Probeset	Entrez	Symbol	Name	Th	Th act	Tc	Tc act	B	B act	B aIgM	Mem IgG	Mem IgM	PC	NK	NK act	mono	mono act	DC	DC act	neutro
242904_x_at				5292.9	104.6	191.29	41.553	671.33	125.9	78.485	1355.6	2167.6	575.7	234.22	193.38	60.43	94.563	44.379	42.333	528.59
232165_at				605.1	57.129	19.282	23.457	19.076	22.296	18.152	51.534	45.244	24.699	22.594	25.374	15.907	23.304	17.794	14.671	49.81
201369_s_at	678	ZFP36L2	zinc finger protein 36, C3	5213.9	625.69	860.42	267.09	1040.9	563.83	248.18	422.51	430.73	170.09	944.54	829.88	947.23	303.25	768	224.49	685.29
233127_at				1710.6	222.72	147.83	79.293	319.04	236.36	156.19	134.89	262.19	128.64	118.91	74.872	49.12	36.745	26.144	19.642	24.981
204621_s_at	4929	NR4A2	nuclear receptor subfami	1407.8	247.46	19.61	143.09	50.766	112.38	322.4	202.34	365.71	42.58	16.994	108.1	62.992	220.11	49.329	100.96	106.7
202861_at	5187	PER1	period homolog 1 (Droso	1079.4	104.49	189.22	154.5	57.445	81.581	162.01	48.368	116.49	31.034	102.67	116.49	189.11	128.72	103	82.624	72.219
222862_s_at	26289	AK5	adenylate kinase 5	1379.6	330.29	203.27	52.567	50.005	48.406	38.379	69.45	65.663	55.256	199.32	175.49	27.348	33.242	26.782	42.736	114.44
235213_at				9321.4	1863.7	1799	613.54	1615.5	2073.6	2786.8	231.11	151.04	294.67	2300	1248	134.46	135.6	290.04	94.781	610.74
235739_at	4929	NR4A2	nuclear receptor subfami	6951	389.67	369.9	408.86	1149.1	388.4	630.63	1875.9	3071.5	448.05	192.38	340.33	209.66	263.31	151.63	182.99	1779.9
231798_at	9241	NOG	noggin	1205.8	216.29	175.73	61.715	163.99	62.575	71.163	153.18	210.55	226.19	108.3	65.022	67.638	107.4	66.928	68.087	245.57
200664_s_at	3337	DJB1	DnaJ (Hsp40) homolog, s	4361.3	980.92	842.13	919.73	580.86	779.84	629.17	342.02	318.99	888.9	913.3	898.93	1674.3	1546.3	1153.8	890.5	815.13
240452_at				3343.6	1173.4	661.92	751.8	737.37	768.24	920.58	513.34	503.68	565.05	460.41	371.28	148.31	138.81	121.14	117.47	234.13
236907_at				14624	2674.3	1387.4	3255.3	5202	1429.3	2599.7	3060.4	3890.8	1513.1	773.73	774.08	460.05	327.9	234.19	363.43	1545.7
204622_x_at	4929	NR4A2	nuclear receptor subfami	1142.3	259.03	75.925	204.75	125.36	209.68	312.83	502.61	820.81	97.28	167.22	211.2	153.38	325.82	153.57	127.94	217.77
227307_at				1948.2	400.21	234	129.54	353.61	210.34	149.64	323.63	239.4	712.5	93.089	110.13	83.97	121.68	134.12	124.52	230.54
227641_at	146330	FBXL16	F-box and leucine-rich re	3811.3	1576.5	1403.5	697.89	732.34	672.19	494.3	833.77	550.58	1161	311.67	323.29	224.19	245.56	351.13	401.22	1170.8
201367_s_at	678	ZFP36L2	zinc finger protein 36, C3	3793.8	620.67	771.55	219.34	1023.5	486.97	268.09	366.91	415.21	172.63	893.78	701.58	1109.2	276.4	869.58	278.41	706.3
234408_at	112744	IL17F	interleukin 17F	75.661	9648.6	96.514	126.73	146.75	99.437	125.95	181.58	145.47	169.89	132.08	57.788	66.779	62.439	48.914	67.21	192.07
