

2460.seq	613	gi 665587 dbj D45887 HUMALDN	Human mRNA for calmodulin, complete cds	
2461.seq	878	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-135
2462.seq	529	gi 4929694 gb AF151871.1 AF151871	Homo sapiens CGI-113 protein mRNA, complete cds	0
2463.seq	427	gi 5835387 ref NC_001807.1	Human mitochondrion, complete genome >gi 1944628 gb J01415 HUMMTCG	0
2464.seq	476	gi 4753248 gb AC007033.3 AC007033	Homo sapiens BAC clone NH0028D14 from 7p15.1-p13, complete sequence	0.5
2465.seq	864	gi 45112277 dbj AB019438.1 AB019438	Homo sapiens DNA for immunoglobulin heavy-chain variable region, complete cds	E-122
2466.seq	391	gi 5815507 gb AC006509.1 AC006509	Homo sapiens Zp16 PAC RPCIS-960D23 (Roswell Park Cancer Institute Human)	5.00E-96
2468.seq	355	gi 5174470 ref NM_005896.1 IDH1	Homo sapiens isocitrate dehydrogenase 1 (NADP+), soluble (IDH1) mRNA	0
2470.seq	417	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	E-112
2471.seq	571	gi 32487 emb X15183 HSHSP90R	Human mRNA for 90-kDa heat-shock protein	0
2472.seq	469	gi 4914589 emb ALD050386.1 HSM800533	Homo sapiens mRNA; cDNA DKFZp564G0422 (from clone DKFZp564G0422)	E-148
2473.seq	560	gi 388031 gb L05092 HUMRIBPROC	Homo sapiens ribosomal protein L11 mRNA, complete cds	0
2474.seq	693	gi 4826705 ref NM_004951.1 EBI2	Homo sapiens Epstein-Barr virus induced gene 2	4.00E-21
2475.seq	593	gi 3242769 gb AC005166 AC005166	Homo sapiens PAC clone DJ0905M06 from 7q31, complete sequence [Homo]	0.63
2476.seq	605	gi 6002584 gb AF091035.1 AF091035	Homo sapiens GTP-binding protein RAB21 (RAB21) mRNA, complete cds	0
2477.seq	872	gi 5453905 ref NM_006319.1 PIS1	Homo sapiens phosphatidylinositol synthase (PIS) mRNA	0
2478.seq	663	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-174
2479.seq	213	gi 4507512 ref NM_000362.1 TIMP3	Homo sapiens tissue inhibitor of metalloproteinase 3 (Sorsby fundus)	4.00E-46
2480.seq	426	gi 3133268 gb U20352 HSU20352	Homo sapiens malate dehydrogenase (MDHA) mRNA, complete cds	0
2481.seq	270	gi 2293555 gb AF012086 AF012086	Homo sapiens Ran binding protein 2 (RanBP2alpha) mRNA, partial cds	E-148
2482.seq	679	gi 4502950 ref NM_000900.1 COL3A1	Homo sapiens collagen, type III, alpha 1 (Ehlers-Danlos syndrome type I)	0
2483.seq	564	gi 5834027 gb AF174034.1 AF174034	Homo sapiens clone 77u-c28 immunoglobulin heavy chain variable	E-139
2485.seq	621	gi 4508141 gb AC006017.2 AC006017	Homo sapiens PAC clone DJ0981007 from 7q34-q36, complete sequence	0.042
2486.seq	643	gi 4506822 ref NM_002979.1 SCP2	Homo sapiens sterol carrier protein 2 (SCP2) mRNA	0
2487.seq	598	gi 3881948 emb ALQ031774 HS298L15	Human DNA sequence from clone 298L15 on chromosome 6p22.3-23 Contain	0
2488.seq	654	gi 2969925 emb AJ000534 HEPSSARC	Homo sapiens mRNA for epsilon-sarcoglycan	0
2490.seq	552	gi 4502598 ref NM_001757.1 CBR1	Homo sapiens carbonyl reductase 1 (CBR1) mRNA	0
2491.seq	554	gi 4758041 ref NM_004718.1 COX7RP1	Homo sapiens cytochrome c oxidase subunit VII-related protein	0
2492.seq	603	gi 5835387 ref NC_001807.1	Human mitochondrion, complete genome >gi 1944628 gb J01415 HUMMTCG	0
2493.seq	521	gi 190978 gb M18963 HUMREGA	Human islet of Langerhans regenerating protein (reg) mRNA, complete	0
2494.seq	608	gi 456345 emb X73079 HSP1R	Homo sapiens encoding Polymeric immunoglobulin receptor	5.00E-17
2495.seq	613	gi 33709 emb X57806 HSIGVL006	Human rearranged immunoglobulin lambda light chain mRNA	0
2496.seq	446	gi 2347181 gb AC002112 AC002112	Genomic sequence from Human 6, complete sequence [Homo sapiens]	0.002
2497.seq	640	gi 755555 gb U21852 ETU21852	Erythrodermis tralilin ribulose-1,5-bisphosphate carboxylase	0.17
2498.seq	525	gi 5306223 gb AC006255.9 AC006255	Homo sapiens 3p21.1-9 PAC RPCIS-1087L12 (Roswell Park Cancer Institute)	3.00E-67
2499.seq	637	gi 405193 gb L10811 HUMHCCB	Homo sapiens splicing factor (CC1.4) mRNA, complete cds.	0
2500.seq	627	gi 4557374 ref NM_001206.1 BTEB1	Homo sapiens basic transcription element binding protein 1 (BTEB1)	0
2502.seq	887	gi 184918 gb M60192 HUMIGHA2A	Human germline IgA2 chain C region.	E-169
2503.seq	585	gi 2801527 gb AF035397 AF035397	Homo sapiens cosmid Qc15C1 and 94B6 from Xq28, complete sequence [Hor	0
2504.seq	868	gi 179349 gb M33521 HUMBAT3B2	Human HLA-B-associated transcript 3 (BAT3) gene, 3' end.	0
2506.seq	706	gi 4506730 ref NM_001010.1 RPS6I	Homo sapiens ribosomal protein S6 (RPS6) mRNA	0
2507.seq	523	gi 5032036 ref NM_005612.1 REST1	Homo sapiens RE1-silencing transcription factor (REST) mRNA	0
2508.seq	633	gi 4826923 ref NM_005034.1 POLR2K	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide K	E-175
2509.seq	643	gi 4176417 dbj AB004304 AB004304	Homo sapiens mRNA for IgG kappa chain, partial cds	0
2510.seq	618	gi 2583221 gb AF026948 AF026948	Saccharomyces cerevisiae 15S ribosomal RNA gene, mitochondrial gene	0.003
2511.seq	583	gi 6005813 ref NM_007271.1 NDR1	Homo sapiens serine threonine protein kinase (NDR), mRNA	0
2512.seq	660	gi 5005783 ref NM_007265.1 HSGT1	Homo sapiens suppressor of <i>S. cerevisiae</i> gcr2 (HSGT1), mRNA	2.00E-29
2513.seq	269	gi 4502166 ref NM_000484.1 APP1	Homo sapiens amyloid beta (A4) precursor protein (protease nexin-II),	E-138
2515.seq				
2516.seq	490	gi 2580549 gb AF00982 HSAF00982	Homo sapiens dead box, X isoform (DBX) mRNA, alternative transcript	E-157
2517.seq	491	gi 4031443 gb M84379 HUMMHAE	Human MHC class I lymphocyte antigen (HLA-A*0201) mRNA, complete	0
2518.seq	531	gi 180757 gb M92642 HUMCOL16A	Homo sapiens alpha-1 type XVI collagen (COL16A1) mRNA, complete cds.	0
2519.seq	597	gi 3164071 emb Z93242 HS389A20	Human DNA sequence from PAC 389A20 on chromosome X contains ESTs ST	2.5
2521.seq				
2522.seq	454	gi 3123551 emb AL021408 HS523C21	Homo sapiens DNA sequence from PAC 523C21 on chromosome 6q23.1-23.3	0.48
2523.seq	457	gi 4758777 ref NM_004546.1 NDUFB2	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2	0
2524.seq	885	gi 4579986 dbj AF000065.1 AP000065	Homo sapiens genomic DNA of 8011.2 senescence gene, segment 1/19,	0.015
2525.seq	884	gi 602125 gb U09120 HSRNPAIL	Human hnRNP protein A2 gene, promoter region and exon 1	0.95
2526.seq	204	gi 3954884 emb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	E-107
2527.seq	392	gi 4506604 ref NM_002580.1 PAP1	Homo sapiens pancreatitis-associated protein (PAP) mRNA	0
2528.seq	495	gi 1834463 emb Z84816 HSA2A2B	Human DNA sequence from PAC 2A2 on chromosome X contains ESTs	0.002
2530.seq	267	gi 1420884 gb U34333 PVU34333	Phaseolus vulgaris proline-rich 14 kDa protein mRNA, complete cds	0.27
2531.seq				
2532.seq	618	gi 476134 gb U06864 MMU06664	Mus musculus Balb/c mid-G1 serum response (1-8-23) mRNA, 5'	3.00E-37
2533.seq	448	gi 310078 gb L19347 RATADRFLA	Rattus norvegicus DNA sequence with unspecified 5' or 3' flanking	1.9
2534.seq	184	gi 4505316 ref NM_002480.1 MYPT1	Homo sapiens myosin phosphatase, target subunit 1 (MYPT1) mRNA	3.00E-37
2535.seq	608	gi 2295179 emb Z99297.1 HS262D12	Homo sapiens DNA sequence from PAC 262D12 on chromosome 1q23.3-24.3.	0
2537.seq	457	gi 2304970 gb U82828 HSU82828	Homo sapiens ataxia telangiectasia (ATM) gene, complete cds	2.00E-06
2538.seq	372	gi 3483426 gb AF086081 HJMZY83C11	Homo sapiens full length insert cDNA clone Y283C11	1.00E-28
2539.seq	567	gi 5764048 emb AL050322.10 HSJ727110	Human DNA sequence from clone 727110 on chromosome 20, complete	0.01
2540.seq	667	gi 200288 gb M12039 MUSPER	Mouse DNA homologous to the Drosophila per locus.	0.045
2541.seq	297	gi 5931966 gb AF125164.1 AF125164	Bacteroides fragilis 638R polysaccharide B (PS B2) biosynthesis	0.077
2542.seq	563	gi 1407654 gb U58884 MMU58884	Mus musculus SH3-containing protein SH3P7 mRNA, complete cds, similar	4.00E-39
2543.seq	495			
2544.seq	557	gi 4557893 ref NM_000239.1 LYZ	Homo sapiens lysozyme (LYZ) mRNA >gi 187243 gb J03801 HUMLSZ Human	0
2545.seq	163	gi 189383 gb M59907 HUMOGCB	Human melanoma-associated antigen ME491 mRNA, complete cds.	4.00E-33
2547.seq	434	gi 4504618 ref NM_001553.1 IGFBP7	Homo sapiens insulin-like growth factor binding protein 7 (IGFBP7)	E-179
2549.seq	525	gi 4759297 ref NM_004182.1 LXT	Homo sapiens ubiquitously-expressed transcript (LXT) mRNA	0
2550.seq	265	gi 4506634 ref NM_000994.1 RPL32	Homo sapiens ribosomal protein L32 (RPL32) mRNA	E-101
2551.seq	870	gi 3319850 emb AL022164 HS581F7	Human DNA sequence from clone 581F7 on chromosome Xq26.1-27.3 Contain	0.24
2552.seq	575	gi 4506362 ref NM_002492.1 NDUFB5	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 5	0
2554.seq				
2555.seq	607	gi 3954892 emb AJ010446 HSA010446	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	0
2556.seq	522	gi 4506744 ref NM_001013.1 RPS9	Homo sapiens ribosomal protein S9 (RPS9) mRNA	0
2557.seq	870	gi 4467344 emb AL021391.2 HS102D24	Human DNA sequence from clone 102D24 on chromosome 22q13.31-13.32.	0.24
2558.seq	645	gi 609307 gb U17104 HSU17104	Human cytoplasmic chaperonin hTRIC5 mRNA, partial cds	0
2560.seq	381	gi 13683 emb V00710 MIT1HS	Human mitochondrial genes for several tRNAs (Phe, Val, Leu) and 12S	0
2561.seq	584	gi 184743 gb J00220 HUMIGCC8	Homo sapiens immunoglobulin alpha-1 heavy chain constant region	0
2563.seq	583	gi 388031 gb L05092 HUMRIBPROC	Homo sapiens ribosomal protein L11 mRNA, complete cds	0
2565.seq	672	gi 5031752 ref NM_005520.1 HNRPH1	Homo sapiens heterogeneous nuclear ribonucleoprotein H1 (H) (HNRPH1)	0
2566.seq	588	gi 2822174 gb AC004079 AC004079	Homo sapiens PAC clone DJ0167F23 from 7p15, complete sequence [Homo]	0
2567.seq	318	gi 1373420 gb U57847 HSU57847	Human ribosomal protein S27 mRNA, complete cds.	E-176
2568.seq	637	gi 4504374 ref NM_000186.1 HF1	Homo sapiens H factor 1 (complement) (HF1) mRNA	0
2570.seq	530	gi 185363 gb M87790 HUMIGHEPAL	Human (hybridoma H210) anti-hepatitis A immunoglobulin lambda chain	0
2571.seq	296	gi 5360987 emb AL034424.9 HS257E24	Human DNA sequence from clone 257E24 on chromosome 20q12-13.1, comp	3.00E-04
2572.seq	746	gi 5487890 gb AC008017.2 AC008017	Arabidopsis thaliana chromosome I BAC F3N23 genomic sequence, complete	0.051
2574.seq	729	gi 5523838 gb AC007128.2 AC007128	Homo sapiens clone DJ0594A05, complete sequence	6.00E-14
2575.seq	508	gi 4503470 ref NM_001402.1 EEF1A1	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
2576.seq	604	gi 3115230 emb Y08995 HSLAB45V	Homo sapiens mRNA for HLA-B antigen alpha chain (B*5002)	0
2577.seq	405	gi 3126777 gb AC004590 AC004590	Homo sapiens chromosome 17, clone hCIT.22, K.21, complete sequence	0.027
2579.seq	373	gi 184658 gb J03607 HUMIFP	Human 40-kDa keratin intermediate filament precursor gene.	0
2580.seq	666	gi 2957144 gb U90878 HSU90878	Homo sapiens carboxyl terminal LIM domain protein (CLIM1) mRNA,	0.71
2581.seq	692	gi 5881769 emb AL117386.1 AT30A10	Arabidopsis thaliana DNA chromosome 4, BAC clone T30A10 (ESSA project)	0.74
2583.seq	390	gi 181477 gb M14200 HUMDBI	Human diazepam binding inhibitor (DBI) mRNA, complete cds.	E-158
2584.seq	325	gi 4679017 gb AF072202.1 AF072202	Homo sapiens HSPC016 mRNA, complete cds	E-174
2586.seq	536	gi 2731601 gb AC003692 AC003692	Human Chromosome 11 pac pDJ306b4, complete sequence [Homo sapiens]	0.036
2587.seq	555	gi 4406687 gb AF131846 AF131846	Homo sapiens clone 25028 mRNA sequence	0
2588.seq	607	gi 577294 dbj D42041 HUMKIAAE	Human mRNA for KIAA0088 gene, partial cds	0

2589.seq					
2590.seq	396	gi 3659407 gb AC005295 AC005295	Homo sapiens Xp22 bins 3-5 PAC RPC14-617A9 (Roswell Park Cancer		0
2591.seq	614	gi 5689809 emb AL109701.1 JRO45620	Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 45620		0
2592.seq	276	gi 3492855 gb AC002328 AC002328	Genomic sequence for Arabidopsis thaliana BAC F20N2, complete sequence		0.071
2593.seq	447	gi 5420326 emb Z84718.2 HS322B1	Human DNA sequence from clone 322B1 on chromosome 22q11-12, complete		1.00E-84
2594.seq	450	gi 4758509	Homo sapiens H beta 58 homolog (H858) mRNA		0
2595.seq	582	gi 4506674	Homo sapiens ribophorin I (RPN1) mRNA >gi 36052 emb Y00281 HSRIBIR		0
2596.seq	567	gi 1905937 gb U44792 HSU44792	Human Ig kappa L-chain variable region (B6B7K) mRNA, partial cds		6.00E-38
2597.seq	868	gi 4506900	Homo sapiens splicing factor, arginine/serine-rich 3 (SFRS3) mRNA		0
2598.seq	508	gi 4759099	Homo sapiens splicing factor, arginine/serine-rich 11 (SFRS11) mRNA		5.00E-26
2599.seq	496	gi 179363 gb L08634 HUMBCKDHAX	Human branched chain keto acid dehydrogenase E1-alpha subunit		2.00E-80
2601.seq	503				
2602.seq	538	gi 3860076 gb AF097021 AF097021	Homo sapiens GW112 protein (GW112) mRNA, complete cds		E-153
2603.seq	259	gi 870785 gb U26455 HSU26455	Human phosphatidylinositol 3-kinase homolog (ATM) mRNA, complete cds.		E-113
2607.seq	629	gi 5042201	Homo sapiens membrane metallo-endopeptidase (neutral endopeptidase,		0
2608.seq	595	gi 4501888	Homo sapiens actin, gamma 2, smooth muscle, enteric (ACTG2) mRNA		0
2609.seq	500	gi 4826825	Homo sapiens methionine-tRNA synthetase (MARS) mRNA		E-106
2610.seq	570	gi 3355524 gb AC004984 AC004984	Homo sapiens PAC clone DJ1164K10 from Tp21-p22, complete sequence [Hor		0.61
2611.seq	592	gi 35456 emb X54629 HSPHL1	Human pHL-1 gene is a c-myc oncogene containing coxIII sequence		0
2612.seq	483	gi 3435311 gb AF083255 AF083255	Homo sapiens RNA helicase-related protein mRNA, complete cds		0
2613.seq	530	gi 1033989 emb Z61611 HS5587F	H.sapiens CpG island DNA genomic MseI fragment, clone 55b7, forward		0.56
2614.seq	467	gi 4506666	Homo sapiens ribosomal protein, large, PO (RPLPO) mRNA		E-173
2617.seq	513	gi 190978 gb M18963 HUMREGA	Human islet of Langerhans regenerating protein (reg) mRNA, complete		0
2618.seq	432	gi 5454161	Homo sapiens upregulated by 1,25-dihydroxyvitamin D-3 (VDUP1) mRNA		E-156
2619.seq	519	gi 5737839 gb AF149301.1 AF149301	Homo sapiens SOX13 gene, complete cds		2.2
2622.seq	353	gi 4375980 emb AL034406.1 HS571B2	Human DNA sequence from clone 571B2 on chromosome Xq21.1-21.33,		0.37
2623.seq	949	gi 703088 gb M27487 HUMMHDPA	Homo sapiens MHC class II DPw3-alpha-1 chain mRNA, complete cds.		0
2624.seq	477	gi 4503304	Homo sapiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA		0
2625.seq	305	gi 2339843 gb U95742 HSU95742	Human chromosome 16p13.1 BAC clone CIT987SK-551G9 complete sequence		7.00E-98
2626.seq	552	gi 5031962	Homo sapiens heat shock protein A binding protein 2 (c-terminal)		0
2627.seq	260	gi 2341013 gb AC002487 AC002487	Human BAC clone RG369K23 from 7q31, complete sequence [Homo sapiens]		4.1
2629.seq	372	gi 4071020 gb AC006211 AC006211	Homo sapiens chromosome 18, clone hRPK537_E_1, complete sequence [Hom		1.5
2630.seq	534	gi 3201899 gb AF067420 AF067420	Homo sapiens SNC73 protein (SNC73) mRNA, complete cds		0
2631.seq	544	gi 603801 dbj D14694 HUMORF11	Human mRNA for KIAA0024 gene, complete cds		0
2632.seq	352	gi 188542 gb M16279 HUMMIC2A	Human MIC2 mRNA, complete cds.		E-149
2633.seq	888	gi 4757797	Homo sapiens APG5 (autophagy 5, S. cerevisiae)-like (APG5L), mRNA		0
2634.seq	352	gi 4678784 emb AL035610.2 HSM17648	Homo sapiens chromosome 21 PAC LLNLP704M17648Q13, complete sequenc		0.024
2635.seq	496	gi 2280483 dbj AB002370 AB002370	Human mRNA for KIAA0372 gene, complete cds		E-151
2637.seq	537	gi 4759211	Homo sapiens tubulin-specific chaperone a (TBCA) mRNA		0
2638.seq	609	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1		0
2641.seq	238	gi 5689420 dbj AB028965.1 AB028965	Homo sapiens mRNA for KIAA1042 protein, complete cds		2.00E-51
2642.seq	831	gi 2224646 dbj AB002351 AB002351	Human mRNA for KIAA0353 gene, partial cds		0
2643.seq	943	gi 5712732 gb AF154838.1 HSRAB27A4	Homo sapiens Ras-like GTP-binding protein (RAB27A) gene, exon 4		1
2644.seq	453	gi 4263638 gb AC000015 AC000015	Homo sapiens chromosome 4 clone B271E1 map 4q25, complete sequence		0
2646.seq	520	gi 433415 dbj D17554 HUMTR107	Human mRNA for DNA-binding protein, TAXREB107, complete cds		0
2647.seq	382	gi 339688 gb M17733 HUMTHYB4	Human thymosin beta-4 mRNA, complete cds.		0
2648.seq	466	gi 4506652	Homo sapiens ribosomal protein L4 (RPL4) mRNA		0
2650.seq	565	gi 3387930 gb AF070562 AF070562	Homo sapiens clone 24529 mRNA sequence		0
2651.seq	211	gi 13003 emb Y00662 MH5XX	H.sapiens mitochondrial genome		E-109
2652.seq	650	gi 4758713	Homo sapiens microsomal glutathione S-transferase 3 (MGST3) mRNA		0
2653.seq	537	gi 4505704	Homo sapiens phosphoprotein enriched in astrocytes 15 (PEA15) mRNA		0
2654.seq	502	gi 465150 gb L29155 HUMGHZ1	Human immunoglobulin heavy chain variable region (VH III family)		E-102
2655.seq	606	gi 189019 gb M22918 HUMMYLCB	Human non-muscle myosin alkali light chain mRNA, 3' end.		0
2656.seq	496	gi 5032050	Homo sapiens ribosomal protein S14 (RPS14) mRNA		0
2657.seq	288	gi 4835616 dbj AP000247.1 AP000247	Homo sapiens genomic DNA, chromosome 21q22.1, D21S226-AML region,		0.3
2660.seq	561	gi 3928123 gb AC005972 AC005972	Homo sapiens chromosome 17, clone hRPK180_P_8, complete sequence [Hor		E-145
2661.seq	545	gi 5420148 emb AL031587.3 HS1039K5	Human DNA sequence from clone 1039K5 on chromosome 22q12.3-13.2,		0.037
2662.seq	562	gi 2576343 gb AC002041 HUAC002041	Human Chromosome 16 BAC clone CIT987SK-A-17E1, complete sequence [H		2.00E-28
2663.seq	533	gi 546798 gb S71043 S71043	Human alpha-2-immunoglobulin A heavy chain allotype 2 (constant		E-169
2664.seq	331	gi 177922 gb M63473 HUMASCL18	Human alpha-5 collagen type IV gene, exon 1.		E-111
2665.seq	599	gi 3355522 gb AC004992 AC004992	Homo sapiens PAC clone DJ1189K17 from 7q21, complete sequence [Homo		0.15
2666.seq	534	gi 4502208	Homo sapiens ADP-ribosylation factor 5 (ARF5) mRNA		0
2667.seq	611	gi 2337871 gb AC002458 AC002458	Human BAC clone RG098M04 from 7q21-q22, complete sequence [Homo		0.003
2668.seq	420	gi 505472 emb X69181 HSL31	H.sapiens mRNA for ribosomal protein L31		E-145
2669.seq	504	gi 4504954	Homo sapiens laminin, gamma 1 (formerly LAMB2) (LAMC1) mRNA		5.00E-29
2670.seq	573	gi 5835387	Human mitochondrion, complete genome >gi 1944628 gb J01415.1 HUMMTCCG		0
2671.seq	399	gi 4506662	Homo sapiens ribosomal protein L8 (RPL8) mRNA		E-123
2672.seq	478	gi 187701 gb M24194 HUMHBA123	Human MHC protein homologous to chicken B complex protein mRNA,		0
2674.seq	486	gi 261239 gb S50732 S50732	Immunoglobulin M light chain V region:anti-lipid A antibody [Human,		E-153
2675.seq	503	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1		0
2677.seq	505	gi 2588643 gb AC003071 AC003071	Homo sapiens BAC clone BK085E05 from 22q12.1-qter, complete sequence		2.1
2678.seq	429	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.		E-103
2680.seq	912	gi 186914 gb M20206 HUMLAMBA	Human laminin B1 mRNA, complete cds. >gi 591561 gb J04653 Sequence 5		2.00E-64
2681.seq	446	gi 4503142	Homo sapiens cathepsin D (lysosomal aspartyl protease) (CTSD) mRNA		0
2682.seq	498	gi 5835387	Human mitochondrion, complete genome >gi 1944628 gb J01415.1 HUMMTCCG		0
2684.seq	524	gi 3924668 gb AC004990 AC004990	Homo sapiens PAC clone DJ1185K07 from 7q11.23-q21, complete sequence		0.036
2685.seq	933	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.		E-127
2686.seq	427	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.		4.00E-94
2687.seq	527	gi 5836198 gb AC007382.3 AC007382	Homo sapiens clone NH0288C18, complete sequence		0
2688.seq	559	gi 3548789 gb AC005620 AC005620	Homo sapiens chromosome 19, cosmid R33590, complete sequence [Homo		2.00E-65
2692.seq	391	gi 4512311 dbj AB019441.1 AB019441	Homo sapiens DNA for immunoglobulin heavy-chain variable region, complete		0.1
2693.seq	489	gi 3540173 gb AC005618 AC005618	Homo sapiens chromosome 19, cosmid R31372, complete sequence [Homo		1.00E-14
2694.seq	417	gi 4506690	Homo sapiens ribosomal protein S16 (RPS16) mRNA		0
2695.seq	833	gi 4505604	Homo sapiens pancreatic-associated protein (PAP) mRNA		0
2696.seq	456	gi 505097 dbj D31885 HUMORFKG1N	Human mRNA for KIAA0069 gene, partial cds		0
2698.seq	576	gi 1710280 gb U9297 HSU79297	Human clone 23589 mRNA sequence		0
2699.seq	517	gi 2677628 emb Z82206 HS370M22	Human DNA sequence from PAC 370M22 on chromosome 22q12-qter, contain		0.55
2700.seq	579	gi 4205085 gb U79458 HSU79458	Human WW domain binding protein-2 mRNA, complete cds		0
2701.seq	912	gi 4503538	Homo sapiens eukaryotic translation initiation factor 4 gamma, 2		0
2702.seq	598	gi 5923840 gb AC008125.9 AC008125	Homo sapiens 12 BAC RPC11-25E2 (Roswell Park Cancer Institute Human B		0.16
2704.seq	590	gi 4581311 emb ALD34395.6 HS243L18	Human DNA sequence from clone 243L18 on chromosome 1p36.11-36.23.		0.04
2705.seq	519	gi 4502492	Homo sapiens complement component 1, r subcomponent (C1R) mRNA		0
2706.seq	531	gi 3043681 dbj AB011151 AB011151	Homo sapiens mRNA for KIAA0579 protein, partial cds		7.00E-50
2708.seq	504	gi 3882148 dbj AB018257.1 AB018257	Homo sapiens mRNA for KIAA0714 protein, partial cds		0
2709.seq	491	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.		E-110
2710.seq	552	gi 249370 gb S35960 S35960	laminin receptor homolog [3' region] [Human, mRNA Partial, 739 nt]		0
2711.seq	407	gi 4464208 emb AL035447.3 HS1183I21	Human DNA sequence from clone 1183I21 on chromosome 20q12. Contains a		0.43
2712.seq	507	gi 5835387	Human mitochondrion, complete genome >gi 1944628 gb J01415.1 HUMMTCCG		0
2713.seq	507	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1		0
2715.seq	545	gi 4758509	Homo sapiens H beta 58 homolog (H858) mRNA		0
2717.seq	356				
2719.seq	330	gi 4503162	Homo sapiens cullin 2 (CUL2) mRNA, and translated products		7.00E-95
2721.seq	553	gi 2326514 emb Z85999 HS94G16	Human DNA sequence from PAC 94G16 on chromosome 6q21. Contains		0.15
2722.seq	522	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1		0
2724.seq	129	gi 4758039	Homo sapiens cytochrome c oxidase subunit VIc (COX6C), nuclear gene		2.00E-25
2725.seq	329	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1.		E-163
2727.seq	255	gi 5689536 dbj AB029023.1 AB029023	Homo sapiens mRNA for KIAA1100 protein, complete cds		1.00E-71
2729.seq	344	gi 3201899 gb AF067420 AF067420	Homo sapiens SNC73 protein (SNC73) mRNA, complete cds		E-163

