

Probe set	Gene	Accession	Lymphoma #							
			1	2	3	4	5	6	7	8
			Fold upregulation							
1457917_at	complement component 3	K02782	4.5	20.3	37.2	34.8	11.6	4.7	16.9	77.1
1440721_at	Mus musculus cDNA clone IMAGE:30046293, partial cds	AV111366	4.2	13.2	27.7	14.5	23.9	23.1	17.0	33.4
1458058_at	chemokine (C-C motif) ligand 6	BC002073	12.0	17.0	15.0	9.9	5.4	19.8	7.6	42.3
1458309_at	serine (or cysteine) proteinase inhibitor, clade E, member 2	NM_009255	8.6	17.6	16.3	12.4	16.6	15.6	13.1	20.2
1439412_at	POU domain, class 2, associating factor 1	NM_011136	8.2	13.3	20.5	8.5	26.6	23.0	3.3	15.7
1427655_a_at	insulin-like growth factor binding protein 7	AI790558	11.2	18.4	5.7	14.1	19.1	7.1	22.2	14.9
1440188_at	dopa decarboxylase	AF071068	4.7	20.7	4.2	6.1	15.7	19.5	17.4	20.5
1444942_at	ras homolog gene family, member AB	BC018275	22.9	24.8	14.1	9.3	5.4	12.6	6.7	11.6
1429254_at	H19 fetal liver mRNA	NM_023123	29.7	16.2	1.2	8.6	20.7	4.8	6.4	18.3
1427532_at	serum amyloid A 3	NM_011315	2.8	17.7	11.9	5.8	8.2	14.2	9.4	32.7
1439315_at	nuclear factor, erythroid derived 2, like 3	NM_010903	35.0	21.2	3.7	9.6	4.0	2.9	4.8	18.1
1445148_at	hormonally upregulated Neu-associated kinase	NM_015755	19.4	13.0	7.2	14.4	4.6	12.5	12.0	9.4
1457063_at	Mus musculus transcribed sequences	AI450842	1.6	3.2	33.6	16.6	15.7	1.6	7.8	10.6
1420166_at	Mus musculus transcribed sequences	AU067644	7.2	6.5	14.2	12.0	11.1	7.0	8.0	11.8
1418065_at	Fc receptor-like 3	BC027310	1.7	14.7	4.2	4.0	2.1	20.2	1.2	28.2
1438837_at	Mus musculus transcribed sequences	AV222628	5.1	12.6	16.1	5.9	12.1	12.9	0.8	7.9
1458050_at	CD5 antigen-like	NM_009690	2.1	9.3	6.7	5.2	3.4	11.5	4.8	30.0
1447792_x_at	serine (or cysteine) proteinase inhibitor, clade G, member 1	NM_009776	3.0	6.4	14.7	11.6	4.5	2.4	4.0	24.0
1421604_a_at	adenylate kinase 5	AB060081	14.3	11.2	7.2	7.5	8.0	7.2	6.5	5.1
1443894_at	lamin A	NM_019390	8.4	7.9	12.4	8.3	7.1	12.6	4.3	4.8
1451474_a_at	Fc receptor, IgG, low affinity III	NM_010188	4.1	8.8	8.3	5.5	3.5	5.9	3.4	22.7
1457141_at	immunoglobulin joining chain	BC006026	1.2	9.8	22.6	10.7	3.9	0.7	4.3	9.2
1447214_at	myelocytomatosis oncogene	BC006728	10.1	8.0	5.0	7.5	6.5	7.9	7.9	5.8
1439686_at	RIKEN cDNA D330017J20 gene	BB098407	2.7	7.8	5.1	1.9	15.3	13.7	2.5	8.1
1417481_at	phospholipase A2, group IID	AF169407	1.3	6.3	12.1	9.3	3.4	0.9	5.7	17.9
1419028_at	fibrinogen-like protein 2	NM_008013	0.7	7.7	3.3	2.9	17.7	9.8	1.4	11.6
1448929_at	RIKEN cDNA 3110007P09 gene	BC017596	14.1	10.2	8.7	4.2	4.1	1.4	4.3	7.1
1438894_at	cDNA clone MGC:51479 IMAGE:4012958, complete cds	BB404446	9.9	7.1	3.8	4.8	6.8	10.3	4.4	6.2
1439897_at	transferrin receptor	NM_011638	8.2	4.9	7.3	6.8	6.2	3.4	5.2	9.4
1418776_at	apolipoprotein E	AK019319	2.2	4.5	7.6	7.0	5.5	7.6	4.1	12.5
1419480_at	cDNA clone MGC:64659 IMAGE:6402346, complete cds	BF580053	9.8	8.4	4.1	3.7	6.6	9.2	3.9	5.3