216876_s_at	3605	IL17A	interleukin 17A	7.194	809.34	3.396	10.259	7.24	8.1	3.219	24.895	36.767	5.585	6.173	21.649	12.938	15.326	13.615	19.02	18.389
234362_s_at	1493	CTLA4	cytotoxic T-lymphocyte-a	1152.6	5171.8	644.65	1132.7	400.95	331.91	319.33	599.21	486.96	540.88	280.2	272.35	127.25	202.8	219.86	213.03	546.89
221331_x_at	1493	CTLA4	cytotoxic T-lymphocyte-a	456.17	1826.6	170.64	539.95	77.447	57.268	68.635	58.954	76.762	89.278	105.53	117.46	74.196	74.198	166.69	144.29	185.74
211856_x_at	940	CD28	CD28 molecule	484.12	1102.4	350.46	347.35	148.4	101.36	39.628	175.46	291.21	134.16	54.782	99.024	69.36	108.31	103.85	99.261	313.67
219179_at	51339	DACT1	dapper, antagonist of bet	276.97	664.71	99.844	95.204	207.63	228.89	116.55	212.18	274.44	115.61	56.953	114.38	164.18	108.19	238.82	129.36	226.99
221165_s_at	50616	IL22	interleukin 22	195.27	1621.9	140.2	272.41	182.77	177	131.76	305.89	259.44	170.05	130.08	140.77	160.41	208.94	195.76	176.95	458.47
235150_at				848.53	1665.6	388.3	162.9	305.6	519.98	590.79	135.39	151.2	120.16	479.75	251.81	70.653	94.283	209.79	790.54	79.465
235616_at	128553	TSHZ2	teashirt zinc finger home	669.63	1663.3	288.41	312.36	171.51	286.96	117.25	355.25	272.75	201.94	169.31	130.05	81.735	121.52	97.362	109.71	253.85
227262_at	145864	HAPLN3	hyaluronan and proteogly	1946.6	8728.1	1101.6	2305.8	594.52	1180.4	388.18	558.27	536.28	992.28	563.5	2809.2	254.6	430.41	267.09	1769.6	644.6
226913_s_at	30812	SOX8	SRY (sex determining re	298.04	1025	19.781	21.533	30.567	26.109	23.557	28.412	27.116	41.972	39.268	45.119	15.518	25.178	50.898	22.052	25.256
211861_x_at	940	CD28	CD28 molecule	424.1	927.04	284.99	249.87	32.017	33.209	17.453	61.339	66.092	45.075	65.611	35.191	29.141	49.231	19.951	27.802	37.221
231794_at	1493	CTLA4	cytotoxic T-lymphocyte-a	829.63	2074.9	360.49	772.02	172.99	225.83	82.15	277.45	353.17	308.49	124.66	116.43	92.171	92.422	121.75	109.83	501.04
216901_s_at	10320	IKZF1	IKAROS family zinc finge	585.01	1251.5	289.46	343.31	289.61	355.95	186.75	251.73	257.34	202.66	571.49	213.94	254.88	233.08	271.55	439.89	210.19
220892_s_at	29968	PSAT1	phosphoserine aminotrar	346.27	2214.3	394.62	805.9	320.85	337.81	736.12	429.65	309.66	1025.6	199.65	207.83	202.32	287.5	581.74	510.86	477.33
208193_at	3578	IL9	interleukin 9	97.274	1343.5	49.426	410.4	83.163	57.007	22.289	145.32	101.28	54.427	95.923	63.72	52.85	60.811	68.764	52.046	88.46
243940_at	128553	TSHZ2	teashirt zinc finger home	305.87	591.41	112.12	137.34	154.73	110.94	63.719	214.32	219.95	234.12	102.72	72.86	28.29	63.443	42.089	40.373	163.89
215332_s_at	926	CD8B	CD8b molecule	41.855	32.983	1489.1	122.68	37.742	27.152	8.683	93.873	74.936	50.758	66.688	29.283	42.361	57.728	46.062	30.886	58.733
