

In the version of the supplementary information initially published online to accompany this article, a software error in the program that generates gene lists from expression signatures resulted in an incorrect listing of genes that make up the signatures for chemotherapy sensitivity (Supplementary Table 1). This error does not alter the results of the study with respect to the ability of the signatures to predict clinical response. The correct list of genes is contained in this file.

5- FU predictor

Probe Set ID	Gene Title	Chromosomal Location
152_f_at	histone 2, H4a /// histone H4/o /// similar to germinal histone H4 gene	1q21
1712_s_at	mitogen-activated protein kinase kinase 4	17p11.2
1882_g_at	---	---
31322_at	T cell receptor alpha locus	14q11.2
31726_at	gamma-aminobutyric acid (GABA) A receptor, alpha 3	Xq28
32308_r_at	collagen, type I, alpha 2	7q22.1
32318_s_at	actin, beta	7p15-p12
32610_at	PDZ and LIM domain 4	5q31.1
32755_at	actin, alpha 2, smooth muscle, aorta	10q23.3
33437_at	FtsJ homolog 1 (E. coli)	Xp11.23
33444_at	similar to neighbor of BRCA1 gene 1	17q21.31
33659_at	cofilin 1 (non-muscle)	11q13
34377_at	ATPase, Na ⁺ /K ⁺ transporting, alpha 2 (+) polypeptide	1q21-q23
34454_r_at	apolipoprotein C-IV	19q13.2
34545_at	KIAA1509	14q32.12
34843_at	zinc finger protein 516	18q23
34905_at	glutamate receptor, ionotropic, kainate 5	19q13.2
34954_r_at	phosphodiesterase 5A, cGMP-specific	4q25-q27
35056_at	arylsulfatase F	Xp22.3
35144_at	zinc finger CCCH-type containing 7B	22q13.2
35213_at	WW domain binding protein 4 (formin binding protein 21)	13q14.11
35816_at	cystatin B (stefin B)	21q22.3
35929_s_at	testis specific protein, Y-linked 1 /// testis specific protein,	Yp11.2
36245_at	5-hydroxytryptamine (serotonin) receptor 2B	2q36.3-q37.1
36453_at	kelch repeat and BTB (POZ) domain containing 11	8p23.3
36549_at	solute carrier family 25 (mitochondrial carrier; peroxisomal membrane protein, 34kD:	22q13.2
37349_r_at	high mobility group nucleosomal binding domain 3	6q14.1
37361_at	fibroblast growth factor (acidic) intracellular binding protein	11q13.1
37437_at	intraflagellar transport 140 homolog (Chlamydomonas)	16p13.3
37802_r_at	family with sequence similarity 63, member B	15q22.1
37860_at	zinc finger protein 337	20p11.21
39783_at	KIAA0100	17q11.2
39898_at	family with sequence similarity 13, member C1	10q21.1
40104_at	serine/threonine kinase 25 (STE20 homolog, yeast)	2q37.3
40452_at	copine I	20q11.22
40471_at	peroxisomal biogenesis factor 19	1q22
40536_f_at	Eukaryotic translation initiation factor 5B	2p11.1-q11.1
40886_at	eukaryotic translation elongation factor 1 alpha 1 /// apolipoprotein L domain containi	6q14.1 /// 12p13.1 /// 1p21.3
40983_s_at	serine racemase	17p13
41058_g_at	thioesterase superfamily member 2	6p22.2
41536_at	Inhibitor of DNA binding 4, dominant negative helix-loop-helix protein	6p22-p21
41868_at	gamma-glutamyltransferase 1 /// gamma-glutamyltransferase-like 4	22q11.23 /// 22q11.22
427_f_at	interferon, alpha 10	9p22
429_f_at	tubulin, beta 2A /// tubulin, beta 4 /// tubulin, beta 2B	6p25 /// 19p13.3
471_f_at	tubulin, beta 3	16q24.3

