

Cluster ID	Database	Accession	Entrez	Definition
212	U62794	31451_at		Cluster Incl. U62794:Human CDC42 GAP-related protein mRNA, partial cds /cds=(0,671) /gb=U62794 /gi=2326170 /ug=Hs.166138 /len=1378
213	U91640	31452_at		Cluster Incl. U91640:Human survival motor neuron pseudogene /cds=(0,849) /gb=U91640 /gi=2351795 /ug=Hs.166140 /len=850
214	AF025530	31453_s_at		Cluster Incl. AF025530:Homo sapiens leucocyte immunoglobulin-like receptor-6a (LIR-6) mRNA, complete cds /cds=(182,1651) /gb=AF025530 /gi=2653866 /ug=Hs.166156 /len=1910
215	AF034209	31454_f_at		Cluster Incl. AF034209:Homo sapiens RIG-like 5-6 mRNA, complete cds /cds=(720,1070) /gb=AF034209 /gi=2724107 /ug=Hs.166175 /len=1519
216	AF034209	31455_r_at		Cluster Incl. AF034209:Homo sapiens RIG-like 5-6 mRNA, complete cds /cds=(720,1070) /gb=AF034209 /gi=2724107 /ug=Hs.166175 /len=1519
217	AL008627	31456_at		Cluster Incl. AL008627:Human DNA sequence from PAC 130G2 on chromosome 6p22.2-22.3. Contains ribosomal protein L29 pseudogene, ESTs and STSs /cds=(43,528) /gb=AL008627 /gi=2769539 /ug=Hs.166181 /len=648
218	AF042832	31457_at		Cluster Incl. AF042832:Homo sapiens forkhead-related transcription factor FREAC-9 (FKHL17) mRNA, complete cds /cds=(2127,3620) /gb=AF042832 /gi=3090886 /ug=Hs.166188
219	M99063	31458_at		Cluster Incl. M99063:Homo sapiens cytokeratin 2 mRNA, complete cds /cds=(31,1947) /gb=M99063 /gi=181389 /ug=Hs.166189 /len=2513
220	AF043586	31459_i_at		Cluster Incl. AF043586:Homo sapiens clone CPRF1-T2 immunoglobulin lambda chain VJ region, (IGL) mRNA, partial cds /cds=(0,273) /gb=AF043586 /gi=2865483 /ug=Hs.166191 /len=306
221	AF043586	31460_f_at		Cluster Incl. AF043586:Homo sapiens clone CPRF1-T2 immunoglobulin lambda chain VJ region, (IGL) mRNA, partial cds /cds=(0,273) /gb=AF043586 /gi=2865483 /ug=Hs.166191 /len=306
222	AF050199	31461_at		Cluster Incl. AF050199:Homo sapiens putative peroxisome microbody protein 175.1 mRNA, complete cds /cds=(28,684) /gb=AF050199 /gi=2961556 /ug=Hs.166197 /len=1890
223	U90842	31462_f_at		Cluster Incl. U90842:Homo sapiens SSX5 (SSX5) mRNA, complete cds /cds=(0,566) /gb=U90842 /gi=2952024 /ug=Hs.166198
224	AL022097	31463_s_at		Cluster Incl. AL022097:Homo sapiens DNA sequence from PAC 256G22 on chromosome 6p24.1-25.3. Contains a HNRNP Core Protein A1 LIKE pseudogene and an exon with similarity to yeast and fly phenylalanyl tRNA synthetase PHERS. Contains ESTs and GSSs /cds=(63,1031) /gb=AL022097 /gi=3169107 /ug=Hs.166203 /len=1737
225	AJ000503	31464_at		Cluster Incl. AJ000503:Homo sapiens mRNA for tyrosinase-related protein 2 (TRP-2) /cds=(399,1112) /gb=AJ000503 /gi=3319962 /ug=Hs.167384 /len=3525
226	AJ000503	31465_g_at		Cluster Incl. AJ000503:Homo sapiens mRNA for tyrosinase-related protein 2 (TRP-2) /cds=(399,1112) /gb=AJ000503 /gi=3319962 /ug=Hs.167384 /len=3525
227	U33547	31466_at		Cluster Incl. U33547:Homo sapiens MHC class II antigen HLA-DRB6 mRNA, partial cds /cds=(0,106) /gb=U33547 /gi=3320135 /ug=Hs.167385 /len=720
228	M38056	31467_at		Cluster Incl. M38056:Human lymphocyte antigen mRNA, complete cds /cds=(46,249) /gb=M38056 /gi=184159 /ug=Hs.167748 /len=271
229	W26104	31468_f_at		Cluster Incl. W26104:22f6 Homo sapiens cDNA /gb=W26104 /gi=1306650 /ug=Hs.167981 /len=702
230	W27234	31469_s_at		Cluster Incl. W27234:24b9 Homo sapiens cDNA /gb=W27234 /gi=1306750 /ug=Hs.167982 /len=679
231	W27604	31470_at		Cluster Incl. W27604:35b1 Homo sapiens cDNA /gb=W27604 /gi=1307552 /ug=Hs.167991 /len=682

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232	AL031393	31471_at	Cluster Incl. AL031393:Human DNA sequence from clone 733D15 on chromosome Xp11.3. Contains a Zinc-finger (pseudo?) gene and GSSs /cds=(0,1492) /gb=AL031393 /gi=3646112 /ug=Hs.169190 /len=1493
233	AF098641	31472_s_at	Cluster Incl. AF098641:Homo sapiens CD44 isoform RC (CD44) mRNA, complete cds /cds=(18,437) /gb=AF098641 /gi=3832517 /ug=Hs.169200 /len=484
234	AF082558	31473_s_at	Cluster Incl. AF082558:Homo sapiens truncated TRF1-interacting ankyrin-related ADP-ribose polymerase TT7 mRNA, alternatively spliced, partial cds /cds=(0,100) /gb=AF082558 /gi=3929222 /ug=Hs.169213 /len=357
235	AF082558	31474_r_at	Cluster Incl. AF082558:Homo sapiens truncated TRF1-interacting ankyrin-related ADP-ribose polymerase TT7 mRNA, alternatively spliced, partial cds /cds=(0,100) /gb=AF082558 /gi=3929222 /ug=Hs.169213 /len=357
236	AF082559	31475_at	Cluster Incl. AF082559:Homo sapiens truncated TRF1-interacting ankyrin-related ADP-ribose polymerase TT9 mRNA, alternatively spliced, partial cds /cds=(0,206) /gb=AF082559 /gi=3929224 /ug=Hs.169214 /len=523
237	AF082559	31476_g_at	Cluster Incl. AF082559:Homo sapiens truncated TRF1-interacting ankyrin-related ADP-ribose polymerase TT9 mRNA, alternatively spliced, partial cds /cds=(0,206) /gb=AF082559 /gi=3929224 /ug=Hs.169214 /len=523
238	L08044	31477_at	Cluster Incl. L08044:Human intestinal trefoil factor mRNA, complete cds /cds=(1,225) /gb=L08044 /gi=307520
239	M16653	31478_at	Cluster Incl. M16653:Human pancreatic elastase IIB mRNA, complete cds /cds=(25,834) /gb=M16653 /gi=182059 /ug=Hs.169234 /len=922
240	J00146	31479_f_at	Cluster Incl. J00146:Human dihydrofolate reductase pseudogene (psi-hd1) /cds=(23,316) /gb=J00146 /gi=182732 /ug=Hs.169235 /len=964
241	L18877	31480_f_at	Cluster Incl. L18877:Human MAGE-12 protein gene, complete cds /cds=(75,1019) /gb=L18877 /gi=499345 /ug=Hs.169246 /len=1583
242	M92383	31481_s_at	Cluster Incl. M92383:Homo sapiens thymosin beta-10 gene, 3 end /cds=(0,149) /gb=M92383 /gi=339696 /ug=Hs.169250 /len=400
243	L14813	31482_at	Cluster Incl. L14813:Human carboxyl ester lipase like protein (CELL) mRNA, complete cds /cds=(13,192) /gb=L14813 /gi=409423 /ug=Hs.169271 /len=1214
244	L14813	31483_g_at	Cluster Incl. L14813:Human carboxyl ester lipase like protein (CELL) mRNA, complete cds /cds=(13,192) /gb=L14813 /gi=409423 /ug=Hs.169271 /len=1214
245	X72882	31484_at	Cluster Incl. X72882:H.sapiens 14A6CK DNA sequence /cds=(0,131) /gb=X72882 /gi=667005 /ug=Hs.169275 /len=132
246	M57423	31485_at	Cluster Incl. M57423:Homo sapiens phosphoribosylpyrophosphate synthetase subunit III mRNA, 3 end /cds=(81,1037) /gb=M57423 /gi=190521 /ug=Hs.169284 /len=1091
247	Z70218	31486_s_at	Cluster Incl. Z70218:H.sapiens mRNA for MN1 protein (clone ICRFP507I0498) /cds=(0,50) /gb=Z70218 /gi=1235523 /ug=Hs.169287 /len=341
248	L17328	31487_at	Cluster Incl. L17328:Human pre-T/NK cell associated protein (3Ci) mRNA, complete cds /cds=(1296,1451) /gb=L17328 /gi=306331 /ug=Hs.169304 /len=1911
249	S81916	31488_s_at	Cluster Incl. S81916:phosphoglycerate kinase [alternatively spliced] [human, phosphoglycerate kinase deficient patient with episodes of muscle, mRNA Partial Mutant, 307 nt] /cds=(0,143) /gb=S81916 /gi=1470308 /ug=Hs.169313 /len=307
250	U63332	31489_at	Cluster Incl. U63332:Human super cysteine rich protein mRNA, partial cds /cds=(0,141) /gb=U63332 /gi=1480862 /ug=Hs.169325 /len=427