2730.seq	454	gi 45057774	Homo sapiens phosphate carrier, mitochondrial (PHC), splice variant	0
2731.seq	329	gi 409069 dbj J017652 HUMHBL2A	Human mRNA for HbP15/L22, complete cds	E-184
2734.seq	437	gi 306743 gb M97164 HUMFERRH	Human ferritin heavy chain mRNA, complete cds.	E-127
2735.seq	504	gi 3171161 gb AF064866 AF064866	Homo sapiens chromosome 21q22.3 PAC 198E8, complete sequence [Homo	0.53
2736.seq	448	gi 4503304	Homo sapiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	0
2738.seq	551	gi 5835387	Human mitochondrion, complete genome >gi 1944628 gb J01415.1 HUMMTCG	0
2739.seq	255	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	E-105
2740.seq	522	gi 1042081 gb S78569 S78569	laminin alpha 4 chain [human, fetal lung, mRNA, 6204 nt]	0
2741.seq	252	gi 4633837 gb AF139813.1 AF139813	Homo sapiens clone pJ759 12 chromosome 11 map 11q13, complete sequenc	2.00E-85
2743.seq	499	gi 4505604	Homo sapiens parcreatilis-associated protein (PAP) mRNA	0
2744.seq	542	gi 2832592 emb Z95116 HSS57G9	Human DNA sequence from BAC 57G9 on chromosome 22q12.1. Contains EST	0
2748.seq	458	gi 3287468 gb AC004448 AC004448	Homo sapiens chromosome 17, clone HCT187M2, complete sequence [Homo	E-154
2749.seq	548	gi 1944628 gb J01415.1 HUMMTCG	Human mitochondrion, complete genome >gi 1944628 gb J01415.1 HUMMTCG	0
2753.seq	622	gi 31091 emb X16689 HSEF1AC	Human mRNA for elongation factor 1-alpha (clone CEF4)	0
2755.seq	317	gi 441340 emb X72436 HSIGKLV15	H.sapiens mRNA for rearranged Ig kappa light chain variable region	E-130
2756.seq	297	gi 4504278	Homo sapiens H3 histone, family 3A (H3F3A) mRNA	9.00E-88
2757.seq	631	gi 4506606	Homo sapiens ribosomal protein L18 (RPL18) mRNA	0
2758.seq	565	gi 4500166 emb AL049378.1 HSM800163	Homo sapiens mRNA; cDNA DKFZp5861518 (from clone DKFZp5861518)	0
2759.seq	304	gi 4557436	Homo sapiens cell division cycle 20, S.cerevisiae homolog (CDC20)	E-147
2760.seq	497	gi 1430780 emb Z75889 HS267P19	Human DNA sequence from cosmid 267P19, BRCA2 gene region chromosome	0.034
2761.seq	487	gi 3688349 emb AL030996.1 HS1189B24	Human DNA sequence from clone 1189B24 on chromosome Xq25-26.3. Conta	7.00E-93
2762.seq	323	gi 4165005 gb AC005912 AC005912	Homo sapiens 12p13.3 BAC RPC11-543P15 (Roswell Park Cancer Institute	E-179
2763.seq	437	gi 4504900	Homo sapiens karyopherin alpha 4 (Importin alpha 3) (KPNA4) mRNA	0
2764.seq	569	gi 1262342 emb X93334 HSMITG	H.sapiens mitochondrial DNA, complete genome	0
2765.seq	823	gi 3659494 gb AC005332 AC005332	Homo sapiens chromosome 17, clone hRPK.147.L_13, complete sequence	1.00E-39
2766.seq	244			
2767.seq	561	gi 311374 emb X57352 HS18U	Human 1-8U gene from Interferon-inducible gene family	0
2768.seq	301	gi 1213466 gb U29057 HSU29057	Human MHC class I antigen HLA-B*7 variant (HLA-B) mRNA, complete	E-145
2769.seq	631	gi 4210446 dbj AB022785.1 AB022785	Homo sapiens ASH2L gene, complete cds, similar to Drosophila ash2 gene	0.17
2770.seq	229	gi 23690 emb X56932 HS23KDHP	H.sapiens mRNA for 23 kD highly basic protein	E-120
2771.seq	531	gi 5730117	Homo sapiens X-prolyl aminopeptidase (aminopeptidase P)-like	0
2772.seq	479	gi 4506700	Homo sapiens ribosomal protein S23 (RPS23) mRNA	0
2773.seq	494	gi 3694657 gb AC005021 AC005021	Homo sapiens BAC clone GS293C05 from 7q21-q22, complete sequence [Horr	0.13
2774.seq	396	gi 3659494 gb AC005332 AC005332	Homo sapiens chromosome 17, clone hRPK.147.L_13, complete sequence	0.007
2775.seq	890	gi 4758913	Homo sapiens LPS-induced TNF-alpha factor (PIG7) mRNA	0
2776.seq	341	gi 4210952 gb AF092907 HSHUMRPS2	Homo sapiens ribosomal protein S19 (RPS19) gene, exons 3, 4 and 5 and	2.00E-82
2777.seq	565	gi 4586074 gb AC006285.9 AC006285	, complete sequence [Homo sapiens]	0.039
2778.seq	963	gi 13683 emb V00710 MIT1HS	Human mitochondrial genes for several tRNAs (Phe, Val, Leu) and 12S	E-179
2779.seq	533	gi 4557756	Homo sapiens mutL homolog 1 (MLH1) mRNA >gi 463988 gb U07343 HSU0734	0
2781.seq	841	gi 4508141 gb AC006017.2 AC006017	Homo sapiens PAC clone DJ0981007 from 7q34-q36, complete sequence	0.058
2782.seq	283	gi 6012990 emb AL121735.1 HS224A62E	Isoform of human GTP-binding protein G25K	8.00E-48
2783.seq	840	gi 249370 gb S35960 S35960	laminin receptor homolog (3' region) [human, mRNA Partial, 739 nt]	0
2784.seq	438	gi 3338395 gb M81355 HUMSPHINO	Homo sapiens sphingolipid activator proteins 1 and 2 processed mutant	0
2785.seq	152	gi 3650045 gb AC005544 AC005544	Homo sapiens chromosome 17, clone hRPK.349.A_8, complete sequence (Horr	2.00E-59
2787.seq	498	gi 298485 gb S56985 S56985	Ribosomal protein L19 [human, breast cancer cell line, MCF-7, mRNA,	0
2788.seq	497	gi 4097713 gb U67734 HSU67734	Human cPLA2 interacting protein mRNA, complete cds	0
2790.seq	469	gi 13683 emb V00710 MIT1HS	Human mitochondrial genes for several tRNAs (Phe, Val, Leu) and 12S	0
2791.seq	490	gi 3954888 emb AJ010444 HSA010444	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RHD,	0
2792.seq	521	gi 1136401 dbj D79993 D79993	Human mRNA for KIAA0171 gene, complete cds	0
2794.seq	378	gi 4501882	Homo sapiens actin alpha 2 (ACTA2) mRNA >gi 28329 emb X13839 HSACTA	0
2795.seq	453	gi 3702296 gb AC005782 AC005782	Homo sapiens chromosome 19, cosmid R31855, complete sequence [Homo	5.00E-81
2796.seq	517	gi 4758761	Homo sapiens asparaginyl-tRNA synthetase (NARS) mRNA	0
2797.seq	510	gi 532674 dbj D10758 HUMHC3	Human gene for proteasome HC3, 5'-flanking region	6.00E-47
2798.seq	285	gi 4505036	Homo sapiens latent transforming growth factor beta binding protein 4	E-145
2799.seq	511	gi 179274 gb M37104 HUMATPSY	Human mitochondrial ATPase coupling factor 6 subunit (ATP6A) mRNA.	0
2800.seq	388	gi 5174602	Homo sapiens myosin regulatory light chain 2, smooth muscle isoform	0
2801.seq	487	gi 187366 gb M69126 HUMMAOB07	Human monoamine oxidase B (MAOB) gene, exon 7.	0.52
2802.seq	136	gi 115987 emb AL021528 HS394P21	Homo sapiens DNA sequence from clone 394P21 on chromosome 1p36.12-36.	0.13
2803.seq	192	gi 4995304 emb AJ242829.1 HSA242829	Homo sapiens mRNA for PQBP-1 protein	E-100
2805.seq	393	gi 5031644	Homo sapiens human homolog of yeast mitochondrial copper	0
2806.seq	171	gi 184090 gb M11354 HUMHISH3B	Human H3.3 histone, class B mRNA, complete cds.	6.00E-35
2807.seq	383	gi 3171028 gb AF062278 AF062278	Homo sapiens clone Xu-50 immunoglobulin heavy chain variable region	E-139
2808.seq	373	gi 3845417 gb AC004837 AC004837	Homo sapiens PAC clone DJ0570D02 from 7p13-p14, complete sequence [Hor	8.00E-24
2810.seq	316	gi 3150276 gb AC004106 AC004106	Homo sapiens Xp22 BAC GSHB-45N23 (Genome Systems Human BAC library	E-153
2811.seq	342	gi 4503304	Homo sapiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	0
2813.seq	395	gi 2961444 gb AC003109 AC003109	Human DNA from overlapping chromosome 7 PAC and P1 clones containing	0.41
2817.seq	394	gi 5532410 gb AF138300.1 AF138300	Homo sapiens decorin variant A mRNA, complete cds	0
2818.seq	328	gi 404194 emb X75159 HSTEF2PG	H.sapiens DNA for transcriptional elongation factor TFIIS	1.3
2820.seq	205	gi 4506682	Homo sapiens ribosomal protein S12 (RPS12) mRNA	E-102
2825.seq	351	gi 2735714 gb AF004877 AF004877	Homo sapiens pro-alpha 2(I) collagen (COL1A2) gene, complete cds	1.00E-74
2826.seq	444	gi 5803112	Homo sapiens transmembrane protein (63kD), endoplasmic	E-136
2828.seq	406	gi 5230675 gb AF159570.1 AF159570	Homo sapiens regulator of G-protein signalling 5 (RGSS5) mRNA.	E-102
2830.seq	434	gi 3377602 gb AF075061 HUMYP07G10	Homo sapiens full length insert cDNA YP07G10	0
2831.seq	376	gi 4508110 gb AC005076.2 AC005076	Homo sapiens BAC clone RG227L24 from 7q21.1-q21.2, complete sequence	4.00E-72
2833.seq	139	gi 2358019 gb AE006658 HUA006658	Homo sapiens T-cell receptor alpha delta locus from bases 1 to 250529	0.52
2834.seq	152	gi 182904 gb M93036 HUMGA7A08	Human (clone 21726) carcinoma-associated antigen GA733-2 (GA733-2)	9.00E-71
2835.seq	499	gi 4506688	Homo sapiens ribosomal protein, large, P1 (RPLP1) mRNA	0
2836.seq	475	gi 1930412 gb L81394 HSL81394	Homo sapiens (subclone 1_g5 from P1 H39) DNA sequence	E-124
2837.seq	528	gi 1590807 dbj D87914 D87914	Human mRNA for ornithine decarboxylase antizyme, complete cds	0
2838.seq	413	gi 3954884 emb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RHD,	0
2839.seq	351			
2842.seq	275	gi 4503304	Homo sapiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	E-108
2843.seq	348	gi 1620014 dbj D87665 D87665	Human heart mRNA for MHC class I HLA-Bw62, partial cds	E-176
2845.seq	394	gi 557545 gb M28699 HUMB23A	Homo sapiens nucleolar phosphoprotein B23 (NPM1) mRNA, complete cds.	0
2846.seq	87	gi 36540 emb X65965 HSSOD2G	H.sapiens SOD-2 gene for manganese superoxide dismutase	1.00E-12
2848.seq	476	gi 4096058 gb AC006274 AC006274	Homo sapiens chromosome 19, cosmid R29335, complete sequence [Homo	0.13
2849.seq	321	gi 3763941 gb AC005826 AC005826	Homo sapiens clone UWGC:rg041a03 from 7p14-15, complete sequence [Horr	0.33
2850.seq	381	gi 4884375 emb AL050161.1 HSM800459	Homo sapiens mRNA; cDNA DKFZp58680222 (from clone DKFZp58680222)	E-167
2851.seq	389	gi 4914513 emb AL022320.23 HS199H16	Human DNA sequence from clone 199H16 on chromosome 22q12-13. Contains	1.6
2852.seq	421	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	E-100
2858.seq	171	gi 3153101 emb AJ229041 HS229041	Homo sapiens 959 kb contig between AML1 and CBR1 on chromosome 21q22.	0.17
2859.seq	443	gi 4503954	Homo sapiens GCN5 (general control of amino-acid synthesis, yeast,	0
2860.seq	426	gi 13683 emb V00710 MIT1HS	Human mitochondrial genes for several tRNAs (Phe, Val, Leu) and 12S	0
2862.seq	537	gi 5306288 gb AC007938.1 AC007938	Homo sapiens clone UWGC:djs201 from 7q31, complete sequence	0.14
2865.seq	493	gi 3127145 gb AF061298 HSSUR2G15	Homo sapiens sulfonylurea receptor (SUR2) gene, exon 9	2.1
2866.seq	245	gi 3483378 gb AF086033 HUMY26E09	Homo sapiens full length insert cDNA clone YW26E09	6.00E-39
2868.seq	498	gi 5360122 gb AF155114.1 AF155114	Homo sapiens NY-REN-57 antigen mRNA, partial cds	0
2869.seq	174			
2870.seq	494	gi 4502692	Homo sapiens CD9 antigen (p24) (CD9) mRNA	1.00E-94
2872.seq	420	gi 4165210 emb AL021937 HS149A16	Human DNA sequence from clone 149A16 on chromosome 22q12-13. Contain	1.7
2873.seq	423	gi 2282011 gb AC002390 AC002390	Human DNA from overlapping chromosome 19-specific cosmids R30072 and	E-105
2874.seq	575	gi 36405 emb X00457 HSSBA1	Human mRNA for SB classII histocompatibility antigen alpha-chain	0
2875.seq	520	gi 5032050	Homo sapiens ribosomal protein S14 (RPS14) mRNA	0
2876.seq	506	gi 5919146 gb AF182289.1 AF182289	Homo sapiens U6 snRNA-associated Sm-like protein LSm3 mRNA.	0
2877.seq	529	gi 1514580 emb X95747 HSIGKLC5	H.sapiens immunoglobulin kappa chain (clone: 5)	0
2878.seq	489	gi 5729833	Homo sapiens ankyrin repeat-containing protein (G9A) mRNA	6.00E-07
2879.seq	495	gi 4506744	Homo sapiens ribosomal protein S9 (RPS9) mRNA	0
2880.seq	496	gi 4826450 emb Z93241.1 HS222E13	Human DNA sequence from clone 222E13 on chromosome 22. Contains three	0.53

2881.seq	479	gi 506336 dbj D16562 HUMATP SGL	Human mRNA for ATP synthase gamma-subunit (L-type), complete cds	0
2882.seq	481	gi 185287 gb M34024 HUMIGHCXE	Human fetal Ig heavy chain variable region (clone M43) mRNA,	E-136
2883.seq	488	gi 4506678	Homo sapiens ribosomal protein S10 (RPS10) mRNA	0
2884.seq	533	gi 31031 emb Z14182 HSE554A1	H.sapiens rearranged Ig heavy chain variable region (VDJ)	E-152
2886.seq	486	gi 3947784 emb AL031229 HS15D7	Human DNA sequence from clone 15D7 on chromosome 6p22.1-22.3. Contains	0.13
2887.seq	360	gi 5453739	Homo sapiens myosin, light polypeptide, regulatory, non-sarcomeric	E-147
2889.seq	558	gi 6005702	Homo sapiens mayven (ABP-KELCH), mRNA	0
2890.seq	444	gi 4506622	Homo sapiens ribosomal protein L27 (RPL27) mRNA	0
2891.seq	291	gi 4519939 dbj AB019524.1 AB019524	Homo sapiens mRNA for nuclear receptor co-repressor, partial cds	E-124
2894.seq	376	gi 5263001 emb AL035422.12 HS164F3	Human DNA sequence from clone 164F3 on chromosome Xq21.33-23 Contains	E-155
2895.seq	308	gi 3413879 dbj AB007928 AB007928	Homo sapiens mRNA for KIAA0459 protein, partial cds	E-102
2896.seq	243	gi 6013067 emb AL078621.19 HSA395L1	Human DNA sequence from clone 395L14 on chromosome 22, complete	3.00E-87
2897.seq	152	gi 407696 gb L25080.1 HUMRH0AA	Homo sapiens GTP-binding protein (rhoA) mRNA, complete cds	5.00E-20
2898.seq	514	gi 5016088	Homo sapiens actin, beta (ACTB) mRNA	0
2899.seq	441	gi 5835387	Human mitochondrion, complete genome >gi 1944628 gb J01415.1 HUMMTCC	0
2900.seq	479	gi 2058319 emb Z84495 HSLUCA8	Human DNA sequence from cosmid LUCA8 on chromosome 3p21.3 contains	0.51
2901.seq	267	gi 3483804 gb AF086459 HJMZD84G09	Homo sapiens full length insert cDNA clone ZD84G09	E-125
2902.seq	384	gi 4507516	Homo sapiens tight junction protein 1 (zona occludens 1) (TJP1) mRNA	0
2904.seq	538	gi 4505896	Homo sapiens plastin 1 (I isoform) (PLS1) mRNA	0
2906.seq	450	gi 4557808	Homo sapiens ornithine aminotransferase (OAT), nuclear gene encoding	0
2908.seq	525	gi 6094661 gb AC005225.2 AC005225	Homo sapiens clone DJ0414A15, complete sequence	0.56
2909.seq	444	gi 4505604	Homo sapiens pancreatitis-associated protein (PAP) mRNA	0
2910.seq	558	gi 32795 emb X61015 HSIG479	Human immunoglobulin VH mRNA (479 bp)	E-134
2914.seq	97	gi 4469068 emb AL031319.5 HS36411	Human DNA sequence from clone 36411 on chromosome Xq23. Contains a	1.4
2915.seq	406	gi 6001959 gb AC008116.8 AC008116	Homo sapiens 3q25-26 BAC C11B-187P1 (California Institute of Technology	1.00E-75
2916.seq	497	gi 4504766	Homo sapiens integrin, beta 1 (fibronectin receptor, beta	0
2917.seq	513	gi 5106993 gb AF125099.1 AF125099	Homo sapiens HSPC38 protein mRNA, complete cds	0
2918.seq	511	gi 4506666	Homo sapiens ribosomal protein, large, PO (RPLPO) mRNA	0
2921.seq	312	gi 3005703 gb AF054990 AF054990	Homo sapiens clone Z3884 mRNA sequence	E-172
2922.seq	500	gi 2984582 gb AC004472 AC004472	Homo sapiens chromosome 9, P1 clone 11659, complete sequence [Homo	E-135
2927.seq	389	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
2928.seq	533	gi 184743 gb J00220 HUMIGCC8	Homo sapiens immunoglobulin alpha-1 heavy chain constant region	0
2929.seq	250	gi 3947808 emb AL031224.1 HS336H9	Human DNA sequence from clone 336H9 on chromosome 6p12.1-21.1. Contain	1
2930.seq	483	gi 4506750	Homo sapiens restin (Reed-Steinberg cell-expressed intermediate	0
2931.seq	531	gi 4689139 gb AF077046.1 AF077046	Homo sapiens ganglioside expression factor 2 homolog mRNA, complete	0
2932.seq	533	gi 4503832	Homo sapiens frizzled (Drosophila) homolog 7 (FZD7) mRNA	0
2933.seq	374	gi 188231 gb J00194 HUMMHDR	human hla-dr antigen alpha-chain mRNA & lvs fragments.	0
2934.seq	107	gi 4507644	Homo sapiens triosephosphate isomerase 1 (TP1) mRNA	3.00E-51
2937.seq	507	gi 5835387	Human mitochondrion, complete genome >gi 1944628 gb J01415.1 HUMMTCC	0
2938.seq	225	gi 1669374 gb AC000062 HUM2G3A	Human PAC clone 2G3A from 13q12-13q13, complete sequence [Homo sapier	0.9
2939.seq	509	gi 2580477 gb AC002992.1 AC002992	Homo sapiens chromosome Y, clone 203M13, complete sequence	0.009
2940.seq	412	gi 3201899 gb AF064720 AF064720	Homo sapiens SINC73 protein (SINC73) mRNA, complete cds	3.00E-76
2942.seq	532	gi 2995179 emb Z99297.1 HS262D12	Homo sapiens DNA sequence from PAC 262D12 on chromosome 1q23.3-24.3.	0.14
2943.seq	501	gi 3954884 emb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	E-180
2944.seq	467	gi 5817134 emb AL110212.1 HSM80086	Homo sapiens mRNA; cDNA DKFZp664D173 (from clone DKFZp664D173)	7.00E-62
2945.seq	537	gi 3420065 gb AC005490 AC005490	Homo sapiens UWGC.g1397a051 from T7p14-15, complete sequence [Homo	0.15
2946.seq	482	gi 2224636 dbj AB002346 AB002346	Human mRNA for KIAA0348 gene, complete cds	E-141
2947.seq	534	gi 4926891 gb AC007179.3 AC007179	Homo sapiens clone NH0569H17, complete sequence	0.57
2948.seq	570	gi 3483458 gb AF086113 HJMZA06D10	Homo sapiens full length insert cDNA clone ZA06D10	0
2949.seq	498	gi 3810670 gb AC005495 AC005495	Homo sapiens chromosome 17, clone hRPK.293_K_20, complete sequence	0.13
2950.seq	513	gi 1619291 emb Z80902 HSN80H12	Human DNA sequence from cosmid N80H12 on chromosome 22a12-qter cont	0.009
2951.seq	519	gi 1136389 dbj D79986 D79986	Human mRNA for KIAA0164 gene, complete cds	0
2952.seq	528	gi 4501888	Homo sapiens actin, gamma 2, smooth muscle, enteric (ACTG2) mRNA	0
2953.seq	421	gi 6102866 gb AC006449.19 AC006449	Homo sapiens chromosome 17, clone HCIT.58_E_17, complete sequence	0.11
2954.seq	343	gi 5725210 emb AL109758.1 CNS0180V	Human chromosome 14 DNA sequence *** IN PROGRESS *** BAC R-873E2	2.00E-45
2955.seq	490	gi 5708492 gb AC007875.2 AC007875	Homo sapiens clone DJ1145A22, complete sequence	1.00E-04
2956.seq	575	gi 4557494	Homo sapiens cleavage stimulation factor, 3' pre-RNA, subunit 3, 77kD	0
2957.seq	515	gi 5523841 gb AC005280.2 AC005280	Homo sapiens clone J0240K06, complete sequence	0.035
2958.seq	541	gi 3172145 gb AC004782 AC004782	Homo sapiens chromosome 5, BAC clone 205e20 (LBNL H170), complete	4.00E-54
2962.seq	537	gi 307150 gb L19182 HUMMAC25X	Human MAC25 mRNA, complete cds	E-157
2965.seq	523	gi 185845 gb M27025 HUMIGKAI	Human Ig active kappa chain mRNA V-region (V-J1-C), clone NOV.	E-163
2966.seq	523	gi 5453739	Homo sapiens myosin, light polypeptide, regulatory, non-sarcomeric	0
2967.seq	99	gi 4508122 gb AC006379.2 AC006379	Homo sapiens BAC clone NH0456N16 from 7p15.1-p13, complete sequence	0.089
2968.seq	291			
2970.seq	474	gi 4506648	Homo sapiens ribosomal protein L3 (RPL3) mRNA	0
2971.seq	509	gi 2827470 emb Z97205 HS155D22	Human DNA sequence from PAC 155D22 on chromosome 6q27. Contains EST	0.14
2973.seq	530	gi 6049285 gb AF163570.1 AF163570	Homo sapiens DINB protein (DINB1) mRNA, complete cds	8.00E-25
2974.seq	218	gi 4587122 dbj AB019392.1 AB019392	Homo sapiens mRNA of muscle specific gene M9, complete cds	E-115
2975.seq	454	gi 5680371 gb AF158555.1 AF158555	Homo sapiens glutaminase C mRNA, complete cds	E-137
2976.seq	520	gi 4506728	Homo sapiens ribosomal protein S5 (RPS5) mRNA	0
2977.seq	480	gi 184743 gb J00220 HUMIGCC8	Homo sapiens immunoglobulin alpha-1 heavy chain constant region	0
2979.seq	501	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-162
2980.seq	452	gi 4378110 emb AL021578.1 HS453C12	Human DNA sequence from clone 453C12 on chromosome 20q12-13.12, cont	4.00E-66
2982.seq	413	gi 34284 emb X06537.1 HSLCAT1	Human mRNA for lecithin:cholesterol acyltransferase (LCAT): EC	8.00E-49
2983.seq	481	gi 23711 emb X16850 HS241D5	Human mRNA for myoblast cell surface antigen 24.1D5	0
2984.seq	506	gi 4996607 dbj AB022435.1 AB022435	Homo sapiens LG mRNA for E2 ubiquitin-conjugating enzyme, complete	0.54
2985.seq	282	gi 4504618	Homo sapiens insulin-like growth factor binding protein 7 (IGFBP7)	E-107
2986.seq	191	gi 4206702 gb AF082569 AF082569	Homo sapiens D-type cyclin-interacting protein 1 (DIP1) mRNA.	2.00E-76
2987.seq	489	gi 3646117 emb AL031075 HS789O11	Human DNA sequence from clone 789O11 on chromosome Xq25. Contains par	0.53
2988.seq	469	gi 3954884 emb AJ010442 HSA010442	Homo sapiens ribosomal protein L30 mRNA, complete cds	0
2994.seq	489	gi 2529399 gb AC002398 AC002398	Human DNA from chromosome 19-specific cosmid F25965, genomic sequenc	2.00E-53
2995.seq	358	gi 4506646	Homo sapiens ribosomal protein L39 (RPL39) mRNA	0
2997.seq	510	gi 1917006 gb U91903 HSU91903	Human Fritz mRNA, complete cds	0
2999.seq	512	gi 2281039 emb Z95328 HS447B16	Human DNA sequence from PAC 447B16 on chromosome Xq13.1-Xq13.3	2.2
3000.seq	230	gi 3954884 emb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	E-126
3001.seq	150	gi 6103356 gb AF115570.1 HSSMTNS20	Homo sapiens smoothelin (SMTN) gene, exon 20, complete sequence;	4.00E-11
3003.seq	530	gi 1160964 gb L40933 HUMPGMR	Homo sapiens phosphoglucomutase-related protein (PGMRP) gene,	0
3005.seq	203	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	1.00E-85
3006.seq	170	gi 530074 gb U12693 HSU12693	Human Hsa5 mitochondrial cytochrome oxidase subunit II (COII) gene.	8.00E-13
3007.seq	489	gi 4507170	Homo sapiens secreted protein, acidic, cysteine-rich (osteonectin)	0
3008.seq	476	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
3011.seq	366	gi 5174446	Homo sapiens guanine nucleotide binding protein (G protein), beta	0
3014.seq	799	gi 184420 gb M18660 HUMHSP90	Human 90-kDa heat-shock protein gene, cDNA, complete cds.	0
3016.seq	507	gi 5354053 gb AF140360.1 AF140360	Homo sapiens histone acetyltransferase (HBOA) mRNA, complete cds	0
3017.seq	411	gi 4240123 dbj AB020626.1 AB020626	Homo sapiens mRNA for KIAA0819 protein, partial cds	0
3020.seq	506	gi 3170970 gb AF062252 AF062252	Homo sapiens clone Xu-7 Immunoglobulin heavy chain variable region	E-125
3021.seq	196	gi 4758873	Homo sapiens 76 kDa membrane protein (P76) mRNA	4.00E-55
3022.seq	438	gi 732797 emb Z48054 HSP T51R	H.sapiens mRNA for peroxisomal targeting signal 1 (SKL type) receptor	0
3023.seq	554	gi 4038573 emb AL031313 HS581F12	Human DNA sequence from clone 581F12 on chromosome Xq21. Contains	0.038
3024.seq	506	gi 35456 emb X54629 HSPHL1	Human pHL-1 gene is a c-myc oncogene containing cox11 sequence	0
3025.seq	381	gi 2055423 gb U94728 HSCALM2504	Homo sapiens calmodulin (CALM2) gene, exons 3-6, and complete cds	0
3026.seq	518	gi 450780 gb U02570 HSU02570	Human CDC42 GTPase-activating protein mRNA, partial cds.	3.00E-86
3027.seq	501	gi 5016088	Homo sapiens actin, beta (ACTB) mRNA	0
3028.seq	198	gi 4808226 emb AL049176.3 HS141H5	Human DNA sequence from clone 141H5 on chromosome Xq22.1-23. Contains	8.00E-04
3029.seq	519	gi 5578939 emb AL035417.15 HS891H21	Human DNA sequence from clone 891H21 on chromosome 1p32.2-34.2 Cont	0.14
3030.seq	420	gi 4502268	Homo sapiens ATPase, Na+/K+ transporting, alpha 1 polypeptide	0
3031.seq	464	gi 3046309 gb AC004547 AC004547	Homo sapiens PAC clone DJ0982C16 from 7q11.23-q21.1, complete sequenc	0.13
3032.seq	494	gi 4504850	Homo sapiens potassium channel, subfamily K, member 5 (TASK-2)	E-128

3033.seq	411	gi 36135 emb Z12962 HSRPL41	H.sapiens mRNA for homologue to yeast ribosomal protein L41	0
3034.seq	335	gi 31107 emb Z11692 HSEF2MR	H.sapiens mRNA for elongation factor 2	0
3035.seq	501	gi 4726099 gb AC006254.10 AC006254	Homo sapiens 3p21.1-9 PAC RPC15-966M1 (Roswell Park Cancer Institute)	1.00E-04
3036.seq	489	gi 189600 gb M84337 HUMANSECP	Human pancreatitis associated protein mRNA, complete cds.	0
3037.seq	505	gi 3702283 gb AC005788 AC005788	Homo sapiens chromosome 19, cosmid R26652, complete sequence [Homo	2.1
3039.seq	503	gi 1469164 dbj D5091.1 D5091.1	Human mRNA for KIAA0121 gene, complete cds	0
3041.seq	489	gi 3548788 gb AC005621 AC005621	Homo sapiens chromosome 19, cosmid F19678, complete sequence [Homo	0
3043.seq	392	gi 4153878 gb AC004819 AC004819	Homo sapiens PAC clone DJ0015123 from 22, complete sequence [Homo	0.11
3044.seq	502	gi 4506602	Homo sapiens ribosomal protein L15 (RPL15) mRNA	0
3045.seq	451	gi 28403 emb X04299 HSADH3G2	Human mRNA for liver alcohol dehydrogenase (EC 1.1.1.1) gamma 2	0
3046.seq	493	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	0
3047.seq	361	gi 6016927 emb Z84465.5 HS209H1	Human DNA sequence from clone 209H1 on chromosome Xq25-26, complete	3.00E-08
3048.seq	480	gi 4191326 gb AF065391 AF065391	Homo sapiens ZIS1 mRNA, complete cds	0
3049.seq	491	gi 265703 gb S55735 S55735	Ig alpha 1 - alpha 2m-immunoglobulin A1-A2 lambda hybrid GAU heavy	0
3050.seq	499	gi 4588523 gb AF109196.1 AF109196	Homo sapiens Intracellular chloride channel p64H1 mRNA, complete	0
3051.seq	510	gi 4506740	Homo sapiens ribosomal protein S7 (RPS7) mRNA	0
3052.seq	506	gi 598784 dbj D17049 HUMD3G05M5	Human HepG2 partial cDNA, clone hmd3g05m5	E-144
3053.seq	841	gi 4761681 gb AF147330.1 AF147330	Homo sapiens full length insert cDNA clone YB24C12	0
3054.seq	474			
3055.seq	435	gi 33217 emb X67858 HSIGKA	H.sapiens DNA for IG kappa JC region	0
3056.seq	255	gi 1008967 emb Z47034 HSB98D12E	Human partial cDNA sequence, complete sequence of clone 98D12.	E-126
3057.seq	510	gi 5102744 emb AL079283.1 HS1000202	Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 39515	E-179
3058.seq	486	gi 4506718	Homo sapiens ribosomal protein S2 (RPS2) mRNA	0
3060.seq	353	gi 598138 gb L38287 HUMADH2505	Homo sapiens alcohol dehydrogenase beta-3 subunit (ADH2-3) gene.	1.00E-25
3061.seq	517	gi 4502146	Homo sapiens amyloid beta (A4) precursor-like protein 2 (APLP2) mRNA	0
3062.seq	493	gi 4506588	Homo sapiens RNA 3'-terminal phosphate cyclase (RPC) mRNA	E-140
3063.seq	367	gi 2995603 gb AC004231 AC004231	Homo sapiens chromosome 17, clone hRPC.1110 E_20, complete sequence	0.38
3066.seq	471	gi 6006017	Homo sapiens lectin, galactoside-binding, soluble, 4 (galectin 4)	0
3067.seq	539	gi 5306301 gb AC007560.3 AC007560	Homo sapiens clone NH0386G20, complete sequence	E-172
3069.seq	501	gi 5802965	Homo sapiens desrin (actin depolymerizing factor) (ADF), mRNA	0
3070.seq	477	gi 1136403 dbj D79994 D79994	Human mRNA for KIAA0172 gene, partial cds	2
3071.seq	485	gi 5733683 gb AF166330.1 AF166330	Homo sapiens stratum corneum chymotryptic enzyme gene, complete cds	2.1
3072.seq	495	gi 5306220 gb AC007564.9 AC007564	Homo sapiens 12q22 BAC RPC11-513P18 (Roswell Park Cancer Institute	0.53
3073.seq	488	gi 4758901	Homo sapiens 6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase 4	0
3074.seq	481	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
3075.seq	497			
3076.seq	516	gi 1196441 gb M35368 HUMMAC2	Human macrophage galactose-specific lectin (hMAC-2) mRNA, complete	0
3078.seq	492	gi 4759301	Homo sapiens VAMP (vesicle-associated membrane protein)-associated	2.1
3080.seq	480	gi 2828786 gb AC004111 AC004111	Homo sapiens BAC clone RG103H13 from 7q31, complete sequence [Homo	0.008
3081.seq	492	gi 5803112	Homo sapiens transmembrane protein (63kD), endoplasmic	0
3083.seq	346	gi 3242759 gb AC005156 AC005156	Homo sapiens PAC clone DJ1099C19 from 7q21-q22, complete sequence [Hor	0
3084.seq	500	gi 4503048	Homo sapiens cysteine-rich protein 2 (CRIP2) mRNA	0
3085.seq	367	gi 1322149 emb Z73358 HS65O19	Human DNA sequence from cosmid 65O19, BRC42 gene region chromosome	0.39
3086.seq	512	gi 4503592	Homo sapiens epidermal growth factor receptor pathway substrate 15	0
3089.seq	423	gi 1488611 emb X51985 HSLAG3	Human LAG-3 mRNA for CD4-related protein involved in lymphocyte	E-120
3090.seq	513	gi 4572584 emb AL032821.2 HS55C23	Human DNA sequence from clone 55C23 on chromosome 6q22.2-23.3 contain	0
3092.seq	527	gi 6048963 gb AF103798.1 AF103798	Homo sapiens clone C40 unknown mRNA	4.00E-42
3093.seq	494	gi 5106790 gb AF083248.1 AF083248	Homo sapiens ribosomal protein L26 homolog mRNA, complete cds	0
3094.seq	509	gi 1944628 gb J01415.1 HUMMTCG	Human mitochondrion, complete genome	0
3095.seq	475	gi 57234 emb X16359 RNSEPI3	Rat mRNA for SPI-3 serine protease inhibitor	0.13
3097.seq	473	gi 2804235 dbj D78195 D78195	Homo sapiens J-chain gene promoter region	3.00E-70
3098.seq	487	gi 218405 dbj D12918 YSCBEL2	Yeast BEL2 gene for ORF, complete cds.	0.52
3099.seq	469	gi 495865 gb L23982 HUMCOL7A1X	Homo sapiens (clones: CW52-2, CW27-6, CW15-2, CW26-5, 11-67) collagen	0.032
3100.seq	495	gi 2865235 gb U89705 U89705	Uroleptus gallina macronuclear DNA polymerase alpha gene, partial	2.1
3101.seq	479	gi 1915966 gb U51678 HSU51678	Homo sapiens small acidic protein mRNA, complete cds	0
3102.seq	532	gi 4506886	Homo sapiens selenoprotein W, 1 (SEPW1) mRNA	0
3103.seq	518	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-118
3104.seq	500	gi 249370 gb S35960 S35960	tannin receptor homolog (3' region) (human, mRNA Partial, 739 nt)	0
3105.seq	416	gi 5803004	Homo sapiens hepatitis delta antigen-interacting protein A (DIPA),	0
3106.seq	404	gi 3483426 gb AF086081 HUMYZ83C11	Homo sapiens full length insert cDNA clone YZ83C11	4.00E-26
3107.seq	476	gi 5901925	Homo sapiens pre-mRNA cleavage factor I _m (25kD) (CFIM25), mRNA	0
3108.seq	375	gi 4888411 emb AL050091.1 HSM80016C	Homo sapiens mRNA: cDNA DKFZp586F1918 (from clone DKFZp586F1918)	0
3110.seq	272	gi 1679960 gb S42658 S42658	S3 ribosomal protein [human, colon, mRNA, 826 nt]	E-149
3111.seq	519	gi 3256174 emb AL022069.1 HS168L15	Human DNA sequence from clone 168L15 on chromosome 6q26-27 Contains p	6.00E-78
3112.seq	490	gi 30371 emb X09607 HSCYSTCR	Human mRNA for cysteine proteinase inhibitor precursor cystatin C	0
3114.seq	372	gi 4508118 gb AC005041.2 AC005041	Homo sapiens clone NH0523H20, complete sequence	4.00E-04
3115.seq	528	gi 5725511 gb AF072097.1 AF072097	Homo sapiens beta-2 microglobulin gene, complete cds	0
3116.seq	449			
3117.seq	393	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	E-113
3118.seq	503	gi 4680716 gb AF132973.1 AF132973	Homo sapiens CGI-39 protein mRNA, complete cds	0
3119.seq	534	gi 33330 emb X56510 HSIGKVLU	Human mRNA for immunoglobulin variable kappa region	E-175
3120.seq	437	gi 3152704 gb AF065414 AF065414	Homo sapiens COBW-like placental protein mRNA, partial cds	5.00E-44
3121.seq	487	gi 4505604	Homo sapiens pancreatitis-associated protein (PAP) mRNA	0
3122.seq	421	gi 4753259 gb AC006323.3 AC006323	Homo sapiens PAC clone DJ1151M05 from 7q35-q36, complete sequence	0
3123.seq	461	gi 4506686	Homo sapiens ribosomal protein S15 (RPS15) mRNA	0
3127.seq	535	gi 6010168 emb AL078596.8 HSDJ429C4	Human DNA sequence from clone 429G5 on chromosome 6q21-22.1, complete	0
3128.seq	483	gi 2935595 gb AC003036 AC003036	Homo sapiens Xp22 PAC RPC11-10B6 (Roswell Park Cancer Center PAC	0.033
3129.seq	485	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-140
3130.seq	491	gi 4753230 gb AC007001.2 AC007001	Homo sapiens BAC clone NH0486P11 from 7p21-p15.1, complete sequence	0
3131.seq	378	gi 4505270	Homo sapiens metallothionein 1L (MT1L) mRNA	E-157
3132.seq	428	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	1.00E-93
3134.seq	499	gi 4902741 emb AL049548.6 HSDJ398G3	Human DNA sequence from clone 398G3 on chromosome 6q25.1-25.3. Contain	2.1
3135.seq	516	gi 4071022 gb AC006236 AC006236	Homo sapiens chromosome 17, clone hCIT.162 E_12, complete sequence [Homo	0.036
3136.seq	480	gi 4504164	Homo sapiens gelsolin (amyloidosis, Finnish type) (GSN) mRNA	0
3138.seq	404	gi 31914 emb X04408 HSGSA2R	Human mRNA for coupling protein G(s) alpha subunit (alpha-S2)	0
3139.seq	288	gi 476704 gb L26336 HUMHSPA2A	Human heat shock protein HSPA2 gene, complete cds.	E-133
3141.seq	491	gi 4757817	Homo sapiens vacuolar H(+) -ATPase subunit (ATP6G) mRNA	0
3142.seq	215			
3143.seq	484	gi 1167502 dbj D50525 HUMTI227HC	Human mRNA for TI-227H	0
3145.seq	524	gi 5531840 gb AF078863.1 AF078863	Homo sapiens PTD010 mRNA, complete cds	0
3146.seq	518	gi 6005763	Homo sapiens GABA(A) receptor-associated protein (GABARAP), mRNA	0
3147.seq	362	gi 4165210 emb AL021937 HS149A16	Human DNA sequence from clone 149A16 on chromosome 22q12-13. Contains	2.00E-39
3148.seq	508	gi 3983571 gb AC004839 AC004839	Homo sapiens PAC clone DJ0593H12 from 7p31, complete sequence [Homo	5.00E-63
3151.seq	473	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	0
3152.seq	339	gi 5922006	Homo sapiens fibulin 1 (FBLN1), splice variant B, mRNA	0.023
3153.seq	551	gi 4506678	Homo sapiens ribosomal protein S10 (RPS10) mRNA	0
3154.seq	503	gi 285940 dbj D14658 HUMORFO2	Human mRNA for KIAA0102 gene, complete cds	0
3155.seq	495	gi 190978 gb M18863 HUMREGA	Human islet of Langerhans regenerating protein (reg) mRNA, complete	0
3157.seq	371	gi 4502678	Homo sapiens CD63 antigen (melanoma 1 antigen) (CD63) mRNA	0
3158.seq	480	gi 4503480	Homo sapiens eukaryotic translation elongation factor 1 gamma	0
3159.seq	235	gi 2342696 gb AC002507 AC002507	Human Cosmid g5129z132 from 7q31.3, complete sequence [Homo sapiens]	3.8
3160.seq	507	gi 178854 gb J02908 HUMAPCJ	Human apolipoprotein J mRNA, complete cds.	0
3161.seq				
3162.seq	446	gi 4504900	Homo sapiens karyopherin alpha 4 (importin alpha 3) (KPNA4) mRNA	0
3163.seq	390	gi 291855 gb L09209 HUMAMYLOID	Homo sapiens amyloid protein homologue mRNA, complete cds	0
3165.seq	489	gi 3449301 dbj AB011538 AB011538	Homo sapiens mRNA for MEGF5, partial cds	2.00E-53
3166.seq	515	gi 4581270 emb AL035419.9 HS1100H13	Human DNA sequence from clone 1100H13 on chromosome 20q11.2 Contains	0.55
3167.seq	429	gi 537525 gb L29766 HUMEPHYDD	Homo sapiens epoxide hydrolase (EPHX) gene, complete cds	E-104