Adriamycin Predictor

Probe Set ID	Gene Title	Chromosomal Location
101_at	dual-specificity tyrosine-(Y)-phosphorylation regulate	12p13.32
1051_g_at	melan-A	9p24.1
1259_at	zinc finger protein 19 /// zinc finger protein 23 (KOX 1	16q22
1319_at	discoïdin domain receptor family, member 2	1q12-q23
1519_at	v-ets erythroblastosis virus E26 oncogene homolog 2	21q22.3 21q22.2
1537_at	epidermal growth factor receptor (erythroblastic leuk	7p12
1848_at	RAP1A, member of RAS oncogene family	1p13.3
1911_s_at	growth arrest and DNA-damage-inducible, alpha	1p31.2-p31.1
191_at	mucin 8	12q24.3
2011_s_at	BCL2-interacting killer (apoptosis-inducing)	22q13.31
2035_s_at	enolase 1, (alpha)	1p36.3-p36.2
32139_at	zinc finger protein 185 (LIM domain)	Xq28
32168_s_at	Down syndrome critical region gene 1	21q22.1-q22.2 21q22.12
32612_at	gelsolin (amyloidosis, Finnish type)	9q33
32718_at	tyrosylprotein sulfotransferase 1	7q11.21
32821_at	lipocalin 2 (oncogene 24p3)	9q34
32967_at	Fas apoptotic inhibitory molecule 3	1q32.1
33004_g_at	NCK adaptor protein 2	2q12
33240_at	PDZ domain containing RING finger 3	3p13
33409_at	FK506 binding protein 2, 13kDa	11q13.1-q13.3
33824_at	keratin 8	12q13
33853_s_at	neuropilin 2	2q33.3
33892_at	plakophilin 2	12p11
33904_at	claudin 3	7q11.23
33908_at	calpain 1, (mu/l) large subunit	11q13
33942_s_at	syntaxin binding protein 1	9q34.1
33956_at	lymphocyte antigen 96	8q21.11
34213_at	WW, C2 and coiled-coil domain containing 1	5q35.1
34303_at	chromosome 10 open reading frame 56	10q22.3
34348_at	serine peptidase inhibitor, Kunitz type, 2	19q13.1
34859_at	melanoma antigen family D, 2	Xp11.2
34885_at	synaptogyrin 2	17q25.3
34993_at	sarcoglycan, delta (35kDa dystrophin-associated gly	5q33-q34
35280_at	laminin, gamma 2	1q25-q31
35444_at	chromosome 19 open reading frame 21	19p13.3
35681_r_at	zinc finger homeobox 1b	2q22
35766_at	keratin 18	12q13
35807_at	cytochrome b-245, alpha polypeptide	16q24
36133_at	desmoplakin	6p24
36618_g_at	inhibitor of DNA binding 1, dominant negative helix-lc	20q11
36795_at	prosaposin (variant Gaucher disease and variant me	10q21-q22 /// 17p13.1
36826_at	general transcription factor IIF, polypeptide 1, 74kDa	19p13.3
37327_at	epidermal growth factor receptor (erythroblastic leuk	7p12
37345_at	calumenin	7q32
37695_at	ring finger protein 144	2p25.2-p25.1
37743_at	fasciculation and elongation protein zeta 1 (zygin I)	11q24.2
37749_at	mesoderm specific transcript homolog (mouse)	7q32
37926_at	Kruppel-like factor 5 (intestinal)	13q22.1
38004_at	chondroitin sulfate proteoglycan 4 (melanoma-assoc	15q24.2
38078_at	filamin B, beta (actin binding protein 278)	3p14.3
38110_at	syndecan binding protein (syntenin)	8q12
38119_at	glycophorin C (Gerbich blood group)	2q14-q21
38122_at	solute carrier family 23 (nucleobase transporters), m	20p13
38297_at	phosphatidylinositol transfer protein, membrane-assc	11q13
38379_at	glycoprotein (transmembrane) nmb	7p15
38653_at	peripheral myelin protein 22	17p12-p11.2

