

Table 9. Probesets (545) representing 488 transcripts identified by SAM that are enriched ≥ 2 -fold in poor-prognosis NE tumors compared to good-prognosis NE tumors

| Probeset ID | GenBank/TIGR ID | Gene Name | Gene Symbol | LocusLink ID | SAM Score | SAM Fold Change |
|-------------|-----------------|--|--------------|--------------|-----------|-----------------|
| 37625_at | U52682 | interferon regulatory factor 4 | IRF4 | 3662 | 5.865 | 196.2 |
| 38850_at | M11119 | Human endogenous retrovirus envelope region mRNA (PL1) | --- | --- | 4.910 | 100.0 |
| 41165_g_at | X67301 | immunoglobulin heavy constant mu | IGHM | 3507 | 5.534 | 83.8 |
| 36105_at | M18728 | carcinoembryonic antigen-related cell adhesion molecule 6 (non-specific cross reacting antigen) | CEACAM6 | 4680 | 9.560 | 53.5 |
| 36299_at | X02330 | calcitonin/calcitonin-related polypeptide, alpha | CALCA | 796 | 6.160 | 48.9 |
| 35822_at | L15702 | B-factor, properdin | BF | 629 | 3.022 | 45.5 |
| 32821_at | A1762213 | lipocalin 2 (oncogene 24p3) | LN2 | 3634 | 2.456 | 44.0 |
| 1490_at | M19720 | v-myc myelocytomatosis viral oncogene homolog 1, lung carcinoma derived (avian) | MYC1 | 4610 | 3.036 | 35.5 |
| 41164_at | X67301 | immunoglobulin heavy constant mu | IGHM | 3507 | 5.856 | 30.0 |
| 1582_at | M29540 | carcinoembryonic antigen-related cell adhesion molecule 5 | CEACAM5 | 1048 | 8.310 | 26.5 |
| 31579_at | AF005082 | skin-specific protein | xp33 | 388699 | 6.447 | 25.0 |
| 39842_at | AF059293 | cytokine receptor-like factor 1 | CRLF1 | 9244 | 1.848 | 23.2 |
| 37865_at | J05081 | endothelin 3 | EDN3 | 1908 | 3.659 | 22.5 |
| 34566_at | X02404 | calcitonin-related polypeptide, beta | CALCB | 797 | 3.945 | 21.7 |
| 40035_at | AB012917 | kallikrein 11 | KLK11 | 11012 | 7.269 | 21.5 |
| 37186_s_at | U11863 | amiloride binding protein 1 (amine oxidase (copper-containing)) | ABP1 | 26 | 4.669 | 21.1 |
| 37533_r_at | D86980 | tetratricopeptide repeat domain 9 | TTC9 | 23508 | 4.217 | 18.0 |
| 40398_s_at | A1743406 | mesenchyme homeo box 2 (growth arrest-specific homeo box) | MEOX2 | 4223 | 3.348 | 15.6 |
| 35832_at | AB029000 | sulfatase 1 | SULF1 | 23213 | 7.824 | 15.3 |
| 38582_at | A1961220 | serine protease inhibitor, Kazal type 1 | SPINK1 | 6690 | 3.837 | 15.1 |
| 33714_at | Y10043 | high-mobility group box 3 | HMG3 | 3149 | 5.705 | 14.9 |
| 31536_f_at | X72475 | immunoglobulin kappa constant | IGKC | 3514 | 6.682 | 14.7 |
| 38524_at | U48194 | occludin | OCLN | 4950 | 5.288 | 14.2 |
| 33383_f_at | A1820718 | surfactant, pulmonary-associated protein B | SFTPB | 6439 | 5.290 | 13.8 |
| 31859_at | J05070 | matrix metalloproteinase 9 (gelatinase B, 92kDa gelatinase, 92kDa type IV collagenase) | MMP9 | 4318 | 6.573 | 13.8 |
| 36503_at | AB002409 | chemokine (C-C motif) ligand 21 | CCL21 | 6366 | 2.566 | 13.6 |
| 31315_at | D84143 | Immunoglobulin light chain lambda variable region [Homo sapiens], mRNA sequence | --- | --- | 6.660 | 13.6 |
| 37322_s_at | X82460 | hydroxyprostaglandin dehydrogenase 15-(NAD) | HPGD | 3248 | 3.699 | 13.2 |
| 37637_at | U27655 | regulator of G-protein signalling 3 | RGS3 | 5998 | 6.022 | 13.0 |
| 41294_at | AJ238246 | keratin 7 | KRT7 | 3855 | 8.151 | 12.9 |
| 1069_at | U04636 | prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase) | PTGS2 | 5743 | 5.001 | 12.9 |
| 34105_f_at | A1147237 | immunoglobulin heavy constant gamma 1 (G1m marker) | IGHG1 | 3500 | 6.409 | 12.9 |
| 36300_at | X15943 | calcitonin/calcitonin-related polypeptide, alpha | CALCA | 796 | 5.813 | 12.7 |
| 31319_at | M20707 | Clone 2-12 immunoglobulin light chain mRNA, partial cds | --- | --- | 5.100 | 11.8 |
| 39945_at | U09278 | fibroblast activation protein, alpha | FAP | 2191 | 6.957 | 11.6 |
| 931_at | L08177 | Epstein-Barr virus induced gene 2 (lymphocyte-specific G protein-coupled receptor) | EBI2 | 1880 | 4.401 | 11.4 |
| 37006_at | A1660656 | immunoglobulin J polypeptide, linker protein for immunoglobulin alpha and mu polypeptides | IGJ | 3512 | 6.584 | 11.3 |
| 32305_at | K01079 | collagen, type I, alpha 2 | COL1A2 | 1278 | 5.987 | 11.1 |
| 41416_at | D14446 | fibronectin-like 1 | FN1 | 2267 | 7.794 | 11.0 |
| 40041_at | AF017790 | kinetochore associated 2 | KNTC2 | 10403 | 4.826 | 11.0 |
| 34095_f_at | U80114 | immunoglobulin heavy locus | IGH@ | 3492 | 7.063 | 10.8 |
| 37971_at | AL050089 | bromodomain adjacent to zinc finger domain, 1A | BAZ1A | 11177 | 3.048 | 10.3 |
| 41827_f_at | A1932613 | similar to bK246H3.1 (immunoglobulin lambda-like polypeptide 1, pre-B-cell specific) | LOC91316 | 91316 | 7.780 | 10.1 |
| 33070_at | AF041243 | hypocretin (orexin) receptor 1 | HCRTR1 | 3061 | 6.951 | 10.0 |
| 34193_at | AF002246 | cell adhesion molecule with homology to L1CAM (close homolog of L1) | CHL1 | 10752 | 4.914 | 9.8 |
| 35566_f_at | AF015128 | immunoglobulin heavy constant gamma 1 (G1m marker) | IGHG1 | 3500 | 6.347 | 9.7 |
| 40328_at | X99268 | twist homolog 1 (acrocephalosyndactyly 3; Saethre-Chotzen syndrome) (Drosophila) | TWIST1 | 7291 | 3.742 | 9.6 |
| 34094_i_at | U80114 | immunoglobulin heavy constant gamma 1 (G1m marker) | IGHG1 | 3500 | 6.802 | 9.6 |
| 37864_s_at | AJ001564 | immunoglobulin heavy constant gamma 4 (G4m marker) | IGHG4 | 3503 | 7.975 | 9.5 |
| 37422_at | S71018 | peptidylprolyl isomerase C (cyclophilin C) | PPIC | 5480 | 6.709 | 9.5 |
| 39515_s_at | W27099 | processing of precursor 5, ribonuclease P/MRP subunit (S. cerevisiae) | POP5 | 51367 | 5.919 | 9.5 |
| 37425_g_at | AB029343 | chromosome 6 open reading frame 18 | C6orf18 | 54535 | 3.781 | 9.3 |
| 40385_at | U64197 | chemokine (C-C motif) ligand 20 | CCL20 | 6364 | 3.129 | 9.2 |
| 33274_f_at | X79783 | immunoglobulin lambda joining 3 | IGLJ3 | 28831 | 7.967 | 9.1 |
| 36879_at | M63193 | endothelial cell growth factor 1 (platelet-derived) | ECGF1 | 1890 | 7.018 | 9.0 |
| 38087_s_at | W72186 | S100 calcium binding protein A4 (calcium protein, calvasculin, metastasin, murine placental homolog) | S100A4 | 6175 | 5.844 | 8.7 |
| 370129_at | J030129 | fibronectin, B beta polypeptide | FN2 | 5244 | 3.244 | 8.5 |
| 33273_f_at | J03011 | immunoglobulin lambda joining 3 | IGLJ3 | 28831 | 7.973 | 8.5 |
| 31888_s_at | AF001294 | pleckstrin homology-like domain, family A, member 2 | PHLDA2 | 7262 | 5.986 | 8.3 |
| 37497_at | L16499 | hematopoietically expressed homeobox | HHEX | 3087 | 3.601 | 8.2 |
| 36239_at | Z49194 | POU domain, class 2, associating factor 1 | POU2AF1 | 5450 | 5.406 | 8.1 |