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251	M77235	31490_at	Cluster Incl. M77235:Human cardiac tetrodotoxin-insensitive voltage-dependent sodium channel alpha subunit (HH1) mRNA, complete cds /cds=(150,6200) /gb=M77235 /gi=184038 /ug=Hs.169331 /len=8491
252	X98175	31491_s_at	Cluster Incl. X98175:H.sapiens mRNA for MACH-beta-2 protein /cds=(0,277) /gb=X98175 /gi=1403326 /ug=Hs.169669 /len=334
253	AB019392	31492_at	Cluster Incl. AB019392:Homo sapiens mRNA of muscle specific gene M9, complete cds /cds=(171,827) /gb=AB019392 /gi=4587122 /ug=Hs.170120 /len=859
254	J03071	31493_s_at	Cluster Incl. J03071:Human growth hormone (GH-1 and GH-2) and chorionic somatomammotropin (CS-1, CS-2 and CS-5) genes, complete cds /cds=(0,653) /gb=J03071 /gi=183148 /ug=Hs.173250 /len=654
255	D25272	31494_at	Cluster Incl. D25272:Homo sapiens mRNA, clone-RES4-16 /cds=UNKNOWN /gb=D25272 /gi=464187 /ug=Hs.174225
256	D63789	31495_at	Cluster Incl. D63789:Homo sapiens DNA for SCM-1beta precursor, complete cds /cds=(21,365) /gb=D63789 /gi=1754608 /ug=Hs.174228 /len=485
257	D63789	31496_g_at	Cluster Incl. D63789:Homo sapiens DNA for SCM-1beta precursor, complete cds /cds=(21,365) /gb=D63789 /gi=1754608 /ug=Hs.174228 /len=485
258	U19142	31497_at	Cluster Incl. U19142:Human GAGE-1 protein mRNA, complete cds /cds=(48,464) /gb=U19142 /gi=914898 /ug=Hs.176660 /len=634
259	U19147	31498_f_at	Cluster Incl. U19147:Human GAGE-6 protein mRNA, complete cds /cds=(81,434) /gb=U19147 /gi=914908 /ug=Hs.176662 /len=527
260	X16863	31499_s_at	Cluster Incl. X16863:Human Fc-gamma RIII-1 cDNA for Fc-gamma receptor III-1 (CD 16) /cds=(33,734) /gb=X16863 /gi=31321 /ug=Hs.176663 /len=887
261	W26594	31500_at	Cluster Incl. W26594:33g6 Homo sapiens cDNA /gb=W26594 /gi=1307455 /ug=Hs.177484 /len=670
262	W27645	31501_at	Cluster Incl. W27645:37d5 Homo sapiens cDNA /gb=W27645 /gi=1307719 /ug=Hs.177488 /len=742
263	W27953	31502_at	Cluster Incl. W27953:39h7 Homo sapiens cDNA /gb=W27953 /gi=1307901 /ug=Hs.177489 /len=758
264	W28732	31503_at	Cluster Incl. W28732:50h7 Homo sapiens cDNA /gb=W28732 /gi=1308680 /ug=Hs.177496 /len=818
265	M64098	31504_at	Cluster Incl. M64098:Human high density lipoprotein binding protein (HBP) mRNA, complete cds /cds=(154,3960) /gb=M64098 /gi=183891 /ug=Hs.177516 /len=4354
266	Z28407	31505_at	Cluster Incl. Z28407:H.sapiens mRNA for ribosomal protein L8 /cds=(43,816) /gb=Z28407 /gi=433898 /ug=Hs.178551 /len=852
267	L12691	31506_s_at	Cluster Incl. L12691:Human neutrophil peptide-3 gene, complete cds /cds=(50,334) /gb=L12691 /gi=292364 /ug=Hs.178741
268	U80756	31507_at	Cluster Incl. U80756:Homo sapiens CAGL114 mRNA, partial cds /cds=(0,509) /gb=U80756 /gi=2565081 /ug=Hs.178746 /len=533
269	S73591	31508_at	Cluster Incl. S73591:brain-expressed HHCPA78 homolog [human, HL-60 acute promyelocytic leukemia cells, mRNA, 2704 nt] /cds=(221,1396) /gb=S73591 /gi=688296 /ug=Hs.179526
270	X64707	31509_at	Cluster Incl. X64707:H.sapiens BBC1 mRNA /cds=(51,686) /gb=X64707 /gi=29382 /ug=Hs.180842 /len=942
271	Z48950	31510_s_at	Cluster Incl. Z48950:H.sapiens hH3.3B gene for histone H3.3 /cds=(10,420) /gb=Z48950 /gi=761715 /ug=Hs.180877 /len=1555
272	U14971	31511_at	Cluster Incl. U14971:Human ribosomal protein S9 mRNA, complete cds /cds=(35,619) /gb=U14971 /gi=550022 /ug=Hs.180920
273	Z00010	31512_at	Cluster Incl. Z00010:H.sapiens germ line pseudogene for immunoglobulin kappa light chain leader peptide and variable region (subgroup V kappa I) /cds=(0,170) /gb=Z00010 /gi=33146 /ug=Hs.37089 /len=351

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274	X90828	31513_at	Cluster Incl. X90828:H.sapiens mRNA for transcription factor, Lbx1 /cds=(141,983) /gb=X90828 /gi=1617115 /ug=Hs.37128
275	AF034970	31514_at	Cluster Incl. AF034970:Homo sapiens docking protein (DOK-2) mRNA, complete cds /cds=(0,1238) /gb=AF034970 /gi=3043918 /ug=Hs.181515 /len=1239
276	AF053356	31515_at	Cluster Incl. AF053356:Homo sapiens chromosome 7q22 sequence /cds=(0,809) /gb=AF053356 /gi=3135305 /ug=Hs.181516 /len=810
277	M90354	31516_f_at	Cluster Incl. M90354:Human BTF3 protein homologue gene, complete cds /cds=(0,335) /gb=M90354 /gi=179571 /ug=Hs.181965 /len=336
278	M90355	31517_f_at	Cluster Incl. M90355:Human BTF3 protein homologue gene, complete cds /cds=(0,203) /gb=M90355 /gi=1435190 /ug=Hs.181966 /len=204
279	M90356	31518_j_at	Cluster Incl. M90356:Human BTF3 protein homologue gene, complete cds /cds=(0,644) /gb=M90356 /gi=179575 /ug=Hs.181967 /len=645
280	M90356	31519_f_at	Cluster Incl. M90356:Human BTF3 protein homologue gene, complete cds /cds=(0,644) /gb=M90356 /gi=179575 /ug=Hs.181967 /len=645
281	X77824	31520_at	Cluster Incl. X77824:H.sapiens chromobox 2 gene /cds=(0,236) /gb=X77824 /gi=773207 /ug=Hs.181970 /len=237
282	X60484	31521_f_at	Cluster Incl. X60484:H.sapiens H4/e gene for H4 histone /cds=(0,311) /gb=X60484 /gi=32000 /ug=Hs.181972 /len=312
283	Z80779	31522_f_at	Cluster Incl. Z80779:H.sapiens H2B/g gene /cds=(0,380) /gb=Z80779 /gi=1568548 /ug=Hs.182137 /len=381
284	Z80780	31523_f_at	Cluster Incl. Z80780:H.sapiens H2B/h gene /cds=(0,380) /gb=Z80780 /gi=1568550 /ug=Hs.182138 /len=381
285	Z80782	31524_f_at	Cluster Incl. Z80782:H.sapiens H2B/k gene /cds=(0,380) /gb=Z80782 /gi=1568554 /ug=Hs.182140 /len=381
286	J00153	31525_s_at	Cluster Incl. J00153:Human alpha globin gene cluster on chromosome 16- zeta gene /cds=(0,428) /gb=J00153 /gi=183794 /ug=Hs.182374 /len=429
287	X63547	31526_f_at	Cluster Incl. X63547:H.sapiens mRNA for tre oncogene (clone 213) /cds=UNKNOWN /gb=X63547 /gi=37332 /ug=Hs.182377
288	X17206	31527_at	Cluster Incl. X17206:Human mRNA for LLRep3 /cds=(240,905) /gb=X17206 /gi=34391 /ug=Hs.182426 /len=934
289	Z83738	31528_f_at	Cluster Incl. Z83738:H.sapiens hH2B/e gene /cds=(0,380) /gb=Z83738 /gi=1749801 /ug=Hs.182432 /len=381
290	X99141	31529_at	Cluster Incl. X99141:H.sapiens mRNA for hair keratin, hHb3 /cds=(52,1533) /gb=X99141 /gi=1903215 /ug=Hs.182506
291	U89344	31530_at	Cluster Incl. U89344:Human acetyl-CoA carboxylase (ACC2) mRNA, complete cds /cds=(0,7451) /gb=U89344 /gi=2138329 /ug=Hs.183857 /len=7452
292	U89344	31531_g_at	Cluster Incl. U89344:Human acetyl-CoA carboxylase (ACC2) mRNA, complete cds /cds=(0,7451) /gb=U89344 /gi=2138329 /ug=Hs.183857 /len=7452
293	U43292	31532_at	Cluster Incl. U43292:Human MDS1B (MDS1) mRNA, complete cds /cds=(307,816) /gb=U43292 /gi=1294814 /ug=Hs.54504 /len=1277
294	AF048713	31533_s_at	Cluster Incl. AF048713:Homo sapiens Kv4.3 potassium channel long splice variant (Kv4.3) mRNA, complete cds /cds=(0,1967) /gb=AF048713 /gi=2935435 /ug=Hs.184889 /len=1968
295	M30607	31534_at	Cluster Incl. M30607:Human zinc finger protein Y-linked (ZF) mRNA, complete cds /cds=(0,2405) /gb=M30607 /gi=340435 /ug=Hs.184915 /len=2659
296	W27858	31535_i_at	Cluster Incl. W27858:39e3 Homo sapiens cDNA /gb=W27858 /gi=1307869 /ug=Hs.186617 /len=821
297	AB020693	31536_at	Cluster Incl. AB020693:Homo sapiens mRNA for KIAA0886 protein, complete cds /cds=(134,3712) /gb=AB020693 /gi=4240260 /ug=Hs.65450 /len=4053

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298	S74683	31537_at	Cluster Incl. S74683:ADP-ribosyltransferase [human, skeletal muscle, mRNA, 1334 nt] /cds=(101,1084) /gb=S74683 /gi=807099 /ug=Hs.73139 /len=1311
299	M17885	31538_at	Cluster Incl. M17885:Human acidic ribosomal phosphoprotein P0 mRNA, complete cds /cds=(77,1030) /gb=M17885 /gi=190231 /ug=Hs.73742 /len=1097
300	L23852	31539_r_at	Cluster Incl. L23852:Homo sapiens (clone Z146) retinal mRNA, 3 end and repeat region /cds=(0,241) /gb=L23852 /gi=393126 /ug=Hs.73838 /len=1711
301	U03397	31540_at	Cluster Incl. U03397:Human receptor protein 4-1BB mRNA, complete cds /cds=(119,886) /gb=U03397 /gi=571320 /ug=Hs.73895 /len=1399
302	X98307	31541_at	Cluster Incl. X98307:H.sapiens mRNA for UV-B repressed sequence, HUR 7 /cds=(0,169) /gb=X98307 /gi=1524079 /ug=Hs.84168 /len=394
303	M60502	31542_at	Cluster Incl. M60502:Human profilaggrin mRNA, 3 end /cds=(0,2393) /gb=M60502 /gi=190403 /ug=Hs.73995 /len=2930
304	S83198	31543_at	Cluster Incl. S83198:BPLP=basic proline-rich protein [human, lacrimal gland, mRNA, 947 nt] /cds=(80,685) /gb=S83198 /gi=1836021 /ug=Hs.87198 /len=947
305	L13203	31544_at	Cluster Incl. L13203:Human HNF-3/fork-head homolog-3 HFH-3 mRNA, complete cds /cds=(78,1133) /gb=L13203 /gi=1911184 /ug=Hs.87236 /len=2089
306	AL031228	31545_at	Cluster Incl. AL031228:dJ1033B10.4 (40S ribosomal protein S18 (RPS18, KE-3)) /cds=(39,497) /gb=AL031228 /gi=3646023 /ug=Hs.75362 /len=541
307	L11566	31546_at	Cluster Incl. L11566:Homo sapiens ribosomal protein L18 (RPL18) mRNA, complete cds /cds=(15,581) /gb=L11566 /gi=337492 /ug=Hs.75458 /len=630
308	D37933	31547_at	Cluster Incl. D37933:Human mRNA for SYNTAXIN1B, complete cds /cds=(17,883) /gb=D37933 /gi=531249 /ug=Hs.99880
309	L13291	31548_at	Cluster Incl. L13291:Human ADP-ribosylarginine hydrolase mRNA, complete cds /cds=(105,1178) /gb=L13291 /gi=402477 /ug=Hs.99884 /len=1365
310	M13150	31549_at	Cluster Incl. M13150:Human mas proto-oncogene mRNA, complete cds /cds=(267,1244) /gb=M13150 /gi=187388 /ug=Hs.99900 /len=1388
311	J03019	31550_at	Cluster Incl. J03019:Human beta-1-adrenergic receptor mRNA, complete cds /cds=(86,1519) /gb=J03019 /gi=178199 /ug=Hs.99913 /len=1723
312	M86868	31551_at	Cluster Incl. M86868:Human gamma amino butyric acid (GABA rho2) gene mRNA, complete cds /cds=(80,1477) /gb=M86868 /gi=182912 /ug=Hs.99927 /len=1628
313	M28219	31552_at	Cluster Incl. M28219:Homo sapiens low density lipoprotein receptor (FH 10 mutant causing familial hypercholesterolemia) mRNA, 3 end /cds=(0,182) /gb=M28219 /gi=619785 /ug=Hs.99940
314	X78931	31553_at	Cluster Incl. X78931:H.sapiens HZF8 mRNA for zinc finger protein /cds=(0,524) /gb=X78931 /gi=498733 /ug=Hs.99971 /len=746
315	X55019	31554_at	Cluster Incl. X55019:H.sapiens mRNA for acetylcholine receptor delta subunit /cds=(4,1557) /gb=X55019 /gi=297401 /ug=Hs.99975 /len=1741
316	X16666	31555_at	Cluster Incl. X16666:Human HOX2I mRNA from the Hox2 locus /cds=(6,911) /gb=X16666 /gi=32383 /ug=Hs.99992 /len=1021
317	X61072	31556_at	Cluster Incl. X61072:Human mRNA for T cell receptor, clone IGRA17 /cds=(0,200) /gb=X61072 /gi=33511 /ug=Hs.99996
318	M17733	31557_at	Cluster Incl. M17733:Human thymosin beta-4 mRNA, complete cds /cds=(77,211) /gb=M17733 /gi=339688 /ug=Hs.75968 /len=556
319	X91103	31558_at	Cluster Incl. X91103:H.sapiens mRNA for Hr44 protein /cds=(0,1016) /gb=X91103 /gi=1054775 /ug=Hs.102131 /len=1182