1418912_at	naked cuticle 1 homolog (Drosophila)	NM_027280	9.4	6.4	5.4	3.6	7.8	7.7	1.6	7.6
1429794_a_at	mucolin 2	AK014467	5.1	7.2	6.9	5.4	7.0	8.0	2.7	6.6
1447272_s_at	complement component 1, q subcomponent, polypeptide	NM_007572	4.0	7.5	4.6	4.9	3.3	6.3	3.3	14.7
1443772_at	lysozyme	AW208566	3.0	6.3	7.7	6.6	4.4	2.7	4.4	11.4
1449038_at	RIKEN cDNA 2610307O08 gene	AV300716	1.7	2.8	14.2	7.8	4.3	4.5	2.2	8.7
1422494_s_at	kelch domain containing 2	AV298107	7.4	9.1	5.4	4.6	4.5	5.7	4.2	4.6
1441068_at	serine hydroxymethyl transferase 1 (soluble)	NM_009171	7.1	6.6	5.4	5.8	4.8	5.6	4.9	4.6
1434793_at	lysosomal membrane glycoprotein 2	NM_010685	4.8	4.4	4.1	5.2	9.2	6.3	1.7	9.0
1439449_at	dCMP deaminase	BG069699	4.1	4.3	5.6	6.3	5.3	6.8	5.4	4.3
1425569_a_at	RIKEN cDNA E130012A19 gene	BC006054	8.8	6.0	4.0	4.4	4.2	6.4	3.5	4.5
1443192_at	tubulin, beta 3	NM_023279	10.2	5.9	4.5	2.0	3.8	6.9	0.6	7.0
1440372_at	RWD domain containing 2	AK006533	8.0	7.8	4.8	6.8	2.9	2.5	2.8	4.7
1446088_at	cadherin 22	AB019618	3.3	4.1	6.9	3.9	5.4	8.4	1.7	6.3
1458589_at	X-ray repair compl. defective repair in Chin. hamster cells 5	AF166486	4.3	5.0	7.4	4.3	6.6	3.7	2.0	5.9
1433988_s_at	RIKEN cDNA B230114H05 gene	BM214769	4.6	5.3	5.6	3.9	7.0	5.4	2.6	4.8
1444159_at	expressed sequence AI324154	BQ031094	3.9	5.1	5.3	4.4	5.4	3.1	6.1	4.6
1420841_at	DNA segment Chr13, Wayne State University 177, expressed	BC012213	4.4	5.4	3.7	3.6	4.1	6.2	3.3	3.8
1452928_at	SET and MYND domain containing 2	BC023119	6.6	5.7	2.9	3.1	3.3	4.7	3.4	4.4
1457656_s_at	retinoblastoma binding protein 4	AU040918	5.0	6.0	3.8	4.2	3.2	3.5	4.1	3.9
1443284_at	leucine aminopeptidase 3	AK010384	4.3	5.1	4.0	3.6	3.9	5.5	3.0	3.5
1438570_at	RNA polymerase 1-4	NM_009088	4.4	4.2	3.9	3.6	3.8	4.6	3.3	4.2
1446284_at	RIKEN cDNA 1110004L07 gene	AK003432	6.2	6.2	3.1	3.1	3.1	3.3	3.0	3.5
1459674_at	MYB binding protein (P160) 1a	BM227247	4.3	4.0	3.4	3.6	3.8	4.9	3.1	3.1

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			Fold downregulation							
1457917_at	Mus musculus transcribed sequences	BB200032	96.4	-	45.2	15.1	104.1	152.5	14.2	41.6
1440721_at	RIKEN full-length enriched library, clone:5930433N17	C85455	-	-	22.7	65.7	22.0	-	14.7	25.7
1458058_at	cDNA, RIKEN full-length enriched library, clone:7030407E18	AV352368	318.4	-	74.9	-17.8	28.1	-	4.3	1123.7
1458309_at	RIKEN cDNA 9030605E16 [Mus musculus]	BM122422	78.7	29.2	-	-	-	-	2.6	-
1439412_at	Mus musculus transcribed sequences	BB167922	-	55.8	-	148.4	-	71.0	6.1	120.4
1427655_a_at	RIKEN cDNA A630038E17 gene	M37694	-98.4	-993.0	33.8	18.2	-177.6	44.3	56.5	10.1
1440188_at	gb:BE980962	BE980962	-	1069.7	179.7	320.1	-	-	2.6	12.1
1444942_at	Mus musculus mRNA for mKIAA0054 protein	BB277106	8.3	-	-	-	17.5	-	4.1	74.4
1429254_at	aquaporin 11	BF472491	-	-	84.6	2.1	-	-	3.5	-
1427532_at	RIKEN cDNA C030046M14 gene	BB622792	50.4	26.6	20.1	19.5	21.6	16.3	5.4	19.6
