	Alcohol dehydrogenase 1C (class I), gamma polypeptide
R45266	R45266	Transcribed locus
BF243303	BF243303	601878855F1 NIH_MGC_55 Homo sapiens cDNA clone IMAGE:4107828 5', mRNA sequence.
MFAP3L	NM_021647	Microfibrillar-associated protein 3-like
PRO3121	AF130082	Homo sapiens clone FLC1492 PRO3121 mRNA, complete cds.
C14orf121	BF346216	Chromosome 14 open reading frame 121
TMEM50B	AL355685	Transmembrane protein 50B
lactate dehydrogenase B	BC008952	Homo sapiens, lactate dehydrogenase B, clone MGC:3600 IMAGE:3028947, mRNA, complete cds.
AA021278	AA021278	Transcribed locus
RPH3AL	AK054983	Rabphilin 3A-like (without C2 domains)
SLC6A15	AI368909	Solute carrier family 6, member 15
IGHG1	AF067420	Immunoglobulin heavy constant gamma 1 (G1m marker)
FMN2	AI356773	Formin 2
AB019409	AB019409	Homo sapiens mRNA, expressed in fibroblasts of periodontal ligament, complete cds, clone:PDL-108.
CHRD12	NM_015424	Chordin-like 2
BE856278	BE856278	7f91g02.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:3304370 3' similar to contains L1.b1 L1 repetitive element ; mRNA sequence.
DNAH8	NM_001371	Dynein, axonemal, heavy polypeptide 8
LOC339483	AK055560	hypothetical LOC339483

OSBPL8	AI869983	Oxysterol binding protein-like 8
C14orf118	BI550029	Chromosome 14 open reading frame 118
COL6A1	NM_001848	Collagen, type VI, alpha 1
ASB13	NM_024701	Ankyrin repeat and SOCS box-containing 13
KIAA0409	AL515775	KIAA0409
RGS10	NM_002925	Regulator of G-protein signalling 10
CCDC102B	NM_024781	Chromosome 18 open reading frame 14
AA481806	AA481806	aa93f03.s1 Stratagene fetal retina 937202 Homo sapiens cDNA clone IMAGE:838877 3', mRNA sequence.
SLC35F4	H15157	Transcribed locus, strongly similar to XP_292260.4 PREDICTED: similar to solute carrier family 35, member F3 [Homo sapiens]
NRCAM	R40058	Neuronal cell adhesion molecule
CTNNA2	AA081905	Catenin (cadherin-associated protein), alpha 2
BF437925	BF437925	Transcribed locus
FLJ34870	AK057980	FLJ34870 protein
AI732969	AI732969	Transcribed locus
TNS1	AW888222	Tensin 1
GRIK5	AA977136	Glutamate receptor, ionotropic, kainate 5
SLC5A6	AL096737	Solute carrier family 5 (sodium-dependent vitamin transporter), member 6
ZNF283	AW080602	Zinc finger protein 283
MFAP4	L38486	Microfibrillar-associated protein 4
DPYSL3	AL134820	Dihydropyrimidinase-like 3
MTPN	AK055660	Myotrophin
C3orf26	F33061	Transcribed locus
LOC285711	BF511077	hypothetical LOC285711
DDX24	AL119608	DEAD (Asp-Glu-Ala-Asp) box polypeptide 24
CCT5	BC002971	Chaperonin containing TCP1, subunit 5 (epsilon)
BG185052	BG185052	RST3990 Athersys RAGE Library Homo sapiens cDNA, mRNA sequence.
CDC34	NM_004359	Cell division cycle 34
SYNPR	BI755007	Synaptoporin
C3orf52	NM_024616	Chromosome 3 open reading frame 52
AW148846	AW148846	xf05c09.x1 NCI_CGAP_Brn35 Homo sapiens cDNA clone IMAGE:2617168 3', mRNA sequence.
CES1	NM_001266	Carboxylesterase 1 (monocyte/macrophage serine esterase 1)
ARMC9	NM_025139	Armadillo repeat containing 9
OLFM2	NM_058164	Olfactomedin 2
TMTC3	BF115629	Transmembrane and tetratricopeptide repeat containing 3
LOC400614	BG985119	Transcribed locus, strongly similar to XP_378693.2 PREDICTED: hypothetical protein XP_378693 [Homo sapiens]
WDR19	NM_025132	WD repeat domain 19
SEMA3A	BG251582	Transcribed locus
DYNC2LI1	AW131874	Dynein, cytoplasmic 2, light intermediate chain 1

NCOA5	NM_020967	Nuclear receptor coactivator 5
BI554066	BI554066	603235466F1 NCI_CGAP_Brn67 Homo sapiens cDNA clone IMAGE:5309537 5', mRNA sequence.