HG_U95A

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320	U26209	31559_at	Cluster Incl. U26209:Human renal sodium/dicarboxylate cotransporter (NADC1) mRNA, complete cds /cds=(68,1846) /gb=U26209 /gi=1098556 /ug=Hs.102307 /len=1953
321	U49065	31560_at	Cluster Incl. U49065:Human interleukin-1 receptor-related protein mRNA, complete cds /cds=(134,1822) /gb=U49065 /gi=1236078 /ug=Hs.102865 /len=1971
322	S81591	31561_at	Cluster Incl. S81591:SLC9A2=Na ⁺ /H ⁺ exchanger [human, liver, mRNA, 2665 nt] /cds=(418,2514) /gb=S81591 /gi=1478393 /ug=Hs.103132 /len=2665
323	U63973	31562_at	Cluster Incl. U63973:Human rhodopsin kinase mRNA, complete cds /cds=(108,1799) /gb=U63973 /gi=1488315 /ug=Hs.103501 /len=1952
324	U03115	31563_at	Cluster Incl. U03115:Human V beta T-cell receptor (TCRBV) gene locus /cds=(0,149) /gb=U03115 /gi=467918 /ug=Hs.103945 /len=342
325	S65761	31564_at	Cluster Incl. S65761:anti-colorectal carcinoma heavy chain=glycoprotein CANAG-50 specific IgG1 kappa [human, 19.9 hybridoma, antibody 1116NS19.9, mRNA, 1515 nt] /cds=(44,1420) /gb=S65761 /gi=425517 /ug=Hs.103995 /len=1515
326	S65921	31565_at	Cluster Incl. S65921:anti-colorectal carcinoma light chain=glycoprotein CANAG-50 specific IgG1 kappa [human, 19.9 hybridoma, antibody 1116NS19.9, mRNA, 998 nt] /cds=(79,789) /gb=S65921 /gi=425519 /ug=Hs.103996 /len=998
327	U00956	31566_at	Cluster Incl. U00956:Human clone KDB5.1 (CAC)n/(GTG)n repeat-containing mRNA /cds=UNKNOWN /gb=U00956 /gi=405058 /ug=Hs.104112 /len=1424
328	S82769	31567_at	Cluster Incl. S82769:GABAA receptor gamma 3 subunit [human, fetal brain, mRNA Partial, 1536 nt] /cds=(32,1435) /gb=S82769 /gi=1754748 /ug=Hs.104133 /len=1536
329	U14972	31568_at	Cluster Incl. U14972:Human ribosomal protein S10 mRNA, complete cds /cds=(16,513) /gb=U14972 /gi=550024
330	U76362	31569_at	Cluster Incl. U76362:Human retinal glutamate transporter EAAT5 mRNA, complete cds /cds=(188,1870) /gb=U76362 /gi=2076761 /ug=Hs.104637 /len=1996
331	U70212	31570_at	Cluster Incl. U70212:Human SIM1 mRNA, complete cds /cds=(216,2516) /gb=U70212 /gi=2245351 /ug=Hs.105925 /len=4007
332	U93868	31571_at	Cluster Incl. U93868:Human RNA polymerase III subunit (RPC32) mRNA, complete cds /cds=(162,863) /gb=U93868 /gi=2228749 /ug=Hs.105929 /len=1000
333	AF011757	31572_at	Cluster Incl. AF011757:Homo sapiens RAGE binding protein (P12) mRNA, complete cds /cds=(0,278) /gb=AF011757 /gi=2293532 /ug=Hs.105937 /len=388
334	M64716	31573_at	Cluster Incl. M64716:Human ribosomal protein S25 mRNA, complete cds /cds=(71,448) /gb=M64716 /gi=337507 /ug=Hs.113029 /len=497
335	M14087	31574_i_at	Cluster Incl. M14087:Human HL14 gene encoding beta-galactoside-binding lectin, 3 end, clone 2 /cds=(0,205) /gb=M14087 /gi=187113 /ug=Hs.113212 /len=229
336	M14087	31575_f_at	Cluster Incl. M14087:Human HL14 gene encoding beta-galactoside-binding lectin, 3 end, clone 2 /cds=(0,205) /gb=M14087 /gi=187113 /ug=Hs.113212 /len=229
337	M20919	31576_at	Cluster Incl. M20919:Human DNA with a hepatitis B virus surface antigen (HBsAg) gene (complete cds) insertion /cds=(0,593) /gb=M20919 /gi=184331 /ug=Hs.113220 /len=594
338	D38466	31577_at	Cluster Incl. D38466:Homo sapiens gene for a1 chain of type XIX collagen /cds=(0,89) /gb=D38466 /gi=1468954 /ug=Hs.113225 /len=721

Cluster ID	Database Accession	Affymetrix Accession	Entrez Definition
339	M96936	31578_at	Cluster Incl. M96936:Homo sapiens cystic fibrosis transmembrane conductance regulator (CTFR) gene, exons 23 (3 end), 24a, and 24 (5 end) /cds=(0,74) /gb=M96936 /gi=180293 /ug=Hs.113235
340	AF005082	31579_at	Cluster Incl. AF005082:Homo sapiens skin-specific protein (xp33) mRNA, partial cds /cds=(0,287) /gb=AF005082 /gi=2589191 /ug=Hs.113261 /len=303
341	AF028840	31580_at	Cluster Incl. AF028840:Homo sapiens Kruppel-associated box protein mRNA, complete cds /cds=(42,1385) /gb=AF028840 /gi=2623621 /ug=Hs.113263 /len=1834
342	U96291	31581_at	Cluster Incl. U96291:Homo sapiens Ig kappa light chain variable region (VklI-A23) mRNA, complete cds /cds=(9,110) /gb=U96291 /gi=2345027 /ug=Hs.113273 /len=336
343	U93163	31582_at	Cluster Incl. U93163:Homo sapiens MAGE-B2 (MAGE-B2), MAGE-B3 (MAGE-B3), MAGE-B4 (MAGE-B4), and MAGE-B1 (MAGE-B1) genes, complete cds /cds=(60,1100) /gb=U93163 /gi=3228377 /ug=Hs.113291 /len=1727
344	X67247	31583_at	Cluster Incl. X67247:H.sapiens rpS8 gene for ribosomal protein S8 /cds=(23,649) /gb=X67247 /gi=36149 /ug=Hs.118690 /len=705
345	X16064	31584_at	Cluster Incl. X16064:Human mRNA for translationally controlled tumor protein /cds=(94,612) /gb=X16064 /gi=37495 /ug=Hs.119252 /len=830
346	X94552	31585_at	Cluster Incl. X94552:H.sapiens mRNA for metabotropic glutamate receptor type 7 /cds=(238,2985) /gb=X94552 /gi=1370110 /ug=Hs.83407 /len=3021
347	X72475	31586_f_at	Cluster Incl. X72475:H.sapiens mRNA for rearranged Ig kappa light chain variable region (L114) /cds=(0,421) /gb=X72475 /gi=441418 /ug=Hs.192989 /len=454
348	X96969	31587_at	Cluster Incl. X96969:H.sapiens mRNA for urea transporter /cds=(273,1466) /gb=X96969 /gi=1483515 /ug=Hs.193377 /len=2053
349	AL034450	31588_at	Cluster Incl. AL034450:Human DNA sequence from clone 115K14 on chromosome Xq22.3-23 Contains high mobility group protein 2a, ESTs, STS /cds=(0,605) /gb=AL034450 /gi=4210359 /ug=Hs.194749 /len=730
350	AL031983	31589_at	Cluster Incl. AL031983:dJ271M21.2 (hs6M1-12 (7 transmembrane receptor (Rhodopsin family) (Olfactory Receptor Like) protein)) /cds=(497,1435) /gb=AL031983 /gi=4160199 /ug=Hs.194767 /len=1591
351	AL031983	31590_g_at	Cluster Incl. AL031983:dJ271M21.2 (hs6M1-12 (7 transmembrane receptor (Rhodopsin family) (Olfactory Receptor Like) protein)) /cds=(497,1435) /gb=AL031983 /gi=4160199 /ug=Hs.194767 /len=1591
352	X98337	31591_s_at	Cluster Incl. X98337:H.sapiens mRNA for complement factor H-related protein 4 /cds=(85,1080) /gb=X98337 /gi=1524057 /ug=Hs.194776 /len=1305
353	AJ002425	31592_at	Cluster Incl. AJ002425:Homo sapiens mRNA for p65 protein /cds=(60,1904) /gb=AJ002425 /gi=4753767 /ug=Hs.194790 /len=2077
354	Z70200	31593_at	Cluster Incl. Z70200:H.sapiens gene for U5 snRNP-specific 200kD protein /cds=(0,5105) /gb=Z70200 /gi=3255964 /ug=Hs.197800 /len=5106
355	Y16788	31594_at	Cluster Incl. Y16788:Homo sapiens hHa3-I gene, complete CDS /cds=(0,1214) /gb=Y16788 /gi=3724100 /ug=Hs.197874 /len=1215
356	AJ223317	31595_at	Cluster Incl. AJ223317:Homo sapiens mRNA for sarcosine dehydrogenase, complete cds /cds=(132,1559) /gb=AJ223317 /gi=4454258 /ug=Hs.198003 /len=1970
357	L02326	31596_f_at	Cluster Incl. L02326:Homo sapiens (clone Hu lambda-17) lambda-like gene, complete cds /cds=(0,94) /gb=L02326 /gi=292400 /ug=Hs.198118 /len=95