| 1197_at | D00654 | actin, gamma 2, smooth muscle, enteric | ACTG2 | 72 | 2.908 | 8.0 |
| 32368_at | AF061573 | protocadherin 8 | PCDH8 | 5100 | 2.754 | 7.9 |
| 1147_at | HG3510-HT3704 | V-Erba Related Ear-3 Protein | --- | --- | 2.078 | 7.7 |
| 658_at | L12350 | thrombospondin 2 | THBS2 | 7058 | 5.997 | 7.6 |
| 37231_at | D13633 | discs, large homolog 7 (Drosophila) | DLG7 | 9787 | 4.262 | 7.6 |
| 33500_i_at | S71043 | hypothetical protein MGC27165 | MGC27165 | 283650 | 5.764 | 7.5 |
| 41109_at | M31452 | complement component 4 binding protein, alpha | C4BPA | 722 | 2.515 | 7.4 |
| 36070_at | AL049389 | KIAA1199 protein | KIAA1199 | 57214 | 6.626 | 7.3 |
| 34342_s_at | AF052124 | secreted phosphoprotein 1 (osteopontin, bone sialoprotein I, early T-lymphocyte activation 1) | SPP1 | 6696 | 6.045 | 7.3 |
| 33904_at | AB000714 | claudin 3 | CLDN3 | 1365 | 6.347 | 7.2 |
| 14105_l_at | M21121 | chemokine (C-C motif) ligand 5 | CCL5 | 6362 | 5.469 | 7.2 |
| 1482_g_at | L23808 | matrix metalloproteinase 12 (macrophage elastase) | MMP12 | 4321 | 6.015 | 7.2 |
| 38194_s_at | M63438 | immunoglobulin kappa variable 3D-15 | IGKV3D-15 | 28875 | 8.366 | 7.2 |
| 34033_s_at | U82278 | leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2 | LILRA2 | 11027 | 2.349 | 7.1 |
| 2092_s_at | J04765 | secreted phosphoprotein 1 (osteopontin, bone sialoprotein I, early T-lymphocyte activation 1) | SPP1 | 6696 | 5.284 | 7.1 |
| 33143_s_at | U81800 | solute carrier family 16 (monocarboxylic acid transporters), member 3 | SLC16A3 | 9123 | 5.980 | 6.9 |
| 37892_at | J04177 | collagen, type XI, alpha 1 | COL11A1 | 1301 | 2.721 | 6.9 |
| 33499_s_at | AF067420 | hypothetical protein MGC27165 | MGC27165 | 283650 | 5.838 | 6.8 |
| 1451_s_at | D13666 | periostin, osteoblast specific factor | POSTN | 10631 | 7.646 | 6.8 |
| 38583_at | Y15014 | UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 2 | B3GALT2 | 8707 | 3.443 | 6.8 |
| 38469_at | M35252 | transmembrane 4 superfamily member 3 | TM4SF3 | 7103 | 4.817 | 6.7 |
| 35902_at | AJ132820 | a disintegrin and metalloproteinase domain 5 | ADAM5 | 255926 | 6.164 | 6.6 |
| 32306_g_at | K01079 | collagen, type I, alpha 2 | COL1A2 | 1278 | 7.993 | 6.6 |
| 38324_at | AD000684 | liver-specific bHLH-Zip transcription factor | LISCH7 | 51599 | 5.853 | 6.6 |
| 33501_r_at | S71043 | hypothetical protein MGC27165 | MGC27165 | 283650 | 5.775 | 6.5 |
| 1481_at | L23808 | matrix metalloproteinase 12 (macrophage elastase) | MMP12 | 4321 | 5.748 | 6.4 |
| 40456_at | AL049863 | solute carrier family 39 (zinc transporter), member 8 | SLC39A8 | 64116 | 6.519 | 6.4 |
| 41470_at | AF027208 | prominin 1 | PROM1 | 8842 | 4.972 | 6.4 |
| 35735_at | M65542 | guanylate binding protein 1, interferon-inducible, 67kDa | GBP1 | 2633 | 7.454 | 6.3 |
| 40145_at | A1375913 | topoisomerase (DNA) II alpha 170kDa | TOP2A | 7153 | 5.574 | 6.3 |
| 36861_at | AL049946 | adican | DKFZp5641922 | 25878 | 4.551 | 6.3 |
| 859_at | U03688 | cytochrome P450, family 1, subfamily B, polypeptide 1 | CYP1B1 | 1545 | 8.108 | 6.2 |
| 38111_at | X15998 | chondroitin sulfate proteoglycan 2 (versican) | CSPG2 | 1462 | 7.162 | 6.1 |
| 40071_at | U03688 | cytochrome P450, family 1, subfamily B, polypeptide 1 | CYP1B1 | 1545 | 8.171 | 6.1 |
| 35927_r_at | AF090005 | leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 1 | LILRB1 | 10859 | 6.401 | 6.1 |
| 703_at | HG3731-HT4001 | Immunoglobulin Heavy Chain, Vdjrj Regions | --- | --- | 6.060 | 6.1 |
| 31720_s_at | M10905 | fibronectin 1 | FN1 | 2335 | 9.146 | 6.0 |
| 33754_at | U43203 | thyroid transcription factor 1 | TTF1 | 7080 | 8.314 | 6.0 |
| 32488_at | X06700 | collagen, type III, alpha 1 (Ehlers-Danlos syndrome type IV, autosomal dominant) | COL3A1 | 1281 | 8.360 | 6.0 |
| 904_s_at | L47276 | topoisomerase (DNA) II alpha 170kDa | TOP2A | 7153 | 5.493 | 5.8 |
| 33133_at | U80184 | flightless I homolog (Drosophila) | FLII | 2314 | 6.337 | 5.6 |
| 40519_at | Y00638 | protein tyrosine phosphatase, receptor type, C | PTPRC | 5788 | 5.588 | 5.6 |
| 38077_at | X52022 | collagen, type VI, alpha 3 | COL6A3 | 1293 | 7.459 | 5.4 |
| 32239_at | U69263 | matrilin 2 | MATN2 | 4147 | 3.038 | 5.4 |
| 31460_f_at | AF043586 | immunoglobulin lambda joining 3 | IGLJ3 | 28831 | 4.754 | 5.4 |
| 35590_s_at | X81832 | gastric inhibitory polypeptide receptor | GIPR | 2696 | 6.659 | 5.4 |
| 34319_at | AA131149 | S100 calcium binding protein P | S100P | 6286 | 3.251 | 5.4 |
| 36133_at | AL031058 | desmoplakin | DSP | 1832 | 7.345 | 5.4 |
| 38299_at | X04430 | interleukin 6 (interferon, beta 2) | IL6 | 3569 | 4.476 | 5.3 |
| 37141_at | U39840 | forkhead box A1 | FOXA1 | 3169 | 7.060 | 5.2 |

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|------------|---------------|--|----------|--------|--------|-----|
| 41239_r_at | M90696 | cathepsin S | CTSS | 1520 | 4.702 | 5.1 |
| 35372_r_at | M28130 | interleukin 8 | IL8 | 3576 | 4.304 | 5.1 |
| 31719_at | X02761 | fibronectin 1 | FN1 | 2335 | 9.607 | 5.1 |
| 38017_at | U05259 | CD79A antigen (immunoglobulin-associated alpha) | CD79A | 973 | 5.321 | 5.1 |
| 33998_at | U91618 | neurotensin | NTS | 4922 | 2.565 | 5.0 |
| 40518_at | Y00062 | protein tyrosine phosphatase, receptor type, C | PTPRC | 5788 | 5.448 | 5.0 |
| 35926_s_at | AF009005 | leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 1 | LILRB1 | 10859 | 7.143 | 5.0 |
| 291_s_at | J04152 | tumor-associated calcium signal transducer 2 | TACSTD2 | 4070 | 4.585 | 4.9 |
| 36500_at | AF030186 | glypican 4 | GPC4 | 2239 | 4.058 | 4.9 |
| 38976_at | D44497 | coronin, actin binding protein, 1A | CORO1A | 11151 | 6.920 | 4.9 |
| 38112_g_at | X15998 | chondroitin sulfate proteoglycan 2 (versican) | CSPG2 | 1462 | 6.816 | 4.8 |
| 38853_at | X81892 | G protein-coupled receptor 64 | GPR64 | 10149 | 3.618 | 4.8 |
| 32307_s_at | V00503 | collagen, type I, alpha 2 | COL1A2 | 1278 | 8.262 | 4.8 |
| 32818_at | X78565 | tenascin C (hexabrachion) | TNC | 3371 | 4.270 | 4.8 |
| 32570_at | L76465 | hydroxyprostaglandin dehydrogenase 15-(NAD) | HPGD | 3248 | 4.105 | 4.7 |
| 41048_at | D90070 | phorbol-12-myristate-13-acetate-induced protein 1 | PMAIP1 | 5366 | 4.786 | 4.7 |
| 33641_g_at | Y14768 | allograft inflammatory factor 1 | AIF1 | 199 | 4.144 | 4.7 |
| 38827_at | AF038451 | anterior gradient 2 homolog (Xenopus laevis) | AGR2 | 10551 | 4.021 | 4.7 |
| 311_s_at | HG3044-HT3742 | Fibronectin, Alt. Splice 1 | --- | --- | 8.613 | 4.7 |
| 37215_at | AF046798 | phosphorylase, glycogen; liver (Hers disease, glycogen storage disease type VI) | PYGL | 5836 | 3.491 | 4.6 |
