

20000238

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「研究成果の刊行に関する一覧表」をご参照ください。

「研究成果の刊行に関する一覧表」

【V】小腸EST情報

09.seq	378	gi 4506692 ref NM_001021.1 RPSP17	Homo sapiens ribosomal protein S17 (RPS17) mRNA	0
11.seq	444	gi 532596 gb U12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-151
14.seq	358	gi 5821114 dbj AB017710.1 AB0177	Homo sapiens U50HG genes for U50' snoRNA and U50 snoRNA, complete	1.00E-37
16.seq	525	gi 2576344 gb AC002400 HUAC0024	Human Chromosome 16 BAC clone CIT9875K-A-73566, complete sequen	E-130
17.seq	441	gi 4314418 gb AC006487 AC006487	Homo sapiens chromosome 17, clone hRPC.1029_K_10, complete sequen	0
18.seq	385	gi 4506696 ref NM_001023.1 RPS20	Homo sapiens ribosomal protein S20 (RPS20) mRNA	0
19.seq	336	gi 2529723 gb AF001893 BETA2	Human MEN1 region clone epsilon/beta mRNA, 3' fragment	E-174
20.seq	344	gi 5032086 ref NM_005877.1 SF3A1	Homo sapiens pre-mRNA splicing factor SF3a (120 kDa subunit),	E-172
22.seq	298	gi 4505656 ref NM_002599.1 PDE2A	Homo sapiens phosphodiesterase 2A, cGMP-stimulated (PDE2A) mRNA	5.00E-86
25.seq	471	gi 33949 emb X68742 HSINTASA	H.sapiens mRNA for integrin, alpha subunit	6.00E-87
27.seq	254	gi 150120 gb K02075 MXARSSMS	Myxococcus xanthus satellite multicopy single-stranded DNA (msDNA).	0.25
28.seq	305	gi 4775623 emb AL035594.7 HS67A	Human DNA sequence from clone 67A5 on chromosome 6q22.31-22.33	E-130
29.seq	422	gi 3970932 gb AC006130 AC006130	Homo sapiens chromosome 19, BAC CIT-8-191n6, complete sequence [H	5.00E-13
31.seq	862	gi 5032088 ref NM_005626.1 SFRS4	Homo sapiens splicing factor, arginine/serine-rich 4 (SFRS4) mRNA	2.00E-94
32.seq	410	gi 1834938 emb Z85227 HSZ85227	H.sapiens Ig lambda light chain variable region gene (9-150IA119)	E-135E
34.seq	460	gi 5262556 emb AL080123.1 HSM80	Homo sapiens mRNA; cDNA DKFZp569D2231 (from clone DKFZp569D22	0
35.seq	569	gi 434050 gb L12535 HUMRSU1A	Human RSU-1/RSP-1 mRNA, complete cds.	0
36.seq	406	gi 186628 gb U77663 HUMK10A	Human keratin 10 mRNA, 3' end.	E-169
37.seq	297	gi 13003 emb V00662 MHSXX	H.sapiens mitochondrial genome	E-160
40.seq	431	gi 468242 gb U07989 HSU07989	Human Burkitt's lymphoma immunoglobulin kappa light chain mRNA,	E-148
41.seq	524	gi 532596 gb U12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-137
43.seq	479	gi 4506638 ref NM_000996.1 RPL35	Homo sapiens ribosomal protein L35a (RPL35A) mRNA	0
44.seq	437	gi 4808971 gb AF128894.1 HSTERT2	Homo sapiens telomerase reverse transcriptase (TERT) gene, exons 7-16	0.44
45.seq	415	gi 4506700 ref NM_001025.1 RPS23	Homo sapiens ribosomal protein S23 (RPS23) mRNA	0
47.seq	527	gi 3165402 gb AC004752 AC004752	Homo sapiens chromosome 5, BAC clone Be5 (LBNL H167), complete seq	0
48.seq	414	gi 4506596 ref NM_000976.1 RPL12	Homo sapiens ribosomal protein L12 (RPL12) mRNA	0
51.seq	273	gi 4753237 gb AC006372.2 AC00637	Homo sapiens clone NH0331D05, complete sequence	E-127
52.seq	494	gi 4914614 emb AL050393.1 HSM80	Homo sapiens mRNA; cDNA DKFZp586F0420 (from clone DKFZp586F042	0
53.seq	490	gi 4506696 ref NM_001023.1 RPS20	Homo sapiens ribosomal protein S20 (RPS20) mRNA	0
54.seq	372	gi 2358100 gb AE000665 MMAF0006	Mus musculus TCR beta locus from bases 501860 to 700960 (section 3 c	0.37
55.seq	367	gi 5231140 gb AF153608.1 AF15360	Homo sapiens sin3 associated polypeptide (SAP18) mRNA, complete cds	0
57.seq	455	gi 5016088 ref NM_001101.2 ACTB	Homo sapiens actin, beta (ACTB) mRNA	0
59.seq	355	gi 2429080 gb U087675 U087675	Homo sapiens DNA for amyloid precursor protein, complete cds	0.36
60.seq	580	gi 4507030 ref NM_000343.1 SLCSA	Homo sapiens solute carrier family 5 (sodium/glucose	0
61.seq	861	gi 5174726 ref NM_005998.1 TRIC5	Homo sapiens TCP1 (t-complex-1) ring complex, polypeptide 5 (TRIC5)	0
64.seq	399	gi 307150 gb L19182 HUMMAC25X	Human MAC25 mRNA, complete cds.	E-124
66.seq	524	gi 179831 gb U94859 HUMCALIEF	Human calnexin mRNA, complete cds.	0
67.seq	492	gi 4589591 dbj AB023191.1 AB0231	Homo sapiens mRNA for KIAA0974 protein, partial cds	0
68.seq	857	gi 307150 gb L19182 HUMMAC25X	Human MAC25 mRNA, complete cds.	E-117
69.seq	851	gi 4760290 emb AL031346.8 HS597	Human DNA sequence from clone 597B2 on chromosome 22q13.2-13.32	0.057
70.seq	471	gi 992692 gb U33324 CCU33324	Caulobacter crescentus RNase PH (rph) gene, Grp (grpE) gene and	0.031
72.seq	523	gi 348908 gb L22548 HUMCOL18AX	Human collagen type XVIII alpha 1 (COL18A1) mRNA, partial cds.	E-159
73.seq	454	gi 186042 gb U74019 HUMIGKW	Human Ig rearranged light-chain mRNA V region.	0
77.seq	409	gi 244237 gb S78624 S78624	YCR591...YCR522 [Saccharomyces cerevisiae=yeast, Genomic, 4 genes,	0.1
80.seq	368	gi 5565976 gb AF152462.1 AF1524	Homo sapiens transmembrane protein BRI (BRI) mRNA, complete cds	E-139
81.seq	538	gi 4679019 gb JAF07203.1 JAF0720	Homo sapiens HSPC017 mRNA, complete cds	0
82.seq	483	gi 2656027 dbj JAB00905.1 JAB00905	Arabidopsis thaliana genomic DNA, chromosome 5, P1 clone: MJH22	0.12
83.seq	419	gi 31091 emb J16869 HSEF1AC	Human mRNA for elongation factor 1-alpha (clone CEF4)	0
85.seq	404	gi 434764 dbj D21262 HUMORFGA	Human mRNA for KIAA0035 gene, partial cds	0
86.seq	537	gi 3165402 gb AC004752 AC004752	Homo sapiens chromosome 5, BAC clone Be5 (LBNL H167), complete seq	2.00E-44
88.seq	544	gi 1590807 dbj U087914 U087914	Human mRNA for ornithine decarboxylase antizyme, complete cds	0
89.seq	300	gi 181532 gb U97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	E-110
91.seq	496	gi 5679530 emb AL050305.9 HSDA6	Human DNA sequence from clone 60B16 on chromosome Xp11.21-22.2,	0.008
92.seq	838	gi 532596 gb U12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-174
94.seq	838	gi 4505604 ref NM_002580.1 PAP	Homo sapiens pancreatitis-associated protein (PAP) mRNA	0
95.seq	831	gi 202922 gb L11002 RATANKBIND	Rat ankyrin binding glycoprotein-1 related mRNA sequence.	5.00E-61
96.seq	851	gi 1009231 gb L38828 TOBSUP1NT	Nicotiana tabacum EF-1-alpha-related GTP-binding protein (SUP1) mRNA,	0.22
97.seq	853	gi 532596 gb U12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-103
98.seq	797	gi 23690 emb X56932 HS23KDHP	H.sapiens mRNA for 23 kD highly basic protein	E-154
99.seq	887	gi 181070 gb U33146 HUMCRPR	Human cysteine-rich peptide mRNA, complete cds.	0
100.seq	821	gi 13003 emb V00662 MHSXX	H.sapiens mitochondrial genome	0
101.seq	861			
103.seq	875	gi 4504618 ref NM_001553.1 IGFBP7	Homo sapiens insulin-like growth factor binding protein 7 (IGFBP7)	0
107.seq	919	gi 3962496 gb AC006117 AC006117	Homo sapiens chromosome 19, cosmid F20720, complete sequence [Homo	1E-123
108.seq	858	gi 1262342 emb X93334 HSMITG	H.sapiens mitochondrial DNA, complete genome	0
109.seq	879	gi 292446 gb U77234 HUMRPSA3A	Human ribosomal protein S3a mRNA, complete cds.	0
110.seq	848			
111.seq	898	gi 1405347 dbj D45132 HUMHOXY1	Homo sapiens mRNA for zinc-finger DNA-binding protein, complete cds	1E-154
112.seq	828	gi 181532 gb U97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	1E-132
114.seq	879			
115.seq	848	gi 530070 gb U12691 HSU12691	Human Hsa3 mitochondrion cytochrome oxidase subunit II (COII) gene.	1E-175
118.seq	863			
119.seq	898	gi 1209254 gb L20311 BOVIGJMRNA	Bos taurus mu- and alpha-immunoglobulin-associated J-chain mRNA,	5.00E-95
120.seq	973	gi 181532 gb U97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	1E-101
122.seq	924	gi 4505408 ref NM_002512.1 NME2	Homo sapiens non-metastatic cells 2, protein (NM23B) expressed in	0
124.seq	892	gi 4929722 gb AF151885.1 AF151885	Homo sapiens CGI-127 protein mRNA, complete cds	0
126.seq	897			
130.seq	888	gi 4506622 ref NM_000988.1 RPL27	Homo sapiens ribosomal protein L27 (RPL27) mRNA	0
131.seq	887	gi 2055294 dbj D87438 D87438	Human mRNA for KIAA0251 gene, partial cds	0
133.seq	868	gi 187762 gb U26432 HUMMHC1B	Human MHC class I HLA-C.1 gene, complete cds.	0
134.seq	988	gi 4506834 ref NM_002989.1 SCYA21	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member	0
135.seq	1001			
136.seq	966	gi 598836 dbj D16992 HUMD2D02M5	Human HepG2 partial cDNA, clone hmd2d02m5	1E-109
137.seq	874			
140.seq	876			
141.seq	972			
142.seq	982	gi 3928036 gb AC005865 AC005865	Homo sapiens 12p13.3 PAC RPC15-1063M23 (Roswell Park Cancer Institute	0
143.seq	886	gi 4503456 ref NM_001401.1 EDG2	Homo sapiens endothelial differentiation, lysophasphatidic acid	0
145.seq	912	gi 30091 emb X03963 HSCOLA1C	Human procollagen alpha-1 (IV) mRNA 3' untranslated region (UTR)	0
148.seq	881			
149.seq	835	gi 4503184 ref NM_001916.1 CYC1	Homo sapiens cytochrome c-1 (CYC1) mRNA >gi 30302 emb X06994 HSCYC	0
150.seq	845			
151.seq	878	gi 4506684 ref NM_001017.1 RPS13	Homo sapiens ribosomal protein S13 (RPS13) mRNA	0
152.seq	972	gi 927447 emb X88832 MTHSLAS44	H.sapiens mitochondrial DNA for loop attachment sequence (clone	1E-156
153.seq	862	gi 1495465 emb X97260 HSMTISO2	H.sapiens mRNA for metallothionein isoform 2	1E-172
154.seq	862			
155.seq	865	gi 261239 gb U50732 U50732	immunoglobulin M light chain V region=anti-lipid A antibody (human,	0
156.seq	850			
157.seq	857	gi 4507894 ref NM_003380.1 VIM	Homo sapiens vimentin (VIM) mRNA >gi 37851 emb Z19554 HSVIMENTA	0
158.seq	865	gi 4758543 ref NM_004500.1 HNRPC	Homo sapiens heterogeneous nuclear ribonucleoprotein C (C1/C2)	0
160.seq	856	gi 4506722 ref NM_001006.1 RPS3A	Homo sapiens ribosomal protein S3A (RPS3A) mRNA	0
161.seq				
162.seq	862	gi 4929576 gb AF151812.1 AF151812	Homo sapiens CGI-54 protein mRNA, complete cds	0
163.seq	839	gi 2335046 dbj D17409.1 D17409	Homo sapiens mRNA for SM22 alpha, complete cds	0

164.seq	841					
165.seq	873	gi 5870425 emb AJ249366.1 HSA249366	Homo sapiens mRNA for epsilon-COP protein and epsilon-COP protein			0
166.seq						
167.seq	830	gi 2583174 gb AF026844.1 AF026844	Homo sapiens ribosomal protein L41 mRNA, complete cds			0
169.seq	837	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.			0
170.seq	868	gi 4504964 ref NM_002298.1 LCP1	Homo sapiens lymphocyte cytosolic protein 1 (L-plastin) (LCP1) mRNA			0
171.seq	840	gi 2339843 gb U95742 HSU95742	Human chromosome 16p13.1 BAC clone CIT9875K-551G9 complete sequence			0
172.seq	865					
173.seq	817					
175.seq	833	gi 2335046 dbj D17409.1 D17409	Homo sapiens mRNA for SM22 alpha, complete cds			0
176.seq	853	gi 2894828 gb U91942 U91942	Homo sapiens immunoglobulin light chain mRNA, partial cds			0
178.seq						
179.seq	826	gi 1770459 emb X98261 HSMPP5	H.sapiens mRNA for M-phase phosphoprotein, mpp5			1E-158
180.seq	828					
181.seq	827	gi 5453671 ref NM_006418.1 GW112	Homo sapiens differentially expressed in hematopoietic lineages			0
182.seq	834					
183.seq	831	gi 186969 gb M21731 HUMLC5	Human lipocortin-V mRNA, complete cds.			0
184.seq	844	gi 4509304 ref NM_001926.1 DEFA6	Homo sapiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA			0
185.seq	823					
186.seq	828	gi 30029 emb X15880 HSCOL1C	Human mRNA for collagen VI alpha-1 C-terminal globular domain			0
187.seq	816	gi 4759055 ref NM_004587.1 RRBP1	Homo sapiens ribosome binding protein 1 (dog 180kD homolog) (RRBP1)			2.00E-54
188.seq	804	gi 5032050 ref NM_005617.1 RPS14B	Homo sapiens ribosomal protein S14 (RPS14) mRNA			0
189.seq	822	gi 31091 emb X18689 HSEF1AC	Human mRNA for elongation factor 1-alpha (clone CEF4)			0
191.seq	818	gi 184656 gb M77804 HUMIFNWR5	Human tryptophanyl tRNA synthetase (IFNWR5) mRNA, complete cds.			0
192.seq	839	gi 2281706 gb AF014838 AF014838	Homo sapiens galectin-4 mRNA, complete cds			0
193.seq	309	gi 4506899 ref NM_003011.1 SET	Homo sapiens SET translocation (myeloid leukemia-associated) (SET)			1E-123
194.seq	439	gi 4506636 ref NM_000995.1 RPL34	Homo sapiens ribosomal protein L34 (RPL34) mRNA			0
195.seq	381	gi 5911928 emb AL117465.1 HSM80097	Homo sapiens mRNA; cDNA DKFZp586i0322 (from clone DKFZp586i0322)			0
196.seq	480					
197.seq	414					
198.seq	496	gi 288099 emb X55733 HSINTFA4B	H.sapiens initiation factor 4B cDNA			1E-171
199.seq	464	gi 2341021 gb AC002477 AC002477	Human PAC clone DJ327A19 from Xq25-q26, complete sequence [Homo sapiens]			9.00E-55
200.seq	454					
201.seq	405	gi 3043681 dbj AB011151 AB011151	Homo sapiens mRNA for KIAA0579 protein, partial cds			0
202.seq	360	gi 13003 emb V00662 MHSSXX	H.sapiens mitochondrial genome			1E-116
203.seq	410					
204.seq	333	gi 36506 emb X69292 HSSMMHC	H.sapiens mRNA for smooth muscle myosin			0
205.seq	433	gi 4503470 ref NM_001402.1 EEF1A1	Homo sapiens eukaryotic translation elongation factor 1 alpha 1			0
206.seq	475	gi 1209254 gb M12031 BOVIGJMRNA	Bos taurus mu- and alpha-immunoglobulin-associated J-chain mRNA.			1E-102
207.seq	449	gi 1237214 emb X96754 HSAKAPALC	H.sapiens gene encoding kappa light chain constant region			1E-177
209.seq	186	gi 340233 gb M25246 HUMVIM3	Human vimentin (HuVim3) mRNA, 3' end.			6.00E-60
210.seq	406	gi 4758873 ref NM_004800.1 P76	Homo sapiens 76 kDa membrane protein (P76) mRNA			0
211.seq	356	gi 4504278 ref NM_002107.1 H3F3A	Homo sapiens H3 histone, family 3A (H3F3A) mRNA			1E-139
212.seq	280	gi 182355 gb M10050 HUMFABPL	Human liver fatty acid binding protein (FABP) mRNA, complete cds.			1E-148
213.seq	356	gi 13003 emb V00662 MHSSXX	H.sapiens mitochondrial genome			1E-135
214.seq	153	gi 4502959 ref NM_000090.1 COL3A1	Homo sapiens collagen, type III, alpha 1 (Ehlers-Danlos syndrome type I)			1.00E-51
217.seq	419	gi 4689129 gb AF07041.1 AF07041	Homo sapiens SH002 mRNA, complete cds			0
218.seq	380	gi 4505316 ref NM_002480.1 MYPT1	Homo sapiens myosin phosphatase, target subunit 1 (MYPT1) mRNA			1E-167
219.seq	365	gi 1834859 emb J285148 HSZ85148	H.sapiens Ig lambda light chain variable region gene (4'-29OI(A)95)			1E-151
221.seq	420					
222.seq	119	gi 285940 dbj D14658 HUMORF02	Human mRNA for KIAA0102 gene, complete cds			8.00E-49
223.seq	377	gi 4062176 gb AC006121 AC006121	Homo sapiens chromosome X, clone hCIT200_L_4, complete sequence			1E-121
224.seq	395	gi 644480 dbj D38112.1 HUMMTA	Homo sapiens mitochondrial DNA, complete sequence			1E-154
227.seq	128	gi 291843 gb L13434 HUMAGCGB	Human chromosome 3p21.1 gene sequence, complete cds.			2.00E-21
229.seq	502					
230.seq	431	gi 1136389 dbj D79986 D79986	Human mRNA for KIAA0164 gene, complete cds			1E-179
231.seq	402	gi 4502959 ref NM_000090.1 COL3A1	Homo sapiens collagen, type III, alpha 1 (Ehlers-Danlos syndrome)			1E-174
232.seq	457	gi 1913895 gb U90915 HSU90915	Human clone 23600 cytochrome c oxidase subunit IV mRNA, complete			1E-174
234.seq	398					
235.seq	428					
236.seq	401	gi 4502302 ref NM_001697.1 ATP5O1	Homo sapiens ATP synthase, H+ transporting, mitochondrial F1			1E-155
237.seq	408	gi 1021480 emb Z55439 HS40B5F	H.sapiens CpG island DNA genomic MseI fragment, clone 40B5, forward			6.00E-99
238.seq	440	gi 3327155 dbj AB014571 AB014571	Homo sapiens mRNA for KIAA0671 protein, complete cds			0
239.seq	472	gi 1209254 gb M2031 BOVIGJMRNA	Bos taurus mu- and alpha-immunoglobulin-associated J-chain mRNA.			2.00E-95
240.seq	384	gi 3183974 emb Y10183.1 HSMEMD	H.sapiens mRNA for MEMD protein			1E-128
241.seq	604					
244.seq	319	gi 4092059 gb AF054187 AF054187	Homo sapiens alpha NAC mRNA, complete cds			1E-166
245.seq	460	gi 189019 gb M22918 HUMMYLCB	Human non-muscle myosin alkali light chain mRNA, 3' end.			0
246.seq	325	gi 5174556 ref NM_005928.1 MFGE8B	Homo sapiens milk fat globule-EGF factor 8 protein (MFGE8) mRNA			1E-173
248.seq	428	gi 4972127 emb AL048829.2 CNS0000B	Human chromosome 14 DNA sequence *** IN PROGRESS *** BAC 124D2 of			0
249.seq	506	gi 339442 gb M81601 HUMTEFSII	Human transcription elongation factor (SII) mRNA, complete cds.			0
251.seq	481	gi 2662290 dbj AB009282 AB009282	Homo sapiens mRNA for cytochrome b5, partial cds			0
252.seq	424	gi 5835387 ref NC_001807.1	Human mitochondrial, complete genome >gi 1944628 gb J01415 HUMMTCCG			0
254.seq	485	gi 2832592 emb Z95116 HSU57G9	Human DNA sequence from BAC 57G9 on chromosome 22q12.1 Contains EST			0
255.seq	459	gi 4884267 emb AL050028.1 HSM800362	Homo sapiens mRNA; cDNA DKFZp566C0424 (from clone DKFZp566C0424)			0
256.seq						
258.seq	446	gi 3483018 emb AJ010569 HSA010569	Homo sapiens mRNA for GEF-2 protein			0
259.seq	481	gi 1049079 gb U30826 HSU30826	Human splicing factor Srp40-1 (Srp40) mRNA, complete cds.			1E-167
260.seq	154	gi 3859769 emb AJ011930 HSS1171	Homo sapiens chromosome 21q22.3, PAC clones 314N7, 225L15, BAC clone 7			2.00E-81
261.seq	535	gi 475791 ref NM_004346.1 CASP3	Homo sapiens caspase 3, apoptosis-related cysteine protease (CASP3)			1E-171
262.seq	491	gi 3287468 gb AC004448 AC004448	Homo sapiens chromosome 17, clone hCIT187M2, complete sequence [Homo			0
263.seq	438	gi 5763746 emb AL049766.14 HSDJ686	Human DNA sequence from clone 886N3 on chromosome 20q13.2-13.2.			1E-180
264.seq	519	gi 4504766 ref NM_002211.1 TGB1	Homo sapiens integrin, beta 1 (fibronectin receptor, beta			0
265.seq	549					
266.seq	443	gi 4757925 ref NM_004902.1 CC1.3f	Homo sapiens splicing factor (CC1.3) (CC1.3) mRNA			0
267.seq	450	gi 1663763 gb U66359 HSU66359	Human T54 protein (T54) mRNA, complete cds			0
268.seq	480	gi 2072143 gb U67093 HSU67093	Human ataxia-telangiectasia locus protein (ATM) mRNA, 3'UTR sequence			0
269.seq	553					
270.seq	451	gi 3954884 emb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-rhD,			0
271.seq	527	gi 4377992 gb AF075242 AF075242	Homo sapiens tumor transforming protein 1 (TUTR1) mRNA, complete			0
272.seq	375	gi 4757799 ref NM_004318.1 ASPH	Homo sapiens aspartate beta-hydroxylase (ASPH) mRNA			4.00E-44
273.seq	314	gi 189596 gb M55409 HUMPANCAN	Homo sapiens pancreatic tumor-related protein mRNA, partial cds			1.00E-93
275.seq	414					
276.seq	445	gi 4758949 ref NM_000942.1 PPIB	Homo sapiens peptidylprolyl isomerase B (cyclophilin B) (PPIB) mRNA			0
277.seq	509	gi 4507446 ref NM_003223.1 TFAP4	Homo sapiens transcription factor AP-4 (activating enhancer-binding			1E-177
278.seq	521	gi 181966 gb M29548 HUMEF1AB	Human elongation factor 1-alpha (EF1A) mRNA, partial cds.			0
279.seq	469	gi 180614 gb M55593 HUMCLG4Q13	Human collagenase type IV (CLG4) gene, exon 13.			0
280.seq	290					
281.seq	547	gi 5738434 emb AL079333.7 HSDJ82F3	Human DNA sequence from clone 82F3 on chromosome Xq21.33-22.3.			0
282.seq	493					
285.seq	515	gi 2121233 gb U94585 HSU94585	Homo sapiens requiem homolog (hsReq) mRNA, complete cds			0
286.seq	508					
287.seq	546	gi 285992 dbj D13634 HUMR3C14	Human mRNA for KIAA0009 gene, complete cds			1E-159
288.seq	344	gi 572991 ref NM_006691.1 LYVE-1	Homo sapiens lymphatic vessel endothelial hyaluronan receptor 1			1E-176
289.seq	405	gi 580491 gb AL080250.11 HJ234P1	Human DNA sequence from clone 234P15 on chromosome 6q13-14.3, comple			4.00E-53
290.seq	446	gi 4506706 ref NM_001028.1 RPS25	Homo sapiens ribosomal protein S25 (RPS25) mRNA			0
291.seq	486					
292.seq	368	gi 2462497 emb Y11843 HSY11843	H.sapiens mRNA for MHC class I human leukocyte antigen, allele			1E-141