HG_U95A

Cluster ID	Database Accession	Affymetrix Accession	Entrez Definition
358	L36055	31597_r_at	Cluster Incl. L36055:Human 4E-binding protein 1 mRNA, complete cds /cds=(0,356) /gb=L36055 /gi=561629 /ug=Hs.198144
359	L41668	31598_s_at	Cluster Incl. L41668:Homo sapiens UDP-galactose-4-epimerase (GALE) mRNA, complete cds /cds=(93,1139) /gb=L41668 /gi=2623737 /ug=Hs.198245 /len=1488
360	U10691	31599_f_at	Cluster Incl. U10691:Human MAGE-6 antigen (MAGE6) gene, complete cds /cds=(65,1009) /gb=U10691 /gi=533522 /ug=Hs.198263 /len=1564
361	D38435	31600_s_at	Cluster Incl. D38435:Homo sapiens hPMS3 mRNA, partial cds /cds=(0,772) /gb=D38435 /gi=600590 /ug=Hs.198283 /len=846
362	U36610	31601_s_at	Cluster Incl. U36610:Human Y-chromosome RNA recognition motif protein (YRRM) gene /cds=(0,498) /gb=U36610 /gi=1033164 /ug=Hs.198290 /len=768
363	AJ007989	31602_at	Cluster Incl. AJ007989:Homo sapiens mRNA for transcription factor TBX6 /cds=(24,1334) /gb=AJ007989 /gi=4049462 /ug=Hs.198301 /len=1798
364	W27974	31603_at	Cluster Incl. W27974:40c1 Homo sapiens cDNA /gb=W27974 /gi=1307922 /ug=Hs.198819 /len=751
365	AJ009849	31604_at	Cluster Incl. AJ009849:Homo sapiens GNAS1 gene encoding NESP55 /cds=(384,1121) /gb=AJ009849 /gi=4092645 /ug=Hs.199297 /len=1164
366	U72518	31605_at	Cluster Incl. U72518:Human destrin-2 pseudogene mRNA, complete cds /cds=(268,798) /gb=U72518 /gi=1673523 /ug=Hs.199299 /len=1057
367	AB002338	31606_at	Cluster Incl. AB002338:Human mRNA for KIAA0340 gene, partial cds /cds=(0,3163) /gb=AB002338 /gi=2224620 /ug=Hs.199301 /len=6395
368	AJ002428	31607_at	Cluster Incl. AJ002428:Homo sapiens VDAC1 pseudogene /cds=(0,853) /gb=AJ002428 /gi=3183956 /ug=Hs.201553 /len=854
369	AJ002428	31608_g_at	Cluster Incl. AJ002428:Homo sapiens VDAC1 pseudogene /cds=(0,853) /gb=AJ002428 /gi=3183956 /ug=Hs.201553 /len=854
370	L33799	31609_s_at	Cluster Incl. L33799:Human procollagen C-proteinase enhancer protein (PCOLCE) mRNA, complete cds /cds=(60,1409) /gb=L33799 /gi=642907 /ug=Hs.202097 /len=1480
371	U21049	31610_at	Cluster Incl. U21049:Human DD96 mRNA, complete cds /cds=(0,344) /gb=U21049 /gi=722243 /ug=Hs.202557 /len=345
372	AF032457	31611_s_at	Cluster Incl. AF032457:Homo sapiens BimEL mRNA, complete cds /cds=(0,596) /gb=AF032457 /gi=2895495 /ug=Hs.202657 /len=597
373	AJ224171	31612_at	Cluster Incl. AJ224171:Homo sapiens mRNA for lipophilin A /cds=(47,319) /gb=AJ224171 /gi=4107228 /ug=Hs.202686
374	AF029325	31613_at	Cluster Incl. AF029325:Homo sapiens laminin beta-4 chain precursor (LAMB4) mRNA, alternatively spliced short variant, partial cds /cds=(0,157) /gb=AF029325 /gi=4003504
375	U88048	31614_at	Cluster Incl. U88048:Human clone KiSS-16 unknown product mRNA, complete cds /cds=(263,427) /gb=U88048 /gi=4099534 /ug=Hs.202694 /len=630
376	U94902	31615_i_at	Cluster Incl. U94902:Human soluble CD44 (CD44) mRNA, with exon v9 extension, partial cds /cds=(0,95) /gb=U94902 /gi=4100167 /ug=Hs.202695 /len=127
377	U94902	31616_r_at	Cluster Incl. U94902:Human soluble CD44 (CD44) mRNA, with exon v9 extension, partial cds /cds=(0,95) /gb=U94902 /gi=4100167 /ug=Hs.202695 /len=127
378	AF030427	31617_at	Cluster Incl. AF030427:Homo sapiens lung type-I cell membrane-associated protein hT1a-1 (hT1a-1) mRNA, complete cds /cds=(342,836) /gb=AF030427 /gi=4103966 /ug=Hs.202697 /len=1172

Cluster ID	Database Accession	Entrez Definition
379	S66666	31618_at Cluster Incl. S66666:p53=tumor suppressor (alternatively spliced, exon 9-10) [human, Molt-4, T-lymphoblastic leukemia cell line, mRNA Partial Mutant, 160 nt] /cds=(0,47) /gb=S66666 /gi=436292 /ug=Hs.202717 /len=160
380	K00627	31619_at Cluster Incl. K00627:human kpni repeat mrna (cdna clone pcd-kpni-8), 3 end /cds=UNKNOWN /gb=K00627 /gi=337653 /ug=Hs.203776 /len=2126
381	AF033579	31620_at Cluster Incl. AF033579:untitled /cds=(0,633) /gb=AF033579 /gi=3213224 /ug=Hs.203847 /len=634
382	M36860	31621_s_at Cluster Incl. M36860:Human elastin mRNA, complete cds /cds=(49,2241) /gb=M36860 /gi=182061 /ug=Hs.203917 /len=2242
383	M10943	31622_f_at Cluster Incl. M10943:Human metallothionein-I β gene (hMT-I β) /cds=(0,185) /gb=M10943 /gi=187540 /ug=Hs.203936 /len=186
384	K01383	31623_f_at Cluster Incl. K01383:Human metallothionein-I α gene, complete coding sequence /cds=(0,185) /gb=K01383 /gi=187536 /ug=Hs.203967 /len=186
385	AF008203	31624_at Cluster Incl. AF008203:Homo sapiens homeobox protein (Alx3) mRNA, complete cds /cds=(0,1031) /gb=AF008203 /gi=4102042 /ug=Hs.204039 /len=1800
386	AB007921	31625_at Cluster Incl. AB007921:Homo sapiens mRNA for KIAA0452 protein, partial cds /cds=(0,1362) /gb=AB007921 /gi=3413865 /ug=Hs.207796 /len=6256
387	AF047485	31626_i_at Cluster Incl. AF047485:Homo sapiens amine oxidase pseudogene mRNA, splice variant HLAO1 /cds=(134,808) /gb=AF047485 /gi=3851493 /ug=Hs.207798 /len=2073
388	AF047485	31627_f_at Cluster Incl. AF047485:Homo sapiens amine oxidase pseudogene mRNA, splice variant HLAO1 /cds=(134,808) /gb=AF047485 /gi=3851493 /ug=Hs.207798 /len=2073
389	AB001328	31628_at Cluster Incl. AB001328:Homo sapiens mRNA for pH-sensing regulatory factor of peptide transporter, complete cds /cds=(66,692) /gb=AB001328 /gi=2506042 /ug=Hs.207825
390	U51713	31629_at Cluster Incl. U51713:HSU51713 Homo sapiens cDNA /gb=U51713 /gi=1806581 /ug=Hs.208322 /len=707
391	AF083384	31630_at Cluster Incl. AF083384:Homo sapiens 45kDa splicing factor mRNA, complete cds /cds=(0,1217) /gb=AF083384 /gi=3746839 /ug=Hs.209450 /len=1218
392	AF095288	31631_f_at Cluster Incl. AF095288:Homo sapiens pituitary tumor transforming gene protein 2 (PTTG2) mRNA, complete cds /cds=(0,575) /gb=AF095288 /gi=3766237 /ug=Hs.211506 /len=576
393	Z95118	31632_at Cluster Incl. Z95118:Human DNA sequence from clone 354J5 on chromosome 6q21-22. Contains pseudogene similar to zinc finger protein (ZPR1), EST, STS, GSS /cds=(0,1364) /gb=Z95118 /gi=3821018 /ug=Hs.211509 /len=1365
394	Z95118	31633_g_at Cluster Incl. Z95118:Human DNA sequence from clone 354J5 on chromosome 6q21-22. Contains pseudogene similar to zinc finger protein (ZPR1), EST, STS, GSS /cds=(0,1364) /gb=Z95118 /gi=3821018 /ug=Hs.211509 /len=1365
395	M13057	31634_at Cluster Incl. M13057:Human acidic proline-rich protein (PRH1) gene, complete cds /cds=(92,592) /gb=M13057 /gi=190511 /ug=Hs.211591 /len=611
396	M13057	31635_g_at Cluster Incl. M13057:Human acidic proline-rich protein (PRH1) gene, complete cds /cds=(92,592) /gb=M13057 /gi=190511 /ug=Hs.211591 /len=611
397	U09210	31636_s_at Cluster Incl. U09210:Human vesicular acetylcholine transporter mRNA, complete cds /cds=(442,2040) /gb=U09210 /gi=507743 /ug=Hs.211598 /len=2421
398	X72631	31637_s_at Cluster Incl. X72631:H.sapiens mRNA encoding Rev-ErbAalpha /cds=UNKNOWN /gb=X72631 /gi=732801 /ug=Hs.211606

HG_U95A

Cluster ID	Database Accession	Affymetrix Accession	Entrez Definition
399	AC005329	31638_at	Cluster Incl. AC005329:Homo sapiens chromosome 19, cosmid R34382 /cds=(0,476) /gb=AC005329 /gi=3342732 /ug=Hs.211914 /len=477
400	Y13436	31639_f_at	Cluster Incl. Y13436:Homo sapiens sox1 gene /cds=(0,1163) /gb=Y13436 /gi=4128158 /ug=Hs.211945 /len=1164
401	Y13436	31640_r_at	Cluster Incl. Y13436:Homo sapiens sox1 gene /cds=(0,1163) /gb=Y13436 /gi=4128158 /ug=Hs.211945 /len=1164
402	AL031282	31641_s_at	Cluster Incl. AL031282:dJ283E3.2.4 (Matrix Metalloproteinase MMP21/22C) /cds=(0,727) /gb=AL031282 /gi=3860395 /ug=Hs.214290 /len=750
403	AB007970	31642_at	Cluster Incl. AB007970:Homo sapiens mRNA, chromosome 1 specific transcript KIAA0501 /cds=UNKNOWN /gb=AB007970 /gi=3413945 /ug=Hs.223020 /len=4756
404	AB008185	31643_at	Cluster Incl. AB008185:Homo sapiens mRNA for ganglioside sialidase, complete cds /cds=(201,1487) /gb=AB008185 /gi=5631292 /ug=Hs.225947 /len=2083
405	AJ243542	31644_at	Cluster Incl. AJ243542:Homo sapiens mRNA for CCL27 chemokine /cds=(24,362) /gb=AJ243542 /gi=5457353 /ug=Hs.225948
406	AB020625	31645_at	Cluster Incl. AB020625:Homo sapiens mRNA for butyrophilin like receptor, complete cds /cds=(158,1456) /gb=AB020625 /gi=4587208 /ug=Hs.225949 /len=1964
407	AJ006054	31646_at	Cluster Incl. AJ006054:Homo sapiens mRNA for UDP glucuronosyltransferase /cds=(63,1646) /gb=AJ006054 /gi=4753765 /ug=Hs.225950 /len=1766
408	AJ242832	31647_at	Cluster Incl. AJ242832:Homo sapiens mRNA for calpain /cds=(103,2211) /gb=AJ242832 /gi=5689775 /ug=Hs.225953 /len=2338
409	AB016899	31648_at	Cluster Incl. AB016899:Homo sapiens HGC6.1.1 mRNA, complete cds /cds=(618,1163) /gb=AB016899 /gi=5006254 /ug=Hs.225962 /len=2785
410	AB016900	31649_at	Cluster Incl. AB016900:Homo sapiens HGC6.1.2 mRNA, complete cds /cds=(336,827) /gb=AB016900 /gi=5006256 /ug=Hs.225963 /len=1407
411	AB016900	31650_g_at	Cluster Incl. AB016900:Homo sapiens HGC6.1.2 mRNA, complete cds /cds=(336,827) /gb=AB016900 /gi=5006256 /ug=Hs.225963 /len=1407
412	AB016901	31651_at	Cluster Incl. AB016901:Homo sapiens HGC6.2 mRNA, complete cds /cds=(280,453) /gb=AB016901 /gi=5006258 /ug=Hs.225964 /len=1197
413	AB023217	31652_at	Cluster Incl. AB023217:Homo sapiens mRNA for KIAA1000 protein, partial cds /cds=(0,2675) /gb=AB023217 /gi=4589643 /ug=Hs.225968 /len=3852
414	AB022847	31653_at	Cluster Incl. AB022847:Homo sapiens mRNA for norepinephrine transporter isoform 2, partial cds /cds=(0,161) /gb=AB022847 /gi=5360690 /ug=Hs.225971 /len=1028
415	AB028982	31654_at	Cluster Incl. AB028982:Homo sapiens mRNA for KIAA1059 protein, partial cds /cds=(0,3895) /gb=AB028982 /gi=5689454 /ug=Hs.225976 /len=5757
416	AL031737	31655_at	Cluster Incl. AL031737:Human DNA sequence from clone 8B22 on chromosome 1p35.1-36.21 Contains gene similar to cytoplasmic dynein light chain 1, GSSs /cds=(92,361) /gb=AL031737 /gi=4464258 /ug=Hs.225980 /len=371
417	AL050263	31656_at	Cluster Incl. AL050263:Homo sapiens mRNA; cDNA DKFZp547J0410 (from clone DKFZp547J0410) /cds=(651,926) /gb=AL050263 /gi=4886484 /ug=Hs.225985 /len=2336
418	AL050153	31657_at	Cluster Incl. AL050153:Homo sapiens mRNA; cDNA DKFZp586K1520 (from clone DKFZp586K1520) /cds=(0,679) /gb=AL050153 /gi=4884364 /ug=Hs.225987 /len=2642