209840_s_at	54674	LRRN3	leucine rich repeat neuro	735.62	503.37	6792.8	1198.6	62.117	40.136	28.776	157.23	118.26	33.943	46.41	34.332	99.24	152.85	121.16	87.014	107.64
205758_at	925	CD8A	CD8a molecule	310.06	405.99	9957.8	3315.8	611.63	460.91	106.98	363.04	243.73	125.54	1747.3	1021.5	176.49	358.05	191.57	212.95	659.97
207979_s_at	926	CD8B	CD8b molecule	138.63	177.5	5811.1	2622.3	115.71	130.58	54.383	281.08	335.21	118.48	239.29	169.83	138.51	232.77	128.76	116.27	422.64
209841_s_at	54674	LRRN3	leucine rich repeat neuro	706.71	581.81	8644.5	2115.9	159.72	143.34	90.669	332.35	344.48	173.99	111.6	105.81	175.1	267.95	204.23	201.49	495.58
227915_at	51676	ASB2	ankyrin repeat and SOCS	79.337	112.51	2639.9	927.9	38.727	46.681	31.775	53.498	43.509	219.16	31.89	23.954	18.171	22.51	23.759	68.321	32.828
207232_s_at	9666	DZIP3	DAZ interacting protein 3	63.449	15.003	194.1	12.941	59.71	26.541	25.563	37.349	36.696	17.2	35.794	21.865	21.678	28.811	39.256	44.789	30.134
234427_at				246.02	176.44	613.76	234.81	24.22	30.823	24.564	39.521	42.166	31.209	23.72	13.456	19.854	21.666	14.435	20.995	53.299
226423_at	85315	PAQR8	progestin and adipoQ rec	786.88	1044.9	2261.4	883.48	460.26	520.17	258.38	424.19	344.92	557.11	945.21	671.94	472.6	224.57	882.61	233.68	556.69
208406_s_at				102.7	186.89	569.46	181.94	24.59	16.219	12.373	19.32	20.428	13.39	189.79	247.42	18.973	24.599	22.105	22.97	27.632
217119_s_at	2833	CXCR3	chemokine (C-X-C motif)	232.2	123.87	1592.3	282.94	73.401	52.68	25.821	101.19	79.587	242.77	602.71	345.71	49.709	78.357	64.907	28.424	62.056
242628_at				237.41	181.43	750.95	276.46	270.68	110.13	115.56	162.12	76.456	206.92	278.26	253.8	41.45	39.372	63.174	30.798	241.79
201414_s_at	4676	P1L4	nucleosome assembly pr	1095.9	1054.6	1961.3	1146	851.07	858.5	642.55	569.35	587.16	796.58	1028.9	774.41	1198.6	1045.5	896.23	1096.3	451.85
227552_at	1731	38230	septin 1	1853.2	1078	3456.8	1540.5	1236.1	1119.3	944.65	371.66	442.15	1079.7	1159.2	975.22	21.243	61.854	26.316	69.847	148.47
241871_at	814	CAMK4	calcium/calmodulin-depe	1278.2	978.5	2274.6	1265.9	238.29	296.08	68.922	188	231.35	301.28	139.73	101.14	25.46	75.621	50.817	142.6	204.37
219025_at	57124	CD248	CD248 molecule, endosia	366.65	216.83	1028.2	288.42	329.16	282.39	195.08	324.75	255.77	252.92	213.8	213.93	317.03	255.8	371.85	289.84	453.31
234398_at				116.38	108.86	526.68	257.03	74.15	61.25	39.666	104.62	117.43	122.23	49.748	27.014	31.759	55.044	32.022	36.939	217.53
243110_x_at	283869	NPW	neuropeptide W	5.177	5.275	4.551	539.25	6.061	5.692	446.58	11.427	16.884	12.242	4.098	4.461	7.843	11.752	8.392	9.127	7.039
227617_at	199953	RP13-15M	hypothetical protein LOC	192.86	235.07	220.53	3249.7	173.54	170.17	95.56	141.98	147.22	290.42	134.75	135.83	100.88	112.42	107.51	102.01	213.66