39214_at	plexin B3 /// SFRS protein kinase 3	Xq28
39271_at	melanoma inhibitory activity	19q13.32-q13.33
39316_at	RAB40C, member RAS oncogene family	16p13.3
39386_at	MAD2L1 binding protein	6p21.1
39801_at	procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3	7q22
40088_at	nuclear receptor interacting protein 1	21q11.2
40103_at	villin 2 (ezrin)	6q25.2-q26
40202_at	Kruppel-like factor 9	9q13
40434_at	podocalyxin-like	7q32-q33
40568_at	ATPase, H+ transporting, lysosomal 56/58kDa, V1 s1	8p22-p21
40926_at	solute carrier family 6 (neurotransmitter transporter, c)	Xq28
41158_at	proteolipid protein 1 (Pelizaeus-Merzbacher disease, Xq22	Xq22
41294_at	keratin 7	12q12-q13
41359_at	plakophilin 3	11p15
41378_at	MRNA from chromosome 5q31-33 region	---
41453_at	discs, large homolog 3 (neuroendocrine-dlg, Drosophila)	Xq13.1
41503_at	zinc fingers and homeoboxes 2	8q24.13
41610_at	laminin, alpha 5	20q13.2-q13.3
41644_at	SAM and SH3 domain containing 1	6q24.3
41839_at	growth arrest-specific 1	9q21.3-q22
575_s_at	tumor-associated calcium signal transducer 1	2p21
661_at	growth arrest-specific 1	9q21.3-q22
953_g_at	---	---
999_at	cytochrome P450, family 27, subfamily A, polypeptide	2q33-qter

Docetaxel Predictor

Probe Set ID	Gene Title	Chromosomal Location
1259_at	zinc finger protein 19 /// zinc finger protein 23 (KOX 16)	16q22
1410_at	vitamin D (1,25- dihydroxyvitamin D3) receptor	12q13.11
1567_at	fms-related tyrosine kinase 1 (vascular endothelial growth factor receptor)	13q12
1752_at	calreticulin	19p13.3-p13.2
1804_at	kallikrein 3, (prostate specific antigen)	19q13.41
1879_at	related RAS viral (r-ras) oncogene homolog	19q13.3-qter
1998_i_at	BCL2-associated X protein	19q13.3-q13.4
2086_s_at	TYRO3 protein tyrosine kinase	15q15.1-q21.1
31432_g_at	Fc fragment of IgG, receptor, transporter, alpha	19q13.3
31433_at	olfactory receptor, family 2, subfamily J, member 2	6p21.31-22.2
31639_f_at	SRY (sex determining region Y)-box 1	13q34
32085_at	phosphatidylinositol-3-phosphate/phosphatidylinositol 5-kinase, type III	2q33.3
321_at	Wiskott-Aldrich syndrome-like	7q31.3
32218_at	CDNA: FLJ22515 fis, clone HRC12122,	---
32238_at	bridging integrator 1	2q14
32524_s_at	Junctional adhesion molecule 3	11q25
32828_at	branched chain ketoacid dehydrogenase kinase	16p11.2
32844_at	eukaryotic translation initiation factor 4 gamma, 1	3q27-qter
33048_at	EF-hand calcium binding protein 2	16q23.3
33134_at	adenylate cyclase 3	2p24-p22
33204_at	Forkhead box D1	5q12-q13
33215_g_at	mitochondrial ribosomal protein S12	19q13.1-q13.2
33286_at	endonuclease G	9q34.1
33372_at	RAB31, member RAS oncogene family	18p11.3
33388_at	testis expressed sequence 261	2p13.3
33434_at	BET1 homolog (S. cerevisiae)	7q21.1-q22
34647_at	DEAD (Asp-Glu-Ala-Asp) box polypeptide 5	17q21
34773_at	tubulin-specific chaperone a	5q14.1
34801_at	ubiquitin specific peptidase 52	12q13.2-q13.3
34804_at	Solute carrier family 25, member 36	3q23
35018_at	calcium binding protein P22	15q13.3
35655_at	ankyrin repeat domain 28	3p24.3
35714_at	pyridoxal (pyridoxine, vitamin B6) kinase	21q22.3
35770_at	ATPase, H+ transporting, lysosomal accessory protein 1	Xq28
35815_at	SET domain containing 2	3p21.31
36209_at	bromodomain containing 2	6p21.3
36250_at	aspartate beta-hydroxylase domain containing 1	16p11.2
36395_at	Transcribed locus	---
36528_at	argininosuccinate lyase	7cen-q11.2
36642_at	pepsinogen 5,	11q13 /// 11q12.2
36663_at	natriuretic peptide precursor A	1p36.21
38831_f_at	guanine nucleotide binding protein (G protein), beta polypeptide 2	7q21.3-q22.1 7q22
39199_at	activin A receptor, type IB	12q13
40568_at	ATPase, H+ transporting, lysosomal 56/58kDa, V1 subunit B2	8p22-p21
41063_g_at	polycomb group ring finger 1	2p13.1
41077_at	hypothetical protein LOC643641	7q36.1
41285_at	inositol polyphosphate-5-phosphatase, 40kDa	10q26.3
41689_at	plasma membrane proteolipid (plasmolipin)	16q13
41713_at	zinc finger with KRAB and SCAN domains 1	7q21.3-q22.1
941_at	proteasome (prosome, macropain) subunit, beta type, 6	17p13