HG_U95A

Cluster ID	Database Accession	Affymetrix Accession	Entrez Definition
419	AJ133776	31658_at	Cluster Incl. AJ133776:Homo sapiens gene for neurogenin 3 /cds=(158,802) /gb=AJ133776 /gi=5123782 /ug=Hs.225996 /len=1167
420	AL080175	31659_at	Cluster Incl. AL080175:Homo sapiens mRNA; cDNA DKFZp434K091 (from clone DKFZp434K091) /cds=(0,966) /gb=AL080175 /gi=5262647 /ug=Hs.226000 /len=1300
421	AL080191	31660_at	Cluster Incl. AL080191:Homo sapiens mRNA; cDNA DKFZp434A062 (from clone DKFZp434A062) /cds=(0,585) /gb=AL080191 /gi=5262671 /ug=Hs.226001 /len=2924
422	AJ243936	31661_at	Cluster Incl. AJ243936:Homo sapiens mRNA for G16 protein (G16 gene located in the class III region of the major histocompatibility complex) /cds=(100,642) /gb=AJ243936 /gi=5578772 /ug=Hs.226019 /len=1109
423	U35246	31662_at	Cluster Incl. U35246:Human vacuolar protein sorting homolog h-vps45 mRNA, complete cds /cds=(43,1755) /gb=U35246 /gi=1477465 /ug=Hs.226025 /len=1987
424	AB016902	31663_at	Cluster Incl. AB016902:Homo sapiens HGC6.3 mRNA, complete cds /cds=(287,802) /gb=AB016902 /gi=5030941 /ug=Hs.227569 /len=1016
425	W25940	31664_at	Cluster Incl. W25940:15c1 Homo sapiens cDNA /gb=W25940 /gi=1306063 /ug=Hs.228064 /len=712
426	W27675	31665_s_at	Cluster Incl. W27675:36b3 Homo sapiens cDNA /gb=W27675 /gi=1307623 /ug=Hs.228067 /len=748
427	W28731	31666_f_at	Cluster Incl. W28731:50h6 Homo sapiens cDNA /gb=W28731 /gi=1308679 /ug=Hs.229142 /len=806
428	W27698	31667_r_at	Cluster Incl. W27698:36f8 Homo sapiens cDNA /gb=W27698 /gi=1307664 /ug=Hs.230638 /len=688
429	W28193	31668_f_at	Cluster Incl. W28193:43d12 Homo sapiens cDNA /gb=W28193 /gi=1308141 /ug=Hs.231096 /len=981
430	AF039307	31669_s_at	Cluster Incl. AF039307:Homo sapiens homeobox A11 (HOXA11) gene, complete cds /cds=(0,941) /gb=AF039307 /gi=2745850 /ug=Hs.231795 /len=942
431	U81554	31670_s_at	Cluster Incl. U81554:Homo sapiens CaM kinase II isoform mRNA, complete cds /cds=(212,931) /gb=U81554 /gi=2275253 /ug=Hs.231812 /len=972
432	D82351	31671_at	Cluster Incl. D82351:Human retropseudogene MSSP-1 DNA, complete cds /cds=(0,1121) /gb=D82351 /gi=1139532 /ug=Hs.233347 /len=1122
433	D82351	31672_g_at	Cluster Incl. D82351:Human retropseudogene MSSP-1 DNA, complete cds /cds=(0,1121) /gb=D82351 /gi=1139532 /ug=Hs.233347 /len=1122
434	X65784	31673_s_at	Cluster Incl. X65784:H.sapiens CAR gene /cds=(0,428) /gb=X65784 /gi=452752 /ug=Hs.233451 /len=429
435	Z81330	31674_s_at	Cluster Incl. Z81330:Human DNA sequence from cosmid H2A1KLT on chromosome 9q34 contains ORFX gene, STS /cds=(0,1094) /gb=Z81330 /gi=1648871 /ug=Hs.233452 /len=1095
436	AF019083	31675_s_at	Cluster Incl. AF019083:Homo sapiens phosphatase and tensin homolog 2 (PTH2) mRNA, partial cds /cds=(0,1211) /gb=AF019083 /gi=2655024 /ug=Hs.233527 /len=1212
437	AC003973	31676_at	Cluster Incl. AC003973:Homo sapiens DNA from chromosome 19, BAC 33152 /cds=(0,3503) /gb=AC003973 /gi=2739352 /ug=Hs.233550 /len=3504
438	U28131	31677_at	Cluster Incl. U28131:Human HMGI-C chimeric transcript mRNA, partial cds /cds=(0,212) /gb=U28131 /gi=1054886 /ug=Hs.234740 /len=312
439	X66610	31678_at	Cluster Incl. X66610:H.sapiens mRNA for enolase /cds=(39,1415) /gb=X66610 /gi=31178 /ug=Hs.234748 /len=1733

Cluster ID	Database Accession	Affymetrix Accession	Entrez Definition
440	U43279	31679_at	Cluster Incl. U43279:Human nucleoporin nup 36 mRNA, complete cds /cds=(322,1290) /gb=U43279 /gi=1565299 /ug=Hs.234750 /len=3980
441	M55630	31680_at	Cluster Incl. M55630:Human topoisomerase I pseudogene 2 /cds=UNKNOWN /gb=M55630 /gi=339812 /ug=Hs.234751
442	X57282	31681_at	Cluster Incl. X57282:H.sapiens mRNA for soluble erythropoietin receptor /cds=(0,185) /gb=X57282 /gi=36426 /ug=Hs.234752 /len=200
443	D32039	31682_s_at	Cluster Incl. D32039:Human pgH3 mRNA for proteoglycan PG-M(V3), complete cds /cds=(105,2072) /gb=D32039 /gi=1008912 /ug=Hs.234753 /len=2087
444	S71020	31683_at	Cluster Incl. S71020:THRA1/BTR=fusion gene {3 region, deletion/fusion rearrangement} [human, breast cancer cell line BT474, mRNA Partial Mutant, 543 nt] /cds=(0,47) /gb=S71020 /gi=546111 /ug=Hs.234755 /len=526
445	M62896	31684_at	Cluster Incl. M62896:Human lipocortin (LIP) 2 pseudogene mRNA, complete cds-like region /cds=UNKNOWN /gb=M62896 /gi=187146 /ug=Hs.234757 /len=1342
446	Y08976	31685_at	Cluster Incl. Y08976:H.sapiens mRNA for FEV-protein /cds=(584,1300) /gb=Y08976 /gi=1834508 /ug=Hs.234759 /len=1881
447	U83668	31686_at	Cluster Incl. U83668:Human beta-tubulin (TUB4q) gene, complete cds /cds=(293,1594) /gb=U83668 /gi=1857525 /ug=Hs.234761 /len=1747
448	M25079	31687_f_at	Cluster Incl. M25079:Human sickle cell beta-globin mRNA, complete cds /cds=(0,443) /gb=M25079 /gi=179408
449	AF005080	31688_at	Cluster Incl. AF005080:Homo sapiens skin-specific protein (xp5) mRNA, complete cds /cds=(51,383) /gb=AF005080 /gi=2589187 /ug=Hs.234766 /len=606
450	AL096715	31689_at	Cluster Incl. AL096715:Homo sapiens mRNA; cDNA DKFZp564G092 (from clone DKFZp564G092) /cds=(1733,2065) /gb=AL096715 /gi=5419848 /ug=Hs.234796 /len=2437
451	U08997	31690_at	Cluster Incl. U08997:Human glutamate dehydrogenase gene, complete cds /cds=(0,1676) /gb=U08997 /gi=478987 /ug=Hs.239377 /len=1677
452	U08997	31691_g_at	Cluster Incl. U08997:Human glutamate dehydrogenase gene, complete cds /cds=(0,1676) /gb=U08997 /gi=478987 /ug=Hs.239377 /len=1677
453	M59830	31692_at	Cluster Incl. M59830:Human MHC class III HSP70-2 gene (HLA), complete cds /cds=(0,1925) /gb=M59830 /gi=188489 /ug=Hs.239378 /len=1926
454	Z80776	31693_f_at	Cluster Incl. Z80776:H.sapiens H2A/g gene /cds=(0,392) /gb=Z80776 /gi=1568542 /ug=Hs.239458 /len=393
455	X82877	31694_at	Cluster Incl. X82877:H.sapiens Na ⁺ -D-glucose cotransport regulator gene /cds=(0,1853) /gb=X82877 /gi=1617112 /ug=Hs.239459 /len=1854
456	X82877	31695_g_at	Cluster Incl. X82877:H.sapiens Na ⁺ -D-glucose cotransport regulator gene /cds=(0,1853) /gb=X82877 /gi=1617112 /ug=Hs.239459 /len=1854
457	X52987	31696_at	Cluster Incl. X52987:Human mRNA for rap2b gene /cds=(0,551) /gb=X52987 /gi=35862 /ug=Hs.239527 /len=552
458	J04755	31697_s_at	Cluster Incl. J04755:Human ferritin H processed pseudogene, complete cds /cds=UNKNOWN /gb=J04755 /gi=182512 /ug=Hs.239542 /len=2083
459	Y08223	31698_at	Cluster Incl. Y08223:H.sapiens MFH-1 gene /cds=(0,1505) /gb=Y08223 /gi=1869804 /ug=Hs.239571 /len=1506
460	S67334	31699_at	Cluster Incl. S67334:phosphatidylinositol 3-kinase p110 beta isoform=110 kda catalytic subunit [human, mRNA Partial, 3213 nt] /cds=(0,3212) /gb=S67334 /gi=455759 /ug=Hs.239818 /len=3213