1439315_at	cDNA, RIKEN full-length enriched library, clone:A630042G22	BB233192	15.5	11.5	2124.1	15.4	18.7	-	3.3	34.4
1445148_at	cDNA, RIKEN full-length enriched library, clone:C230091K16	BB041366	10.2	43.0	5.8	10.2	14.5	32.8	12.3	12.4
1457063_at	Mus musculus transcribed sequences	AI447357	41.6	10.7	16.1	10.5	14.0	10.5	8.3	9.6
1420166_at	deoxynucleotidyltransferase, terminal	BB160593	5.0	4.7	-	20.4	75.1	7.8	150.1	14.6
1418065_at	recombination activating gene 2	NM_009020	4.4	6.1	-	39.3	-	60.8	4.0	31.6
1438837_at	Mus musculus transcribed sequences	BB198675	103.8	32.2	14.3	11.8	7.5	13.6	4.4	11.3
1458050_at	Mus musculus transcribed sequences	BB243259	13.9	14.4	13.6	20.8	8.1	62.5	3.8	17.6
1447792_x_at	Mus musculus transcribed sequences	BB241847	-	35.7	3.8	4.9	28.4	-	6.2	6.4
1421604_a_at	Kruppel-like factor 3 (basic)	NM_008453	25.7	8.9	7.7	22.0	105.0	-	7.0	3.7
1443894_at	Mus musculus transcribed sequences	BB236216	279.1	-	7.8	8.4	16.5	-	2.8	10.0
1451474_a_at	DNA segment, Chr 13, ERATO Doi 275, expressed	BC022679	45.7	10.8	8.5	11.2	7.1	14.2	8.7	7.1
1457141_at	Mus musculus transcribed sequences	BB229969	8.9	-	85.8	3.9	14.5	82.5	2.8	21.5
1447214_at	Mus musculus transcribed sequences	BG094486	9.8	8.4	13.1	40.9	6.9	17.8	3.8	58.1
1439686_at	Mus musculus transcribed sequences	BB235876	20.4	14.4	10.0	7.5	13.9	21.5	5.1	7.3
1417481_at	receptor (calcitonin) activity modifying protein 1	NM_016894	12.5	7.6	44.1	6.6	22.2	14.1	4.4	12.3
1419028_at	protein phosphatase 1, regulatory (inhibitor) subunit 1C	NM_033264	16.4	12.6	23.1	9.2	26.9	20.1	2.7	17.5
1448929_at	coagulation factor XIII, alpha subunit	NM_028784	21.7	11.3	22.6	19.4	10.5	5.7	19.0	3.8
1438894_at	Mus musculus transcribed sequences	BB371643	16.9	13.6	26.2	16.9	13.0	20.1	2.6	11.1
1439897_at	transcribed sequence weak sim. to protein sp:O76041 (H.sapiens)	BE457796	9.5	7.3	19.4	8.1	18.6	88.9	3.5	9.8
1418776_at	RIKEN cDNA 5830443L24 gene	NM_029509	60.4	30.2	5.5	7.3	14.3	37.0	10.2	3.3
1419480_at	selectin, lymphocyte	NM_011346	30.3	12.6	5.0	6.3	15.7	47.1	6.2	6.3
1418912_at	plexin domain containing 2	NM_026162	9.3	6.6	11.6	9.2	17.2	18.7	5.1	8.0
1429794_a_at	purinergic receptor P2X, ligand-gated ion channel, 1	AK007650	4.3	2.8	587.3	33.0	41.8	25.8	5.0	455.9
1447272_s_at	ATPase, class V, type 10A	BM249532	30.7	15.5	15.1	7.9	14.1	-	2.1	11.4
1443772_at	cDNA, RIKEN full-length enriched library, clone:B130053A22	BB202611	9.1	5.5	-	67.4	13.4	5.5	3.0	35.3
1449038_at	hydroxysteroid 11-beta dehydrogenase 1	NM_008288	6.4	7.9	23.4	12.5	15.5	23.7	8.8	3.4
1422494_s_at	RIKEN cDNA 4932702G04 gene	NM_053180	13.3	6.8	8.6	9.7	7.3	12.8	5.9	10.7
1441068_at	RIKEN cDNA A130001G05 gene	BB631473	25.5	20.3	14.0	13.7	9.9	28.4	2.0	15.7
1434793_at	hypothetical protein MGC38950	AI851014	17.5	10.0	10.8	7.4	6.4	26.1	3.9	9.6