Etoposide Predictor

Probe Set ID	Gene Title	Chromosomal Location
1015_s_at	LIM domain kinase 1	7q11.23
1188_g_at	ligase III, DNA, ATP-dependent	17q11.2-q12
1233_s_at	AXL receptor tyrosine kinase	19q13.1
1456_s_at	interferon, gamma-inducible protein 16	1q22
1590_s_at	v-Ha-ras Harvey rat sarcoma viral oncogene homolog	11p15.5
1670_at	transcription factor Dp-1	13q34
1704_at	vav 2 oncogene	9q34.1
1963_at	fms-related tyrosine kinase 1	13q12
2047_s_at	junction plakoglobin	17q21
296_at	---	---
297_g_at	---	---
311_s_at	---	---
31719_at	fibronectin 1	2q34
31720_s_at	fibronectin 1	2q34
32378_at	pyruvate kinase, muscle	15q22
32387_at	lysophospholipase 3 (lysosomal phospholipase A2)	16q22.1
32593_at	raft-linking protein	3p24.3
33282_at	ladinin 1	1q25.1-q32.3
33448_at	serine peptidase inhibitor, Kunitz type 1	15q15.1
33904_at	claudin 3	7q11.23
34320_at	polymerase I and transcript release factor	17q21.31
34348_at	serine peptidase inhibitor, Kunitz type, 2	19q13.1
34747_at	matrix metalloproteinase 14 (membrane-inserted)	14q11-q12
34769_at	fatty acid amide hydrolase	1p35-p34
35276_at	claudin 4	7q11.23
35309_at	suppression of tumorigenicity 14 (colon carcinoma)	11q24-q25
35444_at	chromosome 19 open reading frame 21	19p13.3
35541_r_at	KIAA0506 protein	---
35630_at	MAP-kinase activating death domain	17q24-q25 /// 11p11.2
35669_at	cordon-bleu homolog (mouse)	7p12.1
35681_r_at	zinc finger homeobox 1b	2q22
35735_at	guanylate binding protein 1, interferon-inducible, 67kDa	1p22.2
36097_at	immediate early response 2	19p13.13
36890_at	periplakin	16p13.3
37934_at	transmembrane protein 30B	14q23.1
38221_at	connector enhancer of kinase suppressor of Ras 1	1p36.11
38482_at	claudin 7	17p13
38759_at	butyrophilin, subfamily 3, member A2	6p22.1
38760_f_at	butyrophilin, subfamily 3, member A2	6p22.1
39331_at	tubulin, beta 2A	6p25
39732_at	microtubule-associated protein 7	6q23.3
39870_at	Testes-specific heterogenous nuclear ribonucleoprotein G	11p15
40215_at	UDP-glucose ceramide glucosyltransferase	9q31
40225_at	cyclin G associated kinase	4p16
41359_at	plakophilin 3	11p15
41872_at	deafness, autosomal dominant 5	7p15
479_at	disabled homolog 2, mitogen-responsive phosphoprotein	5p13
575_s_at	tumor-associated calcium signal transducer 1	2p21
671_at	secreted protein, acidic, cysteine-rich (osteonectin)	5q31.3-q32
903_at	protein phosphatase 2, regulatory subunit B (B56), alpha i	1q32.2-q32.3