294.seq	398	gi 4504964 ref NM_002298.1 LCP1	Homo sapiens lymphocyte cytosolic protein 1 (L-plastin) (LCP1) mRNA	0
295.seq	420	gi 4506666 ref NM_001002.1 RPLP0	Homo sapiens ribosomal protein, large, P0 (RPLP0) mRNA	0
296.seq				
297.seq	264	gi 441374 emb X72453 HSIGLV32	H.sapiens mRNA for rearranged Ig kappa light chain variable region	1E-125
298.seq	362	gi 4506834 ref NM_002989.1 SCYA21	Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member	0
299.seq	365	gi 1911773 gb S83364 S83364	putative Rab5-interacting protein (clone L1-57) [human, HeLa cells,	2.00E-86
300.seq	449	gi 4506680 ref NM_001015.1 RPS11	Homo sapiens ribosomal protein S11 (RPS11) mRNA	0
301.seq	403			
302.seq	429			
303.seq	408	gi 1834892 emb Z85181 HSZ85181	H.sapiens Ig lambda light chain variable region gene (6-090IIA61)	1E-109
304.seq	409			
306.seq	337			
307.seq	325	gi 5453671 ref NM_006418.1 GW112	Homo sapiens differentially expressed in hematopoietic lineages	1E-153
308.seq	267	gi 3859064 emb AJ006031 MMU6031	Mus musculus IHABP gene, promoter	4.00E-25
309.seq	358	gi 32220 emb X64881 HSHLAEUT3	H.sapiens gene for HLA-E heavy chain (3' untranslated region)	8.00E-30
310.seq	398	gi 3152832 gb AF062327 AF062327	Homo sapiens p120 catenin isoform 2A (CTNND1) mRNA, alternatively	1E-153
311.seq	399	gi 388034 gb L05095 HUMRIBPROF	Homo sapiens ribosomal protein L30 mRNA, complete cds	1E-144
312.seq	332	gi 2114292 db D87023 D87023	Human (lambda) DNA for immunoglobulin light chain	1E-117
313.seq	251	gi 4154282 gb AF089744 AF089744	Homo sapiens xenotropic and polytropic murine leukemia virus receptor	3.00E-47
315.seq	351			
316.seq	404	gi 5678818 gb AF146191.1 AF146191	Homo sapiens FRG1 (FRG1) gene, complete cds; 5S ribosomal RNA gene,	2.00E-12
317.seq	402	gi 4506722 ref NM_001006.1 RPS3A	Homo sapiens ribosomal protein S3A (RPS3A) mRNA	0
320.seq	242	gi 2558844 gb AF017732 AF017732	Homo sapiens chromosome 2 cosmid cos111 and cos 607/6 containing	1E-120
321.seq	481	gi 4504340 ref NM_003642.1 HAT1	Homo sapiens histone acetyltransferase 1 (HAT1) mRNA	1E-157
322.seq	327	gi 4502012 ref NM_001625.1 AK2	Homo sapiens adenylyate kinase 2 (AK2), nuclear gene encoding	1E-153
323.seq	331			
324.seq	385	gi 185267 gb M34024 HUMIGHCXE	Human fetal Ig heavy chain variable region (clone M43) mRNA,	1E-142
325.seq	434			
327.seq	875	gi 4884392 emb ALD50179.1 HSM800475	Homo sapiens mRNA: cDNA DKFZp586K2222 (from clone DKFZp586K2222)	1E-130
328.seq	319	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	3.00E-75
330.seq	388	gi 23221 emb X56468 HS1433	Human mRNA for 14.3.3 protein, a protein kinase regulator	0
331.seq	407	gi 291855 gb L09209 HUMAMYLOID	Homo sapiens amyloid protein homologue mRNA, complete cds	3.00E-88
333.seq	422	gi 312806 emb X68641 HSHIP	H.sapiens HIP mRNA	1E-173
334.seq	483	gi 4503140 ref NM_001814.1 CTSC	Homo sapiens cathepsin C (CTSC) mRNA >gi 1006656 emb X87212 HSCATH	1E-123
336.seq	134	gi 5174470 ref NM_005896.1 IDH1	Homo sapiens isocitrate dehydrogenase 1 (NADP+), soluble (IDH1) mRNA	4.00E-57
337.seq	404	gi 3928123 gb AC005972 AC005972	Homo sapiens chromosome 17, clone hRPK.180_P_8, complete sequence [Ho	2.00E-86
338.seq	426	gi 4335942 gb AF061739 AF061739	Homo sapiens unknown mRNA	0
339.seq	395	gi 4501888 ref NM_001615.1 ACTG2	Homo sapiens actin, gamma 2, smooth muscle, enteric (ACTG2) mRNA	0
340.seq	933	gi 178698 gb M82609 HUMANX4A	Human annexin IV (ANX4) mRNA, complete cds.	1.00E-64
341.seq	254	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	1E-113
342.seq	467			
343.seq	350	gi 3646023 emb AL031228 HS1033B10	Human DNA sequence from clone 1033B10 on chromosome 6p21.2-21.31.	1.00E-19
344.seq	456	gi 337454 gb M29063 HUMRNPC2A	Human hnRNP C2 protein mRNA.	0
345.seq	440	gi 266703 gb S55735 S55735	Ig alpha 1- alpha 2m-immunoglobulin A1-A2 lambda hybrid GAU heavy	0
347.seq	383	gi 510270 gb U09500 HSU09500	Human mitochondrion cytochrome b gene, partial cds.	1E-176
348.seq	489	gi 3882314 gb AB018340.1 AB018340	Homo sapiens mRNA for KIAA0797 protein, partial cds.	0
349.seq	253	gi 4895146 gb AC007115.1 AC007115	Homo sapiens chromosome 12 clone 91705, complete sequence	1E-138
350.seq	833			
351.seq	830	gi 3201899 gb AF067420 AF067420	Homo sapiens SNC73 protein (SNC73) mRNA, complete cds	1E-139
352.seq	473			
353.seq	487	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	1E-120
354.seq	446	gi 4506668 ref NM_001003.1 RPLP1	Homo sapiens ribosomal protein, large, P1 (RPLP1) mRNA	0
356.seq	267	gi 1469307 gb U57456 HSU57456	Human transforming growth factor-beta signaling protein-1 (bsp-1)	1E-110
357.seq	509	gi 4506146 ref NM_002770.1 PRSS2	Homo sapiens protease, serine, 2 (trypsin 2) (PRSS2) mRNA	0
358.seq	334	gi 5835387 ref NC_001807.1	Human mitochondrion, complete genome >gi 1944628 gb J01415 HUMMTCG	1E-150
359.seq	398			
360.seq	519	gi 4507668 ref NM_003295.1 TPT1	Homo sapiens tumor protein, translationally-controlled 1 (TPT1)	0
361.seq	540	gi 2280483 gb AB002370 AB002370	Human mRNA for KIAA0372 gene, complete cds	0
362.seq	532			
363.seq	488	gi 2576344 gb AC002400 HUAC002400	Human Chromosome 16 BAC clone CIT987SK-A-735G6, complete sequence	0
366.seq	464			
368.seq	454	gi 452059 gb L27560 HUMIGFBP5X	Human insulin-like growth factor binding protein 5 (IGFBP5) mRNA.	0
370.seq	392	gi 4504748 ref NM_000885.1 ITGA4	Homo sapiens Integrin, alpha 4 (antigen CD49D, alpha 4 subunit of	8.00E-95
371.seq	509	gi 4507558 ref NM_003278.1 TNA	Homo sapiens tetranectin (plasminogen-binding protein) (TNA) mRNA	0
372.seq	448	gi 184847 gb M63438 HUMIGGK	Human Ig rearranged gamma chain mRNA, V-J-C region and complete cds.	0
373.seq	599	gi 5835387 ref NC_001807.1	Human mitochondrion, complete genome >gi 1944628 gb J01415 HUMMTCG	0
374.seq	534	gi 4506242 ref NM_002819.1 PTB	Homo sapiens polypyrimidine tract binding protein (heterogeneous	1E-171
375.seq	536			
376.seq	475	gi 1778152 gb U66839 HSU66839	Human MAP kinase 3b mRNA, complete cds	0
377.seq	813	gi 4826871 ref NM_005124.1 NUP153	Homo sapiens nucleoporin 153kD (NUP153) mRNA	0
378.seq	486	gi 4454701 gb AF070663 AF070663	Homo sapiens HSPC007 mRNA, complete cds	1E-142
379.seq	483	gi 432438 gb U03863 HSU03863	Human MHC Class I HLA-A-0203 mRNA, complete cds.	0
380.seq	507	gi 3789115 gb AC005799 AC005799	Homo sapiens chromosome 17, clone hRPK.82_F_10, complete sequence [Ho	1E-154
381.seq	557	gi 178276 gb M61831 HUMAHCY	Human S-adenosylhomocysteine hydrolase (AHCY) mRNA, complete cds.	0
383.seq				
384.seq	526	gi 4505922 ref NM_002687.1 PNN1	Homo sapiens pinn, desmosome associated protein (PNN) mRNA	1E-109
385.seq	545			
388.seq	487	gi 5453671 ref NM_006418.1 GW112	Homo sapiens differentially expressed in hematopoietic lineages	2.00E-89
390.seq	329	gi 1562509 gb U68105 HSPABP513	Human poly(A)- binding protein (PABP) gene, exon 15	1E-122
391.seq	443	gi 35237 emb X52112.1 HSPAG1L	Human mRNA for immunoglobulin light chain VJ region, lambda-V	1E-130
392.seq	435	gi 4506670 ref NM_001004.1 RPLP2	Homo sapiens ribosomal protein, large P2 (RPLP2) mRNA	0
395.seq	487	gi 4929684 gb AF151866.1 AF151866	Homo sapiens CGI-108 protein mRNA, complete cds	9.00E-15
397.seq	516	gi 5835387 ref NC_001807.1	Human mitochondrion, complete genome >gi 1944628 gb J01415 HUMMTCG	0
399.seq	397			
400.seq	444	gi 347963 gb L22453 HUMTAPR	Homo sapiens HIV-1 TAR RNA binding protein (TARBP-b) mRNA, complete	0
401.seq	199			
402.seq	375	gi 184819 gb M65062 HUMIGFBP6	Human insulin-like growth factor binding protein 5 (IGFBP-5) mRNA,	1E-152
404.seq	324			
406.seq	379	gi 5835387 ref NC_001807.1	Human mitochondrion, complete genome >gi 1944628 gb J01415 HUMMTCG	1E-142
408.seq	369	gi 2920708 emb AJ224143.1 HSAJ4143	Homo sapiens mRNA variant alfa for RNA polymerase II subunit 3	0
409.seq	410	gi 1209254 gb L20311 BOVIGJMRNA	Bos taurus mu- and alpha-immunoglobulin-associated J-chain mRNA,	1.00E-87
410.seq	415	gi 4557576 ref NM_001443.1 FABP1	Homo sapiens fatty acid binding protein 1, liver (FABP1) mRNA	1E-154
411.seq	442	gi 4503304 ref NM_001926.1 DEFA6	Homo sapiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	0
415.seq	418			
418.seq	372	gi 1197535 gb U49020 HSMFE2A10	Human myocyte-specific enhancer factor 2A (MEF2A) gene, last coding	5.00E-53
419.seq	451	gi 184847 gb M63438 HUMIGGK	Human Ig rearranged gamma chain mRNA, V-J-C region and complete	0
420.seq	399			
421.seq	365	gi 285970 db D13510.1 HUMPAPC	Homo sapiens mRNA for PAP homologous protein, complete cds	0
422.seq	413	gi 984142 emb X89401 HSL21PROT	H.sapiens mRNA for large subunit of ribosomal protein L21	0
423.seq	207			
426.seq	342	gi 4557454 ref NM_001277.1 CHK1	Homo sapiens choline kinase (CHK) mRNA >gi 219540 db D10704 HUMCK1	1E-110
427.seq	908	gi 4506658 ref NM_000971.1 RPL7	Homo sapiens ribosomal protein L7 (RPL7) mRNA	1E-173
432.seq	297	gi 3970963 gb AC004999 AC004999	Homo sapiens PAC clone DJ215P15 from 5a12, complete sequence [Homo	1.00E-92
435.seq	435			
438.seq	437	gi 4557800 ref NM_000270.1 NP1	Homo sapiens nucleoside phosphorylase (NP) mRNA	0
439.seq	472			
440.seq	485	gi 23690 emb X56932 HS23KDHBP	H.sapiens mRNA for 23 kD highly basic protein	0
441.seq				

446.seq	528	gi 2224716 dbj AB002386 AB002386	Human mRNA for KIAA0388 gene, complete cds	0
447.seq	449	gi 3882244 dbj AB018305.1 AB018305	Homo sapiens mRNA for KIAA0762 protein, partial cds	0
448.seq	248	gi 4753237 gb AC006372.2 AC006372	Homo sapiens clone NH0331D05, complete sequence	1.00E-15
449.seq	511	gi 2887498 gb AC004143 AC004143	Homo sapiens chromosome 19, DNA from overlapping cosmids R29893 and	1E-127
450.seq	536	gi 4506618 ref NM_000986.1 RPL24	Homo sapiens ribosomal protein L24 (RPL24) mRNA	0
452.seq	482	gi 5453671 ref NM_006418.1 GW112	Homo sapiens differentially expressed in hematopoietic lineages	0
454.seq	502	gi 4758157 ref NM_004044.1 DIFP6	Homo sapiens differentiation 6 (deoxyguanosine triphosphate	0
455.seq	521	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	1E-131
456.seq	565	gi 5802969 ref NM_006796.1 AFG3L2	Homo sapiens AFG3 (ATPase family gene 3, yeast)-like 2 (AFG3L2), mRNA	0
457.seq	515	gi 4885240 ref NM_005252.1 FOS	Homo sapiens v-fos FBJ murine osteosarcoma viral oncogene homolog	0
458.seq	531	gi 2341021 gb AC002477 AC002477	Human PAC clone DJ327A19 from Xq25-q26, complete sequence [Homo sapi	1.00E-94
459.seq	575	gi 2822459 gb AF030234 AF030234	Homo sapiens splicing factor Sfp1 mRNA, complete cds	0
460.seq	290			
461.seq	612	gi 190978 gb M18963 HUMREGA	Human islet of Langerhans regenerating protein (reg) mRNA, complete	0
462.seq	523	gi 5868946 gb AC008101.9 AC008101	Homo sapiens, complete sequence	1E-158
463.seq	509	gi 185361 gb M87789 HUMIGHEPAH	Human (hybridoma H210) anti-hepatitis A IgG variable region,	0
464.seq	433	gi 31912 emb X04409 HSGSA1R	Human mRNA for coupling protein G(s) alpha-subunit (alpha-S1)	0
465.seq	517	gi 4507812 ref NM_003359.1 UGDH	Homo sapiens UDP-glucose dehydrogenase (UGDH) mRNA, and translated	0
466.seq	883	gi 32804 emb X61012 HSIG580	Human immunoglobulin VH mRNA (580 bp)	0
467.seq	572			
470.seq	284	gi 5174628 ref NM_006099.1 PIAS3	Homo sapiens protein inhibitor of activated STAT3 (PIAS3) mRNA	1E-151
471.seq	524	gi 5912164 emb AL117596.1 HSMB01125	Homo sapiens mRNA; cDNA DKFZp564C2163 (from clone DKFZp564C2163)	0
472.seq	504			
473.seq	516	gi 13683 emb V00710 MIT1HS	Human mitochondrial genes for several tRNAs (Phe, Val, Leu) and 12S	0
474.seq	571			
475.seq	504	gi 259140 gb S49975 S49975	glutathione transferase A1-1 [human, HepG2 hepatoma cells, mRNA,	0
477.seq	490			
478.seq	479			
479.seq	571	gi 1262342 emb X93334 HSMITG	H.sapiens mitochondrial DNA, complete genome	0
480.seq	480	gi 2076723 gb AC002064 HSAC002064	Human BAC clone RGO16J04 from 7q21, complete sequence [Homo sapiens]	0
481.seq	491	gi 2599233 gb AC002994 AC002994	Homo sapiens chromosome 17, clone HRPC897K16, complete sequence [Hom	0
482.seq	601	gi 5138925 gb AF092138.1 AF092138	Homo sapiens HSPC033 mRNA, complete cds	0
483.seq	491	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	1E-145
486.seq	524	gi 4503744 ref NM_001456.1 FLNA	Homo sapiens filamin A, alpha (actin-binding protein-280) (FLNA) mRNA	0
487.seq	524	gi 13003 emb V00662 MIHSXX	H.sapiens mitochondrial genome	0
488.seq	476	gi 30204 emb X06409 HSCRAF1R	Human mRNA fragment for activated c-raf-1 (exons 8-17)	0
489.seq	492	gi 3483595 gb AF086250 HUMZD39H03	Homo sapiens full length insert cDNA clone ZD39H03	1E-147
490.seq	553	gi 28251 emb X00351 HSAC07	Human mRNA for beta-actin	0
491.seq	417			
492.seq	473	gi 2245670 gb AF006516 AF006516	Homo sapiens eps8 binding protein e3B1 mRNA, complete cds	0
493.seq	405	gi 3228200 emb AL023578 HS248E1	Human DNA sequence from clone 248E1 on chromosome 6q23.1-23.3 Contain	0
494.seq	562	gi 4759095 ref NM_004261.1 SEP15	Homo sapiens 15 kDa selenoprotein (SEP15) mRNA	0
497.seq	555			
498.seq	321	gi 2558844 gb AF017732 AF017732	Homo sapiens chromosome 2 cosmids cos111 and cos 607/6 containing	1E-133
499.seq	432	gi 598852 gb J17249 HUMDSA09M3	Human HepG2 3' region Mbol cDNA, clone hmd5a09m3	1E-137
500.seq	415	gi 4210498 gb AC006486 AC006486	Homo sapiens chromosome 19, BAC CIT-147B23 (BC85722), complete	2.00E-68
501.seq	561	gi 1042081 gb 578569 S78569	laminin alpha 4 chain [human, fetal lung, mRNA, 6204 nt]	0
502.seq	403	gi 5763746 emb AL049786.1 HSDJ686	Human DNA sequence from clone 686N3 on chromosome 20q13.2-13.2,	0
503.seq	399	gi 5821152 dbj AB016092.1 AB016092	Homo sapiens mRNA for RNA binding protein, complete cds	1E-103
504.seq	553	gi 1508218 emb Z78940 HSPA19G2	H.sapiens flow-sorted chromosome 6 HindIII fragment, SC6A19G2	1E-153
505.seq	524			
506.seq	457	gi 388034 gb L05095 HUMRIBPROF	Homo sapiens ribosomal protein L30 mRNA, complete cds	0
509.seq	417	gi 4506730 ref NM_001010.1 RPS6	Homo sapiens ribosomal protein S6 (RPS6) mRNA	0
510.seq	473	gi 4826429 emb AL035551.6 HS1100E15	Human DNA sequence from clone 1100E15 on chromosome Xq22.1-22.3	3.00E-76
511.seq	447	gi 4505046 ref NM_002345.1 LUM	Homo sapiens lumican (LUM) mRNA > gi 699576 gb U21128 HSU21128 Human	0
512.seq	530			
513.seq	500	gi 5802967 ref NM_006818.1 AF1Q	Homo sapiens transmembrane protein (AF1Q), mRNA	0
514.seq	534	gi 3287156 emb AL031005 HS329E20	Homo sapiens DNA sequence from PAC 329E20 on chromosome 1p34.4-36.1	0
515.seq				
516.seq	355	gi 264772 gb S54005 S54005	thyrosin beta-10 [human, metastatic melanoma cell line, mRNA, 453	0
517.seq	430	gi 4510439 gb AC007065.5 AC007065	Homo sapiens, clone hRPK.58.A.1, complete sequence	0
518.seq	488	gi 1924939 emb X9841 HSMYOSIE	H.sapiens mRNA for myosin-IE	0
519.seq	493			
520.seq	415	gi 493150 gb L33038 HUMIGKVRG	Human immunoglobulin kappa light chain (IGKV) mRNA variable region,	1E-177
521.seq	475	gi 5771401 gb AF096771.1 HSMCLKKRF	Homo sapiens myosin light chain kinase (MYLK) gene, exon 31, and	0
522.seq	540	gi 3688107 gb AC004477 AC004477	Homo sapiens chromosome 17, clone HRPC89E16, complete sequence [Hom	0
523.seq	510			
524.seq	355	gi 4928275 gb AF131913.1 AF131913	Homo sapiens alpha-(1.3/1.4)-fucosyltransferase (FT3B) mRNA, complete	2.00E-30
525.seq	860	gi 348706 gb L22569 HUMCACTH8S	Homo sapiens cathepsin B mRNA, 3' UTR with a stem-loop structure	0
526.seq	541			
527.seq	472	gi 4505184 ref NM_002415.1 MIF	Homo sapiens macrophage migration inhibitory factor	0
529.seq	510			
530.seq	556	gi 5262538 emb AL080111.1 HSM800618	Homo sapiens mRNA; cDNA DKFZp586G2222 (from clone DKFZp586G2222)	0
531.seq	550	gi 38334 emb X62125 HSWLRG	H.sapiens rearranged WLR gene for lambda-immunoglobulin subgroup II	0
532.seq	528	gi 13000 emb X62996 MIHSGENOM	H.sapiens mitochondrial genome (consensus sequence)	0
533.seq	520	gi 39511 emb X65912 HSIGHXX30	H.sapiens mRNA for XLA IG heavy chain VDJ region (LE 5-11)	1.00E-94
534.seq	533			
535.seq				
536.seq	838			
537.seq				
538.seq				
539.seq	191	gi 4506614 ref NM_000984.1 RPL23A	Homo sapiens ribosomal protein L23a (RPL23A) mRNA	1.00E-82
540.seq	504	gi 3201899 gb AF067420 AF067420	Homo sapiens SNC73 protein (SNC73) mRNA, complete cds	0
541.seq	331			
542.seq	546	gi 185539 gb M18513 HUMIGHYAK	Human (fetal) rearranged IgM chain VDJ-region mRNA, clone 30P1.	1E-122
543.seq	519	gi 23690 emb X56932 HS23KDHP	H.sapiens mRNA for 23 kD highly basic protein	0
544.seq	495	gi 339899 gb M23161 HUMTRANSC	Human transposon-like element mRNA.	1E-134
545.seq	582			
546.seq	570	gi 4503470 ref NM_001402.1 EEF1A1	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
548.seq	534	gi 4502294 ref NM_001686.1 ATP5B	Homo sapiens ATP synthase, H+-transporting, mitochondrial F1	0
549.seq	560			
551.seq	186	gi 13000 emb X62996 MIHSGENOM	H.sapiens mitochondrial genome (consensus sequence)	1.00E-94
552.seq	573	gi 5803154 ref NM_006842.1 SAP145	Homo sapiens spliceosome associated protein 145, SF3b subunit	0
553.seq	504	gi 38336 emb X62125 HSWHRG	H.sapiens rearranged WHR gene for immunoglobulin heavy-chain	1E-112
554.seq	497			
557.seq	498	gi 1000861 gb L47647 HUMCKBR	Homo sapiens creatine kinase B mRNA, complete cds.	0
558.seq	495	gi 576474 gb M28203 HUMMHBW62	Homo sapiens (clone pMF18) MHC class I HLA-Bw62 mRNA, 3' end,	0
559.seq	546	gi 4557494 ref NM_001326.1 CSTF3	Homo sapiens cleavage stimulation factor, 3' pre-RNA, subunit 3.	0
560.seq	476	gi 2662156 dbj AB007898 AB007898	Homo sapiens KIAA0438 mRNA, complete cds	1E-136
561.seq	540	gi 1490418 gb U49869 HSLU49869	Human ubiquitin gene, complete cds	0
562.seq	442	gi 5042403 gb AC007785.1 AC007785	Homo sapiens chromosome 19, BAC 282485 (CIT-8-344H19), complete sequ	2.00E-34
563.seq	542	gi 4503452 ref NM_003792.1 EDF1	Homo sapiens endothelial differentiation-related factor 1 (EDF1)	0
565.seq	438			
566.seq	536	gi 4186004 emb AJ007680 HSA007680	Homo sapiens ZNF198 exon 8	1E-117
567.seq	475	gi 404781 gb L20681 RATETSONCA	Rat proto-oncogene (Ets-1) mRNA, complete cds.	3.00E-39
568.seq	829			
569.seq	509			
570.seq	514	gi 4753238 gb AC006039.2 AC006039	Homo sapiens clone NH0319F03, complete sequence	1.00E-26

572.seq	298	g 2673898 gb AC002536 AC002536	Human Chromosome 11 pac pDJ1075f20, complete sequence [Homo sapiens]	1E-164
573.seq	542	g 4506598 ref NM_000977.1 RPL13	Homo sapiens ribosomal protein L13 (RPL13) mRNA	0
574.seq	548	g 4503470 ref NM_001402.1 EEF1A1	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
575.seq	436	g 4049267 gb U17714 HSU17714	Homo sapiens putative tumor suppressor ST13 (ST13) mRNA, complete cds	0
576.seq	352	g 4502950 ref NM_00090.1 COL3A1	Homo sapiens collagen, type III, alpha 1 (Ehlers-Danlos syndrome type	0
577.seq	437	g 2781391 gb U63387 MMU63387	Mus musculus transcriptional repressor Mpc2 mRNA, complete cds	3.00E-39
579.seq	256	g 5031778 ref NM_005531.1 IFI16	Homo sapiens interferon, gamma-inducible protein 16 (IFI16) mRNA	7.00E-85
580.seq	428	g 4503538 ref NM_001418.1 EIF4G2	Homo sapiens eukaryotic translation initiation factor 4 gamma, 2	0
583.seq	322	g 402482 gb L15203 HUMPIB	Human secretory protein (P1.B) mRNA, complete cds.	1E-129
584.seq	228	g 3873504 gb AC004471 AC004471	Homo sapiens Chromosome 22q11.2 Cosmid Clone 111f11 In DGCR Region,	1E-115
585.seq	406	g 338060 gb M33651 HUMSGP3	Human (clone lambda-PG(6,7,8)) secretory granule proteoglycan core	1E-118
586.seq	283			
588.seq	452	g 1562509 gb U68105 HSPABP13	Human poly(A)-binding protein (PABP) gene, exon 15	1E-137
591.seq	319			
593.seq	270	g 180500 gb M77829 HUMCHIP28A	Human channel-like integral membrane protein (CHIP28) mRNA,	1E-127
594.seq	274			
598.seq	244	g 4557394 ref NM_000067.1 CA2	Homo sapiens carbonic anhydrase II (CA2) mRNA	1E-109
599.seq	215	g 5640003 gb AF165522.1 AF165522	Homo sapiens ras-related GTP-binding protein 4b (RAB4B) mRNA,	1E-108
604.seq	357	g 5523811 gb AC006333.3 AC006333	Homo sapiens clone NH0390E23, complete sequence	1E-111
607.seq	255	g 4506706 ref NM_001028.1 RPS25	Homo sapiens ribosomal protein S25 (RPS25) mRNA	1E-101
611.seq	350			
612.seq	326	g 3928291 emb Z68992 HSPOM16	H.sapiens mRNA for immunoglobulin kappa light chain VJ region (iD	1E-111
613.seq	351			
614.seq				
615.seq	231			
617.seq	221			
618.seq	344	g 32797 emb X61011 HSIG486	Human Immunoglobulin VH mRNA (486 bp)	1E-111
619.seq	343	g 187625 gb M84378 HUMMHAD	Human MHC class I lymphocyte antigen (HLA-A*0212) mRNA, complete	0
621.seq	408			
622.seq	337	g 4336348 gb AF082871 AF082871	Homo sapiens arsenate resistance protein ARS2 mRNA, complete cds	1E-168
623.seq	506	g 6137796 ref NC_001807.2	Human mitochondrion, complete genome	0
624.seq	209	g 1619959 gb U72063 HSU72063	Human immunoglobulin kappa chain constant region gene, partial cds	1.00E-98
626.seq	413	g 546798 gb S71043 S71043	Ig alpha 2-immunoglobulin A heavy chain allotype 2 (constant region,	0
627.seq	373	g 4309874 gb AC006501 AC006501	Homo sapiens, clone hRPK.35_A_1, complete sequence [Homo sapiens]	1E-154
628.seq	400	g 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	1E-105
629.seq	432	g 5052332 gb AF126743.1 AF126743	Homo sapiens DNAJ domain-containing protein MCJ (MCJ) mRNA, complete	0
630.seq	338			
631.seq	359			
632.seq	229	g 4589625 db AB023208.1 AB023208	Homo sapiens mRNA for KIAA0991 protein, complete cds	1E-102
634.seq	317	g 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	1E-113
635.seq	457	g 3288442 emb AL022394 HS511B24	Homo sapiens DNA sequence from clone 511B24 on chromosome 20q11.2-12	0
636.seq	480	g 5835387 ref NC_001807.1	Human mitochondrion, complete genome >g 1944628 gb J01415 HUMMTCG	0
637.seq	399	g 3066552 gb L13802 HUMCH13C3A	Homo sapiens (clone O3) liver expressed ribosomal protein S3	0
638.seq	453	g 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	1E-127
639.seq	418	g 4504164 ref NM_000177.1 GSN	Homo sapiens gelsolin (amyloidosis, Finnish type) (GSN) mRNA	0
641.seq	224			
642.seq	435	g 1370102 emb X74107 HSAMAK67	H.sapiens gene for antigen of the monoclonal antibody KI-67	0
643.seq	400			
644.seq	331	g 2460036 gb AF014837 AF014837	Homo sapiens mRNA for antigen of the monoclonal antibody KI-67	0
646.seq	418			
647.seq	381			
648.seq	209	g 184517 gb M13560 HUMIAIG6	Human Ia-associated invariant gamma-chain gene, exon 8, clones	2.00E-75
650.seq	409	g 5454127 ref NM_006388.1 TIP60	Homo sapiens Tat interactive protein (60kD) (TIP60) mRNA	0
651.seq	424			
653.seq	369	g 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	0
654.seq	471	g 5729967 ref NM_006713.1 PC4	Homo sapiens activated RNA polymerase II transcription cofactor 4	0
655.seq	266	g 23238 emb X15256 HS14LECT	Homo sapiens mRNA for 14kDa beta-galactoside-binding lectin	1E-125
657.seq	403	g 395086 emb X74070 HSBTF3	H.sapiens mRNA for transcription factor BTF 3	0
658.seq				
659.seq	321	g 1110519 gb U40490 HSU40490	Human nicotinamide nucleotide transhydrogenase mRNA, nuclear gene	1E-168
660.seq	423			
661.seq	376	g 5912158 emb AL117594.1 HSM801122	Homo sapiens mRNA: cDNA DKFZp564C1563 (from clone DKFZp564C1563)	0
663.seq	464	g 4507584 ref NM_003823.1 TNFRSF6B	Homo sapiens tumor necrosis factor receptor superfamily, member 6b,	0
664.seq	424	g 3126777 gb AC004590 AC004590	Homo sapiens chromosome 17, clone hCIT.22.K.21, complete sequence [Homo	1E-107
665.seq	406	g 1834980 emb Z85269 HSZ85269	H.sapiens Ig lambda light chain variable region gene	3.00E-94
667.seq	455	g 3540146 gb AC002395 AC002395	Homo sapiens 12q24.1 PAC RPC11-315L5 (Roswell Park Cancer Institute Hun	7.00E-31
668.seq	405			
669.seq	403	g 5835387 ref NC_001807.1	Human mitochondrion, complete genome >g 1944628 gb J01415 HUMMTCG	0
670.seq				
671.seq	451	g 187783 gb M24097 HUMMHACA	Human MHC class I HLA-C-alpha-2 chain and alternative mRNA, complete	0
672.seq	436	g 4314324 gb AC006101 AC006101	Homo sapiens chromosome 10 clone CIT-HSP-1338F24 map 10q11.2-10q12,	4.00E-97
673.seq	252			
674.seq	325	g 4501930 ref NM_000668.1 ADH2	Homo sapiens alcohol dehydrogenase 2 (class I), beta polypeptide	1.00E-43
675.seq	373	g 5678818 gb AF146191.1 AF146191	Homo sapiens FRG1 (FRG1) gene, complete cds: S5 ribosomal RNA gene,	1E-120
676.seq	211	g 13683 emb V00710 MIT1HS	Human mitochondrial genes for several tRNAs (Phe, Val, Leu) and 12S	1E-100
678.seq	317	g 4506648 ref NM_000967.1 RPL3	Homo sapiens ribosomal protein L3 (RPL3) mRNA	4.00E-56
679.seq	347	g 3335147 gb AF055376 AF055376	Homo sapiens short form transcription factor C-MAF (c-maf) mRNA,	1E-117
680.seq	384	g 4454485 gb AC006599 AC006599	Homo sapiens PAC clone pDJ593b2, complete sequence [Homo sapiens]	0
681.seq	350	g 338008 gb M28226 HUMSECPA	Human JE gene encoding a monocyte secretory protein mRNA, complete	1E-123
682.seq	314			
683.seq	316	g 1037301 emb Z64479 HS127C4R	H.sapiens CpG Island DNA genomic MseI fragment, clone 127c4,	3.00E-72
685.seq	423	g 4426566 gb AF034607 AF034607	Homo sapiens chloride channel ABP mRNA, complete cds	1E-164
688.seq	437	g 2707904 gb AF038404 AF038404	Homo sapiens homolog of Nedd5 (hNedd5) mRNA, complete cds	0
690.seq	372			
693.seq	402	g 35324 emb X01742 HSPCAIII	Human mRNA for pro alpha 1 (III) collagen C-terminal propeptide	0
696.seq	388	g 339688 gb M17733 HUMTHYB4	Human thymosin beta-4 mRNA, complete cds.	1E-171
697.seq	343	g 4503056 ref NM_001885.1 CRYAB	Homo sapiens crystallin, alpha B (CRYAB) mRNA	1E-135
698.seq	282	g 4502208 ref NM_001662.1 ARF5	Homo sapiens ADP-ribosylation factor 5 (ARF5) mRNA	1E-113
699.seq	363	g 4507294 ref NM_003569.1 STX7	Homo sapiens syntaxin 7 (STX7) mRNA >g 2337919 gb U77942 HSU77942	1E-135
700.seq	362	g 202446 gb M21829 MUSZFP	Mouse Evi-1 locus zinc finger protein mRNA, partial cds.	7.00E-18
701.seq	300	g 599787 emb Z46810 HSHLACW7	H.sapiens HLA-Cw*0701 mRNA for HLA-Cw7	1E-129
703.seq	375	g 2662152 db AB007896 AB007896	Homo sapiens KIAA0436 mRNA, partial cds.	2.00E-58
705.seq	480	g 2245670 gb AF006516 AF006516	Homo sapiens eps8 binding protein e3B1 mRNA, complete cds	0
706.seq	384	g 306552 gb L13802 HUMCH13C3A	Homo sapiens (clone O3) liver expressed ribosomal protein S3	1E-170
708.seq	338	g 2880082 gb AC004140 AC004140	Homo sapiens PAC clone DJ0841B21 from 7q21.1-q31.1, complete sequence	1E-136
709.seq	151	g 3954884 emb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	1.00E-63
710.seq	346	g 598572 db D16869 HUMD1E05	Human HepG2 3' region cDNA, clone hmd1e05	2.00E-58
711.seq	366	g 1256836 gb U53209 HSU53209	Human transformer-2 alpha (htra-2 alpha) mRNA, complete cds	1E-127
712.seq	333			
713.seq	387	g 4507216 ref NM_003133.1 SRP9	Homo sapiens signal recognition particle 9kD (SRP9) mRNA	0
714.seq	335			
715.seq	429	g 4768676 gb AF113887.1 AF113887	Homo sapiens kappa 1 immunoglobulin light chain mRNA, complete cds	1E-164
716.seq	495	g 4505278 ref NM_002454.1 MTRR	Homo sapiens 5-methyltetrahydrofolate-homocysteine methyltransferase	0
717.seq	494	g 5262631 emb AL080166.1 HSM800685	Homo sapiens mRNA: cDNA DKFZp564O2082 (from clone DKFZp564O2082)	0
718.seq	389	g 4009459 gb AF039401 AF039401	Homo sapiens calcium-dependent chloride channel-1 (hCLCA1) gene,	1E-174
720.seq	307	g 5668757 gb AC008009.4 AC008009	Homo sapiens 3q26.2-27 BAC RPC111-436A20 (Roswell Park Cancer Institut	3.00E-38
723.seq	566	g 4506598 ref NM_000977.1 RPL13	Homo sapiens ribosomal protein L13 (RPL13) mRNA	0
724.seq	439			