235238_at	399694	SHC4	SHC (Src homology 2 do	59.433	60.769	40.265	570.24	33.133	62.257	34.752	92.555	87.079	47.668	21.764	31.468	18.077	29.599	26.114	25.334	60.573
205266_at	3976	LIF	leukemia inhibitory facto	31.201	1084.5	36.214	5823.8	25.66	21.031	19.177	49.072	46.371	25.724	20.468	84.797	104.49	625.31	55.962	48.533	66.225
209000_s_at	23176		septin 8	11.013	20.212	16.933	126.6	10.758	11.885	13.505	25.628	24.617	18.216	13.231	10.118	20.444	23.372	23.637	21.66	24.944
203967_at	990	CDC6	cell division cycle 6 homo	93.088	194.93	220.38	1223.1	107.62	112.04	83.857	174.97	175.32	310.89	80.572	41.998	52.73	66.45	58.133	41.306	272.02
223490_s_at	51010	EXOSC3	exosome component 3	118.67	789.6	738.75	2638.3	174.02	567.47	822.37	48.067	79.427	299.82	499.31	691.75	285.35	269.96	183.8	226.82	93.13
224851_at	1021	CDK6	cyclin-dependent kinase	3018.2	4621.4	3289.6	19509	2481.6	2276.6	1490.8	493.55	258.8	3295	1248.6	830.99	475.77	235.09	1131.8	1444.6	318.97

218529_at	51293	CD320	CD320 molecule	144.21	273.65	232.18	1229.6	68.268	144.65	283.68	59.468	53.224	315.2	113.03	29.97	59.036	145.06	75.737	63.493	39.181
202613_at	1503	CTPS	CTP synthase	233.48	828.46	608.83	2722.2	268.05	532.91	994.27	373.66	397.97	386.96	223.21	303.98	430.55	239.9	573.47	332.52	293.59
224523_s_at	84319	C3orf26	chromosome 3 open reac	844.34	1873.3	1306.2	6709.1	935.86	1139.8	3038.4	411.69	346.84	1622.2	427.68	588.48	366.89	183.34	348.61	236.71	368.07
206653_at	10622	POLR3G	polymerase (R) III (DNA	111.28	129.25	80.188	499.74	110.15	90.279	191.06	105.25	95.599	122.2	65.923	66.872	62.421	96.927	95.214	73.507	98.346
228671_at	199953	RP13-15M	hypothetical protein LOC	169.99	261.11	249.09	1122.7	140.65	160.81	171.54	123.62	129.9	114.47	115.85	116.68	56.981	51.74	109.25	33.679	89.197
224848_at	1021	CDK6	cyclin-dependent kinase	1750.7	2549.3	1791.3	9645	1502.4	1408.9	897.3	752.27	632.6	2131.3	830.37	552.98	397.79	296.18	766.85	720.85	518.97
203270_at	1841	DTYMK	deoxythymidylate kinase	108.09	195.01	278.64	1007.2	163.8	160.07	273.64	133.27	153.84	292.14	178.37	112.05	210.83	157.39	223.51	179.2	45.123
205164_at	23464	GCAT	glycine C-acetyltransfera	20.973	14.22	28.833	191.85	16.681	11.739	14.28	22.541	21.448	16.567	17.535	9.065	24.936	33.227	23.741	24.881	37.288
205133_s_at	3336	HSPE1	heat shock 10kDa protei	844.06	2476.6	1665.5	7532.5	669.89	1505.3	2903.3	846.93	783.39	1572.6	605.63	767.69	1753.7	1231	2327	1763.6	188.67
236820_at				59.386	43.757	62.947	48.205	1755	162.4	482.29	413.11	121.42	128.14	60.648	39.065	34.083	55.897	28.843	37.901	114.38
215925_s_at	971	CD72	CD72 molecule	58.037	36.506	120.52	81.1	2131.4	429.63	938.19	245.96	573.81	17.596	140.01	70.636	86.43	25.64	98.177	38.692	20.212