Paclitaxel Predictor

Probe	Description	Chromosomal location
1228_s_at	CTAGE family, member 5	14q13.3
1356_at	death associated protein 3	1q21-q22
243_g_at	microtubule-associated protein 4	3p21
250_at	nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3	16q22.2
31463_s_at	heterogeneous nuclear ribonucleoprotein A1	12q13.1
31511_at	ribosomal protein S9	19q13.4
31546_at	ribosomal protein L18	19q13
31867_at	ARTC1 mRNA, complete sequence	3q13.1-q13.12
32226_at	microtubule-associated protein 4	3p21
32378_at	pyruvate kinase, muscle	15q22
33452_at	plasminogen activator, tissue	8p12
34091_s_at	vimentin	10p13
34246_at	chromosome 6 open reading frame 145	6p25.2
35352_at	aryl-hydrocarbon receptor nuclear translocator 2	15q24
356_at	kinesin family member 22	16p11.2
35763_at	neurobeachin-like 2	3p21.31
36192_at	secernin 1	7p14.3-p14.1
36989_at	dystroglycan 1 (dystrophin-associated glycoprotein 1)	3p21
37345_at	calumenin	7q32
37747_at	annexin A5	4q26-q28 4q28-q32
38120_at	polycystic kidney disease 2 (autosomal dominant)	4q21-q23
38405_at	fragile X mental retardation, autosomal homolog 1	3q28
38479_at	acidic (leucine-rich) nuclear phosphoprotein 32 family, member B	9q22.32
39019_at	lysosomal-associated protein transmembrane 4 alpha	2p24.1
39077_at	DR1-associated protein 1 (negative cofactor 2 alpha)	11q13.3
39170_at	CD59 molecule, complement regulatory protein	11p13
39750_at	zinc finger, DHHC-type containing 3	3p21.31
39798_at	ribosomal protein S28	19p13.2
40255_at	DEAD (Asp-Glu-Ala-Asp) box polypeptide 28	16q22.1
40822_at	nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3	16q22.2
40865_at	thymine-DNA glycosylase	12q24.1
41403_at	small nuclear ribonucleoprotein polypeptide F	12q22
41758_at	chromosome 22 open reading frame 5	22q12
41834_g_at	jumping translocation breakpoint	1q21
591_s_at	protein tyrosine phosphatase, non-receptor type 1	20q13.1-q13.2
798_at	primase, polypeptide 1, 49kDa	12q13

36 Genes

Topotecan Predictor

Probe ID	Gene Title	Chromosomal Location
1005_at	dual specificity phosphatase 1	5q34
116_at	protein kinase, cAMP-dependent, regulatory, type II, alpha	3p21.3-p21.2
1233_s_at	AXL receptor tyrosine kinase	19q13.1
1251_g_at	RAP1 GTPase activating protein	1p36.1-p35
1257_s_at	quiescin Q6	1q24
1278_at	---	---
1368_at	interleukin 1 receptor, type I	2q12
1385_at	transforming growth factor, beta-induced, 68kDa	5q31
1491_at	pentraxin-related gene, rapidly induced by IL-1 beta	3q25
1544_at	Bloom syndrome	15q26.1
1563_s_at	tumor necrosis factor receptor superfamily, member 1A	12p13.2
1593_at	fibroblast growth factor 2 (basic)	4q26-q27
160_at	signal transducing adaptor molecule (SH3 domain and ITAM motif) 1	10p14-p13
160044_g_at	aconitase 2, mitochondrial	22q11.2-q13.31 22q13.2-q13.31
1751_g_at	phenylalanine-tRNA synthetase-like, alpha subunit	19p13.2
1783_at	Ras and Rab interactor 2	20p11.22
1828_s_at	fibroblast growth factor 2 (basic)	4q26-q27
1879_at	related RAS viral (r-ras) oncogene homolog	19q13.3-qter
1958_at	c-fos induced growth factor (vascular endothelial growth factor D)	Xp22.31
2042_s_at	v-myb myeloblastosis viral oncogene homolog (avian)	6q22-q23
2053_at	cadherin 2, type 1, N-cadherin (neuronal)	18q11.2
2056_at	fibroblast growth factor receptor 1	8p11.2-p11.1
2057_g_at	fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2)	8p11.2-p11.1
232_at	laminin, gamma 1 (formerly LAMB2)	1q31
31521_f_at	histone 1, H4k /// histone 1, H4j	6p22-p21.3
32098_at	collagen, type VI, alpha 2	21q22.3
32116_at	transmembrane channel-like 6	17q25.3
32260_at	phosphoprotein enriched in astrocytes 15	1q21.1
32434_at	myristoylated alanine-rich protein kinase C substrate	6q22.2
32529_at	cytoskeleton-associated protein 4	12q23.3
32531_at	gap junction protein, alpha 1, 43kDa (connexin 43)	6q21-q23.2
32535_at	fibrillin 1	15q21.1
32606_at	Transcribed locus	---
32607_at	brain abundant, membrane attached signal protein 1	5p15.1-p14
32673_at	butyrophilin, subfamily 2, member A1	6p22.1
32808_at	integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29)	10p11.2
32812_at	hypothetical protein	4p13
32847_at	myosin, light polypeptide kinase	3q21
33127_at	lysyl oxidase-like 2	8p21.3-p21.2
33328_at	HEG homolog 1 (zebrafish)	3q21.2
33337_at	degenerative spermatocyte homolog 1, lipid desaturase (Drosophila)	1q42.11
33404_at	CAP, adenylate cyclase-associated protein, 2 (yeast)	6p22.3
33405_at	CAP, adenylate cyclase-associated protein, 2 (yeast)	6p22.3
33440_at	---	---
33772_at	prostaglandin E receptor 4 (subtype EP4)	5p13.1
33785_at	brain-specific angiogenesis inhibitor 2	1p35
33787_at	NUAK family, SNF1-like kinase, 1	12q23.3
33791_at	deleted in lymphocytic leukemia, 1 /// SPANX family, member C	13q14.3 /// Xq27.1
33800_at	adenylate cyclase 9	16p13.3
33882_at	RAB11 family interacting protein 5 (class I)	2p13-p12
33994_g_at	myosin, light polypeptide 6, alkali, smooth muscle and non-muscle	12q13.2
34091_s_at	vimentin	10p13
34106_at	guanine nucleotide binding protein (G protein) alpha 12	7p22.2
34218_at	SPEG complex locus	2q35
34320_at	polymerase I and transcript release factor	17q21.31
34375_at	chemokine (C-C motif) ligand 2	17q11.2-q12