| 37099_at | A1806222 | arachidonate 5-lipoxygenase-activating protein | ALOX5AP | 241 | 4.714 | 4.6 |
| 548_s_at | S80267 | spleen tyrosine kinase | SYK | 6850 | 9.851 | 4.6 |
| 37343_at | U01062 | inositol 1,4,5-triphosphate receptor, type 3 | ITPR3 | 3710 | 6.462 | 4.6 |
| 37187_at | M36820 | chemokine (C-X-C motif) ligand 2 | CXCL2 | 2920 | 4.670 | 4.5 |
| 40079_at | AA156240 | retinoic acid induced 3 | RAI3 | 9052 | 4.522 | 4.5 |
| 40544_g_at | L08424 | achaete-scute complex-like 1 (Drosophila) | ASCL1 | 429 | 5.876 | 4.5 |
| 645_at | L26336 | heat shock 70kDa protein 2 | HSPA2 | 3306 | 2.928 | 4.5 |
| 1343_s_at | S66896 | serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 3 | SERPINF3 | 6317 | 1.793 | 4.5 |
| 35530_f_at | X92997 | Immunoglobulin lambda light chain variable and constant region mRNA, partial cds | --- | --- | 6.461 | 4.4 |
| 31811_r_at | L11667 | peptidylprolyl isomerase D (cyclophilin D) | PPID | 5481 | 1.670 | 4.4 |
| 40757_at | M18737 | granzyme A (granzyme 1, cytotoxic T-lymphocyte-associated serine esterase 3) | GZMA | 3001 | 6.765 | 4.4 |
| 128_at | X82153 | cathepsin K (pseudosostosis) | CTSK | 1513 | 4.461 | 4.4 |
| 32715_at | AF053233 | vesicle-associated membrane protein 8 (endobrevin) | VAMP8 | 8673 | 10.338 | 4.3 |
| 32916_at | X54134 | protein tyrosine phosphatase, receptor type, E | PTPRE | 5791 | 4.631 | 4.3 |
| 41069_at | AB006000 | leukocyte cell derived chemotaxin 1 | LECT1 | 11061 | 2.648 | 4.3 |
| 33933_at | X63187 | WAP four-disulfide core domain 2 | WFDC2 | 10406 | 4.094 | 4.3 |
| 37168_at | AB013924 | lysosomal-associated membrane protein 3 | LAMP3 | 27074 | 4.625 | 4.3 |
| 37714_at | M25667 | growth associated protein 43 | GAP43 | 2596 | 2.904 | 4.2 |
| 37899_at | X02308 | thymidylate synthetase | TYMS | 7298 | 5.547 | 4.2 |
| 38116_at | D14657 | KIAA0101 gene product | KIAA0101 | 9768 | 7.244 | 4.2 |
| 649_s_at | L06797 | chemokine (C-X-C motif) receptor 4 | CXCR4 | 7852 | 6.262 | 4.1 |
| 39260_at | U91485 | solute carrier family 16 (monocarboxylic acid transporters), member 4 | SLC16A4 | 9122 | 3.586 | 4.1 |
| 195_s_at | U28014 | casease 4, apoptosis-related cysteine protease | CASP4 | 837 | 6.842 | 4.1 |
| 34245_at | W25917 | chromosome 6 open reading frame 145 | C6orf145 | 221749 | 8.435 | 4.1 |
| 700_s_at | HG371-HT26388 | Mucin 1, Epithelial, Alt. Splice 9 | --- | --- | 4.033 | 4.1 |
| 38420_at | Y14690 | collagen, type V, alpha 2 | COL5A2 | 1290 | 5.198 | 4.1 |
| 37552_at | U33632 | potassium channel, subfamily K, member 1 | KCNK1 | 3775 | 4.318 | 4.1 |
| 35249_at | AF102778 | cyclin E2 | CNE2 | 9134 | 3.731 | 4.1 |
| 41592_at | AB005043 | suppressor of cytokine signaling 1 | SOCS1 | 8651 | 4.312 | 4.0 |
| 1385_at | M77349 | transforming growth factor, beta-induced, 68kDa | TGFB1 | 7045 | 5.541 | 4.0 |
| 38466_at | X82153 | cathepsin K (pseudosostosis) | CTSK | 1513 | 5.936 | 4.0 |
| 39605_at | X74142 | forkhead box G1B | FOXP1 | 2290 | 3.067 | 4.0 |
| 35766_at | X12883 | keratin 18 | KRT18 | 3875 | 7.363 | 4.0 |
| 41819_at | U93049 | FYN binding protein (FYB-120/130) | FYB | 2533 | 5.962 | 4.0 |
| 38391_at | M94345 | capping protein (actin filament), gelsolin-like | CAPG | 822 | 7.577 | 3.9 |
| 36533_at | D83402 | prostaglandin I2 (prostaglyclin) synthase | PTGIS | 5740 | 5.054 | 3.9 |
| 38379_at | X76534 | glycoprotein (transmembrane) nmb | GNPMB | 10457 | 5.259 | 3.9 |
| 31798_at | AA314625 | trefoil factor 1 (breast cancer, estrogen-inducible sequence expressed in) | TFF1 | 7031 | 3.064 | 3.9 |
| 41036_at | AB011421 | neutrophil cytosolic factor 2 (65kDa, chronic granulomatous disease, autosomal 2) | NCF2 | 4688 | 4.703 | 3.9 |
| 37524_at | U28014 | serine/threonine kinase 17b (apoptosis-inducing) | STK17B | 9262 | 6.877 | 3.9 |
| 37897_s_at | A1985964 | trefoil factor 3 (intestinal) | TFF3 | 7033 | 5.750 | 3.8 |
| 41433_at | M73255 | vascular cell adhesion molecule 1 | VCAM1 | 7412 | 4.239 | 3.8 |
| 31477_at | L08044 | trefoil factor 3 (intestinal) | TFF3 | 7033 | 6.322 | 3.8 |
| 37759_at | U51240 | Lysosomal-associated multispansing membrane protein-5 | LAPTM5 | 7805 | 7.051 | 3.8 |
| 38578_at | M63928 | tumor necrosis factor receptor superfamily, member 7 | TNFRSF7 | 939 | 5.375 | 3.8 |
| 1404_r_at | M21121 | chemokine (C-C motif) ligand 5 | CCL5 | 6352 | 2.387 | 3.8 |
| 39421_at | D43969 | runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene) | RUNX1 | 861 | 8.487 | 3.8 |
| 1123_at | L01406 | growth hormone releasing hormone receptor | GHRHR | 2692 | 6.804 | 3.8 |
| 35362_at | AB018342 | myosin X | MYO10 | 4651 | 4.180 | 3.8 |
| 36108_at | M16276 | major histocompatibility complex, class II, DQ beta 1 | HLA-DQB1 | 3119 | 4.087 | 3.7 |
| 38404_at | M55153 | transglutaminase 2 (C polypeptide, protein-glutamine-gamma-glutamyltransferase) | TGM2 | 7052 | 6.441 | 3.7 |
| 774_g_at | D10667 | myosin, heavy polypeptide 11, smooth muscle | MYH11 | 4629 | 2.366 | 3.7 |
| 36889_at | M33195 | Fc fragment of IgE, high affinity I, receptor for; gamma polypeptide | FCER1G | 2207 | 7.332 | 3.7 |
| 38784_g_at | J05581 | mucin 1, transmembrane | MUC1 | 4582 | 4.029 | 3.7 |
| 182_at | U01062 | inositol 1,4,5-triphosphate receptor, type 3 | ITPR3 | 3710 | 5.886 | 3.7 |
| 406_at | X54489 | chemokine (C-X-C motif) ligand 2 | CXCL2 | 2920 | 3.162 | 3.7 |
| 943_at | D43968 | runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene) | RUNX1 | 861 | 9.283 | 3.7 |
| 33956_at | AB018549 | lymphocyte antigen 96 | LY96 | 23643 | 7.483 | 3.7 |
| 40642_at | A1312646 | nuclear factor I/B | NFIB | 4781 | 6.658 | 3.6 |
| 1262_s_at | M19154 | transforming growth factor, beta 2 | TGFB2 | 7042 | 3.530 | 3.6 |
| 32314_g_at | M75165 | tropomyosin 2 (beta) | TPM2 | 7169 | 4.846 | 3.6 |
| 39008_at | M13699 | ceruloplasmin (ferroxidase) | CP | 1356 | 3.864 | 3.6 |
| 34678_at | AL096713 | fer-1-like 3, myoferlin (C. elegans) | FER1L3 | 26509 | 7.463 | 3.6 |
| 32083_at | AF027826 | transmembrane 7 superfamily member 1 (upregulated in kidney) | TM7SF1 | 7107 | 3.262 | 3.6 |
| 36689_at | AL040446 | oxysterol binding protein-like 1A | OSBPL1A | 114876 | 3.399 | 3.5 |
| 33131_at | X70683 | SRY (sex determining region Y)-box 4 | SOX4 | 6659 | 7.140 | 3.5 |
| 38138_at | AC004668 | S100 calcium binding protein A11 (calgizzarin) | S100A11 | 6282 | 5.565 | 3.5 |
| 36397_at | U62434 | cholinergic receptor, nicotinic, alpha polypeptide 5 | CHRNA5 | 1138 | 3.441 | 3.4 |
| 36496_at | AF014398 | inositol(myo)-1(or 4)-monophosphatase 2 | IMPA2 | 3613 | 4.542 | 3.4 |