204581_at	933	CD22	CD22 molecule	15.819	9.314	10.67	9.456	4220.4	899.99	1028.4	411.99	546.56	11.084	17.766	12.93	105.23	68.891	59.415	28.518	31.877
216766_at				163.92	86.114	48.873	16.798	1334.1	244.78	244.54	171.12	212.56	22.94	70.721	45.52	19.657	34.482	25.068	27.183	27.135
210432_s_at	6328	SCN3A	sodium channel, voltage-	9.889	14.549	3.845	5.263	722.63	159.39	75.757	124.23	81.037	22.318	9.035	16.917	20.771	16.759	15.008	14.52	30.192
224329_s_at	84518	CNFN	cornifelin	17.525	20.474	15.638	15.914	223.47	42.35	25.314	47.823	161.18	31.814	13.975	11.258	14.451	23.411	16.705	19.943	27.996
39318_at	8115	TCL1A	T-cell leukemia/lymphom	413.37	244.87	53.831	83.168	11712	2673.8	2889.4	498.98	831.45	293.26	180.04	170.35	119.83	297.68	185.86	228.53	348.28
220068_at	29802	VPREB3	pre-B lymphocyte gene 3	155.11	68.624	42.995	57.886	2850.9	755.75	264.61	1466.2	1287.6	178.78	61.713	94.775	90.293	96.477	78.715	97.26	123.4
235460_at	79856	SNX22	sorting nexin 22	47.638	48.233	40.53	35.855	607.56	124.46	101.79	348.5	401.1	104.37	44.03	51.925	27.59	49.441	36.764	35.319	111.72
230385_at				81.96	51.288	73.467	72.773	345.25	86.994	110.34	75.808	75.208	61.51	39.875	54.774	28.204	37.371	32.484	30.718	88.844
209995_s_at	8115	TCL1A	T-cell leukemia/lymphom	641.02	399.85	114.45	76.966	13310	3401.9	2895.9	575.35	902.55	275.69	286.46	251.84	247	312.07	191.15	185.85	392.58
239655_at				915.56	281.78	499.02	263.74	3325	798.08	831.82	778	802.7	432.14	407.7	331.35	113.97	184.05	84.152	122.42	456.53
239292_at				221.07	128.11	19.172	32.89	1833	502.73	422.8	785.39	690.67	272.74	31.909	47.045	11.475	22.142	16.219	13.423	120.6
230245_s_at	283663	LOC28366	hypothetical protein LOC	726.74	166.83	123	126.47	23022	6478.5	4462	6503.1	7265.8	226.35	265.95	221.32	914.55	1401.3	239.32	423.65	219.41
230648_at	283663	LOC28366	hypothetical protein LOC	692.64	453.05	520.19	392.15	13389	4016.6	2195.3	3613.9	4563.9	556.32	476.24	428.56	669.75	1095.6	309.15	406.34	897.64
38521_at	933	CD22	CD22 molecule	390.89	254.78	175.49	153.74	5953.5	1829.8	1631.8	1793.3	2011	266.97	243.9	251.42	697.13	618.3	649.58	454.74	714.06
218949_s_at	55278	QRSL1	glutaminyl-tR synthase (	387.5	465.19	526.61	783.83	2643	875.1	727.76	1464.3	1714.3	621.96	354.9	363.68	464.82	337.89	791.38	630.87	176.49
234284_at	94235	GNG8	guanine nucleotide bindi	53.579	611.56	23.299	184.36	51.681	8705.2	681.42	43.5	64.063	57.214	16.072	16.477	17.853	61.13	14.916	18.059	41.406
212094_at	23089	PEG10	paternally expressed 10	46.498	31.167	66.482	18.336	253.9	2036.6	121.93	81.024	103.77	73.413	54.027	25.162	34.173	64.445	37.859	21.48	90.119
227193_at				743.61	680.67	534.2	492.15	1318.3	7943.9	326.7	1065.5	710.81	847.84	805.39	1089.5	85.611	279.09	740.4	422.82	497.14