34795_at	procollagen-lysine, 2-oxoglutarate 5-dioxygenase 2	3q23-q24
34802_at	collagen, type VI, alpha 2	21q22.3
34811_at	ATP synthase, H+ transporting, mitochondrial F0 complex	2q31.1
35130_at	glutathione reductase	8p21.1
35264_at	NADH dehydrogenase (ubiquinone)	11p11.11
35309_at	suppression of tumorigenicity 14 (colon carcinoma)	11q24-q25
35366_at	nidogen 1	1q43
35729_at	myosin ID	17q11-q12
36057_at	armadillo repeat containing, X-linked 2	Xq21.33-q22.2
36119_at	caveolin 1, caveolae protein, 22kDa	7q31.1
36149_at	dihydropyrimidinase-like 3	5q32
36369_at	polymerase I and transcript release factor	17q21.31
36525_at	F-box and leucine-rich repeat protein 2	3p22.3
36550_at	Ras and Rab interactor 2	20p11.22
36577_at	pleckstrin homology domain containing, family C	14q22.2
36638_at	connective tissue growth factor	6q23.1
36659_at	collagen, type IV, alpha 2	13q34
36790_at	tropomyosin 1 (alpha)	15q22.1
36791_g_at	tropomyosin 1 (alpha)	15q22.1
36792_at	tropomyosin 1 (alpha)	15q22.1
36799_at	frizzled homolog 2 (Drosophila)	17q21.1
36811_at	lysyl oxidase-like 1	15q24-q25 15q22
36885_at	spleen tyrosine kinase	9q22
36952_at	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/en	2p23
36988_at	tumor necrosis factor, alpha-induced protein 1 (endothelial)	17q22-q23
37032_at	nicotinamide N-methyltransferase	11q23.1
37322_s_at	hydroxyprostaglandin dehydrogenase 15-(NAD)	4q34-q35
37408_at	mannose receptor, C type 2	17q23.2
37486_f_at	Meis1 homolog 3 (mouse) pseudogene 1	17p12
37599_at	aldehyde oxidase 1	2q33
377_g_at	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (ser	7q21-q31
378_s_at	GPI anchored molecule like protein	8q24.3
38113_at	spectrin repeat containing, nuclear envelope 1	6q25
38125_at	serpin peptidase inhibitor, clade E (nexin, plasminogen activator inhibitor type	7q21.3-q22
38299_at	interleukin 6 (interferon, beta 2)	7p21
38338_at	related RAS viral (r-ras) oncogene homolog	19q13.3-qter
38394_at	glycerol-3-phosphate dehydrogenase 1-like	3p22.3
38396_at	3'UTR of hypothetical protein (ORF1)	---
38433_at	AXL receptor tyrosine kinase	19q13.1
38449_at	WD repeat domain 23	14q12
38482_at	claudin 7	17p13
38488_s_at	Interleukin 15	4q31
38631_at	tumor necrosis factor, alpha-induced protein 2	14q32
38772_at	cysteine-rich, angiogenic inducer, 61	1p31-p22
38775_at	low density lipoprotein-related protein 1	12q13-q14
38842_at	angiomin like 2	3q21-q22
38921_at	phosphodiesterase 1B, calmodulin-dependent	12q13
39000_at	N-myristoyltransferase 1	17q21.31
39254_at	retinoic acid induced 14	5p13.3-p13.2
39277_at	---	---
39327_at	peroxidasin homolog (Drosophila)	2p25
39333_at	collagen, type IV, alpha 1	13q34
39409_at	complement component 1	12p13 /// 12p13.31
39614_at	KIAA0802 /// chromosome 21 open reading frame 57	18p11.22 /// 21q22.3
39710_at	chromosome 5 open reading frame 13	5q22.1
39867_at	Tu translation elongation factor, mitochondrial	16p11.2
39901_at	EGF-like repeats and discoidin I-like domains 3	5q14
40023_at	brain-derived neurotrophic factor	11p13