| 34203_at | D17408 | calponin 1, basic, smooth muscle | CNN1 | 1264 | 3.861 | 3.4 |
| 38038_at | U21128 | lumican | LUM | 4080 | 8.584 | 3.4 |
| 36681_at | J02911 | apolipoprotein D | APOD | 347 | 3.354 | 3.4 |
| 925_at | J03909 | interferon, gamma-inducible protein 30 | IFI30 | 10437 | 9.043 | 3.4 |
| 1371_s_at | M29874 | cytochrome P450, family 2, subfamily B, polypeptide 6 | CYP2B6 | 1555 | 3.444 | 3.4 |
| 675_at | J04164 | interferon induced transmembrane protein 1 (9-27) | IFITM1 | 8519 | 6.128 | 3.4 |
| 32451_at | L35848 | membrane-spanning 4-domains, subfamily A, member 3 (hematopoietic cell-specific) | MS4A3 | 932 | 5.668 | 3.4 |
| 38013_at | AL096842 | mitochondrial tumor suppressor 1 | MTUS1 | 57509 | 4.513 | 3.4 |
| 40522_at | X59834 | glutamate-ammonia ligase (glutamine synthase) | GLUL | 2752 | 8.219 | 3.4 |
| 32404_at | AF065314 | cyclic nucleotide gated channel alpha 3 | CNGA3 | 1261 | 3.816 | 3.4 |
| 39728_at | J03909 | interferon, gamma-inducible protein 30 | IFI30 | 10437 | 9.556 | 3.3 |
| 36575_at | S59049 | regulator of G-protein signalling 1 | RGS1 | 5996 | 6.548 | 3.3 |
| 37407_s_at | AF013570 | myosin, heavy polypeptide 11, smooth muscle | MYH11 | 4629 | 2.748 | 3.3 |
| 1501_at | X57025 | insulin-like growth factor 1 (somatomedin C) | IGF1 | 3479 | 3.031 | 3.3 |
| 659_g_at | L12350 | thrombospondin 2 | THBS2 | 7058 | 5.066 | 3.3 |
| 32454_at | AL080215 | MRNA; cDNA DKFZp586J0323 (from clone DKFZp586J0323) | --- | --- | 2.390 | 3.3 |
| 32081_at | AB023166 | citron (rho-interacting, serine/threonine kinase 21) | CIT | 11113 | 3.426 | 3.3 |
| 36543_at | J02931 | coagulation factor III (thromboplastin, tissue factor) | F3 | 2152 | 4.152 | 3.3 |
| 32985_at | AF017655 | sal-like 1 (Drosophila) | SALL1 | 6299 | 3.334 | 3.3 |
| 38422_s_at | U29332 | four and a half LIM domains 2 | FHL2 | 2274 | 5.225 | 3.3 |
| 40543_at | L08424 | achaete-scute complex-like 1 (Drosophila) | ASCL1 | 429 | 4.370 | 3.2 |
| 38754_at | A1557295 | ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeier-Vogt disease) | CLN3 | 1201 | 6.740 | 3.2 |
| 40715_at | D83597 | lymphocyte antigen 64 homolog, radioprotective 105kDa (mouse) | LY64 | 4064 | 1.753 | 3.2 |
| 39695_at | M31516 | decay accelerating factor for complement (CD55, Cromer blood group system) | DAF | 1604 | 5.928 | 3.2 |
| 37061_at | U29615 | chitinase 1 (chitotriosidase) | CHIT1 | 1118 | 3.701 | 3.2 |

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|------------|---------------|--|---------------|-------|-------|-----|
| 34878_at | AB019987 | SMC4 structural maintenance of chromosomes 4-like 1 (yeast) | SMC4L1 | 10051 | 5.363 | 3.2 |
| 37208_at | AJ001612 | phosphoserine phosphatase | PSPH | 5723 | 2.130 | 3.2 |
| 35367_at | AB006780 | lectin, galactoside-binding, soluble, 3 (galectin 3) | LGALS3 | 3958 | 6.051 | 3.2 |
| 38691_s_at | J03553 | surfactant, pulmonary-associated protein C | SFTPC | 6440 | 2.106 | 3.2 |
| 35168_f_at | M92642 | collagen, type XVI, alpha 1 | COL16A1 | 1307 | 5.214 | 3.2 |
| 34007_at | AJ236922 | glutamate receptor, metabotropic 8 | GRM8 | 2918 | 2.149 | 3.2 |
| 31682_s_at | D32039 | chondroitin sulfate proteoglycan 2 (versican) | CSPG2 | 1462 | 7.100 | 3.2 |
| 37021_at | X16832 | cathepsin H | CTSH | 1512 | 7.046 | 3.2 |
| 36227_at | AF043129 | interleukin 7 receptor | IL7R | 3575 | 5.796 | 3.2 |
| 1280_i_at | HG2702-HT2798 | Serine/Threonine Kinase | --- | --- | 5.638 | 3.2 |
| 37688_f_at | M31932 | Fc fragment of IgG, low affinity Ila, receptor for (CD32) | FCGR2A | 2212 | 5.664 | 3.2 |
| 34378_at | X97324 | adipose differentiation-related protein | ADFP | 123 | 6.629 | 3.2 |
| 33244_at | U07223 | chimerin (chimaerin) 2 | CHN2 | 1124 | 3.029 | 3.2 |
| 35964_at | AJ224741 | matrilin 3 | MATN3 | 4148 | 2.705 | 3.2 |
| 40496_at | J04080 | complement component 1, s subcomponent | C1S | 716 | 6.992 | 3.2 |
| 33331_at | U17077 | BENE protein | BENE | 7851 | 4.861 | 3.1 |
| 38059_g_at | Z22865 | dermatopontin | DPT | 1805 | 2.640 | 3.1 |
| 36059_at | AB011540 | low density lipoprotein receptor-related protein 4 | LRP4 | 4038 | 3.424 | 3.1 |
| 39176_f_at | M54994 | carboxyl ester lipase (bile salt-stimulated lipase) | CEL | 1056 | 2.277 | 3.1 |
| 31344_at | AF063725 | Clone BCSynL38 immunoglobulin lambda light chain variable region mRNA, partial cds | --- | --- | 4.724 | 3.1 |
| 38378_at | M37033 | CD53 antigen | CD53 | 963 | 6.393 | 3.1 |
| 36203_at | X16277 | ornithine decarboxylase 1 | ODC1 | 4953 | 4.144 | 3.1 |
| 37645_at | Z22576 | CD69 antigen (p60, early T-cell activation antigen) | CD69 | 969 | 3.327 | 3.1 |
| 41176_at | AF052162 | hypothetical protein FLJ12443 | FLJ12443 | 79888 | 6.430 | 3.1 |
| 41353_at | Z29574 | tumor necrosis factor receptor superfamily, member 17 | TNFRSF17 | 608 | 5.799 | 3.1 |
| 41177_at | AW024285 | hypothetical protein FLJ12443 | FLJ12443 | 79889 | 6.093 | 3.1 |
| 41073_at | AI743745 | G protein-coupled receptor 49 | GPR49 | 8549 | 3.528 | 3.1 |
| 34304_s_at | AL050290 | spermidine/spermine N1-acetyltransferase | SAT | 6303 | 9.191 | 3.0 |
| 1599_at | L25876 | cyclin-dependent kinase inhibitor 3 (CDK2-associated dual specificity phosphatase) | CDKN3 | 1033 | 2.082 | 3.0 |
| 35479_at | AJ242015 | a disintegrin and metalloproteinase domain 28 | ADAM28 | 10863 | 4.905 | 3.0 |
| 32134_at | AL050162 | testis derived transcript (3 LIM domains) | TES | 26136 | 6.289 | 3.0 |
| 40848_g_at | AB018293 | flavoprotein oxidoreductase MICAL2 | MICAL2 | 9645 | 6.480 | 3.0 |
| 35937_at | U65416 | MHC class I polypeptide-related sequence B | MICB | 4277 | 5.991 | 3.0 |
| 32873_at | AC004888 | potassium voltage-gated channel, Shal-related subfamily, member 2 | KCNQ2 | 3751 | 2.991 | 3.0 |
| 368_at | Z29083 | trophoblast glycoprotein | TPBG | 7162 | 4.091 | 3.0 |
| 38734_at | M63603 | phospholamban | PLN | 5350 | 4.216 | 3.0 |
| 41632_at | D38550 | E2F transcription factor 3 | E2F3 | 1871 | 7.699 | 3.0 |
| 34770_at | Z14138 | mitogen-activated protein kinase kinase kinase 8 | MAP3K8 | 1326 | 4.572 | 3.0 |
| 32034_at | AF041259 | zinc finger protein 217 | ZNF217 | 7784 | 6.640 | 3.0 |
| 32535_at | U63556 | fibrillin 1 (Marfan syndrome) | FBN1 | 2200 | 4.670 | 3.0 |
| 35431_at | U05291 | fibromodulin | FMDQ | 2331 | 5.288 | 3.0 |
| 39690_at | AF032822 | PDZ and LIM domain 3 | PDLIM3 | 27295 | 3.692 | 3.0 |
| 542_at | S74445 | cellular retinoic acid binding protein 1 | CRABP1 | 1381 | 2.258 | 2.9 |