3168.seq	503	gij532596[gb]M12759[HUMIG.J02	Human Ig J chain gene, exons 3 and 4.	E-145
3169.seq	328	gij50683530[dbj]AP000615.1[AF000615	Oryza sativa genomic DNA, chromosome 3, clone:P0043E01	1.4
3170.seq	350	gij3947437[gb]AC005034[AC005034	Homo sapiens BAC clone NH0342K06 from 2, complete sequence [Homo sapie	3.00E-94
3173.seq	281	gij2854102[gb]AF045448[AF045448	Homo sapiens chromosome 21q22.3 PAC 191P10, complete sequence [Homo	4.6
3175.seq	357	gij3336977[emb]AL008626[HS1114G22	Human DNA sequence from clone 1114G22 on chromosome 1q24-25. Contains	0.38
3177.seq	293	gij4467309[emb]AL035423.4[HS2013	Human DNA sequence from clone 2013 on chromosome Xq25-26. Contains a	1.2
3178.seq	401	gij4504278	Homo sapiens H3 histone, family 3A (H3F3A) mRNA	E-104
3179.seq	393	gij4455339[emb]AL035522[ATT12J5	Arabidopsis thaliana DNA chromosome 4, BAC clone T12J5 (ESSAII	0.42
3180.seq	374	gij3080378[emb]AL022575[HS159A15	Homo sapiens DNA sequence from PAC 159A15 on chromosome	0.4
3182.seq	428	gij704346[gb]U03878[XL003878	Xenopus laevis thyroid hormone binding protein/pyruvate kinase type	0.007
3184.seq	403	gij1834699[emb]Z84988[HS284988	H.sapiens Ig lambda light chain variable region gene (18-26[IIA40)	E-170
3185.seq	405	gij3123551[emb]AL021408[HS523C21	Homo sapiens DNA sequence from PAC 523C21 on chromosome 6q23.1-23.3.	0.028
3186.seq	390			
3187.seq	406	gij5669089[gb]AF123094.1[AF123094	Homo sapiens API2-MLT fusion protein (API2-MLT) mRNA, complete cds	0
3188.seq	295	gij4972121[emb]AL049776.2[CNS00001	Human chromosome 14 DNA sequence *** IN PROGRESS *** BAC R-85K15	1.00E-74
3189.seq	411	gij181517[gb]L01131[HUMDCN07	Human decorin (DCN) gene, exon 8, complete cds.	0
3190.seq	291	gij4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	E-102
3191.seq	449	gij388034[gb]L05095[HUMRIBPROF	Homo sapiens ribosomal protein L30 mRNA, complete cds	0
3192.seq	373	gij28401[emb]X04350[HSADH3G1	Human mRNA for liver alcohol dehydrogenase (EC 1.1.1.1) gamma 1	0
3193.seq	427	gij4506700	Homo sapiens ribosomal protein S23 (RPS23) mRNA	0
3195.seq	385	gij339688[gb]M17733[HUMTHYB4	Human thymosin beta-4 mRNA, complete cds.	0
3196.seq	382	gij1773228[gb]U26312[HSU26312	Human heterochromatin protein HP1Hs-gamma mRNA, complete cds	E-163
3197.seq	391	gij4506624	Homo sapiens ribosomal protein L27a (RPL27A) mRNA	0
3198.seq	415	gij1947070[gb]U79415[HSU79415	Homo sapiens prepro dipeptidyl peptidase 1 (DPP-1) gene, complete cds	0
3199.seq	388	gij3954888[emb]AJ010444[HSAA010444	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	E-154
3200.seq	409	gij1806077[emb]Z84526[HS2428D7	H.sapiens flow-sorted chromosome 6 HindIII fragment, SC6pA28D7	1.7
3202.seq	401	gij4454514[gb]AC005536[AC005536	Homo sapiens BAC clone GS115E06 from 7p21-p15.3, complete sequence [Ho	1.7
3203.seq	313	gij5803047	Homo sapiens KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein	E-157
3204.seq	415	gij5454151	Homo sapiens ubiquinol-cytochrome c reductase binding protein	0
3207.seq	334	gij4502202	Homo sapiens ADP-ribosylation factor 3 (ARF3) mRNA	1.4
3208.seq	375			
3209.seq				
3210.seq	396	gij3417305[gb]AC004584[AC004584	Homo sapiens chromosome 17, clone hRPC1107_A_17, complete sequence	8.00E-12
3211.seq	377	gij37847[emb]X16478[HSVIMCD	Human mRNA 5'-fragment for vimentin N-terminal fragment	0
3212.seq	381	gij4699956[gb]AC005377.2[AC005377	Homo sapiens PAC clone DJ1136G02 from 7q32-q34, complete sequence	0.4
3213.seq	400	gij506340[dbj]J031888[HUMORFKG1Q	Human mRNA for KIAA0071 gene, partial cds.	0
3214.seq	416	gij3170740[gb]AF062139[AF062139	Homo sapiens clone 45u-12 immunoglobulin heavy chain variable	1.00E-96
3215.seq	404	gij4507668	Homo sapiens tumor protein, translationally-controlled 1 (TPT1)	0
3217.seq	330	gij3297863[emb]AL02239[HS3E5	Homo sapiens DNA sequence from PAC 3E5 on chromosome 20q12-13.12.	0.35
3218.seq	388	gij5931373[gb]AC008122.15[AC008122	Homo sapiens 2q31-33 BAC RPC111-1021H16 (Roswell Park Cancer Institute	8.00E-21
3219.seq	386	gij3098781[gb]AC003663[AC003663	Homo sapiens chromosome 17, clone HCIT87G17, complete sequence [Homo	E-158
3220.seq	416	gij4504170	Homo sapiens glutathione S-transferase A2 (GSTA2) mRNA	0
3221.seq	356	gij3845348[gb]AC005933[AC005933	Homo sapiens chromosome 19, cosmid F15472, complete sequence [Homo	0.38
3222.seq	408	gij4454683[gb]AF070654[AF070654	Homo sapiens comichon protein mRNA, complete cds	0
3223.seq	405	gij6180044[gb]AF192529.1[AF192529	Homo sapiens RPA-binding trans-activator (RBT1) mRNA, complete cds	0
3224.seq	379	gij5919146[gb]AF182289.1[AF182289	Homo sapiens U6 snRNA-associated 5m-like protein L3m3 mRNA,	0
3225.seq	435	gij32265[emb]X00452[HSILDC2	Human mRNA for DC classII histocompatibility antigen alpha-chain	0
3226.seq	408	gij4502472	Homo sapiens B-cell translocation protein 1 (BTG1) mRNA	0
3228.seq	422	gij2370083[emb]Z95703[HS433M19	Human DNA sequence from PAC 433M19 on chromosome Xq26.3-Xq27.1.	0.11
3229.seq	387	gij5679536[emb]AL096827.3[HSUJ596C	Human DNA sequence from clone 596C3 on chromosome Xq13.2-21.2. Contain	0.41
3231.seq	430	gij4929630[gb]AF151839.1[AF151839	Homo sapiens CGI-81 protein mRNA, complete cds	0
3232.seq	386	gij5306225[gb]AC005902.7[AC005902	Homo sapiens 3p21.3-4 PAC RPC4-544D10 (Roswell Park Cancer Institute	0.41
3233.seq	379	gij532596[gb]M12759[HUMIG.J02	Human Ig J chain gene, exons 3 and 4.	E-168
3234.seq	368	gij3355875[emb]AL023284[HS406A7	Human DNA sequence from clone 406A7 on chromosome 6q23-24. Contains	E-157
3235.seq	395	gij4505812	Homo sapiens dynein, cytoplasmic, light polypeptide (PIN) mRNA	0
3236.seq	416	gij4490865[emb]Z98882.4[HS356B8	Human DNA sequence from clone 356B8 on chromosome 16, complete sequen	0.44
3237.seq	373	gij5032212	Homo sapiens UDP-galactose transporter related 1 (UGTREL1) mRNA	0
3239.seq	405	gij6015478[dbj]AP000561.1[AP000561	Homo sapiens genomic DNA, chromosome 21q21.2, clone:T48, LL56-APP	1.7
3240.seq	372	gij3860076[gb]AF097021[AF097021	Homo sapiens GW112 protein (GW112) mRNA, complete cds	0
3241.seq	384	gij4503938	Homo sapiens guanylate binding protein 1, interferon-inducible, 67kD	2.00E-49
3242.seq	406	gij3046270[gb]AC004074[AC004074	Homo Sapiens Chromosome X clone BWXD759, complete sequence [Homo	0.43
3243.seq	409	gij4504900	Homo sapiens karyopherin alpha 4 (Importin alpha 3) (KPNA4) mRNA	0
3244.seq	411	gij4507668	Homo sapiens tumor protein, translationally-controlled 1 (TPT1)	0
3246.seq	388	gij38422[emb]X69150.1[HSRPS18	Homo sapiens mRNA for ribosomal protein S18	0
3247.seq	118	gij940043[gb]U28250[HSU28250	Human novel transcript from adenocarcinoma cell line, clone	5.00E-47
3248.seq	371	gij531170[gb]U12404[HSU12404	Human Csa-19 mRNA, complete cds.	0
3249.seq	380	gij5138905[gb]AF092128.1[AF092128	Homo sapiens putative transmembrane protein E3-16 mRNA, complete cds	0
3250.seq	422	gij33731[emb]X57816[HSIGVL026	Human rearranged immunoglobulin lambda light chain mRNA	0
3251.seq	394	gij3258613[gb]AC004687[AC004687	Homo sapiens chromosome 17, clone hRPC.111.1.10, complete sequence	E-140
3253.seq	398	gij4689145[gb]AF077049.1[AF077049	Homo sapiens lambda-crystallin mRNA, complete cds	0
3254.seq	377	gij4581462[emb]Y09703.1[HSMEMA	H.sapiens mRNA for MEMA protein	0
3255.seq	397	gij3598729[gb]AC002483[AC002483	Homo sapiens PAC clone 248015 from 13q12-q13, complete sequence [Homo	1.7
3256.seq	385	gij5678818[gb]AF146191.1[AF146191	Homo sapiens FRG1 (FRG1) gene, complete cds: 5S ribosomal RNA gene,	2.00E-06
3257.seq	351	gij4220441[gb]AC006505[AC006505	Homo sapiens chromosome 19, fosmid 39524, complete sequence [Homo	1.5
3258.seq	311	gij2276311[emb]Z85996.1[HS431A14	Human DNA sequence from PAC 431A14 on chromosome 6p21. Contains	E-170
3259.seq	420	gij5002511[emb]Z98266.1[HSZ98266	Homo sapiens gene encoding plakophilin (exons 1-13)	1.8
3260.seq	421	gij34469[emb]X66867[HSMAXG	H.sapiens max gene	0
3261.seq	328	gij4508118[gb]AC005041.2[AC005041	Homo sapiens clone NH0523H20, complete sequence	0.35
3262.seq	383	gij5917928[gb]AC008055.6[AC008055	Homo sapiens 12q22-103.4-106.5 BAC RPC111-718L23 (Roswell Park Cancer	0.026
3263.seq	384	gij4885066	Homo sapiens ras homolog gene family, member C (ARHC) mRNA	E-173
3264.seq	418	gij4503658	Homo sapiens Finkel-Biskis-Reilly murine sarcoma virus (FBR-MuSV)	E-172
3265.seq	530	gij541612[gb]U14650[HSU14650	Human platelet-endothelial tetraspan antigen 3 mRNA, complete cds.	0
3266.seq	519	gij5708477[gb]AC007463.3[AC007463	Homo sapiens BAC clone NH0244L12 from 2, complete sequence	0.14
3268.seq	476	gij4558768[gb]AC007199.1[AC007199	Homo sapiens chromosome 5 BAC clone 111n13, complete sequence	4.00E-91
3270.seq	512	gij180242[gb]M80651[HUMCE16A	Hum ORF (CE15) mRNA, 3' flank.	E-138
3271.seq	418	gij4503486	Homo sapiens ephrin-A5 (EFNA5) mRNA >gij1019430[gb]U26403[HSU26403	0.45
3273.seq	499	gij4507356	Homo sapiens transgelin 2 (TAGLN2) mRNA	0
3274.seq	526	gij33689[emb]X12683[HSIGVKA10	Human germ-line immunoglobulin kappa light chain variable region	E-127
3275.seq	489	gij531170[gb]U12404[HSU12404	Human Csa-19 mRNA, complete cds.	0
3276.seq	474	gij4502922	Homo sapiens acidic calponin 3 (CNN3) mRNA	5.00E-29
3277.seq	497	gij4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
3278.seq	490	gij5031748	Homo sapiens high-mobility group (nonhistone chromosomal) protein	0
3280.seq	485	gij5757526[gb]AC006371.2[AC006371	Homo sapiens BAC clone NH0304C24 from Y, complete sequence	0.52
3281.seq	489	gij4506660	Homo sapiens ribosomal protein L7a (RPL7A) mRNA	0
3282.seq	499	gij1944528[gb]J01415.1[HUMMTCG	Human mitochondrion, complete genome	0
3283.seq	495	gij4506616	Homo sapiens ribosomal protein L17 (RPL17) mRNA	0
3284.seq	482			
3287.seq	498	gij3861482[gb]AF042089[AF042089	Homo sapiens chromosome 3, olfactory receptor pseudogene cluster 1.	0.14
3289.seq	201	gij5803178	Homo sapiens sperm specific antigen 2 (SSFA2), mRNA	6.00E-11
3290.seq	306	gij2992477[gb]AC004478[AC004478	Homo Sapiens Chromosome X clone BWXD312, complete sequence [Homo	0.081
3291.seq	497			
3292.seq	473			
3293.seq	104	gij5830428[emb]AL110502.1[HSJDJ635E	Human DNA sequence from clone 635E8 on chromosome 1p33-34.3, complete	1.5
3294.seq	354	gij3413268[emb]Z93244[HS116F5	Human DNA sequence from clone 116F5 on chromosome 22q13. Contains part	0.37
3295.seq	511	gij1399085[gb]U43701[HSU43701	Human ribosomal protein L23a mRNA, complete cds.	0
3296.seq	438	gij249370[gb]S35960[S35960	laminin receptor homolog [3' region] (human, mRNA Partial, 739 nt)	0
3297.seq	143	gij3292944[emb]AL022317[HS140L1	Human DNA sequence from clone 140L1 on chromosome 20q13.1-13.31.	0.55
3299.seq	508	gij333211[gb]M73236[PPHPAPV42A	Human papillomavirus ORF E6, ORF E7, ORF E1, ORF E2, ORF E4, ORF E5,	0.55

3300.seq	500	gi 4505864	Homo sapiens plasminogen activator, urokinase receptor (PLAUR) mRNA	0
3302.seq	386	gi 2343108 gb AF004230 AF004230	Homo sapiens monocyte/macrophage Ig-related receptor MIR-7 (MIR cl-7)	0
3304.seq	320	gi 4506644	Homo sapiens ribosomal protein L38 (RPL38) mRNA	E-172
3305.seq	516	gi 2197000 gb U95367 HSU95367	Human GABA-A receptor α 1 subunit mRNA, complete cds	0
3306.seq	508	gi 4503538	Homo sapiens eukaryotic translation initiation factor 4 gamma, 2	0
3307.seq	522	gi 4885590	Homo sapiens selenoprotein P, plasma, 1 (SEPP1) mRNA	E-165
3308.seq	481			
3310.seq	436	gi 5102616 emb AL034417.14 HS215D11	Human DNA sequence from clone 215D11 on chromosome 1p36.12-36.33	0.12
3311.seq	503	gi 4505896	Homo sapiens plastin 1 (I isoform) (PLS1) mRNA	0
3312.seq	508	gi 4507126	Homo sapiens small nuclear ribonucleoprotein polypeptide C (SNRPC)	0
3313.seq	186			
3314.seq	526	gi 3954884 emb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	0
3315.seq	502	gi 219427 dbj D00137 HUMADH21C	Homo sapiens mRNA for alcohol dehydrogenase beta 1, complete cds	0
3316.seq	335	gi 190978 gb M18963 HUMREGA	Human islet of Langerhans regenerating protein (reg) mRNA, complete	0
3317.seq	495	gi 5803049	Homo sapiens KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein	0
3318.seq	174			
3319.seq	481	gi 5912095 emb AL117557.1 HSM801085	Homo sapiens mRNA: cDNA DKFZp564D177 (from clone DKFZp564D177)	0
3321.seq	502	gi 5080755 emb AC007842.1 AC007842	Homo sapiens chromosome 19, BAC 331191 (CIT-B-47113), complete sequence	0
3322.seq	501	gi 5791509 emb AL050341.18 HSDJ39G2	Human DNA sequence from clone 39G22 on chromosome 1p32.1-34.3, complete	E-150
3323.seq	485	gi 4505396	Homo sapiens serine/threonine protein-kinase (NIK) mRNA	0
3324.seq	507	gi 4503530	Homo sapiens eukaryotic translation initiation factor 4A, isoform 2	1.00E-91
3325.seq	496	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
3326.seq	520	gi 6137796	Human mitochondrion, complete genome	0
3327.seq	490	gi 4501888	Homo sapiens actin, gamma 2, smooth muscle, enteric (ACTG2) mRNA	0
3329.seq	451	gi 4501888	Homo sapiens actin, gamma 2, smooth muscle, enteric (ACTG2) mRNA	0
3330.seq	428	gi 339503 gb M16553 HUMTFPA	Human tissue factor mRNA, complete cds, with an Alu repeat in the 3'	5.00E-50
3331.seq	434	gi 339688 gb M17733 HUMTHYB4	Human thymosin beta-4 mRNA, complete cds.	0
3332.seq	215			
3333.seq	443	gi 4914518 emb Z83840.7 HS216E10	Human DNA sequence from clone 216E10 on chromosome 22 Contains genes	0.47
3335.seq	409	gi 5326822 gb AF050641.1 AF050641	Homo sapiens NADH-ubiquinone oxidoreductase 39kDa subunit mRNA,	0
3336.seq	487	gi 4049267 gb U17714 HSU17714	Homo sapiens putative tumor suppressor ST13 (ST13) mRNA, complete cds	0
3337.seq	482	gi 4507960	Homo sapiens zinc finger protein homologous to Zfp-36 in mouse	0
3338.seq	401	gi 6137796	Human mitochondrion, complete genome	E-170
3339.seq	501	gi 1164908 emb Z68868 HSJ3E10	Human DNA sequence from PAC 3E10, between markers DXS366 and DXS87	0.14
3341.seq	482	gi 4885094	Homo sapiens murine leukemia viral (bmi-1) oncogene homolog (BMI1)	0
3342.seq	288	gi 4503724	Homo sapiens FK506-binding protein 1A (12kD) (FKBP1A) mRNA	E-129
3343.seq	460	gi 388034 gb L05095 HUMRIBPROF	Homo sapiens ribosomal protein L30 mRNA, complete cds	0
3344.seq	486	gi 339688 gb M17733 HUMTHYB4	Human thymosin beta-4 mRNA, complete cds.	0
3345.seq	486	gi 550014 gb U14967 HSU14967	Human ribosomal protein L21 mRNA, complete cds.	0
3346.seq	402	gi 4505288	Homo sapiens mevalonate (diphospho) decarboxylase (MVD), mRNA	0
3347.seq	491	gi 3954888 emb AJ010444 HSA010444	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	0
3350.seq	518	gi 6005987	Homo sapiens acid phosphatase 1, soluble (ACP1), transcript variant	0
3351.seq	365	gi 4775614 emb AL035079.14 HS53C18	Human DNA sequence from clone 53C18 on chromosome 11q12-13. Contains	0.098
3352.seq	487	gi 1353817 gb U43766 HSU43766	Human immunoglobulin light chain variable region mRNA, cell line	0
3354.seq	295			
3355.seq	467	gi 4506652	Homo sapiens ribosomal protein L4 (RPL4) mRNA	0
3356.seq	424	gi 31091 emb X16869 HSEF1AC	Human mRNA for elongation factor 1-alpha (clone CEF4)	0
3357.seq	472	gi 1752745 dbj D86566 DB6566	Human DNA for NOTCH4, partial cds	0.13
3358.seq	507	gi 872118 emb Z49112 HSHLXACW7	H.sapiens gene for leukocyte antigen-Cw7 variant (HLA-Cw*0702)	0
3360.seq	502	gi 2155224 gb AC002094 AC002094	Genomic sequence from Human 17, complete sequence [Homo sapiens]	2.00E-46
3362.seq	460	gi 4506116	Homo sapiens protein S (alpha) (PROS1) mRNA	0.49
3363.seq	486	gi 4504446	Homo sapiens heterogeneous nuclear ribonucleoprotein A2/B1 (HNRPA2B1)	0
3364.seq	398	gi 5114052 gb AF085359.1 AF085359	Homo sapiens HSPC030 mRNA, complete cds	0
3366.seq	494	gi 3688349 emb AL030996.1 HS1189B24	Human DNA sequence from clone 1189B24 on chromosome Xq25-26.3. Conta	0.53
3368.seq	299	gi 292446 gb M77234 HUMRPSA3A	Human ribosomal protein S3a mRNA, complete cds.	E-154
3369.seq	513	gi 5174538	Homo sapiens malate dehydrogenase 1, NAD (soluble) (MDH1) mRNA	0
3370.seq	477	gi 1944628 gb J01415.1 HUMMTCG	Human mitochondrion, complete genome	0
3371.seq	506	gi 4894373 gb AF067170.1 AF067170	Homo sapiens alpha endosulfine mRNA, complete cds	0
3372.seq	205	gi 4885128	Homo sapiens CCAAT/enhancer binding protein (C/EBP), beta (CEBPB)	3.00E-34
3373.seq	349	gi 4506890	Homo sapiens SET translocation (myeloid leukemia-associated) (SET)	3.00E-63
3375.seq	470	gi 4757993	Homo sapiens clathrin-associated/assembly/adaptor protein, medium 1	0
3378.seq	446	gi 5689486 dbj AB028998.1 AB028998	Homo sapiens mRNA for KIAA1075 protein, partial cds	0
3379.seq	399	gi 4176360 gb AC06406 AC06406	Homo sapiens, clone hRPC.25_A_1, complete sequence [Homo sapiens]	0.42
3380.seq	373	gi 3097841 gb AC004643 AC004643	Homo sapiens chromosome 16, cosmid clone 363E3 (LANL), complete	4.00E-47
3381.seq	480	gi 4454709 gb AF070667 AF070667	Homo sapiens NG,NG-dimethylarginine dimethylaminohydrolase homolog	0
3382.seq	461	gi 2343108 gb AF004230 AF004230	Homo sapiens monocyte/macrophage Ig-related receptor MIR-7 (MIR cl-7)	E-158
3383.seq	511	gi 180990 gb M81478 HUMCPTP	Human cytoplasmic phosphotyrosine phosphatase mRNA.	E-158
3384.seq	208	gi 5738137 gb AF165124.1 AF165124	Homo sapiens chromosome 5q31.1-q33.1 clone BAC dir082c10 containing	0.21
3385.seq	227	gi 4378311 gb AF103491 AF103491	Homo sapiens isolate donor N clone N229K immunoglobulin kappa light	2.00E-63
3386.seq	440	gi 6137796	Human mitochondrion, complete genome	0
3387.seq	409	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	2.00E-92
3388.seq	412	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	E-114
3389.seq	310	gi 2772533 gb AC003992 AC003992	Human BAC clone RG250D13 from 7q31, complete sequence [Homo sapiens]	0.33
3391.seq	426	gi 4506664	Homo sapiens ribosomal protein L9 (RPL9) mRNA	0
3394.seq	422	gi 3282154 gb AC004241 AC004241	Homo sapiens 12q13.1 PAC RPC13-197B17 (Roswell Park Cancer Institute	0.45
3395.seq	495	gi 2804346 gb AC004029 AC004029	Human BAC clone RG035E18 from 7q31, complete sequence [Homo sapiens]	0.034
3396.seq	503	gi 5106997 gb AF125101.1 AF125101	Homo sapiens HSPC040 protein mRNA, complete cds	0
3397.seq	503	gi 4506602	Homo sapiens ribosomal protein L15 (RPL15) mRNA	0
3398.seq	346	gi 29548 emb Y00706 HSC4A1	Human mRNA for alpha 1 chain of basement membrane (type IV) collagen	0
3399.seq	434	gi 1514560 emb Z69780 HSHAL1X37	H.sapiens ALL-1 gene exon 37	8.00E-86
3400.seq	469	gi 3950886 emb X74070 HSBTF3	H.sapiens mRNA for transcription factor BTF 3	0
3401.seq	311	gi 3327205 dbj AB014596 AB014596	Homo sapiens mRNA for KIAA0696 protein, partial cds	3.00E-94
3402.seq	294	gi 5174428	Homo sapiens acetyl-Coenzyme A acyltransferase 2 (mitochondrial	E-159
3403.seq	189	gi 4200326 emb AL022313 HS1119A7	Human DNA sequence from clone 1119A7 on chromosome 22q12.2-12.3	2.00E-04
3404.seq	186			
3405.seq	207	gi 181572 gb M34482 HUMDKERB	Human cyokeratin 8 (CK8) gene, complete cds.	E-112
3406.seq	374	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	7.00E-80
3407.seq	252	gi 190161 gb M73547 HUMPOLLA	Human polyposis locus (DP1 gene) mRNA, complete cds.	E-113
3408.seq	495	gi 4007385 gb AF036715 AF036715	Homo sapiens syntaxin 8 mRNA, complete cds.	0
3409.seq	315	gi 5689442 dbj AB028976.1 AB028976	Homo sapiens mRNA for KIAA1053 protein, partial cds	E-146
3410.seq	474	gi 13003 emb V00662 MIH5XX	H.sapiens mitochondrial genome	0
3411.seq	499			
3412.seq	445	gi 5817150 emb AL110218.1 HSM80087	Homo sapiens mRNA: cDNA DKFZp434A163 (from clone DKFZp434A163)	E-122
3413.seq	409	gi 4003387 gb AB020867.1 AB020867	Homo sapiens genomic DNA of Bp21.3-p22 anti-oncogene of hepatocellular	0.44
3414.seq	482	gi 2769692 gb AC003985 AC003985	Human PAC clone DJ097B12 from 7p15-p21, complete sequence [Homo	0.033
3415.seq	520	gi 3717976 emb AL031466 HS653H13	Human DNA sequence from clone 653H13 on chromosome Xp11.4-21.3 Conta	0.56
3416.seq	462	gi 3063487 gb AC004103 AC004103	Homo sapiens Xp22 BAC GS-619J3 (Genome Systems Human BAC library)	0.13
3417.seq	372	gi 4557836	Homo sapiens procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine	0
3420.seq	491	gi 13000 emb X62996 MIHSGENOM	H.sapiens mitochondrial genome (consensus sequence)	0
3421.seq	501	gi 5902059	Homo sapiens retinitis pigmentosa 2 (X-linked recessive) (RP2), mRNA	6.00E-84
3423.seq	494	gi 5360678 gb AB022656.1 AB022656	Homo sapiens mRNA for anti-Entamoeba histolytica immunoglobulin	E-174
3424.seq	496	gi 4726099 gb AC006254.10 AC006254	Homo sapiens 3p21.1-9 PAC RPC15-966M1 (Roswell Park Cancer Institute	3.00E-67
3425.seq	479	gi 5803101	Homo sapiens MORF-related gene 15 (MRG15), mRNA	0
3426.seq	310	gi 4502248	Homo sapiens ADP-ribosylation factor[arf]-directed GTPase activating	E-160
3427.seq	225	gi 4507668	Homo sapiens tumor protein, translationally-controlled 1 (TPT1)	E-122
3428.seq	459	gi 1944628 gb J01415.1 HUMMTCG	Human mitochondrion, complete genome	0
3429.seq	495	gi 4455411 emb AL035450.1 HS1057D18	Human DNA sequence from clone 1057D18 on chromosome 22q13.31-13.33	0.002
3430.seq	497	gi 4506702	Homo sapiens ribosomal protein S24 (RPS24a) mRNA	0

