

AffyProbe Set	LocusLink	ABREV	Gene Class/Name	Normal	Tumor	FoldDiff	DeltaGPT	Permax	Chromosome
Cell Cycle									
D13639_at	894	CCND2	cyclin D2	1766	175	10.1	1590.5	0.468	12p13
U15642_s_at	1875	E2F5	E2F transcription factor 5, p130-binding	33	146	4.4	113.1	0.2448	8
Z24725_at	10979	MIG2	mitogen inducible 2	818	205	4.0	613.2	0.431	14
X98260_at	27000	ZRF1	zuotin related factor 1	349	25	13.9	323.7	0.1508	7q22-q32
Cell-Cell interaction									
D13626_at	9934	KIAA0001	G protein coupled receptor for UDP-glucose	145	20	7.2	124.9	0.3447	3q21-q25
J04456_at	3956	LGALS1	lectin, galactoside-binding, soluble, 1 (galectin 1)	8915	1537	5.8	7378.3	0.2478	22q13.1
L36033_at	6387	SDF1	stromal cell-derived factor 1	677	49	13.9	628.3	0.3417	10q11.1
M93036_at	4072	TACSTD1	tumor-associated calcium signal transducer 1	583	2192	3.8	1609.5	0.1299	4q
Cytoskeleton/adhesion									
D21255_at	1009	CDH11	cadherin 11 (OB-cadherin, osteoblast)	504	57	8.8	446.4	0.5	16q22.1
U83192_at	1742	DLG4	discs, large (Drosophila) homolog 4	278	20	13.9	258.1	0.028	17p13.1
HG4535-HT4940_s_at	2039	EPB49	erythrocyte membrane protein band 4.9 (dematin)	242	1321	5.5	1078.6	0.2448	8p21.1
M36284_s_at	2995	GYPC	glycophorin C (Gerbich blood group)	330	20	16.5	310.3	0.014	2q14-q21
X12876_s_at	3875	KRT18	keratin 18	1115	4151	3.7	3035.7	0.1728	12q13
X74929_s_at	3856	KRT8	keratin 8	521	5338	10.2	4816.8	0.0669	12q13
L14837_at	7082	TJP1	tight junction protein 1 (zona occludens 1)	988	321	3.1	666.3	0.2228	15q13
Extracellular Matrix									
J04177_at	1301	COL11A1	collagen, type XI, alpha 1	287	32	8.9	255.1	0.2587	1p21
Y11710_ma1_at	7373	COL14A1	collagen, type XIV, alpha 1; undulin	446	33	13.3	412.2	0.1938	8q23
Z48481_at	4323	MMP14	matrix metalloproteinase 14 (membrane-inserted)	844	46	18.2	797.4	0.2038	14q11-q12
M30269_at	4811	NID	nidogen (entactin)	1204	54	22.3	1149.7	0.2458	1q43
M24486_s_at	5033	P4HA1	procollagen-proline, 2-oxoglutarate 4-dioxygenase, alpha 1	41	245	6.0	204.3	0.2987	10q21.3-q23.1
L33799_at	5118	PCOLCE	procollagen C-endopeptidase enhancer	4331	353	12.3	3977.7	0.2038	7q22
Hormone Response/Pathway									
D11151_at	1909	EDNRA	endothelin receptor type A	656	58	11.2	597.5	0.446	4
X97249_at	2491	FSHPRH1	FSH primary response (LRPR1, rat) homolog 1	277	32	8.7	245.1	0.2128	Xq22
X77307_at	3357	HTR2B	5-hydroxytryptamine (serotonin) receptor 2B	434	39	11.2	395.1	0.2388	2q37.1-q36.3
HG3510-HT3704_at	7025	NR2F1	nuclear receptor subfamily 2, group F, member 1	793	68	11.6	724.8	0.3946	5q14
HG721-HT4827_s_at	5047	PAEP(alt2)	progesterone-associ endometrial prot.: Alt. Splice 2	4998	32	158.2	4966.1	0.2478	9q34
Metabolism, amino acid									
D88213_at	314	AOC2	amine oxidase, copper containing 2 (retina-specific)	177	20	8.9	157.3	0.042	17q21
U61263_at	10994	ILVBL	ilvB (bacterial acetolactate synthase)-like	501	81	6.2	420.4	0.5	19p13.1
M16447_at	5860	QDPR	quinoid dihydropteridine reductase	233	20	11.6	212.7	0.0789	4p15.31
M12625_at	3931	LCAT	lecithin-cholesterol acyltransferase	873	20	43.7	853.4	0.035	16q22.1
D00860_at	5631	PRPS1	phosphoribosyl pyrophosphate synthetase 1	150	29	5.1	120.6	0.2068	Xq21-q27
Y00971_at	5634	PRPS2	phosphoribosyl pyrophosphate synthetase 2	204	20	10.2	183.7	0.1099	Xp22.3-p22.2
Miscellaneous									
L76568_xpt3_f_at	2072	ERCC4	excision repair cross-complementing rodent repair deficiency	20	509	25.4	489.0	0.0959	16p13.3-p13.11
J00212_f_at	3452	IFNA21	interferon, alpha 21	236	30	7.9	206.0	0.2108	9p22
X15949_at	3660	IRF2	interferon regulatory factor 2	208	20	10.4	188.0	0.1808	4q34.1-q35.1
Oncogene/Tumor Suppressor									
HG162-HT3165_at	558	AXL (as2)	AXL receptor tyrosine kinase (Alternate Splice 2)	377	32	11.6	344.8	0.2218	19q13.1
M91083_at	8045	C11ORF13	chromosome 11 open reading frame 13	20	918	45.9	898.1	0.2727	11p15.5
X68277_at	1843	DUSP1	dual specificity phosphatase 1	4140	640	6.5	3499.8	0.426	5q34
X57025_at	3479	IGF1	insulin-like growth factor 1 (somatomedin C)	1490	35	42.3	1454.6	0.1618	12q22-q23
M21574_at	5156	PDGFRA	platelet-derived growth factor receptor, alpha polypeptide	1167	65	17.8	1101.4	0.3077	4q11-q13
X69950_s_at	51352	WIT-1	Wilms tumor associated protein	260	20	13.0	239.8	0.038	11p13
Toxin Scavenger/Activator									
U03688_at	1545	CYP1B1	cytochrome P450, subfamily I, polypeptide 1	179	20	8.9	158.8	0.2038	2p21
M13450_at	2098	ESD	esterase D/formylglutathione hydrolase	1103	353	3.1	750.5	0.3057	13q14.1-q14.2
Z11793_at	6414	SEPP1	selenoprotein P, plasma, 1	408	110	3.7	297.9	0.1988	5q31
Vesicle/synapse associated									
U49082_at	10991	G17	G17 transporter protein	20	781	39.1	761.3	0.3247	3p21.3
M85276_at	10578	GNLY	granulysin	3585	64	55.8	3521.2	0.2218	2p12-q11
U50078_at	8925	HERC1	hect domain and RCC1-like domain 1	149	20	7.5	129.1	0.1359	15q22
U44111_at	3176	HNMT	histamine N-methyltransferase	205	20	10.3	185.2	0.0599	2
X98001_at	5876	RABGGTB	Rab geranylgeranyltransferase, beta subunit	204	37	5.6	167.7	0.4076	1p31-p22
X92396_at	6845	SYBL1	synaptobrevin-like 1	245	30	8.3	215.8	0.2218	Xq28