1439449_at	transcribed sequence str. Sim. to protein pir:A43314 (H.sapiens)	BB724383	7.9	11.0	65.9	13.7	11.9	92.1	2.0	21.8
1425569_a_at	signaling lymphocyte activation molecule	AF149791	17.4	12.2	10.7	6.3	14.7	25.2	3.6	6.0
1443192_at	cDNA, RIKEN full-length enriched library, clone:5830487P08	BB033993	13.4	15.8	10.9	13.0	5.1	23.3	2.8	17.5
1440372_at	Mus musculus transcribed sequences	AI785253	14.9	11.4	24.4	38.2	5.2	16.3	2.2	14.5
1446088_at	RIKEN cDNA 9430081I23 gene	BB102291	7.5	13.1	15.3	14.0	9.4	11.8	2.8	9.7
1458589_at	cDNA, RIKEN full-length enriched library, clone:D430030N05	BB485297	7.9	11.9	8.6	6.5	11.6	9.9	4.0	11.9
1433988_s_at	RIKEN cDNA C230098O21 gene	BG075755	10.5	12.5	10.4	9.1	8.2	6.3	4.8	7.4
1444159_at	cDNA, RIKEN full-length enriched library, clone:8030444E01	BM243903	13.8	13.1	18.2	19.8	8.3	24.1	1.9	13.9
1420841_at	protein tyrosine phosphatase, receptor type, F	NM_011213	8.2	11.4	29.3	5.4	15.2	25.0	2.7	9.6
1452928_at	RIKEN cDNA 2210414K06 gene	AK008928	13.3	6.3	6.3	8.0	12.2	9.1	7.2	5.7
1457656_s_at	Mus musculus transcribed sequences	BB235011	6.5	9.8	88.6	9.8	11.9	60.2	2.0	20.4
1443284_at	Mus musculus transcribed sequences	BB198943	5.5	8.2	-	10.1	10.0	74.7	2.1	17.5
1438570_at	Mus musculus transcribed sequences	BE631083	6.5	7.5	40.4	8.9	9.1	26.8	2.5	11.9
1446284_at	cDNA, RIKEN full-length enriched library, clone:A130035K06	BB157298	15.1	17.3	-	13.1	7.1	5.3	2.1	15.0
1459674_at	Mus musculus transcribed sequences	AW907492	12.4	13.5	43.5	10.0	5.5	11.1	2.3	13.6
1456235_at	Mus musculus transcribed sequences	BB198406	11.2	12.9	8.8	11.4	4.7	18.7	2.8	13.0
1441614_at	gb:BB229543	BB229543	8.3	12.7	11.0	11.2	5.6	18.5	2.9	8.9
1437699_at	RIKEN cDNA E430014B02 gene	BB031353	6.5	8.6	59.4	10.1	10.1	44.4	2.0	15.4
1439203_at	Mus musculus transcribed sequences	BB206010	8.5	10.5	21.9	7.7	8.7	25.3	2.2	12.4
1426438_at	RIKEN cDNA 8030469F12 gene	BB667072	-	67.0	1.7	2.3	-	-	-	22.9

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			1	2	3	4	5	6	7	8
			Fold downregulation							
1438125_at	RIKEN cDNA C230085N15 gene	BB393897	5.2	8.4	61.5	8.2	8.4	38.6	2.3	16.5
1454295_at	Mus musculus transcribed sequences	BB041915	20.5	15.0	5.0	9.0	4.2	31.7	3.2	12.1
1441769_at	Mus musculus transcribed sequences	BB731108	16.6	9.8	15.2	19.6	4.8	11.9	2.4	9.8
1440326_at	Mus musculus transcribed sequences	AV312674	26.1	34.0	8.8	13.0	6.7	173.4	1.6	22.6
1457582_at	cDNA, RIKEN full-length enriched library, clone:D930045J11	BB528233	33.0	9.8	2.5	4.1	8.9	20.7	9.1	17.8
1441418_at	Mus musculus transcribed sequences	BM240473	27.6	28.3	7.9	10.4	4.8	50.3	1.9	14.6
1457549_at	Mus musculus transcribed sequences	BB043576	6.4	18.9	17.7	17.9	4.6	20.1	2.0	23.1
1460251_at	tumor necrosis factor receptor superfamily, member 6	NM_007987	12.4	3.8	12.2	11.2	13.6	172.5	2.4	8.2
1416503_at	latexin	NM_016753	9.9	3.9	8.9	5.7	-	-	6.5	2.8