3431.seq	471	gi 441312 emb X72422 HSIGKLV01	H.sapiens mRNA for rearranged Ig kappa light chain variable region	0
3432.seq	506	gi 3288456 emb AJ00008 HSC2P13K1	Homo sapiens mRNA for C2 domain containing P13-kinase	0.14
3433.seq	445	gi 23221 emb X56468 HS1433	Human mRNA for 14.3.3 protein, a protein kinase regulator	0
3434.seq	494	gi 4512311 dbj AB019441.1 AB019441	Homo sapiens DNA for immunoglobulin heavy-chain variable region.	0
3435.seq	194	gi 34717 emb X64899 HSM21HOM	H.sapiens mRNA homologous to mouse P21 mRNA	E-105
3436.seq	466	gi 3170672 gb AF062105 AF062105	Homo sapiens clone 21u-19 immunoglobulin heavy chain variable	E-131
3437.seq	325	gi 1945756 emb Z82899 HS181N1	Human DNA sequence from PAC 181N1 on chromosome X contains ESTs, STS	0.34
3438.seq	504	gi 3882200 dbj AB018283.1 AB018283	Homo sapiens mRNA for KIAA0740 protein, complete cds	0.54
3439.seq	238	gi 1710247 gb U9278 HSU97278	Human protein disulfide isomerase-related protein P5 mRNA, partial	3.00E-99
3440.seq	343	gi 190980 gb J05412 HUMREGB	Human regenerating protein (reg) gene, complete cds.	0
3441.seq	510	gi 2957144 gb U90878 HSU90878	Homo sapiens carboxyl terminal LIM domain protein (CLIM1) mRNA,	0
3442.seq	418	gi 13683 emb V00110 MIT1HS	Human mitochondrial genes for several tRNAs (Phe, Val, Leu) and 12S	E-177
3443.seq	491	gi 1483496 emb Z77998 HSPA6H11	H.sapiens flow-sorted chromosome 6 HindIII fragment, SC6pA6H11	0.13
3444.seq	508	gi 2967847 gb AF052578.1 AF052578	Homo sapiens androgen receptor associated protein 24 (ARA24) mRNA,	0
3445.seq	500	gi 5281313 gb AF133123.1 AF133123	Homo sapiens transcription factor IIIC102 mRNA, complete cds	0
3446.seq	382	gi 4689133 gb AF077043.1 AF077043	Homo sapiens 60S ribosomal protein L36 mRNA, complete cds	0
3448.seq	505	gi 185401 gb L03166 HUMIGHHE	Human Ig rearranged gamma-chain mRNA V-region, partial cds.	E-127
3450.seq	499	gi 189019 gb M22918 HUMMYLCB	Human non-muscle myosin alkali light chain mRNA, 3' end.	0
3452.seq	501	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
3453.seq	330	gi 1679919 gb S40022 S40022	Pcp-2=Putkinje cell protein 2 [mice, BALB C, liver, Genomic, 3935 nt]	7.00E-15
3454.seq	480	gi 1870243 gb U46689 HSU46689	Human microsomal aldehyde dehydrogenase (ALD10) mRNA, complete cds	0
3457.seq	729	gi 190978 gb M18963 HUMREGA	Human islet of Langerhans regenerating protein (reg) mRNA, complete	0
3458.seq	697	gi 2623584 gb AF027158 AF027158	Homo sapiens immunoglobulin kappa light chain (T6J/k) mRNA, partial	0
3459.seq	470	gi 3882264 gb AB018315.1 AB018315	Homo sapiens mRNA for KIAA0772 protein, complete cds	0.13
3460.seq	657	gi 3947836 emb AL024509 HS522P13	Human DNA sequence from clone S22P13 on chromosome 6p21.31-22.3.	0.18
3461.seq	176	gi 2343108 gb AF004230 AF004230	Homo sapiens monocyte/macrophage Ig-related receptor MIR-7 (MIR cl-7)	1.00E-39
3462.seq	618	gi 4768676 gb AF113887.1 AF113887	Homo sapiens kappa 1 immunoglobulin light chain mRNA, complete cds	E-143
3463.seq	461	gi 4506692	Homo sapiens ribosomal protein S17 (RPS17) mRNA	0
3464.seq	670	gi 4503056	Homo sapiens crystallin, alpha B (CRYAB) mRNA	0
3465.seq	689	gi 4826558	Homo sapiens capping protein (actin filament) muscle Z-line, beta	0
3466.seq	599	gi 5002378 gb AF153821.1 AF153821	Homo sapiens alcohol dehydrogenase beta2 subunit mRNA, complete cds	E-167
3467.seq	648	gi 3387966 gb AF070591 AF070591	Homo sapiens clone 24649 mRNA sequence	0
3468.seq	654	gi 4506662	Homo sapiens ribosomal protein LB (RPLB) mRNA	0
3469.seq	525	gi 1460066 emb Z77249 HS358H7	Human DNA sequence from PAC 358H7 on chromosome X	0.14
3470.seq	670	gi 5931507 dbj AP000529.1 AP000529	Homo sapiens genomic DNA, chromosome 22q11.2, Cat Eye Syndrome region,	0.18
3472.seq	579	gi 5708468 gb AC006373.3 AC006373	Homo sapiens clone NH0352E12, complete sequence	0
3473.seq	538	gi 5731112 gb AF156965.1 AF156965	Homo sapiens translocan-associated protein alpha subunit mRNA,	0
3474.seq	700	gi 5836158 gb AC006466.3 AC006466	Homo sapiens clone RG471E18, complete sequence	E-107
3475.seq	652	gi 3638954 gb AC004877 AC004877	Homo sapiens PAC clone DJ0751H13 from 7q35-qrter, complete sequence [Ho	0.18
3476.seq	713	gi 3861482 gb AF042089 AF042089	Homo sapiens chromosome 3, olfactory receptor pseudogene cluster 1,	E-162
3477.seq	368	gi 4502572	Homo sapiens caspase 1, apoptosis-related cysteine protease	E-143
3478.seq	581	gi 3790137 emb AL022151 HS199L16	Human DNA sequence from clone 199L16 on chromosome Xq22.1-22.3,	0.04
3480.seq	583	gi 4503654	Homo sapiens Fanconi anemia, complementation group A (FANCA) mRNA	0
3481.seq	353	gi 5002609 emb AL031390.4 HS197L1	Human DNA sequence from clone 197L1 on chromosome 6q24.1-25.2. Contain	E-163
3483.seq	873	gi 3201899 gb AF067420 AF067420	Homo sapiens SINC73 protein (SINC73) mRNA, complete cds	E-168
3484.seq	650	gi 102632 gb AC011493.2 AC011493	Homo sapiens chromosome 19 clone CIT978SKB_2819, complete sequence	1.00E-70
3485.seq	662	gi 33737 emb X57819 HSIGVL029	Human rearranged immunoglobulin lambda light chain mRNA	0
3486.seq	656			
3487.seq	183	gi 644480 dbj D38112.1 HUMMTA	Homo sapiens mitochondrial DNA, complete sequence	4.00E-30
3488.seq	626	gi 2935593 gb AC003657 AC003657	Homo sapiens Xp22 BAC GS-602M16 (Genome Systems Human BAC library)	E-177
3490.seq	791	gi 339692 gb M14483 HUMTHYMAA	Human prothymosin alpha mRNA, complete cds.	E-135
3491.seq	542	gi 435637 dbj D16111 HUMHRPBP	Human mRNA for human homologue of rat phosphatidylethanolamine	0
3492.seq	878	gi 4502204	Homo sapiens ADP-ribosylation factor 4 (ARF4) mRNA	0
3493.seq	614	gi 4507206	Homo sapiens sorcin (SRI) mRNA >gi 338481 gb M32886 HUMSRICPA Human	0
3494.seq	674	gi 4758515	Homo sapiens hepatoma-derived growth factor (high-mobility group)	0
3495.seq	543	gi 4589587 dbj AB023189.1 AB023189	Homo sapiens mRNA for KIAA0972 protein, complete cds	4.00E-11
3496.seq	643	gi 496901 emb X7538 HST1F	H.sapiens nuk_34 mRNA for translation initiation factor	0
3497.seq	628	gi 2828774 gb AC003664 AC003664	Homo sapiens chromosome 17, clone HCT54K19, complete sequence [Homo	0.17
3498.seq	3			
3499.seq	848	gi 1457953 gb U57091 HSU57091	Human small GTP-binding protein rab22b mRNA, complete cds	0
3500.seq	441	gi 2347079 gb AC002044 HUAC002044	Human Chromosome 16 BAC clone CIT987SK-A-418G10, complete sequence	0.12
3502.seq	661	gi 4503066	Homo sapiens crystallin, zeta (quinone reductase) (CRYZ) mRNA	0
3503.seq	545	gi 5542034 gb AC006536.2 AC006536	Homo sapiens chromosome 14 clone BAC257P13 map 14q31, complete sequer	0.15
3505.seq	470	gi 1786217 gb AE000114 AE000114	Escherichia coli K-12 MG1655 section 4 of 400 of the complete genome	0
3506.seq	57	gi 4503720	Homo sapiens four and a half LIM domains 1 (FHLL1) mRNA	2.00E-16
3507.seq	568	gi 13003 emb V00692 MIHSXX	H.sapiens mitochondrial genome	0
3508.seq	691	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-174
3509.seq	613	gi 476867 gb AF114264.1 AF114264	Homo sapiens clone HH409 unknown mRNA	0
3510.seq	362	gi 4758039	Homo sapiens cytochrome c oxidase subunit Vlc (COX6C), nuclear gene	0
3511.seq	602	gi 2815069 emb AL021451 HS188H10	Human DNA sequence from PAC 188H10 on chromosome 6q22. Contains STS	0
3512.seq	529	gi 6005738	Homo sapiens NRAS-related gene (D1S155E), mRNA	0
3513.seq	612	gi 36139 emb X52967 HSRPL7	Human mRNA for ribosomal protein L7	0
3515.seq	548	gi 182786 gb M80815 HUMFUCAS07	H.sapiens a-L-fucosidase gene, exon 7 and 8, and complete cds.	0
3517.seq	73			
3518.seq	388	gi 3482999 emb Z95889 HS211A9	Human DNA sequence from clone 211A9 on chromosome 22q12.1. Contains ES	0.1
3519.seq	818	gi 4156190 gb AC004880 AC004880	Homo sapiens PAC clone DJ0759N11 from 7q31, complete sequence [Homo	0
3520.seq	633			
3522.seq	612	gi 4926908 gb AC004895.2 AC004895	Homo sapiens clone DJ081DE06, complete sequence	0.66
3523.seq	533	gi 5578939 emb AL035417.15 HS891H21	Human DNA sequence from clone 891H21 on chromosome 1p32.2-34.2 Contain	0
3524.seq	532	gi 4156200 gb AC004860 AC004860	Homo sapiens clone DJ0691L07, complete sequence [Homo sapiens]	0.037
3525.seq	653	gi 4582483 gb AC006312.8 AC006312	Homo sapiens chromosome 9, clone hRPK.401_G_18, complete sequence	2.00E-13
3526.seq	591	gi 3005722 gb AF055003 AF055003	Homo sapiens clone 24414 unknown mRNA, partial cds	0
3528.seq	730	gi 2580673 gb AF009944 HSAF009944	Homo sapiens ubiquitous TPR motif, Y isoform (UTY) mRNA, alternative	0
3529.seq	46			
3530.seq	36	gi 183181 gb J03004 HUMGIAA	Human guanine nucleotide-binding regulatory protein (G)	3.00E-04
3532.seq	586	gi 3005926 emb AJ015611 HSAJ15611	Homo sapiens repetitive sequences, alphoid DNA, 1427bp	2.5
3533.seq	466	gi 2593559 gb AF015812 AF015812	Homo sapiens RNA helicase p68 (HUMP68) gene, complete cds	0
3534.seq	621	gi 1665778 dbj D87445 D87445	Human mRNA for KIAA0256 gene, complete cds	3.00E-95
3535.seq	339	gi 184918 gb M60192 HUMIGHA2A	Human germline IgA2 chain C region.	E-114
3536.seq	274	gi 505101 dbj D31887 HUMORFKG1P	Human mRNA for KIAA0062 gene, partial cds	E-122
3538.seq	79			
3541.seq	563	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	0
3542.seq	401	gi 5052312 gb AF116272.1 AF116272	Homo sapiens T-cell activation protein (PGR1) gene, complete cds	0
3543.seq	540	gi 4506618	Homo sapiens ribosomal protein L24 (RPL24) mRNA	0
3546.seq	851	gi 4503348	Homo sapiens deoxyribonuclease II, lysosomal (DNASE2) mRNA, and	0
3547.seq	624	gi 4079627 gb AC006050 AC006050	Homo sapiens chromosome 17, clone hRPK.268_F_2, complete sequence [Hori	0
3548.seq	633	gi 4503574	Homo sapiens glutaryl aminopeptidase (aminopeptidase A) (ENPEP) mRNA	0
3551.seq	345	gi 292440 gb L11567 HUMRPL37Z	Homo sapiens ribosomal protein L37 mRNA, complete cds.	0
3554.seq	449	gi 5031930	Homo sapiens nascent-polyptide-associated complex alpha	0
3556.seq	504	gi 2961198 gb AF051160 AF051160	Homo sapiens tyrosine phosphatase (PRL-1) gene, complete cds	0
3557.seq	186	gi 185977 gb M23091 HUMIGKVAD	Human germline pseudo-IgK chain gene, V3-region, clone Humkv329.	3.00E-49
3558.seq	471	gi 4039013 gb AF037338 AF037338	Homo sapiens cleft lip and palate transmembrane protein 1 (CLPTM1)	0.13
3559.seq	527	gi 185538 gb M18512 HUMIGHVAJ	Human (fetal) rearranged IgM chain VDJ-region mRNA, clone 20P3.	E-144
3560.seq	506	gi 1805302 gb HU04143 HU04143	Human D-dopachrome tautomerase mRNA, complete cds	0
3561.seq	443	gi 184743 gb J00220 HUMGCC8	Homo sapiens immunoglobulin alpha-1 heavy chain constant region	0
3562.seq	495	gi 190978 gb M18963 HUMREGA	Human islet of Langerhans regenerating protein (reg) mRNA, complete	0
3563.seq	213	gi 5031956	Homo sapiens nuclear protein containing WW domain 38 kD (NPW38) mRNA	E-115
3564.seq	492	gi 4758543	Homo sapiens heterogeneous nuclear ribonucleoprotein C (C1/C2)	0

3565.seq	493	gi 4501888	Homo sapiens actin, gamma 2, smooth muscle, enteric (ACTG2) mRNA	0
3566.seq	487	gi 3721650 dbj AB012910 AB012910	Homo sapiens mRNA for anti-HBsAg immunoglobulin Fab kappa chain.	0
3567.seq	484	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
3568.seq	382	gi 4071052 emb AL032822.1 HS59B16	Human DNA sequence from clone 59B16 on chromosome 6p22.1-22.3. Contains	0.41
3569.seq	396	gi 5565976 gb AF152462.1 AF152462	Homo sapiens transmembrane protein BR1 (BR1) mRNA, complete cds	0
3570.seq	281			
3572.seq	488	gi 664880 gb S73804 S73804	ND2=NADH dehydrogenase subunit I (heteroplasmic G->A transition in	6.00E-13
3573.seq	497			
3574.seq	237	gi 5932521 gb AC007061.3 AC007061	Leishmania major chromosome 3 clone L1559 strain Friedlin, complete	0.016
3575.seq	465	gi 38405 emb X67292 HSIGMHC61	H.sapiens mRNA for IgM heavy chain constant region (mAb61)	0
3576.seq	429	gi 186042 gb M74019 HUMIGKW	Human Ig rearranged light-chain mRNA V region.	E-164
3577.seq	238	gi 2982498 emb AL021407 HS13D10	Homo sapiens DNA sequence from PAC 13D10 on chromosome 6p22.3-23.	5.00E-98
3578.seq	489	gi 1255794 gb U53331.1 U53331	Homo sapiens cosmid clone UT29A10 from Xp22.1-22.2, complete sequence	8.00E-74
3580.seq	478	gi 4759281	Homo sapiens ubiquitin-conjugating enzyme E2L 6 (UBE2L6) mRNA	E-140
3583.seq	470	gi 5042231 emb Y13323.2 HSDISPRO	Homo sapiens mRNA for disintegrin-protease	0
3584.seq	334	gi 3702265 gb AC005797 AC005797	Homo sapiens chromosome 19, cosmid F13825, complete sequence [Homo	1.4
3586.seq	111			
3588.seq	461	gi 3328191 gb AF074604 AF074604	Homo sapiens connective tissue growth factor-related protein	0
3589.seq	480	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-172
3590.seq	421	gi 4506638	Homo sapiens ribosomal protein L35a (RPL35A) mRNA	0
3591.seq	189	gi 4503070	Homo sapiens cold shock domain protein A (CSDA) mRNA	4.00E-27
3592.seq	342	gi 546798 gb S71043 S71043	Ig alpha 2-immunoglobulin A heavy chain allotype 2 (constant	E-141
3593.seq	475	gi 4502492	Homo sapiens complement component 1, r subcomponent (C1R) mRNA	0
3594.seq	506	gi 3242766 gb AC005153 AC005153	Homo sapiens PAC clone DJ0537P09 from 7p11.2-p12, complete sequence [H	1.00E-04
3596.seq	502			
3597.seq	413	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	2.00E-71
3598.seq	463	gi 349475 gb L16785 HUMPUF	Homo sapiens c-myc transcription factor (puf) mRNA, complete cds.	0
3600.seq	290	gi 6137796	Human mitochondrion, complete genome	E-134
3601.seq	133	gi 1256836 gb U53209 HSU53209	Human transformer-2 alpha (htra-2 alpha) mRNA, complete cds	2.00E-53
3602.seq	490	gi 3954884 emb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	0
3606.seq	463	gi 3954884 emb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	E-157
3607.seq	420	gi 1054740 emb X87344 HSEVMHC	H.sapiens DMA, DMB, HLA-Z1, IPP2, LMP2, TAP1, LMP7, TAP2, DOB, DQB2	0
3608.seq	493	gi 13683 emb V00710 MIT1HS	Human mitochondrial genes for several tRNAs (Phe, Val, Leu) and 12S	0
3609.seq	468	gi 4210508 gb AC006287 AC006287	Homo sapiens, clone hRPK22_A_1, complete sequence [Homo sapiens]	2
3610.seq	354	gi 2665724 gb AF035429 AF035429	Homo sapiens cytochrome oxidase subunit I (COI) and subunit II (COII)	0
3611.seq	480	gi 4506700	Homo sapiens ribosomal protein S23 (RPS23) mRNA	0
3612.seq	513	gi 13000 emb X62996 MIHSGENOM	H.sapiens mitochondrial genome (consensus sequence)	0
3613.seq	484	gi 3387979 gb AF070600 AF070600	Homo sapiens clone 24464 beta-tubulin mRNA, complete cds	0
3614.seq	474	gi 5729747	Homo sapiens butyrophilin-like receptor (BTNLR) mRNA	5.00E-04
3615.seq	464	gi 4505604	Homo sapiens pancreatitis-associated protein (PAP) mRNA	0
3616.seq	424			
3617.seq	436	gi 4506700	Homo sapiens ribosomal protein S23 (RPS23) mRNA	0
3619.seq	481	gi 2665724 gb AF035429 AF035429	Homo sapiens cytochrome oxidase subunit I (COI) and subunit II (COII)	0
3621.seq	365	gi 2323213 gb AF004709 AF004709	Homo sapiens stress-activated protein kinase 4 mRNA, complete cds	0.39
3623.seq	359	gi 6017903 gb AF140690.1 AF140690	Homo sapiens melusin mRNA, complete cds	0.096
3624.seq	461	gi 2780183 emb Z99496 HS509L4	Human DNA sequence from PAC 509L4 on chromosome 6q22.1-6q22.33.	0.49
3625.seq	493	gi 4906969 dbj D17652 HUMHBL2A	Human mRNA for HbpsiL22, complete cds	0
3626.seq	466	gi 3387917 gb AF070553 AF070553	Homo sapiens clone 24782 unknown mRNA	0
3627.seq	466	gi 5102744 emb AL079283.1 HS2000202	Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 39515	0
3628.seq	538	gi 4902677 emb AL049801.1 HS650191	Novel human gene mapping to chromosome 13, similar to rat HOGAP	0.15
3629.seq	504	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
3630.seq	491	gi 4501886	Homo sapiens actin, gamma 1 (ACTG1) mRNA	0
3631.seq	508	gi 4557230	Homo sapiens acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight	0
3632.seq	489	gi 184233 gb M12623 HUMHMG17	Human non-histone chromosomal protein HMG-17 mRNA, complete cds.	0
3633.seq	307	gi 2765422 emb Y14736 HSIGG1KL	Homo sapiens mRNA for immunoglobulin kappa light chain	E-170
3634.seq	519	gi 4827106 dbj AP000140.1 AP000140	Homo sapiens genomic DNA, chromosome 21q21.2, LL56-APP region, clone	0.036
3635.seq	512	gi 34137 emb Y07511 HHSKOAT	Human mRNA for kidney ornithine aminotransferase (EC 2.6.1.13)	0
3636.seq	199	gi 5912009 emb AL117502.1 HSM801015	Homo sapiens mRNA; cDNA DKFZp434D0935 (from clone DKFZp434D0935)	E-102
3637.seq	423	gi 546798 gb S71043 S71043	Ig alpha 2-immunoglobulin A heavy chain allotype 2 (constant region,	0
3638.seq	498	gi 33200 emb Z11894 HSIGK37S4	H.sapiens rearranged mRNA for immunoglobulin kappa chain (VNJ)	0
3640.seq	503	gi 2665724 gb AF035429 AF035429	Homo sapiens cytochrome oxidase subunit I (COI) and subunit II	0
3642.seq	553	gi 5757537 gb AC005682.2 AC005682	Homo sapiens clone NH084K19, complete sequence	E-158
3643.seq	487	gi 2853173 emb Y13286 HSY13286	Homo sapiens mRNA for GDP dissociation inhibitor beta	0
3644.seq	509	gi 3954892 emb AJ010446 HSA010446	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	0
3645.seq	494	gi 4759097	Homo sapiens splicing factor, arginine/serine-rich (transformer 2)	0
3646.seq	495	gi 3132352 dbj AP000042.1 AP000042	Homo sapiens genomic DNA, chromosome 21q22.1, segment 13/28, complete	0.14
3647.seq	496	gi 4505604	Homo sapiens pancreatitis-associated protein (PAP) mRNA	0
3648.seq	295	gi 1296606 emb X97074 HSAP17	H.sapiens mRNS for clathrin-associated protein	E-142
3650.seq	330	gi 2275182 gb AC002385 AC002385	Human BAC clone RG222A16 from 7q31, complete sequence [Homo sapiens]	0.088
3651.seq	393	gi 5174430	Homo sapiens DNA segment on chromosome X (unique) 648 expressed	0
3652.seq	612	gi 3922819 emb Z68925 HSKPOLC01	H.sapiens mRNA for immunoglobulin kappa light chain VJ region (ID	0
3655.seq	659	gi 5304861 emb Z98884.11 HS467L1	Human DNA sequence from clone 467L1 on chromosome 1p36.21-36.33,	0
3656.seq	660	gi 35456 emb X54629 HSPHL1	Human pHL-1 gene is a c-myc oncogene containing coxIII sequence	0
3657.seq	482	gi 1916025 gb U77376 OLU77376	Oryzias latipes zinc finger protein sal mRNA, partial cds	0.13
3658.seq	583	gi 4504898	Homo sapiens karyopherin alpha 3 (importin alpha 4) (KPNA3) mRNA	0
3659.seq	384	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	0
3660.seq	666	gi 340184 gb L08599 HUMUVOECAD	Human uvomorulin (E-cadherin) (UVO) mRNA, complete cds.	0
3661.seq	503	gi 4506702	Homo sapiens ribosomal protein S24 (RPS24a) mRNA	0
3662.seq	642	gi 4835595 dbj AP000226.1 AP000226	Homo sapiens genomic DNA, chromosome 21q21.2, LL56-APP region,	0
3663.seq	742	gi 3513299 gb AC005595 AC005595	Homo sapiens chromosome 19, cosmid F16601, complete sequence [Homo	0.052
3664.seq	637	gi 434050 gb L12535 HUMRSU1A	Human RSU-1/RSP-1 mRNA, complete cds.	0
3665.seq	666	gi 5001540 gb AC004961.2 AC004961	Homo sapiens clone DJ1098J04, complete sequence	0.046
3666.seq	604	gi 3080543 dbj AB001451 AB001451	Homo sapiens mRNA for Skc, partial cds	0
3667.seq	637	gi 5306303 gb AC007240.2 AC007240	Homo sapiens clone NH0083M08, complete sequence	0.69
3670.seq	631	gi 4581418 emb AL035402.7 HS88J8	Human DNA sequence from clone 88J8 on chromosome 6p21.31-21.33.	E-174
3671.seq	678	gi 4502162	Homo sapiens apolipoprotein D (APOD) mRNA	0
3672.seq	821	gi 184918 gb M60192 HUMIGHA2A	Human germline IgA2 chain C region.	E-169
3673.seq	228	gi 4506006	Homo sapiens protein phosphatase 1, catalytic subunit, gamma	3.00E-25
3674.seq	571	gi 2665724 gb AF035429 AF035429	Homo sapiens cytochrome oxidase subunit I (COI) and subunit II (COII)	0
3675.seq	733	gi 190978 gb M18963 HUMREGA	Human islet of Langerhans regenerating protein (reg) mRNA, complete	0
3676.seq	672	gi 562764 emb X80169 MMR200KP	M.musculus mRNA for 200 kD protein	0
3677.seq	682	gi 182927 gb M37131.1 HUMGACTA2	Human gamma-actin pseudogene with retroviral insertion site, 3'	0
3678.seq	569	gi 4585468 gb AF127036.1 AF127036	Homo sapiens calcium-activated chloride channel protein 1 (CaCC1)	0
3679.seq	629	gi 11946342 gb U88567 MMU88567	Mus musculus secreted frizzled related protein sFRP-2 (Sfrp2) mRNA,	7.00E-35
3680.seq	571	gi 1246237 gb L48722 HUMPTPIVGA	Homo sapiens (clone hh18) protein tyrosine phosphatase (ptp-1v1r)	0
3681.seq	182	gi 1871209 gb U91327 HSU91327	Human chromosome 12p15 BAC clone CIT987SK-99D8 complete sequence	0.012
3682.seq	400	gi 5763740 emb AL050350.14 HSDJ261K	Human DNA sequence from clone 261K5 on chromosome 6p21-22.1. Contains	0.11
3683.seq	364	gi 3281967 emb Z99129 HS425C14	Human DNA sequence from clone 425C14 on chromosome 6q22 Contains the	0
3684.seq	626	gi 5360215	Homo sapiens iduronate 2-sulfatase (Hunter syndrome) (IDS).	0
3685.seq	548	gi 182059 gb M16653 HUMELAP2B	Human pancreatic elastase IIb mRNA, complete cds.	6.00E-35
3686.seq	637	gi 1263363 gb L77612 HUM17QYCAN	Homo sapiens (clone SEL366) 17q YAC (368C7) RNA.	9.00E-93
3687.seq	626	gi 3954892 emb AJ010446 HSA010446	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	0
3688.seq	575	gi 1035862 emb Z63484 HS85A9R	H.sapiens CpG island DNA genomic MseI fragment, clone 85a9, reverse	0.04
3690.seq	642	gi 790818 gb L39891 HUMPKD1GEN	Homo sapiens polycystic kidney disease-associated protein (PKD1)	3.00E-09
3691.seq	734	gi 4557662	Homo sapiens immunoglobulin-binding protein 1 (IGBP1) mRNA	0
3692.seq	628	gi 4106997 gb AC006299 AC006299	Homo sapiens, clone hRPK17_A_1, complete sequence [Homo sapiens]	0.68
3693.seq	571	gi 2623584 gb AF027158 AF027158	Homo sapiens immunoglobulin kappa light chain (T6J/k) mRNA, partial	E-132
3694.seq	958	gi 432358 dbj D23660 HUMRSP	Human mRNA for ribosomal protein, complete cds	E-139

3695.seq	299	gi 3451027 gb IM88649 DOGADNCCYC	Canis familiaris adenylyl cyclase type V mRNA, partial cds	2.00E-70
3696.seq	455	gi 5453560	Homo sapiens ATP synthase, H ⁺ transporting, mitochondrial F1F0	0
3697.seq	528	gi 49721121 emb AL049776.2 CNS00001	Human chromosome 14 DNA sequence *** IN PROGRESS *** BAC R-85K15	0.14
3699.seq	596	gi 2623584 gb AF027158 AF027158	Homo sapiens immunoglobulin kappa light chain (T6J/k) mRNA, partial	0
3702.seq	583	gi 2952305 gb AF049587 AF049587	Canis familiaris tropomyosin (TPM1) gene, partial cds	0.19
3703.seq	543	gi 13683 emb V00710 MIT1HS	Human mitochondrial genes for several tRNAs (Phe, Val, Leu) and 12S	0
3704.seq	822	gi 4758633	Homo sapiens Sec24b, S. Cerevisiae, homolog of (KIAA0079) mRNA	0
3705.seq	252	gi 5174474	Homo sapiens interferon-stimulated transcription factor 3, gamma	2.00E-85
3706.seq	639	gi 49644 emb X62678 MAP5PROMR	M. auratus mRNA for P5 protein	1.00E-05
3707.seq	373	gi 32998 emb X07868 HSIGF27	Human DNA for insulin-like growth factor II (IGF-2): exon 7 and	E-126
3708.seq	615	gi 3342263 gb AF051561 AF051561	Rattus norvegicus Na-K-C1 cotransporter (Nkcc1) mRNA, complete cds	8.00E-53
3709.seq	575	gi 5578945 emb AL080273.10 HSA259F	Human DNA sequence from clone 259P1 on chromosome 22. Contains STSS,	0.62
3710.seq	499	gi 4895146 gb AC007115.1 AC007115	Homo sapiens chromosome 12 clone 91705, complete sequence	E-156
3712.seq	335	gi 1769890 emb X99747 ATATB2	A.thaliana ATB2 gene	0.09
3713.seq	584	gi 5453558	Homo sapiens ATP synthase, H ⁺ transporting, mitochondrial F1F0,	0
3714.seq	511	gi 1591300 gb U67508 U67508	Methanococcus jannaschii section 50 of 150 of the complete genome	0.55
3715.seq	622	gi 4884198 emb AL049951.1 HSM80029	Homo sapiens mRNA; cDNA DKFZp564O0122 (from clone DKFZp564O0122)	E-140
3716.seq	728	gi 2271030 gb AF008933 AF008933	Homo sapiens MHC class I antigen HLA-A heavy chain (HLA-A*0226) mRNA,	0
3717.seq	602	gi 986883 gb M25388 HUMMTNUBA	Human nuclear-encoded mitochondrial NADH-ubiquinone reductase 24Kd	0
3718.seq	663	gi 187131 gb M25627 HUMJLGTI1	Human liver glutathione S-transferase subunit 1 mRNA, complete cds.	0
3719.seq	428	gi 5065863 emb AL035455.29 HS1018E9	Human DNA sequence from clone 1018E9 on chromosome 20q13.31-13.33,	0.12
3720.seq	610	gi 4502692	Homo sapiens CD9 antigen (p24) (CD9) mRNA	0
3721.seq	580	gi 4502032	Homo sapiens aldehyde dehydrogenase 2, mitochondrial (ALDH2) mRNA	0
3722.seq	444	gi 460086 gb U01925 HSU01925	Human BTK region clone 2F10-rpi mRNA.	9.00E-86
3723.seq	871	gi 499183 emb X7682F DNEUR	F.domestica mRNA for neuronal protein	2.00E-54
3724.seq	348	gi 5257006 gb AF084457.1	Homo sapiens beta-cop homolog mRNA, complete cds	0
3725.seq	600	gi 2780183 emb Z99496 HS509L4	Human DNA sequence from PAC 509L4 on chromosome 6q22.1-6q22.33. Con	0.65
3726.seq	644	gi 4544348 gb AC006059.3 AC006059	Homo sapiens PAC RPC14-613B23 (Roswell Park Cancer Institute Human PA	0
3727.seq	360	gi 4505134	Homo sapiens midkine (neurite growth-promoting factor 2) (MDK) mRNA	0
3730.seq	705	gi 28761 emb X13629 HSAPOA4	Human intestinal mRNA for apolipoprotein A-IV	0
3731.seq	567	gi 557545 gb M28699 HUMB23A	Homo sapiens nucleolar phosphoprotein B23 (NPM1) mRNA, complete	0
3733.seq	612	gi 4757825	Homo sapiens beta-2-microglobulin (B2M) mRNA	0
3734.seq	312	gi 4263638 gb AC000015 AC000015	Homo sapiens chromosome 4 clone B271E1 map 4q25, complete sequence	E-160
3735.seq	497	gi 3387972 gb AF070595 AF070595	Homo sapiens clone 24583 mRNA sequence	0
3738.seq	356	gi 2576343 gb AC002041 HUAC002041	Human Chromosome 16 BAC clone CIT87SK-A-17E1, complete sequence [H	4.00E-84
3739.seq	423	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	9.00E-95
3740.seq	621	gi 1469888 gb D63486 D63486	Human mRNA for KIAA0152 gene, complete cds	0
3741.seq	650	gi 1236166 emb Z70272 HSJ30E17	Human DNA sequence from cosmid J30E17, between markers DXS366 and DX	0.71
3742.seq	862	gi 468256 gb U08032 HSU08032	Human thermolabile (monoamine, M form) phenol sulfotransferase	0
3743.seq	628	gi 38407 emb X67301 HSIGMHC63	H.sapiens mRNA for IgM heavy chain constant region (Ab63)	0
3744.seq	660	gi 5921382 emb AL034552.22 HS824J5	Human DNA sequence from clone B24J5 on chromosome 20q11.23-12. Contai	0.003
3747.seq	617	gi 4757825	Homo sapiens beta-2-microglobulin (B2M) mRNA	0
3748.seq	32	gi 3094988 emb AL008628 HS140C12	Homo sapiens DNA sequence from PAC 140C12 on chromosome 6q26-q27	2.00E-08
3749.seq	191	gi 4503658	Homo sapiens Finkel-Biskis-Relly murine sarcoma virus (FBR-MuSV)	4.00E-12
3750.seq	592	gi 3860070 gb AF097021 AF097021	Homo sapiens GW112 protein (GW112) mRNA, complete cds	0
3752.seq	478	gi 37564 emb X56998 HSUBA52A	Human Uba52 adrenal mRNA for ubiquitin-52 amino acid fusion protein	0
3755.seq	675	gi 2337866 gb AC002463 AC002463	Human BAC clone RG302F04 from 7q31, complete sequence [Homo sapiens]	0.047
3756.seq	579	gi 4507356	Homo sapiens transgelin 2 (TAGLN2) mRNA	0
3758.seq	572	gi 4680714 gb AF132972.1 AF132972	Homo sapiens CGI-38 protein mRNA, complete cds	0
3760.seq	544	gi 4826658	Homo sapiens capping protein (actin filament) muscle Z-line, beta	0
3761.seq	605	gi 3335023 gb U41448 HSU41448	Homo sapiens ribosomal protein S26 (RPS26) gene, complete cds	4.00E-36
3762.seq	121	gi 976086 emb X89763 MTHSDNAL2	H.sapiens mitochondrial DNA for DNA loop attachment sequences	4.00E-60
3764.seq	528	gi 3646129 emb AJ010842 HSAJ010842	Homo sapiens mRNA for putative ATP(GTP)-binding protein, partial	0
3766.seq	421	gi 2653426 emb Z99127 HS102G20	Human DNA sequence from PAC 102G20 on chromosome 1q24-q25. Contains	0.11
3767.seq	569	gi 185786 gb U1410 HUMIGHZ2A	Homo sapiens Ig rearranged gamma-1 chain (IGHG1) mRNA subgroup	E-144
3768.seq	647	gi 4557374	Homo sapiens basic transcription element binding protein 1 (BTEB1)	0
3769.seq	239	gi 4506648	Homo sapiens ribosomal protein L3 (RPL3) mRNA	E-113
3770.seq	511	gi 2765426 emb Y14738 HSIGG1LL	Homo sapiens mRNA for immunoglobulin lambda light chain	0
3771.seq	502	gi 5771401 gb AF096771.1 HSMCLKKRF	Homo sapiens myosin light chain kinase (MYLK) gene, exon 31, and	0
3772.seq	497	gi 188469 gb K01144 HUMMHGM	Human major histocompatibility class II antigen gamma chain mRNA,	0
3773.seq	362	gi 1754834 gb U19495 HSU19495	Human intercrine-alpha (hIRH) mRNA, complete cds	E-147
3774.seq	426	gi 185549 gb M18518 HUMIGHVAP	Human Ig rearranged H-chain VDJ-region mRNA, clone 58P2.	E-144
3776.seq	488			
3777.seq	227	gi 28930 emb X05606 HSA TPB	Human mRNA fragment for ATP synthetase beta-subunit	E-117
3778.seq	225	gi 4506430	Homo sapiens RAS p21 protein activator (GTPase activating protein) 1	7.00E-48
3779.seq	409	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	3.00E-91
3780.seq	195	gi 4884135 emb AL050107.1 HSM80019	Homo sapiens mRNA; cDNA DKFZp5861419 (from clone DKFZp5861419)	1.00E-95
3781.seq	411	gi 2632101 emb Z93943 HSU235H3	Human DNA sequence from cosmid U235H3 on chromosome X	1.7
3782.seq	339	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	8.00E-70
3783.seq	550	gi 4504444	Homo sapiens heterogeneous nuclear ribonucleoprotein A1 (HNRPA1)	0
3784.seq	12			
3785.seq	510			
3786.seq	344	gi 3005086 gb AF044924 AF044924	Homo sapiens hook2 protein (HOOK2) mRNA, complete cds	E-152
3787.seq	451	gi 2358130 gb AC000084 AC000084	Homo sapiens Chromosome 22q11.2 Cosmid Clone 68f in DGCR Region,	0.12
3789.seq	361	gi 5031570	Homo sapiens ARP2 (actin-related protein 2, yeast) homolog (ACTR2)	E-120
3790.seq	525	gi 5231227	Homo sapiens ribonuclease 6 precursor (RNASE6PL) mRNA	0
3791.seq	523	gi 1944628 gb J01415.1 HUMMTCG	Human mitochondrion, complete genome	0
3792.seq	458	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
3795.seq	417	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, complete cds.	2.00E-95
3796.seq	567	gi 4503252	Homo sapiens defender against cell death 1 (DAD1) mRNA	0
3797.seq	433	gi 28335 emb X63342 HSACTB	H.sapiens ACTB mRNA for mutant beta-actin (beta-actin)	0
3798.seq	478	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-112
3799.seq	518	gi 2337882 gb AC002451 AC002451	Human BAC clone GS306C12 from 7q21-q22, complete sequence [Homo	4.00E-85
3800.seq	653	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-174
3801.seq	379	gi 187538 gb M10942 HUMMETIE	Human metallothionein-le gene (hMT-le).	E-118
3802.seq	322	gi 3387987 gb AF070606 AF070606	Homo sapiens clone 24411 mRNA sequence	E-155
3803.seq	558	gi 265703 gb S55735 S55735	Ig alpha 1 - alpha 2m-immunoglobulin A1-A2 lambda hybrid GAU heavy	0
3804.seq	511	gi 4506666	Homo sapiens ribosomal protein, large, P0 (RPLP0) mRNA	0
3805.seq	492	gi 4501888	Homo sapiens actin, gamma 2, smooth muscle, enteric (ACTG2) mRNA	0
3806.seq	533	gi 339688 gb M17733 HUMTHYB4	Human thymosin beta-4 mRNA, complete cds.	0
3807.seq	399			
3808.seq	643	gi 1213466 gb U29057 HSU29057	Human MHC class I antigen HLA-B7 variant (HLA-B) mRNA, complete	0
3809.seq	647	gi 340233 gb M25246 HUMVIM3	Human vimentin (HuVim3) mRNA, 3' end.	0
3810.seq	637	gi 3138923 gb AF002282 AF002282	Homo sapiens alpha-actinin-2 associated LIM protein mRNA,	0
3811.seq	283	gi 4558768 gb AC007199.1 AC007199	Homo sapiens chromosome 5 BAC clone 111n13, complete sequence	E-145
3812.seq	415	gi 555853 gb U13369 HSU13369	Human ribosomal DNA complete repeating unit	E-132
3813.seq	294	gi 5802987	Homo sapiens B-cell translocation gene 2 (pheochromocytoma cell-3)	E-139
3814.seq	599	gi 4868168 gb AF127026.1 AF127026	Homo sapiens brush border myosin I (BBMI) mRNA, complete cds	0
3815.seq	569	gi 4758767	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 10	0
3817.seq	597	gi 185538 gb M18512 HUMIGHYAJ	Human (fetal) rearranged IgM chain VDJ-region mRNA, clone 20P3.	E-162
3818.seq	369	gi 1262342 emb X93334 HSMITG	H.sapiens mitochondrial DNA, complete genome	0
3819.seq	578	gi 4506616	Homo sapiens ribosomal protein L17 (RPL17) mRNA	0
3820.seq	511	gi 3005936 emb AJ004956 HSA004956	Homo sapiens Immunoglobulin kappa light chain variable region	0
3824.seq	568	gi 5031664	Homo sapiens Dmx-like 1 (DMXL1) mRNA >gi 3123571 emb AJ005821 HSA58	2.00E-93
3825.seq	537	gi 4506672	Homo sapiens ribosomal protein, mitochondrial, L12 (RPLM12) mRNA	0
3826.seq	519	gi 4156194 gb AC004869 AC004869	Homo sapiens PAC clone DJ0732C22 from 7p11.2-p13, complete sequence [H	2.2
3827.seq	591	gi 3851580 gb AF091215 AF091215	Mus musculus Wrm protein (Wrm) gene, complete cds	0.041
3828.seq	592	gi 3882204 gb AB018285.1 AB018285	Homo sapiens mRNA for KIAA0742 protein, partial cds	0