Cluster ID	Database Accession	Affymetrix Accession	Entrez Definition
461	AF027957	31700_at	Cluster Incl. AF027957:Homo sapiens G protein-coupled receptor (GPR35) gene, complete cds /cds=(0,929) /gb=AF027957 /gi=2739108 /ug=Hs.239891 /len=930
462	X83412	31701_r_at	Cluster Incl. X83412:H.sapiens B1 mRNA for mucin /cds=(0,515) /gb=X83412 /gi=853955 /ug=Hs.100015 /len=780
463	U57029	31702_at	Cluster Incl. U57029:Human T-cell leukemia virus enhancer factor (HTLF) mRNA, complete cds /cds=(281,1306) /gb=U57029 /gi=1405357 /ug=Hs.103126 /len=1429
464	U62435	31703_at	Cluster Incl. U62435:Human nicotinic acetylcholine receptor alpha6 subunit precursor, mRNA, complete cds /cds=(142,1626) /gb=U62435 /gi=1458117 /ug=Hs.103128 /len=1743
465	U62647	31704_at	Cluster Incl. U62647:Human DNase 1 homolog (DNAS1L2) mRNA, complete cds /cds=(92,991) /gb=U62647 /gi=1518783 /ug=Hs.103503 /len=1183
466	X99977	31705_at	Cluster Incl. X99977:H.sapiens ARS gene, component B /cds=(26,337) /gb=X99977 /gi=1536901 /ug=Hs.103505 /len=528
467	L13283	31706_at	Cluster Incl. L13283:Homo sapiens (clone MG2-5-12) mucin (MG2) mRNA, complete polyA site /cds=UNKNOWN /gb=L13283 /gi=292518 /ug=Hs.103944 /len=2350
468	X52332	31707_at	Cluster Incl. X52332:Homo sapiens mRNA for zinc finger protein 10 /cds=(70,1389) /gb=X52332 /gi=34141 /ug=Hs.104115
469	L05095	31708_at	Cluster Incl. L05095:Human ribosomal protein L30 mRNA, complete cds /cds=(54,401) /gb=L05095 /gi=388034 /ug=Hs.111222 /len=477
470	AI812033	31709_at	Cluster Incl. AI812033:tw46d03.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-2262725 /clone_end=3 /gb=AI812033 /gi=5398599 /ug=Hs.113252 /len=623
471	AE000659	31710_at	Cluster Incl. AE000659:Homo sapiens T-cell receptor alpha delta locus from bases 250472 to 501670 (section 2 of 5) of the Complete Nucleotide Sequence /cds=(0,337) /gb=AE000659 /gi=2358025 /ug=Hs.113284 /len=338
472	U77783	31711_at	Cluster Incl. U77783:Homo sapiens N-methyl-D-aspartate receptor 2D subunit precursor (NMDAR2D) mRNA, complete cds /cds=(88,4098) /gb=U77783 /gi=2444025 /ug=Hs.113286
473	AF009204	31712_at	Cluster Incl. AF009204:Homo sapiens PSD-95/SAP90-associated protein-2 mRNA, partial cds /cds=(0,2832) /gb=AF009204 /gi=2454509 /ug=Hs.113287 /len=3412
474	AB000275	31713_s_at	Cluster Incl. AB000275:Homo sapiens mRNA for DAP-2, partial cds /cds=(0,1616) /gb=AB000275 /gi=2588973 /ug=Hs.113287 /len=1740
475	U89293	31714_at	Cluster Incl. U89293:Human MSH4 (HMSH4) mRNA, complete cds /cds=(40,2851) /gb=U89293 /gi=2463652 /ug=Hs.115246
476	U93720	31715_at	Cluster Incl. U93720:Homo sapiens TEX28 mRNA, complete cds /cds=(144,1376) /gb=U93720 /gi=2443443 /ug=Hs.115365 /len=1525
477	L11369	31716_at	Cluster Incl. L11369:Human protocadherin 42 mRNA, 3 end of cds for alternative splicing PC42-8 /cds=(0,776) /gb=L11369 /gi=307323 /ug=Hs.115642 /len=1321
478	U82382	31717_at	Cluster Incl. U82382:Homo sapiens PIN1 peptidyl-prolyl cis/trans isomerase-like (PIN1L) mRNA, complete cds /cds=(35,337) /gb=U82382 /gi=2501784 /ug=Hs.116993 /len=996
479	AJ000327	31718_at	Cluster Incl. AJ000327:Homo sapiens mRNA for adrenoleukodystrophy related protein (ALDR) /cds=(135,2357) /gb=AJ000327 /gi=2584766 /ug=Hs.117852 /len=3526
480	X02761	31719_at	Cluster Incl. X02761:Human mRNA for fibronectin (FN precursor) /cds=(0,6987) /gb=X02761 /gi=31396 /ug=Hs.118162 /len=7680
481	M10905	31720_s_at	Cluster Incl. M10905:Human cellular fibronectin mRNA /cds=(0,2383) /gb=M10905 /gi=182696 /ug=Hs.118162 /len=2384

Cluster ID	Database Accession	Affymetrix Accession	Entrez Definition
482	X06290	31721_at	Cluster Incl. X06290:Human mRNA for apolipoprotein(a) /cds=(45,13691) /gb=X06290 /gi=28619 /ug=Hs.119520
483	AL022326	31722_at	Cluster Incl. AL022326:dJ333H23.1.1 (60S Ribosomal Protein L3) /cds=(6,1217) /gb=AL022326 /gi=3550039 /ug=Hs.119598 /len=1274
484	X70377	31723_at	Cluster Incl. X70377:H.sapiens mRNA for cystatin D /cds=(4,432) /gb=X70377 /gi=398710 /ug=Hs.121489 /len=465
485	L38518	31724_at	Cluster Incl. L38518:Homo sapiens sonic hedgehog protein (SHH) mRNA, complete cds /cds=(151,1539) /gb=L38518 /gi=663156 /ug=Hs.121539 /len=1576
486	U18235	31725_s_at	Cluster Incl. U18235:Human ATP-binding cassette protein (ABC2) mRNA HFBCD04 clone, partial cds /cds=(2,451) /gb=U18235 /gi=609355 /ug=Hs.121561 /len=525
487	S62908	31726_at	Cluster Incl. S62908:gamma-aminobutyric acidA receptor alpha 3 subunit [human, fetal brain, mRNA, 1637 nt] /cds=(85,1563) /gb=S62908 /gi=386423 /ug=Hs.123024 /len=1637
488	U91510	31727_at	Cluster Incl. U91510:Human CD39L1 mRNA, complete cds /cds=(21,1439) /gb=U91510 /gi=2522323 /ug=Hs.123036
489	M29335	31728_at	Cluster Incl. M29335:Human MHC class II DO-alpha mRNA, partial cds /cds=(0,161) /gb=M29335 /gi=188002 /ug=Hs.123041
490	X84003	31729_at	Cluster Incl. X84003:H.sapiens TAFII18 mRNA for transcription factor TFIID /cds=(11,385) /gb=X84003 /gi=791052 /ug=Hs.123057 /len=413
491	M60094	31730_at	Cluster Incl. M60094:Human testicular H1 histone (H1) gene, complete cds /cds=(43,666) /gb=M60094 /gi=183750 /ug=Hs.123064 /len=725
492	AF013956	31731_at	Cluster Incl. AF013956:Homo sapiens Polycomb 2 homolog (hPc2) mRNA, complete cds /cds=(166,1842) /gb=AF013956 /gi=2317722 /ug=Hs.123085 /len=1867
493	X00948	31732_at	Cluster Incl. X00948:Human mRNA for prepro-relaxin H2 /cds=(0,557) /gb=X00948 /gi=35926 /ug=Hs.127032 /len=558
494	Y07683	31733_at	Cluster Incl. Y07683:H.sapiens mRNA for P2X3 purinoceptor /cds=(165,1358) /gb=Y07683 /gi=2370146 /ug=Hs.127430 /len=1456
495	AJ000041	31734_at	Cluster Incl. AJ000041:Homo sapiens mRNA for HOXC11 /cds=(44,958) /gb=AJ000041 /gi=2661372 /ug=Hs.127562 /len=1981
496	Z94277	31735_at	Cluster Incl. Z94277:Human DNA sequence from PAC 154K9 on chromosome Xp11.3-Xp11.4. Contains protein similar to protein phosphatase inhibitor 2 (IPP-2) ESTs and STS /cds=(0,608) /gb=Z94277 /gi=2578060 /ug=Hs.127689 /len=609
497	AA975427	31736_at	Cluster Incl. AA975427:oaq28g02.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-1587698 /clone_end=3 /gb=AA975427 /gi=3151219 /ug=Hs.128434 /len=398
498	J00068	31737_at	Cluster Incl. J00068:Human adult skeletal muscle alpha-actin mRNA /cds=(103,1236) /gb=J00068 /gi=178028 /ug=Hs.1288
499	W28912	31738_at	Cluster Incl. W28912:53f11 Homo sapiens cDNA /gb=W28912 /gi=1308860 /ug=Hs.129019 /len=975
500	AA977513	31739_at	Cluster Incl. AA977513:on60e11.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-1561100 /clone_end=3 /gb=AA977513 /gi=3154959 /ug=Hs.129019 /len=464
501	AB008913	31740_s_at	Cluster Incl. AB008913:Homo sapiens mRNA for Pax-4, complete cds /cds=(0,1052) /gb=AB008913 /gi=2809074 /ug=Hs.129706 /len=1088
502	AF043978	31741_at	Cluster Incl. AF043978:Homo sapiens paired-box transcription factor (PAX4) mRNA, complete cds /cds=(206,1237) /gb=AF043978 /gi=4105183 /ug=Hs.129706 /len=2009

Cluster ID	Database Accession	Affymetrix Accession	Entrez Definition
503	AF064090	31742_at	Cluster Incl. AF064090:Homo sapiens ligand for herpesvirus entry mediator (HVEM-L) mRNA, complete cds /cds=(382,1104) /gb=AF064090 /gi=3283355 /ug=Hs.129708 /len=1474
504	AF041410	31743_at	Cluster Incl. AF041410:Homo sapiens malignancy-associated protein mRNA, partial cds /cds=(0,199) /gb=AF041410 /gi=2827287 /ug=Hs.129710 /len=555
505	D89479	31744_at	Cluster Incl. D89479:Homo sapiens mRNA for ST1B2, complete cds /cds=(24,914) /gb=D89479 /gi=2826145 /ug=Hs.129742
506	AF007194	31745_at	Cluster Incl. AF007194:Homo sapiens mucin (MUC3) mRNA, partial cds /cds=(0,2874) /gb=AF007194 /gi=2853300 /ug=Hs.129782 /len=4707
507	AI017213	31746_at	Cluster Incl. AI017213:ou32b10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-1627963 /clone_end=3 /gb=AI017213 /gi=3231549 /ug=Hs.131150 /len=462
508	AI017213	31747_g_at	Cluster Incl. AI017213:ou32b10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-1627963 /clone_end=3 /gb=AI017213 /gi=3231549 /ug=Hs.131150 /len=462
509	AL021808	31748_at	Cluster Incl. AL021808:Human DNA sequence from clone 24o18 on chromosome 6p21.31-22.2 Contains zinc finger protein pseudogene, VNO-type olfactory receptor pseudogene, nuclear envelope pore membrane protein, EST, STS, GSS /cds=(0,2915) /gb=AL021808 /gi=3395513 /ug=Hs.131150 /len=2916
510	Z98744	31749_f_at	Cluster Incl. Z98744:histone H2A /cds=(7,399) /gb=Z98744 /gi=3080457 /ug=Hs.131954 /len=499
511	Z98744	31750_at	Cluster Incl. Z98744:histone H1.5 /cds=(118,798) /gb=Z98744 /gi=3080457 /ug=Hs.131956 /len=854
512	Z98744	31751_f_at	Cluster Incl. Z98744:histone H4 /cds=(4,312) /gb=Z98744 /gi=3080457 /ug=Hs.131957 /len=355
513	W25935	31752_at	Cluster Incl. W25935:15b4 Homo sapiens cDNA /gb=W25935 /gi=1306058 /ug=Hs.132341 /len=701
514	W28887	31753_at	Cluster Incl. W28887:53b4 Homo sapiens cDNA /gb=W28887 /gi=1308835 /ug=Hs.133142 /len=935
515	AL080207	31754_at	Cluster Incl. AL080207:Homo sapiens mRNA; cDNA DKFZp434G232 (from clone DKFZp434G232) /cds=(0,591) /gb=AL080207 /gi=5262695 /ug=Hs.134585 /len=921
516	AI094859	31755_at	Cluster Incl. AI094859:qa09a09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-1686232 /clone_end=3 /gb=AI094859 /gi=3433835 /ug=Hs.134989 /len=399
517	AL049328	31756_at	Cluster Incl. AL049328:Homo sapiens mRNA; cDNA DKFZp564E026 (from clone DKFZp564E026) /cds=UNKNOWN /gb=AL049328 /gi=4500102 /ug=Hs.135642 /len=1785
518	W27128	31757_at	Cluster Incl. W27128:22g7 Homo sapiens cDNA /gb=W27128 /gi=1306662 /ug=Hs.135884 /len=701
519	W28876	31758_at	Cluster Incl. W28876:52h7 Homo sapiens cDNA /gb=W28876 /gi=1308824 /ug=Hs.136085 /len=781
520	W26220	31759_at	Cluster Incl. W26220:22d9 Homo sapiens cDNA /gb=W26220 /gi=1306631 /ug=Hs.136089 /len=687
521	W27203	31760_at	Cluster Incl. W27203:24a10 Homo sapiens cDNA /gb=W27203 /gi=1306737 /ug=Hs.136089 /len=650
522	W27916	31761_at	Cluster Incl. W27916:39f6 Homo sapiens cDNA /gb=W27916 /gi=1307882 /ug=Hs.136152 /len=831
523	W28605	31762_at	Cluster Incl. W28605:49a7 Homo sapiens cDNA /gb=W28605 /gi=1308553 /ug=Hs.136178 /len=843
524	AC005255	31763_at	Cluster Incl. AC005255:Homo sapiens chromosome 19, CIT-HSP-146e8 /cds=(223,1182) /gb=AC005255 /gi=3289998 /ug=Hs.137573 /len=2138
525	AB014586	31764_at	Cluster Incl. AB014586:Homo sapiens mRNA for KIAA0686 protein, partial cds /cds=(0,980) /gb=AB014586 /gi=3327185 /ug=Hs.137578 /len=4299