725.seq	485	gi 854178 emb X87373 HSRPS3AGE	H.sapiens RPS3a gene	0
727.seq	310	gi 3676217 emb AL022726 HS625H18	Human DNA sequence from clone 625H18 on chromosome 6p22.2-23. Contains	1E-143
728.seq	404	gi 312806 emb X68641 HSHIP	H.sapiens HIP mRNA	0
731.seq	362			
732.seq	359	gi 407467 emb X75316 MMRNASEB4	M.musculus seb4 mRNA	2.00E-21
733.seq	529			
734.seq	479	gi 4503580 ref NM_001978.1 EPB49	Homo sapiens erythrocyte membrane protein band 4.9 (dematin) (EPB49)	9.00E-83
736.seq	328	gi 1773069 gb U82615 HSU82615	Human mRNA downregulated in adenovirus 5-infected cells	1E-111
739.seq	497	gi 2462374 emb Z94721 HHS167A14	Human DNA sequence from PAC 167A14 on chromosome 6q27. Contains EST	0
740.seq	425	gi 3687213 gb AC005738 AC005738	Homo sapiens chromosome 5, BAC clone 7g12 (LBNL HI26), complete sequer	1E-142
741.seq	461			
742.seq	344			
745.seq	539	gi 995938 gb U34343 HSU34343	Human 13kD differentiation-associated protein mRNA, partial cds.	0
746.seq	399			
748.seq	397	gi 4506800 ref NM_003469.1 JSCG2	Homo sapiens secretogranin II (chromogranin C) (SCG2) mRNA	0
749.seq	558	gi 3327109 gb AB014548 AB014548	Homo sapiens mRNA for KIAA0648 protein, partial cds	0
751.seq	517	gi 4503470 ref NM_001402.1 EEF1A1	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
752.seq	483	gi 893287 gb L38951 HUMIMP90A	Homo sapiens importin beta subunit mRNA, complete cds	0
753.seq	567	gi 1663703 gb D87684 D87684	Human mRNA for KIAA0242 gene, partial cds	0
754.seq	312			
755.seq	470	gi 5803210 ref NM_006759.1 UGP2	Homo sapiens UDP-glucose pyrophosphorylase 2 (UGP2), mRNA	0
756.seq	447	gi 4503744 ref NM_001456.1 FLNA	Homo sapiens filamin A, alpha (actin-binding protein-280) (FLNA) mRNA	0
757.seq	518	gi 555940 gb U14990 HSU14990	Human XP1PO ribosomal protein S3 (rpS3) mRNA, complete cds.	0
758.seq	432	gi 4506688 ref NM_001019.1 RPS15A	Homo sapiens ribosomal protein S15a (RPS15A) mRNA	0
759.seq	261	gi 3043681 gb AB011151 AB011151	Homo sapiens mRNA for KIAA0579 protein, partial cds	2.00E-79
760.seq	442	gi 4506651 ref NM_002923.1 RGS2	Homo sapiens regulator of G-protein signalling 2, 24kD (RGS2) mRNA	1E-169
761.seq	609			
762.seq	579	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	0
763.seq	543			
764.seq	265	gi 4505752 ref NM_002629.1 PGAM1	Homo sapiens phosphoglycerate mutase 1 (brain) (PGAM1) mRNA	1E-132
765.seq	527	gi 30308 emb Y00052 HSCYCR	Human mRNA for T-cell cyclophilin	0
766.seq	552	gi 4758647 ref NM_004521.1 KIF5B	Homo sapiens kinesin family member 5B (KIF5B) mRNA	0
767.seq	431	gi 5579476 ref NM_006233.2 POLR2I	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide I	0
768.seq	489			
769.seq	461	gi 4502950 ref NM_000090.1 COL3A1	Homo sapiens collagen, type III, alpha 1 (Ehlers-Dankos syndrome)	0
770.seq	453			
771.seq	509	gi 349475 gb L16785 HUMPUF	Homo sapiens c-myc transcription factor (puf) mRNA, complete cds.	0
772.seq	472	gi 3483527 gb AF086182 HUMZB97G06	Homo sapiens full length insert cDNA clone ZB97G06	3.00E-33
773.seq	501			
774.seq	537			
775.seq	422	gi 2224646 gb AB002351 AB002351	Human mRNA for KIAA0353 gene, partial cds	0
778.seq	357	gi 1839516 gb S82240 S82240	RhoE-26 kDa GTPase homolog [human, HeLa cell line, mRNA, 833 nt]	1E-104
780.seq	483	gi 2822137 gb AC002418 AC002418	Human Chromosome X, complete sequence [Homo sapiens]	1E-100
781.seq	511	gi 4504444 ref NM_002136.1 HNRPA1	Homo sapiens heterogeneous nuclear ribonucleoprotein A1 (HNRPA1)	1E-114
786.seq	458	gi 36139 emb X52967 HSRPL7	Human mRNA for ribosomal protein L7	0
787.seq	510			
789.seq	489			
791.seq	429	gi 4104866 gb AF040639.1 AF040639	Homo sapiens aflatoxin B1-aldehyde reductase mRNA, complete cds	1E-179
792.seq	353	gi 4758647 ref NM_004521.1 KIF5B	Homo sapiens kinesin family member 5B (KIF5B) mRNA	1E-161
793.seq	435	gi 189019 gb M22918 HUMMYLCB	Human non-muscle myosin alkali light chain mRNA, 3' end.	0
795.seq	407			
796.seq	317	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	1E-144
797.seq	363	gi 4929137 gb AF141349.1 AF141349	Homo sapiens beta-tubulin mRNA, complete cds	2.00E-92
798.seq	467	gi 4884205 emb AL049955.1 HSM800300	Homo sapiens mRNA; cDNA DKFZp564J0123 (from clone DKFZp564J0123)	0
799.seq	488	gi 2058316 emb Z84488 HS93H18	Human DNA sequence from PAC 93H18 on chromosome 6 contains ESTs	0
800.seq	433	gi 3043685 gb AB011153 AB011153	Homo sapiens mRNA for KIAA0581 protein, partial cds	0
801.seq	376	gi 184743 gb J00220 HUMIGCC8	Homo sapiens immunoglobulin alpha-1 heavy chain constant region	0
803.seq	380	gi 5815340 gb AF176006.1 AF176006	Homo sapiens mitochondrial carrier homolog 1 mRNA, complete cds;	0
804.seq	330			
805.seq	400	gi 1665786 gb D87449 D87449	Human mRNA for KIAA0260 gene, partial cds	1E-155
807.seq	534			
810.seq	461	gi 5442419 gb AF159246.1 AF159246	Bos taurus calpastatin mRNA, complete cds	7.00E-28
812.seq	113			
814.seq	109	gi 4503270 ref NM_001920.1 DCN	Homo sapiens decorin (DCN) mRNA >gi 181169 gb M14219 HUMCSPG1A	7.00E-18
815.seq	542	gi 1430779 emb Z75888 HS130N4B	Human DNA sequence from cosmid 130N4, BRCA2 gene region chromosome	0
817.seq	328	gi 1878541 gb M57648 HUMMHCDRZ	Human MHC HLA-Dw12 mRNA, complete cds.	1E-161
818.seq	437	gi 28335 emb X63432 HSACTB	H.sapiens ACTB mRNA for mutant beta-actin (beta'-actin)	0
819.seq	516	gi 33733 emb X57817 HSIGVL027	Human rearranged immunoglobulin lambda light chain mRNA	0
820.seq	563			
823.seq	306	gi 4506678 ref NM_001014.1 RPS10	Homo sapiens ribosomal protein S10 (RPS10) mRNA	1E-133
824.seq	226	gi 4761682 gb AF147331.1 AF147331	Homo sapiens full length insert cDNA clone YB24D08	1E-103
825.seq	361	gi 4503248 ref NM_003472.1 D6S231E	Homo sapiens DEK gene (D6S231E) mRNA >gi 30502 emb X64229 HSDEK9	0
826.seq	480	gi 3881948 emb AL031774 HS298J15	Human DNA sequence from clone 298J15 on chromosome 6p22.3-23 Contains	0
827.seq	547	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	1E-149
828.seq	547	gi 1262342 emb X93334 HSMITG	H.sapiens mitochondrial DNA, complete genome	0
829.seq	283			
830.seq	424	gi 4063630 gb AF090095 AF090095	Homo sapiens clone IMAGE 286356	5.00E-47
831.seq	501	gi 184847 gb M63438 HUMIGGK	Human Ig rearranged gamma chain mRNA, V-J-C region and complete cds.	0
832.seq	505	gi 38434 emb X70326 HSMMAR	H.sapiens MacMarcks mRNA	1E-171
833.seq	448			
838.seq	230	gi 4502268 ref NM_000701.1 ATP1A1	Homo sapiens ATPase, Na+/K+ transporting, alpha 1 polypeptide	9.00E-72
839.seq	462			
840.seq	429	gi 4502268 ref NM_000701.1 ATP1A1	Homo sapiens ATPase, Na+/K+ transporting, alpha 1 polypeptide	1E-145
841.seq	500	gi 181485 gb M24070 HUMDBPB	Human DNA-binding protein B (dbpB) gene, 3' end.	1E-171
842.seq	471	gi 5453671 ref NM_006418.1 GW112	Homo sapiens differentially expressed in hematopoietic lineages	2.00E-55
843.seq	483	gi 4314324 gb AC006101 AC006101	Homo sapiens chromosome 10 clone CIT-HSP-1338F24 map 10p11.2-10p12.	1E-124
844.seq	460	gi 3043729 gb AB011175 AB011175	Homo sapiens mRNA for KIAA0603 protein, complete cds	0
845.seq	440	gi 4506658 ref NM_000971.1 RPL7	Homo sapiens ribosomal protein L7 (RPL7) mRNA	0
846.seq	446	gi 1072387 emb X92665 MMUBCM3	M.musculus mRNA for ubiquitin-conjugating enzyme Ubcm3	6.00E-28
847.seq	491	gi 13000 emb X62996 MHSGENOM	H.sapiens mitochondrial genome (consensus sequence)	0
848.seq	472			
850.seq	414	gi 337764 gb M60257 HUMSAPC1	Human mutant cerebroside sulfate activator protein (SAP-MU-0) mRNA.	0
851.seq	446	gi 4894853 gb AF135157.1 AF135157	Homo sapiens complement C1q A chain precursor, mRNA, complete cds	0
852.seq	383	gi 181615 gb M31767 HUMDNAMET	Human O6-methylguanine-DNA methyltransferase mRNA, complete cds.	1E-133
853.seq	562	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	0
854.seq	442			
855.seq	442	gi 220021 gb D00759 HUMPCSC2	Human mRNA for proteasome subunit HC2 >gi 2171628 gb E03412 E03412	0
856.seq	497	gi 2558667 gb U78525 HSU78525	Homo sapiens eukaryotic translation initiation factor (eIF3) mRNA,	0
857.seq	477			
858.seq	476	gi 4092080 gb AF098642 AF098642	Homo sapiens phospholipid scramblase mRNA, complete cds	1E-131
859.seq	413			
860.seq	479	gi 4929600 gb AF151824.1 AF151824	Homo sapiens CGI-66 protein mRNA, complete cds	0
861.seq	431	gi 184743 gb J00220 HUMIGCC8	Homo sapiens immunoglobulin alpha-1 heavy chain constant region	0
862.seq	492			
863.seq	267	gi 5817037 emb AL110142.1 HSM800786	Homo sapiens mRNA; cDNA DKFZp564D0264 (from clone DKFZp564D0264)	1.00E-80
864.seq	499			
865.seq	353	gi 562073 gb U12465 HSU12465	Human ribosomal protein L35 mRNA, complete cds.	1E-130
866.seq	480	gi 5531804 gb AF078845.1 AF078845	Homo sapiens 16.7kD protein mRNA, complete cds	0

867.seq	521				
868.seq	519				
871.seq	444	gi 5815499 gb AC007688.15 AC007688	Homo sapiens 12p12-27.2-31.7 BAC RPC11-392P7 (Roswell Park Cancer		0
873.seq	393	gi 1256836 gb U53209 HSU53209	Human transformer-2 alpha (htra-2 alpha) mRNA, complete cds	1.00E-84	
874.seq	532				
876.seq	441	gi 4503470 ref NM_001402.1 E1F1A1	Homo sapiens eukaryotic translation elongation factor 1 alpha 1		0
877.seq	551				
878.seq	508	gi 4757825 ref NM_004048.1 B2M	Homo sapiens beta-2-microglobulin (B2M) mRNA		0
879.seq	508	gi 4758041 ref NM_004718.1 COX7RP	Homo sapiens cytochrome c oxidase subunit VII-related protein		0
880.seq	393	gi 4758543 ref NM_004500.1 HNRPC	Homo sapiens heterogeneous nuclear ribonucleoprotein C (C1/C2)		0
881.seq	300				
883.seq	434	gi 2894828 gb U91942 U91942	Homo sapiens immunoglobulin light chain mRNA, partial cds	1E-163	
884.seq	403	gi 2662156 dbj AB007898 AB007898	Homo sapiens KIAA0438 mRNA, complete cds		0
885.seq	446				
886.seq	340	gi 1079782 gb U40975 HSGHMLH16	Human DNA mismatch repair protein homolog (hMLH1) gene, exon 16.	7.00E-58	
887.seq	413	gi 5803174 ref NM_006780.1 SMA3	Homo sapiens SMA3 (SMA3), mRNA >gi 603027 emb X83299 HSSMA3 H.sapi	1.00E-10	
891.seq	447	gi 4240246 dbj AB020686.1 AB020686	Homo sapiens mRNA for KIAA0879 protein, complete cds		0
892.seq	424	gi 23878 emb Z11695 HS40KDA	H.sapiens 40 kDa protein kinase related to rat ERK2		0
894.seq	435	gi 339755 gb MG3121 HUMTNFR	Human tumor necrosis factor receptor (TNF receptor) mRNA, complete		0
895.seq	400	gi 4502920 ref NM_001299.1 CNN1	Homo sapiens calponin 1, basic, smooth muscle (CNN1) mRNA		0
896.seq	428	gi 292434 gb L07287 HUMRPL26X	Human ribosomal protein L26 (RPL26) gene exon 2, complete cds.		0
899.seq	466	gi 4557386 ref NM_00587.1 C7	Homo sapiens complement component 7 (C7) mRNA	1E-120	
900.seq	429	gi 4464283 gb AC007136 AC007136	Homo sapiens chromosome 19, cosmid F20063, complete sequence [Homo	1E-146	
901.seq	458				
903.seq	209	gi 577294 dbj D42041 HUMKIAAE	Human mRNA for KIAA088 gene, partial cds	3.00E-40	
906.seq	388	gi 4501888 ref NM_001615.1 ACTG2	Homo sapiens actin, gamma 2, smooth muscle, enteric (ACTG2) mRNA		0
907.seq	396				
908.seq	497				
909.seq	225				
910.seq	456	gi 5453671 ref NM_006418.1 GW112	Homo sapiens differentially expressed in hematopoietic lineages		0
911.seq	408	gi 5457169 emb AL031320.6 HS20N2	Human DNA sequence from clone 20N2 on chromosome 6q24. Contains the	1E-115	
913.seq	266	gi 4506622 ref NM_00988.1 RPL27	Homo sapiens ribosomal protein L27 (RPL27) mRNA	9.00E-91	
914.seq	489	gi 4506620 ref NM_00987.1 RPL26	Homo sapiens ribosomal protein L26 (RPL26) mRNA		0
916.seq	537	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	1E-145	
917.seq	498				
918.seq	179	gi 903978 gb U29589 HSU29589	Human m3 muscarinic acetylcholine receptor (CHRM3) gene, complete	4.00E-67	
919.seq	441	gi 38405 emb X67292 HSGIMHC61	H.sapiens mRNA for IgM heavy chain constant region (mAb61)		0
920.seq	223	gi 396704 emb Z25521 HSINASPA	H.sapiens Integrin associated protein mRNA, complete CDS,	1E-118	
922.seq	357	gi 4559317 gb AC007204.1 AC007204	Homo sapiens chromosome 19, BAC 273239 cCIT-B-320G13), complete	1E-161	
924.seq	424	gi 4159882 gb AC005630 AC005630	Homo sapiens clone DU1129D05, complete sequence [Homo sapiens]		0
925.seq	390	gi 5031692 ref NM_005758.1 FBRNP	Homo sapiens heterogeneous nuclear protein similar to rat helix		0
926.seq	416	gi 181962 gb J04617 HUMEF1A	Human elongation factor EF-1-alpha gene, complete cds.	1E-151	
927.seq	489	gi 2133864 gb AC00403 AC00403	Genomic sequence from Human 13, complete sequence [Homo sapiens]		0
928.seq	530	gi 2224704 dbj AB002380 AB002380	Human mRNA for KIAA0382 gene, partial cds	1E-165	
930.seq	425	gi 3413936 dbj AB007962 AB007962	Homo sapiens mRNA, chromosome 1 specific transcript KIAA0493		0
931.seq	406	gi 1262342 emb X93334 HSMITG	H.sapiens mitochondrial DNA, complete genome		0
932.seq	495	gi 165703 gb M7233 RABSMMLCKR	Rabbit smooth muscle myosin light chain kinase mRNA, complete CDS.	3.00E-21	
933.seq	413	gi 2583174 gb AF026844.1 AF026844	Homo sapiens ribosomal protein L41 mRNA, complete cds		0
934.seq	405	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	1E-104	
935.seq	434	gi 183657 gb M15872 HUMGST2	Human glutathione S-transferase 2 (GST) mRNA, complete cds.		0
936.seq	455	gi 4507218 ref NM_003137.1 SRPK1	Homo sapiens SFRS protein kinase 1 (SRPK1) mRNA		0
938.seq	505				
940.seq	537				
941.seq	502	gi 3413792 emb AJ010118 HSA010118	Homo sapiens mRNA for mutant NADH-cytochrome b5 reductase, partial.		0
944.seq	494				
945.seq	433	gi 4128014 emb AJ011497.1 HSA011497	Homo sapiens mRNA for Claudin-7		0
946.seq	183				
947.seq	485	gi 5801654 gb AC008115.3 AC008115	Homo sapiens 12p12-27.2-31.7 BAC RPC11-180M15 (Roswell Park Cancer		0
948.seq	519	gi 1101779 gb U40260 RNU40260	Rattus norvegicus tissue carboxypeptidase inhibitor (TCI) mRNA,	8.00E-74	
949.seq	494	gi 3357906 dbj AB016496 AB016496	Mus musculus mRNA for intelectin, complete cds	1E-105	
951.seq	477				
952.seq	471	gi 463124 gb L25275 HUMPSULTRA	Human estrogen sulfotransferase mRNA, complete cds.		0
953.seq	203	gi 3882210 dbj AB018288.1 AB018288	Homo sapiens mRNA for KIAA0745 protein, partial cds	3.00E-71	
956.seq	486	gi 1899202 gb M11968 HUMPGKA1	Human X-linked phosphoglycerate kinase gene, exon 11.	1E-173	
957.seq	464	gi 2343108 gb AF004230 AF004230	Homo sapiens monocyte/macrophage Ig-related receptor MIR-7 (MIR cl-7)	1E-167	
959.seq	531	gi 2665741 gb AF035537 AF035537	Homo sapiens DNA polymerase zeta (REV3) mRNA, complete cds		0
960.seq	422	gi 184743 gb J00220 HUMIGCC8	Homo sapiens immunoglobulin alpha-1 heavy chain constant region		0
961.seq	435	gi 505540 emb X56740 HSMRAB11	H.sapiens mRNA for Rab11 gene		0
962.seq	506	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	1E-133	
963.seq	481	gi 5689428 dbj AB028969.1 AB028969	Homo sapiens mRNA for KIAA1046 protein, complete cds		0
964.seq	471	gi 3043595 dbj AB011108 AB011108	Homo sapiens mRNA for KIAA0536 protein, partial cds		0
965.seq	475	gi 5453671 ref NM_006418.1 GW112	Homo sapiens differentially expressed in hematopoietic lineages		0
966.seq	457	gi 4503308 ref NM_001928.1 DF	Homo sapiens adiponin/complement factor D precursor (DF) mRNA		0
967.seq	275	gi 4503520 ref NM_001568.1 E1F3S6	Homo sapiens eukaryotic translation initiation factor 3, subunit 6	1E-131	
969.seq	471	gi 5453671 ref NM_006418.1 GW112	Homo sapiens differentially expressed in hematopoietic lineages		0
970.seq	164				
971.seq	372	gi 4505604 ref NM_002580.1 PAP	Homo sapiens pancreatitis-associated protein (PAP) mRNA		0
972.seq	460	gi 2950165 emb Z84477 HS265J14	Human DNA sequence from PAC 265J14 on chromosome 6 contains	1E-100	
973.seq	402	gi 2224646 dbj AB002351 AB002351	Human mRNA for KIAA0353 gene, partial cds		0
974.seq	475	gi 4884218 emb AL049969.1 HSM800316	Homo sapiens mRNA; cDNA DKFpZ564A072 (from clone DKFpZ564A072)	1E-115	
975.seq	405	gi 2583174 gb AF026844.1 AF026844	Homo sapiens ribosomal protein L41 mRNA, complete cds		0
976.seq	468	gi 3954884 emb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,		0
977.seq	430				
978.seq	490	gi 3201899 gb AF067420 AF067420	Homo sapiens SNCF3 protein (SNCF3) mRNA, complete cds	1E-128	
979.seq	497	gi 2582056 gb AF014882.1 AF014882	Homo sapiens NADH dehydrogenase subunit 2 (ND2) gene, mitochondrial		0
980.seq	478				
981.seq	494				
982.seq	459				
983.seq	458	gi 186042 gb M74019 HUMIGKW	Human Ig rearranged light-chain mRNA V region.	1E-149	
984.seq	439	gi 184483 gb M55618 HUMHXB	Homo sapiens hexabrachion (HXB) mRNA, complete cds.		0
986.seq	138	gi 5531251 emb AL035659.2 HS979N1	Human DNA sequence from clone 979N1 on chromosome 22q13.1-13.32,	3.00E-70	
987.seq	443	gi 2341008 gb U96629 HSU96629	Human chromosome 8 BAC clone C1T987SK-2A8 complete sequence [Homo	8.00E-80	
988.seq	458	gi 4902640 emb Z98946.15 HS376D21	Human DNA sequence from clone 376D21 on chromosome Xq11.1-12 Contain		0
989.seq	502	gi 458663 gb U04245 HSU04245	Human MHC class I antigen HLA-B (HLA-B-0704 allele) mRNA, complete		0
990.seq	433	gi 1620408 emb Z80843 HSZ80843	H.sapiens rearranged immunoglobulin heavy chain variable region	1E-159	
991.seq	475	gi 4335942 gb AF061739 AF061739	Homo sapiens unknown mRNA	1E-122	
992.seq	474				
993.seq	505	gi 5835387 ref NC_001807.1	Human mitochondrion, complete genome >gi 1944628 gb J01415 HUMMTGC		0
994.seq	347	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	1E-100	
995.seq	508	gi 2224636 dbj AB002346 AB002346	Human mRNA for KIAA0348 gene, complete cds	1E-148	
996.seq	459				
997.seq	480	gi 2291005 gb AF006011.1 AF006011	Homo sapiens dsisveiled 1 (DVL1) mRNA, complete cds		0
998.seq	496	gi 2326514 emb Z85999 HS94G16	Human DNA sequence from PAC 94G16 on chromosome 6q21. Contains		0
999.seq	514	gi 5835387 ref NC_001807.1	Human mitochondrion, complete genome >gi 1944628 gb J01415 HUMMTGC		0
1000.seq	494	gi 4506712 ref NM_002954.1 RPS27A	Homo sapiens ribosomal protein S27a (RPS27A) mRNA		0
1001.seq	460	gi 395086 emb X74070 H8BTF3	H.sapiens mRNA for transcription factor BTF 3		0
1002.seq	288	gi 5764721 gb AC007731.1 AC007731	Homo sapiens, complete sequence	3.00E-78	
1003.seq	487	gi 5453683 ref NM_006344.1 HML2	Homo sapiens macrophage lectin 2 (calcium dependent) (HML2) mRNA		0