3829.seq	566	gi 1944628 gb J01415.1 HUMMTCG	Human mitochondrion, complete genome	0
3830.seq	562	gi 5360972 emb AL035563.1 HS1068E	Human DNA sequence from clone 1068E13 on chromosome 20p11.21-12.3.	0.61
3832.seq	524	gi 37570 emb X63237 HSUBA80R	H.sapiens Uba80 mRNA for ubiquitin	0
3834.seq	457	gi 4505356	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 4	0
3835.seq	528	gi 5454161	Homo sapiens upregulated by 1,25-dihydroxyvitamin D-3 (VDUPT1) mRNA	0
3836.seq	507	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-116
3837.seq	466	gi 1926407 gb U20938 HSU20938	Human lymphocyte dihydropyrimidine dehydrogenase mRNA, complete cds	0
3838.seq	489	gi 414346 db J014531 HUMHRRPL9	Human mRNA for human homologue of rat ribosomal protein L9.	0
3839.seq	450	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	4.00E-97
3840.seq	500	gi 4506730	Homo sapiens ribosomal protein S6 (RPS6) mRNA	0
3843.seq	546	gi 4758767	Homo sapiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 10	0
3844.seq	580	gi 2280486 db AB002375 AB002375	Human mRNA for KIAA0377 gene, complete cds	0
3846.seq	534	gi 5771401 gb AF096771.1 HSMCLKKRF	Homo sapiens myosin light chain kinase (MYLK) gene, exon 31, and	E-158
3847.seq	549	gi 4884115 emb AL050093.1 HSM80016E	Homo sapiens mRNA; cDNA DKFZp586L0518 (from clone DKFZp586L0518)	E-165
3848.seq	499	gi 36888360 emb AL031803 HS22C1D	Human DNA sequence from clone 27C10 on chromosome Xp11.4-21.3. Contain	0.14
3849.seq	565	gi 179295 gb L08441 HUMAUTONH	Human autonomously replicating sequence (ARS) mRNA.	0
3850.seq	642	gi 4914577 gb emb AL050373.1 HSM80015E	Homo sapiens mRNA; cDNA DKFZp586F1318 (from clone DKFZp586F1318)	0
3851.seq	514	gi 4502596	Homo sapiens collagen-binding protein 2 (collgen 2) (CBP2) mRNA	0
3852.seq	625	gi 285982 db D13635 HUMRSC1083	Human mRNA for KIAA0010 gene, complete cds	0
3853.seq	678	gi 4579988 db JAP000067.1 JAP000067	Homo sapiens genomic DNA of 8p11.2 senescence gene, segment 3/19.	2.9
3854.seq	602	gi 6016843 db JAP000352.2 JAP000352	Homo sapiens genomic DNA, chromosome 22q11.2, clone KB1561E1	0.65
3855.seq	661	gi 4467346 emb AL022323.7 HS243E7	Human DNA sequence from clone 243E7 on chromosome 22q12.1. Contains E5	0.18
3856.seq	529	gi 4506618	Homo sapiens ribosomal protein L24 (RPL24) mRNA	0
3858.seq	606	gi 388185 gb L22473 HUMBAXA	Human Bax alpha mRNA, complete cds.	0
3859.seq	334	gi 3135974 emb Z99128 HS422H11	Homo sapiens DNA sequence from PAC 422H11 on chromosome 6p21.1-22.2.	0.023
3860.seq	568	gi 189019 gb M22918 HUMMYLCB	Human non-muscle myosin alkali light chain mRNA, 3' end.	0
3861.seq	525	gi 6102666 gb ACD06449.19 ACD06449	Homo sapiens chromosome 17, clone hCIT.58_E.17, complete sequence	2.2
3862.seq	589	gi 5733597 gb AF159456.1 AF159456	Homo sapiens gp-340 variant protein (DMBT1) mRNA, complete cds	0
3863.seq	248	gi 187245 gb M21119 HUMLSZA	Human lysozyme mRNA, complete cds.	1.00E-64
3864.seq	604	gi 5881960 gb AF064257.1	Homo sapiens Dhm1-like protein mRNA, complete cds	0
3865.seq	207	gi 5901941	Homo sapiens E1B-55kDa-associated protein 5 (E1B-AP5), mRNA	2.00E-91
3866.seq	427	gi 4503304	Homo sapiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	0
3867.seq	480	gi 4885262	Homo sapiens GTP-binding protein overexpressed in skeletal muscle	E-138
3868.seq	645	gi 5924054 emb AL096709.1 HSDJ20M	Human DNA sequence *** SEQUENCING IN PROGRESS *** from clone dJ20M	0.045
3869.seq	568	gi 183655 gb J03746 HUMGST	Human glutathione S-transferase mRNA, complete cds.	0
3871.seq	370	gi 4689113 gb AF077033.1 AF077033	Homo sapiens putative ATP-dependent RNA helicase ROK1 mRNA,	0.39
3872.seq	626	gi 2599359 gb AF015812 AF015812	Homo sapiens RNA helicase p68 (HUMP68) gene, complete cds	0
3873.seq	262	gi 4240220 db AB020673.1 AB020673	Homo sapiens mRNA for KIAA0866 protein, complete cds	2.00E-42
3874.seq	444	gi 6273777 gb AF141968.1 AF141968	Homo sapiens trabeculin-alpha mRNA, complete cds	0
3875.seq	314	gi 32822 emb X57331 HSIGCMUDE	Human immunoglobulin C (mu) and C(delta) heavy chain genes (constant	E-167
3876.seq	564	gi 4505940	Homo sapiens polymerase (RNA) II (DNA directed) polypeptide B (140kD)	0
3878.seq	474	gi 3849824 gb ACD05900 ACD05900	Homo sapiens chromosome 17, clone hRPK.998_F_8, complete sequence [Homo	0.13
3879.seq	534	gi 4588663 gb U04245 HSU04245	Human MHC class I antigen HLA-B (HLA-B*0704 allele) mRNA, complete	0
3880.seq	618	gi 249370 gb S35960 S35960	laminin receptor homolog (3' region) [human, mRNA Partial, 739 nt]	0
3881.seq	413	gi 4666255 db JAP000009.2 JAP000009	Homo sapiens genomic DNA of 21q22.2 Down Syndrome region, segment 2/12	0
3882.seq	637	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
3883.seq	577	gi 2772533 gb ACD003992 ACD003992	Human BAC clone RG250D13 from 7q31, complete sequence [Homo sapiens]	0.04
3884.seq	603	gi 1663703 db D87684 D87684	Human mRNA for KIAA0242 gene, partial cds	0
3885.seq	387	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
3886.seq	642	gi 4885590	Homo sapiens selenoprotein P, plasma, 1 (SEPP1) mRNA	0
3887.seq	166	gi 3283903 gb AF070634 AF070634	Homo sapiens clone 24725 antizyme inhibitor mRNA, partial cds	7.00E-84
3889.seq	303			0
3890.seq	461	gi 4506706	Homo sapiens ribosomal protein S25 (RPS25) mRNA	0
3891.seq	645	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-141
3892.seq	606	gi 3882246 db JAB018306.1 JAB018306	Homo sapiens mRNA for KIAA0763 protein, complete cds	0
3893.seq	629	gi 4758541	Homo sapiens heterogeneous nuclear ribonucleoprotein A/B (HNRPAB)	0
3894.seq	581	gi 432974 gb M75883 HUMSCP2A	Human sterol carrier protein X/sterol carrier protein 2 mRNA,	0
3895.seq	661	gi 6048973 gb AF103803.1 AF103803	Homo sapiens clone H41 unknown mRNA	0
3896.seq	631	gi 2052040 emb J284483 HS46H23	Human DNA sequence from PAC 46H23, BRCA2 gene region chromosome 13c	0.69
3897.seq	613	gi 6094661 gb AC005225.2 AC005225	Homo sapiens clone J0414A15, complete sequence	2.00E-13
3898.seq	530	gi 4502134	Homo sapiens N-acetylaminoacyl-peptide hydrolase (APEH) mRNA	0
3899.seq	563			0
3900.seq	869	gi 12583 emb X15759 HSCOXII	H.sapiens mitochondrial mRNA for cytochrome c oxidase subunit II	0
3901.seq	576	gi 4758137	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 5 (RNA	0
3902.seq	609	gi 3043595 db JAB011108 JAB011108	Homo sapiens mRNA for KIAA0536 protein, partial cds	2.00E-62
3903.seq	583	gi 5174520	Homo sapiens mannosidase, alpha, class 1A, member 1 (MAN1A1) mRNA	0
3904.seq	627	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
3905.seq	501	gi 4902764 emb AL049550.5 HSDJ90L14	Human DNA sequence from clone 90L14 on chromosome 6q14.1-15 Contains	0.14
3906.seq	593	gi 5803099	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 18	0
3907.seq	389	gi 4507512	Homo sapiens tissue inhibitor of metalloproteinase 3 (Sorsby fundus	0
3909.seq	608	gi 4468336 emb AL033533.5 HS973M2	Human DNA sequence from clone 973M2 on chromosome 1q24.3-31.1 Contain	0.66
3910.seq	608	gi 5020252 gb AF150962.1 AF150962	Homo sapiens repressor of estrogen receptor activity (REA) mRNA,	0
3911.seq	661	gi 5453805	Homo sapiens NS1-associated protein 1 (NSAP1) mRNA	0
3913.seq	561	gi 4925990 gb AF151819.1 AF151819	Homo sapiens CGI-61 protein mRNA, complete cds	E-160
3914.seq	552	gi 3522917 gb AC005601 AC005601	Homo sapiens chromosome 5, BAC clone 343g16 (LBNL H180), complete	2.4
3915.seq	613	gi 3688087 gb AC005758 AC005758	Homo sapiens chromosome 19, cosmid R27946, complete sequence [Homo	6.00E-32
3916.seq	488	gi 186522 gb L10284 HUMIP90	Homo sapiens integral membrane protein, calnexin, (IP90) mRNA,	1.00E-38
3917.seq	463	gi 5031924	Homo sapiens megakaryocyte stimulating factor (MSF) mRNA	3.00E-21
3918.seq	306	gi 3540152 gb AC005613 AC005613	Homo sapiens chromosome 5, BAC clone 114k9 (LBNL H94), complete	0.021
3919.seq	553	gi 186840 gb J03799 HUMLAMB	Human collagen laminin-binding protein mRNA, complete cds.	0
3920.seq	540	gi 546798 gb S710433 S71043	Ig alpha 2-immunoglobulin A heavy chain allotype 2 (constant	E-152
3921.seq	583	gi 265703 gb S55735 S55735	Ig alpha 1 - alpha 2m-immunoglobulin A1-A2 lambda hybrid GAU heavy	0
3922.seq	731	gi 3169112 emb AL021997 HSB74C20	Homo sapiens DNA sequence from PAC 874C20 on chromosome 6p22.1-22.3.	0.8
3923.seq	590	gi 5123976 gb AC006552.7 AC006552	Homo sapiens chromosome 4 clone C0084L10 map 4p16, complete sequence	0.16
3925.seq	736	gi 2394273 gb AF017061 AF017061	Homo sapiens vasopressin-activated calcium mobilizing putative	0
3926.seq	598	gi 1469196 db J050927 J050927	Human mRNA for KIAA0137 gene, complete cds	0
3927.seq	526	gi 2735037 gb U82696 U82696	Homo sapiens chromosome X from Xq28, complete sequence [Homo sapiens]	0.036
3930.seq	565	gi 12583 emb X15759 HSCOXII	H.sapiens mitochondrial mRNA for cytochrome c oxidase subunit II	0
3931.seq	571	gi 3264535 emb AL021920 HS163M9	Homo sapiens DNA sequence from PAC 163M9 on chromosome 1p35.1-p36.21	E-136
3932.seq	623	gi 5174602	Homo sapiens myosin regulatory light chain 2, smooth muscle isoform	0
3933.seq	543	gi 3165402 gb AC004752 AC004752	Homo sapiens chromosome 5, BAC clone 8e5 (LBNL H167), complete sequenc	0
3934.seq	599	gi 4504618	Homo sapiens insulin-like growth factor binding protein 7 (IGFBP7)	0
3935.seq	669	gi 167414 gb M67448 CRECYTC6	C.reinhardtii cytochrome c6 gene, complete cds.	0.18
3936.seq	549	gi 3204432 emb AL021878 HS257120	Homo sapiens DNA sequence from PAC 257120 on chromosome 22q13.1-13.2.	E-135
3938.seq	619	gi 4502920	Homo sapiens calponin 1, basic, smooth muscle (CNN1) mRNA	0
3939.seq	543	gi 5123960 gb AC007527.6 AC007527	Homo sapiens 12p12 BAC RPC111-4N23 (Roswell Park Cancer Institute Huma	0.59
3940.seq	317	gi 3892081 gb AC005951 AC005951	Homo sapiens chromosome 17, clone hRPK.372_K_20, complete sequence [Ho	0.084
3941.seq	177	gi 340874 gb M23323 HUMCD3E08	Human membrane protein (CD3-epsilon) gene, exon 9.	2.00E-32
3942.seq	566	gi 184918 gb M60192 HUMIGHA2A	Human germline IgA2 chain C region.	E-169
3943.seq	494	gi 292434 gb L07287 HUMRPL26X	Human ribosomal protein L26 (RPL26) gene exon 2, complete cds.	0
3944.seq	629	gi 1031821 emb J25908 HS17B10R	H.sapiens CpG island DNA genomic MseI fragment, clone 17b10.	2.00E-07
3946.seq	724	gi 1246239 gb L48723 HUMPTPIVGB	Homo sapiens (clone h113) protein tyrosine phosphatase (ptp-IV1a,	0
3947.seq	611	gi 4557500	Homo sapiens cathepsin O (CTSO) mRNA >gi 574803 emb X77383 HSCATHO	0
3948.seq	573	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-174
3949.seq	606	gi 182864 gb J02939 HUMG4F2	Human membrane glycoprotein 4F2 antigen heavy chain mRNA.	0
3950.seq	461	gi 6005797	Homo sapiens synaptotagmin (KIAA1029), mRNA	0
3951.seq	498	gi 1399085 gb U43701 HSU43701	Human ribosomal protein L23a mRNA, complete cds.	0
3952.seq	652	gi 1730464 emb Z83307 HSA1280	Human DNA sequence from cosmid A1280, chromosome region 11p13 Contain	0

3953.seq	685	gi 3860076 gb AF097021 AF097021	Homo sapiens GW112 protein (GW112) mRNA, complete cds	2.00E-07
3954.seq	358	gi 3165399 gb AC003684 AC003684	Homo sapiens Xp22 BAC GSHB-519E5 (Genome Systems Human BAC library)	0.024
3955.seq	545	gi 3483898 gb AF086553 HUMZE12H05	Homo sapiens full length insert cDNA clone ZE12H05	0
3956.seq	620	gi 4506666	Homo sapiens ribosomal protein, large, PO (RPLP0) mRNA	0
3957.seq	705	gi 4503182	Homo sapiens cytochrome b-5 (CYB5), nuclear gene encoding	0
3958.seq	513	gi 4759099	Homo sapiens splicing factor, arginine/serine-rich 11 (SFRS11) mRNA	2.00E-80
3959.seq	651	gi 4826462 emb AL033377.2 HS287G14	Human DNA sequence from clone 287G14 on chromosome 6q23.1-24.3. Contain	0
3960.seq	465	gi 5263010 emb AL022328.21 HS402G11	Human DNA sequence from clone 402G11 on chromosome 22q13.31-13.33	2.00E-86
3961.seq	195	gi 4502162	Homo sapiens apolipoprotein D (APOD) mRNA	2.00E-78
3962.seq	573	gi 13000 emb X62996 MIHSGENOM	H.sapiens mitochondrial genome (consensus sequence)	0
3963.seq	565	gi 185397 gb U03162 HUMIGHHA	Human Ig rearranged gamma-chain mRNA V-region, partial cds.	0
3964.seq	164	gi 4895146 gb AC007115.1 AC007115	Homo sapiens chromosome 12 clone 91705, complete sequence	3.00E-61
3965.seq	530	gi 38407 emb X67301 HSIGMHC63	H.sapiens mRNA for IgM heavy chain constant region (Ab63)	0
3966.seq	560	gi 3482959 gb AC004675 AC004675	Homo sapiens chromosome 17, clone hCIT.186_H_2, complete sequence [Hom	0.15
3967.seq	254	gi 1944628 gb J01415.1 HUMMTCG	Human mitochondrion, complete genome	E-131
3968.seq	283			
3969.seq	504	gi 4758713	Homo sapiens microsomal glutathione S-transferase 3 (MGST3) mRNA	0
3970.seq	379	gi 185538 gb M18512 HUMIGHVAJ	Human (fetal) rearranged IgM chain VDJ-region mRNA, clone 20P3.	E-149
3971.seq	592	gi 6137796	Human mitochondrion, complete genome	0
3972.seq	535	gi 2326193 gb AF000137 BTAF000137	Bos taurus connective tissue growth factor precursor (CTGF) mRNA,	0.58
3973.seq	532	gi 4589679 db JAB023232.1 AB023232	Homo sapiens mRNA for KIAA1015 protein, complete cds	0.58
3974.seq	341	gi 37564 emb X56998 HSUBA52A	Human Uba52 adrenal mRNA for ubiquitin-52 amino acid fusion protein	0
3975.seq	292	gi 2224590 db AB002323 AB002323	Human mRNA for KIAA0325 gene, partial cds	E-133
3976.seq	531	gi 3954884 gb AJ010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	0
3977.seq	565	gi 33330 emb X56510 HSIGKVLU	Human mRNA for Immunoglobulin variable kappa region	0
3978.seq	537	gi 1262342 emb X93334 HSMITG	H.sapiens mitochondrial DNA, complete genome	0
3980.seq	543	gi 475005 db D29808 HUMTALLA1	Human mRNA for T-cell acute lymphoblastic leukemia associated antigen	0
3982.seq	763	gi 5523803 gb AC007367.3 AC007367	Homo sapiens BAC clone NH0518G12 from 2, complete sequence	0.053
3983.seq	324	gi 4680704 gb AF132967.1 AF132967	Homo sapiens CG1-33 protein mRNA, complete cds	E-111
3984.seq	204	gi 5649387 gb AC005998.3 AC005998	Homo sapiens clone DJ0622E21, complete sequence	0.82
3985.seq	268	gi 4506636	Homo sapiens ribosomal protein L34 (RPL34) mRNA	1.00E-62
3986.seq	278	gi 3201899 gb AF067420 AF067420	Homo sapiens SNCT3 protein (SNCT3) mRNA, complete cds	E-146
3987.seq	683	gi 3005747 gb AF055020 AF055020	Homo sapiens clone 24722 unknown mRNA, partial cds	2.9
3988.seq	181	gi 4309945 gb AC006251 AC006251	Homo sapiens chromosome 17, clone hRPK.692_E_18, complete sequence	0.012
3989.seq	637	gi 4835815 gb AC007566.1 AC007566	Homo sapiens clone RG010G05, complete sequence	0
3991.seq	596	gi 4502018	Homo sapiens aldo-keto reductase family 1, member C3 (3-alpha	0
3992.seq	696	gi 249370 gb S35960 S35960	laninin receptor homolog (3' region) [human, mRNA Partial, 739 nt]	0
3993.seq	337	gi 4877279 emb AJ006591.1 HSA6591	Homo sapiens mRNA for cysteine-rich protein	0
3994.seq	611	gi 5902109	Homo sapiens SRY (sex-determining region Y)-box 22 (SOX22), mRNA	0
3995.seq	681	gi 262368 gb S51768 S51768	PAP-H-pancreatitis-associated protein [human, pancreas, mRNA, 797	0
3996.seq	626	gi 13000 emb X62996 MIHSGENOM	H.sapiens mitochondrial genome (consensus sequence)	0
3997.seq	395	gi 6138757 emb Z98752.13 HS13887	Human DNA sequence from clone 13887 on chromosome 20q13.12, complete	2.00E-30
3998.seq	645	gi 1399645 gb U53454 HSU53454	Human chloride ion current inducer protein (CLIC1) mRNA, complete	3.00E-55
3999.seq	348	gi 4506146	Homo sapiens protease, serine, 2 (trypsin 2) (PRSS2) mRNA	0
4000.seq	348	gi 1839333 gb S79979 S79979	ribosomal protein L37 [human, HeLa cells, Genomic/mRNA, 754 nt]	0
4001.seq	563	gi 182513 gb M11147 HUMFERL	Human ferritin L chain mRNA, complete cds.	0
4002.seq	622	gi 6138746 emb AL049828.3 CNS00008	Human chromosome 14 DNA sequence *** IN PROGRESS *** BAC R-124D2	0.68
4003.seq	662	gi 5262703 emb AL080213.1 HSM800740	Homo sapiens mRNA; cDNA DKFZp586i1823 (from clone DKFZp586i1823)	0
4004.seq	427	gi 33140 emb U00557 HSIGK1	C-terminal part of human kappa-immunoglobulin gene coding for amino	0
4005.seq	682	gi 1514580 emb X95747 HSIGKLC5	H.sapiens immunoglobulin kappa chain (clone: 5)	0
4006.seq	396	gi 1281706 gb AF014838 AF014838	Homo sapiens galectin-4 mRNA, complete cds	0
4007.seq	677	gi 3560556 gb AF053641 AF053641	Homo sapiens brain cellular apoptosis susceptibility protein (CSE1)	0
4008.seq	701	gi 5802973	Homo sapiens antioxidant protein 1 (AOP1), mRNA	0
4009.seq	573	gi 5803071	Homo sapiens LIM domain only 4 (LMO4), mRNA	E-168
4010.seq	374	gi 2094790 emb Z83827 HS473J6	Human DNA sequence from PAC 473J6 on chromosome X contains STS	0.1
4011.seq	432	gi 4502472	Homo sapiens B-cell translocation protein 1 (BTGL1) mRNA	0
4012.seq	346	gi 4914506 emb AL031589.10 HS11007	Human DNA sequence from clone 1107 on chromosome 22q13.1-13.2. Contain	0.023
4013.seq	615	gi 4757825	Homo sapiens beta-2-microglobulin (B2M) mRNA	0
4014.seq	450	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	0
4016.seq	240	gi 181962 gb J04617 HUMEF1A	Human elongation factor EF-1-alpha gene, complete cds.	E-132
4018.seq	602	gi 2168138 emb Z96050 HS105D12	Human DNA sequence from PAC 105D12 on chromosome 1, contains FASL	0.042
4019.seq	628	gi 1136401 db D79993 D79993	Human mRNA for KIAA0171 gene, complete cds	0
4020.seq	529	gi 3892081 gb AC005951 AC005951	Homo sapiens chromosome 17, clone hRPK.372_K_20, complete sequence [Ho	0.037
4021.seq	682	gi 29469 emb X62515 HSBMHSP	H.sapiens mRNA for basement membrane heparan sulfate proteoglycan	0.74
4023.seq	527	gi 4827116 db JAP000150.1 JAP000150	Homo sapiens genomic DNA, chromosome 21q22.2, DSCR region, clone	0.14
4024.seq	501	gi 4508123 gb AC006378.2 AC006378	Homo sapiens BAC clone NH0455K9 from 7q21.2-q22, complete sequence	0.54
4026.seq	580	gi 3360413 gb AF052106 AF052106	Homo sapiens clone 23781 mRNA sequence	0.04
4027.seq	682	gi 4508162 gb AC007161.1 AC007161	Homo sapiens BAC clone GS165B14 from 7p22-p21, complete sequence	0.19
4028.seq	429	gi 4757725	Homo sapiens ALL1 fused gene from chromosome 10 (AF10) mRNA	0.12
4029.seq	641	gi 4835680 db JAP000311.1 JAP000311	Homo sapiens genomic DNA, chromosome 21q22.1, D21S226-AML region,	E-121
4030.seq	761	gi 2887430 db AB007888 AB007888	Homo sapiens KIAA0428 mRNA, complete cds	0
4031.seq	674	gi 4176355 gb AC005509 AC005509	Homo sapiens chromosome 4 clone B20N5 map 4q25, complete sequence [Ho	0.19
4032.seq	637	gi 5052312 gb AF116272.1 AF116272	Homo sapiens T-cell activation protein (PGR1) gene, complete cds	0
4034.seq	606	gi 5931954 gb AF124145.1 AF124145	Homo sapiens autocrine motility factor receptor (AMFR) mRNA,	0
4034.seq	697	gi 2765422 emb Y14736 HSIGG1KL	Homo sapiens mRNA for Immunoglobulin kappa light chain	0
4036.seq	606	gi 5410281 gb AF100748.1 HSPC035	Homo sapiens HSPC035 protein mRNA, complete cds	0
4036.seq	670	gi 3738108 gb AC005722 AC005722	Homo sapiens chromosome 17, clone hRPK.311_F_12, complete sequence [Ho	0.047
4037.seq	195	gi 1332489 emb Z70699 HSB2X2G	H.sapiens mRNA (fetal brain cDNA b2_2q)	1.00E-05
4037.seq	697	gi 3668102 gb AC005691 AC005691	Homo sapiens chromosome 17, clone hRPK.28_A_22, complete sequence	3
4038.seq	498	gi 4557386	Homo sapiens complement component 7 (C7) mRNA	0
4038.seq	637	gi 23690 emb X56932 HS23KDHP	H.sapiens mRNA for 23 kD highly basic protein	0
4039.seq	405	gi 36135 emb Z12962 HSRPL41	H.sapiens mRNA for homologue to yeast ribosomal protein L41	0
4042.seq	659	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	E-174
4043.seq	737	gi 5679459 emb AL031057.10 HS498I24	Human DNA sequence from clone 498I24 on chromosome 6p22.1-22.3 Contain	0.81
4044.seq	433	gi 475633 emb AL031681.13 HS862K6	Human DNA sequence from clone 862K6 on chromosome 20q12-13.13. Contain	0
4045.seq	389	gi 2780183 emb Z99496 HS509L4	Human DNA sequence from PAC 509L4 on chromosome 6q22.1-6q22.33.	0
4046.seq	589	gi 4263688 gb AC006502 AC006502	Homo sapiens, clone hRPK.36_A_1, complete sequence [Homo sapiens]	0.64
4047.seq	251	gi 6175373 gb AC004820.2 AC004820	Homo sapiens clone DJ0038I10, complete sequence	0.066
4048.seq	601	gi 339688 gb M17733 HUMTHYB4	Human thymosin beta-4 mRNA, complete cds.	0
4049.seq	642	gi 5360678 db JAB022656.1 AB022656	Homo sapiens mRNA for anti-Entamoeba histolytica immunoglobulin	0
4050.seq	518	gi 4730900 db JAP000131.1 JAP000131	Homo sapiens genomic DNA of 21q22.1, GART and AML1 f43D11-11988 regio	8.00E-99
4051.seq	680	gi 261239 gb S50732 S50732	immunoglobulin M light chain V region-anti-lipid A antibody [human,	0
4052.seq	535	gi 4758203	Homo sapiens dual specificity phosphatase 1 (DUSP1) mRNA	0
4053.seq	188	gi 4467295 emb AL021919.4 HS1045J21	Human DNA sequence from clone 1045J21 on chromosome 1q23.3-24.3. Cont	0.75
4054.seq	591	gi 184847 gb M63438 HUMIGGK	Human Ig rearranged gamma chain mRNA, V-J-C region and complete	0
4055.seq	353	gi 4506636	Homo sapiens ribosomal protein L34 (RPL34) mRNA	0
4056.seq	590	gi 4503470	Homo sapiens eukaryotic translation elongation factor 1 alpha 1	0
4057.seq	584	gi 1000861 gb L47647 HUMCKBR	Homo sapiens creatine kinase B mRNA, complete cds.	0
4058.seq	346	gi 4092478 gb AC006257 AC006257	Homo sapiens, clone hRPK.11_A_1, complete sequence [Homo sapiens]	0.37
4059.seq	604	gi 4519416 db JAB024327.1 AB024327	Homo sapiens pt-wd mRNA for WD-40 repeat protein, complete cds	0
4060.seq	568	gi 1944628 gb J01415.1 HUMMTCG	Human mitochondrion, complete genome	0
4061.seq	535	gi 452059 gb L27580 HUMIGFBPX	Human insulin-like growth factor binding protein 5 (IGFBP5) mRNA.	E-163
4062.seq	533	gi 3492893 gb AC005288 AC005288	Homo sapiens chromosome 17, clone hCIT.131_K_11, complete sequence [Hom	2.00E-25
4063.seq	491	gi 3341671 gb AC003672 ATAC003672	Arabidopsis thaliana chromosome II BAC F16622 genomic sequence.	0.13
4064.seq	592	gi 1478196 emb X97321 HSHLACW07	H.sapiens mRNA for human leukocyte antigen C alpha chain	0
4065.seq	571	gi 3483627 gb AF086282 HUMZD46B07	Homo sapiens full length insert cDNA clone ZD46B07	0
4066.seq	515	gi 5836157 gb AC006146.2 AC006146	Homo sapiens clone DJ0693M11, complete sequence	0
4067.seq	690	gi 4138921 gb AF060568.1 AF060568	Homo sapiens promyelocytic leukemia zinc finger protein (PLZF) gene.	0.75