HG_U95A

Cluster ID	Database Accession	Affymetrix Accession	Entrez Definition
526	AB014594	31765_at	Cluster Incl. AB014594:Homo sapiens mRNA for KIAA0694 protein, complete cds /cds=(603,2144) /gb=AB014594 /gi=3327201 /ug=Hs.137579 /len=4132
527	AI653069	31766_s_at	Cluster Incl. AI653069:wb23b02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-2306475 /clone_end=3 /gb=AI653069 /gi=4737048 /ug=Hs.137579 /len=437
528	AB015046	31767_at	Cluster Incl. AB015046:Homo sapiens XYLb mRNA for xylulokinase, complete cds /cds=(120,1703) /gb=AB015046 /gi=3298501 /ug=Hs.137580 /len=1963
529	AL009179	31768_at	Cluster Incl. AL009179:dJ97D16.5 (Histone H2A) /cds=(11,403) /gb=AL009179 /gi=3217024 /ug=Hs.137592 /len=503
530	Y11094	31769_at	Cluster Incl. Y11094:H.sapiens mRNA for WNT-8B protein /cds=(137,1192) /gb=Y11094 /gi=3219593 /ug=Hs.137595 /len=2119
531	W27119	31770_at	Cluster Incl. W27119:20e3 Homo sapiens cDNA /gb=W27119 /gi=1306491 /ug=Hs.140106 /len=711
532	U80770	31771_at	Cluster Incl. U80770:Human EST clone 251800 mariner transposon Hsmar1 sequence /cds=UNKNOWN /gb=U80770 /gi=2921608 /ug=Hs.143439 /len=1134
533	AD001527	31772_at	Cluster Incl. AD001527:Homo sapiens DNA from chromosome 19-cosmid f24590 containing CAPNS and POL2RI, genomic sequence /cds=(0,655) /gb=AD001527 /gi=1905899 /ug=Hs.143920 /len=656
534	U06715	31773_at	Cluster Incl. U06715:Human cytochrome B561, HCYTO B561, mRNA, partial cds /cds=(422,1159) /gb=U06715 /gi=476590 /ug=Hs.153028 /len=2643
535	AF068863	31774_at	Cluster Incl. AF068863:Homo sapiens oligodendrocyte-specific protein (OSP) mRNA, complete cds /cds=(19,675) /gb=AF068863 /gi=3283414 /ug=Hs.153297 /len=2243
536	X65018	31775_at	Cluster Incl. X65018:H.sapiens mRNA for lung surfactant protein D /cds=(171,1298) /gb=X65018 /gi=34766 /ug=Hs.153415 /len=1400
537	AI446234	31776_at	Cluster Incl. AI446234:tj25g10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-2142594 /clone_end=3 /gb=AI446234 /gi=4293782 /ug=Hs.156072 /len=368
538	AF006464	31777_at	Cluster Incl. AF006464:Homo sapiens muscle specific tyrosine kinase receptor (MUSK) mRNA, complete cds /cds=(46,2655) /gb=AF006464 /gi=2253311 /ug=Hs.156465 /len=2666
539	U34802	31778_at	Cluster Incl. U34802:Human intrinsic membrane protein MP70 (Cx50) gene, complete cds /cds=(0,1298) /gb=U34802 /gi=1002998 /ug=Hs.157433 /len=1299
540	AF084199	31779_s_at	Cluster Incl. AF084199:Homo sapiens beta-interferon gene positive-regulatory domain I binding factor (BLIMP1) mRNA, complete cds /cds=(223,2592) /gb=AF084199 /gi=3493157 /ug=Hs.158303 /len=2615
541	AC006293	31780_f_at	Cluster Incl. AC006293:Homo sapiens chromosome 19, cosmid F15658 /cds=(508,1740) /gb=AC006293 /gi=4106979 /ug=Hs.158310 /len=2113
542	U65652	31781_at	Cluster Incl. U65652:Homo sapiens HREP protein (C17ORF1) mRNA, complete cds /cds=(221,979) /gb=U65652 /gi=2909658 /ug=Hs.158313 /len=1298
543	U31099	31782_at	Cluster Incl. U31099:Human DP prostanoid receptor (PTGDR) mRNA, partial cds /cds=(0,435) /gb=U31099 /gi=940376 /ug=Hs.158326 /len=495
544	U52112	31783_at	Cluster Incl. U52112:renin-binding protein /cds=(191,1444) /gb=U52112 /gi=1302657 /ug=Hs.158331 /len=1461
545	U86759	31784_at	Cluster Incl. U86759:Human netrin-2 like protein (NTN2l) mRNA, complete cds /cds=(203,1945) /gb=U86759 /gi=2052394 /ug=Hs.158336 /len=1986

Cluster ID	Database Accession	Affymetrix Accession	Entrez Definition
546	U92817	31785_f_at	Cluster Incl. U92817:Homo sapiens unnamed HERV-H protein mRNA, complete cds /cds=(154,306) /gb=U92817 /gi=2465327 /ug=Hs.158340 /len=458
547	AF040630	31905_at	Cluster Incl. AF040630:Homo sapiens galanin receptor GalR2 mRNA, complete cds /cds=(112,1275) /gb=AF040630 /gi=2921759 /ug=Hs.158351 /len=1340
548	AF068754	31906_at	Cluster Incl. AF068754:Homo sapiens heat shock factor binding protein 1 HSBP1 mRNA, complete cds /cds=(54,284) /gb=AF068754 /gi=3283408 /ug=Hs.158675 /len=534
549	D87735	31907_at	Cluster Incl. D87735:Homo sapiens mRNA for ribosomal protein L14, complete cds /cds=(17,679) /gb=D87735 /gi=1620021 /ug=Hs.158675 /len=722
550	U94354	31908_at	Cluster Incl. U94354:Human lunatic fringe mRNA, partial cds /cds=(0,775) /gb=U94354 /gi=2204350 /ug=Hs.159142 /len=1063
551	AB018297	31909_at	Cluster Incl. AB018297:Homo sapiens mRNA for KIAA0754 protein, partial cds /cds=(0,2105) /gb=AB018297 /gi=3882228 /ug=Hs.159183 /len=4044
552	U92021	31910_at	Cluster Incl. U92021:Human clone 255298 defective mariner transposon Hsmar2 mRNA sequence /cds=UNKNOWN /gb=U92021 /gi=2052463 /ug=Hs.159192 /len=1268
553	AF000989	31911_at	Cluster Incl. AF000989:Homo sapiens thymosin beta 4 Y isoform (TB4Y) mRNA, complete cds /cds=(767,901) /gb=AF000989 /gi=2580563 /ug=Hs.159201 /len=1161
554	AF076292	31912_at	Cluster Incl. AF076292:Homo sapiens TGF-beta/activin signal transducer FAST-1p (FAST1) mRNA, complete cds /cds=(579,1676) /gb=AF076292 /gi=3523161 /ug=Hs.159251
555	AF049703	31913_at	Cluster Incl. AF049703:Homo sapiens E74-like factor 5 (ELF5) mRNA, complete cds /cds=(109,876) /gb=AF049703 /gi=3928981 /ug=Hs.159267 /len=898
556	AF054177	31914_at	Cluster Incl. AF054177:Homo sapiens chromodomain-helicase-DNA-binding protein mRNA, complete cds /cds=(98,760) /gb=AF054177 /gi=3341997 /ug=Hs.159273 /len=1147
557	U43519	31915_at	Cluster Incl. U43519:Human dystrophin-related protein 2 (DRP2) mRNA, complete cds /cds=(327,3191) /gb=U43519 /gi=1353781 /ug=Hs.159291 /len=3499
558	AF047445	31916_at	Cluster Incl. AF047445:Homo sapiens Ly-49L (LY49L) mRNA, complete cds /cds=(96,743) /gb=AF047445 /gi=3608369 /ug=Hs.159297 /len=1243
559	AC005609	31917_at	Cluster Incl. AC005609:Homo sapiens chromosome 5, BAC clone 203o13 (LBNL H155), complete sequence /cds=(0,2423) /gb=AC005609 /gi=3540156 /ug=Hs.159402 /len=2424
560	U44060	31918_at	Cluster Incl. U44060:Human homeodomain protein (Prox 1) mRNA, complete cds /cds=(606,2816) /gb=U44060 /gi=1511629 /ug=Hs.159437 /len=2924
561	AF002986	31919_at	Cluster Incl. AF002986:Homo sapiens platelet activating receptor homolog (H963) mRNA, complete cds /cds=(219,1178) /gb=AF002986 /gi=2580587 /ug=Hs.159545 /len=1249
562	AF004841	31920_at	Cluster Incl. AF004841:Homo sapiens GDO mRNA, complete cds /cds=(0,3722) /gb=AF004841 /gi=3559765 /ug=Hs.159565 /len=3986
563	U56421	31921_at	Cluster Incl. U56421:Human olfactory receptor (OLF3) gene, complete cds /cds=(0,953) /gb=U56421 /gi=1336042 /ug=Hs.159898 /len=954
564	U60269	31922_i_at	Cluster Incl. U60269:Human endogenous retrovirus HERV-K(HML6) proviral clone HML6.17 putative polymerase and envelope genes, partial cds, and 3LTR /cds=(0,491) /gb=U60269 /gi=1408208 /ug=Hs.159902 /len=492

Cluster ID	Database Accession	Accession	Entrez Definition
565	U60269	31923_f_at	Cluster Incl. U60269:Human endogenous retrovirus HERV-K(HML6) proviral clone HML6.17 putative polymerase and envelope genes, partial cds, and 3LTR /cds=(0,491) /gb=U60269 /gi=1408208 /ug=Hs.159902 /len=492
566	AL035122	31924_at	Cluster Incl. AL035122:H.sapiens gene from PACs 295C6 and 313L4 /cds=(949,2067) /gb=AL035122 /gi=4153950 /ug=Hs.163045 /len=2567
567	L26584	31925_s_at	Cluster Incl. L26584:Human (CDC25) mRNA, complete cds /cds=(0,3827) /gb=L26584 /gi=433719 /ug=Hs.1636 /len=3972
568	M93133	31926_at	Cluster Incl. M93133:Human cholesterol 7-alpha hydroxylase (CYP7) mRNA, complete cds /cds=(59,1573) /gb=M93133 /gi=181318 /ug=Hs.1644 /len=2869
569	D86062	31927_s_at	Cluster Incl. D86062:Human mRNA for KNP-1b, complete cds /cds=(18,731) /gb=D86062 /gi=1545814 /ug=Hs.166012 /len=898
570	X63454	31928_at	Cluster Incl. X63454:H.sapiens hst-2 (FGF-6) mRNA /cds=(74,670) /gb=X63454 /gi=32490 /ug=Hs.166015 /len=744
571	X76092	31929_at	Cluster Incl. X76092:H.sapiens HRFX3 mRNA /cds=(8,2131) /gb=X76092 /gi=452403 /ug=Hs.166019 /len=2514
572	X63096	31930_f_at	Cluster Incl. X63096:H.sapiens mRNA for rhesus polypeptide (RhVIII) /cds=(6,806) /gb=X63096 /gi=36044 /ug=Hs.166021
573	AI632247	31931_f_at	Cluster Incl. AI632247:tt20f10.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-2241355 /clone_end=3 /gb=AI632247 /gi=4683577 /ug=Hs.166021 /len=482
574	M90357	31932_f_at	Cluster Incl. M90357:Human basic transcription factor 3a (BTF3a) gene /cds=(0,476) /gb=M90357 /gi=457435 /ug=Hs.166033 /len=487
575	M90357	31933_r_at	Cluster Incl. M90357:Human basic transcription factor 3a (BTF3a) gene /cds=(0,476) /gb=M90357 /gi=457435 /ug=Hs.166033 /len=487
576	Z48614	31934_at	Cluster Incl. Z48614:H.sapiens MADM gene (exon 1) /cds=UNKNOWN /gb=Z48614 /gi=728589 /ug=Hs.166038
577	U75968	31935_s_at	Cluster Incl. U75968:Human clone C3 CHL1 protein (CHLR1) mRNA, alternatively spliced, complete cds /cds=(220,1086) /gb=U75968 /gi=1666894 /ug=Hs.166048 /len=4182
578	AB007890	31936_s_at	Cluster Incl. AB007890:Homo sapiens KIAA0430 mRNA, complete cds /cds=(0,3172) /gb=AB007890 /gi=2887438 /ug=Hs.166163 /len=6011
579	AA526011	31937_at	Cluster Incl. AA526011:ni58d01.s1 Homo sapiens cDNA /clone=IMAGE-981025 /gb=AA526011 /gi=2268080 /ug=Hs.166163 /len=506
580	AA526011	31938_g_at	Cluster Incl. AA526011:ni58d01.s1 Homo sapiens cDNA /clone=IMAGE-981025 /gb=AA526011 /gi=2268080 /ug=Hs.166163 /len=506
581	AB011122	31939_at	Cluster Incl. AB011122:Homo sapiens mRNA for KIAA0550 protein, complete cds /cds=(729,3683) /gb=AB011122 /gi=3043623 /ug=Hs.166201 /len=5749
582	AJ010228	31940_s_at	Cluster Incl. AJ010228:Homo sapiens mRNA for RET finger protein-like 1, long variant /cds=(296,1162) /gb=AJ010228 /gi=3417311 /ug=Hs.167750 /len=1496
583	AJ010232	31941_s_at	Cluster Incl. AJ010232:Homo sapiens mRNA for RET finger protein-like 3 /cds=(292,1158) /gb=AJ010232 /gi=3417318 /ug=Hs.167751 /len=1488
584	AF045583	31942_at	Cluster Incl. AF045583:Homo sapiens tubby like protein 3 (TULP3) mRNA, complete cds /cds=(45,1373) /gb=AF045583 /gi=3372492 /ug=Hs.169084 /len=1482
585	AF045583	31943_g_at	Cluster Incl. AF045583:Homo sapiens tubby like protein 3 (TULP3) mRNA, complete cds /cds=(45,1373) /gb=AF045583 /gi=3372492 /ug=Hs.169084 /len=1482