1004.seq	442	gi 190978 gb U18963 HUMREGA	Human islet of Langerhans regenerating protein (reg) mRNA, complete	0
1005.seq	470	gi 452059 gb U27560 HUMIGFBP5X	Human insulin-like growth factor binding protein 5 (IGFBP5) mRNA.	0
1006.seq	518	gi 1669383 gb AC000057 HSAC000057	Human BAC clone RG067M09 from Tq21-Tq22, complete sequence [Homo]	0.54
1007.seq	507	gi 3367616 emb AL022722 HS354N19	Human DNA sequence from clone 354N19 on chromosome 6q22. Contains the	0.034
1008.seq	492	gi 2766638 gb AF040901 AF040901	Ovis aries immunoglobulin V lambda chain (V lambda 1.3) gene.	0.002
1011.seq	485	gi 903741 gb L3861 HUMHLA B	Human MHC class I lymphocyte antigen (HLA-B*3801) mRNA, complete	0
1013.seq	416	gi 532596 gb U12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-130
1014.seq	469	gi 4071041 emb AL031775 HS30M3	Human DNA sequence from clone 30M3 on chromosome 6p22.1-22.3. Contain	0
1015.seq	496	gi 4506662 ref NM_000973.1 RPL8	Homo sapiens ribosomal protein L8 (RPL8) mRNA	0
1016.seq	280	gi 4914577 emb AL050373.1 HSM800156	Homo sapiens mRNA; cDNA DKFZp586F1318 (from clone DKFZp586F1318)	E-150
1017.seq	502	gi 4768674 gb AF114264.1 AF114264	Homo sapiens clone HI409 unknown mRNA	E-137
1018.seq				
1020.seq	478	gi 3980553 gb AC004849 AC004849	Homo sapiens PAC clone DJ0659J06 from 7q33-q35, complete sequence [Homo]	0.032
1021.seq	508	gi 23690 emb X56932 HS23KDHP	H.sapiens mRNA for 23 kD highly basic protein	0
1023.seq	460	gi 5835387 ref NC_001807.1	Human mitochondrion, complete genome >gi 1944628 gb J01415 HUMMTCG	0
1024.seq	454	gi 2745966 gb U83999.1 HSTS BIN1	Homo sapiens bridging integrator protein-1 (BIN1) promoter region and	0.12
1025.seq	463	gi 4503273 ref NM_001923.1 DDB1	Homo sapiens damage-specific DNA binding protein 1 (127kD) (DDB1)	0
1026.seq	496	gi 34504 emb Y00651 HSMCP	Human mRNA for membrane cofactor protein >gi 2081922 gb J41332 J41332	0
1027.seq	357	gi 4753238 gb AC006463.3 AC006463	Homo sapiens clone NH0395G17, complete sequence	4.00E-44
1028.seq	475	gi 5803158 ref NM_006746.1 SCML1	Homo sapiens sex comb on midleg (Drosophila)-like 1 (SCML1), mRNA	0
1029.seq	461	gi 339505 gb J02846 HUMTFPB	Human tissue factor gene, complete cds.	0
1030.seq	485	gi 5805075 emb AJ249201.1 PAL249201	Prevotella albensis ftsQ (partial), ftsA and ftsZ genes and ORF -fts	0.008
1031.seq	454	gi 5103672 db JAB010427.2 JAB010427	Homo sapiens mRNA for NORI-1, complete cds	0
1033.seq	491	gi 23221 emb X56468 HS1433	Human mRNA for 14.3.3 protein, a protein kinase regulator	0
1034.seq	494	gi 4758773 ref NM_004548.1 INDUFB10	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 10	0
1036.seq	460	gi 4507668 ref NM_003295.1 TPT1	Homo sapiens tumor protein, translationally-controlled 1 (TPT1)	0
1037.seq	473	gi 532603 gb U35920 HUMIGVL4E	Human immunoglobulin lambda light chain variable region (IGLV)	E-158
1038.seq	491	gi 2911718 gb AC004227 AC004227	Homo sapiens chromosome 5, P1 clone 356a8 (LBNL.H32), complete	0
1039.seq	485	gi 5002644 emb AJ006776.1 HSA6776	Homo sapiens mRNA for translation initiation factor 2 (IF2 gene)	3.00E-05
1040.seq	503	gi 184847 gb U63438 HUMIGGK	Human Ig rearranged gamma chain mRNA, V-J-C region and complete cds.	0
1041.seq	490	gi 2887308 emb Z99943 HS313L4	Human DNA sequence from PAC 313L4 on chromosome 12q24. Contains ESTs	0
1044.seq	476	gi 3550455 emb AJ005073.1 MMAJ5073	Mus musculus mRNA for Alx (ALX-2-interacting protein X), complete	E-134
1045.seq	412	gi 3264850 gb AE001274.1 AE001274	Leishmania major chromosome 1, complete sequence	1.00E-04
1046.seq	452	gi 2780183 emb Z99496 HS509L4	Human DNA sequence from PAC 509L4 on chromosome 6q22.1-6q22.33. Cor	E-149
1047.seq	471	gi 2661084 gb AF035320 AF035320	Homo sapiens clone 23731 peptidylglycine alpha-amidating	0
1049.seq	446	gi 36541 emb X02317 HSSODR1	Human mRNA for Cu/Zn superoxide dismutase (SOD)	0
1050.seq	490	gi 4506602 ref NM_002948.1 RPL15	Homo sapiens ribosomal protein L15 (RPL15) mRNA	0
1051.seq	483	gi 3236226 gb AC004550 AC004550	Homo sapiens 12q13.1 Cosmid C174F5 (Lawrence Livermore LL12NC01 or	0.5
1052.seq	394	gi 339688 gb U17733 HUMTHYB4	Human thymosin beta-4 mRNA, complete cds.	0
1054.seq	491	gi 5031990 ref NM_005806.1 PRSC1	Homo sapiens protease, cysteine, 1 (legumain) (PRSC1) mRNA	0
1056.seq	103	gi 2791860 gb AF041428 AF041428	Homo sapiens ribosomal protein s4 X isoform gene, complete cds	1.00E-50
1057.seq	339	gi 4582128 emb Z97630.1 HSA466N1	Human DNA sequence from clone 466N1 on chromosome 22q12-13 Contains	1.00E-06
1058.seq	470	gi 4503842 ref NM_003917.1 G2AD	Homo sapiens gamma2-adaptin (G2AD) mRNA	0
1060.seq	458	gi 1438780 gb S81752 S81752	DPH2L=candidate tumor suppressor gene [ovarian cancer critical region	e-152
1061.seq	416	gi 5705994 gb AC006946.1B AC006946	Homo sapiens Chromosome 22q11 P.ac Clone p1093 in CES region, complete	0.002
1062.seq	473	gi 4758203 ref NM_004417.1 DUSP1	Homo sapiens dual specificity phosphatase 1 (DUSP1) mRNA	0
1063.seq	410	gi 5453991 ref NM_006264.1 PTPN13	Homo sapiens protein tyrosine phosphatase, non-receptor type 13	0
1064.seq	463	gi 5732140 gb AC006089.2 AC006089	Homo sapiens clone RG315H11, complete sequence	e-130
1065.seq	224	gi 188712 gb U03910 HUMMT2A	Human (clone 14VS) metallothionein-Ig (MT1G) gene, complete cds.	5.00E-64
1066.seq	327	gi 401844 gb U05093 HUMRIBPROD	Homo sapiens ribosomal protein L18a mRNA, complete cds	e-175
1067.seq	491	gi 185397 gb U03162 HUMIGH4A	Human Ig rearranged gamma-chain mRNA V-region, partial cds.	e-165
1068.seq	286	gi 4507762 ref NM_003334.1 UBE1	Homo sapiens ubiquitin-activating enzyme E1 (A1S9T and BN75	e-160
1069.seq	241	gi 4533671 ref NM_006418.1 GW112	Homo sapiens differentially expressed in hematopoietic lineages	e-116
1070.seq	491	gi 261239 gb S50732 S50732	immunoglobulin M light chain V region:anti-lipid A antibody [human,	e-172
1071.seq	458	gi 5836198 gb AC007382.3 AC007382	Homo sapiens clone NH0288C18, complete sequence	0.12
1072.seq	468	gi 5924006 emb AL024498.12 HS417M14	Human DNA sequence from clone 417M14 on chromosome 6p23-25.1, complete	e-166
1073.seq	461	gi 577314 db U0420841 HUMKIAAO	Human mRNA for KIAA0094 gene, partial cds.	0
1074.seq	489	gi 177828 gb U01396 HUMA1ATM	Human alpha-1-antitrypsin mRNA, complete cds.	0
1075.seq	509	gi 541612 gb U14650 HUSU14650	Human platelet-endothelial tetraspan antigen 3 mRNA, complete cds.	0
1076.seq	495	gi 337186 gb U25171 HUMMTCDK	Human mitochondrion cytochrome oxidase subunit II (COII), Asp-tRNA	0
1077.seq	473	gi 4758791 ref NM_004553.1 INDUF5B	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 6 (13kD)	0
1078.seq	497	gi 13683 emb V00710 MIT1HS	Human mitochondrial genes for several tRNAs (Phe, Val, Leu) and 12S	0
1079.seq	481	gi 2621443 gb AE000824 AE000824	Methanobacterium thermoautotrophicum from bases 328735 to 340997	2
1080.seq	496	gi 881458 gb U27682 PLU27682	Paeonia lactiflora Internal transcribed spacer 1 (ITS1), 5.8S	0.52
1081.seq	508	gi 5835387 ref NC_001807.1	Human mitochondrion, complete genome >gi 1944628 gb J01415 HUMMTCG	0
1082.seq	461	gi 4506602 ref NM_002948.1 RPL15	Homo sapiens ribosomal protein L15 (RPL15) mRNA	0
1083.seq				
1084.seq	512	gi 3319121 gb AC003042 AC003042	Homo sapiens chromosome 17, clone HCIT75G16, complete sequence [Homo]	e-101
1085.seq	484	gi 1636191 gb U18396 BOVPROA	Bovine protamine gene P1 allele BP 1.3, complete cds.	0.5
1086.seq	504	gi 5931802 emb AL049765.16 HS1022J1	Human DNA sequence *** SEQUENCING IN PROGRESS *** from clone dj10	5.00E-04
1087.seq	487	gi 298485 gb S56985 S56985	ribosomal protein L19 [human, breast cancer cell line, MCF-7, mRNA,	e-163
1090.seq	495	gi 5689428 db JAB028969.1 JAB028969	Homo sapiens mRNA for KIAA1046 protein, complete cds	0
1091.seq	449	gi 4506668 ref NM_001003.1 RPLP1	Homo sapiens ribosomal protein, large, P1 (RPLP1) mRNA	0
1092.seq	442	gi 2121300 gb AF002992 HSAFD02992	Homo sapiens cosmid from Xq28, complete sequence [Homo sapiens]	6.00E-25
1093.seq	513	gi 4503470 ref NM_001402.1 EEF1A1	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
1094.seq	493	gi 4826450 emb Z93241.1 HS222E13	Human DNA sequence from clone 222E13 on chromosome 22. Contains three	0
1095.seq	467	gi 870744 db U05572 HUMDCB	Human mRNA for heterogeneous nuclear ribonucleoprotein D (hnRNP D),	0
1096.seq	274	gi 4506890 ref NM_003011.1 SET1	Homo sapiens SET translocation (myeloid leukemia-associated) (SET),	e-126
1098.seq	494	gi 3483490 gb AF086145 HUMZA99C08	Homo sapiens full length insert cDNA clone ZA99C08	0
1099.seq	439	gi 4336324 gb AF081282 AF081282	Homo sapiens small membrane protein 1 (SMP1) mRNA, complete cds	e-139
1101.seq	511	gi 4588491 gb AF108826.1 AF108826	Terebra subulata 16S ribosomal RNA gene, mitochondrial gene	0.034
1102.seq	451	gi 4826548 emb AL049764.4 HS362J20	Human DNA sequence from clone 362J20 on chromosome 22q13.1-13.31	0.008
1103.seq	407	gi 14015 emb X55654 MTSCOXII	H.sapiens mitochondrial coxII mRNA for cytochrome C oxidase II	0
1104.seq	464	gi 33290 emb X06763 HSIGKVAH	Human mRNA for Ig kappa L-chain variable region (HumkvHAH)	0
1106.seq	463	gi 4506722 ref NM_001006.1 RPS3A	Homo sapiens ribosomal protein S3A (RPS3A) mRNA	0
1107.seq	479	gi 3386533 gb AF078934 AF078934	Glycine max mariner element Soymer1 transposase gene, complete cds	0.13
1108.seq	482	gi 3080352 emb AL025880.1 ATT5K18	Arabidopsis thaliana DNA chromosome 4, BAC clone TSK18 (ESSA project)	0.5
1109.seq	485	gi 337755 gb J03015 HUMSAP	Homo sapiens sphingolipid activator protein 1 mRNA, 3' end.	0
1110.seq	488	gi 3893858 gb AF051941 AF051941	Homo sapiens type 6 nucleoside diphosphate kinase NM23-H6 (NM23-H6)	e-109
1111.seq	423	gi 4504900 ref NM_002268.1 KPNA4	Homo sapiens karyopherin alpha 4 (importin alpha 3) (KPNA4) mRNA	0
1113.seq	479	gi 3341979 gb AF049523 AF049523	Homo sapiens huntingtin-interacting protein HYPA/FBP11 (HYPA) mRNA,	e-118
1114.seq	475	gi 4759269 ref NC_004622.1	Homo sapiens tranlsin (TSN) mRNA >gi 607129 emb X78627 HS-TRANSJ	0.57
1115.seq	496	gi 3954884 emb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	e-171
1118.seq	470	gi 4406648 gb AF131814 AF131814	Homo sapiens clone 24983 mRNA sequence	0
1120.seq	476	gi 5668629 emb DAL109619.1 ATT19F6	Arabidopsis thaliana DNA chromosome 4, BAC clone T19F6, partial	5.00E-04
1122.seq	432	gi 385936 gb S61826 S61826	hinge-OXPHOS system complex III mitochondrial subunit 5' region, Alu	0
1123.seq	473	gi 3702281 gb AC005789 AC005789	Homo sapiens chromosome 19, cosmid F5960, complete sequence [Homo]	e-111
1124.seq	467	gi 3483780 gb AF086435 HUMZD80D09	Homo sapiens full length insert cDNA clone ZD80D09	5.00E-47
1126.seq	469	gi 4506954 ref NM_003896.1 SIATGM3S	Homo sapiens sialyltransferase 9 (CMP-NeuAc: lactosylceramide	0
1127.seq	427	gi 577294 db U042041 HUMKIAAE	Human mRNA for KIAA0088 gene, partial cds.	e-177
1128.seq	476	gi 3510226 gb AC005575 AC005575	Homo sapiens chromosome 5, BAC clone 246J10 (LBNL.H186), complete	0
1129.seq	497	gi 4503360 ref NM_001383.1 DPH2L1	Homo sapiens diptheria toxin resistance protein required for	0
1130.seq	435	gi 5912120 emb AL117568.1 HSM801095	Homo sapiens mRNA; cDNA DKFZp564C1216 (from clone DKFZp564C1216)	0
1131.seq	457	gi 4507762 ref NM_003334.1 UBE1	Homo sapiens ubiquitin-activating enzyme E1 (A1S9T and BN75	0
1133.seq	446	gi 3097871 gb AC003663 AC003663	Homo sapiens chromosome 17, clone HCIT87G17, complete sequence [Homo]	0
1134.seq	468	gi 3201899 gb AF067420 AF067420	Homo sapiens SNCT3 protein (SNCT3) mRNA, complete cds	0
1135.seq	451	gi 3287468 gb AC004448 AC004448	Homo sapiens chromosome 17, clone HCIT187M2, complete sequence [Homo]	0
1136.seq	347	gi 3805922 emb AL03120 HS459L4	Human DNA sequence from clone 459L4 on chromosome 6p22.3-24.1 Contain	0.35

1137.seq						
1138.seq	451	g 4406571 gb AF131753 AF131753	Homo sapiens clone 24859 mRNA sequence			
1139.seq	492	g 3337392 gb AC004531 HUAC004531	Homo sapiens Chromosome 16 BAC clone CIT987SK-A-67A.1, complete sequ		4.00E-85	
1140.seq	486	g 4501882 ref NM_001613.1 ACTA2	Homo sapiens actin alpha 2 (ACTA2) mRNA >g 283292 emb X13839 HSACTA			
1142.seq	459	g 33479 emb X57086 HSHGMHCC	H.sapiens mRNA for IgM heavy chain constant domain			
1143.seq	462	g 5042231 emb Y13323.2 HSDJSPRO	Homo sapiens mRNA for disintegrin-protease			
1144.seq	485	g 4557576 ref NM_001443.1 FABP1	Homo sapiens fatty acid binding protein 1, liver (FABP1) mRNA			
1146.seq	464	g 23505 emb Z16917 HS205WH10	H.sapiens (D5S416) DNA segment containing (CA) repeat: clone		0.48	
1147.seq	323	g 189446 gb M17252 HUMP45C21	Human cytochrome P450c21 mRNA, 3' end.		4.00E-99	
1148.seq	476	g 4185272 gb AF080157 AF080157	Homo sapiens I-kB kinase-a (IKK-alpha) mRNA, complete cds		1.00E-81	
1149.seq	388	g 4504618 ref NM_001553.1 IGFBP7	Homo sapiens insulin-like growth factor binding protein 7 (IGFBP7)			
1150.seq	530	g 484292 dbj D16237 RAICDC25B	Rat mRNA for cdc25B, complete cds		0.14	
1151.seq	496	g 4809277 ref NM_001156.2 ANX7A	Homo sapiens annexin VII (synexin) (ANX7), splice variant 1, mRNA			
1152.seq	476	g 4506730 ref NM_001010.1 RPS6I	Homo sapiens ribosomal protein S6 (RPS6) mRNA			
1156.seq	468	g 916 emb X56388 CFRAB11	Canine rab11 mRNA for ras-related GTP-binding protein			
1157.seq	182	g 33082 emb X69860 HSHGHZCA	H.sapiens lg rearranged heavy chain mRNA VDJ region		4.00E-36	
1158.seq	195	g 31091 emb J16869 HSEFTAC	Human mRNA for elongation factor 1-alpha (clone CEF4)		4.00E-55	
1159.seq	286	g 5678818 gb AF146191.1 AF146191	Homo sapiens FRG1 (FRG1) gene, complete cds; 5S ribosomal RNA gene,		2.00E-94	
1160.seq	319	g 4507876 ref NM_003373.1 VCLL	Homo sapiens vinculin (VCL) mRNA >g 340236 gb M33308 HUMVINC Human	e-118		
1161.seq	311	g 4589272 gb AF135023.1 AF135023	Homo sapiens kallikrein-like protein 1 KLK-L1 gene, partial cds		0.02	
1163.seq	206	g 5454161 ref NM_006472.1 VDUP1	Homo sapiens upregulated by 1,25-dihydroxyvitamin D-3 (VDUP1) mRNA	e-112		
1164.seq	200	g 3859054 emb AL033403 HS88D7	Human DNA sequence from clone 88D7 on chromosome Xq25-26.3 Contains		0.05	
1165.seq	339	g 4506606 ref NM_000979.1 RPL18	Homo sapiens ribosomal protein L18 (RPL18) mRNA			
1166.seq	367	g 4505434 ref NM_000906.1 NPR1	Homo sapiens natriuretic peptide receptor A/guanylate cyclase A	e-173		
1167.seq						
1168.seq	176	g 2832592 emb Z95116 HS57G9	Human DNA sequence from BAC 57G9 on chromosome 22q12.1 Contains EST		5.00E-79	
1169.seq	379	g 4511885 gb AF044418.1 INDUFAV4	Homo sapiens NADH-ubiquinone oxidoreductase 13 kDa subunit (NDUFA5)	e-180		
1170.seq	179	g 2335061 gb AF001550 HSAF001550	Homo sapiens chromosome 16 BAC clone CIT987SK-334D11 complete sequ		0.003	
1171.seq	355	g 2243159 gb M83233 HUMHTF4A	Homo sapiens transcription factor (HTF4A) mRNA, complete cds			
1173.seq						
1175.seq	189	g 5817212 emb AL110253.1 HSM800915	Homo sapiens mRNA: cDNA DKFZp566B1346 (from clone DKFZp566B1346)		3.00E-43	
1179.seq	209	g 5454027 ref NM_006270.1 RRAS	Homo sapiens related RAS viral (v-ras) oncogene homolog (RRAS) mRNA		3.00E-56	
1180.seq	344	g 804748 gb L42024 HUMHLABH	Homo sapiens MHC HLA-B39 mRNA, complete cds.	e-138		
1181.seq	385	g 3779035 gb AC004526 AC004526	Homo sapiens chromosome 17, Neurofibromatosis 1 locus, complete sequence	e-102		
1182.seq	225	g 5454161 ref NM_006472.1 VDUP1	Homo sapiens upregulated by 1,25-dihydroxyvitamin D-3 (VDUP1) mRNA		2.00E-57	
1183.seq	391	g 984508 gb U07802 HJU07802	Human Tis1d gene, complete cds.	e-148		
1184.seq	464	g 1786921 gb M18217 HUMANTNCB	Human DNA of undetermined origin found 5' to NCA mRNA.		0	
1185.seq	486	g 160324 gb M88110 PFAHGPTA	Plasmodium falciparum hypoxanthine-guanine phosphoribosyl transferase		0.13	
1186.seq	465	g 2977767 emb X67813 CFSPRP72	C.familialis SRP72 mRNA for signal recognition particle		7.00E-28	
1188.seq	230	g 3283923 gb AF070649 AF070649	Homo sapiens clone 24452 mRNA sequence	e-125		
1189.seq	458	g 5931523 dbj AP000545.1 AP000545	Homo sapiens genomic DNA, chromosome 22q11.2, Cat Eye Syndrome region		0.48	
1190.seq	453	g 4971212 emb AL049776.2 CNS00001	Human chromosome 14 DNA sequence *** IN PROGRESS *** BAC R-85K15		0.12	
1191.seq	358	g 4504966 ref NM_002299.1 LCT	Homo sapiens lactase (LCT) mRNA >g 34399 emb X07994 HSLPH Human m			
1192.seq	391	g 185267 gb M34024 HUMHGHCXE	Human fetal lg heavy chain variable region (clone M43) mRNA.	e-176		
1194.seq	393	g 4505894 ref NM_002669.1 PRRG1	Homo sapiens pleiotropic regulator 1 (PR1, Arabidopsis homolog).	e-160		
1195.seq	322	g 4759027 ref NM_004164.1 RBP2	Homo sapiens retinol-binding protein 2, cellular (RBP2) mRNA	e-150		
1196.seq	327	g 3885371 dbj AB019568 AB019568	Homo sapiens mRNA expressed only in placental villi, clone SMAP83	e-174		
1197.seq	310	g 4894853 gb AF135157.1 AF135157	Homo sapiens complement C1q A chain precursor, mRNA, complete cds	e-174		
1198.seq	413	g 5668757 gb AC008009.4 AC008009	Homo sapiens 3q26.2-27 BAC RPC11-436A20 (Roswell Park Cancer Institut		1.00E-75	
1199.seq						
1201.seq	283	g 3954884 emb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,		2.00E-85	
1202.seq	341	g 3005935 emb AJ004955 HSA004955	Homo sapiens immunoglobulin kappa light chain variable region	e-170		
1203.seq	456	g 4154164 gb U88086 HUMZF4	Human zinc finger protein (ZNF165), gene, exon 5, and complete cds		8.00E-06	
1204.seq	349	g 402644 emb X73501 HSCYTKO20	H.sapiens gene for cytokeratin 20		0	
1205.seq	311	g 4263636 gb AC005150 AC005150	Homo sapiens chromosome 4 clone B55824 map 4q25, complete sequence		0.08	
1206.seq	489	g 4885220 ref NM_005239.1 ETS2	Homo sapiens v-ets avian erythroblastosis virus E26 oncogene			
1208.seq	351	g 4506516 ref NM_002923.1 RGS2	Homo sapiens regulator of G-protein signalling 2, 24kd (RGS2) mRNA	e-158		
1210.seq	448	g 5454047 ref NM_006379.1 SEMA3C	Homo sapiens sema domain, immunoglobulin domain (Ig), short basic		0	
1211.seq	374	g 6006017 ref NM_006149.2 LGALS4	Homo sapiens lectin, galactoside-binding, soluble, 4 (galectin 4)			
1212.seq	122	g 339722 gb J03592 HUMTLCA	Human ADP/ATP translocase mRNA, 3' end, clone pHAT8.		2.00E-34	
1213.seq	392	g 1478196 emb X97321 HSHLACW07	H.sapiens mRNA for human leukocyte antigen C alpha chain		0	
1214.seq	388	g 3025444 gb AC004528 AC004528	Homo sapiens chromosome 19, cosmid R32184, complete sequence [Homo	e-150		
1215.seq	365	g 3287468 gb AC004448 AC004448	Homo sapiens chromosome 17, clone HCIT187M2, complete sequence [Homo	e-177		
1217.seq	493	g 5701616 gb AC006160.9 AC006160	Homo sapiens chromosome 4 clone C0481P14 map 4p16, complete sequence		0	
1218.seq	420	g 5803180 ref NM_006819.1 STIP1	Homo sapiens stress-induced-phosphoprotein 1 (Hsp70/Hsp90-organizing		0	
1219.seq	489	g 190978 gb M18963 HUMREGA	Human islet of Langerhans regenerating protein (reg) mRNA, complete			
1223.seq	454	g 3355518 gb AC005029 AC005029	Homo sapiens BAC clone GS588G18 from 7p12-p14, complete sequence [Hor		0.47	
1224.seq	265	g 5114052 gb AF085359.1 AF085359	Homo sapiens HSPC030 mRNA, complete cds		1.00E-64	
1225.seq	440	g 4240298 dbj AB020712.1 AB020712	Homo sapiens mRNA for KIAA0905 protein, complete cds		0	
1226.seq	445	g 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.		0	
1227.seq	349	g 4503056 ref NM_001885.1 CRYAB	Homo sapiens crystallin, alpha B (CRYAB) mRNA		0	
1228.seq	453	g 1698575 gb U60415 HSLU60415	Human bHLH-PAS protein JAP3 mRNA, complete cds		0	
1229.seq	434	g 189347 gb M23204 HUMOATC	Human ornithine-oxo-acid aminotransferase mRNA, exons 3-11, complete		0	
1230.seq	83	g 3256174 emb AL022069 HS168L15	Homo sapiens DNA sequence from PAC 168L15 on chromosome 6q26-27. Co		3.00E-32	
1231.seq	445	g 13000 emb X62996 MIHSGENOM	H.sapiens mitochondrial genome (consensus sequence)			
1232.seq	399	g 1151150 gb M62866 PSEC0BGEN	P.denitrificans cobN, cobO, cobP, cobQ, cobW, and ORF6-9 genes.		0.1	
1233.seq	411	g 4455632 emb J97353.3 HS90L6	Human DNA sequence from clone 90L6 on chromosome 22q11.21-11.23. Con		0.11	
1234.seq	452	g 665587 dbj D45887 HUMALDN	Human mRNA for calmodulin, complete cds		0	
1235.seq	490	g 4503800 ref NM_003902.1 FUBP1	Homo sapiens far upstream element binding protein (FUBP) mRNA		0	
1238.seq	375	g 5138905 gb AF092128.1 AF092128	Homo sapiens putative transmembrane protein E3-16 mRNA, complete		0	
1239.seq	469	g 5019423 emb AL049775.2 CNS00006	Human chromosome 14 DNA sequence *** IN PROGRESS *** BAC R-497E1		8.00E-06	
1240.seq	237	g 1088447 dbj D49387 HUMNADPDLB	Human mRNA for NADP dependent leukotriene b4		7.00E-48	
1250.seq	444	g 31974 emb X52317 HSH2AZ	Human mRNA for histone H2A.Z			
1251.seq	502	g 5689544 dbj AB029027.1 AB029027	Homo sapiens mRNA for KIAA1104 protein, complete cds	E-159		
1252.seq	359	g 31974 emb X52317 HSH2AZ	Human mRNA for histone H2A.Z	E-168		
1253.seq	472	g 747649 dbj D49839 ASNPEPAB	Aspergillus niger pepA gene for aspergillopepsin A precursor.		1.9	
1256.seq	454	g 4454331 emb AJ224356 SLY224356	Solanum lycopersicon tDET1 gene		0.03	
1258.seq	473	g 5689226 dbj AB021226.1 AB021226	Mus musculus mRNA for membrane-type-5 matrix metalloproteinase,		3.00E-67	
1259.seq	448	g 31988 emb X05857 HSH33G4	Human H3.3 gene exon 4	E-125		
1261.seq						
1262.seq	368	g 5739179 gb AF120982.1 HSM1MBD4	Homo sapiens methyl-CpG binding protein 1 (MBD1) gene, exon 15b		5.00E-96	
1263.seq	505	g 5453897 ref NM_006221.1 PIN1	Homo sapiens protein (peptidyl-prolyl cis/trans isomerase)			
1264.seq	435	g 665924 gb U17248 HSUT17248	Human succinate dehydrogenase iron-protein subunit (sdhB) gene.		0	
1265.seq	427	e-110				
1266.seq	491	g 3990474 gb AC004195 AC004195	Homo sapiens clone UWGC-y24c027 from 6p21, complete sequence [Homo		0.51	
1268.seq	458	g 4758345 ref NM_004107.1 FCGR1	Homo sapiens Fc fragment of IgG, receptor, transporter, alpha		0	
1269.seq	319	g 4826450 emb Z93241.1 HS222E13	Human DNA sequence from clone 22E13 on chromosome 22. Contains three	E-169		
1270.seq	315	g 5823975 emb AJ238395.1 HSA238395	Homo sapiens mRNA for RPGR protein, retina specific isoform		0.32	
1271.seq	494	g 4501886 ref NM_001614.1 ACTG1	Homo sapiens actin, gamma 1 (ACTG1) mRNA		0	
1272.seq	437	g 425353 gb U03643 HSU03643	Human leukoaphysin (LKP) mRNA, complete cds.		0	
1275.seq	334	g 2224704 dbj AB002380 AB002380	Human mRNA for KIAA0382 gene, partial cds		E-176	
1277.seq	472	g 337297 gb M10546 HUMMTM1	Human mitochondrial DNA, fragment M1, encoding transfer RNAs.		0	
1278.seq	417	g 1035240 emb Z62862 HS7481R	H.sapiens CpG island DNA genomic MseI fragment, clone 74b1, reverse	E-114		
1279.seq	515	g 4506800 ref NM_003469.1 SCG2	Homo sapiens secretogranin II (chromogranin C) (SCG2) mRNA		0	
1280.seq	481	g 37847 emb X16478 HSMVIMCD	Human mRNA 5'-fragment for vimentin N-terminal fragment		0	
1281.seq	476	g 1507673 dbj D87120 D87120	Homo sapiens mRNA for GS3786, complete cds	E-151		
1282.seq	360	g 5031714 ref NM_005716.1 GLUT1CBF	Homo sapiens GLUT1 C-terminal binding protein (GLUT1CBP) mRNA		8.00E-67	
1283.seq	529	g 2623584 gb AF027158 AF027158	Homo sapiens immunoglobulin kappa light chain (T61/k) mRNA, partial		0	