1439839_at	RIKEN cDNA D130051D11 gene	BB815394	6.3	11.0	9.7	6.1	5.5	5.9	6.8	7.9
1419091_a_at	annexin A2	NM_007585	22.6	9.4	4.4	6.1	11.8	3.4	11.9	6.7
1440900_at	cDNA, RIKEN full-length enriched library, clone:5830404C02	AV312603	13.5	7.6	4.3	5.3	9.4	9.1	6.2	6.3
1441589_at	Mus musculus transcribed sequences	BB634277	27.5	28.8	8.0	12.6	5.7	50.7	1.6	15.7
1441552_at	cDNA, RIKEN full-length enr. library, cl.:5830474E17 product:T-cell receptor alpha chain precursor V-J region(TA72)	BB032962	12.1	9.9	17.0	4.9	7.9	4.5	15.0	3.2
1425792_a_at	RAR-related orphan receptor gamma	AF163668	3.3	4.9	1404. 0	6.3	22.1	95.1	2.6	15.3
1438392_at	Mus musculus transcribed sequences	BM239011	10.0	11.2	9.2	7.9	6.0	9.8	2.6	9.4
1458605_at	Mus musculus transcribed sequences	AW107249	13.6	11.3	11.7	13.0	4.2	8.6	2.4	10.2
1453595_at	RIKEN cDNA 2900064B18 gene	AV154597	12.6	28.0	8.3	9.0	4.2	9.4	2.6	9.0
1440586_at	RIKEN cDNA B430203I24 gene	BB326079	13.8	10.4	12.6	15.7	4.2	9.1	2.2	12.7
1458974_at	Mus musculus transcribed sequences	BM245368	9.1	9.0	11.9	10.5	5.1	14.4	2.5	7.8
1438981_at	Mus musculus transcribed sequences	BB160675	9.4	15.4	9.3	13.0	6.1	10.8	2.0	12.2
1426906_at	interferon activated gene 205	M74124	127.9	14.0	3.7	3.2	9.2	26.0	7.1	3.8
1448355_at	protease, serine, 16 (thymus)	NM_019429	6.7	5.6	7.1	5.3	6.3	6.1	7.0	9.9
1459202_at	cDNA, RIKEN full-length enriched library, clone:C130034D14	BB366565	4.1	5.6	13.6	12.3	4.8	8.6	4.6	8.4
1457377_at	Mus musculus transcribed sequences	BB253657	10.0	12.3	6.1	9.1	5.3	11.2	2.5	8.9
1444224_at	Mus musculus transcribed sequences	AI550170	30.1	97.0	8.5	9.4	4.9	10.2	2.1	4.9
1433719_at	hypothetical protein 9930105B05	AI854429	4.6	7.2	6.6	5.9	11.5	11.1	5.3	4.7
1456678_at	cDNA, RIKEN full-length enriched library, clone:A130058I09	AV314270	15.1	24.1	6.3	7.7	5.0	25.9	1.9	12.1
1459362_at	Mus musculus transcribed sequences	BB033588	10.7	13.0	7.6	8.9	5.4	14.5	2.0	11.5
1438452_at	RIKEN cDNA A630080F05 gene	BM121794	7.7	5.1	16.8	6.8	10.9	10.7	2.6	6.4
1441762_at	cDNA, RIKEN full-length enriched library, clone:5830468D02	BM877183	4.0	6.9	15.7	9.1	8.2	8.9	2.4	26.1
1439895_at	transcribed seq. mod. sim. to protein pir:S12207 (M.musculus)	BG069874	11.1	10.5	9.7	12.3	4.1	10.1	2.3	8.0
1446711_at	RIKEN cDNA A130040G06 gene	BB165609	7.5	10.2	8.1	5.7	6.2	11.7	2.5	9.3
1460745_at	RIKEN cDNA A630098A13 gene	AI327345	2.4	5.8	8.3	6.7	11.3	10.6	7.4	8.7
1457184_at	Mus musculus transcribed sequences	BB472173	5.6	10.8	7.3	8.0	4.1	9.2	2.9	17.8
1431118_at	RIKEN cDNA 6720427H10 gene	BB664131	10.8	9.1	9.9	11.1	3.8	11.7	2.3	7.7
1444293_at	Mus musculus, clone IMAGE:1328084, mRNA	BB204970	3.5	5.5	17.2	7.5	13.4	8.7	3.1	6.8
1441058_at	RIKEN cDNA E130307H12 gene	BB434111	11.2	8.3	7.9	9.1	5.8	8.4	2.2	7.6
1456426_at	Mus musculus transcribed sequences	BE686253	7.7	3.7	4.0	5.7	11.9	105.3	6.5	4.0