POR	AF258341	P450 (cytochrome) oxidoreductase
MGC39497	BG201172	hypothetical protein MGC39497
TIGA1	NM_053000	TIGA1
KSP37	NM_031950	Ksp37 protein
MAF	AF055376	V-maf musculoaponeurotic fibrosarcoma oncogene homolog (avian)
BG927468	BG927468	Transcribed locus, strongly similar to NP_115626.1 ankyrin repeat domain 20A [Homo sapiens]
FBXW10	NM_031456	F-box and WD-40 domain protein 10
TCR-alpha chain	M12959	Human mRNA for T-cell receptor alpha chain (TCR-alpha).
NTNG1	AV722613	Netrin G1
BG205612	BG205612	Transcribed locus
MGC4655	NM_033309	hypothetical protein MGC4655
ZNF704	BC004287	Zinc finger protein 704
TGOLN2	AK025557	Trans-golgi network protein 2
CD59	AL049957	CD59 molecule, complement regulatory protein
HOM-TES-103	NM_015438	HOM-TES-103 tumor antigen-like
RHBDF1	NM_022450	Rhomboid 5 homolog 1 (Drosophila)
HSPA1A	NM_005345	Heat shock 70kDa protein 1A
RTBDN	NM_031429	Retbindin
NDRG4	NM_020465	NDRG family member 4
NEFH	NM_021076	Neurofilament, heavy polypeptide 200kDa
LOC84661	NM_032574	dpy-30-like protein
SGCE	NM_003919	Sarcoglycan, epsilon
PLXNA1	NM_032242	Plexin A1
TTY14	NM_031932	Homo sapiens testis transcript Y 14 (TTY14), mRNA.
GBP2	NM_004120	Guanylate binding protein 2, interferon-inducible
MGC5139	NM_032661	Homo sapiens hypothetical protein MGC5139 (MGC5139), mRNA.
FLJ22332	NM_024724	Homo sapiens hypothetical protein FLJ22332 (FLJ22332), mRNA.
CNTD2	NM_024877	V-akt murine thymoma viral oncogene homolog 2
SIVA	NM_006427	CD27-binding (Siva) protein
GUP1	NM_020707	GUP1, glycerol uptake/transporter homolog (yeast)
NKAP	NM_024528	NF-kappaB activating protein
FAM57B	NM_031478	Family with sequence similarity 57, member B
AMY1A	NM_004038	Amylase, alpha 1A; salivary
CKMT1B	NM_020990	Creatine kinase, mitochondrial 1A
FHL2	NM_001450	Four and a half LIM domains 2
CANX	NM_001746	Calnexin
MAP2K2	NM_030662	Mitogen-activated protein kinase kinase 2
RTN1	NM_021136	Reticulon 1
LIN7B	NM_022165	Lin-7 homolog B (C. elegans)
FKBP10	NM_021939	FK506 binding protein 10, 65 kDa

C15orf48	NM_032413	Chromosome 15 open reading frame 48
ADAMTS1	NM_006988	ADAM metallopeptidase with thrombospondin type 1 motif, 1
SVIL	NM_021738	Supervillin
SNRPN	NM_022807	Small nuclear ribonucleoprotein polypeptide N
DPYSL4	NM_006426	Dihydropyrimidinase-like 4
TUBA6	NM_032704	Tubulin, alpha 6
C3orf14	NM_020685	Chromosome 3 open reading frame 14
C1QTNF3	NM_030945	C1q and tumor necrosis factor related protein 3
HIF3A	NM_022462	Hypoxia inducible factor 3, alpha subunit
HNRPH3	NM_012207	Heterogeneous nuclear ribonucleoprotein H3 (2H9)
MCM7	NM_005916	MCM7 minichromosome maintenance deficient 7 (S. cerevisiae)
CD59	NM_000611	CD59 molecule, complement regulatory protein
ZNF179	NM_007148	Zinc finger protein 179
PTTG1IP	NM_004339	Pituitary tumor-transforming 1 interacting protein
AVPI1	NM_021732	Arginine vasopressin-induced 1
HIST1H3F	NM_021018	Histone 1, H3f
GNGT1	NM_021955	Guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 1
GSG1	NM_031289	Germ cell associated 1
FBXO18	NM_032807	F-box protein, helicase, 18
SPATA20	NM_022827	Spermatogenesis associated 20
BXDC5	NM_025065	Brix domain containing 5
LGALS3	NM_002306	Lectin, galactoside-binding, soluble, 3 (galectin 3)
C6orf62	NM_030939	TRAF and TNF receptor associated protein
CHCHD6	NM_032343	Coiled-coil-helix-coiled-coil-helix domain containing 6
MYO10	NM_012334	Myosin X
TF	NM_001063	Transferrin
RHOA	NM_001664	Ras homolog gene family, member A
MFRP	NM_031433	Melanoma cell adhesion molecule
RAP1A	NM_002884	RAP1A, member of RAS oncogene family
FLJ14327	NM_024912	Homo sapiens hypothetical protein FLJ14327 (FLJ14327), mRNA.