Accession	Gene/Feature	Species	Description	Value
1284.seq	451 gi 401844 gb L05093 HUMR1BPROD	Homo sapiens	ribosomal protein L18a mRNA, complete cds	0
1285.seq	506 gi 5103017 db J AF000354.1 JAP000354	Homo sapiens	genomic DNA, chromosome 22q11.2, clone KB1674E1	2.1
1287.seq	516 gi 306741 gb J 2006 HUMERCC5A	Human	ERCC5 excision repair protein mRNA, complete cds	0
1288.seq	311 gi 386007 gb AF097021 AF097021	Homo sapiens	GW112 protein (GW112) mRNA, complete cds	E-143
1291.seq	470 gi 3413792 emb AJ010118 HSA010118	Homo sapiens	mRNA for mutant NADH-cytochrome b5 reductase, partial	0
1293.seq	332 gi 4505022 ref NM_003720.1 DSCR2	Homo sapiens	Down syndrome critical region gene 2 (DSCR2), mRNA	3.00E-72
1294.seq	454 gi 5454053 ref NM_006276.1 SFRS7	Homo sapiens	splicing factor, arginine/serine-rich 7 (35kD) (SFRS7)	E-102
1295.seq	498 gi 5921996 ref NM_004343.2 CALR	Homo sapiens	calreticulin (CALR), mRNA	0
1296.seq	451 gi 476704 gb L26336 HUMHSPA2A	Human	heat shock protein HSPA2 gene, complete cds	E-158
1297.seq	439 gi 1469867 db D63477 D63477	Human	mRNA for KIAA0143 gene, partial cds	E-156
1299.seq	458 gi 506340 db J D31888 HUMORFKG1Q	Human	mRNA for KIAA0071 gene, partial cds	E-179
1300.seq	475 gi 5835387 ref NC_018007.1	Human	mitochondrion, complete genome >gi 1944628 gb J01415 HUMMTCG	0
1301.seq	427 gi 2502049 gb AF022104 AFD22104	Neochanna cleaveri	16S ribosomal RNA gene, mitochondrial gene for	1.8
1302.seq	234 gi 3978547 gb AF109718.1 HSCOS10	Homo sapiens	chromosome 3 subtelomeric region	E-104
1304.seq	478 gi 1302649 gb U52111 HSU52111	Homo sapiens	Xq28 genomic DNA in the region of the ALD locus containing	3.00E-61
1306.seq	271 gi 1517817 gb U33834 HSU33834	Human	CHL1-related helicase (CHLR2) mRNA, partial cds	E-123
1307.seq	254 gi 4375969 emb AL031432 HS465N24	Human	DNA sequence from clone 465N24 on chromosome 1p35.1-36.13	1.00E-52
1308.seq	491 gi 5729910 ref NM_006691.1 LYVE-1	Homo sapiens	lymphatic vessel endothelial hyaluronan receptor 1	0
1311.seq	422 gi 4062176 gb AC006121 AC006121	Homo sapiens	chromosome X, clone hCIT.200_L_4, complete sequence	0
1312.seq	480 gi 1262342 emb X93334 HSMITG	H.sapiens	mitochondrial DNA, complete genome	0
1314.seq	410 gi 3287320 db AF014891.1 AF014891	Homo sapiens	NADH dehydrogenase subunit 2 (ND2) gene, mitochondrial	0
1315.seq	502 gi 341790 gb M27877 M27877	Figure 2	Nucleotide and translated protein sequences of HPF1, -2,	4.00E-17
1317.seq	459 gi 181532 gb M97925 HUMDEF5A	H.sapiens	defensin 5 gene, complete cds	E-108
1319.seq	442 gi 4557520 ref NM_000398.2 DIA1	Homo sapiens	diaphorase (NADH) (cytochrome b-5 reductase) (DIA1),	0
1320.seq	196 gi 181962 gb J04617 HUMEF1A	Human	elongation factor EF-1-alpha gene, complete cds	4.00E-30
1321.seq	258 gi 3126781 gb AC003957 AC003957	Homo sapiens	chromosome 17, clone hCIT.457_L_16, complete sequence	1
1322.seq	463 gi 3201899 gb AF067420 AF067420	Homo sapiens	SNC73 protein (SNC73) mRNA, complete cds	0
1323.seq	485 gi 4504164 ref NM_000177.1 GSN	Homo sapiens	gelsolin (amyloidosis, Finnish type) (GSN) mRNA	0
1324.seq	509 gi 3327141 db J A014564 A014564	Homo sapiens	mRNA for KIAA0664 protein, partial cds	0
1326.seq	476 gi 2765422 emb Y14736 HISGIG1KL	Homo sapiens	mRNA for immunoglobulin kappa light chain	E-151
1327.seq	344 gi 5901895 ref NM_006886.1 ATP5E	Homo sapiens	ATP synthase, H+ transporting, mitochondrial F1	E-171
1328.seq	163 gi 4633837 gb AF139813.1 AF139813	Homo sapiens	clone pDJ759J12 chromosome 11 map 11q13, complete sequen	1.00E-85
1329.seq	439 gi 5729876 ref NM_006597.1 HSPA10	Homo sapiens	heat shock 70kD protein 10 (HSC71) (HSPA10), mRNA	0
1331.seq	453 gi 31091 emb X16869 HSEFIAC	Human	mRNA for elongation factor 1-alpha (clone CEF4)	0
1332.seq	456 gi 2661047 gb AF035293 AF035293	Homo sapiens	clones 23797 and 23917 mRNA, partial cds	E-126
1334.seq	502 gi 4680716 gb AF132973.1 AF132973	Homo sapiens	CGI-39 protein mRNA, complete cds	0
1335.seq	469 gi 532596 gb M12759 HUMIGJ02	Human	Ig J chain gene, exons 3 and 4.	0
1336.seq	502 gi 4506310 ref NM_002840.1 PTPRF	Homo sapiens	protein tyrosine phosphatase, receptor type, F (PTPRF)	E-162
1339.seq	527 gi 5835387 ref NC_018007.1	Human	mitochondrion, complete genome >gi 1944628 gb J01415 HUMMTCG	0
1340.seq	514 gi 1389630 db J D7836 HUMODAZ	Human	mRNA for ornithine decarboxylase antizyme, ORF 1 and ORF 2	0
1342.seq	510 gi 4506744 ref NM_001013.1 RPS9	Homo sapiens	ribosomal protein S9 (RPS9) mRNA	0
1343.seq	447 gi 3449326 db J A016885.1 A016885	Arabidopsis thaliana	genomic DNA, chromosome 5, TAC clone: K19M22,	0.12
1345.seq	520 gi 4506742 ref NM_001012.1 RPS8	Homo sapiens	ribosomal protein S8 (RPS8) mRNA	0
1346.seq	231 gi 1200181 gb U33317 HUS33317	Human	defensin 6 (HD-6) gene, complete cds	E-116
1347.seq	488 gi 190978 gb M18963 HUMREGA	Human	islet of Langerhans regenerating protein (reg) mRNA, complete	0
1348.seq	452 gi 3327191 db J A014589 A014589	Homo sapiens	mRNA for KIAA0689 protein, partial cds	0
1349.seq	518 gi 4321848 gb AF064879 AF064879	Homo sapiens	clone C2 immunoglobulin heavy chain alpha VDJ region	1.00E-75
1350.seq	177 gi 3419836 gb AC005389 AC005389	Homo sapiens	chromosome 17, clone hRPK.601_N_13, complete sequence [H]	0.68
1351.seq	483 gi 431309 gb L23116 HUMGALC	Homo sapiens	galactocerebroside galactosyl transferase (GALC) mRNA, complete cds	0
1352.seq	314 gi 332022 emb J211890 HISGK3BS1	H.sapiens	rearranged mRNA for immunoglobulin kappa chain (VNJ)	4.00E-50
1354.seq	413 gi 4753220 gb AC005095.2 AC005095	Homo sapiens	BAC clone RG332P12 from 7q22-q31.1, complete sequence	E-149
1355.seq	371 gi 1617516 gb U67784 HUSU67784	Human	orphan G protein-coupled receptor (RDC)1 mRNA, partial cds	E-140
1357.seq	313 gi 28401 emb X04353 HSAHDH3G1	Human	mRNA for liver alcohol dehydrogenase (EC 1.1.1.1) gamma	E-170
1358.seq	495 gi 2337864 gb AC002464 AC002464	Human	BAC clone RG331P03, complete sequence [Homo sapiens]	0.002
1359.seq	415 gi 3287468 gb AC004448 AC004448	Homo sapiens	chromosome 17, clone hCIT187M2, complete sequence [Homo	E-180
1360.seq	202 gi 3282156 gb AC005222 AC005222	Homo sapiens	cosmid clone RT163 (LANL), complete	6.00E-88
1361.seq	203 gi 2576344 gb AC002400 HUAAC002400	Human	Chromosome 16 BAC clone CIT987SK-A-735G6, complete sequence	6.00E-14
1362.seq	478 gi 3789975 gb AF097649 AF097649	Homo sapiens	chromosome 7qtel BAC ES, complete sequence [Homo sapiens]	0.032
1363.seq	424 gi 4505896 ref NM_002670.1 PLS1	Homo sapiens	plastin 1 (I isoform) (PLS1) mRNA	0
1364.seq	468 gi 4502204 ref NM_001660.1 ARF4	Homo sapiens	ADP-ribosylation factor 4 (ARF4) mRNA	0
1365.seq	471 gi 4557796 ref NM_002699.1 NME1	Homo sapiens	non-metastatic cells 1 (NME1) mRNA	0
1366.seq	263 gi 185948 gb M11737 HUMIGKCB	Human	Ig gamma chain C-region from kappa-deficient	3.00E-75
1367.seq	533 gi 4757753 ref NM_004306.1 ANXA13	Homo sapiens	annexin A13 (ANXA13), mRNA	0
1369.seq	265 gi 38334 emb X62125 HSWLRG	H.sapiens	rearranged WLR gene for lambda-immunoglobulin subgroup II	3.00E-56
1370.seq	507 gi 34717 emb X64899 HSPM21HOM	H.sapiens	mRNA homologous to mouse P21 mRNA	0
1371.seq	485 gi 3168605 gb AF021937 AF021937	Arabidopsis thaliana	catalase 3 (CAT3) and catalase 1 (CAT1) genes.	0.51
1375.seq	419 gi 262368 gb SS1768 SS1768	PAP-Hpancreatitis-associated protein [Human, pancreas, mRNA, 797		
1376.seq	449 gi 4886224 ref NM_005243.1 EWSR1	Homo sapiens	Ewing sarcoma breakpoint region 1 (EWSR1) mRNA	E-155
1377.seq	397 gi 186267 gb M34024 HUMIGHCXE	Human	fetal Ig heavy chain variable region (clone M43) mRNA,	E-125
1378.seq				
1379.seq	445 gi 3005010 gb AF040759 MMGRK5S05	Mus musculus	G protein-coupled receptor kinase 5 (GRK5) gene, exons	0.12
1381.seq	351 gi 1294809 gb L08086 HUMIGH3AE	Human	(clone MAB57) Ig H-chain (IGH@) mRNA, partial cds, VH-1	6.00E-77
1383.seq	359 gi 790235 gb U21158 OCU21156	Oryctolagus cuniculus	sarcosin-associated protein-2 mRNA, complete	0
1384.seq	454 gi 4503470 ref NM_001402.1 EEF1A1	Homo sapiens	eukaryotic translation elongation factor 1 alpha 1	0
1386.seq	381 gi 5107075 gb AF144700.1 AF144700	Homo sapiens	small zinc finger-like protein (TIM13) mRNA, complete	0
1387.seq	316 gi 4034056 emb Z85986.1 HS108K11	Human	DNA sequence from clone 108K11 on chromosome 6p21 Contains SRF	E-178
1388.seq	446 gi 5733977 gb AF120276.1 AF120276	Oryctolagus cuniculus	sarcosin-associated protein, complete cds	5.00E-13
1389.seq	464 gi B44480 db J D38112.1 HUMMTA	Homo sapiens	mitochondrial DNA, complete sequence	0
1390.seq	374 gi 4506678 ref NM_001014.1 RPS10	Homo sapiens	ribosomal protein S10 (RPS10) mRNA	0
1391.seq	453 gi 2292904 emb Y10344 HSY10344	H.sapiens	GaINAc-T2 gene, 3'UTR	0
1393.seq	345 gi 432358 db J D23660 HUMRSP	Human	mRNA for ribosomal protein, complete cds	E-108
1394.seq	448 gi 488591 gb U09477 HSU09477	Human	clone 53BP1 p53-binding protein mRNA, partial cds.	0.47
1395.seq	516 gi 3650436 db J A017766.1 A017766	Protopterus annectens	mRNA for somatolactin, complete cds	0.14
1396.seq	105 gi 482908 gb L15533 HUMPAP	Homo sapiens	pancreatitis-associated protein (PAP) gene, complete cds.	6.00E-34
1398.seq	442 gi 219897 db J D90453 HUMLCA2	Human	mRNA for l-caldesmon II	4.00E-35
1400.seq	509 gi 4506662 ref NM_000973.1 RPL8	Homo sapiens	ribosomal protein L8 (RPL8) mRNA	0
1401.seq	472 gi 665587 db J D45887 HUMALDN	Human	mRNA for calmodulin, complete cds	E-165
1402.seq	495 gi 5926888 db J A000501.1 A000501	Homo sapiens	genomic DNA, chromosome 8p11.2, clone:91h23 to 9-41	E-176
1403.seq	548 gi 532596 gb M12759 HUMIGJ02	Human	Ig J chain gene, exons 3 and 4.	E-128
1405.seq	460 gi 36578 emb Y00692 HSSPROTR	Human	mRNA for protein S	0
1406.seq	457 gi 572551.1 gb AF072097.1 AF072097	Homo sapiens	beta-2-microglobulin gene, complete cds	0
1408.seq	448 gi 3282164 gb AC005216 AC005216	Homo sapiens	chromosome 5, BAC clone 282B7 (LBNL H192), complete	E-116
1409.seq	446 gi 5729876 ref NM_006597.1 HSPA10	Homo sapiens	heat shock 70kD protein 10 (HSC71) (HSPA10), mRNA	0
1410.seq	480 gi 4758785 ref NM_004550.1 INDUF52	Homo sapiens	NADH dehydrogenase (ubiquinone) F6-S protein 2 (49kD)	0
1412.seq	487 gi 182506 gb M12937 HUMFERHA	Human	ferritin heavy subunit mRNA, complete cds.	E-173
1413.seq	351 gi 1669681 emb Z82901 HSS06G2A	Human	DNA sequence from PAC 506G2 contains ESTs	0.091
1414.seq	463 gi 265703 gb S55735 S55735	Ig alpha 1 - alpha 2m-immunoglobulin A1-A2 lambda	hybrid GAU heavy	0
1415.seq	476 gi 3170864 gb AF062201 AF062201	Homo sapiens	clone 48u-41 immunoglobulin heavy chain variable	E-158
1416.seq	394 gi 532596 gb M12759 HUMIGJ02	Human	Ig J chain gene, exons 3 and 4.	E-130
1417.seq	444 gi 1262342 emb X93334 HSMITG	H.sapiens	mitochondrial DNA, complete genome	E-152
1418.seq	537	Human	DNA sequence from clone 134N	0.14
1419.seq	480 gi 3970934 gb AC006128 AC006128	Homo sapiens	chromosome 19, cosmid F20900, complete sequence [Homo	0.032
1421.seq	448 gi 1177288 emb X54452 DD31	D.discoidium	culmination spA (Dd31) gene	0.008
1422.seq				
1423.seq	491 gi 3201899 gb AF067420 AF067420	Homo sapiens	SNC73 protein (SNC73) mRNA, complete cds	0
1424.seq	473 gi 5821146 db J A016089.1 A016089	Homo sapiens	mRNA for RNA binding protein, partial cds, clone: R52	0
1425.seq	466 gi 3201899 gb AF067420 AF067420	Homo sapiens	SNC73 protein (SNC73) mRNA, complete cds	0
1426.seq	505 gi 2920805 gb AC003037 AC003037	Homo sapiens	Xp22 BAC GSHB 526D21 (Genome Systems Human BAC librar	5.00E-04

1427.seq	520	g 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-130
1428.seq	384	g 4775603 emb Z93930.10 HS292E10	Human DNA sequence from clone 292E10 on chromosome 22q11-12. Contains	E-140
1429.seq	449	g 4757861 ref NM_004332.1 BPHL	Homo sapiens biphenyl hydrolase-like (serine hydrolase) (BPHL) mRNA	6.00E-59
1430.seq				
1432.seq	463	g 4455133 gb AF130248 AF130248	Homo sapiens clone 10J13 chromosome 21 map 21q11.2, complete sequence	0.12
1433.seq	446	g 3646131 emb AJ010952 HSA010952	Homo sapiens mRNA for putative tRNA splicing protein, partial	0
1434.seq	507	g 4503470 ref NM_001402.1 EEF1A1	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
1435.seq	505	g 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-170
1436.seq	406	g 3367616 gb AL022722 HS354N19	Human DNA sequence from clone 354N19 on chromosome 6q22. Contains the	0
1437.seq	483	g 4504444 ref NM_002136.1 HNPA1	Homo sapiens heterogeneous nuclear ribonucleoprotein A1 (HNPA1) mRNA	E-152
1438.seq	323	g 4580400 gb AC006570.4 ATAC006570	Arabidopsis thaliana chromosome II BAC T16121 genomic sequence,	0.33
1439.seq	431	g 181485 gb M24070 HUMDBPB	Human DNA-binding protein B (dnpB) gene, 3' end.	0
1441.seq	493	g 33479 emb X57086 HSIGMHCC	H.sapiens mRNA for IgM heavy chain constant domain	0
1442.seq	501	g 5738607 emb AJ132584.1 HSA132584	Homo sapiens mRNA for HIRA-interacting protein HIRIP5	E-139
1445.seq	539	g 185397 gb L03162 HUMIGHHA	Human Ig rearranged gamma-chain mRNA V-region, partial cds.	E-160
1446.seq	498	g 182482 gb M12670 HUMFC1	Human fibroblast collagenase inhibitor mRNA, complete cds.	0
1447.seq	502	g 4507668 ref NM_003295.1 TPT1	Homo sapiens tumor protein, translationally-controlled 1 (TPT1)	0
1448.seq	498	g 5419676 emb AJ132917.1 HSA132917	Homo sapiens mRNA for methyl-CoG-binding protein 2	0
1449.seq	503	g 4309693 gb AC005717 AC005717	Drosophila melanogaster, chromosome 2R, region 53E6-53F4, P1 clone	0.53
1450.seq	287	g 4468882 emb AL031657.2 HS28F12	Human DNA sequence from clone 28F12 on chromosome 20q11.22-12 Contain	0.019
1452.seq				
1453.seq	494	g 4506582 ref NM_002945.1 RPA1	Homo sapiens replication protein A1 (70kD) (RPA1) mRNA	0
1454.seq	475	g 1834974 emb Z85263 HS285263	H.sapiens Ig lambda light chain variable region gene (19-12OIVG41)	E-136
1456.seq	477	g 261239 gb S50732 S50732	Immunoglobulin M light chain V region:anti-lipid A antibody (human,	0
1457.seq	499	g 4894367 gb AF067167.1 AF067167	Homo sapiens NADH-ubiquinone oxidoreductase B17 subunit homolog	0
1458.seq	492	g 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-150
1460.seq	455	g 5802967 ref NM_006818.1 AF1Q	Homo sapiens transmembrane protein (AF1Q), mRNA	0
1461.seq	497	g 1032859 emb Z60755 HS33G5F	H.sapiens CpG island DNA genomic MseI fragment, clone 33g5, forward	E-154
1462.seq	458	g 200626 gb M79306 MUSRAB16A	Mouse GTP-binding protein (Rab16) mRNA sequence.	3.00E-42
1464.seq	498	g 12998 emb V00660 MIHS01	Human mitochondrial genes for tRNA (Phe) and 12S rRNA (fragment)	E-142
1465.seq	496	g 186076 gb M11937 HUMIGLAC	Human Ig rearranged kappa light chain mRNA V-region (V-J) partial	0
1466.seq	500	g 1161563 emb X85117 HSEPB72EX	H.sapiens epb72 gene exons 2,3,4,5,6,7	0
1469.seq	492	g 4506676 ref NM_002951.1 RPN2	Homo sapiens ribophorin II (RPN2) mRNA	0
1470.seq	465	g 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-123
1471.seq	167	g 37847 emb X16478 HSMVIMCD	Human mRNA 5'-fragment for vimentin N-terminal fragment	1.00E-76
1472.seq	368	g 2765422 emb Y14736 HSIGG1KL	Homo sapiens mRNA for Immunoglobulin kappa light chain	0
1473.seq	401	g 177064 gb M30684 GORMHCBA	Gorilla gorilla beta-2-microglobulin mRNA (GGOB2M).	0
1474.seq	483	g 13003 emb V00662 MIHSXX	H.sapiens mitochondrial genome	0
1476.seq	452	g 33338 emb X51755 HSIGLAMB	Human lambda-immunoglobulin constant region complex (germline)	0
1477.seq	509	g 1209254 gb L20311 BOVIGJMRNA	Bos taurus mu- and alpha-immunoglobulin-associated J-chain mRNA,	E-103
1478.seq	511	g 1783204 db D17406 D17408	Homo sapiens mRNA for calponin, complete cds	0
1479.seq	503	g 35219 emb X52104 HSP68A	Human mRNA for p68 protein	0
1481.seq	457	g 4929768 gb AF151908.1 AF151908	Homo sapiens CGI-150 protein mRNA, complete cds	0
1482.seq	368	g 4557568 ref NM_001433.1 ERN1	Homo sapiens ER to nucleus signalling 1 (ERN1) mRNA	0.096
1484.seq	453	g 5454236 gb AC008039.1 AC008039	Homo sapiens clone UWG.C.rg391h05 from 7q31, complete sequence	1.00E-72
1485.seq	388	g 334433 emb X14008 HSLYSOZY	Human lysozyme gene (EC 3.2.1.17)	0
1486.seq	370	g 4506664 ref NM_000661.1 RPL9	Homo sapiens ribosomal protein L9 (RPL9) mRNA	0
1487.seq	381	g 2736315 gb AF035528 AF035528	Homo sapiens Smad6 mRNA, complete cds	E-109
1488.seq	504	g 409205 gb M80670 HUMHLAB61	Human germline (HLA-B) gene, complete cds without introns.	0
1489.seq	387	g 5006419 gb AF107045.1 AF107045	Homo sapiens ribonucleotide reductase M1 subunit (RRM1) gene, complete	0.4
1490.seq	478	g 4506622 ref NM_000988.1 RPL27	Homo sapiens ribosomal protein L27 (RPL27) mRNA	0
1491.seq	381	g 4809277 ref NM_001156.2 ANX7A	Homo sapiens annexin VII (synexin) (ANX7), splice variant 1, mRNA	E-163
1492.seq	504	g 2924249 emb Z98745.1 HS29K1	Human DNA sequence from PAC 29K1 on chromosome 6p21.3-22.2. Contains	0.53
1493.seq	515	g 4999920 gb AC007461.8 AC007461	Homo sapiens chromosome 17, clone 84_E_24, complete sequence	0.14
1494.seq	497	g 4538895 emb AL049482.1 ATF17A8	Arabidopsis thaliana DNA chromosome 4, BAC clone F17A8 (ESSA project)	0.008
1495.seq	436	g 180708 gb M29551 HUMCNAB	Human calcineurin A2 mRNA, complete cds. >g 1375291 gb G27041 G27041	0
1496.seq	500	g 37849 emb X56134 HSMVIMENT	Human mRNA for vimentin	0
1497.seq	405	g 1561514 db D38526 D38526	Human mRNA for HLA-C*0702, complete cds	0
1498.seq	439	g 5835387 ref NC_001807.1	Human mitochondrion, complete genome >g 1944628 gb J01415 HUMMTCCG	0
1500.seq	494	g 32822 emb X57331 HSIGCMUDE	Human immunoglobulin C(mu) and C(delta) heavy chain genes (constant	0
1501.seq	494	g 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-119
1502.seq	497	g 5453703 ref NM_006407.1 JWA	Homo sapiens vitamin A responsive; cytoskeleton related (JWA) mRNA	E-139
1504.seq	546	g 5763700 emb AL035684.25 HS1114A1	Human DNA sequence from clone 1114A1 on chromosome 20q13.11-13.2,	2.3
1505.seq	441	g 4753254 gb AC005011.2 AC005011	Homo sapiens BAC clone GS111G14 from 7q11, complete sequence	0.12
1506.seq	269	g 215010 gb L04815 ZEFMHDABAK	Brachydanio rerio MHC class II DA-beta-3*01 gene, 3' end.	0.017
1507.seq	355	g 4468801 emb AL035601 ATF6G17	Arabidopsis thaliana DNA chromosome 4, BAC clone FBG17 (ESSA project)	0.002
1509.seq	433	g 4501888 ref NM_001615.1 ACTG2	Homo sapiens actin, gamma 2, smooth muscle, enteric (ACTG2) mRNA	0
1510.seq	468	g 5419832 emb AL049699.8 HSJ47H23	Human DNA sequence from clone 747H23 on chromosome 6q13-15. Contains	0
1511.seq	344	g 4646249 gb AC005529.7 AC005529	.complete sequence [Homo sapiens]	2.00E-61
1512.seq	510	g 5031634 ref NM_005507.1 CFL1	Homo sapiens coflin 1 (non-muscle) (CFL1) mRNA	0
1513.seq	497	g 5835387 ref NC_001807.1	Human mitochondrion, complete genome >g 1944628 gb J01415 HUMMTCCG	0
1514.seq	489	g 3717970 emb AL031771 HS217016	Human DNA sequence from clone 217016 on chromosome 6q24 Contains G5	0.13
1515.seq	512	g 576472 gb M28204 HUMMHBBZ	Homo sapiens (clone pMF28) MHC class I HLA-B*62 mRNA, 3' end.	0
1516.seq	276	g 3483804 gb AF086459 HUMZD84G09	Homo sapiens full length insert cDNA clone ZD84G09	E-138
1517.seq	444	g 4003342 gb AC005875 AC005875	Homo sapiens chromosome 10 clone CIT987SK-1188B12 map 10p12.1, comp	1.00E-04
1518.seq	510	g 2780183 emb Z99498 HS09L4	Human DNA sequence from PAC, 509L4 on chromosome 6q22.1-8q22.33.	0
1519.seq	509	g 4504164 ref NM_000177.1 GSN	Homo sapiens gelsolin (amyloidosis, Finnish type) (GSN) mRNA	0
1520.seq	493	g 4505058 ref NM_002354.1 MAS1	Homo sapiens membrane component, chromosomal 4, surface marker	E-151
1521.seq	492	g 2358042 gb AE000660 HUA000660	Homo sapiens T-cell receptor alpha delta locus from bases 501613 to	0.033
1522.seq	447	g 1160468 emb Z68694 HSCU177E8	Human DNA sequence from cosmid cU177E8, between markers DXS366 and	1.00E-90
1523.seq	512	g 5649377 gb AC007386.3 AC007386	Homo sapiens clone NH0359K10, complete sequence	3.00E-30
1524.seq	492	g 1934877 emb Z82211 HS447N6	Human DNA sequence from PAC 447N6 on chromosome X contains ESTs and	0.008
1525.seq	477	g 13000 emb X62996 MIHSGENOM	H.sapiens mitochondrial genome (consensus sequence)	E-103
1526.seq	486	g 4506689 ref NM_001023.1 RPS20	Homo sapiens ribosomal protein S20 (RPS20) mRNA	0
1527.seq	501	g 5708490 gb AC004985.2 AC004985	Homo sapiens clone DJ1165K10, complete sequence	0
1531.seq	498	g 2133911 gb AC000394 AC000394	Genomic sequence from Human 9q34, complete sequence [Homo sapiens]	0
1532.seq	491	g 5725511 gb AF072097.1 AF072097	Homo sapiens beta-2 microglobulin gene, complete cds	0
1534.seq	510	g 1929409 emb Z93771 JNTNPP4	N.tabacum npp4 mRNA for protein phosphatase type 2A	0.002
1535.seq	503	g 1665794 db D87453 D87453	Human mRNA for KIAA0264 gene, partial cds	0
1536.seq	436	g 183097 gb M14300 HUMGF12A9	Human growth factor-inducible 2A9 gene, complete cds.	0
1538.seq	500	g 5850575 gb AW003659.1 AW003659	wx34h05.x1 NCI_CGAP_Pit1 Homo sapiens cDNA clone IMAGE:2545593 3'	E-106
1540.seq	474	g 6002388 emb AL121655.1 CNSO1DS5	BAC sequence from the SPG4 candidate region at 2p21-2p22, complete	3.00E-83
1542.seq	486	g 5918434 emb AL049796.1 HSJ561L2	Homo sapiens chromosome 1 clone DJ561L2, WORKING DRAFT SEQUENCE	0.037
1543.seq	455	g 5931442 gb AC010988.1 AC010988	Homo sapiens clone NH0570F04, WORKING DRAFT SEQUENCE, 29 unorder	0.55
1545.seq	437	g 5932620 gb AC009570.2 AC009570	Homo sapiens chromosome 4 clone 46_J_23 map 4, WORKING DRAFT SEQU	0
1546.seq	472	g 6006170 gb AC011133.1 AC011133	Homo sapiens chromosome 15 clone 304_J_8 map 15, LOW-PASS SEQUENC	0.14
1547.seq	483	g 5875534 gb AW022004.1 AW022004	df31f10.y1 Morton Fetal Cochlea Homo sapiens cDNA clone	E-171
1549.seq	500	g 5882699 gb AC010233.1 AC010233	Homo sapiens chromosome 5 clone CIT-HSPC_330K4, WORKING DRAFT SE	0.15
1550.seq	449	g 5837345 gb AI990464.1 AI990464	wf74h03.x1 Soares thymus NHFT Homo sapiens cDNA clone	E-172
1551.seq	202	g 5856271 gb AW007493.1 AW007493	w52d1.x1 NCI_CGAP_Brn25 Homo sapiens cDNA clone IMAGE:2500821 3'	4.00E-49
1552.seq	485	g 5836193 gb AC009958.1 AC009958	Homo sapiens clone NH0316G09, WORKING DRAFT SEQUENCE, 4 unorder	1.00E-72
1553.seq	400	g 5838750 gb AI991845.1 AI991845	wu32d03.x1 Soares_Dieckgraebe_colon_NHCD Homo sapiens cDNA clone	E-161
1555.seq	478	g 5856875 gb AW008108.1 AW008108	wy49g07.x1 NCI_CGAP_Gas4 Homo sapiens cDNA clone IMAGE:990895 3'	0
1556.seq	268	g 5878268 gb AW024738.1 AW024738	wu91b07.x1 NCI_CGAP_Kid3 Homo sapiens cDNA clone IMAGE:990728 3'	E-141
1557.seq	513	g 5874137 gb AW020607.1 AW020607	df12a06.y1 Morton Fetal Cochlea Homo sapiens cDNA clone	0
1558.seq	379	g 5922846 emb AL034422.18 HS1141E1	Homo sapiens chromosome 20 clone DJ1141E15, WORKING DRAFT SEQUEN	6.00E-96
1559.seq				
1560.seq	467	g 6006084 gb AC010122.4 AC010122	Drosophila melanogaster chromosome 3 clone BACR06B08 (D1094) RPCI-98	0.56