4068.seq	656	gi 2662100 cbj AB007870 AB007870	Homo sapiens KIAA0410 mRNA, complete cds	0
4069.seq	424	gi 4503304	Homo sapiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	0
4070.seq	644	gi 3882272 dbj AB018319.1 AB018319	Homo sapiens mRNA for KIAA0776 protein, partial cds	0
4071.seq	578	gi 3378627 emb AJ006996.1 HSG11150	Homo sapiens chromosome 21 PAC LLNLP704G1150Q13, complete sequence	2.5
4072.seq	580	gi 31103 emb Z11531 HSEF1GMR	H.sapiens mRNA for elongation factor-1-gamma	0
4073.seq	268	gi 3108042 gb AC004222 AC004222	Homo sapiens chromosome 17, clone HCIT499120, complete sequence [Homo	4.3
4074.seq	511	gi 5419821 emb AL050337.6 HSJ503F13	Human DNA sequence from clone 503F13 on chromosome 6q24.1-25.2.	E-168
4075.seq	567	gi 984955 gb U14750 HSU14750	Human connective tissue growth factor mRNA, partial cds.	0
4076.seq	752	gi 2326514 emb J285999 HS94G16	Human DNA sequence from PAC 94G16 on chromosome 6q21. Contains	0
4077.seq	244	gi 1262342 emb X93334 HSMITG	H.sapiens mitochondrial DNA, complete genome	E-106
4078.seq	647	gi 4128022 emb AJ011737 HSA011737	Homo sapiens mRNA for integrase interactor 1a protein (INI1A)	0
4079.seq	695	gi 5419576 emb AJ132917.1 HSA132917	Homo sapiens mRNA for methyl-CpG-binding protein 2	0
4080.seq	783	gi 4828656	Homo sapiens caldesmon 1 (CALD1) mRNA >gi 306508 gb M83216 HUMCDSM	E-141
4081.seq	696	gi 5725480 emb AL109793.1 IROO51524	Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 51524	0
4082.seq	510	gi 4572584 emb AL032821.2 HS55C23	Human DNA sequence from clone 55C23 on chromosome 6q22.3-23.3 contain	0.14
4083.seq	214	gi 3366562 gb AC005324 AC005324	Homo sapiens chromosome 17, clone hRPK.640.1.15, complete sequence	0.87
4084.seq	551	gi 3387924 gb AF070558 AF070558	Homo sapiens clone 24450 RING zinc finger protein RZF mRNA,	E-139
4085.seq	571	gi 5042474 gb AC007790.1 AC007790	Homo sapiens clone UWGC:djs310all from 7q31, complete sequence	0.62
4086.seq	579	gi 5453739	Homo sapiens myosin, light polypeptide, regulatory, non-sarcomeric	0
4087.seq	381	gi 4929554 gb AF151801.1 AF151801	Homo sapiens CGI-43 protein mRNA, complete cds	0
4088.seq	558	gi 388308 gb L22005 HUMCD34H	Human ubiquitin conjugating enzyme mRNA, partial cds	0.6
4089.seq	565	gi 2224690 dbj AB002373 AB002373	Human mRNA for KIAA0375 gene, complete cds	0.15
4090.seq	380	gi 1619959 gb U72063 HSU72063	Human immunoglobulin kappa chain constant region gene, partial cds	0
4091.seq	599	gi 4757725	Homo sapiens ALL1 fused gene from chromosome 10 (AF10) mRNA	0
4093.seq	645	gi 1143491 emb X87949 HSRNABIP	H.sapiens mRNA for BIP protein	0
4095.seq	617	gi 5579469	Homo sapiens heat shock 70kD protein 1 (HSPA1A) mRNA	0
4096.seq	525	gi 4826855	Homo sapiens NADH dehydrogenase (ubiquinone) Fe-S protein 1 (75kD)	0
4097.seq	485	gi 4827068	Homo sapiens zinc finger protein 262 (ZNF262) mRNA	0
4098.seq	635	gi 4503536	Homo sapiens eukaryotic translation initiation factor 4E binding	0
4099.seq	621	gi 184428 gb J04621 HUMHSPGC	Human heparan sulfate proteoglycan (HSPG) core protein, 3' end.	0
4100.seq	646	gi 2342595 emb Z75746 HSU221F2	Human DNA sequence from cosmid U221F2 on chromosome X. Contains ESTs	5.00E-05
4101.seq	431	gi 1033631 emb Z61253 HS49B10F	H.sapiens CpG island DNA genomic Mse1 fragment, clone 49B10,	1.00E-47
4102.seq	537	gi 179295 gb L08441 HUMAUTONH	Human autonomously replicating sequence (ARS) mRNA.	0
4103.seq	319	gi 3065552 gb L13802 HUMCH13C3A	Homo sapiens (clone 03) liver expressed ribosomal protein S3	E-173
4104.seq	619	gi 1754632 dbj D88010 D88010	Human DNA for ribosomal protein S13, complete cds, U14 small	E-150
4105.seq	679	gi 3091285 gb AF060555 AF060555	Homo sapiens estrogen receptor beta 3 isoform mRNA, complete cds	0.74
4106.seq	510	gi 4557316	Homo sapiens annexin XI (56kD autoantigen) (ANX11) mRNA	0
4107.seq	610	gi 4158181 gb AC006007 AC006007	Homo sapiens PAC clone DJ0819004 from 7q33-q35, complete sequence [Hor	0.66
4108.seq	624	gi 190978 gb M18663 HUMRECA	Human islet of Langerhans regenerating protein (reg) mRNA, complete	0
4109.seq	390	gi 5031972	Homo sapiens protein disulfide isomerase-related protein (P5) mRNA	0
4110.seq	531	gi 3980526 gb AC005483 AC005483	Homo sapiens PAC clone DJ0806A17 from 7p13-p14, complete sequence [Hor	0.037
4111.seq	726	gi 4885072	Homo sapiens activating transcription factor 1 (ATF1) mRNA	0
4112.seq	682	gi 1699367 gb AC000065 HSAC000065	Human BAC clone RG085C06 from 7q21-7q22, complete sequence [Homo	0
4113.seq	617	gi 3869308 gb AF071741 HSTOP2A04	Homo sapiens topoisomerase II alpha (TOP2A) gene, exons 17 through 20	0.67
4114.seq	623	gi 3687210 gb AC005740 AC005740	Homo sapiens chromosome 5p, BAC clone 50621 (LBNL.H154), complete	0
4115.seq	599	gi 3059064 emb Z97181 HS95C20	Homo sapiens DNA sequence from PAC 95C20 on chromosome Xp11.3-11.4.	0.042
4116.seq	685	gi 31348 emb X06292 HSEF5P5	Human c-fes/fps proto-oncogene	0.75
4117.seq	542	gi 1621235 emb Z81009 HS473J10	Human DNA sequence from PAC 473J10, between markers DXS6791 and DX	0.59
4118.seq	562	gi 4092849 dbj AB021866 AB021866	Homo sapiens KIP gene, complete cds	E-158
4119.seq	590	gi 4454525 gb AC006145 AC006145	Homo sapiens PAC clone DJ0560014 from 7q21.1-q21.2, complete sequence	0.041
4120.seq	659	gi 4506196	Homo sapiens proteasome (prosome, macropain) subunit, beta type, 3	0
4122.seq	676	gi 4760262 emb JL049773.1 HS10618A	Human DNA sequence from clone 1061018A on chromosome 22q13.32-13.33	0.047
4123.seq	661	gi 6009489 dbj AB018358.1 AB018358	Homo sapiens mRNA for ABP125, complete cds	0
4124.seq	679	gi 3171157 gb AF064862 AF064862	Homo sapiens chromosome 21q22.3 PAC 31P10, complete sequence [Homo	0.19
4125.seq	493	gi 5725511 gb AF072097.1 AF072097	Homo sapiens beta-2 microglobulin gene, complete cds	0
4126.seq	598	gi 13683 emb V00710 MIT1HS	Human mitochondrial genes for several tRNAs (Phe, Val, Leu) and 12S	1.00E-23
4127.seq	653	gi 4633508 gb AF119297.1 AF119297	Homo sapiens neuroendocrine-specific protein-like protein 1 (NSPL1)	0
4128.seq	137	gi 3327440 gb AF014662.1 AF014662	Human T-cell lymphotropic virus type 1 isolate BRASP23, 5' long	0.52
4130.seq	595	gi 4454488 gb AC004796 AC004796	Homo sapiens chromosome 11p14.3 PAC clone pDJ1034g4, complete sequenc	4.00E-08
4131.seq	606	gi 3242759 gb AC005156 AC005156	Homo sapiens PAC clone DJ1099C19 from 7q21-q22, complete sequence [Hor	0
4132.seq	614	gi 4503270	Homo sapiens decorin (DCN) mRNA >gi 181169 gb M14219 HUMCSPG1A	0
4133.seq	653	gi 31091 emb X18689 HSEF1AC	Human mRNA for elongation factor 1-alpha (clone CEF4)	0
4134.seq	475	gi 4507668	Homo sapiens tumor protein, translationally-controlled 1 (TPT1)	0
4135.seq	593	gi 4164360 emb JL031782.1 HS708F5	Human DNA sequence from clone 708F5 on chromosome 6p11.2-12.3. Contain	0.01
4136.seq	724	gi 394330 emb Z24130 HS294VD5	H.sapiens (D85544) DNA segment containing (CA) repeat; clone	2.00E-63
4137.seq	524	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	0
4139.seq	617	gi 4504168	Homo sapiens glutathione synthetase (GSS) mRNA	0
4140.seq	579	gi 4503116	Homo sapiens cystatin B (stefin B) (CSTB) mRNA	0
4141.seq	644	gi 950211 gb U31345 HSU31345	Human calpastatin mRNA, partial cds, short 3'UTR.	0
4143.seq	647	gi 5101771 emb J242978.1 HSA242978	Homo sapiens partial mRNA for putative p621 protein which interacts	0
4144.seq	651	gi 5668655 emb AL050331.1 HSDJ4861	Human DNA sequence from clone 48613 on chromosome 6q22.1-22.3. Contain	0
4145.seq	652	gi 4758157	Homo sapiens differentiation 6 (deoxyguanosine triphosphate	0
4146.seq	491	gi 4505574	Homo sapiens poly(A)-binding protein-like 1 (PABP1) mRNA	E-113
4147.seq	562	gi 4504164	Homo sapiens gelsolin (amyloidosis, Finnish type) (GSN) mRNA	0
4148.seq	676	gi 4759237	Homo sapiens transmembrane 4 superfamily member 3 (TM4SF3) mRNA	0
4149.seq	195	gi 4504618	Homo sapiens insulin-like growth factor binding protein 7 (IGFBP7)	1.00E-85
4150.seq	636	gi 1901671 gb M73547 HUMPOLLA	Human polyposis locus (DP1) gene) mRNA, complete cds.	0
4151.seq	301	gi 3413951 dbj AB007976 AB007976	Homo sapiens mRNA, chromosome 1 specific transcript KIAA0507	E-144
4152.seq	539	gi 5649377 gb AC007386.3 AC007386	Homo sapiens BAC clone NH0359K10 from 2, complete sequence [Homo sapiens]	2.00E-68
4153.seq	566	gi 3253130 gb AC004167 AC004167	Homo sapiens chromosome 17, clone HRPC109F1, complete sequence [Homo	0.16
4154.seq	608	gi 414346 dbj D14531 HUMHRRPL9	Human mRNA for human homologue of rat ribosomal protein L9.	0
4155.seq	608	gi 4758949	Homo sapiens peptidylprolyl isomerase B (cyclophilin B) (PPIB) mRNA	0
4156.seq	653			0
4157.seq	813	gi 4240232 dbj AB020679.1 AB020679	Homo sapiens mRNA for KIAA0872 protein, complete cds	0
4158.seq	345	gi 4730839 dbj AP000105.1 AP000105	Homo sapiens genomic DNA of 21q22.1, GART and AML related,	E-155
4159.seq	596	gi 3043679 dbj AB011150 AB011150	Homo sapiens mRNA for KIAA0578 protein, partial cds	0
4160.seq	623	gi 4191326 gb AF065391 AF065391	Homo sapiens ZIS1 mRNA, complete cds	0
4161.seq	634	gi 184847 gb M63438 HUMIGGK	Human Ig rearranged gamma chain mRNA, V-J-C region and complete cds.	0
4163.seq	663	gi 2688637 gb AC003075 AC003075	Human PAC clone DJ0658N05 from 7p21, complete sequence [Homo sapiens]	0
4164.seq	506	gi 2564670 gb AF023462 AF023462	Homo sapiens peroxisomal phytanoyl-CoA alpha-hydroxylase (PAHX) mRNA,	0
4165.seq	518	gi 5596978 emb AL049735.4 HSJ735P11	Human DNA sequence from clone 735P11 on chromosome Xp11.3-11.4.	0.56
4167.seq	346	gi 4325309 gb AF124523.1 AF124523	Homo sapiens chromosome 8 clone Kox18-21 containing zinc-finger protein	0.093
4168.seq	434	gi 3043661 dbj AB011141 AB011141	Homo sapiens mRNA for KIAA0569 protein, complete cds	E-115
4169.seq	566	gi 576474 gb M28203 HUMHMBW62	Homo sapiens (clone pMF18) MHC class I HLA-B*62 mRNA, 3' end.	0
4170.seq	736	gi 5453703	Homo sapiens vitamin A responsive; cytoskeleton related (JWA) mRNA	0
4173.seq	220	gi 4758647	Homo sapiens kinesin family member 5B (KIF5B) mRNA	1.00E-80
4174.seq	408	gi 4500154 emb ALD49365.1 HSM80015	Homo sapiens mRNA; cDNA DKFZp586A0618 (from clone DKFZp586A0618)	0
4175.seq	600	gi 598954 dbj D28915 HUMHCAMAP8	Human gene for hepatitis C-associated microtubular aggregate	E-160
4176.seq	507	gi 6005849	Homo sapiens protein kinase Chk2 (RAD53), mRNA	0
4177.seq	330	gi 2098573 gb AC002115 HSAC002115	Human DNA from overlapping chromosome 19 cosmids R31396, F25451, and	0.35
4178.seq	396	gi 5499744 gb AF165138.1 AF165138	Homo sapiens PAC B39112 derived from chromosome 21p11.2, complete	E-109
4179.seq	474	gi 388034 gb L05095 HUMRIBPROF	Homo sapiens ribosomal protein L30 mRNA, complete cds	0
4180.seq	316	gi 758797 gb U23765 HSU23765	Human Bak protein mRNA, complete cds.	E-176
4181.seq	506	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	0
4182.seq	629	gi 2580474 gb AC002558 AC002558	Homo sapiens chromosome 17, clone hRPC867C24, complete sequence [Homo	0
4183.seq	398			0
4184.seq	411	gi 5815507 gb AC006509.15 AC006509	Homo sapiens 2p16 PAC RPC15-960D23 (Roswell Park Cancer Institute Huma	9.00E-98
4185.seq	638	gi 4758293	Homo sapiens glutaryl-prolyl-tRNA synthetase (EPRS) mRNA	0

4187.seq	617	g 4210505 gb AC006305 AC006305	Homo sapiens chromosome 18, clone hRPK.456.O.19, complete sequence	0.17
4188.seq	282	g 506337 db D30648 HUMMFP5	Human mRNA for flavoprotein subunit of complex II, complete cds	E-156
4189.seq	602	g 2992475 gb AC002549 AC002549	Homo sapiens Xp22 BAC GS-377014 (Genome Systems Human BAC library)	0.042
4190.seq	361	g 5306308 gb AC004878.2 AC004878	Homo sapiens clone DJ0754G14, complete sequence	3.00E-17
4191.seq	688	g 2623584 gb AF027158 AF027158	Homo sapiens immunoglobulin kappa light chain (T6J/k) mRNA, partial	0
4192.seq	679	g 337188 gb M2517 HUMMTCDK	Human mitochondrion cytochrome oxidase subunit II (COII), Asp-tRNA	0
4193.seq	651	g 3954884 emb JA010442 HSA010442	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	0
4194.seq	611	g 3043571 db AB011096 AB011096	Homo sapiens mRNA for KIAA0524 protein, partial cds	0
4195.seq	567	g 4506208	Homo sapiens proteasome (prosome, macropain) 26S subunit, ATPase, 2	0
4196.seq	623	g 4557352	Homo sapiens branched chain keto acid dehydrogenase E1, beta	9.00E-56
4197.seq	242	g 1469885 db D63486 D63486	Human mRNA for KIAA0152 gene, complete cds	E-104
4198.seq	685	g 6016975 emb AL109609.5 HJS556A22	Human DNA sequence from clone 556A22 on chromosome Xp21.1-21.3,	2.00E-04
4199.seq	711	g 4506944	Homo sapiens sucrose-isomaltase (SI) mRNA	0
4200.seq	487	g 4502046	Homo sapiens aldehyde dehydrogenase 9 (zamma-aminobutyraldehyde	2.1
4201.seq	658	g 4504504	Homo sapiens hydroxysteroid (17-beta) dehydrogenase 4 (HSD17B4)	0
4202.seq	605	g 4758301	Homo sapiens enhancer of rudimentary (Drosophila) homolog (ERH)	0
4204.seq	675	g 506335 db D14436.1 HUMHHIRE	Human gene for histamine H1-receptor, complete cds	0
4205.seq	686	g 1110519 gb U40490 HSU40490	Human nicotinamide nucleotide transhydrogenase mRNA, nuclear gene	0
4206.seq	683	g 5802965	Homo sapiens destrin (actin depolymerizing factor) (ADF), mRNA	0
4207.seq	553	g 4505742	Homo sapiens prefoldin 5 (PFDN5) mRNA >g 1731808 db D89667 D89667	0
4208.seq	674	g 3041856 gb AC004536 AC004536	Homo sapiens PAC clone DJ0852D15 from 7q31, complete sequence [Homo	0.74
4209.seq	701	g 4713939 gb AC008518.1 AC008518	Homo sapiens 12p13 BAC RPC11-144023 (Roswell Park Cancer Institute Hur	0
4210.seq	638	g 3005738 gb AF05014 AF05014	Homo sapiens clone 24521 mRNA sequence	0
4211.seq	567	g 4758289	Homo sapiens EphB4 (EPHB4) mRNA >g 495472 gb U07695 HSU07695 Huma	0
4212.seq	485	g 5016885 db AP000280.2 AP000280	Homo sapiens genomic DNA, chromosome 21q22.1, clone: S12, D21S226-AML	2.1
4213.seq	624	g 2315183 emb Z82216 HS75N13	Human DNA sequence from PAC 75N13 on chromosome Xq21.1. Contains ZNF	0
4214.seq	805	g 4507668	Homo sapiens tumor protein, translationally-controlled 1 (TPT1)	0
4215.seq	585	g 4406571 gb AF131753 AF131753	Homo sapiens clone 24859 mRNA sequence	0
4216.seq	633	g 6005881	Homo sapiens serine protease, umbilical endothelium (SPUVE), mRNA	0
4218.seq	672	g 414346 db D14531 HUMHRRPL9	Human mRNA for human homologue of rat ribosomal protein L9,	0
4220.seq	681	g 1944628 gb J01415.1 HUMMTCG	Human mitochondrion, complete genome	0
4221.seq	583	g 4506602	Homo sapiens ribosomal protein L15 (RPL15) mRNA	0
4222.seq	204			0
4223.seq	588	g 339688 gb M17733 HUMTHYB4	Human thymosin beta-4 mRNA, complete cds.	0
4224.seq	186	g 187625 gb M84378 HUMMHAD	Human MHC class I lymphocyte antigen (HLA-A*0212) mRNA, complete	2.00E-59
4226.seq	629	g 2565322 gb AF024710.1 AF024710	Homo sapiens vascular endothelial growth factor (VEGF) mRNA, 3'UTR	0
4227.seq	512	g 189600 gb M84337 HUMPANSECP	Human pancreatitis associated protein mRNA, complete cds.	0
4228.seq	348	g 4506644	Homo sapiens ribosomal protein L38 (RPL38) mRNA	0
4229.seq	629	g 5649179 gb AC007860.6 AC007860	Homo sapiens 12p13 BAC RPC11-709E21 (Roswell Park Cancer Institute	0.044
4230.seq	576	g 1488412 gb S82081 S82081	N8-tumor expression-enhanced gene [human, NCI H-89 cell line, mRNA,	0
4232.seq	213			0
4233.seq	546	g 4761653 gb AF147302.1 AF147302	Homo sapiens full length insert cDNA clone YA77A08	0
4234.seq	580	g 31916 emb X56009 HSGSAG	Human GSA mRNA for alpha subunit of GsGTP binding protein	0
4235.seq	609	g 4508126 gb AC006376.2 AC006376	Homo sapiens clone NH0386L03, complete sequence	1.00E-20
4236.seq	713	g 189340 gb M29927 HUMOAT11	Human ornithine aminotransferase gene, exon 11.	0
4237.seq	313	g 4543853	Homo sapiens poly(rC)-binding protein 1 (PCBP1) mRNA	5.00E-77
4239.seq	261	g 337186 gb M2517 HUMMTCDK	Human mitochondrion cytochrome oxidase subunit II (COII), Asp-tRNA	1.00E-83
4240.seq	597	g 30228 emb Y00057 HSCSABR	Human mRNA for beta-subunit of cell surface adhesion glycoproteins	0
4242.seq	624	g 4456531 emb AJ239355 HSA239355	Homo sapiens mRNA for immunoglobulin heavy chain variable region,	2.7
4243.seq	720	g 5830435 emb Y14485.1 HSY14485	Homo sapiens mRNA for cytosolic serine hydroxymethyltransferase,	0.013
4244.seq	576	g 5453889	Homo sapiens PIBF1 gene product (PIBF1) mRNA	3.00E-80
4246.seq	715	g 2342715 gb AC002287 HUAC002287	Homo sapiens Chromosome 16 BAC clone CIT987-SKA-237H1 ~complete	0.2
4247.seq	737	g 492958 gb AF151853.1 AF151853	Homo sapiens CGI-95 protein mRNA, complete cds	0
4248.seq	589	g 5103012 db AP000349.1 AP000349	Homo sapiens genomic DNA, chromosome 22q11.2, clone KB1839H6	3.00E-77
4250.seq	645	g 4500092 emb AL049319.1 HSM800104	Homo sapiens mRNA; cDNA DKFZp564C046 (from clone DKFZp564C046)	0
4251.seq	407	g 4580685 gb AF105999.1 AF105999	Homo sapiens acetylcholine receptor epsilon subunit (CHRNE) gene,	0.44
4252.seq	386	g 2244605 db AB001523 AB001523	Homo sapiens gene for TMEM1 and PWP2, complete and partial cds	E-121
4253.seq	240	g 6102633 gb AC011422.2 AC011422	Homo sapiens chromosome 5 clone CITB-H1_2335E17, complete sequence	2.00E-36
4254.seq	297	g 5262352 emb AL035398.19 HS796117	Human DNA sequence from clone 796117 on chromosome 22q13.2. Contains	1.2
4255.seq	681	g 1808647 emb Y10807 HSY10807	H.sapiens mRNA for arginine methyltransferase, splice variant, 1262	0
4256.seq	635	g 4240244 db AB020685.1 AB020685	Homo sapiens mRNA for KIAA0878 protein, complete cds	0
4257.seq	695	g 5830347 emb AL031667.17 HS620E11	Human DNA sequence from clone 620E11 on chromosome 20q11.2-12. Cortical	0.19
4258.seq	787	g 1060888 db D78152 HUMALV	Human mRNA for annexin IV (carbohydrate-binding protein p33/41),	0
4260.seq	702	g 4506722	Homo sapiens ribosomal protein S3A (RPS3A) mRNA	0
4261.seq	572	g 4406690 gb AF131848 AF131848	Homo sapiens clone 24922 mRNA sequence, complete cds	0
4262.seq	580	g 31912 emb X04409 HSGSA1R	Human mRNA for coupling protein G(s) alpha-subunit (alpha-S1)	0
4263.seq	521	g 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exons 3 and 4.	0
4264.seq	702	g 2394173 gb AF016052 AF016052	Homo sapiens zinc finger protein ZNF191 (ZNF191) gene, complete cds	1.3
4265.seq	201	g 4775617 emb AL035410.7 HS591B8	Human DNA sequence from clone 591B8 on chromosome 1p13.1, complete	1.00E-76
4266.seq	519	g 2398553 gb AC002529 AC002529	Human BAC clone RG161N06 from 7q31, complete sequence [Homo sapiens]	0.036
4267.seq	718	g 4504618	Homo sapiens insulin-like growth factor binding protein 7 (IGFBP7)	0
4268.seq	728	g 3080378 emb AL022575 HS159A15	Homo sapiens DNA sequence from PAC 159A15 on chromosome	0.81
4269.seq	188	g 4454519 gb AC006456 AC006456	Homo sapiens PAC clone DJ0968D04 from 7q33-q35, complete sequence [Hor	0.012
4270.seq	751	g 4835817 gb AC007077.2 AC007077	Homo sapiens BAC clone NH0260K08 from 2, complete sequence	0.053
4271.seq	478	g 1164913 emb Z68873 HSU209G1	Human DNA sequence from cosmid U209G1 on chromosome X	2.00E-22
4272.seq	564	g 3327107 db AB014547 AB014547	Homo sapiens mRNA for KIAA0647 protein, partial cds	0
4273.seq	591	g 4506518	Homo sapiens regulator of G-protein signaling 5 (RGS5) mRNA	0
4274.seq	422	g 517114 db D25215 HUMORFKA	Human mRNA for KIAA0032 gene, complete cds	E-107
4275.seq	423	g 4503304	Homo sapiens defensin, alpha 5, Paneth cell-specific (DEFA5) mRNA	0
4276.seq	310	g 2307001 gb AF012024 AF012024	Homo sapiens integrin cytoplasmic domain associated protein	1.00E-80
4277.seq	540	g 3645947 gb AC002543 AC002543	Homo sapiens BAC clone RG300C03 from 7q31.2, complete sequence	0.038
4278.seq	621	g 4505782	Homo sapiens phosphorylase kinase, beta (PHKB) mRNA	0
4279.seq	416	g 409069 db D17652 HUMHBL2A	Human mRNA for HbP15/L22, complete cds	0
4280.seq	706	g 4885408	Homo sapiens high density lipoprotein binding protein (HDLBP) mRNA	0
4281.seq	472	g 2813964 emb Z97053 HS179M20	Human DNA sequence from PAC 179M20 on chromosome 20q12-13.1. Cortical	0
4282.seq	684	g 4506742	Homo sapiens ribosomal protein S8 (RPS8) mRNA	0
4283.seq	654	g 2145059 gb AF000560 AF000560	Homo sapiens TTF-1 interacting peptide 20 mRNA, partial cds	E-158
4284.seq	622	g 6318543 gb AF126780.1 AF126780	Homo sapiens retinal short-chain dehydrogenase/reductase retSDR2	0
4285.seq	650	g 4506648	Homo sapiens ribosomal protein L3 (RPL3) mRNA	0
4286.seq	660	g 3954892 emb JA010446 HSA010446	Homo sapiens mRNA for immunoglobulin kappa light chain, anti-RhD,	0
4287.seq	591	g 3540146 gb AC002395 AC002395	Homo sapiens 12q24.1 PAC RPC11-315L5 (Roswell Park Cancer Institute	E-109
4288.seq	380	g 4506624	Homo sapiens ribosomal protein L27a (RPL27A) mRNA	0
4289.seq	676	g 5729963	Homo sapiens inactive progesterone receptor, 23 kD (P23) mRNA	0
4290.seq	620	g 5819124 gb AC007286.18 AC007286	Homo sapiens 12q15 BAC RPC11-298M11 (Roswell Park Cancer Institute	0.17
4291.seq	565	g 2765426 emb Y14738 HISIG1LL	Homo sapiens mRNA for immunoglobulin lambda light chain	0
4292.seq	382	g 758075 emb X83545 HSIDD	H.sapiens mRNA for a cell surface protein	E-146
4293.seq	599	g 4503380	Homo sapiens down-regulator of transcription 1, TBP-binding (negative	0
4294.seq	670	g 1039380 gb U16028 HSU16028	Human CRE-BP1 transcription factor mRNA, complete cds	0
4295.seq	418	g 4506708	Homo sapiens ribosomal protein S26 (RPS26) mRNA	0
4296.seq	626			0
4297.seq	707	g 310911 emb X16869 HSEF1AC	Human mRNA for elongation factor 1-alpha (clone CEF4)	0
4298.seq	361	g 3419846 gb AC004982 AC004982	Homo sapiens PAC clone DJ1159C04 from 7p21-p22, complete sequence [Hor	0.39
4299.seq	641	g 4758027	Homo sapiens collagen, type VI, alpha 3 (COL6A3) mRNA	0
4300.seq	537	g 34469 emb X66867 HSMAXG	H.sapiens max gene	0
4301.seq	631	g 5016088	Homo sapiens actin, beta (ACTB) mRNA	0
4302.seq	707	g 190979 gb M18963 HUMREGA	Human Islet of Langerhans regenerating protein (reg) mRNA, complete	0
4303.seq	683	g 4506292	Homo sapiens protein tyrosine phosphatase, non-receptor type 3	0
4305.seq	602	g 1561514 db D38526 D38526	Human mRNA for HLA-Cw*0702, complete cds	0

4306.seq	535	gi 184847 gb M63438 HUMIGGK	rearranged gamma chain mRNA, V-J-C region and complete cds.	0
4307.seq	578	gi 5629939 gb AC007368.1 AC007368	siens 12q24.2 PAC RPC14-809F18 (Roswell Park Cancer institute	0.01
4308.seq	436	gi 264772 gb S54005 S54005	beta-10 [human, metastatic melanoma cell line, mRNA, 453	0
4309.seq	512	gi 5763746 emb AL049766.14 HSDJ686	INA sequence from clone 686N3 on chromosome 20q13.2-13.2.	E-114
4310.seq	727	gi 4503470	siens eukaryotic translation elongation factor 1 alpha 1	0
4311.seq	753	gi 2133864 gb AC000403 AC000403	sequence from Human 13, complete sequence [Homo sapiens]	0
4312.seq	636	gi 4501886	siens actin, gamma 1 (ACTG1) mRNA	0
4313.seq	576	gi 4929578 gb AF151813.1 AF151813	siens CGI-55 protein mRNA, complete cds	0
4314.seq	681	gi 5262614 emb AL080156.1 HSM8006	siens mRNA: cDNA DKFZp434J214 (from clone DKFZp434J214)	0
4315.seq	587	gi 4337095 gb AF129756.1 D2J01G24	siens MSH5 gene, partial cds; and CLIC1, DDAH, G6b, G6c, G5b,	0
4316.seq	560	gi 5821139 dbj AB015639.1 AB015639	siens ASY mRNA, complete cds	1.00E-63
4317.seq	641	gi 4507358	siens transferrin (TAGLN) mRNA >gi 177174 gb M95787 HUM22SM	0
4318.seq	727	gi 1216453 gb U48488 HUMRYR40	keletal muscle ryanodine receptor gene (RYR1), exons 74, 75,	0.013
4319.seq	674	gi 5031608	siens branched chain alpha-ketoacid dehydrogenase kinase	0
4320.seq	380	gi 306367 gb M95786 HAVGLYGBB	lpha-herpesvirus (isolated from Saimiri sciureus)	0.41
4322.seq	585	gi 532596 gb M12759 HUMIGJ02	g J chain gene, exons 3 and 4.	E-174
4323.seq	602	gi 572551 gb AF072097.1 AF072097	siens beta-2 microglobulin gene, complete cds	0
4324.seq	680	gi 5531804 gb AF078845.1 AF078845	siens 16.7Kd protein mRNA, complete cds	0
4327.seq	609	gi 4506730	siens ribosomal protein S6 (RPS6) mRNA	0
4328.seq	584	gi 4850411 gb AC007288.1 ATAC007	sis thalassa chromosome II BAC F10C8 genomic sequence,	0.041
4329.seq	619	gi 182976 gb M33197 HUMGAPDH	lyceraldehyde-3-phosphate dehydrogenase (GAPDH) mRNA,	0
4331.seq	728	gi 1684744 emb Y07783 RNDR142	icus mRNA for ER transmembrane protein	1.00E-05
4332.seq	518			
4333.seq	603	gi 4506090	siens mitogen-activated protein kinase 6 (MAPK6), mRNA	0
4334.seq	575	gi 182864 gb J02939 HUMG4F2	membrane glycoprotein 4F2 antigen heavy chain mRNA,	0
4335.seq	682	gi 4758415	siens golgi-specific brefeldin A-resistance factor 1 (GBF1)	2.00E-50
4336.seq	595	gi 806707 gb U12518 SMU12518	sis macroura transthyretin mRNA, complete cds.	0.17
4337.seq	412	gi 4503304	siens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	0
4338.seq	352	gi 3133268 gb U20352 HSU20352	siens malate dehydrogenase (MDHA) mRNA, complete cds	0
4339.seq	692	gi 3980430 gb AC003061 AC003061	hromosome 6 BAC Clone b245c12, complete sequence [Mus musculus	0.76
4340.seq	515	gi 185538 gb M18512 HUMIGHVAJ	fetal) rearranged IgM chain VDJ-region mRNA, clone 20P3.	0
4341.seq	661	gi 1809237 gb AC000111 HSAC000111	IAC clone 068P20 from 7q31-q32, complete sequence [Homo sapiens]	E-107
4342.seq	490	gi 1737211 gb U80017 HSU80017	siens basic transcription factor 2 p44 (btf2p44) gene, partial	0.002
4343.seq	334	gi 292440 gb L11567 HUMRPL37Z	siens ribosomal protein L37 mRNA, complete cds.	0
4344.seq	242	gi 532596 gb M12759 HUMIGJ02	g J chain gene, exons 3 and 4.	E-119
4345.seq	696	gi 468242 gb U07989 HSU07989	lurkitt's lymphoma immunoglobulin kappa light chain mRNA,	0
4347.seq	382	gi 4759321	siens wingless-type MMTV integration site family, member 2B	0
4349.seq	225	gi 2745720 gb U89321 HSMNPP25	siens nucleophosmin phosphoprotein (NPM) gene, exon 12,	1.00E-86
4350.seq	383	gi 3478668 gb AC005550 AC005550	siens PAC clone DJ0620P06 from 7p21-p22, complete sequence [Homo	E-179
4351.seq	633	gi 475207 emb X78989 MMTEST	siens mRNA for testin	E-112
4352.seq	605	gi 4506730	siens ribosomal protein S6 (RPS6) mRNA	0
4353.seq	372	gi 4501932	siens alcohol dehydrogenase 3 (class I), gamma polypeptide	0
4354.seq	446	gi 3399669 gb AC005392 AC005392	siens chromosome 19, CIT-HSP BAC 49q23 (BC338531), complete	0.031
4355.seq	513	gi 176828 gb M30685 CHPMHC8AD	siens MHC class I protein mRNA (MHCATRF1).	0
4358.seq	620	gi 3688107 gb AC004477 AC004477	siens chromosome 17, clone HRPC890E16, complete sequence [Homo	5.00E-14
4359.seq	542	gi 2894828 gb U91942 U91942	siens immunoglobulin light chain mRNA, partial cds	0
4360.seq	775	gi 1633547 gb U47924 HSU47924	hromosome 12p13 sequence, complete sequence [Homo sapiens]	0.004
4361.seq	274	gi 4507668	siens tumor protein, translationally-controlled 1 (TPT1)	E-137
4362.seq	426	gi 181532 gb M97925 HUMDEF5A	s defensin 5 gene, complete cds.	E-100
4363.seq	622	gi 3928250 emb Z68955 HSLAM004	s mRNA for immunoglobulin kappa light chain VJ region (ID	E-162
4364.seq	605	gi 6164703 gb AF167572.1 AF167572	siens protein methyltransferase (JBP1) mRNA, complete cds	0
4365.seq	650	gi 4753253 gb AC006364.3 AC006364	siens BAC clone GS146J04 from 7q31.1-31.3, complete sequence	0.046
4366.seq	561	gi 5834019 gb AF174030.1 AF174030	siens clone 77u-c17 immunoglobulin heavy chain variable	E-156
4367.seq	580	gi 261239 gb S50732 S50732	globulin M light chain V region-anti-lipid A antibody [human,	E-147
4368.seq	622	gi 2832583 emb Z97056 HS434P1	ANA sequence from PAC 434P1 on chromosome 22. Contains inward	0
4370.seq	617	gi 4902597 emb AL034396.8 HS11588	ANA sequence from clone 1158812 on chromosome Xp11.21-11.4	0
4371.seq	441	gi 3366586 gb AC005279 AC005279	siens chromosome 17, clone hRPK.103_M_22, complete sequence [Ho	1.9
4372.seq	704	gi 28335 emb X63432 HSACTB	s ACTB mRNA for mutant beta-actin (beta-actin)	0
4374.seq	253	gi 5454099	siens transforming, acidic coiled-coil containing protein 1	E-120
4375.seq	734	gi 5410325 gb AF106680.1 AF106680	siens RNA helicase mRNA, complete cds	0
4376.seq	698	gi 4239993 dbj AB011785 AB011786	um perfringens genes of hem operon, complete cds	0.049
4377.seq	630	gi 5912178 emb AL117608.1 HSM8011	siens mRNA: cDNA DKFZp564O1863 (from clone DKFZp564O1863)	0
4378.seq	549	gi 6006276 dbj AP00469.2 AP00469	siens genomic DNA, chromosome 21q21.2, clone:8132H24, LL56-APP	0.15
4379.seq	648	gi 1800114 gb U80579 HSUB0579	siens ribosomal protein L7 antisense mRNA gene, partial sequence	0
4380.seq	396	gi 36135 emb Z12962 HSRPL41	s mRNA for homologue to yeast ribosomal protein L41	E-177
4381.seq	652	gi 3882280 dbj AB018323.1 AB018323	siens mRNA for KIAA0780 protein, partial cds	0
4383.seq	623	gi 4505362	siens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 5	E-148
4384.seq	612	gi 3820992 emb AL008728 HS125N5	ANA sequence from clone 125N5 on chromosome 6q26-27. Contains a	0.043
4386.seq	553	gi 4505922	siens plin1, desmosome associated protein (PNN) mRNA	0
4387.seq	727	gi 4506204	siens proteasome (prosome, macropain) subunit, beta type, 9	0
4388.seq	232	gi 1204118 emb Z69720 HSRA36	ANA sequence from cosmid RA36 from a contig from the tip of the	E-128
4389.seq	791	gi 4759095	siens 15 kDa selenoprotein (SEP15) mRNA	0
4390.seq	557	gi 5454161	siens upregulated by 1,25-dihydroxyvitamin D-3 (VDUP1) mRNA	0
4391.seq	564	gi 2313072 gb U90095 U90095	Chromosome 11p15.5 pac pDJ74k15, complete sequence [Homo sapie	0.62
4392.seq	641	gi 3138891 gb AC004700 AC004700	siens chromosome 19, cosmid F14498, complete sequence [Homo	0.71
4393.seq	435	gi 2981253 gb AC003964 HUAAC00396	siens Chromosome 16 BAC clone CIT987SK-A-91E12, complete	0.12
4394.seq	639	gi 5262538 emb AL080111.1 HSM8006	siens mRNA: cDNA DKFZp586G2222 (from clone DKFZp586G2222)	0
4395.seq	525	gi 1665822 dbj D87470 D87470	mRNA for KIAA0280 gene, partial cds	0
4396.seq	632	gi 4502100	siens annexin I (lipocortin I) (ANX1) mRNA	0
4397.seq	474	gi 6119551 gb AC000096.13 AC00009	sculus Chromosome 16 BAC Clone bcd3-6 Syntenic To Homo sapiens	0.033
4398.seq	258	gi 5541861 emb AL034548.24 HS1103	DNA sequence from clone 1103G7 on chromosome 20p12.2-13, compl	5.00E-18
4399.seq	746	gi 4884467 emb AL050225.1 HSM8006	siens mRNA: cDNA DKFZp586M1523 (from clone DKFZp586M1523)	0
4400.seq	704	Homo sapiens ribosomal protein L14 (I		0
4401.seq	510	gi 3169015 emb AJ006268 HSAJ5268	siens mRNA for putative ATPase, partial	1.00E-91
4402.seq	383	gi 28335 emb X63432 HSACTB	is ACTB mRNA for mutant beta-actin (beta-actin)	0
4404.seq	658	gi 558437 gb U13897 HSU13897	homolog of Drosophila discs large protein, isoform 1 (hdlg-1)	3.00E-86
4405.seq	607	gi 4512311 dbj AB019441.1 AB019441	siens DNA for immunoglobulin heavy-chain variable region,	0
4406.seq	471	gi 3283909 gb AF070638 AF070638	siens clone 24448 unknown mRNA, partial cds	0
4407.seq	400	gi 181532 gb M97925 HUMDEF5A	is defensin 5 gene, complete cds.	8.00E-86
4409.seq	700	gi 5650596 emb AL031734.9 HS652L8	DNA sequence from clone 652L8 on chromosome 1q24.1-25.2, comple	0.77
4410.seq	723	gi 181532 gb M97925 HUMDEF5A	is defensin 5 gene, complete cds.	4.00E-98
4411.seq	648	gi 746405 gb U02292 BTU02292	rus link protein mRNA, complete cds	6.00E-17
4412.seq	198	gi 644480 dbj D38112.1 HUMMTA	siens mitochondrial DNA, complete sequence	E-103
4413.seq	723	gi 6137796	mitochondrion, complete genome	0
4414.seq	456	gi 4504164	siens gelsolin (amyloidosis, Finnish type) (GSN) mRNA	0
4415.seq	628	gi 4503470	siens eukaryotic translation elongation factor 1 alpha 1	0
4416.seq	642	gi 183921 gb M63544 HUMHDABCD	DNA sequence of contig comprising 3 cosmids (HDAB, HDAC, HDAD).	0.045
4417.seq	585	gi 265703 gb S55735 S55735	.1-alpha 2m-immunoglobulin A1-A2 lambda hybrid GAU heavy	0
4420.seq	557	gi 4759223	siens programmed cell death 5 (TFAR19), mRNA	0
4422.seq	677	gi 4506660	siens ribosomal protein L7a (RPL7A) mRNA	0
4423.seq	755	gi 5803156	siens sterol-C4-methyl oxidase-like (SC4MOL), mRNA	3.00E-68
4424.seq	584	gi 4506616	siens ribosomal protein L17 (RPL17) mRNA	0
4425.seq	657	gi 2842782 gb AC003103 AC003103	siens chromosome 17, clone HCT1268N12, complete sequence [Homo	0.003
4426.seq	619	gi 603763 gb U18321 HSU18321	ionizing radiation resistance conferring protein mRNA, complete	0
4427.seq	634	gi 4504786	siens Integral transmembrane protein 1 (ITM1) mRNA	0
4428.seq	405	gi 4753275 gb AC006148.3 AC006148	siens PAC clone DJ0902E20 from 7q31.2-q31.3, complete sequence	E-173
4429.seq	555	gi 265703 gb S55735 S55735	.1-alpha 2m-immunoglobulin A1-A2 lambda hybrid GAU heavy	0
4430.seq	351	gi 4758435	siens gastric inhibitory polypeptide (GIP) mRNA	0