MXRA8	NM_032348	Matrix-remodelling associated 8
MRPS35	NM_021821	Mitochondrial ribosomal protein S35
IFI6	NM_022873	Interferon, alpha-inducible protein (clone IFI-6-16)
GLT8D2	NM_031302	Glycosyltransferase 8 domain containing 2
THOC7	NM_025075	THO complex 7 homolog (Drosophila)
BCDO2	NM_031938	Beta-carotene dioxygenase 2
MDS025	NM_021825	hypothetical protein MDS025
MAP2K1IP1	NM_021970	Mitogen-activated protein kinase kinase 1 interacting protein 1
CPSF3L	NM_032179	Cleavage and polyadenylation specific factor 3-like
MGC13114	NM_032366	hypothetical protein MGC13114
AMY2B	NM_020978	Amylase, alpha 1A; salivary
TPT1	NM_003295	Tumor protein, translationally-controlled 1

THRA	NM_003250	Thyroid hormone receptor, alpha (erythroblastic leukemia viral (v-erb-a) oncogene homolog, avian)
FAM117A	NM_030802	C/EBP-induced protein
PRDX2	NM_005809	Peroxiredoxin 2
NCOA4	NM_005437	Nuclear receptor coactivator 4
HAMP	NM_021175	Hepcidin antimicrobial peptide
FAM111A	NM_022074	Family with sequence similarity 111, member A
CHMP5	NM_016410	Chromatin modifying protein 5
IGFBP5	NM_000599	Insulin-like growth factor binding protein 5
FGFR1	NM_023109	Fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2, Pfeiffer syndrome)
AZI2	NM_022461	5-azacytidine induced 2
WFDC1	NM_021197	WAP four-disulfide core domain 1
CRISPLD2	NM_031476	Cysteine-rich secretory protein LCCL domain containing 2
NRD1	NM_002525	Nardilysin (N-arginine dibasic convertase)
PGCP	NM_016134	plasma glutamate carboxypeptidase
HMOX2	NM_002134	Heme oxygenase (decycling) 2
LSMD1	NM_032356	LSM domain containing 1
EEF2	NM_001961	Eukaryotic translation elongation factor 2
PFDN2	NM_012394	Prefoldin subunit 2
TMEM147	NM_032635	Seven transmembrane domain protein
BSCL2	NM_032667	Bernardinelli-Seip congenital lipodystrophy 2 (seipin)
JUND	NM_005354	Jun D proto-oncogene
STK24	NM_003576	Serine/threonine kinase 24 (STE20 homolog, yeast)
MGC4825	NM_024122	hypothetical protein MGC4825
RPL18A	NM_000980	Ribosomal protein L18a
CUL4B	NM_003588	Cullin 4B
MRPS26	NM_030811	Mitochondrial ribosomal protein S26
ACTR2	NM_005722	ARP2 actin-related protein 2 homolog (yeast)
OPN1LW	NM_020061	Opsin 1 (cone pigments), long-wave-sensitive (color blindness, protan)
FXYD6	NM_022003	FXYD domain containing ion transport regulator 6
RPL22	NM_000983	Ribosomal protein L22
USMG5	NM_032747	Upregulated during skeletal muscle growth 5 homolog (mouse)
PIN1	NM_006221	Protein (peptidylprolyl cis/trans isomerase) NIMA-interacting 1
COL1A1	NM_000088	Collagen, type I, alpha 1
LRTM1	NM_020678	Leucine-rich repeats and transmembrane domains 1
INA	NM_032727	Internexin neuronal intermediate filament protein, alpha
PLCD4	NM_032726	Phospholipase C, delta 4
HBB	NM_000518	Hemoglobin, beta
LBH	NM_030915	hypothetical protein DKFZp566J091
RPS13	NM_001017	Ribosomal protein S13