| 39125_at | M14083 | serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1 | SERPINE1 | 5054 | 3.371 | 2.9 |
| 39114_at | AB022718 | chromosome 10 open reading frame 10 | C10orf10 | 11067 | 3.678 | 2.9 |
| 37365_at | X63368 | DnaJ (Hsp40) homolog, subfamily B, member 2 | DNAJB2 | 3300 | 7.054 | 2.9 |
| 36453_at | AB018254 | KIAA0711 gene product | KIAA0711 | 9920 | 3.311 | 2.9 |
| 39143_at | U08015 | nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1 | NFATC1 | 4772 | 4.870 | 2.9 |
| 32773_at | AA868382 | major histocompatibility complex, class II, DQ alpha 1 | HLA-DQA1 | 3117 | 5.277 | 2.9 |
| 34736_at | M25753 | cyclin B1 | CCNB1 | 891 | 5.732 | 2.9 |
| 32227_at | X17042 | proteoglycan 1, secretory granule | PRG1 | 5552 | 6.770 | 2.9 |
| 41229_at | AI222594 | nuclear factor I/B | NFIB | 4781 | 6.347 | 2.9 |
| 41815_at | AL080133 | spectrin repeat containing, nuclear envelope 2 | SYNE2 | 23224 | 4.477 | 2.9 |
| 1891_at | D14497 | mitogen-activated protein kinase kinase kinase 8 | MAP3K8 | 1326 | 5.288 | 2.9 |
| 40915_r_at | Y00272 | cell division cycle 2, G1 to S and G2 to M | CDC2 | 983 | 4.244 | 2.9 |
| 37219_at | X72755 | chemokine (C-X-C motif) ligand 9 | CXCL9 | 4283 | 4.695 | 2.9 |
| 41531_at | AI445461 | transmembrane 4 superfamily member 1 | TM4SF1 | 4071 | 4.958 | 2.9 |
| 35807_at | M21186 | cytochrome b-245, alpha polypeptide | CYBA | 1535 | 9.528 | 2.9 |
| 36376_at | AF030890 | solute carrier family 26, member 4 | SLC26A4 | 5172 | 2.167 | 2.9 |
| 1054_at | M87339 | replication factor C (activator 1) 4, 37kDa | RFC4 | 5984 | 3.608 | 2.9 |
| 37200_at | J04162 | Fc fragment of IgG, low affinity IIIa, receptor for (CD16) | FCGR3A | 2214 | 6.081 | 2.9 |
| 39167_r_at | D83174 | serine (or cysteine) proteinase inhibitor, clade H (heat shock protein 47), member 1, (collagen binding protein 1) | SERPINH1 | 871 | 6.144 | 2.8 |
| 41064_at | AL049296 | likely ortholog of mouse nervous system polycmb 1 | NSPC1 | 84759 | 5.978 | 2.8 |
| 37233_at | AF079167 | oxidized low density lipoprotein (lectin-like) receptor 1 | OLR1 | 4973 | 4.206 | 2.8 |
| 37842_at | AF054589 | I-mfa domain-containing protein | HIC | 29969 | 6.020 | 2.8 |
| 40051_at | D31762 | translocation associated membrane protein 2 | TRAM2 | 9697 | 5.793 | 2.8 |
| 38805_at | X89750 | TGF-beta-induced factor (TAFE family homeobox) | TGIF | 7050 | 7.490 | 2.8 |
| 133_at | X87212 | cathepsin C | CTSC | 1075 | 6.331 | 2.8 |
| 37576_at | U52969 | Purkinje cell protein 4 | PCP4 | 5121 | 2.507 | 2.8 |
| 32582_at | X69292 | myosin, heavy polypeptide 11, smooth muscle | MYH11 | 4629 | 2.903 | 2.8 |
| 39677_at | D80008 | KIAA0186 gene product | KIAA0186 | 9837 | 5.816 | 2.8 |
| 31870_at | X14046 | CD37 antigen | CD37 | 951 | 4.414 | 2.8 |
| 37405_at | U29091 | selenium binding protein 1 | SELENBP1 | 8991 | 6.740 | 2.8 |
| 33279_s_at | X80062 | SA hypertension-associated homolog (rat) | SAH | 6296 | 4.769 | 2.8 |
| 36079_at | AF010309 | tumor protein p53 inducible protein 3 | TP53I3 | 9540 | 4.948 | 2.8 |
| 41401_at | U57646 | cysteine and glycine-rich protein 2 | CGRP2 | 1466 | 5.246 | 2.8 |
| 35961_at | AL049390 | MRNA, cDNA DKFZp586O1318 (from clone DKFZp586O1318) | --- | --- | 3.442 | 2.8 |
| 39069_at | AF053944 | AE binding protein 1 | AEBP1 | 165 | 3.991 | 2.8 |
| 35985_at | AB023137 | A kinase (PRKA) anchor protein 2 | AKAP2 | 11217 | 5.366 | 2.8 |
| 35729_at | AB018270 | myosin ID | MYO1D | 4642 | 6.349 | 2.8 |
| 34267_r_at | U50748 | leptin receptor | LEPR | 3953 | 2.242 | 2.8 |
| 40767_at | M59499 | tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor) | TFPI | 7035 | 4.587 | 2.8 |
| 37749_at | D78611 | mesoderm specific transcript homolog (mouse) | MEST | 4232 | 4.830 | 2.8 |
| 36708_at | J04970 | carboxypeptidase M | CPM | 1368 | 5.578 | 2.7 |
| 41504_s_at | AF055376 | v-maf musculoaponeurotic fibrosarcoma oncogene homolog (avian) | MAF | 4094 | 5.231 | 2.7 |
| 41365_at | Y09788 | mucin 5, subtype B, tracheobronchial | MUC5B | 4587 | 3.024 | 2.7 |
| 575_s_at | M93036 | tumor-associated calcium signal transducer 1 | TACSTD1 | 4072 | 6.000 | 2.7 |
| 2042_s_at | M15024 | v-myb myeloblastosis viral oncogene homolog (avian) | MYB | 4602 | 2.499 | 2.7 |
| 757_at | D28364 | annexin A2 | ANXA2 | 302 | 6.352 | 2.7 |
| 35370_at | AB018307 | SPTF-associated factor 65 gamma | STAF65(gamma) | 9913 | 6.590 | 2.7 |
| 1456_s_at | M63838 | interferon, gamma-inducible protein 16 | IFI16 | 3428 | 5.216 | 2.7 |
| 32112_s_at | AF015732 | absent in melanoma 1 | AIM1 | 202 | 4.587 | 2.7 |
| 38028_at | AL050152 | LIM domain only 3 (rhombotin-like 2) | LMO3 | 55895 | 5.669 | 2.7 |
| 1625_at | HG2309-HT2405 | Insulin-Like Growth Factor I | --- | --- | 2.029 | 2.7 |
| 677_s_at | J04430 | acid phosphatase 5, tartrate resistant | ACP5 | 54 | 7.717 | 2.7 |
| 38363_at | W60864 | TYRO protein tyrosine kinase binding protein | TYROBP | 7305 | 6.211 | 2.7 |
| 1261_i_at | M16594 | glutathione S-transferase A2 | GSTA2 | 2939 | 2.045 | 2.7 |
| 1427_g_at | D89077 | Src-like adaptor | SLA | 6503 | 5.241 | 2.7 |
| 1173_g_at | HG172-HT3924 | Spermidine/Spermine N1-Acetyltransferase, Alt. Splice 2 | --- | --- | 9.337 | 2.7 |
| 36753_at | U82979 | leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 4 | LILRB4 | 11006 | 6.798 | 2.7 |
| 37512_at | U89281 | 3-hydroxysteroid epimerase | RODH | 8630 | 4.309 | 2.7 |
| 160031_at | X63629 | cadherin 3, type 1, P-cadherin (placental) | CDH3 | 1001 | 5.752 | 2.6 |
| 32640_at | M24283 | intercellular adhesion molecule 1 (CD54), human rhinovirus receptor | ICAM1 | 3383 | 4.973 | 2.6 |
| 1761_at | D37965 | platelet-derived growth factor receptor-like | PDGFRL | 5157 | 5.831 | 2.6 |
| 1372_at | M31165 | tumor necrosis factor, alpha-induced protein 6 | TNFAIP6 | 7130 | 5.168 | 2.6 |
| 37018_at | AI189287 | histone 1, H1c | HIST1H1C | 3006 | 3.094 | 2.6 |
| 767_at | AF001548 | myosin, heavy polypeptide 11, smooth muscle | MYH11 | 4629 | 2.585 | 2.6 |
| 1656_s_at | U07563 | v-abl Abelson murine leukemia viral oncogene homolog 1 | ABL1 | 25 | 3.562 | 2.6 |
| 35226_at | U71207 | eyes absent homolog 2 (Drosophila) | EYA2 | 2139 | 4.597 | 2.6 |
| 32186_at | M80244 | solute carrier family 7 (cationic amino acid transporter, y+ system), member 5 | SLC7A5 | 8140 | 6.046 | 2.6 |
| 39710_at | U30521 | chromosome 5 open reading frame 13 | C5orf13 | 9315 | 5.736 | 2.6 |
| 39061_at | D28137 | bone marrow stromal cell antigen 2 | BST2 | 684 | 4.787 | 2.6 |
| 39135_at | AB018310 | KIAA0767 protein | KIAA0767 | 23151 | 4.622 | 2.6 |
| 40516_at | L19872 | aryl hydrocarbon receptor | AHR | 196 | 5.602 | 2.6 |