4431.seq	606	gi 34111162 gb AF075589 AF075589	Homo sapiens MDA-MB-23	-type benzodiazepine receptor	0
4432.seq	656	gi 2738086 gb AF017257 AF017257	Homo sapiens chromosome	BAC containing erythroblastosis	0
4433.seq	594	gi 4504448	Homo sapiens heterogeneou	nucleooreoprotein D (HNRPD) mRNA	0.003
4434.seq	435	gi 2335066 gb AC002430 AC002430	Human BAC clone RG066D1	2, complete sequence [Homo sapiens]	0.12
4436.seq	593	gi 5174602	Homo sapiens myosin regula	nain 2, smooth muscle isoform	0
4437.seq	541	gi 2605593 dbj D50372 D50372	Homo sapiens mRNA for my	tory light chain, complete cds	0
4438.seq	534	gi 3766261 emb AL031904 HS103M22	Human DNA sequence from	t22 on chromosome 5p24. Contains ST	0.037
4440.seq	606	gi 4502598	Homo sapiens carbonyl redu	1R1) mRNA	0
4441.seq	505	gi 3327388 gb AC004483 ATAC004483	Arabidopsis thaliana chromo	C T12J2 genomic sequence, complete	0.55
4442.seq	568	gi 5531251 emb AL035659.2 HS979N1	Human DNA sequence from	11 on chromosome 22q13.1-13.32,	0
4443.seq	670	gi 4101573 gb AF004876 AF004876	Homo sapiens 54TmP (54tm	plete cds	0
4444.seq	651	gi 1930058 gb U95040 HSU95040	Human transcriptional corec	P1/TIF1B mRNA, complete cds	0
4445.seq	589	gi 468242 gb U07989 HSU07989	Human Burkitt's lymphoma i	ulin kappa light chain mRNA,	E-134
4446.seq	342	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exo		E-147
4448.seq	600	gi 5080755 gb AC007842.1 AC007842	Homo sapiens chromosome.	1191 (CIT-B-471f3), complete sequenc	6.00E-97
4450.seq	550	gi 4505018	Homo sapiens low density lip	ceptor-related protein 7	0
4451.seq	678	gi 498724 emb X78926 HSHZF32	H.sapiens HZF3 mRNA for z	rotein	0
4452.seq	505	gi 34433 emb X14008 HSLYSOZY	Human lysozyme gene (EC :		E-121
4453.seq	459	gi 4579999 dbj AP000078.1 AP000078	Homo sapiens genomic DNA	enesence gene, segment 14/19,	0.008
4454.seq	585	gi 1679673 gb U68385 HSU68385	Human Mels1-related prote), mRNA, partial cds	E-120
4455.seq	235	gi 2804346 gb AC004029 AC004029	Human BAC clone RG035E1	1, complete sequence [Homo sapiens]	0.062
4457.seq	295	gi 4502146	Homo sapiens amyloid beta	sor-like protein 2 (APLP2)	E-107
4458.seq	519				
4459.seq	640	gi 1944628 gb J01415.1 HUMMTCCG	Human mitochondrion, com	ne	0
4461.seq	599	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exo		0
4462.seq	675	gi 4503270	Homo sapiens decorin (DCN	ii181169 gb M14219 HUMCSPG1A	0
4463.seq	401	gi 1399085 gb U43701 HSU43701	Human ribosomal protein L2	complete cds.	0
4464.seq	338	gi 30371 emb X05607 HSCYSTCR	Human mRNA for cysteine f	hibitor precursor cystatin C	E-160
4465.seq	614	gi 557545 gb M28699 HUMB23A	Homo sapiens nucleolar pho	B23 (NPM1) mRNA, complete cds.	0
4466.seq	597	gi 3360440 gb AF052131 AF052131	Homo sapiens clone 23930 r	ance	0
4467.seq	417	gi 4164368 emb AL031846 HS742C19	Human DNA sequence from	19 on chromosome 22q12.3-13.1.	0
4468.seq	404	gi 190416 gb M11948 HUMPROL	Human promyelocytic leuke	NA, clones pHH58 and pHH81.	E-159
4469.seq	658	gi 4003388 dbj AB020864.1 AB020864	Homo sapiens genomic DNA	p22 anti-oncogene of	0
4470.seq	673	gi 4507668	Homo sapiens tumor protein	tally-controlled 1 (TPT1)	0
4471.seq	573	gi 190416 gb M11948 HUMPROL	Human promyelocytic leuke	NA, clones pHH58 and pHH81.	0
4472.seq	444	gi 4099322 gb U85976 HSU85976	Human nucleoside diphosph	NDK) pseudogene, partial cds	0
4473.seq	635	gi 4502726	Homo sapiens cadherin 5, VI	(vascular epithelium) (CDH5)	0
4474.seq	290	gi 6136940 emb AL035411.2 HS592A1	Human DNA sequence from	v1 on chromosome 1p31.2-32.3, comple	E-153
4475.seq	580	gi 4886265 emb AL050300.1 ATF2206	Arabidopsis thaliana DNA cl	3, BAC clone F2206	0.16
4476.seq	552	gi 2632099 emb Z73639 HSV389H8	Human DNA sequence from	89H8 on chromosome X. Contains mRN	7.00E-84
4477.seq	655	gi 3721650 dbj AB012910 AB012910	Homo sapiens mRNA for arr	nmunoglobulin Fab kappa chain,	0
4478.seq	633	gi 2335058 gb U95741 HSU95741	Human chromosome 16p13.	rom CIT987SK-777B5 complete sequenc	0.045
4479.seq	669	gi 4827328 gb AC004854.2 AC004854	Homo sapiens PAC clone D.	rom 7p13-p11.2, complete sequence	0
4480.seq	628	gi 6137796	Human mitochondrion, com	ne	0
4481.seq	751	gi 6164987 gb AC007869.2 AC007869	Genomic Sequence for Hom	ione 431C18 From Chromosome 8 (loc	2.00E-04
4482.seq	339	gi 2673897 gb AC000385 AC000385	Human Chromosome 11 par	7, complete sequence [Homo sapiens]	E-124
4483.seq	582	gi 6318317 gb AF100154.1 AF100154	Rattus norvegicus kynurenin	iferase/glutamine transaminase K	2.5
4485.seq	496	gi 3746532 gb AF059202 AF059202	Homo sapiens ACAT relatet	uct 1 mRNA, complete cds	E-108
4486.seq	641	gi 4972129 emb AL049832.2 CN00000G	Human chromosome 14 DN.	*** IN PROGRESS *** BAC R-63D17	7.00E-04
4487.seq	643	gi 4503470	Homo sapiens eukaryotic tri	ngation factor 1 alpha 1	0
4488.seq	589	gi 4503480	Homo sapiens eukaryotic tri	ngation factor 1 gamma (EEF1G)	0
4489.seq	473	gi 4758789	Homo sapiens NADH dehyd	biquinone) Fe-S protein 5 (15kD)	0
4490.seq	648	gi 178746 gb M81955 HUMAPEA	Human apurinic/apyrimidin	ase (HAP1h) mRNA, complete	0
4491.seq	534	gi 1944628 gb J01415.1 HUMMTCCG	Human mitochondrion, com	ne	0
4492.seq	414	gi 505472 emb X69181 HSL31	H.sapiens mRNA for ribosor	L31	0
4493.seq	453	gi 4160143 gb AC006213 AC006213	Homo sapiens, clone hRPK.	nplete sequence [Homo sapiens]	0.032
4494.seq	608	gi 3461801 gb AC005542 AC005542	Homo sapiens chromosome	R32516, complete sequence [Homo	0.17
4495.seq	691	gi 546798 gb S71043 S71043	Ig alpha 2-immunoglobulin /	in allotype 2 (constant	E-124
4496.seq	561	gi 2765422 emb Y14736 HSIGG1KL	Homo sapiens mRNA for im	in kappa light chain	0
4497.seq	256	gi 13000 emb X62996 MH5GENOM	H.sapiens mitochondrial ger	nisus sequence)	8.00E-91
4498.seq	601	gi 2832593 emb Z97056 HS434P1	Human DNA sequence from	1 on chromosome 22. Contains inward	0
4499.seq	648	gi 4506662	Homo sapiens ribosomal prc	L8) mRNA	0
4500.seq	559	gi 5305595 gb AF102989.1 AF102989	Homo sapiens Ca2+-indepe	pholipase A2 long isoform (IPLA2)	0
4501.seq	589	gi 5031692	Homo sapiens heterogeneo	rotein similar to rat helix	E-149
4502.seq	579	gi 4836804 gb AF146793.1 AF146793	Mus musculus protein B ger	ts, and CLOCK (Clock), PFT27	0.64
4504.seq	394	gi 3810574 gb AC004671 AC004671	Homo sapiens 12p13.3 PAC	113 (Roswell Park Cancer Institute	0.11
4505.seq	228	gi 5174730	Homo sapiens translin-asso	cy X (TSNAX) mRNA	3.00E-56
4506.seq	604	gi 536067 dbj AB022656.1 AB022656	Homo sapiens mRNA for an	iba histolytica Immunoglobulin	0
4508.seq	585	gi 498014 gb L27479 HUMX123A	Human X123 mRNA, 3' end.		0
4509.seq	590	gi 2801527 gb AF035397 AF035397	Homo sapiens cosmids Qc:1	B6 from Xq28, complete sequence	0.16
4510.seq	552	gi 536067 dbj AB022656.1 AB022656	Homo sapiens mRNA for an	iba histolytica Immunoglobulin	0
4511.seq	554	gi 2605593 dbj D50372 D50372	Homo sapiens mRNA for m	tory light chain, complete cds	0
4512.seq	642	gi 6013496 gb AC008000.7 AC008000	Homo sapiens 3q25-26 BAC	Q18 (California Institute of	0.71
4513.seq	516	gi 4503286	Homo sapiens growth arres	damage-inducible, alpha	0
4516.seq	618	gi 4506800	Homo sapiens secretogranli	granin C) (SCG2) mRNA	0
4517.seq	752	gi 5757524 gb AC007385.3 AC007385	Homo sapiens BAC clone N	from 2, complete sequence	3.3
4519.seq	418	gi 181532 gb M97925 HUMDEFA	H.sapiens defensin 5 gene.	ts.	6.00E-96
4520.seq	570	gi 950414 gb L37516 HUMPSDBPB	Homo sapiens DNA-binding	pseudogene (PSDBPB1) gene.	0
4521.seq	605	gi 5032148	Homo sapiens TATA box bl	in (TBP)-associated factor. RNA	0
4523.seq	416	gi 3885376 dbj AB019573 AB019573	Homo sapiens mRNA expre	placental villi, clone SMAP31	5.00E-81
4524.seq	610	gi 4759095	Homo sapiens 15 kDa selen	EP15) mRNA	0
4526.seq	507	gi 188483 gb M80469 HUMMHHLAJB	Human MHC class I HLA-J	s 1-8 and complete cds.	0
4527.seq	476	gi 5174744	Homo sapiens ubiquinol-cyl	eductase hinge protein (UQCRH)	0
4528.seq	365	gi 337513 gb J03537 HUMRPS6	Human ribosomal protein S	mplete cds.	0
4529.seq	542	gi 1314309 gb U41743 HSU41743	Human nucleophosmin-reti	ceptor alpha fusion protein	0
4531.seq	627	gi 6049285 gb AF163570.1 AF163570	Homo sapiens DINB protein	RNA, complete cds	9.00E-25
4532.seq	584	gi 4758011	Homo sapiens clathrin, hea	ide-like 2 (CLTCL2) mRNA	0
4533.seq	859	gi 5729729	Homo sapiens APIS-like 1 (?NA	1.00E-24
4534.seq	522	gi 5280990 emb Z97338.2 ATFCA3	Arabidopsis thaliana DNA c	4, ESSA1 FCA contig fragment No. 3	0.037
4535.seq	544	gi 1944628 gb J01415.1 HUMMTCCG	Human mitochondrion, com	ne	0
4536.seq	628	gi 577310 dbj D42054 HUMKIAAM	Human mRNA for KIAA009	nplete cds	0
4539.seq	583	gi 4502744	Homo sapiens cyclin-deper	8 (CDK8) mRNA	E-109
4540.seq	623	gi 4505574	Homo sapiens poly(A)-bind	-like 1 (PABPL1) mRNA	0
4541.seq	410	gi 181532 gb M97925 HUMDEFA	H.sapiens defensin 5 gene.	ts.	5.00E-90
4542.seq	539	gi 4884260 emb AL050005.1 HSM80035	Homo sapiens mRNA: cDN	4A153 (from clone DKFZp564A153)	0
4543.seq	865	gi 4884467 emb AL050225.1 HSM80052	Homo sapiens mRNA: cDN	6M1523 (from clone DKFZp586M1523)	0
4544.seq	363	gi 4507132	Homo sapiens small nuclea	protein polypeptide G (SNRPG)	0
4545.seq	525	gi 2076720 gb AC002067 HSAC002067	Human BAC clone RG126h	i21-a22, complete sequence [Homo	0.15
4546.seq	634	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exo		E-171
4547.seq	593	gi 6005977	Homo sapiens zinc finger p	ZNF258), mRNA	2.00E-78
4548.seq	445	gi 186840 gb J03799 HUMLAMB	Human colon carcinoma lan	g protein mRNA, complete cds.	0
4549.seq	706	gi 1688100 gb U68299 MCU68299	Mouse cytomegalovirus 1 c	normic sequence	0.003
4550.seq	579	gi 1668818 gb L29074.1 HUMFMFR1S	Homo sapiens fragile X me	tion syndrome protein (FMR1) gene.	0
4551.seq	505	gi 6226515	Saccharomyces cerevisiae	ion, complete genome	0.035
4552.seq	413	gi 4455632 emb Z97353.3 HS90L6	Human DNA sequence from	6 on chromosome 22q11.21-11.23.	7.00E-71
4553.seq	473	gi 6325471	Homo sapiens ribosomal pr	(RPL10A), mRNA	0
4554.seq	93	gi 4761685 gb AF147334.1 AF147334	Homo sapiens full length in	clone YB24H02	8.00E-33
4556.seq	708	gi 6137796	Human mitochondrion, com	ne	0

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4557.seq	585	gi 3047019 gb AF057177 AF057177	apiens T-cell receptor gamma V1 gene region	0.041
4558.seq	559	gi 3483501 gb AF086156 HUMZB55H	apiens full length insert cDNA clone ZB55H12	0
4559.seq	350	gi 13153224 gb AF034899 AF034899	norvegicus olfactory receptor-like protein (SCR D-9) gene,	0.095
4560.seq	569	gi 6118386 gb AF187991.1 AF187991	apiens zinc finger protein ZNF225 mRNA, complete cds	0
4562.seq	517	gi 532956 gb J12759 HUMIGJ02	Ig J chain gene, exons 3 and 4.	0
4564.seq	871	gi 4505774	apiens phosphate carrier, mitochondrial (PHC), splice variant	E-132
4565.seq	509	gi 5668757 gb AC008009.4 AC008009	apiens 3q26.2-27 BAC RPC111-436A20 (Roswell Park Cancer Institute)	0
4566.seq	655	gi 4557520	apiens diaphorase (NADH) cytochrome b-5 reductase (DIA1),	0
4567.seq	551	gi 4730826 dbj AP000085.1 AP000085	apiens genomic DNA of 21q22.1, APP related, B2291C14-T1533	0.01
4569.seq	595	gi 5803200	apiens transmembrane trafficking protein (TMP21), mRNA	0
4570.seq	499	gi 2335186 gb AF011889 AF011889	Xq28 cosmids U126G1, U142F2, U69B6, U145C10, U169A5, U84H1, U	0.14
4571.seq	584	gi 4506730	apiens ribosomal protein S6 (RPS6) mRNA	0
4572.seq	552	gi 434060 gb L12535 HUMRSU1A	RSU-1/RSP-1 mRNA, complete cds.	0
4574.seq	874	gi 13701112 emb Z73963 HNS62C4	DNA sequence from cosmid N62C4, on chromosome 22 Contains ESTs	0.97
4575.seq	601	gi 23902 emb X59727 HS63KAP	ns 63 kDa protein kinase related to rat ERK3	0
4576.seq	564	gi 4185494 gb AF095891 AF095891	apiens calpastatin mRNA, partial cds and 3' UTR	5.00E-17
4577.seq	533	gi 1008967 emb Z47034 HS98D12E	partial cDNA sequence, complete sequence of clone 98D12:	0
4579.seq	444	gi 306548 gb L13799 HUMCH13C1A	apiens (clone 01) liver expressed protein mRNA, 3'end.	0
4582.seq	499	gi 2935477 gb AF048836 AF048836	dium falciparum guanidine nucleotide exchange factor (PfRCC1)	2.2
4583.seq	548	gi 2935477 gb AF048836 AF048836	dium falciparum guanidine nucleotide exchange factor (PfRCC1)	0.6
4584.seq	633	gi 2623584 gb AF027158 AF027158	apiens immunoglobulin kappa light chain (T6J/k) mRNA, partial	0
4585.seq	862	gi 5262473 emb AL080066.1 HSM800	apiens mRNA; cDNA DKFZp564J142 (from clone DKFZp564J142)	0
4586.seq	588	gi 4507466	apiens transforming growth factor, beta-induced, 68kD (TGFB1)	0
4587.seq	444	gi 1944628 gb J01415.1 HUMMTCG	mitochondrion, complete genome	0
4588.seq	497	gi 5803071	apiens LIM domain only 4 (LMO4), mRNA	0
4589.seq	605			
4590.seq	533	gi 1591687 gb U67546 U67546	ococcus jannaschii section 88 of 150 of the complete genome	0.037
4593.seq	659	gi 1145815 gb U42031 HSU42031	54 kDa progesterone receptor-associated immunophilin FKBP54	0
4594.seq	660	gi 546798 gb S71043 S71043	gamma 2-immunoglobulin A heavy chain allotype 2 (constant region,	0
4595.seq	416	gi 4503304	apiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	0
4597.seq	604	gi 4502302	apiens ATP synthase, H+ transporting, mitochondrial F1	0
4598.seq	621	gi 31091 emb X16869 HSEF1AC	mRNA for elongation factor 1-alpha (clone CEFA)	0
4599.seq	568	gi 4557492	apiens cleavage stimulation factor, 3' pre-RNA, subunit 2, 64kD	0
4600.seq	440	gi 5708471 gb AC007041.3 AC007041	apiens clone NH0327N17, complete sequence	0
4603.seq	568	gi 222559 dbj D00736 RSHBCNP	respiratory syncytial virus 1B, 1C, N, and P genes	0.16
4604.seq	871	gi 32797 emb X61011 HSIG486	Immunoglobulin VH mRNA (486 bp)	0
4605.seq	474	gi 386007 gb AF097021 AF097021	apiens GW112 protein (GW112) mRNA, complete cds	E-112
4606.seq	518	gi 177830 gb K02212 HUMA1ATP	alpha-1-antitrypsin gene (S variant), complete cds.	0.57
4607.seq	507	gi 1663691 dbj D80011 D80011	mRNA for KIAA0189 gene, complete cds	0
4608.seq	482	gi 5803130	apiens RAB31, member RAS oncogene family (RAB31), mRNA	0
4609.seq	508	gi 6086099 gb AB027251.1 AB027251	apiens mRNA for zinc finger protein (ZF25), complete cds	E-100
4610.seq	510	gi 5565976 gb AF152462.1 AF152462	apiens transmembrane protein BRI (BRI) mRNA, complete cds	0
4611.seq	495	gi 31896 emb X58295 HSGXP	GPx-3 mRNA for plasma glutathione peroxidase	0
4612.seq	469	gi 5306223 gb AC006255.9 AC006255	apiens 3p21.1-9 PAC RPC15-1087L12 (Roswell Park Cancer Institute)	E-134
4613.seq	512	gi 3387884 gb AF070529 AF070529	apiens clone 24525 mRNA sequence	0
4614.seq	490	gi 4506232	apiens proteasome (prosome, macropain) 26S subunit,	0
4615.seq	510	gi 632994 gb L39063 HUMIL9RPS	apiens interleukin 9 receptor (IL9R) pseudogenes, exons 1-9	2.2
4616.seq	458	gi 4503304	apiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	0
4617.seq	514	gi 3132318 dbj AP000008.1 AP000008	apiens genomic DNA of 21q22.2 Down Syndrome region, segment 1/15	7.00E-96
4618.seq	475	gi 5468516 gb AF107885.2 DU270M1	apiens chromosome 14q24.3 clone BAC270M14 transforming growth	0
4619.seq	525	gi 2088550 gb U91328 HSU91328	hereditary haemochromatosis protein, histone 2A-like protein gene,	E-171
4621.seq	340	gi 3659494 gb AC005332 AC005332	apiens chromosome 17, clone hRPK.147.L.13, complete sequence	0.024
4623.seq	511	gi 5902013	apiens nuclear pore complex interacting protein (NPIP), mRNA	0
4625.seq	409	gi 184743 gb J00220 HUMIGCC8	apiens immunoglobulin alpha-1 heavy chain constant region	0
4626.seq	526	gi 414346 dbj D14531 HUMHRRPL5	mRNA for human homologue of rat ribosomal protein L9.	0
4627.seq	397	gi 693936 emb Z48501 HSPABP11	ns mRNA for polyadenylate binding protein II	0
4629.seq	505	gi 414346 dbj D14531 HUMHRRPL5	mRNA for human homologue of rat ribosomal protein L9.	0
4630.seq	471	gi 4503470	apiens eukaryotic translation elongation factor 1 alpha 1	0
4631.seq	497	gi 2335063 gb AC002433 AC002433	BAC clone RG317M02 from 7p15-p21, complete sequence [Homo	0.14
4632.seq	512	gi 4699970 gb AC005997.2 AC005997	apiens PAC clone DJ0558L10 from 7q34-q36, complete sequence	0.036
4633.seq	496	gi 5668655 emb AL050331.11 HSDJ4	1 DNA sequence from clone 48613 on chromosome 6q22.1-22.3, Contain	6.00E-04
4634.seq	513	gi 4758179	apiens cell cycle progression 3 protein (DNJ3) mRNA	0
4635.seq	519	gi 1850553 gb U88032 SCU88032	cereale Ty-1 copia retrotransposon reverse transcriptase	0.037
4637.seq	525	gi 3201899 gb AF067420 AF067420	apiens SNC73 protein (SNC73) mRNA, complete cds	E-110
4638.seq	503	gi 2853186 emb AL021786 HSB96H2	1 DNA sequence from PAC 696H22 on chromosome Xq21.1-21.2.	E-168
4639.seq	502	gi 5174602	apiens myosin regulatory light chain 2, smooth muscle isoform	0
4640.seq	501	gi 4507668	apiens tumor protein, translationally-controlled 1 (TPT1)	0
4641.seq	553	gi 189021 gb M22920 HUMMYLCC	1 smooth muscle myosin alkali light chain (MLC 1sm) mRNA,	0
4642.seq	502	gi 2293555 gb AF012086 AF012086	apiens Ran binding protein 2 (RanBP2aloha) mRNA, partial cds	0
4643.seq	509	gi 4240294 dbj AB020710.1 AB020710	apiens mRNA for KIAA0903 protein, partial cds	0
4644.seq	515	gi 4506660	apiens ribosomal protein L7a (RPL7a) mRNA	0
4645.seq	484	gi 3954884 emb J010442 HSA0104	apiens mRNA for immunoglobulin kappa light chain, anti-RhD,	E-165
4647.seq	518	gi 262368 gb S51768 S51768	1-pancreatitis-associated protein [human, pancreas, mRNA, 797	0
4648.seq	508	gi 2911720 gb AC004226 AC004226	apiens chromosome 5, BAC clone 313n8 (LBXN H146), complete	0.56
4650.seq	305	gi 2895757 gb AF045022 AF045022	urus phosphatidic acid-preferring phospholipase A1 mRNA,	2.00E-27
4651.seq	512	gi 1685101 gb U47101 HSLU47101	1 NIFU-like protein (hNIFU) mRNA, partial cds	0
4652.seq	516	gi 5103850 gb AC007651.2 AC007651	iosis thalana chromosome 1 BAC F20D23 genomic sequence, complete	0.009
4654.seq	382	gi 2735711 gb AF004023 AF004023	usculus cell surface molecule OX-2 mRNA, complete cds	8.00E-12
4655.seq	14			
4656.seq	464	gi 4506356	apiens NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 4	0
4659.seq	517	gi 4769003 gb AF140598.1 AF140598	apiens ring-box protein 1 (RBX1) mRNA, complete cds	0
4660.seq	312	gi 5091653 gb AC005519.2 AC005519	apiens clone DJ0919J22, complete sequence	5.00E-37
4661.seq	499	gi 181915 gb M91670 HUME2EPI	1 ubiquitin carrier protein (E2-EPF) mRNA, complete cds.	8.00E-68
4663.seq	102	gi 190416 gb M11948 HUMPROL	1 promyelocytic leukemia cell mRNA, clones pH58 and pHH81.	3.00E-17
4664.seq	521	gi 339688 gb M17733 HUMTHYB4	1 thymosin beta-4 mRNA, complete cds.	0
4665.seq	497	gi 5650629 emb AL078593.9 HSDJ2	1 DNA sequence from clone 21081 on chromosome 6q23.3-25.1. Contain	0.55
4666.seq	506	gi 5656681 gb AC007297.22 AC007297	apiens 12q22 BAC RPC111-690J15 (Roswell Park Cancer Institute)	0.002
4668.seq	339	gi 5123989 gb AC006987.2 AC006987	apiens clone NH0160K17, complete sequence	1.4
4669.seq	505	gi 5669135 gb AF152365.1 AF152365	apiens constitutive fragile region FRA3B sequence	0.14
4670.seq	531	gi 1478196 emb X97321 HSHLACWC	ens mRNA for human leukocyte antigen C alpha chain	0
4671.seq	512	gi 6137796	1 mitochondrion, complete genome	0
4672.seq	524	gi 5174722	apiens DNA segment on chromosome 19 (unique) 1177 expressed	0
4673.seq	21	gi 3980107 emb AL020993.1 HS506	1 DNA sequence from clone 506 on chromosome 22q12 Contains the 5'	0.001
4674.seq	321	gi 181532 gb M97925 HUMDEFA5A	ens defensin 5 gene, complete cds.	7.00E-89
4676.seq	489	Homo sapiens small inducible cytokin		0
4679.seq	517	gi 292301 gb L14299 HUMMF2D55	apiens (chromosome 1) DNA sequence, (CA)n dinucleotide	0.14
4680.seq	501	gi 4504766	apiens integrin, beta 1 (fibronectin receptor, beta	0
4681.seq	466	gi 1906003 gb U90540 DMU90540	phila melanogaster chromosomal form domain cut site	0.13
4682.seq	377	gi 961441 dbj D63875 D63875	1 nRNA for KIAA0155 gene, complete cds	0
4683.seq	504	gi 4884198 emb AL049951.1 HSM8C	apiens mRNA; cDNA DKFZp564A00122 (from clone DKFZp564A00122)	0
4684.seq	479	gi 3483353 gb AF086008 HUMYU69	apiens full length insert cDNA clone YU69D10	2.00E-93
4685.seq	493	gi 498730 emb X78929 HSHZFS	ens HZF6 mRNA for zinc finger protein	8.00E-65
4686.seq	503			
4688.seq	398	gi 1183970 emb Z69381 SCCXIV39H	1 vislae 38,855 bp segment of chromosome XIV	0.43
4691.seq	510	gi 5832741 dbj AP000420.1 AP0004	iosis thalana genomic DNA, chromosome 5, P1 clone: MZN14,	2.2
4692.seq	462	gi 3980553 gb AC004849 AC004849	1 sapiens PAC clone DJ0659J06 from 7q33-q35, complete sequence [Homo	0.032
4694.seq	320	gi 2921499 gb AF037222 AF037222	1 DNA from chromosome 14-specific cosmid containing XRCC3 DNA	E-151
4695.seq	503	gi 1035472 emb Z63094 HST8C5F	1 ens CpG island DNA genomic MseI fragment, clone 78c5, forward	1.00E-85