40078_at	protease, serine, 23	11q14.1
40096_at	ATP synthase, H ⁺ transporting, mitochondrial F1 complex	18q12-q21
40171_at	frequently rearranged in advanced T-cell lymphomas 2	10q24.1
40341_at	chromosome 16 open reading frame 51	16p13.13-p12.3
40497_at	tumor suppressor candidate 4	3p21.3
40564_at	nucleoporin 50kDa	22q13.31
40567_at	tubulin, alpha 3	12q12-12q14.3
40642_at	nuclear factor I/B	9p24.1
40692_at	transducin-like enhancer of split 4 (E(sp1) homolog, Drosophila)	9q21.31
40781_at	V-akt murine thymoma viral oncogene homolog 3 (protein kinase B, gamma)	1q43-q44
40936_at	cysteine rich transmembrane BMP regulator 1 (chordin-like)	2p21
41097_at	telomeric repeat binding factor 2	16q22.1
41223_at	cytochrome c oxidase subunit Va	15q25
41236_at	Smith-Magenis syndrome chromosome region, candidate 7-like	22q13
41273_at	matrix-remodelling associated 7	17q25.1-q25.2
41295_at	START domain containing 7	2q11.2
41354_at	stanniocalcin 1	8p21-p11.2
41478_at	tetratricopeptide repeat domain 28	22q12.1
41544_at	polo-like kinase 2 (Drosophila)	5q12.1-q13.2
41667_s_at	TDP-glucose 4,6-dehydratase	13q32.1
41738_at	caldesmon 1	7q33
41744_at	optineurin	10p13
41745_at	interferon induced transmembrane protein 3 (1-8U)	11p15.5
41872_at	deafness, autosomal dominant 5	7p15
424_s_at	fibroblast growth factor receptor 1	8p11.2-p11.1
465_at	HIV-1 Tat interacting protein, 60kDa	11q13
548_s_at	spleen tyrosine kinase	9q22
581_at	laminin, beta 1	7q22
628_at	frizzled homolog 2 (Drosophila)	17q21.1
672_at	serpin peptidase inhibitor, clade E	7q21.3-q22
867_s_at	thrombospondin 1	15q15
875_g_at	chemokine (C-C motif) ligand 2	17q11.2-q12
884_at	integrin, alpha 3 (antigen CD49C, alpha 3 subunit of VLA-3 receptor)	17q21.33
885_g_at	integrin, alpha 3 (antigen CD49C, alpha 3 subunit of VLA-3 receptor)	17q21.33
890_at	ubiquitin-conjugating enzyme E2A (RAD6 homolog)	Xq24-q25
919_at	---	---