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|------------|----------|--|----------|--------|-------|-----|
| 38247_at | U67058 | coagulation factor II (thrombin) receptor-like 1 | F2RL1 | 2150 | 3.857 | 2.6 |
| 32847_at | U48959 | myosin, light polypeptide kinase | MYLK | 4638 | 3.692 | 2.6 |
| 40726_at | U37426 | kinesin family member 11 | KIF11 | 3832 | 4.419 | 2.6 |
| 40091_at | U00115 | B-cell CLL/lymphoma 6 (zinc finger protein 51) | BCL6 | 604 | 6.263 | 2.6 |
| 1495_at | M34057 | latent transforming growth factor beta binding protein 1 | LTBP1 | 4052 | 5.464 | 2.6 |
| 38065_at | X62534 | high-mobility group box 2 | HMGB2 | 3148 | 4.834 | 2.6 |
| 38288_at | U69196 | snail homolog 2 (Drosophila) | SNAI2 | 6591 | 2.429 | 2.6 |
| 38661_at | X75314 | RNA-binding region (RNP1, RRM) containing 1 | RNFC1 | 55544 | 5.749 | 2.6 |
| 36643_at | L20817 | discoilin domain receptor family, member 1 | DDR1 | 780 | 5.694 | 2.6 |
| 35794_at | AB023159 | ADP-ribosylation factor guanine nucleotide factor 6 | EFAR8 | 23362 | 5.971 | 2.6 |
| 41409_at | AF044896 | chromosome 1 open reading frame 38 | C1orf38 | 9473 | 8.029 | 2.5 |
| 39319_at | U20158 | lymphocyte cytosolic protein 2 (SH2 domain containing leukocyte protein of 76kDa) | LCP2 | 3937 | 5.701 | 2.5 |
| 36906_at | U73304 | cannabinoid receptor 1 (brain) | CNR1 | 1268 | 1.950 | 2.5 |
| 33876_at | AL050107 | transcriptional co-activator with PDZ-binding motif (TAZ) | TAZ | 25937 | 5.967 | 2.5 |
| 1478_at | L10717 | IL2-inducible T-cell kinase | ITK | 3702 | 2.988 | 2.5 |
| 1633_g_at | U77735 | pim-2 oncogene | PIM2 | 11040 | 4.290 | 2.5 |
| 32113_at | U83115 | absent in melanoma 1 | AIM1 | 202 | 3.485 | 2.5 |
| 39333_at | M11315 | collagen, type IV, alpha 1 | COL4A1 | 1282 | 4.199 | 2.5 |
| 34202_at | AL050071 | poliovirus receptor-related 3 | PVR3 | 25945 | 2.815 | 2.5 |
| 33390_at | AA203487 | serine/threonine kinase 17b (apoptosis-inducing) | STK17B | 9262 | 6.001 | 2.5 |
| 31897_at | U53445 | downregulated in ovarian cancer 1 | DOC1 | 11259 | 3.781 | 2.5 |
| 39593_at | AI432401 | fibrinogen-like 2 | FGL2 | 10875 | 4.352 | 2.5 |
| 272_at | K02054 | gastrin-releasing peptide | GRP | 2922 | 2.184 | 2.5 |
| 1654_at | X13293 | v-myb myeloblastosis viral oncogene homolog (avian)-like 2 | MYBL2 | 4605 | 5.096 | 2.5 |
| 33324_s_at | D89357 | cell division cycle 2, G1 to S and G2 to M | CD22 | 983 | 5.512 | 2.5 |
| 34822_at | U58334 | tumor protein p53 binding protein, 2 | TP53BP2 | 7159 | 9.873 | 2.5 |
| 1945_at | M25753 | cyclin B1 | CCNB1 | 891 | 5.027 | 2.5 |
| 393_s_at | X90976 | runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene) | RUNX1 | 861 | 7.163 | 2.5 |
| 339_at | AF035752 | caveolin 2 | CAV2 | 858 | 4.120 | 2.5 |
| 1960_at | L22967 | L22967/FEATURE= /DEFINITION=HUMC Homo sapiens acidic growth fibroblast growth factor (aFGF) gene sequence | --- | --- | 2.117 | 2.5 |
| 1062_g_at | U00672 | interleukin 10 receptor, alpha | IL10RA | 3587 | 6.006 | 2.5 |
| 1984_s_at | X69549 | Rho GDP dissociation inhibitor (GDI) beta | ARHGDB | 397 | 5.973 | 2.5 |
| 977_s_at | Z35402 | cadherin 1, type 1, E-cadherin (epithelial) | CDH1 | 999 | 5.006 | 2.5 |
| 33849_at | U02020 | pre-B-cell colony enhancing factor 1 | PBEF1 | 10135 | 5.832 | 2.5 |
| 37324_at | X01060 | transferrin receptor (p90, CD71) | TFRC | 7037 | 6.047 | 2.5 |
| 37170_at | AB015331 | BMP2 inducible kinase | BMP2K | 55589 | 5.194 | 2.5 |
| 404_at | X52425 | interleukin 4 receptor | IL4R | 3566 | 5.819 | 2.5 |
| 40412_at | AA203476 | pituitary tumor-transforming 1 | PTTG1 | 9232 | 5.984 | 2.5 |
| 33462_at | D13626 | G protein-coupled receptor 105 | GPR105 | 9934 | 3.034 | 2.5 |
| 287_at | L19871 | activating transcription factor 3 | ATF3 | 467 | 3.199 | 2.5 |
| 41700_at | M62424 | coagulation factor II (thrombin) receptor | F2R | 2149 | 3.502 | 2.5 |
| 38416_at | X97978 | cyclin D1 (PRAD1; parathyroid adenomatosis 1) | CND1 | 595 | 4.118 | 2.5 |
| 32240_at | M65292 | H factor (complement)-like 1 | HFL1 | 3078 | 3.901 | 2.5 |
| 40720_at | AL022398 | TRAF3-interacting Jun N-terminal kinase (JNK)-activating modulator | TJAM | 80342 | 5.109 | 2.5 |
| 1733_at | M60315 | bone morphogenetic protein 6 | BMP6 | 654 | 3.289 | 2.4 |
| 37403_at | X05908 | annexin A1 | ANXA1 | 301 | 3.711 | 2.4 |
| 40698_at | X96719 | C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 2 (activation-induced) | CLEC5F2 | 9976 | 5.096 | 2.4 |
| 37024_at | AF010312 | lipopolysaccharide-induced TNF factor | LITAF | 9516 | 5.428 | 2.4 |
| 2002_s_at | U27467 | BCL2-related protein A1 | BCL2A1 | 597 | 3.976 | 2.4 |
| 2020_at | M73554 | cyclin D1 (PRAD1; parathyroid adenomatosis 1) | CCND1 | 595 | 3.972 | 2.4 |
| 38740_at | X79067 | zinc finger protein 36, C3H type-like 1 | ZFP36L1 | 677 | 6.885 | 2.4 |
| 34857_at | Z24724 | ATPase family homolog up-regulated in senescence cells | AFURS1 | 79572 | 4.789 | 2.4 |
| 34715_at | U74612 | forkhead box M1 | FOXM1 | 2305 | 5.540 | 2.4 |
| 41723_s_at | M32578 | major histocompatibility complex, class II, DR beta 1 | HLA-DRB1 | 3123 | 5.154 | 2.4 |
| 40856_at | U29953 | serine (or cysteine) proteinase inhibitor, clade F (alpha-2 antipainin, pigment epithelium derived factor), member 1 | SERPINF1 | 5176 | 6.873 | 2.4 |
| 33213_g_at | AF006751 | ribosome binding protein 1 homolog 180kDa (dog) | RRBP1 | 6238 | 6.638 | 2.4 |
| 1909_at | M14745 | B-cell CLL/lymphoma 2 | BCL2 | 596 | 4.220 | 2.4 |
| 35474_s_at | Y15915 | chimaeric transcript of collagen type 1 alpha 1 and platelet derived growth factor beta | --- | --- | 6.356 | 2.4 |
| 531_at | U18907 | GLI pathogenesis-related 1 (glioma) | GLIPR1 | 11010 | 4.336 | 2.4 |
| 33690_at | M18667 | prolactinase (pepsinogen C) | PGC | 5225 | 2.018 | 2.4 |
| 36659_at | X05610 | collagen, type IV, alpha 2 | COL4A2 | 1284 | 4.237 | 2.4 |
| 36341_s_at | X64877 | H factor (complement)-like 1 | HFL1 | 3078 | 3.901 | 2.4 |
| 35699_at | AF046918 | BUB1 budding uninhibited by benzimidazoles 1 homolog beta (yeast) | BUB1B | 701 | 4.314 | 2.4 |
| 38796_at | X03084 | complement component 1, q subcomponent, beta polypeptide | C1QB | 713 | 5.113 | 2.4 |
| 37310_at | X02419 | plasminogen activator, urokinase | PLAU | 5328 | 4.103 | 2.4 |
| 1237_at | S81914 | immediate early response 3 | IER3 | 8870 | 4.748 | 2.4 |
| 35357_at | W26640 | PP784 protein | PP784 | 114932 | 3.186 | 2.4 |
| 1368_at | M27492 | interleukin 1 receptor, type I | IL1R1 | 3554 | 5.651 | 2.4 |