1420515_a_at	RIKEN cDNA C730002N09 gene	NM_021319	9.1	6.5	8.5	2.1	22.7	-	4.9	4.2
1449831_at	RIKEN cDNA 1700055O19 gene	NM_026100	5.3	4.7	5.2	6.6	8.5	13.0	3.7	8.6
1423549_at	solute carrier family1 (glutamate/neutral aa transporter), member 4	BB762524	2.6	4.7	18.6	17.6	6.3	8.1	7.5	4.6
1446950_at	Mus musculus transcribed sequences	BM124834	7.3	10.6	9.0	7.8	5.0	10.7	2.4	6.4
1443534_at	Mus musculus transcribed sequences	BM201095	19.2	13.4	4.3	5.8	5.4	26.5	2.4	5.8
1451846_at	RIKEN cDNA 1200007O21 gene	BC025863	20.8	8.5	11.9	4.1	8.4	7.3	2.6	4.7
1424903_at	selected mouse cDNA on the Y	AF127244	27.9	22.2	1.6	2.3	20.9	23.4	29.4	11.3
1443349_at	Mus musculus transcribed sequences	BB211894	246.8	16.0	2.5	3.5	6.5	30.1	4.0	5.3
1458263_at	Mus musculus transcribed sequences	BE979587	25.9	15.5	3.2	5.0	3.7	415.8	2.6	8.8
1431110_at	RIKEN cDNA 5430431D22 gene	AK017369	6.6	3.4	23.4	4.7	6.4	-	2.5	5.5
1443094_at	expressed sequence BB031773	BB031311	5.5	45.2	6.2	8.7	3.6	7.0	2.5	11.8
1451475_at	Mus musculus mRNA for mKIAA0620 protein	BC019530	26.1	7.7	5.6	5.5	33.1	-	2.0	2.8
1421386_at	ankyrin repeat domain 6	NM_080471	9.5	7.1	7.2	4.3	5.6	8.7	3.0	6.1
1456685_at	Mus musculus transcribed sequences	BB227199	13.2	6.2	3.2	5.8	5.9	8.9	4.2	5.7
1446476_at	cDNA, RIKEN full-length enriched library, clone:5830404G23 product:T-cell receptor alpha chain precursor V-J region (TA72)	AV312643	6.6	12.5	3.8	3.5	2.4	34.6	31.0	6.2
1418777_at	chemokine (C-C motif) ligand 25	NM_009138	7.6	5.0	12.3	3.1	7.7	13.3	3.4	5.1
1426721_s_at	expressed sequence AW558171	BB667071	5.6	3.4	5.5	7.0	6.7	13.2	3.7	7.2
1456650_at	cDNA, RIKEN full-length enriched library, clone:A030010B11	BB150912	2.8	10.0	6.5	7.0	5.8	75.0	3.5	4.6

Probe set	Gene	Accession	Lymphoma #							
			1	2	3	4	5	6	7	8
			Fold downregulation							
1445426_at	gb:BB457090	BB457090	10.0	4.9	6.9	26.1	10.5	8.1	1.8	5.2
1446272_at	PCTAIRE-motif protein kinase 2	BB200545	2.8	6.2	16.2	6.5	8.2	26.8	2.2	9.8
1434473_at	Mus musculus cDNA clone IMAGE:3471887, partial cds	AI647939	3.9	6.1	8.2	4.1	11.0	12.3	3.3	5.1
1431014_at	cat eye syndrome chrom. region, candidate 2 homolog (human)	AK013061	4.6	6.0	13.4	7.3	3.7	6.2	3.8	6.0
1447458_at	RIKEN cDNA 9330161A08 gene	BE954474	5.7	14.5	5.9	9.6	4.5	4.3	2.5	10.2
1443947_at	RIKEN cDNA C430041B13 gene	BB415206	41.8	3.5	3.1	6.4	3.2	-	2.6	11.7
1419226_at	T cell activation, increased late expression	NM_032465	7.7	4.8	14.1	4.9	6.5	8.3	2.8	4.2
1427184_at	T-cell receptor beta, variable 8.2	BF318536	2.5	6.3	6.0	6.5	5.6	-	7.8	3.5
1436154_at	Mus musculus transcribed sequences	BE457918	2.6	5.2	14.7	5.5	9.2	8.3	3.3	8.7
1437584_at	Mus musculus transcribed sequences	BE685667	5.6	11.3	10.4	8.9	3.9	9.6	1.7	12.1
1438960_at	Mus musculus transcribed sequences	BB164652	21.5	10.5	4.0	11.8	2.9	15.5	1.8	14.6
1417141_at	interferon gamma induced GTPase	NM_018738	10.6	5.8	5.1	3.7	11.1	10.0	6.8	2.2