Cluster ID	Database Accession	Affymetrix Accession	Entrez Definition
586	AI028290	31944_at	Cluster Incl. AI028290:ov84f11.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-1644045 /clone_end=3 /gb=AI028290 /gi=3245599 /ug=Hs.169084 /len=512
587	AF100539	31945_s_at	Cluster Incl. AF100539:Homo sapiens SH2 domain protein 1A isoform B (SH2D1A) mRNA, complete cds /cds=(54,284) /gb=AF100539 /gi=3928893 /ug=Hs.169209 /len=483
588	X74143	31946_s_at	Cluster Incl. X74143:H.sapiens HBF-2 mRNA for transcription factor /cds=(201,1610) /gb=X74143 /gi=516382 /ug=Hs.169277
589	X74143	31947_r_at	Cluster Incl. X74143:H.sapiens HBF-2 mRNA for transcription factor /cds=(201,1610) /gb=X74143 /gi=516382 /ug=Hs.169277
590	X79563	31948_at	Cluster Incl. X79563:H.sapiens 8.2kDa differentiation factor mRNA /cds=(60,353) /gb=X79563 /gi=499069 /ug=Hs.169301 /len=465
591	S62035	31949_at	Cluster Incl. S62035:Ras-specific guanine nucleotide-releasing factor=CDC25 homolog [human, brain stem, mRNA, 2861 nt] /cds=(1182,2651) /gb=S62035 /gi=386046 /ug=Hs.169350 /len=2861
592	Y00345	31950_at	Cluster Incl. Y00345:Human mRNA for polyA binding protein /cds=(502,2403) /gb=Y00345 /gi=35569 /ug=Hs.172182 /len=2848
593	Z48501	31951_s_at	Cluster Incl. Z48501:H.sapiens mRNA for polyadenylate binding protein II /cds=(0,1568) /gb=Z48501 /gi=693936 /ug=Hs.172182 /len=1569
594	X69391	31952_at	Cluster Incl. X69391:H.sapiens mRNA for ribosomal protein L6 /cds=(26,892) /gb=X69391 /gi=36137 /ug=Hs.174131 /len=926
595	U19144	31953_f_at	Cluster Incl. U19144:Human GAGE-3 protein mRNA, complete cds /cds=(99,455) /gb=U19144 /gi=914902 /ug=Hs.176661 /len=552
596	AA447559	31954_f_at	Cluster Incl. AA447559:zw81e11.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-782636 /clone_end=3 /gb=AA447559 /gi=2161229 /ug=Hs.176661 /len=489
597	X65923	31955_at	Cluster Incl. X65923:H.sapiens fau mRNA /cds=(56,457) /gb=X65923 /gi=31302 /ug=Hs.177415 /len=508
598	M17886	31956_f_at	Cluster Incl. M17886:Human acidic ribosomal phosphoprotein P1 mRNA, complete cds /cds=(129,473) /gb=M17886 /gi=190233 /ug=Hs.177592 /len=512
599	M17886	31957_r_at	Cluster Incl. M17886:Human acidic ribosomal phosphoprotein P1 mRNA, complete cds /cds=(129,473) /gb=M17886 /gi=190233 /ug=Hs.177592 /len=512
600	AA347932	31958_i_at	Cluster Incl. AA347932:EST54334 Homo sapiens cDNA, 3 end /clone=ATCC-149018 /clone_end=3 /gb=AA347932 /gi=2000189 /ug=Hs.177820 /len=258
601	AF029900	31959_at	Cluster Incl. AF029900:Homo sapiens ADAM 21 mRNA, partial cds /cds=(0,1628) /gb=AF029900 /gi=2739136 /ug=Hs.178748 /len=1850
602	U19143	31960_f_at	Cluster Incl. U19143:Human GAGE-2 protein mRNA, complete cds /cds=(83,433) /gb=U19143 /gi=914900 /ug=Hs.180976 /len=530
603	AF070579	31961_r_at	Cluster Incl. AF070579:Homo sapiens clone 24487 mRNA sequence /cds=UNKNOWN /gb=AF070579 /gi=3387951 /ug=Hs.181544 /len=1565
604	L06499	31962_at	Cluster Incl. L06499:Homo sapiens ribosomal protein L37a (RPL37A) mRNA, complete cds /cds=(17,295) /gb=L06499 /gi=292438 /ug=Hs.184109 /len=357
605	M19684	31963_at	Cluster Incl. M19684:Human alpha-1-antitrypsin-related protein gene /cds=(0,1262) /gb=M19684 /gi=177813 /ug=Hs.184929 /len=1263
606	AF003738	31964_at	Cluster Incl. AF003738:Homo sapiens chromosome 21q22.1 anonymous mRNA sequence /cds=UNKNOWN /gb=AF003738 /gi=2197087 /ug=Hs.184943 /len=1356
607	AC004262	31965_at	Cluster Incl. AC004262:Homo sapiens chromosome 19, cosmid R29368 /cds=(0,1034) /gb=AC004262 /gi=2935596 /ug=Hs.184950 /len=1035

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608	S48220	31966_at	Cluster Incl. S48220:type I 5 iodothyronine deiodinase [human, mRNA, 2222 nt] /cds=(6,755) /gb=S48220 /gi=257451 /ug=Hs.186571 /len=2222
609	AF035835	31967_at	Cluster Incl. AF035835:Homo sapiens nephrin (NPHS1) mRNA, complete cds /cds=(0,3725) /gb=AF035835 /gi=3025698 /ug=Hs.190311 /len=4285
610	U46120	31968_at	Cluster Incl. U46120:Human expressed unknown mRNA /cds=UNKNOWN /gb=U46120 /gi=1184779 /ug=Hs.193392 /len=1767
611	AF023456	31969_i_at	Cluster Incl. AF023456:Homo sapiens protein phosphatase with EF-hands-2 long form (PPEF-2) mRNA, complete cds /cds=(367,2628) /gb=AF023456 /gi=2586412 /ug=Hs.193558
612	AF023456	31970_r_at	Cluster Incl. AF023456:Homo sapiens protein phosphatase with EF-hands-2 long form (PPEF-2) mRNA, complete cds /cds=(367,2628) /gb=AF023456 /gi=2586412 /ug=Hs.193558
613	AF008192	31971_at	Cluster Incl. AF008192:Homo sapiens putative GR6 protein (GR6) mRNA, complete cds /cds=(967,1416) /gb=AF008192 /gi=2291083 /ug=Hs.194283 /len=2700
614	U92971	31972_at	Cluster Incl. U92971:Human protease-activated receptor 3 (PAR3) mRNA, complete cds /cds=(144,1268) /gb=U92971 /gi=1938374 /ug=Hs.194351 /len=1830
615	AB012043	31973_at	Cluster Incl. AB012043:Homo sapiens mRNA for NBR13, complete cds /cds=(28,4695) /gb=AB012043 /gi=4115531 /ug=Hs.194746 /len=4903
616	AJ132445	31974_at	Cluster Incl. AJ132445:Homo sapiens CLDN14 gene /cds=(0,719) /gb=AJ132445 /gi=4186039 /ug=Hs.194758 /len=720
617	X55503	31975_at	Cluster Incl. X55503:H.sapiens pseudogene for metallothionein and AG/CT repetitive element /cds=(0,157) /gb=X55503 /gi=38328 /ug=Hs.194762 /len=503
618	X81333	31976_at	Cluster Incl. X81333:H.sapiens mRNA for PPH beta subunit protein /cds=(46,2148) /gb=X81333 /gi=557645 /ug=Hs.194777
619	M92432	31977_at	Cluster Incl. M92432:Homo sapiens retinal guanylyl cyclase (retGC) mRNA, complete cds /cds=(73,3384) /gb=M92432 /gi=623414 /ug=Hs.1974 /len=3621
620	AB012722	31978_at	Cluster Incl. AB012722:Homo sapiens gene for kinesin-like protein, complete cds /cds=(94,1248) /gb=AB012722 /gi=4115550 /ug=Hs.198256 /len=1342
621	D49818	31979_at	Cluster Incl. D49818:Homo sapiens mRNA for 6-phosphofructo-2-kinase/fructose-2, 6-bisphosphatase, complete cds /cds=(26,1435) /gb=D49818 /gi=1905760 /ug=Hs.198278
622	Y11739	31980_at	Cluster Incl. Y11739:H.sapiens mRNA for whn transcription factor /cds=(29,1975) /gb=Y11739 /gi=2315191 /ug=Hs.198313
623	Z99916	31981_at	Cluster Incl. Z99916:Human DNA sequence from clone 221G9 on chromosome 22q11.21-12.2. Contains the CRYBB3 and CRYBB2 genes for Beta Crystallin B3 and B2, and a Stathmin family (SCG10-like, SCLIP, Neuroplasticin-2, NCP2) pseudogene. Contains ESTs and GSSs /cds=(72,707) /gb=Z99916 /gi=4158122 /ug=Hs.198968 /len=853
624	AB020701	31982_at	Cluster Incl. AB020701:Homo sapiens mRNA for KIAA0894 protein, complete cds /cds=(168,1769) /gb=AB020701 /gi=4240276 /ug=Hs.199117 /len=4113
625	X68880	31983_at	Cluster Incl. X68880:H.sapiens EMX2 mRNA /cds=(0,476) /gb=X68880 /gi=31141 /ug=Hs.202095 /len=477
626	AC006293	31984_at	Cluster Incl. AC006293:Homo sapiens chromosome 19, cosmid F15658 /cds=(378,1484) /gb=AC006293 /gi=4106979 /ug=Hs.202680 /len=1683
627	AL049228	31985_at	Cluster Incl. AL049228:Homo sapiens mRNA; cDNA DKFZp564N1716 (from clone DKFZp564N1716) /cds=UNKNOWN /gb=AL049228 /gi=4499959 /ug=Hs.202681 /len=1619