4696.seq	496					
4697.seq	501	gi 4506846	Homo sapiens small inducibl	A5 (RANTES) (SCYA5) mRNA	0	
4698.seq	521	gi 5032050	Homo sapiens ribosomal prc	PS14) mRNA	0	
4699.seq	500	gi 4557590	Homo sapiens fibrillin 1 (Mar	me) (FBN1) mRNA	0	
4700.seq	15					
4701.seq	465	gi 37847 emb X16478 HSVIMCD	Human mRNA 5'-fragment	7 N-terminal fragment	0	
4703.seq	494	gi 3281985 emb AL022165 HS71L16	Human DNA sequence from	6 on chromosome Xp11. Contains a	0	
4704.seq	497	gi 183677 gb M14777 HUMGSTH	Human glutathione S-trans	f)-subunit mRNA, complete cds.	0	
4705.seq	516	gi 2773059 gb AF039018 AF039018	Homo sapiens actinin- assoc	rotein mRNA, complete cds	E-165	
4706.seq	487	gi 5802983	Homo sapiens i-beta-1,3-N	osaminyltransferase	E-103	
4707.seq	250	gi 307378 gb L10376 HUMRNAE	Human (clone CTG-B33) m	nce.	E-103	
4708.seq	270	gi 4826869	Homo sapiens nucleobindin:	mRNA	E-133	
4710.seq	512	gi 4309780 gb AC006566 AC006566	Homo sapiens chromosome	RPK.376_P_6, complete sequence [Hom	0.56	
4711.seq	526	gi 2055296 dbj D87687 D87687	Xenopus laevis mRNA for L1	ete cds	0.58	
4712.seq	225	gi 5031814	Homo sapiens lysyl-tRNA sy	(ARS) mRNA	E-120	
4713.seq	503	gi 2911509 gb AF036364 AF036364	Homo sapiens epslon-sarcoc	G) mRNA, complete cds	0	
4714.seq	516	gi 4506730	Homo sapiens ribosomal prc	'S6) mRNA	0	
4715.seq	513	gi 4506470	Homo sapiens regenerating	ad 1 alpha (pancreatic stone	0	
4716.seq	174	gi 598165 gb L33854 HUMIGKZM	Human immunoglobulin ligh	able region (Vk-IV subgroup)	7.00E-69	
4718.seq	275	gi 5636191 gb AC007250.2 AC007250	Homo sapiens clone NH033-	lete sequence	E-106	
4719.seq	491					
4720.seq	211	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene. c	ls.	2.00E-85	
4721.seq	323	gi 4506698	Homo sapiens ribosomal prc	PS21) mRNA	E-176	
4722.seq	146	gi 4507574	Homo sapiens tumor necros	ceptor superfamily, member 1A	0.15	
4725.seq	257	gi 2230868 emb Y09188 HSFERRL	H.sapiens mRNA for ferritin		E-124	
4727.seq	302	gi 2224590 dbj ABD02323 ABD02323	Human mRNA for KIAA032:	bial cds	E-151	
4728.seq	182					
4729.seq	219	gi 1223861 gb U35430 HSU35430	Human cytochrome c oxidase	ii (COIII) pseudogene.	2.00E-91	
4730.seq	446	gi 2815069 emb AL021451 HS188H10	Human DNA sequence from	10 on chromosome 6q22. Contains STS	0	
4735.seq	429	gi 4758125	Homo sapiens DEAD/H (Asi	Asp/His) box polypeptide, Y	0	
4739.seq	521	gi 190978 gb M18963 HUMREGA	Human islet of Langerhans I	g protein (reg, complete	0	
4740.seq	143	gi 2281706 gb AF014838 AF014838	Homo sapiens galectin-4 m	lete cds	9.00E-74	
4741.seq	300	gi 5730026	Homo sapiens GAP- associa	phosphoprotein p62 (Sam68)	E-135	
4742.seq	298	gi 3522915 gb AC005145 AC005145	Homo sapiens Xp22-166-16	23A23 (Genome Systems Human BAC	0.081	
4743.seq	292	gi 307194 gb L09260 HUMMEMPROT	Human (chromosome 3p25)	:protein mRNA.	E-135	
4744.seq	485	gi 5596765 emb AL049557.19 HSJ773A1	Human DNA sequence from	118 on chromosome 1p13.2-21.1 Contai	0	
4746.seq	413	gi 4240160 dbj AB020643.1 AB020643	Homo sapiens mRNA for Kl	tein, partial cds	0	
4748.seq	370	gi 3420063 gb AC005492 AC005492	Homo sapiens clone UWGC	from 7p14-15, complete sequence [Ho	0.1	
4750.seq	323	gi 644480 dbj D38112.1 HUMMTA	Homo sapiens mitochondrial	plete sequence	E-108	
4751.seq	523	gi 2385366 emb X80199 HSMNLN51	H.sapiens MLN51 mRNA		E-151	
4752.seq	154	gi 3928512 dbj AB005144 AB005144	Magnaporthe grisea mRNA	ive oxidase, complete cds	0.15	
4754.seq	252	gi 2828601 gb AF026126 AF026126	Homo sapiens heterogeneo	ribonucleoprotein D (HNRPD) gene.	E-139	
4755.seq	285	gi 2439515 gb AC002563 AC002563	Human PAC clone 127H14	omplete sequence [Homo sapiens]	1.2	
4756.seq	531	gi 1944628 gb J01415.1 HUMMTCG	Human mitochondrion, com	ne	0	
4757.seq	501	gi 186840 gb J03799 HUMLAMB	Human colln carcinoma lam	protein mRNA, complete cds.	E-148	
4758.seq	502	gi 3044065 gb AF052389 AF052389	Homo sapiens LIM domain E	sin (LDB1) mRNA, complete cds	E-148	
4759.seq	262	gi 4885524	Homo sapiens novel SH2-c	rotein 1 (NSP1) mRNA	8.00E-14	
4760.seq	496	gi 5107001 gb AF125103.1 AF125103	Homo sapiens neuroendocr	rotein c homolog mRNA, complete	0	
4761.seq	345	gi 3970970 gb AC004812 AC004812	Homo sapiens PAC clone 26	12, complete sequence [Homo sapiens]	E-105	
4762.seq	501	gi 4544348 gb AC006059.3 AC006059	Homo sapiens PAC RPC14-	swell Park Cancer Institute Human PA	E-153	
4763.seq	293	gi 1944628 gb J01415.1 HUMMTCG	Human mitochondrion, com	ne	E-145	
4765.seq	532	gi 733134 gb U22384 HSU22384	Human lysyl oxidase gene. c		0	
4766.seq	282	gi 5304892 emb AL049575.7 HSJ316D	Human DNA sequence from	27 on chromosome 11p13. Contains par	0.076	
4767.seq	498	gi 5679789 emb AL049643.12 HSJ672M	Human DNA sequence from	415 on chromosome Xp21.1-21.3.	2.2	
4768.seq	499	gi 4506666	Homo sapiens ribosomal prc	PD (RPLP0) mRNA	0	
4769.seq	502	gi 4507108	Homo sapiens synuclein, alp	component of amyloid	0	
4770.seq	495	gi 4505034	Homo sapiens lymphotoxin I	xuperfamily, member 3) (LTB)	0	
4771.seq	483	gi 4502100	Homo sapiens annexin I (lipo	NX1) mRNA	0	
4772.seq	500	gi 5881950 gb AF064257.1	Homo sapiens Dhml- like pr	%, complete cds	0	
4774.seq	508	gi 4503470	Homo sapiens eukaryotic tr.	ngation factor 1 alpha 1	0	
4776.seq	184	gi 5679750 emb AL023807.6 HS525L6	Human DNA sequence from	.6 on chromosome 6p22.3-23, complete	9.00E-53	
4777.seq	500	gi 4506370	Homo sapiens RAB5B, merr	xogene family (RAB5B) mRNA	2.00E-84	
4778.seq	469	gi 4263802 gb AC006602 AC006602	Homo sapiens Chromosome	3) PAC clone pDJ47619 from the	0.13	
4779.seq	498	gi 3518151 emb AL031114 HS468H17	Human DNA sequence from	417 on chromosome Xq21.3-22.1.	0.035	
4780.seq	501	gi 3327109 dbj AB014548 AB014548	Homo sapiens mRNA for Kl	tein, partial cds	0	
4781.seq	468	gi 1620014 dbj D87665 D87665	Human heart mRNA for MH-	A-Bw62, partial cds	0	
4783.seq	505	gi 2606593 dbj D50372 D50372	Homo sapiens mRNA for m	itory light chain, complete cds	0	
4785.seq	507	gi 2565045 gb U80735 HSU80735	Homo sapiens CAGF28 mR	cds	0	
4789.seq	503	gi 6272679 gb AF181252.1 AF181252	Mus musculus testis TPR-c	rotein involved in	7.00E-59	
4790.seq	508	gi 4503470	Homo sapiens eukaryotic tr.	ngation factor 1 alpha 1	0	
4791.seq	500	gi 3360399 gb AF052093 AF052093	Homo sapiens clone 23685	ence	0	
4792.seq	400	gi 3550044 emb AL022577 HS353H6	Human DNA sequence from	46 on chromosome Xq25-26.2. Contains	2.00E-98	
4794.seq	501	gi 6331419 dbj AB033115.1 AB033115	Homo sapiens mRNA for Kl	tein, partial cds	0	
4795.seq	512	gi 1537069 gb U63840 RNU63840	Rattus norvegicus nucleopo	NA, complete cds	1.00E-14	
4796.seq	536	gi 181572 gb M34482 HUMDKERB	Human cytokerin 8 (CK8)	cds.	E-163	
4797.seq	452	gi 4504478	Homo sapiens hydroxyprost	hydrogenase 15- (NAD) (HPGD) mRNA	0	
4798.seq	520	gi 13000 emb X62996 MIHSGENOM	H.sapiens mitochondrial ger	ensus sequence)	7.00E-93	
4799.seq	568	gi 5817115 emb AL110197.1 HSM80084	Homo sapiens mRNA; cDN	3J021 (from clone DKFZp586J021)	0	
4800.seq	521	gi 2746542 gb AF037643 AF037643	Homo sapiens 60S ribosom	12 (RPL12) pseudogene, partial	0	
4801.seq	626	gi 3165402 gb AC004752 AC004752	Homo sapiens chromosome	8e5 (LBNL H167), complete sequenc	0	
4802.seq	417	gi 3402263 emb X94253.1 SS-NRNP	S.scrofa mRNA for heterog	lear ribonucleoprotein	E-161	
4803.seq	576	gi 3954892 emb AJ010446 HSA010446	Homo sapiens mRNA for irr	lin kappa light chain,anti-RhD.	0	
4804.seq	694	gi 5031856	Homo sapiens lactate dehyx	ase (LDHA), nuclear gene encoding	0	
4806.seq	600	gi 1537067 gb U63839 RNU63839	Rattus norvegicus nucleopo	NA, complete cds	2.00E-16	
4807.seq	626	gi 4506724	Homo sapiens ribosomal pri	linked (RPS4X) mRNA	0	
4808.seq	449	gi 4758203	Homo sapiens dual specifical	ase 1 (DUSP1) mRNA	0	
4809.seq	769	gi 4884176 emb AL049932.1 HSM80027	Homo sapiens mRNA; cDN	4H2416 (from clone DKFZp564H2416)	0	
4810.seq	558	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exc		E-172	
4811.seq	614	gi 1888539 gb U89355 HSU89355	Homo sapiens clone cRT16	fling protein mRNA, partial cds	0	
4812.seq	496	gi 1827474 dbj D50683 D50683	Homo sapiens mRNA for T(alpha, complete cds	0	
4813.seq	142	gi 4336689 gb AF101037 AF101037	Morone americana gamma:	ric-acid receptor rho-3 subunit	0.55	
4814.seq	246	gi 2653365 emb AL008710 HS292H14	Human DNA sequence from	114 on chromosome Xp21. Contains STS	4	
4815.seq	653	gi 160466 emb Z68886 HSN2E9C	Human DNA sequence from	4E9 on chromosome 22. Contains EST.	4.00E-30	
4816.seq	452	gi 4506656	Homo sapiens ribosomal pri	'L6) mRNA	0	
4817.seq	661	gi 4506666	Homo sapiens ribosomal pri	PD (RPLP0) mRNA	E-179	
4818.seq	630	gi 3646023 emb AL031228 HS1033B10	Human DNA sequence from	3B10 on chromosome 6p21.2-21.31.	0.011	
4819.seq	558	gi 189019 gb M22918 HUMMYLCB	Human non-muscle myosin	chain mRNA, 3' end.	0	
4821.seq	572	gi 5725210 emb AL109758.1 CNS0180V	Human chromosome 14 DN	*** IN PROGRESS *** BAC R-873E2	0	
4823.seq	592	gi 31916 emb X56009 HSGSAG	Human GSA mRNA for alpl	if GsGTP binding protein	0	
4825.seq	506	gi 35456 emb X54629 HSPHL1	Human pHL-1 gene is a c-	ne containing coxII sequence	0	
4826.seq	412	gi 5031740	Homo sapiens HIRA interac	14 (dnaJ-like) (HIRP4) mRNA	E-179	
4827.seq	370	gi 4557306	Homo sapiens aldolase B, fi	phosphate (ALDOB) mRNA	0	
4828.seq	474	gi 5689570 dbj AB029040.1 AB029040	Homo sapiens mRNA for Kl	tein, partial cds	0	
4829.seq	435	gi 598859 dbj D17252 HUMD5C03M3	Human HepG2 3' region Mt	lone hmd5c03m3	2.00E-46	
4830.seq	425	gi 445632 emb J297353.3 HS90L6	Human DNA sequence fro	6 on chromosome 22q11.21-11.23.	3.00E-36	
4831.seq	234	gi 178678 gb M88499 HUMANTIGPA	Human anti-glycoprotein 1	nal antibody variable region and	4.00E-83	
4832.seq	621	gi 4503470	Homo sapiens eukaryotic tr	longation factor 1 alpha 1	0	
4835.seq	546	gi 5262922 emb AL050308.9 HSJ164F24	Human DNA sequence fro	64F24 on chromosome Xq26.3-27.3.	6.00E-04	

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4837.seq	505	gi 3659494 gb AC005332 AC005332	apiens chromosome 17, clone hRPK.147.L.13, complete sequence [Homo	0
4838.seq	339	gi 4504864	apiens KH-type splicing regulatory protein (KHSRP) mRNA, and	4.00E-84
4839.seq	580	gi 2979567 gb AC002301 HUAC002301	apiens Chromosome 16 BAC clone CIT987SK-A-328A3, complete	0.01
4840.seq	181	gi 1710211 gb U97258 HSU79258	clone 23732 mRNA, partial cds	3.00E-68
4841.seq	202	gi 5901877 gb AF182645.1 AF182645	apiens chondrosarcoma-associated protein 2 (CSA2) mRNA,	7.00E-94
4842.seq	426	gi 307150 gb L19182 HUMMAC25X	MAC25 mRNA, complete cds	E-161
4843.seq	490	gi 4156182 gb AC004899 AC004899	apiens PAC clone DJ0815D20 from 7p11-p13, complete sequence [Homo	9.00E-09
4845.seq	577	gi 6137796	mitochondrion, complete genome	0
4846.seq	644	gi 4826474 emb AL031716.8 HS360B	DNA sequence from clone 360B4 on chromosome 16. Contains part	0.71
4847.seq	657	gi 4646258 gb AC007157.6 AC007157	apiens, clone hRPK.78.A.1, complete sequence	0.73
4848.seq	605	gi 3738102 gb AC005772 AC005772	apiens chromosome 17, clone hRPK.924.A.14, complete sequence [Homo	0.011
4850.seq	434	gi 4469083 emb AL031670.6 HS581N	DNA sequence from clone 681N20 on chromosome 20p12.1-13. Contain	E-176
4852.seq	595	gi 3386333 gb AF035120 AF035120	amilaris type 1 procollagen pro-alpha 2 chain (COL1A2)	0
4853.seq	474	gi 6358759 gb AC011665.5 ATAC011	opsis thalana chromosome I BAC T6L1 genomic sequence, complete	0.033
4855.seq	489	gi 4759017	apiens Rab geranylgeranyltransferase, beta subunit (RABGGTB)	0
4856.seq	727	gi 4079626 gb AC005839 AC005839	apiens chromosome 17, clone hRPK.481.C.4, complete sequence	E-138
4857.seq	625	gi 4885404	apiens hematopoietic cell-specific Lyn substrate 1 (HCLS1)	0
4858.seq	610	gi 4056515 gb AC006080 AC006080	apiens chromosome 17, clone hRPK.394.K.10, complete sequence	0.17
4859.seq	681			
4860.seq	505	gi 465173 gb L29157 HUMIGLZH	Immunoglobulin light chain variable region (lambda IIIB)	0
4861.seq	554	gi 4753268 gb AC006352.3 AC006352	apiens clone DJ1044J09, complete sequence	0.002
4867.seq	426	gi 2347130 gb AF019413.1 HSMHC1T	apiens HLA class III region containing tenascin X (tenascin-X)	E-164
4868.seq	548	gi 5532293 gb AF129332.1 AF129332	apiens MUM2 (MUM2) gene, complete cds	E-107
4869.seq	631	gi 4156188 gb AC004886.1 AC004886	apiens PAC clone DJ0784G16 from 7q11.23-q21.1, complete sequence	0.18
4870.seq	597	gi 5031750	apiens 3-hydroxy-3-methylglutaryl-Coenzyme A synthase 2	0
4871.seq	621	gi 4165210 emb AL021937 HS149A11	DNA sequence from clone 149A16 on chromosome 22q12-13. Contains	E-109
4872.seq	646	gi 4506656	apiens ribosomal protein L6 (RPL6) mRNA	0
4873.seq	288	gi 4156134 gb AC005091 AC005091	apiens BAC clone RG318C11 from 7p14-p15, complete sequence [Homo	0.31
4875.seq	210	gi 30031 emb X15879 HSCOL1N	mRNA for collagen VI alpha-1 N-terminal globular domain	E-112
4876.seq	438	gi 4503304	apiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	0
4877.seq	611	gi 4505028	apiens leukotriene A4 hydrolase (LTA4H) mRNA	0
4879.seq	691	gi 13000 emb X62996 MIHSGENOM	ris mitochondrial genome (consensus sequence)	0
4880.seq	541	gi 2992475 gb AC002549 AC002549	apiens Xp22 BAC GS-377014 (Genome Systems Human BAC library)	0.59
4881.seq	678	gi 4757825	apiens beta-2-microglobulin (B2M) mRNA	0
4882.seq	659	gi 4464234 emb AL009178.4 HS431P	DNA sequence from clone 431P23 on chromosome 6q27. Contains the	0.73
4883.seq	539	gi 179994 gb L00354 HUMCCK03	cholecystokinin (CCK) gene, exon 3.	0
4885.seq	411	gi 5830437 emb Y14486.1 HSY14486	apiens mRNA for cytosolic serine hydroxymethyltransferase,	0
4886.seq	593	gi 5032200	apiens tetraspan 3 (TSPAN-3) mRNA	0
4887.seq	485	gi 3367505 gb AC005369 AC005369	apiens chromosome 5, BAC clone 119j3 (LBNL H175), complete	0.53
4888.seq	630	gi 4507668	apiens tumor protein, translationally-controlled 1 (TPT1)	0
4890.seq	208	gi 5689414 dbj AB028962.1 AB02896	apiens mRNA for KIAA1039 protein, partial cds	3.00E-65
4891.seq	364	gi 5107001 gb AF125103.1 AF125103	apiens neuroendocrine specific protein c homolog mRNA, complete	0
4892.seq	699	gi 1944628 gb J01415.1 HUMMTCG	mitochondrion, complete genome	0
4893.seq	458	gi 4678254 emb AJ228697.1 AJL228	zomonon flos-aquae strain NIVA-CYA 142 trnL(UAA) gene, group	2
4894.seq	118			
4895.seq	59	gi 3004574 gb AC004519 AC004519	apiens BAC clone RG041H04 from 7q21-q22, complete sequence [Homo	0.18
4896.seq	425	gi 4759237	apiens transmembrane 4 superfamily member 3 (TM4SF3) mRNA	0
4897.seq	584	gi 3873461 emb AL031278 HS154A11	DNA sequence from clone 154A10 on chromosome 1q24 Contains EST	0.64
4899.seq	437	gi 6015527 emb AL034549.16 HS112	DNA sequence from clone 1125A11 on chromosome 20q11.1-11.23.	0.12
4900.seq	351	gi 5803087	apiens mitogen-activated protein kinase kinase 4	E-175
4902.seq	475	gi 4519622 dbj AB017616.1 AB01761	sculus YGR163w mRNA homolog, complete cds	0
4903.seq	345	gi 4501906	apiens a disintegrin and metalloproteinase domain 15	2.00E-36
4904.seq	241	gi 337297 gb M10546 HUMMTM1	mitochondrial DNA, fragment M1, encoding transfer RNAs,	E-126
4906.seq	290	gi 4557798	apiens Notch (Drosophila) homolog 3 (NOTCH3) mRNA	E-143
4907.seq	434	gi 6137796	mitochondrion, complete genome	0
4909.seq	400	gi 2351064 dbj AB006699.1 AB00669	opsis thalana genomic DNA, chromosome 5, P1 clone: MDJ22,	0.028
4910.seq	197	gi 5031930	apiens nascent polypeptide-associated complex alpha	7.00E-97
4911.seq	399	gi 5002627 emb AL034373.13 HS424	DNA sequence from clone 424L16 on chromosome 6q12-13. Contains	0.11
4912.seq	503	gi 5031748	apiens high-mobility group (nonhistone chromosomal) protein	0
4914.seq	435	gi 465173 gb L29157 HUMIGLZH	Immunoglobulin light chain variable region (lambda IIIB)	E-127
4915.seq	546	gi 4566745 gb AF115384.1 AF115384	apiens LR8 (LR8) mRNA, complete cds	0
4916.seq	580	gi 4502804	apiens chromogranin A (parathyroid secretory protein 1) (CHGA)	0
4917.seq	566	gi 179349 gb M33521 HUMBAT3B2	HLA-B-associated transcript 3 (BAT3) gene, 3' end.	0
4919.seq	472	gi 4806281	apiens bone morphogenetic protein 6 (BMP6) mRNA	0
4920.seq	465	gi 4506668	apiens ribosomal protein, large, P1 (RPLP1) mRNA	0
4922.seq	548	gi 262368 gb S51768 S51768	Pancreatitis-associated protein (human, pancreas, mRNA, 797	0
4923.seq	389	gi 1067154 emb Z67979 NPMSOD1	baginifolia soda1 gene promoter	0.002
4924.seq	623	gi 4753285 gb AC006453.3 AC00645	apiens clone DJ0614C10, complete sequence	0
4925.seq	439	gi 3482904 gb AC005551 AC005551	apiens chromosome 19, cosmid R26529, complete sequence [Homo	4.00E-57
4926.seq	378	gi 5102744 emb AL079283.1 HST000	apiens mRNA full length insert cDNA clone EUROIMAGE 39515	0
4928.seq	437	gi 4884091 emb AL050022.1 HSM80X	apiens mRNA: cDNA DKFzP564D116 (from clone DKFzP564D116)	0
4931.seq	281	gi 4827324 gb AC006975.2 AC00697	apiens PAC clone DJ1026B21 from X, complete sequence	0.001
4932.seq	220	gi 1944628 gb J01415.1 HUMMTCG	mitochondrion, complete genome	E-120
4933.seq	448	gi 2463264 emb AJ001616 MMMYEL	usculus mRNA for myeloid associated differentiation protein	5.00E-04
4934.seq	354	gi 30044 emb X15882 HSCOL2C1	mRNA for collagen VI alpha-2 C-terminal globular domain	E-153
4935.seq	608	gi 1370122 emb Z73965 HSV857G6	DNA sequence from cosmid V857G6, between markers DXS366 and D	0.17
4936.seq	404	gi 6174873 dbj AP000557.2 AP00055	apiens genomic DNA, chromosome 22q11.2, BCRL2 sequence	1.7
4937.seq	290	gi 4506700	apiens ribosomal protein S23 (RPS23) mRNA	E-123
4938.seq	527	gi 4503030	apiens CASP2 and RIPK1 domain containing adaptor with death	3.00E-98
4939.seq	532	gi 4099175 gb U84616 DRU84616	erio E74-like factor-2 (ZEF-2) mRNA, complete cds	0.15
4940.seq	271	gi 5102746 emb AL079286.1 HST000	apiens mRNA full length insert cDNA clone EUROIMAGE 26173	E-123
4941.seq	171	gi 1663703 dbj D87684 D87684	mRNA for KIAA0242 gene, partial cds	1.00E-60
4942.seq	531	gi 4506834	apiens small inducible cytokine subfamily A (Cys-Cys), member	0
4944.seq	51	gi 3978243 gb AF070674.1 AF07067	apiens inhibitor of apoptosis protein-1 (MIHC) mRNA, complete	1.00E-17
4945.seq	482	gi 3165402 gb AC004752 AC004752	apiens chromosome 5, BAC clone 8e5 (LBNL H167), complete sequenc	0
4946.seq	188	gi 190980 gb J05412 HUMREGB	regenerating protein (reg) gene, complete cds.	E-100
4947.seq	373	gi 6013154 gb AF129290.1 AF12929	apiens p35srj gene, alternatively spliced product and complete	E-124
4948.seq	599			
4950.seq	577	gi 4454695 gb AF070660 AF070660	apiens HSPC004 mRNA, complete cds	0
4951.seq	588	gi 4885084	apiens ATPase, H+ transporting, lysosomal (vacuolar proton	0
4953.seq	361	gi 3046306 gb AC004549 AC004549	apiens BAC clone RG459N13 from 7p15, complete sequence [Homo	7.00E-89
4954.seq	642	gi 532596 gb M12759 HUMIGJ02	1 ig J chain gene, exons 3 and 4.	E-174
4955.seq	485	gi 31091 emb X16869 HSEF1AC	mRNA for elongation factor 1-alpha (clone CEF4)	0
4956.seq	518	gi 2734091 gb AC003965 AC003965	apiens chromosome 16, cosmid clone 325D7, complete sequence	0.57
4957.seq	540	gi 4506380	apiens ras-related C3 botulinum toxin substrate 2 (rho	0
4958.seq	466	gi 1663703 dbj D87684 D87684	mRNA for KIAA0242 gene, partial cds	0
4959.seq	591	gi 5174602	apiens myosin regulatory light chain 2, smooth muscle Isoform	0
4960.seq	145	gi 4503304	apiens defensin, alpha 6, Paneth cell-specific (DEFA6) mRNA	2.00E-50
4961.seq	392	gi 2420 emb X07071 JANTRPC	illus niger trpC gene	1.7
4962.seq	572	gi 2104552 gb AF001548 HUAF0015	1 Chromosome 16 BAC clone CIT987SK-A-815A9, complete sequence [0
4963.seq	570	gi 3935222 gb AC005899 AC005899	apiens chromosome 17, clone hRPK.227.G.15, complete sequence	0
4964.seq	300	gi 4757047 emb AL049544.4 HSDJ2	DNA sequence from clone 240B8 on chromosome 6p11.2-q12. Contain	0.32
4965.seq	566	gi 437811 emb AL021578.1 HS453C	DNA sequence from clone 453C12 on chromosome 20q12-13.12.	1.00E-76
4966.seq	562	gi 388031 gb L05092 HUMRIBPROC	apiens ribosomal protein L11 mRNA, complete cds	0
4967.seq	257	gi 30167 emb X52851 HSCPH70	cyclophilin gene for cyclophilin (EC 5.2.1.8)	E-142
4968.seq	281	gi 4582132 emb AL021707.2 HS5081	DNA sequence from clone 508115 ori chromosome 22q12-13 Contains	E-150
4969.seq	327	gi 6330689 dbj AB033054.1 AB03305	apiens mRNA for KIAA1228 protein, partial cds	E-179
4970.seq	519			

4971.seq	535	gi 2842785 gb AC004130 AC004130	Homo sapiens BAC clone Rf	om 7p15-p21, complete sequence [Homo	0.038
4973.seq	221	gi 44309957 gb AC006032 AC006032	Homo sapiens BAC clone Ni	rom Y, complete sequence [Homo	0.23
4974.seq	671	gi 4506062	Homo sapiens protein kinase	pendent, regulatory, type I,	0
4976.seq	687	gi 4507668	Homo sapiens tumor protein	nally-controlled 1 (TPT1)	0
4977.seq	595	gi 3264580 gb AC005189 AC005189	Homo sapiens PAC clone D,	om 7p12-p14, complete sequence [Homo	0.042
4978.seq	631	gi 4826813	Homo sapiens kinectin 1 (kir	tor) (KTN1) mRNA	0
4979.seq	667	gi 1215741 gb U49083 HSU49083	Human cell surface heparin	tein HIP mRNA, complete cds	0
4980.seq	291	gi 5262587 emb ALD80141.1 HSM80065	Homo sapiens mRNA; cDNA	IM183 (from clone DKFZp434M183)	5.00E-77
4981.seq	708	gi 182077 gb M88458 HUMELP1A	Human ELP-1 mRNA sequ	:NA >gi 1794183 gb U24488 HSU24488	0
4982.seq	571	gi 6005907	Homo sapiens tenascin XA (0
4983.seq	600				0
4984.seq	666	gi 4506742	Homo sapiens ribosomal prc	'S8) mRNA	0
4986.seq	656	gi 220073 dbj D10537 HUMSPM	Homo sapiens mRNA for ms	ral protein of myelin, complete	0.18
4987.seq	569	gi 5420148 emb AL031587.3 HS1039KS	Human DNA sequence from	KS on chromosome 22q12.3-13.2, com	0
4988.seq	504	gi 2988486 gb AF047825 AF047825	Homo sapiens PAC 50H2 in	locus, complete sequence [Homo	0.14
4990.seq	511	gi 6226959	Homo sapiens alpha-2-mac	A2M) mRNA	0
4991.seq	538	gi 4755130	Homo sapiens cholecystokin	RNA	0
4992.seq	504	gi 5725511 gb AF072097.1 AF072097	Homo sapiens beta-2 micro	ve, complete cds	0
4993.seq	162	gi 452238 dbj D26442 YSCPES4	Yeast gene for PES4 PAB-1	complete cds	0.16
4994.seq	511	gi 5532410 gb AF138300.1 AF138300	Homo sapiens decorin varia	, complete cds	0
4995.seq	511	gi 452447 gb U04241 HSU04241	Human homolog of Drosoph	r of split m9/m10 mRNA, complete	0
4996.seq	373	gi 5801655 gb AC008134.3 AC008134	Homo sapiens 3q26.2-27 B/	-263C16 (Roswell Park Cancer Institute	0.1
4998.seq	351	gi 177206 gb J02769 HUM4F2A	Human 4F2 antigen heavy c	u, complete cds	E-171
4999.seq	399	gi 2853318 gb AF009341 HSUBE6AP3	Homo sapiens E6-AP ubiq	ligase (UBE3A) exon 16 sequence	E-170
5000.seq	452	gi 4753261 gb AC004975.2 AC004975	Homo sapiens PAC clone D,	om 7q35-q36, complete sequence	4.00E-85
5001.seq	500	gi 6031211 gb U40992.2 HSU40992	Homo sapiens heat shock pr	omolog mRNA, complete cds	E-146
5002.seq	493	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exo		E-117
5003.seq	458	gi 4579990 dbj AP000069.1 AP000069	Homo sapiens genomic DNA	senescence gene, segment 5/19,	0.13
5005.seq	175				0
5006.seq	497	gi 5360674 dbj AB022654.1 AB022654	Homo sapiens mRNA for an	ta histolytica immunoglobulin	0
5007.seq	391	gi 5174898	Homo sapiens stimulated tr	factor (50 kDa) (STAF50) mRNA	E-158
5008.seq	492	gi 546798 gb S71043 S71043	Ig alpha 2-immunoglobulin J	in allotype 2 (constant	E-169
5009.seq	504	gi 2765422 emb Y14736 HSIGG1KL	Homo sapiens mRNA for Im	in kappa light chain	0
5010.seq	132	gi 4758137	Homo sapiens DEAD/H (Asp	Asp/His) box polypeptide 5 (RNA	4.00E-63
5012.seq	161	gi 5714428 gb AF115343.1 RNCSD1	Rattus norvegicus cytochr	o c decarboxylase (CSAD) gene,	0.64
5013.seq	434	gi 4894853 gb AF135157.1 AF135157	Homo sapiens cysteine t	precursor, mRNA, complete cds	0
5015.seq	501	gi 238235 gb S62403 S62403	transmembrane secretory c	human, colonic adenocarcinoma	0
5016.seq	375	gi 5031702	Homo sapiens Ras-GTPase	protein SH3-domain-binding	0
5017.seq	466	gi 4176417 dbj AB004304 AB004304	Homo sapiens mRNA for Ig	in, partial cds	E-154
5018.seq	501	gi 2588627 gb AC003080 AC003080	Human BAC clone GS36B1	.1, complete sequence [Homo sapiens]	0.14
5019.seq	433	gi 2656145 gb AF020503 AF020503	Homo sapiens FRA3B comm	region, diadenosine triphosphate	0
5020.seq	115	gi 5230675 gb AF159570.1 AF159570	Homo sapiens regulator of	gnalling 5 (RGS5) mRNA,	1.00E-16
5021.seq	369	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene, c	ds,	E-108
5023.seq	479	gi 5678818 gb AF146191.1 AF146191	Homo sapiens FRG1 (FRG1	plete cds; 5S ribosomal RNA gene,	3.00E-58
5024.seq	504	gi 4506470	Homo sapiens regenerati	on 1 alpha (pancreatic stone	0
5026.seq	518	gi 29628 emb X52956 HSCALM3	Human CAMII-ps1 calmodi	ropene	0
5027.seq	509	gi 3041843 gb AC004544 AC004544	Homo sapiens BAC clone Rf	om 4, complete sequence [Homo	0.14
5028.seq	501	gi 4768678 gb AF113887.1 AF113887	Homo sapiens kappa.1 imm	light chain mRNA, complete cds	E-180
5029.seq	518	gi 5107005 gb AF125466.1 AF125466	Homo sapiens nuclear distri	ctin C (NUDCP1) pseudogene sequence	0
5030.seq	48				0
5031.seq	498	gi 3894210 gb AC004934 AC004934	Homo sapiens PAC clone D,	om 7p12-p14, complete sequence [Homo	0.14
5032.seq	403	gi 2695566 gb AC002312 AC002312	Human Chromosome 15 pac	3.1, complete sequence [Homo sapiens]	0.007
5033.seq	398	gi 505472 emb X69181 HSL3	H.sapiens mRNA for ribosor	L3)	0
5034.seq	510	gi 441402 emb X72467 HSIGKLV46	H.sapiens mRNA for rearr	pa light chain variable region	0
5035.seq	349	gi 4165332 gb AC006422 AC006422	Drosophila melanogaster, ct	2R, region 50A2-50A3, P1 clones	0.024
5037.seq	510	gi 185363 gb M87790 HUMIGHEPAL	Human (hybridoma H210) a	s A Immunoglobulin lambda chain	0
5038.seq	200	gi 2832603 emb AL021367 HS443E24	Human DNA sequence from	24 on chromosome 6q16. Contains EST	0.053
5039.seq	498	gi 339840 gb M10036 HUMTP1	Human triosephosphate isoz	ym, complete cds.	0
5041.seq	494	gi 5080755 gb AC007842.1 AC007842	Homo sapiens chromosome	1191 (CIT-B-471f3), complete sequen	1.00E-66
5042.seq	303	gi 186618 gb M32219 HUMJNLTRA	Human DNA/endogenous ri	g terminal repeat (LTR) junction	5.00E-06
5043.seq	527	gi 4506680	Homo sapiens ribosomal prc	PS11) mRNA	0
5044.seq	498				0
5046.seq	19				0
5047.seq	279	gi 5912165 emb AL117597.1 HSM80113	Homo sapiens mRNA; cDNA	IG2463 (from clone DKFZp564G2463)	E-123
5048.seq	433	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, exo		0
5049.seq	483	gi 6330892 dbj AB033080.1 AB033080	Homo sapiens mRNA for Kl	stein, partial cds	0.13
5050.seq	503	gi 4506648	Homo sapiens ribosomal prc	'L3) mRNA	0
5051.seq	493	gi 2894518 emb J223812 HSA223812	Homo sapiens mRNA for ca	plete cds	0
5052.seq	504	gi 1665768 dbj D87437 D87437	Human mRNA for KIAA025	melanoma cell line, mRNA, 453	3.00E-42
5053.seq	116	gi 264772 gb U54005 S54005	thymosin beta-10 [human,]	:6584 (CIT-B-459F4), complete sequen	0.47
5054.seq	426	gi 4220442 gb AC006504 AC006504	Homo sapiens chromosome	family member 3 (TM4SF3) mRNA	0
5055.seq	511	gi 4759237	Homo sapiens transmembra	R28154, complete sequence [Homo	0.53
5056.seq	481	gi 3451032 gb AC005523 AC005523	Homo sapiens chromosome	(Ehlers-Danlos syndrome type	0
5057.seq	505	gi 4502950	Homo sapiens collagen, typ	I (L1-beta regulated HSOC1	E-103
5058.seq	496	gi 4838408 gb AF124819.1 AF124819	Homo sapiens T84 colon cat	h, testis, mRNA, 1784 nt)	2.00E-59
5059.seq	496	gi 250802 gb S39127 S39127	Homo sapiens cullin 2 (CUL	nd translated products	0
5060.seq	495	gi 4503162	Homo sapiens chromosome	:1307e8 (LBNL H60), complete	0.005
5061.seq	292	gi 3366557 gb AC005355 AC005355	Homo sapiens chromosome	315 on chromosome 11p13, complete	0.14
5062.seq	513	gi 5596763 emb AL079340.7 HSJ760G15	Human DNA sequence from	clone ZD39H03	3.00E-61
5063.seq	458	gi 3483595 gb AF086250 HUMZD39H03	Homo sapiens full length ins	: Clone p1093 in CES region, complete	1.8
5064.seq	423	gi 5705994 gb AC006946.18 AC006946	Homo sapiens Chromosome	3H0723 (from clone DKFZp586H0723)	E-135
5065.seq	289	gi 5262701 emb ALD80212.1 HSM800735	Homo sapiens mRNA; cDNA	ve receptor-associated protein 1	0
5066.seq	504	gi 4758955	Homo sapiens peripheral be	ne	0
5067.seq	459	gi 1944628 gb J01415.1 HUMMTCC	Human mitochondrion, com	4A, complete cds	3.00E-22
5068.seq	188	gi 6434899 gb AF133423.1 AF133423	Homo sapiens tetraspanin 1	C 55C20 on chromosome 6. Contains a	0.008
5069.seq	435	gi 3080468 emb AL021368 HS55C20	Homo sapiens DNA sequen	ancer cell line, MCF-7, mRNA,	0
5070.seq	501	gi 298485 gb S56985 S56985	ribosomal protein L19 [hum	3olgI intermediate compartment	E-103
5071.seq	257	gi 4894445 gb AF081885.1 HSERIGCP2	Homo sapiens endoplasmic	me exons 1-27, complete cds.	E-123
5072.seq	501	gi 292420 dbj L11910 HUMRETLAS	Human retinoblastoma susc	4C186 (from clone DKFZp564C186)	0
5073.seq	499	gi 4884087 emb AL050019.1 HSM800100	Homo sapiens mRNA; cDNA/	p53) mRNA, complete cds.	0
5074.seq	503	gi 555940 gb U14990 HSU14990	Human XP1PO ribosomal p	ngation factor 1 alpha 1	0
5075.seq	514	gi 4503470	Homo sapiens eukaryotic tr	41138 (from clone DKFZp4341138)	E-154
5076.seq	501	gi 6102948 emb AL122121.1 HSM80128	Homo sapiens mRNA; cDNA/	ferase mRNA, exons 3-11, complete	E-135
5077.seq	498	gi 189347 gb M23204 HUMOATC	Human ornithine-oxo-acid	ts,	E-102
5078.seq	472	gi 181532 gb M97925 HUMDEF5A	H.sapiens defensin 5 gene,	rNA >gi 632963 gb L29219 HUMCLK1	E-156
5079.seq	499	gi 4758007	Homo sapiens CDC-like kin	A16 on chromosome 6p24.1-25.3,	0
5080.seq	500	gi 5679748 emb AL022725.8 HS430A16	Human DNA sequence from) gene locus	0.14
5082.seq	500	gi 467918 gb U03115 HSTCRBV	Human V beta T-cell recep	ge dependent K+ channel (KvLQT1)	0
5083.seq	499	gi 2465530 gb AF000571 AF000571	Homo sapiens kidney and c,	'serine-rich 1 (splicing	E-150
5089.seq	472	gi 5902075	Homo sapiens splicing fact	rPK.640_L15, complete sequence [Homo	0.035
5090.seq	499	gi 3365852 gb AC005324 AC005324	Homo sapiens chromosome	tial cds	1.00E-20
5091.seq	497	gi 2224546 dbj AB002301 AB002301	Human mRNA for KIAA03C	BR1) mRNA	0
5092.seq	508	gi 4502598	Homo sapiens carbonyl red		0
5094.seq	243				0
5095.seq	506	gi 3123607 emb Y17214 HSY17214	Homo sapiens intergenic Di	1 SURF-2 and SURF-4	0
5097.seq	439	gi 5802965	Homo sapiens destrin (actr	izing factor) (ADF), mRNA	3.00E-98
5098.seq	458	gi 532596 gb M12759 HUMIGJ02	Human Ig J chain gene, ex		E-146
5100.seq	502	gi 3366671 gb AF039594 HSDHCG	Homo sapiens succinate de	e subunit C (SDHC) gene, exon 6	E-140