| 160025_at | X70340 | transforming growth factor, alpha | TGFA | 7039 | 3.848 | 2.4 |
| 40078_at | AF015287 | protease, serine, 23 | SPUVE | 11098 | 4.697 | 2.4 |
| 40074_at | X16396 | methylene tetrahydrofolate dehydrogenase (NAD+ dependent), methenyltetrahydrofolate cyclohydrolase | MTHFD2 | 10797 | 5.942 | 2.4 |
| 37544_at | X64318 | nuclear factor, interleukin 3 regulated | NFIL3 | 4783 | 4.622 | 2.4 |
| 32749_s_at | AL050396 | filamin A, alpha (actin binding protein 280) | FLNA | 2316 | 6.125 | 2.4 |
| 33255_at | M97856 | nuclear autoantigenic sperm protein (histone-binding) | NASP | 4678 | 5.915 | 2.4 |
| 31496_g_at | X39798 | chemokine (C motif) ligand 1 | XCL1 | 6375 | 3.903 | 2.3 |
| 1592_at | J04088 | topoisomerase (DNA) II, alpha 170kDa | TOP2A | 7153 | 5.214 | 2.3 |
| 34389_at | Y11710 | collagen, type XIV, alpha 1 (undulin) | COL14A1 | 7373 | 3.939 | 2.3 |
| 32332_at | X69433 | isocitrate dehydrogenase 2 (NADP+), mitochondrial | IDH2 | 3418 | 6.328 | 2.3 |
| 33051_at | M62400 | gamma-aminobutyric acid (GABA) receptor, rho 1 | GABRR1 | 2569 | 1.925 | 2.3 |
| 37918_at | M15395 | integrin, beta 2 (antigen CD18 (p95), lymphocyte function-associated antigen 1; macrophage antigen 1 (mac-1) beta subunit) | ITGB2 | 3689 | 6.923 | 2.3 |
| 41140_at | U05875 | interferon gamma receptor 2 (interferon gamma transducer 1) | IFNGR2 | 3460 | 4.362 | 2.3 |
| 36909_at | X62048 | WEE1 homolog (S. pombe) | WEE1 | 7465 | 4.147 | 2.3 |
| 36849_at | U90920 | PTPL1-associated RhoGAP 1 | PARG1 | 9411 | 3.623 | 2.3 |
| 1348_s_at | S79219 | propionyl Coenzyme A carboxylase, alpha polypeptide | PCCA | 5095 | 1.640 | 2.3 |
| 947_at | D55716 | MCM7 minichromosome maintenance deficient 7 (S. cerevisiae) | MCM7 | 4176 | 6.440 | 2.3 |
| 36908_at | M93221 | mannose receptor, C type 1 | MRC1 | 4360 | 3.410 | 2.3 |
| 37225_at | D79994 | ankyrin repeat domain 15 | ANKRD15 | 23189 | 6.628 | 2.3 |
| 41215_s_at | L13891 | inhibitor of DNA binding 2, dominant negative helix-loop-helix protein | ID2 | 3398 | 6.078 | 2.3 |
| 31631_f_at | AF095288 | pituitary tumor-transforming 1 | PTTG1 | 9232 | 6.194 | 2.3 |
| 1365_at | M26062 | interleukin 2 receptor, beta | IL2RB | 3560 | 5.278 | 2.3 |
| 1057_at | M97815 | cellular retinoic acid binding protein 2 | CRABP2 | 1382 | 3.779 | 2.3 |
| 41585_at | AB019289 | KIAA0746 protein | KIAA0746 | 23231 | 7.476 | 2.3 |
| 31884_at | M62896 | annexin A2 pseudogene 1 | ANXA2P1 | 303 | 5.330 | 2.3 |
| 2017_s_at | M64349 | cyclin D1 (PRAD1; parathyroid adenomatosis 1) | CCND1 | 595 | 4.462 | 2.3 |
| 32263_at | AL080146 | cyclin B2 | CCNB2 | 9133 | 5.192 | 2.3 |
| 33238_at | U23852 | lymphocyte-specific protein tyrosine kinase | LCK | 3932 | 5.587 | 2.3 |
| 1715_at | U37518 | tumor necrosis factor (ligand) superfamily, member 10 | TNFSF10 | 8743 | 5.174 | 2.3 |
| 33026_at | W28250 | Transcribed sequences | --- | --- | 4.637 | 2.3 |
| 31512_at | Z00010 | immunoglobulin kappa constant | IGKC | 3514 | 2.385 | 2.3 |
| 37305_at | U61145 | enhancer of zeste homolog 2 (Drosophila) | EZH2 | 2146 | 5.425 | 2.3 |
| 41216_r_at | D13891 | inhibitor of DNA binding 2, dominant negative helix-loop-helix protein | ID2 | 3398 | 5.440 | 2.3 |
| 37039_at | J00194 | major histocompatibility complex, class II, DR alpha | HLA-DRA | 3122 | 7.267 | 2.3 |
| 39409_at | M14058 | complement component 1, r subcomponent | C1R | 715 | 5.531 | 2.3 |
| 39951_at | L20826 | plastin 1 (I isoform) | PLS1 | 5357 | 4.511 | 2.3 |
| 33371_s_at | U59877 | RAB31, member RAS oncogene family | RAB31 | 11031 | 5.975 | 2.3 |
| 38992_at | X64229 | DEK oncogene (DNA binding) | DEK | 7913 | 6.563 | 2.3 |
| 38772_at | Y11307 | cysteine-rich, angiogenic inducer, 61 | CYR61 | 3491 | 3.081 | 2.3 |
| 37280_at | U59912 | SMAD, mothers against DPP homolog 1 (Drosophila) | SMAD1 | 4086 | 4.812 | 2.2 |
| 33121_g_at | AF045229 | regulator of G-protein signaling 10 | RGS10 | 6001 | 4.095 | 2.2 |
| 31355_at | U77629 | achaete-scute complex-like 2 (Drosophila) | ASCL2 | 430 | 4.038 | 2.2 |
| 417_at | X65293 | protein kinase C, epsilon | PRCKE | 5581 | 5.397 | 2.2 |
| 35012_at | M81750 | myeloid cell nuclear differentiation antigen | MNDA | 4332 | 3.792 | 2.2 |
| 34514_at | S49816 | thyroid stimulating hormone receptor | TSHR | 7253 | 2.305 | 2.2 |
| 40803_at | AL050161 | pro-oncosis receptor inducing membrane injury gene | PORIMIN | 114908 | 5.362 | 2.2 |

| | | | | | | |
|------------|----------|--|-------------|--------|-------|-----|
| 892_at | M90657 | transmembrane 4 superfamily member 1 | TM4SF1 | 4071 | 5.071 | 2.2 |
| 39113_at | AI262789 | protein disulfide isomerase related protein (calcium-binding protein, intestinal-related) | ERP70 | 9601 | 5.849 | 2.2 |
| 35995_at | AF067656 | ZW10 interactor | ZWINT | 11130 | 6.833 | 2.2 |
| 34795_at | U84573 | procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine hydroxylase) 2 | PLOD2 | 5352 | 5.379 | 2.2 |
| 37831_at | AB011117 | signal-induced proliferation-associated 1 like 3 | SIPA1L3 | 23094 | 6.898 | 2.2 |
| 36736_f_at | Y10275 | L-3-phosphoserine phosphatase | --- | --- | 2.300 | 2.2 |
| 33322_i_at | X57348 | stratifin | SFN | 2810 | 7.732 | 2.2 |
| 37209_g_at | AJ001612 | phosphoserine phosphatase | PSPH | 5723 | 2.310 | 2.2 |
| 37762_at | Y07939 | epithelial membrane protein 1 | EMP1 | 2012 | 3.156 | 2.2 |
| 41536_at | AL022726 | inhibitor of DNA binding 4, dominant negative helix-loop-helix protein | ID4 | 3400 | 3.479 | 2.2 |
| 2089_s_at | H06628 | v-erb-b2 erythroblastic leukemia viral oncogene homolog 3 (avian) | ERBB3 | 2065 | 4.035 | 2.2 |
| 33127_at | U89942 | lysyl oxidase-like 2 | LOXL2 | 4017 | 5.246 | 2.2 |
| 36007_at | AL050137 | olfactomedin-like 2B | OLFM2B | 25903 | 5.233 | 2.2 |
| 39154_at | AI952982 | growth arrest and DNA-damage-inducible, gamma | GADD45G | 10912 | 3.538 | 2.2 |
| 37203_at | L07765 | carboxylesterase 1 (monocyte/macrophage serine esterase 1) | CES1 | 1066 | 1.968 | 2.2 |
| 34407_at | U77594 | retinoic acid receptor responder (tazarotene induced) 2 | RARRES2 | 5919 | 3.309 | 2.2 |
| 40161_at | L32137 | cartilage oligomeric matrix protein | COMP | 1311 | 3.532 | 2.2 |
| 1190_at | Z48541 | protein tyrosine phosphatase, receptor type, O | PTPRO | 5800 | 2.000 | 2.2 |
| 36550_at | AL049538 | Ras and Rab interactor 2 | RIN2 | 54453 | 6.409 | 2.2 |
| 1951_at | AF004327 | angiotensin II | ANGPT2 | 285 | 3.613 | 2.2 |
| 1721_g_at | U65410 | MAD2 mitotic arrest deficient-like 1 (yeast) | MAD2L1 | 4085 | 3.221 | 2.2 |
| 38545_at | M31682 | inhibin, beta B (activin AB beta polypeptide) | INHBB | 3625 | 3.398 | 2.2 |