1561.seq	366	gij5924454 gb AC009973.2 AC009973	Homo sapiens clone NH0457P23, WORKING DRAFT SEQUENCE, 8 unorder	1.7
1562.seq	274	gij6010191 emb AL121717.1 HJ5214H13	Homo sapiens chromosome 6 clone dJ1214H13, WORKING DRAFT SEQUENCE	0.081
1563.seq	454	gij6013035 emb AJ389927.1 HSA389927	Homo sapiens IGVH gene for immunoglobulin heavy chain variable	E-103
1564.seq	307	gij5878344 gb AW024814.1 AW024814	wu69c07.x1 NCI_CGAP_Kid3 Homo sapiens cDNA clone IMAGE:990943 3'	E-108
1565.seq	433	gij5866319 gb AW014562.1 AW014562	UI-H-B10p-aaaw-c-05-0-UI.s1 NCI_CGAP_Sub2 Homo sapiens cDNA clone	9.00E-12
1566.seq	377	gij4297834 gb AJ446813.1 AJ446813	tj97f07.x1 NCI_CGAP_Gas4 Homo sapiens cDNA clone IMAGE:2139973 3'	E-132
1567.seq	273	gij5932620 gb AC009570.2 AC009570	Homo sapiens chromosome 4 clone 46_J_23 map 4, WORKING DRAFT SEQU	E-125
1568.seq	453	gij5882635 gb AC010298.1 AC010298	Homo sapiens chromosome 5 clone CIT978SKB_163N20, WORKING DRAFT	1.00E-38
1569.seq	443	gij5886304 gb AW027548.1 AW027548	wv74e09.x1 Soares_thymus_NHFTTh Homo sapiens cDNA clone IMAGE:99097	4.00E-88
1570.seq	479	gij5848852 gb AW001936.1 AW001936	wt82a09.x1 Soares_thymus_NHFTTh Homo sapiens cDNA clone	E-165
1571.seq	438	gij5853431 gb AW004733.1 AW004733	ws77g10.x1 NCI_CGAP_Co3 Homo sapiens cDNA clone IMAGE:2504034 3'	E-167
1572.seq	374	gij6013598 gb AC011342.1 AC011342	Homo sapiens chromosome 5 clone CIT-HSPC_276H5, WORKING DRAFT SE	0
1574.seq	311	gij6007929 gb AW060163.1 AW060163	UI-M-BH1-armu-c-04-0-UI.s1 NIH_BMAP_M_S2 Mus musculus cDNA clone	3.00E-57
1575.seq	496	gij5936560 emb AL048547.2 AL048547	DKFZp58600824_r1 586 (synonym: hute1) Homo sapiens cDNA clone	E-152
1576.seq	244	gij5875806 gb AW022276.1 AW022276	df38c05.y1 Morton Fetal Cochlea Homo sapiens cDNA clone	E-105
1577.seq	435	gij5920242 emb AL118403.1 AL118403	w9107b07 Bedlington mouse dissected endoderm Mus musculus cDNA	3.00E-18
1578.seq	461	gij5972957 gb AR050232.1 AR050232	Sequence 147 from patent US 5827516	5.00E-35
1579.seq	483	gij5858363 gb AW009585.1 AW009585	ws84f08.x1 NCI_CGAP_Co3 Homo sapiens cDNA clone IMAGE:2504679 3'	E-115
1580.seq	397	gij4285396 gb AJ444926.1 AJ444926	tj89e12.x1 NCI_CGAP_Co14 Homo sapiens cDNA clone IMAGE:2154286 3'	0
1581.seq	395	gij5827307 emb AL121306.1 AL121306	DKFZp762M176_r1 762 (synonym: hmet2) Homo sapiens cDNA clone	E-130
1582.seq	413	gij5922687 db AP000500.1 AP000500	Homo sapiens genomic DNA, chromosome 3p21.3, clone:603 to 320,	0.008
1583.seq	464	gij5953129 gb AR036461.1 AR036461	Sequence 147 from patent US 5872215	E-111
1584.seq	497	gij5936436 emb AL048319.2 AL048319	DKFZp586B1224_r1 586 (synonym: hute1) Homo sapiens cDNA clone	2.00E-10
1585.seq	422	gij5856334 gb AW007630.1 AW007630	wt57c05.x1 NCI_CGAP_Pan1 Homo sapiens cDNA clone IMAGE:2511580 3'	0
1586.seq	477	gij5887836 gb AW029080.1 AW029080	wx05a06.x1 NCI_CGAP_Gas4 Homo sapiens cDNA clone IMAGE:2542738 3'	8.00E-25
1587.seq	489	gij5936503 emb AL048440.2 AL048440	DKFZp586K0224_r1 586 (synonym: hute1) Homo sapiens cDNA clone	0
1588.seq	454	gij5920829 gb AW055126.1 AW055126	wz01a03.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:2556748 3'	0
1589.seq	427	gij5876180 gb AW022650.1 AW022650	df41g11.y1 Morton Fetal Cochlea Homo sapiens cDNA clone	E-117
1590.seq	472	gij5923004 gb G32483.1 G32483	A009E35 Human Homo sapiens STS genomic, sequence tagged site	1.00E-88
1591.seq	457	gij5888135 gb AW029379.1 AW029379	wx09g11.x1 NCI_CGAP_Gas4 Homo sapiens cDNA clone IMAGE:2543204 3'	1.00E-72
1592.seq	479	gij5858323 gb AW009545.1 AW009545	ws84b08.x1 NCI_CGAP_Co3 Homo sapiens cDNA clone IMAGE:2504631 3'	0
1593.seq	364	gij6014386 emb AL121572.4 HS111822	Homo sapiens chromosome X clone BA512H23, WORKING DRAFT SEQUENCE	1.7
1595.seq	507	gij6002294 emb Z98200.8 HS111822	Human DNA sequence from clone 111822 on chromosome 6q16-21, complete	0
1596.seq	451	gij5925261 emb AL119362.1 AL119362	DKFZp761F021_r1 761 (synonym: hamy2) Homo sapiens cDNA clone	E-176
1597.seq	457	gij5931470 gb AC010971.1 AC010971	Homo sapiens clone NH0122G11, WORKING DRAFT SEQUENCE, 7 unorder	2.2
1598.seq	376	gij5831612 gb AC007746.2 AC007746	Homo sapiens clone NH0520M23, WORKING DRAFT SEQUENCE, 1 unorder	0.11
1599.seq	469	gij5862748 gb AW013991.1 AW013991	UI-H-B10-aaah-c-08-0-UI.s1 NCI_CGAP_Sub1 Homo sapiens cDNA clone	6.00E-04
1600.seq	444	gij5870173 gb AC009773.2 AC009773	Homo sapiens, WORKING DRAFT SEQUENCE, 51 unorderd pieces	0.53
1601.seq	364	gij5922456 gb AW055745.1 AW055745	ESTO8990526 Rat brain ESTs Rattus norvegicus cDNA clone	8.00E-12
1602.seq	432	gij5854951 gb AW006173.1 AW006173	wz93b08.x1 NCI_CGAP_Brn25 Homo sapiens cDNA clone IMAGE:2566359 3'	0
1603.seq	494	gij5960045 gb AR040682.1 AR040682	Sequence 5 from patent US 5807993	0
1604.seq	453	gij5877428 gb AW023818.1 AW023818	df62e03.y1 Morton Fetal Cochlea Homo sapiens cDNA clone	0
1605.seq	476	gij5922058 gb AC010577.2 AC010577	Drosophila melanogaster chromosome 4 clone BACR22J20 (D1076) RPCI-98	0.037
1606.seq	449	gij5921382 emb AL034552.22 HS824J5	Human DNA sequence from clone 824J5 on chromosome 20q11.23-12,	0.54
1607.seq	426	gij5875963 gb AW026033.1 AW026033	wv06b04.x1 NCI_CGAP_Kid3 Homo sapiens cDNA clone IMAGE:990914 3'	9.00E-12
1608.seq	447	gij5884631 gb AJ1981603.1 AJ1981603	pat.pK060.h7.7 chicken activated T cell cDNA Gallus gallus cDNA	1.00E-41
1609.seq	495	gij5882314 gb AF010192.1 AF010192	Homo sapiens clone RPCI11-141M3, WORKING DRAFT SEQUENCE, 29 unord	0.01
1610.seq	451	gij5917697 db AF170095.1 AF170095	Homo sapiens isolate MCL-BV/128 immunoglobulin heavy chain VDJ	E-142
1611.seq	496	gij5865957 emb AL110349.1 AL110349	DKFZp564K1678_r1 564 (synonym: hfr2) Homo sapiens cDNA clone	E-156
1612.seq	431	gij5927801 emb AL036554.2 AL036554	DKFZp564J2262_r1 564 (synonym: hfr2) Homo sapiens cDNA clone	2.00E-40
1613.seq	298	gij5916455 gb AC010747.1 AC010747	Homo sapiens clone NH0555H09, WORKING DRAFT SEQUENCE, 13 unorder	0.088
1614.seq	397	gij5862546 gb AW013768.1 AW013768	BSBmMF5Z34N4SK Brugia malayi microfilaria cDNA (SAW94L5-BmMF)	2.00E-77
1615.seq	383	gij6006206 gb AC009624.2 AC009624	Homo sapiens chromosome 8 clone 298_E_18 map 8, WORKING DRAFT SEQ	0.12
1616.seq	513	gij5876180 gb AW022650.1 AW022650	df41g11.y1 Morton Fetal Cochlea Homo sapiens cDNA clone	0
1617.seq	366	gij5870989 gb AC010145.1 AC010145	Homo sapiens clone NH0355H10, WORKING DRAFT SEQUENCE, 19 unorder	0.43
1618.seq	504	gij5984498 gb AR058921.1 AR058921	Sequence 2 from patent US 5837838	0
1619.seq	471	gij5876784 gb AW023254.1 AW023254	df52d06.y1 Morton Fetal Cochlea Homo sapiens cDNA clone	0
1620.seq	473	gij5857390 gb AW008612.1 AW008612	ws70h04.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:2502583 3'	E-175
1621.seq	415	gij6007872 gb AC007376.1 AC007376	Homo sapiens chromosome 14 clone BAC 99E15 map 14q24.3, WORKING DR	0
1622.seq	434	gij6006912 gb AC005817.6 AC005817	Mus musculus, complete sequence	0.033
1623.seq	520	gij6006248 gb AC011091.1 AC011091	Homo sapiens clone 1_A_16, WORKING DRAFT SEQUENCE, 12 ordered piece	0.16
1624.seq	467	gij5932571 gb AC010002.2 AC010002	Homo sapiens chromosome 7, WORKING DRAFT SEQUENCE, 55 unorderd p	0.56
1625.seq	485	gij5915183 gb AC007956.2 AC007956	Homo sapiens chromosome 14 clone BAC 3211F8 map 14q24.3, WORKING DR	0.037
1626.seq	458	gij5853733 gb AW004870.1 AW004870	wv02g12.x1 NCI_CGAP_Kid3 Homo sapiens cDNA clone IMAGE:991001 3'	E-105
1627.seq	422	gij5933854 gb AW058215.1 AW058215	wx17c05.x1 NCI_CGAP_Gas4 Homo sapiens cDNA clone IMAGE:2543912 3'	0
1628.seq	356	gij5870350 emb AL031726.7 HS1018K9	Homo sapiens chromosome 1 clone dJ1018K9, WORKING DRAFT SEQUENCE	3.00E-39
1629.seq	449	gij5865405 gb AW016648.1 AW016648	UI-H-B10p-abe-a-01-0-UI.s1 NCI_CGAP_Sub2 Homo sapiens cDNA clone	E-145
1630.seq	492	gij6015480 db AP000563.1 AP000563	Homo sapiens genomic DNA, chromosome 21q22.1, clone:PQ8P9, D21S226-A	0.15
1631.seq	493	gij5918427 emb AL109938.3 HSJ324N14	Homo sapiens chromosome 6 clone dJ324N14, WORKING DRAFT SEQUENCE	0
1632.seq	501	gij6013768 gb AW062383.1 AW062383	RC0-C70037-050899-002-D08 CT0037 Homo sapiens cDNA, mRNA sequen	E-163
1633.seq	460	gij5978971 gb AR054109.1 AR054109	Sequence 1 from patent US 5834441	0
1634.seq	496	gij6007579 gb AF188032.1 AF188032	Homo sapiens chromosome 8 clone BAC 456C06, WORKING DRAFT SEQUEN	0.038
1635.seq	309	gij5951904 gb AR035236.1 AR035236	Sequence 53 from patent US 5871732	E-167
1636.seq	489	gij6007575 gb AF188026.1 AF188026	Homo sapiens chromosome 8 clone BAC 2379L20 map 8q24, WORKING DRA	0.15
1639.seq	497	gij5882380 gb AC010250.1 AC010250	Homo sapiens chromosome 5 clone CIT-HSPC_425H14, WORKING DRAFT	4.00E-45
1640.seq	408	gij5857376 gb AW008598.1 AW008598	ws70f08.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:2502567 3'	0
1641.seq	441	gij5882337 gb AC010172.1 AC010172	Homo sapiens clone RPCI11-551G13, WORKING DRAFT SEQUENCE, 48 unord	0
1642.seq	494	gij6014375 emb AL121766.1 CNS01DSA	Homo sapiens chromosome 14 clone R-412H8, WORKING DRAFT SEQUENC	E-130
1643.seq	481	gij6006108 gb AC011192.1 AC011192	Homo sapiens clone hRPK53_A_1, WORKING DRAFT SEQUENCE, 9 ordered	0
1644.seq	477	gij5875117 gb AW021587.1 AW021587	df25f02.y1 Morton Fetal Cochlea Homo sapiens cDNA clone	0
1645.seq	474	gij5925871 emb AL119972.1 AL119972	DKFZp761E142_r1 761 (synonym: hamy2) Homo sapiens cDNA clone	2.00E-41
1646.seq	484	gij5905466 gb AW044937.1 AW044937	um15f12.y1 Sugano mouse kidney mRka Mus musculus cDNA clone	6.00E-04
1647.seq	478	gij5998689 gb AR067467.1 AR067467	Sequence 16 from patent US 5851764	0
1649.seq	167	gij5902023 ref NM_006902.1 PMX1	Homo sapiens paired mesoderm homeo box 1 (PMX1), mRNA	0.19
1650.seq	477	gij5870342 emb AL079307.2 CNS00M8N	Homo sapiens chromosome 14 clone R-280K24, WORKING DRAFT SEQUENC	0.15
1651.seq	409	gij5928378 gb AL038115.2 AL038115	DKFZp566N191_r1 566 (synonym: hfk2) Homo sapiens cDNA clone	E-106
1652.seq	282	gij5873722 gb AW020192.1 AW020192	df05g05.y1 Morton Fetal Cochlea Homo sapiens cDNA clone	E-137
1653.seq	515	gij6013568 gb AC011373.1 AC011373	Homo sapiens chromosome 5 clone CIT978SKB_10806, WORKING DRAFT	0.04
1654.seq	484	gij5877565 gb AW024035.1 AW024035	df64f02.y1 Morton Fetal Cochlea Homo sapiens cDNA clone	0
1655.seq	494	gij5912651 gb AC008146.2 AC008146	Trypanosoma brucei chromosome V1 clone RPCI93-30P15, WORKING DRAFT	0.6
1656.seq	204	gij4292004 gb AJ445980.1 AJ445980	tj32b05.x1 NCI_CGAP_Pan1 Homo sapiens cDNA clone IMAGE:2143185 3'	E-112
1657.seq	249	gij5919867 emb AL118028.1 AL118028	r9107b56 Bedlington mouse dissected endoderm Mus musculus cDNA	2.00E-39
1658.seq	479	gij5838925 gb AI992020.1 AI992020	ws44b01.x1 NCI_CGAP_Brn25 Homo sapiens cDNA clone IMAGE:2500009 3'	6.00E-47
1659.seq	522	gij5923645 gb AC010865.1 AC010865	Homo sapiens clone RP11-67C2, WORKING DRAFT SEQUENCE, 35 unorder	0.04
1660.seq	457	gij5927696 emb AL036237.2 AL036237	DKFZp564J202_s1 564 (synonym: hfr2) Homo sapiens cDNA clone	0
1661.seq	218	gij5904241 gb AW043712.1 AW043712	wy80d12.x1 Soares_NSF_F8_9W_OT_PA_P_S1 Homo sapiens cDNA clone	E-118
1662.seq	424	gij5906423 gb AW044894.1 AW044894	UI-M-BH1-aku-a-08-0-UI.s1 NIH_BMAP_M_S2 Mus musculus cDNA clone	2.00E-25
1663.seq	495	gij5926613 emb AL120714.1 AL120714	DKFZp762A131_r1 762 (synonym: hmet2) Homo sapiens cDNA clone	2.00E-68
1664.seq	416	gij5862077 gb AW013299.1 AW013299	Sp130f Winter flounder spleen Pleuronectes americanus cDNA clone	0
1667.seq	498	gij5853711 gb AW005007.1 AW005007	wv02e03.x1 NCI_CGAP_Kid3 Homo sapiens cDNA clone IMAGE:990959 3'	E-131
1669.seq	455	gij5932620 gb AC009570.2 AC009570	Homo sapiens chromosome 4 clone 46_J_23 map 4, WORKING DRAFT SEQU	4.00E-91
1671.seq	477	gij5931908 emb AL096829.10 HSJ595K1	Homo sapiens chromosome 1 clone dJ595K12, WORKING DRAFT SEQUENC	E-151
1674.seq	527	gij5923640 gb AC008125.9 AC008125	Homo sapiens 12 BAC RPCI11-25E2 (Roswell Park Cancer Institute Human	0.64
1675.seq				
1676.seq	499	gij6002294 emb Z98200.8 HS111822	Human DNA sequence from clone 111822 on chromosome 6q16-21, complete	0
1677.seq	192	gij5873809 gb AW020279.1 AW020279	df07d06.y1 Morton Fetal Cochlea Homo sapiens cDNA clone	3.00E-53
1678.seq	504	gij5877241 gb AW023711.1 AW023711	df59c02.y1 Morton Fetal Cochlea Homo sapiens cDNA clone	E-103
1679.seq	504	gij5925723 emb AL119824.1 AL119824	DKFZp761H1124_r1 761 (synonym: hamy2) Homo sapiens cDNA clone	0

1680.seq	326	gi 5879594 gb AW026064.1 AW026064	wv06f12.x1 NCI_CGAP_Kid3 Homo sapiens cDNA clone IMAGE:990978 3'	2.00E-98
1681.seq	158	gi 5856875 gb AW008108.1 AW008108	wv49g07.x1 NCI_CGAP_Gas4 Homo sapiens cDNA clone IMAGE:990895 3'	2.00E-44
1682.seq	507	gi 5921610 gb AF035029.1 AF035029	Homo sapiens clone MC58K myosin-reactive immunoglobulin heavy chain	E-123
1684.seq	482	gi 5881357 dbj AP000441.1 AP000441	Homo sapiens chromosome 11 clone B5B24 map 11q12, WORKING DRAFT	0
1685.seq	417	gi 5926703 dbj AP000516.1 AP000516	Homo sapiens genomic DNA, chromosome 6p21.3, HLA Class I region,	0.13
1686.seq	261	gi 5922093 gb AC010858.1 AC010858	Homo sapiens chromosome 15 clone 396_B_8 map 15, LOW-PASS SEQUENC	0.3
1687.seq	506	gi 5931439 gb AC007285.3 AC007285	Homo sapiens clone NH0576P10, complete sequence	0.61
1688.seq	496	gi 5935387 emb AL041391.2 AL041391	DKFZp434B0617_r1 434 (synonym: htes3) Homo sapiens cDNA clone	0
1689.seq	372	gi 4285897 gb AI444995.1 AI444995	t92b01.x1 NCI_CGAP_Co14 Homo sapiens cDNA clone IMAGE:2154505 3'	0
1691.seq	365	gi 5847789 gb AW000873.1 AW000873	wu46f02.x1 Soares_Dieckgraefe_colon_NHCD Homo sapiens cDNA clone	4.00E-63
1692.seq				
1693.seq	467	gi 6013767 gb AW062382.1 AW062382	RC0-CT0037-050899-002-D07 CT0037 Homo sapiens cDNA, mRNA sequenc	0
1694.seq	506	gi 5882617 gb AC010317.1 AC010317	Homo sapiens chromosome 5 clone CIT9785KB_8B14, WORKING DRAFT	0.61
1695.seq	506	gi 5878778 gb AW025248.1 AW025248	wu95d10.x1 NCI_CGAP_Kid3 Homo sapiens cDNA clone IMAGE:990758 3',	0.01
1696.seq	339	gi 6006221 gb AC011107.1 AC011107	Homo sapiens clone 115_L_3, LOW-PASS SEQUENCE SAMPLING	0.1
1697.seq	495	gi 5836192 gb AC009486.2 AC009486	Homo sapiens clone NH0333001, WORKING DRAFT SEQUENCE, 2 unorder	0.038
1698.seq	438	gi 5878255 gb AW024725.1 AW024725	wu91a01.x1 NCI_CGAP_Kid3 Homo sapiens cDNA clone IMAGE:990715 3'	E-137
1699.seq	498	gi 5921614 gb AF035031.1 AF035031	Homo sapiens clone MC57K myosin-reactive immunoglobulin light chain	8.00E-96
1700.seq	386	gi 5880085 gb AW026555.1 AW026555	wv14h10.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:990988 3'	0
1701.seq	438	gi 6006519 emb AL121584.2 H5BA388K	Homo sapiens chromosome 20 clone BA388K2, WORKING DRAFT SEQUENCI	0.53
1702.seq	289	gi 4297834 gb AI446813.1 AI446813	t97f07.x1 NCI_CGAP_Gas4 Homo sapiens cDNA clone IMAGE:2139973 3'	E-145
1703.seq	480	gi 5922074 gb AC008056.4 AC008056	Homo sapiens chromosome 14 clone BAC 242P2 map 14q31, WORKING DRA	0.58
1704.seq	317	gi 6007595 gb AC011216.1 AC011216	Homo sapiens clone 6_F_18, LOW-PASS SEQUENCE SAMPLING	0.094
1705.seq	186	gi 5928576 emb AL110304.2 AL110304	DKFZp564H2078_r1 564 (synonym: hfr2) Homo sapiens cDNA clone	E-101
1706.seq	504	gi 5870305 gb AC010093.1 AC010093	Homo sapiens clone NH0323005, WORKING DRAFT SEQUENCE, 45 unorder	0.15
1707.seq	316	gi 6015390 gb AC009678.2 AC009678	Homo sapiens chromosome 8 clone 56_A_4 map 8, WORKING DRAFT SEQUE	0.024
1709.seq				
1710.seq	507	gi 6015176 gb AC007291.19 AC007291	Drosophila melanogaster chromosome 2 clone BACR02I05 (D571) RPCI-98	6.00E-04
1711.seq				
1712.seq	503	gi 5837998 gb AI991095.1 AI991095	wu38a04.x1 Soares_Dieckgraefe_colon_NHCD Homo sapiens cDNA clone	0
1713.seq	459	gi 5886181 gb AW027425.1 AW027425	wv73f04.x1 Soares_thymus_NHFTt Homo sapiens cDNA clone IMAGE:99086	E-169
1714.seq	375	gi 5857811 gb AW009033.1 AW009033	ws76f02.x1 NCI_CGAP_Co3 Homo sapiens cDNA clone IMAGE:2503899 3'	0
1715.seq	503	gi 6001955 gb AC010205.3 AC010205	Homo sapiens clone RPCI11-285E23, WORKING DRAFT SEQUENCE, 66 un	E-159
1716.seq	500	gi 5883634 gb AI980606.1 AI980606	pat.pk0027.h11.f chicken activated T cell cDNA Gallus gallus cDNA	9.00E-74
1717.seq	409	gi 5928390 emb AL038233.2 AL038233	DKFZp566D202_r1 566 (synonym: hfrd2) Homo sapiens cDNA clone	0
1718.seq	477	gi 5836193 gb AC009958.1 AC009958	Homo sapiens clone NH0316G09, WORKING DRAFT SEQUENCE, 4 unorder	E-136
1719.seq	197	gi 4284541 gb AI445228.1 AI445228	t88a10.x1 NCI_CGAP_Co14 Homo sapiens cDNA clone IMAGE:2154138 3'	E-107
1720.seq	454	gi 5978971 gb AR054109.1 AR054109	Sequence 1 from patent US 5834441	0
1722.seq	499	gi 5874181 gb AW020651.1 AW020651	df12f05.y1 Morton Fetal Cochlea Homo sapiens cDNA clone	0
1723.seq	512	gi 6010197 emb AL121718.1 H5J390M24	Homo sapiens chromosome 6 clone dJ390M24, WORKING DRAFT SEQUENC	0.16
1724.seq	502	gi 6936618 emb AL042879.3 AL042879	DKFZp43411622_r1 434 (synonym: htes3) Homo sapiens cDNA clone	0
1725.seq	403	gi 5867133 gb AW008355.1 AW008355	wv53h08.x1 NCI_CGAP_Gas4 Homo sapiens cDNA clone IMAGE:990898 3',	E-150
1727.seq	545	gi 6014404 emb AL121759.1 H5DJ872K	Homo sapiens chromosome 20 clone dJ872K7, WORKING DRAFT SEQUENC	0.011
1728.seq	363	gi 6015252 gb AC011500.1 AC011500	Homo sapiens chromosome 19 clone CIT9785KB_60E11, WORKING DRAFT S	0.002
1729.seq	510	gi 1901611 gb M73547 HUMPOLLA	Human polyposis locus (DP1 gene) mRNA, complete cds.	0
1729.seq	510	gi 1901611 gb M73547 HUMPOLLA	Human polyposis locus (DP1 gene) mRNA, complete cds.	0
1730.seq	469	gi 1814771 gb M14200 HUMDBI	Human diazepam binding inhibitor (DBI) mRNA, complete cds.	0
1731.seq				
1733.seq	483	gi 4713939 gb AC006518.17 AC006518	Homo sapiens 12p13 BAC RPCI11-144023 (Roswell Park Cancer Institute Hu	0.13
1734.seq	426	gi 3873300 gb AC005829 AC005829	Homo	0
1735.seq	468	gi 2133911 gb AC000394 AC000394	Genomic sequence from Human 9q34, complete sequence [Homo sapiens]	0
1737.seq	419	gi 563340 emb X81371 BTPIGR1	B.taurus mRNA for bovine polymeric immunoglobulin receptor	5.00E-04
1738.seq	139	gi 5174560 ref NM_005930.1 MGEA6	Homo sapiens meningioma expressed antigen 6 (colled-coil)	2.00E-34
1740.seq	469	gi 34137 emb Y07511 HSCOAT	Human mRNA for kidney ornithine aminotransferase (EC 2.6.1.13)	0
1741.seq	462	gi 5924019 emb AL049844.7 H5J468K18	Human DNA sequence from clone 468K18 on chromosome 6q24.1-24.3 Cont	6.00E-93
1742.seq	339	gi 4176442 emb AL022238.1 H5I042K10	Human DNA sequence from clone 1042K10 on chromosome 22q13.1-13.2.	1.00E-84
1743.seq	474	gi 984595 dbj D32058 HUM2OD3	Human gene for 2-oxoglutarate dehydrogenase, exon 3 sequence	2
1744.seq	379	gi 5262657 emb AL080181.1 HSM80070	Homo sapiens mRNA, cDNA DKFZp4340111 (from clone DKFZp4340111)	E-105
1745.seq	469			
1746.seq	478	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-111
1747.seq	492	gi 35219 emb X52104 HSP68A	Human mRNA for p68 protein	0
1748.seq				
1750.seq	488	gi 2335068 gb AC002428 AC002428	Human BAC clone GS039E22 from 5q31, complete sequence [Homo sapiens]	0.51
1751.seq	504	gi 4503270 ref NM_001920.1 DCN	Homo sapiens decorin (DCN) mRNA >gi 181169 gb M14219 HUMCSPG1A	0
1752.seq	461	gi 4469083 emb AL031670.6 H5S681N20	Human DNA sequence from clone 681N20 on chromosome 20p12.1-13 Cont	3.00E-05
1753.seq	490	gi 2343108 gb AF004230 AF004230	Homo sapiens monocyte/macrophage Ig-related receptor MIR-7 (MIR cl-7)	1.00E-44
1754.seq	378	gi 4240226 dbj AB020676.1 AB020676	Homo sapiens mRNA for KIAA0869 protein, partial cds	0.39
1755.seq	508	gi 2961443 gb AC002350 AC002350	Homo sapiens 12q24 PAC RPCI3-424M6 (Roswell Park Cancer Institute Hum	E-144
1757.seq	509	gi 3417301 gb AC005323 AC005323	Homo sapiens chromosome 17, clone hRPK150_K_11, complete sequence	0.54
1758.seq	499	gi 4759085 ref NM_004892.1 SEC22B	Homo sapiens vesicle trafficking protein sec22b (SEC22B) mRNA	1.00E-10
1759.seq	217			
1761.seq	437	gi 550014 gb U14967 H5U14967	Human ribosomal protein L21 mRNA, complete cds.	0
1762.seq	387	gi 4827303 gb AC005060.2 AC005060	Homo sapiens clone RG086D03, complete sequence	0.4
1763.seq	220	gi 3947674 emb AL031311 H5I05D16	Human DNA sequence from clone 105D16 on chromosome Xp11.3-11.4 Cont	0.056
1764.seq	452	gi 4753272 gb AC005534.2 AC005534	Homo sapiens PAC clone DJ0982E09 from 7q35-q36, complete sequence	0
1765.seq	249	gi 644480 dbj D38112.1 HUMMTA	Homo sapiens mitochondrial DNA, complete sequence	E-136
1766.seq	319	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-127
1767.seq	485	gi 241477 gb S74678 S74678	heterogeneous nuclear ribonucleoprotein complex K [human, mRNA,	0
1768.seq	430	gi 5729773 ref NM_006660.1 CLPX	Homo	0
1769.seq	367	gi 288099 emb X55733 HSINTFA4B	H.sapiens initiation factor 4B cDNA	0
1770.seq	136			
1771.seq	478	gi 190978 gb M18963 HUMREGA	Human islet of Langerhans regenerating protein (reg) mRNA, complete	0
1772.seq	446	gi 5262709 emb AL080218.1 HSM800746	Homo sapiens mRNA; cDNA DKFZp586N1323 (from clone DKFZp586N1323)	E-124
1773.seq	454	gi 4506902 ref NM_003769.1 SFRS9	Homo sapiens splicing factor, arginine/serine-rich 9 (SFRS9) mRNA	E-164
1774.seq	448			
1775.seq	168	gi 2340098 gb AC002480 AC002480	Human BAC clone GS055K18 from 7p15-p21, complete sequence [Homo sapi	0.01
1776.seq	445	gi 29720 emb X04076 HSCATR	Human kidney mRNA for catalase	0
1777.seq	306	gi 184250 gb L08048 HUMHMG1C	Human non-histone chromosomal protein (HMG-1) retropseudogene.	E-111
1778.seq	413	gi 3483796 gb AF086451 HJMZD8E12	Homo sapiens full length Insert cDNA clone ZD8E12	E-170
1779.seq	459	gi 2887412 dbj AB007879 AB007879	Homo sapiens KIAA0419 mRNA, complete cds	E-155
1780.seq	475	gi 395086 emb X74070 H5BTF3	H.sapiens mRNA for transcription factor BTF 3	0
1781.seq	464	gi 2780782 dbj AB002806 AB002806	Homo sapiens OS-9 mRNA, complete cds	0
1782.seq	149	gi 5836161 gb AC008283.3 AC008283	Homo sapiens clone NH0575G01, complete sequence	0.56
1783.seq	238	gi 337929 gb M22146 HUMSCAR	Human scar protein mRNA, complete cds.	E-128
1784.seq	519	gi 331140 emb V00557 H5IGK1	C-terminal part of human kappa-immunoglobulin gene coding for amino	0
1796.seq	345	gi 5708490 gb AC004985.2 AC004985	Homo sapiens clone DJ1165K10, complete sequence	1.4
1797.seq				
1798.seq	185	gi 2313485 gb AE000555.1 AE000555	Helicobacter pylori 26695 section 33 of 134 of the complete genome	0.046
1800.seq	416	gi 2076716 gb U30897 H5U30897	Human glutathione S-transferase (P1b) mRNA, complete cds	0
1802.seq	482	gi 4826885 ref NM_005016.1 PCBP2	Homo sapiens poly(rC)-binding protein 2 (PCBP2) mRNA	0
1803.seq	486	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-130
1804.seq	531	gi 3766124 gb AC005747 AC005747	Homo sapiens chromosome 17, clone hRPK150_K_15, complete sequence	0.56
1805.seq	583	gi 184847 gb M63438 HUMIGGK	Human Ig rearranged gamma chain mRNA, V-J-C region and complete cds.	0
1807.seq	496	gi 5032026 ref NM_005610.1 RBBP4	Homo sapiens retinoblastoma-binding protein 4 (RBBP4) mRNA	0
1809.seq	461	gi 1262342 emb X93334 H5MITG	H.sapiens mitochondrial DNA, complete genome	0
1810.seq	460	gi 4506686 ref NM_001018.1 RPS15	Homo sapiens ribosomal protein S15 (RPS15) mRNA	0
1811.seq	481	gi 5649181 gb AC008041.5 AC008041	Homo sapiens 3a26.2-27 BAC RPCI11-469J4 (Roswell Park Cancer Institute	8.00E-06
1812.seq	334	gi 5123959 gb AC007437.16 AC007437	Homo sapiens 12q22 BAC RPCI11-541G9 (Roswell Park Cancer Institute Hum	1.4