1454788_at	RIKEN cDNA A630084M22 gene	BQ176306	4.1	5.1	8.7	4.3	16.3	28.9	2.1	6.0
1445612_at	cDNA, RIKEN full-length enriched library, clone:F830035K10	AI114898	4.8	9.1	7.8	7.5	4.1	11.9	2.3	4.9
1443144_at	RIKEN cDNA A130082F03 gene	BB084315	3.5	6.6	6.6	4.0	6.8	6.3	4.3	5.8
1429690_at	RIKEN cDNA 1300003B13 gene	AK004870	6.3	8.8	4.6	5.6	3.6	14.8	2.8	5.4
1449875_s_at	histocompatibility 2, T region locus 10	NM_010395	8.4	5.3	4.7	4.6	8.7	9.3	2.7	4.5
1436864_at	Mus musculus transcribed sequences	BM214060	7.7	4.6	8.2	9.8	4.5	8.4	2.0	6.8
1444005_at	Mus musculus transcribed sequences	AI529516	10.9	9.9	9.8	5.3	4.7	4.2	2.1	6.6
1417293_at	heparan sulfate 6-O-sulfotransferase 1	NM_015818	6.8	4.9	7.5	6.0	4.3	5.6	4.2	3.6
1422064_a_at	zinc finger protein 288	AY028963	6.1	4.4	2.6	4.2	11.1	9.1	4.5	6.9
1425785_a_at	TXK tyrosine kinase	U19607	5.6	8.7	6.1	2.6	8.5	11.7	2.6	6.5
1439981_at	RIKEN cDNA A430001F24 gene	BB198298	6.0	6.3	5.3	5.4	4.2	12.6	2.7	4.8
1456274_at	cDNA, RIKEN full-length enriched library, clone:C230071H18	BB067544	7.3	10.9	4.8	5.3	4.5	7.9	2.2	5.5
1438906_at	cDNA, RIKEN full-length enriched library, clone:A130030I02	BB156385	18.6	22.6	3.5	6.7	3.0	15.3	1.8	7.9
1440103_at	Mus musculus transcribed sequences	AV351137	5.4	7.8	5.2	5.1	4.3	5.1	3.7	4.8
1418633_at	Notch gene homolog 1 (Drosophila)	NM_008714	9.3	11.7	3.4	3.1	5.6	4.9	5.6	4.0
1443012_at	cDNA, RIKEN full-length enriched library, clone:B130008C17	BB291827	10.0	9.9	4.3	6.2	3.9	17.8	1.7	7.5
1456806_at	RIKEN cDNA A130010C12 gene	BB153010	6.9	9.4	6.3	6.1	3.9	5.7	2.1	7.1
1445106_at	Mus musculus transcribed sequences	BB066105	10.2	9.9	4.1	6.9	4.6	17.6	1.8	5.3
1418968_at	RB1-inducible coiled-coil 1	AB050017	4.0	3.1	5.6	6.1	6.0	20.3	3.1	6.0
1444350_at	schlafen 10	BB227643	3.1	9.2	5.3	3.4	7.3	-	2.7	4.6
1431961_at	RIKEN cDNA 5830472H07 gene	AK018051	6.6	6.4	3.4	4.6	4.5	11.5	3.0	5.3
1429196_at	RIKEN cDNA 5830411O09 gene	BB431654	4.3	1.7	14.1	7.3	8.6	11.6	3.6	7.5
1460003_at	Mus musculus transcribed sequences	AV234963	4.6	3.8	5.6	4.9	4.9	7.2	3.6	5.7
1421512_at	centrosomal protein 2	NM_008383	3.5	4.3	5.2	6.7	6.4	11.0	2.9	4.6
1441253_at	cDNA, RIKEN full-length enriched library, clone:D330006I11	BB474427	11.5	7.6	5.0	6.5	2.9	7.5	2.3	5.4
1437136_at	RIKEN cDNA 5830436I19 gene	AV316469	5.5	11.6	9.5	8.4	3.1	6.2	1.7	8.5
1438805_at	RIKEN cDNA 9230106B05 gene	BE691816	2.6	3.7	10.9	6.2	5.5	10.8	2.4	11.2
1437821_at	cDNA, RIKEN full-length enriched library, clone:C430005H19	BB408240	11.8	10.4	3.5	5.5	3.5	14.1	1.9	6.4
1436398_at	cDNA, RIKEN full-length enriched library, clone:A130028J07	BB033554	5.0	6.1	5.6	7.7	3.9	6.3	2.1	7.0
1429060_at	RIKEN cDNA 9430072K23 gene	AK020483	4.6	5.8	5.3	5.0	3.7	11.4	2.7	5.3
1445881_at	RIKEN cDNA 2310035P21 gene	BG230003	6.1	9.7	4.7	6.5	3.0	6.4	2.2	7.8