| 38096_f_at | M63664 | major histocompatibility complex, class II, DP beta 1 | HLA-DPB1 | 3115 | 4.618 | 2.2 |
| 671_at | J03040 | secreted protein, acidic, cysteine-rich (osteonectin) | SPARC | 6678 | 5.184 | 2.2 |
| 41186_at | W28186 | lysosomal associated protein transmembrane 4 beta | LAPTM4B | 55353 | 7.184 | 2.2 |
| 875_g_at | M26883 | chemokine (C-C motif) ligand 2 | CCL2 | 6347 | 4.044 | 2.2 |
| 35016_at | M13560 | CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated) | CD74 | 972 | 6.555 | 2.2 |
| 34793_s_at | M22299 | plastin 3 (T isoform) | PLS3 | 5358 | 4.167 | 2.2 |
| 38589_i_at | M26708 | prothymosin, alpha (gene sequence 28) | PTMA | 5757 | 9.961 | 2.2 |
| 273_g_at | K02054 | gastrin-releasing peptide | GRP | 2922 | 2.033 | 2.2 |
| 39738_at | Z82215 | myosin, heavy polypeptide 9, non-muscle | MYH9 | 4627 | 6.225 | 2.2 |
| 37874_at | Z47553 | flavin containing monooxygenase 5 | FMO5 | 2330 | 3.381 | 2.2 |
| 33400_r_at | AL080066 | chromosome 14 open reading frame 147 | C14orf147 | 171546 | 2.617 | 2.1 |
| 33447_at | X54304 | myosin regulatory light chain MRCL3 | MRCL3 | 10627 | 6.579 | 2.1 |
| 31481_s_at | M92383 | thymosin, beta 10 | TMSB10 | 9168 | 6.979 | 2.1 |
| 38126_at | J04599 | biglycan | BGN | 633 | 5.436 | 2.1 |
| 40441_g_at | AL080119 | PAI-1 mRNA-binding protein | PAI-RBP1 | 26135 | 5.398 | 2.1 |
| 40742_at | M16591 | hemopoietic cell kinase | HCK | 3055 | 5.038 | 2.1 |
| 39457_r_at | AL050355 | sorting nexin 4 | SNX4 | 8723 | 6.072 | 2.1 |
| 38428_at | M13509 | matrix metalloproteinase 1 (interstitial collagenase) | MMP1 | 4312 | 2.626 | 2.1 |
| 35178_at | W27944 | WNT inhibitory factor 1 | WIF1 | 11197 | 1.969 | 2.1 |
| 3631_f_at | U40370 | phosphodiesterase 1A, calmodulin-dependent | PDE1A | 5136 | 3.294 | 2.1 |
| 33613_at | AA806239 | hypothetical protein MGC27165 | MGC27165 | 283650 | 6.664 | 2.1 |
| 1347_at | S78187 | cell division cycle 25B | CDC25B | 994 | 6.173 | 2.1 |
| 32216_r_at | AB020685 | Rho-related BTB domain containing 3 | RHOBTB3 | 22836 | 3.430 | 2.1 |
| 40422_at | L16302 | insulin-like growth factor binding protein 2, 36kDa | IGFBP2 | 3485 | 3.735 | 2.1 |
| 34766_r_at | AL050209 | solute carrier family 25 (mitochondrial carrier; phosphate carrier), member 24 | SLC25A24 | 29957 | 3.753 | 2.1 |
| 41027_at | AF078096 | forkhead box C1 | FOXC1 | 2296 | 3.355 | 2.1 |
| 33399_at | AA142942 | chromosome 14 open reading frame 147 | C14orf147 | 171546 | 2.866 | 2.1 |
| 34898_at | M30704 | amphiregulin (schwannoma-derived growth factor) | AREG | 374 | 2.374 | 2.1 |
| 32098_at | M20777 | collagen, type VI, alpha 2 | COL6A2 | 1292 | 4.234 | 2.1 |
| 37985_at | L37747 | lamin B1 | LMNB1 | 4001 | 4.891 | 2.1 |
| 32068_at | U62027 | complement component 3a receptor 1 | C3AR1 | 719 | 6.682 | 2.1 |
| 32148_at | AI701049 | FERM, RhoGEF (ARHGEF) and pleckstrin domain protein 1 (chondrocyte-derived) | FARP1 | 10160 | 4.529 | 2.1 |
| 33483_at | X76029 | neuromedin U | NMU | 10874 | 4.382 | 2.1 |
| 38847_at | D79997 | maternal embryonic leucine zipper kinase | MELK | 9833 | 6.086 | 2.1 |
| 36140_at | J03827 | nuclease sensitive element binding protein 1 | NSE1 | 4904 | 7.848 | 2.1 |
| 32428_at | AF026564 | RNA binding motif protein, Y-linked, family 2, member B pseudogene | RBM12BP | 375850 | 4.992 | 2.1 |
| 33919_at | AF029113 | transmembrane 4 superfamily member 7 | TM4SF7 | 7106 | 2.773 | 2.1 |
| 31692_at | M59830 | heat shock 70kDa protein 1A | HSPA1A | 3303 | 3.486 | 2.1 |
| 595_at | M59465 | tumor necrosis factor, alpha-induced protein 3 | TNFAIP3 | 7128 | 5.005 | 2.1 |
| 34802_at | X15882 | collagen, type VI, alpha 2 | COL6A2 | 1292 | 4.812 | 2.1 |
| 37756_at | S59184 | RYK receptor-like tyrosine kinase | RYK | 6259 | 6.073 | 2.1 |
| 129_g_at | X82153 | cathepsin K (pycnodysostosis) | CTSK | 1513 | 3.818 | 2.1 |
| 33218_at | M11730 | v-erb-b2 erythroblastic leukemia viral oncogene homolog 2, neuro/glioblastoma derived oncogene homolog (avian) | ERBB2 | 2064 | 5.651 | 2.1 |
| 38593_r_at | KIAA0284 | KIAA0284 | KIAA0284 | 283638 | 3.681 | 2.1 |
| 33894_at | AJ010046 | neuroepithelial cell transforming gene 1 | NET1 | 10276 | 4.632 | 2.1 |
| 38750_at | U97669 | Notch homolog 3 (Drosophila) | NOTCH3 | 4854 | 3.765 | 2.0 |
| 34681_at | AB002329 | sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3E | SEMA3E | 9723 | 3.284 | 2.0 |
| 32683_at | U18271 | thymopoietin | TMPO | 7112 | 2.660 | 2.0 |
| 37558_at | U97188 | IGF-II mRNA-binding protein 3 | IMP-3 | 10643 | 4.877 | 2.0 |
| 32793_at | X00437 | Human T-cell receptor active beta-chain J/C-segment mRNA from thymus cells, clone pT-beta-2. | --- | --- | 4.605 | 2.0 |
| 238_at | M60724 | ribosomal protein S6 kinase, 70kDa, polypeptide 1 | RPS6KB1 | 6198 | 1.905 | 2.0 |
| 34937_at | AF064019 | DNA fragmentation factor, 40kDa, beta polypeptide (caspase-activated DNase) | DFFB | 1677 | 1.766 | 2.0 |
| 663_s_at | L22524 | matrix metalloproteinase 7 (matrilysin, uterine) | MMP7 | 4316 | 1.830 | 2.0 |
| 1899_at | U43746 | breast cancer 2, early onset | BRCA2 | 675 | 1.930 | 2.0 |
| 189_s_at | U09937 | plasminogen activator, urokinase receptor | PLAUR | 5329 | 5.337 | 2.0 |
| 38326_at | M69199 | putative lymphocyte G0/G1 switch gene | G0S2 | 50486 | 2.199 | 2.0 |
| 37424_at | AB029343 | chromosome 6 open reading frame 18 | C6orf18 | 54535 | 5.118 | 2.0 |
| 39350_at | U50410 | glypican 3 | GPC3 | 2719 | 2.418 | 2.0 |
| 39338_at | AI201310 | S100 calcium binding protein A10 (annexin II ligand, calpactin I, light polypeptide (p11)) | S100A10 | 6281 | 4.308 | 2.0 |
| 31420_at | AL008721 | immunoglobulin lambda variable (IV)OR22-1 | IGLV1OR22-1 | 84088 | 3.279 | 2.0 |
| 40936_at | AI651806 | cysteine-rich motor neuron 1 | CRIM1 | 51232 | 3.931 | 2.0 |
| 38737_at | M14156 | insulin-like growth factor 1 (somatomedin C) | IGF1 | 3479 | 3.701 | 2.0 |
| 37345_at | AF013759 | calumenin | CALU | 813 | 4.562 | 2.0 |
| 37302_at | U30872 | centromere protein F, 350/400kDa (mitosin) | CENPF | 1063 | 4.842 | 2.0 |
| 36890_at | AF001691 | periplakin | PPL | 5493 | 3.486 | 2.0 |
| 41745_at | X57352 | interferon induced transmembrane protein 3 (1-8U) | IFITM3 | 10410 | 6.157 | 2.0 |
| 1602_at | L33881 | protein kinase C, iota | PRKCI | 5584 | 6.157 | 2.0 |
| 1990_g_at | U43746 | breast cancer 2, early onset | BRCA2 | 675 | 4.057 | 2.0 |
| 38453_at | X15606 | intercellular adhesion molecule 2 | ICAM2 | 3384 | 4.483 | 2.0 |
| 31751_f_at | Z98744 | histone 1, H4I | HIST1H4L | 8368 | 4.615 | 2.0 |