1813.seq	463	g 190978 gb U18963 HUMREGA	Human islet of Langerhans regenerating protein (reg) mRNA, complete	0
1816.seq	478	g 4502168 ref NM_003905.1 APPBP1	Homo sapiens amyloid beta precursor protein-binding protein 1, 59kD	0
1817.seq	537	g 181966 gb U29549 HUMEF1AB	Human elongation factor 1-alpha (EF1A) mRNA, partial cds.	0
1818.seq	491	g 331912 gb AC003042 AC003042	Homo sapiens chromosome 17, clone HCIT75G16, complete sequence [Homo	E-117
1819.seq	465	g 5419839 emb AL050327.6 HSJ907D15	Human DNA sequence from clone 907D15 on chromosome 20q13.2-13.33	1.9
1820.seq	439	g 4505764 ref NM_002633.1 PGM1	Homo sapiens phosphoglucomutase 1 (PGM1) mRNA	0
1821.seq	353	g 644480 db J038112.1 HUMMTA	Homo sapiens mitochondrial DNA, complete sequence	3.00E-91
1822.seq				
1823.seq	503	g 432358 db D32660 HUMRSP	Human mRNA for ribosomal protein, complete cds	0
1824.seq	503	g 386007 gb AF097021 AF097021	Homo sapiens GW112 protein (GW112) mRNA, complete cds	E-158
1826.seq	586	g 5042474 gb AC007790.1 AC007790	Homo sapiens clone UWGC:djs310all from 7q31, complete sequence	0.62
1827.seq	561	g 555853 gb U13369 HSU13369	Human ribosomal DNA complete repeating unit	2.3
1828.seq	618	g 5926698 db JAP000511.1 JAP000511	Homo sapiens genomic DNA, chromosome 6p21.3, HLA Class I region,	0.85
1831.seq	595	g 4506602 ref NM_002948.1 RPL15	Homo sapiens ribosomal protein L15 (RPL15) mRNA	0
1832.seq	664	g 4589274 gb AF135024.1 AF135024	Homo sapiens kallikrein-like protein 4 KLK-L4 gene, partial cds.	0.71
1833.seq	727	g 583538 ref NC_01807.1	Human mitochondrion, complete genome >g 1944628 gb J01415 HUMMTCG	0
1835.seq	707	g 5360672 db AB022653.1 AB022653	Homo sapiens mRNA for anti-Entamoeba histolytica immunoglobulin	0
1836.seq	879	g 34805 emb X03299 HSMTLK4	Human nuclear mitochondrial-DNA-like sequence (clone Lm H9)	2.00E-76
1837.seq	859	g 4972129 emb AL048832.2 CNS0000G	Human chromosome 14 DNA sequence *** IN PROGRESS *** BAC R-63D17	3.6
1838.seq	858	g 703088 gb M27487 HUMMHDP4	Homo sapiens MHC class II DPw3-alpha-1 chain mRNA, complete cds.	0
1840.seq	640	g 1478196 emb X97321 HSHLACW07	H.sapiens mRNA for human leukocyte antigen C alpha chain	0
1841.seq	290	g 189265 gb M373791 HUMNOVGENE	Human novel gene mRNA, complete cds.	E-158
1844.seq	499	g 5514658 emb AL060315.8 HS1173K17	Human DNA sequence from clone 1173K17 on chromosome 20. Contains part	0.13
1846.seq	568	g 515771 emb X80437 MMTEG27	M.musculus tex27 mRNA	8.00E-99
1847.seq	484	g 4506682 ref NM_001016.1 RPS12	Homo sapiens ribosomal protein S12 (RPS12) mRNA	0
1849.seq	380	g 4886450 emb AL050273.1 HSM80013C	Homo sapiens mRNA; cDNA DKFZp566B023 (from clone DKFZp566B023)	1.00E-65
1853.seq	621	g 4165479 emb AL031682 HS873P14	Human DNA sequence from clone 873P14 on chromosome 20p12 Contains S	0.66
1854.seq	572	g 4885691 gb AC007055.3 AC007055	Homo sapiens chromosome 14 clone BAC 201F1 map 14q24.3, complete	0.15
1855.seq	630	g 5016088 ref NM_001101.2 ACTB	Homo sapiens actin, beta (ACTB) mRNA	0
1856.seq	600	g 4506668 ref NM_001003.1 RPLP1	Homo sapiens ribosomal protein, large, P1 (RPLP1) mRNA	0
1858.seq	301	g 28333 emb X15804 HSACTAR	Human mRNA for alpha-actinin	3.00E-32
1859.seq	627	g 1478196 emb X97321 HSHLACW07	H.sapiens mRNA for human leukocyte antigen C alpha chain	0
1860.seq	346	g 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	E-110
1861.seq	579	g 4503470 ref NM_001402.1 EEF1A1	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
1863.seq	360	g 3702356 gb AC005669 AC005669	Homo sapiens chromosome 3, clone hRPK.165_1_16, complete sequence [Hom	0.37
1864.seq	342	g 4240565 gb AL234622 DUS26N18	Homo sapiens BAC526N18 neurexin III gene, partial cds	E-147
1865.seq	636	g 1028892 emb Z57661 HS189H1R	H.sapiens CoG Island DNA genomic MseI fragment, clone 189h1,	4.00E-76
1866.seq	751	g 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-138
1867.seq	629	g 4508110 gb AC005076.2 AC005076	Homo sapiens BAC clone RG227L24 from 7q21.1-q21.2, complete sequence	0.67
1868.seq	536	g 5523886 gb AC004913.2 AC004913	Homo sapiens clone DJ0876A24, complete sequence	5.00E-63
1869.seq				
1870.seq	730	g 4503704 ref NM_002009.1 FGF7	Homo sapiens fibroblast growth factor 7 (keratinocyte growth factor)	0
1871.seq	731	g 3367616 emb AL022722 HS354N19	Human DNA sequence from clone 354N19 on chromosome 6q22. Contains the	E-155
1872.seq	848	g 4507512 ref NM_000362.1 TIMP3	Homo sapiens tissue inhibitor of metalloproteinase 3 (Sorsby fundus	0
1873.seq	298	g 4506114 ref NM_000312.1 PROC	Homo sapiens protein C (inactivator of coagulation factors Va and	1.00E-06
1874.seq	702	g 34503020 ref NM_001876.1 CPT1A	Homo sapiens carnitine palmitoyltransferase 1, liver (CPT1A)	0.75
1876.seq	433	g 5524737 gb AF111169.2 AF111169	Homo sapiens chromosome 14 BAC containing gene for KIA00759 and other	1.8
1877.seq	876	g 5508502 emb Z79224 HSPA26C5	H.sapiens flow-sorted chromosome 6 TaqI fragment, SC6A26C5	0.015
1878.seq	511	g 190978 gb U18963 HUMREGA	Human islet of Langerhans regenerating protein (reg) mRNA, complete	0
1879.seq	125	g 4512277 db AB019438.1 AB019438	Homo sapiens DNA for immunoglobulin heavy-chain variable region, complete	0.12
1880.seq	569	g 4504913 ref NM_002273.1 KRT8	Homo sapiens keratin 8 (KRT8) mRNA >g 4004515 emb X74929 HSKRT8	0
1881.seq	648	g 184918 gb M60192 HUMHGHA2A	Human germline IgA2 chain C region.	E-169
1882.seq	650	g 5468517 gb AF111168.2 AF111168	Homo sapiens serine palmitoyl transferase, subunit II gene, complete cds;	0.17
1883.seq	867	g 4506804 ref NM_002580.1 PAP1	Homo sapiens pancreatitis-associated protein (PAP) mRNA	0
1885.seq	260	g 384422 emb X69150.1 HSRPS18	Homo sapiens mRNA for ribosomal protein S18	E-112
1886.seq	683	g 3849820 gb AC005837 AC005837	Homo sapiens chromosome 17, clone hRPK.318_A_15, complete sequence	0
1887.seq	693	g 187779 gb M24096 HUMMHCAA	Human MHC class I HLA-C-alpha-2 chain mRNA, partial cds, clone 9.	0
1889.seq	708	g 3580678 db AB022656.1 AB022656	Homo sapiens mRNA for anti-Entamoeba histolytica immunoglobulin	0
1891.seq	437	g 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	E-109
1891.seq	279	g 4185269 emb J225782 HSSYBL16	Homo sapiens SYBL1 gene, exons 6-8.	0.28
1892.seq	673	g 482908 gb L15533 HUMFAP	Homo sapiens pancreatitis-associated protein (PAP) gene, complete cds.	0
1894.seq	871	g 4885599 ref NM_005410.1 SEPP1	Homo sapiens selenoprotein P, plasma, 1 (SEPP1) mRNA	0
1895.seq	532	g 4502284 ref NM_001681.1 ATP2A2	Homo sapiens ATPase, Ca+> transporting, cardiac muscle, slow twitch 2	0
1896.seq	637	g 3319976 emb AL022157 HS323P24	Human DNA sequence from clone 323P24 on chromosome Xp11.21-11.23	E-175
1897.seq	485	g 3090700 db AC004622 AC004622	Homo sapiens chromosome 5, PAC clone 170m10 (LBNL.H89), complete	0
1899.seq	656	g 5016092 ref NM_000108.2 DLD	Homo sapiens dihydrolipamide dehydrogenase precursor (E3 component	0
1900.seq	471	g 2133911 gb AC000394 AC000394	Genomic sequence from Human 9q34, complete sequence [Homo sapiens]	E-174
1901.seq	410	g 5931501 db JAP000523.1 JAP000523	Homo sapiens genomic DNA, chromosome 22q11.2, Cat Eye Syndrom region	0.43
1902.seq	661	g 23690 emb X56932 HS23KDHPB	H.sapiens mRNA for 23 kD highly basic protein	0
1903.seq	536	g 895839 emb X79882 HSLSRPGENE	H.sapiens lrp mRNA >g 3712923 emb A56918.1 A56918 Sequence 1 from	0
1906.seq	640	g 5419825 emb AL049797.8 HSJ661G12	Human DNA sequence from clone 661G12 on chromosome 1q25.3-31.2.	0
1907.seq	611	g 1669548 db DB4108 DB4108	Homo sapiens mRNA for RBP-MS/type 2, complete cds	0
1908.seq	298	g 5901895 ref NM_006886.1 ATP5E	Homo sapiens ATP synthase, H+ transporting, mitochondrial F1	E-141
1909.seq	606	g 4507676 ref NM_003299.1 TRA1	Homo sapiens tumor rejection antigen (gp96) 1 (TRA1) mRNA	E-135
1910.seq	450	g 4503304 ref NM_001926.1 DEFA6	Homo sapiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	0
1911.seq	637	g 441402 emb X72467 HSIGKLV45	H.sapiens mRNA for rearranged Ig kappa light chain variable region	0
1912.seq	605	g 4753268 gb AC006352.3 AC006352	Homo sapiens clone DJ1044J09, complete sequence	E-154
1914.seq	725	g 5442035 gb AF109126.1 AF109126	Homo sapiens stromal cell-derived receptor-1 beta mRNA, complete cds	E-153
1915.seq	768	g 4204701 gb AC006088 AC006088	Homo sapiens 12q24.1 PAC RPC3-305i20 (Roswell Park Cancer Institute	0
1916.seq	351	g 4557228 ref NM_001087.1 AAAMP	Homo sapiens angio-associated, migratory cell protein (AAAMP) mRNA	7.00E-95
1917.seq	844	g 4503306 ref NM_000126.1 ETFA	Homo sapiens electron-transfer-flavoprotein, alpha polypeptide	0
1918.seq	476	g 3882292 db AB018329.1 AB018329	Homo sapiens mRNA for KIAA0786 protein, partial cds	E-127
1919.seq	866	g 4972121 emb AL049776.2 CNS00001	Human chromosome 14 DNA sequence *** IN PROGRESS *** BAC R-85K15	5.00E-05
1920.seq	311	g 4757755 ref NM_004039.1 ANXA2	Homo sapiens annexin A2 (ANXA2), mRNA >g 219909 db D00017 HUMLIC	E-166
1921.seq	295	g 3688100 gb AC005702 AC005702	Homo sapiens chromosome 17, clone hRPK.178_C_3, complete sequence [Hor	0.076
1922.seq	856	g 2961443 gb AC002350 AC002350	Homo sapiens 12q24 PAC RPC3-424M6 (Roswell Park Cancer Institute Hum	0.92
1924.seq	535	g 5540113 gb AC008080.1 AC008080	Homo sapiens clone UWGC:djs94 from 7p14-15, complete sequence	0.56
1926.seq	499	g 4758043 ref NM_004074.1 COX8	Homo sapiens cytochrome c oxidase subunit VIII (COX8), nuclear gene	0
1927.seq	371	g 4557324 ref NM_000041.1 APOE	Homo sapiens apolipoprotein E (APOE) mRNA	E-104
1928.seq	589	g 5050957 emb AL031709.1 HS316G12	Human DNA sequence from clone 316G12 on chromosome 16. Contains the	0
1929.seq	574	g 5725511 gb AF072097.1 AF072097	Homo sapiens beta-2 microglobulin gene, complete cds	0
1930.seq	655	g 5306222 gb AC007541.9 AC007541	Homo sapiens 12q24.1 BAC RPC111-412D9 (Roswell Park Cancer Institute	0
1932.seq	597	g 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-174
1933.seq	597	g 5565654 gb AF077188.1 AF077188	Homo sapiens cullin 4A (CUL4A) mRNA, complete cds	0
1934.seq	844	g 4504482 ref NM_000194.1 HPRT1	Homo sapiens hypoxanthine phosphoribosyltransferase 1 (Lesch-Nyhan	0
1936.seq	581	g 583538 ref NC_01807.1	Human mitochondrion, complete genome >g 1944628 gb J01415 HUMMTCG	0
1937.seq	781	g 5826298 emb AL111679.1 CNS019IF	Botrytis cinerea strain T4 cDNA library under conditions of	3.3
1938.seq	589	g 4630756 gb AC006544.19 AC006544	, complete sequence [Homo sapiens]	0.038
1939.seq	293	g 4503304 ref NM_001926.1 DEFA6	Homo sapiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	3.00E-72
1940.seq	167			
1941.seq	691	g 3043579 db AB011100 AB011100	Homo sapiens mRNA for KIAA0528 protein, complete cds	0
1944.seq	185	g 6005861 ref NM_007208.1 RPLM3	Homo sapiens ribosomal protein, mitochondrial, L3 (RPLM3), mRNA	7.00E-32
1945.seq	272	g 4557520 ref NM_000398.2 DIA1	Homo sapiens diaphorase (NADH) (cytochrome b-5 reductase) (DIA1),	3.00E-35
1946.seq	624	g 4314330 gb AC005874 AC005874	Homo sapiens chromosome 10 clone CIT98TK3-1175G20 map 10q25.2-10q25	0.55
1948.seq	606	g 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-173
1949.seq	240	g 4314424 gb AC006157 AC006157	Homo sapiens BAC clone NH0414C23 from Y, complete sequence [Homo	0.001
1950.seq	834	g 5679750 emb AL023807.6 HS525L6	Human DNA sequence from clone 525L6 on chromosome 6p22.3-23, complet	0.014
1951.seq	95	g 1399085 gb U43701 HSU43701	Human ri	

1952.seq	141	gi 5672603 cb AB023048.1 AB023048	Homo sapiens genomic DNA, chromosome 6p21.3, HLA class I region,	5.00E-72
1953.seq	198	gi 3366582 gb AC005324 AC005324	Homo s	0
1955.seq	536	gi 4503796 ref NM_00146.1 FTL	Homo sapiens ferritin, light polypeptide (FTL) mRNA	0
1956.seq	622	gi 35456 emb X54629 HSPHL1	Human pHL-1 gene is a c-myc oncogene containing coxIII sequence	0
1957.seq	437	gi 4506688 ref NM_001019.1 RPS15A	Homo sapiens ribosomal protein S15a (RPS15A) mRNA	0
1958.seq	761	gi 5901975 ref NM_007069.1 HREV107	Homo sap	0
1959.seq				
1960.seq	126	gi 1037352 emb Z64530 HS133H3R	H.sapiens CpG island DNA genomic MseI fragment, clone 133h3,	3.00E-48
1962.seq	548	gi 4503518 ref NM_003754.1 EIF3S5	Homo sapiens eukaryotic translation initiation factor 3, subunit 5	0
1963.seq	647	gi 4887253 gb AC006559.6 AC006559	Homo sapiens 12p12 BAC RPC111-501E24 (Roswell Park Cancer Institute Hu	0
1965.seq	346	gi 185363 gb M87790 HUMIGHEPAL	Human (hybridoma H210) anti-hepatitis A immunoglobulin lambda chain	E-134
1966.seq	383	gi 6005809 ref NM_007181.1 MAP4K11	Homo sapiens mitogen-activated protein kinase kinase kinase 1	E-107
1968.seq	460	gi 3702283 gb AC005788 AC005788	Homo sapiens chromosome 19, cosmid R26652, complete sequence [Homo	0.12
1969.seq	193	gi 4503698 ref NM_002006.1 FGF2	Homo sapiens fibroblast growth factor 2 (basic) (FGF2) mRNA	3.00E-40
1970.seq	460	gi 4982556 gb AC005921.3 AC005921	Homo sapiens chromosome 17, clone hRPK.294_J_22, complete sequence	E-119
1971.seq	547	gi 4455014 gb AF113131 AF113131	Homo sapiens host cell factor homolog LCP mRNA, complete cds	0
1972.seq	669	gi 886277 gb L40823 HUMDNL1A	Homo sapiens muscle-specific DNase-like (DNL1L) gene, complete cds.	0.09
1973.seq	386	gi 184743 gb J00220 HUMIGCC8	Homo sapiens immunoglobulin alpha-1 heavy chain constant region	0
1974.seq	785	gi 2529723 gb AF001893 BETA2	Human MEN1 region clone epsilon/beta mRNA, 3' fragment	0
1975.seq	655	gi 4884378 emb AL050162.1 HSM800466	Homo sapiens mRNA, cDNA DKFZp586B2022 (from clone DKFZp586B2022)	0
1977.seq	359	gi 3483368 gb AF086023 HUMYW23E08	Homo sapiens full length insert cDNA clone YW23E08	E-114
1978.seq	559	gi 4321975 gb AF068235 AF068235	Homo sapiens barrier-to-autointegration factor mRNA, complete cds	0
1979.seq	569	gi 4502804 ref NM_001275.1 CHGA	Homo sapiens chromogranin A (parathyroid secretory protein 1)	E-121
1980.seq	632	gi 4191318 gb AF044670 AF044670	Homo s	0
1981.seq	644	gi 4826515 emb AL035899.4 HS73H22	Human DNA sequence from clone 73H22 on chromosome 6q23, complete	0.011
1982.seq	625	gi 182976 gb M33197 HUMGAPDH	Human glyceraldehyde-3-phosphate dehydrogenase (GAPDH) mRNA,	0
1983.seq	866	gi 187507 gb M27937 HUMMEA	Human male-enhanced antigen mRNA (Mea), complete cds.	0
1984.seq	589	gi 4758243 ref NM_004105.1 EFEMP1	Homo sapiens EGF-containing fibulin-like extracellular matrix	0
1985.seq	623	gi 2224556 cb AB002306 AB002306	Human mRNA for KIAA0308 gene, partial cds.	0
1986.seq	445	gi 5668665 emb AL109618.1 HSJ775C13	Human DNA sequence from clone 775C13 on chromosome 20p12.1-13, comp	0.47
1988.seq	626	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-174
1989.seq	852	gi 4508148 gb AC004876.2 AC004876	Homo sapiens PAC clone DJ0747G18 from 7q22-31.1, complete sequence	9.00E-69
1990.seq	650	gi 4507358 ref NM_003186.1 TAGLN	Homo sapiens transgelin (TAGLN) mRNA >gi 177174 gb M95787 HUM22SM	0
1991.seq	393	gi 4506622 ref NM_000988.1 RPL27	Homo sapiens ribosomal protein L27 (RPL27) mRNA	0
1992.seq	582	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-166
1993.seq	599	gi 3954892 emb AJ010446 HSA010446	Homo sapiens mRNA for immunoglobulin kappa light chain,anti-RhD,	0
1994.seq	418	gi 2326193 gb AF000137 BTAF000137	Bos taurus connective tissue growth factor precursor (CTGF) mRNA,	0.44
1995.seq	682	gi 531825 gb U12022 HSCALMG2	Human calmodulin (CALM1) gene, exons 2,3,4,5 and 6, and complete cds	E-172
1996.seq	562	gi 4507194 ref NM_003128.1 SPTBN1	Homo sapiens spectrin, beta, non-erythrocytic 1 (SPTBN1) mRNA	0
1997.seq	419	gi 264772 gb S54005 S54005	thymosin beta-10 [human, metastatic melanoma cell line, mRNA, 453	0
1998.seq	549	gi 4884321 emb AL050086.1 HSM800414	Homo sapiens mRNA; cDNA DKFZp434M211 (from clone DKFZp434M211)	0
1998.seq	526	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-136
1999.seq	552	gi 285962 cb D14696 HUMORF13	Human mRNA for KIAA0108 gene, complete cds	0
2000.seq	861	gi 4557808 ref NM_000274.1 OAT	Homo sapiens ornithine aminotransferase (OAT), nuclear gene encoding	0
2001.seq	554	gi 5835387 ref NC_001807.1	Human mitochondrion, complete genome >gi 1944628 gb J01415 HUMMITCG	0
2002.seq	407	gi 4838516 gb AF132047.1 AF132047	Homo sapiens foocen-m mRNA, complete cds	0
2003.seq	706	gi 4557832 ref NM_000282.1 PCCA	Homo sapiens Propionyl-Coenzyme A carboxylase, alpha polypeptide	0
2004.seq	423			
2005.seq	613	gi 3004572 gb AC004520 AC004520	Homo sapiens BAC clone RG119C02 from 7p15, complete sequence [Homo	0
2006.seq	335	gi 361333 emb Z66699 HSRPL37A	H.sapiens mRNA for ribosomal protein L37a	0
2007.seq	587	gi 401844 gb L05093 HUMRIBPROD	Homo sapiens ribosomal protein L18a mRNA, complete cds	0
2008.seq	710	gi 339553 gb M60556 HUMTGFB3B	Human transforming growth factor beta-3 gene, 5' end.	0.76
2010.seq	684	gi 4503470 ref NM_001402.1 EEF1A1	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
2011.seq	556	gi 984142 emb X89401 HSL21PRDT	H.sapiens mRNA for large subunit of ribosomal protein L21	0
2012.seq	602	gi 2588608 gb AC003093 AC003093	Human BAC clone RG385F02 from 7p15, complete sequence [Homo sapiens]	0
2013.seq	628	gi 4506862 ref NM_003001.1 SDHC	Homo sapiens succinate dehydrogenase complex, subunit C, integral	0
2015.seq	623	gi 184847 gb M63438 HUMIGGK	Human Ig rearranged gamma chain mRNA, V-J-C region and complete cds.	0
2016.seq	845	gi 181569 gb M80536 HUMDIPEPIV	H.sapiens dipeptidyl peptidase IV (DPP4) mRNA, complete cds.	0.9
2017.seq	580	gi 189920 gb M11968 HUMPGKA11	Human X-linked phosphoglycerate kinase gene, exon 11.	0
2019.seq	450	gi 3097871 gb AC003663 AC003663	Homo sapiens chromosome 17, clone HCIT87G17, complete sequence [Homo	0
2020.seq	393	gi 4689133 gb AF077043.1 AF077043	Homo sapiens 60S ribosomal protein L36 mRNA, complete cds	0
2022.seq	507	gi 4506880 ref NM_001015.1 RPS11	Homo sapiens ribosomal protein S11 (RPS11) mRNA	0
2023.seq	610	gi 23690 emb X56932 HS23KDHPB	H.sapiens mRNA for 23 kD highly basic protein	0
2024.seq	537	gi 1668818 gb L29074.1 HUMFMR1S	Homo sapiens fragile X mental retardation syndrome protein (FMR1) gene,	0
2026.seq	657	gi 3413288 emb AL021453 HLSB21D11	Human DNA sequence from clone 821D11 on chromosome 22q12.3-13.1.	0.18
2028.seq	213	gi 2429080 cb D87675 D87675	Homo sapiens DNA for amyloid precursor protein, complete cds	0.21
2029.seq	701	gi 13683 emb V00710 MIT1HS	Human mitochondrial genes for several tRNAs (Phe, Val, Leu) and 12S	0
2030.seq	669	gi 5902065 ref NM_007042.1 RPP14	Homo sapiens ribonuclease P (14kD) (RPP14), mRNA	0
2031.seq	421	gi 187701 gb M24194 HUMHBA123	Human MHC protein homologous to chicken B complex protein mRMO,	E-177
2032.seq	611	gi 4505258 ref NM_002444.1 MSN	Homo sapiens moesin (MSN) mRNA >gi 188625 gb M69066 HUMMOESIN Hum	0
2033.seq	495	gi 4885454 ref NM_005475.1 LNK	Homo sapiens lymphocyte adaptor protein (LNK) mRNA	0
2035.seq	245	gi 4506006 ref NM_002710.1 PPP1CC	Homo sapiens protein phosphatase 1, catalytic subunit, gamma	1.00E-70
2036.seq	668	gi 4914350 gb AC006581.16 AC006581	Homo sapiens 12p21 BAC RPC111-259O18 (Roswell Park Cancer Institute	0.18
2037.seq	461	gi 4557770 ref NM_000256.1 MYBPC3	Homo sapiens protein C, cardiac (MYBPC3) mRNA	0
2039.seq	568	gi 3126777 gb AC004590 AC004590	Homo sapiens chromosome 17, clone hCIT 22_K_21, complete sequence [Hom	0.01
2041.seq	371	gi 3201899 gb AF067420 AF067420	Homo sapiens SNC73 protein (SNC73) mRNA, complete cds	5.00E-62
2042.seq	533	gi 4759261 ref NM_004275.1 TRFP1	Homo sapiens TRF-proximal protein (TRFP) mRNA	0
2043.seq	672	gi 5924006 emb AL024498.12 HS417M14	Human DNA sequence from clone 417M14 on chromosome 6p23-25.1, comple	0
2044.seq	618	gi 5689566 cb AB029038.1 AB029038	Homo sapiens mRNA for KIAA1115 protein, complete cds	E-137
2045.seq	606	gi 3059064 emb Z97181 HS95C20	Homo sapiens DNA sequence from PAC 95C20 on chromosome Xp11.3-11.4.	2.00E-22
2046.seq	421			
2047.seq	624	gi 6002954 gb AF171877.1 AF171877	Homo sapiens cleavage and polyadenylation specificity factor 73 kDa.	1.00E-97
2051.seq	472	gi 5835387 ref NC_001807.1	Human mitochondrion, complete genome >gi 1944628 gb J01415 HUMMITCG	0
2052.seq	523	gi 5931439 gb AC007285.3 AC007285	Homo sapiens clone NH0576P10, complete sequence	0.55
2053.seq	597	gi 338738 gb M12423 HUMTCAXE	Human T-cell receptor active alpha-chain mRNA, complete cds, from	0
2054.seq	232	gi 4507358 ref NM_003186.1 TAGLN	Homo sapiens transgelin (TAGLN) mRNA >gi 177174 gb M95787 HUM22SM	1.00E-36
2055.seq	455	gi 4456457 emb Z99716.4 HS250D10	Human DNA sequence from clone 250D10 on chromosome 22. Contains the	0
2056.seq	641	gi 285970 cb D13510.1 HUMPAPC	Homo sapiens mRNA for PAP homologous protein, complete cds	0
2057.seq	214	gi 2343108 gb AF004230 AF004230	Homo sapiens monocyte/macrophage Ig-related receptor MIR-7 (MIR cl-7)	E-109
2058.seq	592	gi 4557316 ref NM_001157.1 ANX11	Homo sapiens annexin XI (56kD autoantigen) (ANX11) mRNA	0
2061.seq	359	gi 4204244 gb AC004801 AC004801	Homo sapiens 12q13.1 PAC RPC11-228P16 (Roswell Park Cancer Institute	9.00E-73
2062.seq	881	gi 4504438 ref NM_002134.1 HMOX2	Homo sapiens heme oxygenase (decycling) 2 (HMOX2), mRNA	E-111
2064.seq	370	gi 337829 gb M22146 HUMSCAR	Human scar protein mRNA, complete cds.	0
2065.seq	276	gi 4926888 gb AC006387.3 AC006387	Homo sapiens clone RG360L10, complete sequence	0.28
2066.seq	597	gi 5668757 gb AC008009.4 AC008009	Homo sapiens 3q26-27 BAC RPC111-436A20 (Roswell Park Cancer Institut	E-106
2067.seq	428	gi 1946346 gb UB8629 HSUB8629	Human RNA polymerase II elongation factor ELL2, complete cds	3.00E-17
2068.seq	686	gi 4504184 ref NM_000853.1 GSTT1	Homo sapiens glutathione S-transferase theta 1 (GSTT1) mRNA	0
2069.seq	562	gi 4240188 cb AB020657.1 AB020657	Homo sapiens mRNA for KIAA0850 protein, complete cds	0
2070.seq	340	gi 3540172 gb AC005606 AC005606	Homo sapiens chromosome 16, P1 clone 109-8C (LANL), complete sequence	E-141
2071.seq	557	gi 5103019 cb AP000356.1 AP000356	Homo sapiens genomic DNA, chromosome 22q11.2, clone KB1995A5	2.3
2072.seq	603	gi 4504918 ref NM_002273.1 KRT8	Homo sapiens keratin 8 (KRT8) mRNA >gi 400415 emb X74929 HSKRT8	0
2073.seq	516	gi 4835608 cb AP000239.1 AP000239	Homo sapiens genomic DNA, chromosome 21q22.1, D21S226-AML region,	E-142
2075.seq	654	gi 633070 gb D38524 HUM5N	Human mRNA for 5'-nucleotidase	0
2077.seq	609	gi 2809270 gb AC002349 AC002349	Homo sapiens Xp22 PAC RPC11-167A22 (from Roswell Park Cancer Center)	2.5
2078.seq	649	gi 337186 gb M25171 HUMMTC DK	Human mitochondrion cytochrome oxidase subunit I (COI), Asp-tRNA	0
2079.seq	85	gi 3043675 cb AB011148 AB011148	Homo sapiens mRNA for KIAA0576 protein, partial cds	1.00E-18
2080.seq	887	gi 3954884 emb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	0
2081.seq	472	gi 4455493 emb AL031316.2 HS28010	Human DNA sequence from clone 28010 on chromosome 1q32.2-41 Contains	2