1443400_at	transcribed seq. mod. sim. to protein pir:S12207 (M.musculus)	BG060869	6.2	8.1	4.4	5.4	3.1	12.0	2.2	5.8
1445615_at	Mus musculus transcribed sequences	BE852824	5.3	3.6	5.6	4.5	3.5	5.8	3.8	6.9
1445252_at	cDNA, RIKEN full-length enriched library, clone:C130070D21	BB373408	3.5	7.4	4.9	5.4	6.4	3.1	3.2	6.6
1439209_at	cDNA, RIKEN full-length enriched library, clone:E130109D23	BB540782	7.2	8.1	4.2	5.1	3.5	17.3	1.7	7.5
1451310_a_at	cathepsin L	J02583	9.5	5.1	3.0	7.1	2.2	6.9	5.9	4.8
1422287_at	per-hexamer repeat gene 2	NM_008833	5.5	8.7	4.5	4.2	4.1	4.8	3.2	4.0
1459695_at	Mus musculus transcribed sequences	AU040352	5.5	7.0	3.5	3.9	3.7	5.6	3.4	5.7
1438762_at	Mus musculus transcribed sequences	AI604411	9.5	9.3	3.7	4.9	3.3	13.0	1.7	7.2
1458947_at	Mus musculus transcribed sequences	AV341977	7.3	11.3	2.5	3.7	4.1	6.8	2.8	6.2
1430585_at	RIKEN cDNA 5930436O19 gene	BG798082	5.4	6.1	4.0	4.6	3.0	8.2	2.7	5.3
1441319_at	RIKEN cDNA D030069N10 gene	BB449095	4.4	8.7	5.2	5.1	3.3	10.8	1.8	5.8
1446551_at	cDNA, RIKEN full-length enriched library, clone:A130081C01	BB165047	3.9	6.1	5.0	4.3	3.2	7.4	2.7	5.8
1457782_at	Mus musculus transcribed sequences	BB045512	6.9	18.0	4.6	4.3	3.5	3.9	2.2	4.7
1444473_at	Mus musculus transcribed sequences	BM240241	5.2	7.8	3.7	6.0	3.8	6.1	2.1	5.2
1446592_at	cDNA, RIKEN full-length enriched library, clone:D230005P03	BB466044	5.5	7.8	4.1	4.3	3.6	8.7	2.1	4.7
1441556_at	Mus musculus transcribed sequences	BB476958	4.5	5.3	7.0	4.0	4.1	4.1	2.1	9.2
1444676_at	cDNA, RIKEN full-length enriched library, clone:A430066N02	BB205578	9.0	9.6	3.7	3.5	3.8	6.4	1.9	6.3
1423597_at	aminophospholipid transporter (APLT), classI, type 8A, member1	BB303874	7.9	3.9	2.1	3.5	6.9	20.0	3.8	3.5

Probe set	Gene	Accession	Lymphoma #							
			1	2	3	4	5	6	7	8
			Fold downregulation							
1417756_at	lymphocyte specific 1	NM_019391	12.4	5.5	2.5	2.6	6.9	10.4	3.1	3.5
1426505_at	ecotropic viral integration site 2B	BB201368	9.8	7.0	4.2	3.3	5.5	7.8	2.3	2.6
1416091_at	microtubule-associated protein 4	NM_008633	5.2	4.1	6.2	3.9	5.5	5.4	1.9	5.0
1441354_at	Mus musculus transcribed sequences	BG069312	3.6	5.5	3.7	6.8	3.0	5.1	2.6	6.4
1430367_at	RIKEN cDNA 8230401J17 gene	BE331859	6.9	6.8	7.1	4.7	3.2	4.1	1.9	4.3
1452145_at	hexose-6-phosphate dehydrogenase (glucose 1-dehydrogenase)	BC027358	3.5	3.7	6.4	4.2	6.6	6.3	2.4	3.4
1441505_at	cDNA, RIKEN full-length enriched library, clone:A430105J09	BB211484	4.7	5.5	4.0	5.5	2.8	6.3	2.3	5.7
1440905_at	Mus musculus transcribed sequences	AI451467	8.5	8.5	4.0	3.5	3.3	6.2	1.9	4.7
1444051_at	RIKEN cDNA 1700019D03 gene	AA215046	4.7	5.6	3.6	4.4	3.7	9.3	2.1	4.2
1419221_at	regulator of G-protein signaling 14	NM_016758	5.2	3.5	4.4	3.9	4.6	3.2	3.3	3.9
1425123_at	cDNA sequence BC025816	BC025816	3.9	3.1	4.3	3.9	5.2	3.5	3.7	3.4
1451234_at	cDNA sequence BC021381	BC021381	3.2	3.3	4.0	4.7	4.2	4.3	2.3	4.9

-: the expression is below background