2082.seq	328	gi 577314 dbj D42084 HUMKIAAO	Human mRNA for KIAA0094 gene, partial cds	E-178
2083.seq	673	gi 4757825 ref NM_004048.1 B2M	Homo sapiens beta-2-microglobulin (B2M) mRNA	0
2084.seq	693	gi 546830 gb S71022 S71022	neoplasm-related C140 product [human, thyroid carcinoma cells,	0
2085.seq	405	gi 285970 dbj D13510.1 HUMPAPC	Homo sapiens mRNA for PAP homologous protein, complete cds	0
2086.seq	723	gi 5901969 ref NM_006895.1 HNM1	Homo sapiens histamine N-methyltransferase (HNMT), mRNA	0
2087.seq	284	gi 5103045 dbj AB028624.1 AB028624	Homo sapiens mRNA for mitochondrial ATP Synthase subunit e, partial	E-151
2088.seq	618	gi 187701 gb M24194 HUMMHBA123	Human MHC protein homologous to chicken B complex protein mRNA,	0
2089.seq	667	gi 5174504 ref NM_005899.1 IM1752	Homo sapiens membrane component, chromosome 17, surface marker 2	0
2090.seq	660	gi 4210521 gb AC004596 AC004596	Homo sapiens chromosome 17, clone hCIT.175_E_5, complete sequence [Hom	0.7
2091.seq	722	gi 181226 gb M22865 HUMCYB5	Human cytochrome b5 mRNA, complete cds.	0
2092.seq	482	gi 4827147 dbj AF000209.1 AP000209	Homo sapiens genomic DNA, chromosome 21q22.1, D21S226-AML region, cl	E-108
2094.seq	487	gi 3366591 gb AC002120 AC002120	Homo sapiens chromosome 9q34, clone 1B2E8, complete sequence [Homo	9.00E-06
2095.seq	479	gi 4504766 ref NM_002211.1 ITGB1	Homo sapiens Integrin, beta 1 (fibronectin receptor, beta	E-153
2096.seq	612	gi 4884398 emb AL050183.1 HSM800479	Homo sapiens mRNA; cDNA DKFZp586A2223 (from clone DKFZp586A2223)	0
2097.seq	569	gi 5123747 emb AL035562.14 HS106502	Human DNA sequence from clone 106502 on chromosome 20p11.21-11.23.	0.6
2098.seq	662	gi 181226 gb M22865 HUMCYB5	Human cytochrome b5 mRNA, complete cds.	0
2099.seq	606	gi 4758027 ref NM_004369.1 COL6A3	Homo sapiens collagen, type VI, alpha 3 (COL6A3) mRNA	0
2100.seq	276	gi 4308817 gb AC006315 AC006315	Homo sapiens PAC clone DJ0630F01 from Tq33-q35, complete sequence [Ho	5.00E-06
2102.seq	637	gi 4507952 ref NM_003406.1 YWHAZ	Homo sapiens tyrosine 3-monoxygenase/tryptophan 5-monoxygenase	E-160
2104.seq	629	gi 3510235 gb AC005580 AC005580	Homo sapiens chromosome 19, cosmid F15394, complete sequence [Homo	0.17
2106.seq	500	gi 452059 gb L27560 HUMIGFBP5X	Human insulin-like growth factor binding protein 5 (IGFBP5) mRNA.	E-100
2107.seq	673	gi 184250 gb L08048 HUMHMG1C	Human non-histone chromosomal protein (HMG-1) retroseudogene.	0
2108.seq	464	gi 450670 ref NM_001025.1 RPS23	Homo sapiens ribosomal protein S23 (RPS23) mRNA	0
2109.seq	603	gi 4501650 gb AC005158.2 AC005158	Homo sapiens clone GS250N06, complete sequence	0.84
2111.seq	417	gi 4503304 ref NM_001926.1 DEFA6	Homo sapiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	0
2112.seq	657	gi 4929666 gb AF151857.1 AF151857	Homo sapiens CGI-99 protein mRNA, complete cds	0
2113.seq	401	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-175
2114.seq	374	gi 4506724 ref NM_001007.1 RPS4X	Homo sapiens ribosomal protein S4, X-linked (RPS4X) mRNA	0
2115.seq	398	gi 2564323 dbj AB006626 AB006626	Homo sapiens mRNA for KIAA0288 gene, complete cds	8.00E-95
2116.seq	389	gi 189019 gb M22918 HUMMYLBC	Human non-muscle myosin alkali light chain mRNA, 3' end.	0
2118.seq	395	gi 1028944 emb Z57713 HS191C11F	H.sapiens CpG island DNA genomic MseI fragment, clone 191c11,	1.00E-50
2119.seq	397	gi 3046306 gb AC004549 AC004549	Homo sapiens BAC clone RG459N13 from 7p15, complete sequence [Homo	0
2120.seq	275	gi 1773032 gb U80924 HSMCPA02	Human monocyte chemoattractant protein 1 receptor gene, two	E-99
2124.seq	387	gi 1665774 gb D87443 D87443	Human mRNA for KIAA0254 gene, complete cds	0
2125.seq	342	gi 189019 gb M22918 HUMMYLBC	Human non-muscle myosin alkali light chain mRNA, 3' end.	E-166
2126.seq	370	gi 337297 gb M10546 HUMMTM1	Human mitochondrial DNA, fragment M1, encoding transfer RNAs,	8.00E-98
2131.seq	371	gi 179023 gb M64497 HUMARP1	Human apolipoprotein AI regulatory protein (ARP-1) mRNA, complete	2.00E-30
2132.seq	347	gi 4506640 ref NM_000997.1 RPL37	Homo sapiens ribosomal protein L37 (RPL37) mRNA	0
2133.seq	395	gi 241477 gb S74678 S74678	heterogeneous nuclear ribonucleoprotein complex K [human, mRNA, 2302	E-169
2134.seq	365	gi 1136435 dbj D80010 D80010	Human mRNA for KIAA0188 gene, partial cds	0
2135.seq	395	gi 3643808 gb AF062346 AF062346	Homo sapiens zinc finger protein 216 splice variant 1 (ZNF216) mRNA.	0
2136.seq	366	gi 4895008 gb AF143866.1 AF143866	Homo sapiens clone IMAGE:110987 mRNA sequence	3.00E-08
2137.seq	371	gi 2781407 gb U92989 HSU92989	Homo sapiens clone DT1P1E11 mRNA, CAG repeat region	0
2141.seq	331	gi 169573 gb M62755 P0YST4C11	Potato 4-coumarate--CoA ligase (S4C1-1) gene, complete cds.	1.3
2144.seq	370	gi 3548785 gb AC005618 AC005618	Homo sapiens chromosome 5, BAC clone 249h5 (LBNL H149), complete	0.097
2146.seq	370	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-124
2147.seq	398	gi 1034701 emb Z62323 HS67B10F	H.sapiens CpG island DNA genomic MseI fragment, clone 67b10,	0.1
2148.seq	367	gi 2104932 gb U95118 MMU95118	Mus musculus Pafaha/Lis1 exon 2 and Pafaha/Lis1 exon 2, complete	1.5
2149.seq	366	gi 3334980 gb AC005306 AC005306	Homo sapiens chromosome 19, cosmid R27216, complete sequence [Homo	1.00E-07
2150.seq	378	gi 1829716 gb M33197 HUMGAPDH	Human glyceraldehyde-3-phosphate dehydrogenase (GAPDH) mRNA,	0
2151.seq	366	gi 4503304 ref NM_001926.1 DEFA6	Homo sapiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	0
2152.seq	366	gi 4503304 ref NM_001926.1 DEFA6	Homo sapiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	0
2160.seq	270	gi 1514698 ref NM_006074.1 STAF50	Homo sapiens stimulated trans-acting factor (50 kDa) (STAF50) mRNA	8.00E-14
2161.seq	211	gi 3046308 gb AC004548 AC004548	Homo sapiens PAC clone DJ1182N03 from 7q11.23-q21.1, complete sequen	0.21
2162.seq	429	gi 2654487 gb U92262 HSU92262	Human dopamine transporter (DAT1) gene, promoter region, exons 1 and	0.45
2164.seq	864	gi 3882272 dbj AB018319.1 AB018319	Homo sapiens mRNA for KIAA0776 protein, partial cds	0
2165.seq	479	gi 5348385 gb AC006111.2 AC006111	Homo sapiens chromosome 16 clone RQC1-1_461A8, complete sequence	0
2166.seq	383	gi 1161564 emb Z68746 HSQ27	Human DNA sequence from cosmid Q27, chromosome region 11p15.5	1.6
2167.seq	485	gi 5001542 gb AC004947.2 AC004947	Homo sapiens clone DJ1007F24, complete sequence	0.033
2168.seq	392	gi 508477 gb L20599 HUMCYPTA01	Homo sapiens cholesterol 7 alpha-hydroxylase (CYP7) gene, 5'	1.6
2170.seq	587	gi 3282167 gb AC005213 AC005213	Homo sapiens chromosome 5, BAC clone 84E9 (LBNL H120), complete	0.002
2171.seq	538	gi 33737 emb X57819 HSGVL029	Human rearranged immunoglobulin lambda4 light chain mRNA	2.00E-44
2172.seq	455	gi 3320449 gb AF059580 AF059580	Homo sapiens histone deacetylase 3 (HDAC3) gene, complete cds	0.48
2173.seq	397	gi 5225311 gb AF072441.1 AF072441	Homo sapiens calcineurin binding protein cabin 1 mRNA, complete cds	0
2174.seq	165			
2175.seq	272	gi 4775633 emb AL031681.13 HS862K6	Human DNA sequence from clone 862K6 on chromosome 20q12-13.13. Conta	1.1
2176.seq	519	gi 13000 emb X62996 MHSGENOM	H.sapiens mitochondrial genome (consensus sequence)	0
2177.seq	435	gi 4507942 ref NM_003400.1 XPO1	Homo sapiens exportin 1 (CRM1), yeast, homolog (XPO1) mRNA	E-176
2178.seq	527	gi 5823553 gb AF180920.1 AF180920	Homo sapiens cyclin anla-6a mRNA, complete cds	mpIEE sEquEncE2.2
2179.seq	373	gi 291899 gb 18991 HUMCDC8X	Human thymidylate kinase (CDC8) mRNA, complete cds.	0.098
2180.seq	556	gi 1514580 emb X95747 HSGKLC5	H.sapiens immunoglobulin kappa chain (clone: 5)	E-154
2181.seq	545	gi 5454611 ref NM_006472.1 VDUP1	Homo sapiens upregulated by 1,25-dihydroxyvitamin D-3 (VDUP1) mRNA	0
2182.seq	118	gi 5531494 emb AL035593.11 HS310J6	Human DNA sequence from clone 310J6 on chromosome 6q22.1-22.3.	1.7
2183.seq	441	gi 5524930 emb AL036842.1 HSM800681	Homo sapiens mRNA; cDNA DKFZp586D1519 (from clone DKFZp586D1519)	0
2184.seq	481	gi 5926677 dbj AP000495.1 AP000495	Homo sapiens genomic DNA, chromosome 3p21.3, clone:301 to 308.	0
2185.seq	654	gi 36139 emb X52967 HRSRPL7	Human mRNA for ribosomal protein L7	0
2186.seq	486	gi 3887213 gb AC005738 AC005738	Homo sapiens chromosome 5, BAC clone 7g12 (LBNL H126), complete sequer	0
2187.seq	262	gi 4512277 dbj AB019438.1 AB019438	Homo sapiens DNA for immunoglobulin heavy-chain variable region,	0.26
2188.seq	511	gi 3396888 gb M17733 HUMTHYB4	Human thymosin beta-4 mRNA, complete cds.	0
2189.seq	448	gi 35972 emb X64634 HRSREP369	H.sapiens DNA repetitive subtelomeric-like sequence (369 bp)	2.00E-09
2190.seq	386	gi 6010168 emb AL078596.8 HSDJ429G5	Human DNA sequence from clone 429G5 on chromosome 6q21-22.1, complet	E-172
2191.seq	892	gi 4757819 ref NM_004231.1 ATP6S14	Homo sapiens ATPase, vacuolar, 14 kD (ATP6S14) mRNA	0
2192.seq	241	gi 2827206 gb AF038969 AF038969	Homo sapiens general transcription factor 2-1 (GTF2I) mRNA,	E-108
2193.seq	520	gi 394630 emb Z24430 HS331YC5	H. sapiens (D11S1353) DNA segment containing (CA) repeat: clone	9.00E-43
2194.seq				
2195.seq	475	gi 4506518 ref NM_003617.1 RGSS5	Homo sapiens regulator of G-protein signalling 5 (RGSS5) mRNA	E-158
2196.seq	495	gi 5016088 ref NM_001101.2 ACTB	Homo sapiens actin, beta (ACTB) mRNA	E-156
2197.seq	545	gi 5262548 emb AL080118.1 HSM800626	Homo sapiens mRNA; cDNA DKFZp564F1123 (from clone DKFZp564F1123)	0
2198.seq	572	gi 3288418 emb AL024506 HS133H11	Human DNA sequence from clone 133H11 on chromosome 6p24. Contains ST	0.039
2199.seq	459	gi 4505886 ref NM_002667.1 PLN	Homo sapiens phospholamban (PLN) mRNA >gi 189942 gb M63603 HUMPHLA	0
2200.seq	464	gi 4505742 ref NM_002624.1 PFDFNS	Homo sapiens prefoldin 5 (PFDFNS) mRNA >gi 1731808 gb D89667 D89667	0
2201.seq	520	gi 4507520 ref NM_001064.1 TKT	Homo sapiens transketolase (Wernicke-Korsakoff syndrome) (TKT) mRNA	0
2203.seq	451	gi 261239 gb S50732 S50732	Immunoglobulin M light chain V region=anti-lipid A antibody [human,	E-143
2204.seq	583	gi 6010305 emb AL079306.2 CNS008MP	Human chromosome 14 DNA sequence *** IN PROGRESS *** BAC R-216F9	0.01
2205.seq	437	gi 4835699 dbj AP000390.1 AP000390	Homo sapiens genomic DNA, chromosome 21q22.1, D21S226-AML region,	1.8
2206.seq	448	gi 4926910 gb AC006476.3 AC006476	Homo sapiens clone DJ0764012, complete sequence	0.47
2207.seq	554	gi 5791509 emb AL050341.18 HSDJ39G2	Human DNA sequence from clone 39G22 on chromosome 1p32.1-34.3, compk	E-149
2209.seq	559	gi 3043669 dbj AB011145 AB011145	Homo sapiens mRNA for KIAA0573 protein, partial cds	0
2210.seq	523	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	0
2211.seq	809	gi 1218057 gb U49114 HSU49114	Human prothormone convertase 5 precursor (PCS) mRNA, partial cds.	4.00E-77
2212.seq	516	gi 3980526 gb AC005483 AC005483	Homo sapiens PAC clone DJ0806A17 from 7p13-p14, complete sequence [Ho	0.14
2213.seq	552	gi 1915975 gb U63963 HSU63963	Human CSF-1 receptor (FMS) gene, complete cds, and (SMF) gene, partial	E-177
2214.seq	503	gi 4753239 gb AC006370.2 AC006370	Homo sapiens clone NH0292P09, complete sequence	0.13
2215.seq	475	gi 1280207 gb L76937 HUMDR	Homo sapiens Werner syndrome gene, complete cds	0.092
2216.seq				
2217.seq	476	gi 4504766 ref NM_002211.1 ITGB1	Homo sapiens Integrin, beta 1 (fibronectin receptor, beta	0
2218.seq	355	gi 4506646 ref NM_001000.1 RPL39	Homo sapiens ribosomal protein L39 (RPL39) mRNA	0
2219.seq	315	gi 4027011 gb L10612 HUMGIA	Human glycosylation-inhibiting factor mRNA, complete cds.	E-177

2220.seq	191	gi 3201899 gb AF067420 AF067420	Homo sapiens SNC73 protein (SNC73) mRNA, complete cds	E-103
2221.seq	824	gi 2270906 gb AC000378 AC000378	Human Chromosome 11 pac pDJ1173a5, complete sequence [Homo sapiens]	7.00E-60
2223.seq	813	gi 224590 dbj AB002323 AB002323	Human mRNA for KIAA0325 gene, partial cds	0
2224.seq	593	gi 5852347 gb AF096774.1 AF096774	Homo sapiens kinase-related protein isoform 1 mRNA, complete cds	0
2225.seq	475	gi 5708093 emb AL049835.3 CNS0000E	Human chromosome 14 DNA sequence *** IN PROGRESS *** BAC R-862P1	0.5
2226.seq	528	gi 464183 dbj D13891 HUMID2HC	Human mRNA for Id-2H, complete cds	E-174
2227.seq	423	gi 537525 gb L29766 HUMEPHYDD	Homo sapiens epoxide hydrolase (EPHX) gene, complete cds	0
2229.seq	505	gi 576474 gb M28203 HUMMBW62	Homo sapiens (clone pMF18) MHC class I HLA-Bw62 mRNA, 3' end,	0
2230.seq	826	gi 3954884 emb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	0
2231.seq	420	gi 3327223 dbj AB014605 AB014605	Homo sapiens mRNA for KIAA0705 protein, complete cds	0.11
2232.seq	496	gi 4071041 emb AL031775 HS30M3	Human DNA sequence from clone 30M3 on chromosome 6p22.1-22.3. Contain	0
2233.seq	515	gi 3702281 gb AC005789 AC005789	Homo sapiens chromosome 19, cosmid F5960, complete sequence [Homo	0
2234.seq	471	gi 340038 gb L05148 HUMTYRKIN	Human protein tyrosine kinase related mRNA sequence.	0
2235.seq	370	gi 189019 gb M22918 HUMMYLCB	Human non-muscle myosin alkali light chain mRNA, 3' end.	0
2236.seq	565	gi 3461879 dbj AB003502 AB003502	Mus musculus mRNA for Guanine Nucleotide Regulatory Protein, partial	0
2237.seq	816	gi 4972128 emb AL049838.2 CNS0000C	Human chromosome 14 DNA sequence *** IN PROGRESS *** BAC R-999E2	0.22
2238.seq	481	gi 13683 emb V00710 MIT1HS	Human mitochondrial genes for several tRNAs (Phe, Val, Leu) and 12S	0
2239.seq	562	gi 4314330 gb AC005874 AC005874	Homo sapiens chromosome 10 clone C17987SH-1175G20 map 10q25.2-10q25	0.59
2240.seq	561	gi 3954892 emb AJ010446 HSA010446	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	0
2242.seq	567	gi 4049267 gb U17714 HSU17714	Homo sapiens putative tumor suppressor ST13 (ST13) mRNA, complete cds	0
2243.seq	510	gi 4506730 ref NM_001010.1 RPS6	Homo sapiens ribosomal protein S6 (RPS6) mRNA	0
2244.seq	486			
2245.seq	443	gi 4504102 ref NM_002083.1 GPX2	Homo sapiens glutathione peroxidase 2 (gastrointestinal) (GPX2)	0
2246.seq	488	gi 339688 gb M17733 HUMTHYB4	Human thymosin beta-4 mRNA, complete cds.	0
2247.seq	270	gi 4156151 gb AC005007 AC005007	Homo sapiens clone G5051M12, complete sequence [Homo sapiens]	6.00E-30
2248.seq	822	gi 3478647 gb AC002554 AC002554	Human Chromosome 11p11.2 PAC clone pDJ404m15, complete sequence [Ho	1.00E-05
2249.seq	560	gi 2996648 gb AC004493 AC004493	Homo sapiens chromosome 16, cosmid clone 373C8 (LANL), complete	0.01
2250.seq	660	gi 4003388 dbj AB020868.1 AB020868	Homo sapiens genomic DNA of 8p21.3-p22 anti-oncogene of hepatocellular	2.00E-77
2251.seq	485	gi 1145798 gb U41806 HSU41806	Human EB13-associated protein p60 mRNA, complete cds.	E-124
2252.seq	809	gi 3114717 emb AL022098 HS510L9	Homo sapiens DNA sequence from PAC 510L9 on chromosome 6p24.1-p25.3	0.004
2253.seq	459	gi 3377550 gb AF075010 HUMYK03D03	Homo sapiens full length insert cDNA YK03D03	0
2254.seq	594	gi 4960207 gb AF154065.1 AF154065	Homo sapiens Inorganic pyrophosphatase (PP) mRNA, complete cds	0
2255.seq	526	gi 189904 gb L00160 HUMPGK2	Human phosphoglycerate kinase (pgk) mRNA, exons 2 to last.	0
2256.seq	538	gi 1381178 gb U58766 HSU58766	Human FX protein mRNA, complete cds	0
2257.seq	481	gi 5815494 gb AC007690.1 AC007690	Homo sapiens 3q27 BAC RPC11-42D20 (Roswell Park Cancer Institute Hum	0.13
2258.seq	519	gi 5263012 emb AL031121.5 HS49501D	Human DNA sequence from clone 49501D on chromosome 6q26-27. Contains	0.14
2259.seq	832			
2260.seq	34			
2261.seq	597	gi 5103023 dbj AP000360.1 AP000360	Homo sapiens genomic DNA, chromosome 22q11.2, clone N81C12	0.63
2262.seq	501	gi 5032050 ref NM_005617.1 RPS14B	Homo sapiens ribosomal protein S14 (RPS14) mRNA	0
2263.seq	553	gi 6005959 ref NM_007222.1 ZHX1	Homo sapiens zinc-fingers and homeoboxes 1 (ZHX1), mRNA	0.15
2264.seq	563	gi 505091 dbj D31767 HUMORFKG1F	Human mRNA for KIAA0058 gene, complete cds	0
2265.seq	869	gi 2102635 gb AB000528 AB000528	Human DNA for thrombopoietin, 5'flanking region	3.7
2266.seq	835	gi 5729777 ref NM_004370.3 COL12A1	Homo sapiens collagen, type XII, alpha 1 (COL12A1) mRNA	0
2267.seq	484	gi 4507484 ref NM_003246.1 THBS1	Homo sapiens thrombospondin 1 (THBS1) mRNA >gi 37464 emb X14787 HST	E-105
2268.seq	835	gi 4503606 ref NM_000126.1 ETFAI	Homo sapiens electron-transfer-flavoprotein, alpha polypeptide	0
2269.seq				
2270.seq	506	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-128
2271.seq	620	gi 5031930 ref NM_005594.1 NACA	Homo sapiens nascent polypeptide-associated complex alpha	0
2272.seq	410	gi 4503248 ref NM_003472.1 D6S231E	Homo sapiens DEK gene (D6S231E) mRNA >gi 30502 emb X64229 HSDEK9	0
2273.seq	497	gi 5174602 ref NM_006097.1 MYRL2	Homo sapiens myosin regulatory light chain 2, smooth muscle isoform	0
2274.seq	600	gi 3928029 gb AC005859 AC005859	Homo sapiens Xp22-83 BAC GSHB-324M7 (Genome Systems Human BAC Li	0.16
2275.seq	608	gi 809026 emb X86693 HSMAS79	H.sapiens mRNA for hevlin like protein	0
2277.seq	537	gi 1514580 emb X95747 HSIGKLC5	H.sapiens immunoglobulin kappa chain (clone: 5)	0
2279.seq	601	gi 2708753 gb AC003955 AC003955	Homo sapiens chromosome 5, Pac clone 236L2 (LBNL H93), complete	0.01
2281.seq	558	gi 1038257 emb Z65435 HS39F9F	H.sapiens CpG island DNA genomic MseI fragment, clone 39F9, forward	3.00E-80
2282.seq	845	gi 6009641 dbj AB025905.1 AB025905	Homo sapiens NAKAP95 mRNA for neighbor of A-kinase anchoring	0
2283.seq	560	gi 2842785 gb AC004130 AC004130	Homo sapiens BAC clone RG293F17 from 7p15-p21, complete sequence [H	0
2284.seq	609	gi 182976 gb M33197 HUMGAPDH	Human glyceraldehyde-3-phosphate dehydrogenase (GAPDH) mRNA,	0
2285.seq	579	gi 3954884 emb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	0
2286.seq	559	gi 31091 emb X16869 HSEF1AC	Human mRNA for elongation factor 1-alpha (clone CEF4)	0
2287.seq	480	gi 4309874 gb AC006501 AC006501	Homo sapiens, clone hRPK.35_A_1, complete sequence [Homo sapiens]	0
2288.seq	419	gi 4506636 ref NM_000995.1 RPL34	Homo sapiens ribosomal protein L34 (RPL34) mRNA	0
2289.seq	557	gi 4582842 emb AL035451.5 HS145B8	Human DNA sequence from clone 145B8 on chromosome Xq21.2-22.3, comp	0
2290.seq	585	gi 6005738 ref NM_007158.1 D1S155E	Homo sapiens NRAS-related gene (D1S155E), mRNA	0
2291.seq	600	gi 3077822 gb AC004603 AC004603	Homo sapiens chromosome 19, cosmid R33496, complete sequence [Homo	0.64
2292.seq	521	gi 3094991 gb AC004636 AC004636	Homo sapiens chromosome 5, P1 clone 1268h6 (LBNL H50), complete	0.035
2293.seq	604	gi 188697 gb M31659 HUMMSCA	Human GT mitochondrial solute carrier protein homologue mRNA,	0
2296.seq	525	gi 4885240 ref NM_005252.1 FOS	Homo sapiens v-fos FBJ murine osteosarcoma viral oncogene homolog	0
2297.seq	539	gi 1669684 emb Y09305 HSDYRK4	H.sapiens mRNA for protein kinase, Dyrk4, partial	0
2298.seq	561	gi 1262342 emb X93334 HSMITG	H.sapiens mitochondrial DNA, complete genome	0
2299.seq	587	gi 4826855 ref NM_005006.1 NDUFS1	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 1 (75kD)	0
2300.seq	529	gi 3860395 emb AL031282 HS283E3	Human DNA sequence from clone 283E3 on chromosome 1p36.21-36.33.	6.00E-84
2301.seq	518	gi 1220360 gb L31610 HUMS29RP	Homo sapiens (clone cor1-cd5) S29 ribosomal protein mRNA, complete	E-142
2302.seq	481	gi 5817087 emb AL110177.1 HSM800824	Homo sapiens mRNA: cDNA DKFZp564C0762 (from clone DKFZp564C0762)	0
2303.seq				
2304.seq	485	gi 4506682 ref NM_001016.1 RPS12	Homo sapiens ribosomal protein S12 (RPS12) mRNA	0
2306.seq	556	gi 2760552 emb Z98753 HS238J17	Human DNA sequence from PAC 238J17 on chromosome 6q22. Contains EST	0.59
2307.seq	570	gi 5835387 ref NC_001807.1	Human mitochondrion, complete genome >gi 1944628 gb J01415 HUMMTCG	0
2308.seq	557	gi 4506690 ref NM_001020.1 RPS16	Homo sapiens ribosomal protein S16 (RPS16) mRNA	0
2309.seq	283	gi 4333491 emb X56841 HSHLAE	H.sapiens HLA-E gene >gi 1376220 gb G26970 G26970 human STS	1.00E-62
2310.seq	522	gi 190978 gb M18963 HUMREGA	Human islet of Langerhans regenerating protein (reg) mRNA, complete	0
2311.seq	810	gi 3821247 emb Y17293 HSY17293	Homo sapiens FL-1 gene, partial	3.4
2312.seq	497	gi 13000 emb X62996 MHSGENOM	H.sapiens mitochondrial genome (consensus sequence)	0
2313.seq	818	gi 872118 emb Z49112 HSHLAXCW7	H.sapiens gene for leukocyte antigen-Cw7 variant (HLA-Cw*0702)	0
2314.seq	459	gi 505101 dbj D31887 HUMORFKG1P	Human mRNA for KIAA0062 gene, partial cds	0
2315.seq	383	gi 190978 gb M18963 HUMREGA	Human islet of Langerhans regenerating protein (reg) mRNA, complete	0
2316.seq	472	gi 1668818 gb L29074.1 HUMFMR1S	Homo sapiens fragile X mental retardation syndrome protein (FMR1) gene,	0.49
2317.seq	561	gi 4758137 ref NM_004396.1 DDX5	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/Hs) box polypeptide 5 (RNA	0
2318.seq	499	gi 4240388 gb AF067845 AF067845	Homo sapiens clone P1-4248, complete sequence [Homo sapiens]	0.13
2319.seq	619	gi 4503014 ref NM_003909.1 CPNE3	Homo sapiens copine III (CPNE3) mRNA	E-141
2320.seq	389	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	0
2321.seq	508	gi 5803180 ref NM_006819.1 STIP1	Homo sapiens stress-induced-phosphoprotein 1 (Hsp70/Hsp90-organizing	0
2322.seq	578	gi 4826845 ref NM_004999.1 MYO6	Homo sapiens myosin VI (MYO6) mRNA >gi 2224718 dbj AB002387 AB002387	0
2324.seq	529	gi 510270 gb U09500 HSU09500	Human mitochondrion cytochrome b gene, partial cds.	0
2325.seq	519	gi 3293212 gb AC005271 AC005271	Homo sapiens chromosome 7 clone UWGC:g1564a040 from 7p14-15, complete	0.55
2326.seq	491	gi 4809272 ref NM_001153.2 ANX4	Homo sapiens annexin IV (placental anticoagulant protein II) (ANX4)	0
2327.seq	542	gi 4758713 ref NM_004528.1 MGST3	Homo sapiens microsomal glutathione S-transferase 3 (MGST3) mRNA	0
2328.seq	341	gi 5453721 ref NM_006330.1 LYPLA1	Homo sapiens lysophospholipase I (LYPLA1), mRNA	0
2329.seq	283	gi 34717 emb X64899 HSMP21HOM	H.sapiens mRNA homologous to mouse P21 mRNA	E-158
2330.seq	443	gi 4505854 ref NM_002655.1 PLAG1	Homo sapiens pleomorphic adenoma gene 1 (PLAG1) mRNA, and translated	0
2332.seq	468	gi 5738629 emb AL049709.15 HSDJ18CS	Human DNA sequence from clone 18C9 on chromosome 20, complete sequen	0.49
2333.seq	516	gi 5360124 gb AF155115.1 AF155115	Homo sapiens NY-REN-58 antigen mRNA, complete cds	0
2334.seq	498	gi 3649719 emb AL031009 HS431H6	Human DNA sequence from clone 431H6 on chromosome 16. Contains a nove	9.00E-80
2335.seq	165	gi 999188 gb S78172 S78172	rRNA [IGS, intergenic spacer, repeats] [Brassica rapa, mRNA Partial,	0.63
2336.seq	448	gi 4753226 gb AC006385.3 AC006385	Homo sapiens clone NH0559J05, complete sequence	0.47
2337.seq	465	gi 292165 gb L13850 HUMHXBP1YR	Homo sapiens hXBP-1 transcription factor DNA.	0
2338.seq	385	gi 1109629 dbj D49726 HUMGPIASL	Human PIG-F gene for GPI-anchor synthesis, intron5-exon6 boundary.	1.00E-44