223241_at	29886	SNX8	sorting nexin 8	72.247	269.15	243.03	479.83	451.44	3818.5	2789.4	150.52	188.22	240.7	122.51	111.59	513.67	723.39	598.57	235.17	29.861
242468_at				121.46	90.123	93.263	76.315	493.97	2956.6	774.48	232.44	167.55	182.61	91.025	91.24	45.43	58.428	42.262	45.095	185.27
227066_at	148932	MOBKL2C	MOB1, Mps One Binder k	879.19	2018.4	1343.5	1029.6	825.81	13894	717.4	604.51	685.16	895.23	1533.5	3005.6	1422	1492.7	1051.1	937.14	980.55
219517_at	80237	ELL3	elongation factor R polyn	156.69	158.02	95.262	104.13	503.88	2672.8	627.44	196.76	187.92	102	132.33	109.34	241.61	198.86	252.73	569.53	256.98
226274_at	1184	CLCN5	chloride channel 5 (neph	757.15	676.29	495.2	584.13	623.52	4105.2	524.36	1044.6	1151.2	764.8	628.32	554.57	767.28	555.77	891.8	452.43	1058.2
235142_at	653121	ZBTB8	zinc finger and BTB doma	163.85	107.16	57.552	97.531	219.78	2444	154.17	222.47	278.34	440.52	133.21	138.24	97.11	73.722	383.62	117.5	352.18
228231_at				53.024	126.27	68.75	269.56	451.93	3470.4	881.09	127.17	231.04	56.611	98.693	36.689	294.02	279.9	911.37	208.28	38.002
232127_at	1184	CLCN5	chloride channel 5 (neph	155.94	162.89	109.18	181.57	159.9	1331.7	179.77	216.7	184.82	145.23	123.85	106.44	244.96	215.04	243.33	146.17	260.36
227006_at	94274	PPP1R14A	protein phosphatase 1, r	30.346	9.07	9.096	9.518	392.81	2681.6	109.95	196.29	198.03	100.7	11.632	11.142	9.568	26.704	96.818	610.68	22.289
219518_s_at	80237	ELL3	elongation factor R polyn	159.7	130.7	92.114	118.93	420.44	1706	524.31	229.49	264.89	170.12	146.24	169.55	197.8	185.91	202.84	505.74	244.67
209101_at	1490	CTGF	connective tissue growth	22.249	22.905	6.739	3.515	43.341	265.32	34.172	29.803	13.406	12.285	15.781	21.534	11.813	18.578	17.114	41.291	35.862
243052_at	148932	MOBKL2C	MOB1, Mps One Binder k	588.25	1078.3	640.07	622.65	514.18	4626	496.97	611.27	530.52	453.65	829.5	1255.6	710.25	607.66	533.21	447.56	630.08
235278_at	140733	C20orf133	MACRO domain containin	210.22	110.16	58.043	68.735	3510.6	11697	5016.8	124.2	181.28	644.92	79.989	66.195	42.426	76.844	58.566	40.426	140.71
241483_at				177.14	95.259	75.054	51.036	183.49	657.76	44.223	102.93	129.5	188.89	144.92	97.92	42.783	47.617	43.627	37.469	164
223565_at	51237		hypothetical protein MGC	768.21	1072	205.82	123.61	2046.6	1716.7	785.93	740.4	838.1	37183	52.632	104.09	19.182	52.139	32.979	18.784	67.427
211835_at	652070		single-chain Fv fragment	17.024	12.551	9.378	10.146	26.928	11.946	14.494	23.924	21.578	677.99	15.836	17.732	23.55	32.221	33.423	26.31	32.073
233969_at				27.368	28.913	16.45	26.404	135.1	40.005	74.637	57.895	39.922	5202.1	22.518	18.29	18.087	22.137	20.973	43.664	43.749
234792_x_at				476.89	528.5	108.54	42.032	957.48	592.6	712.36	296.25	472.3	32592	92.062	77.439	21.719	38.212	25.846	35.352	64.939
221004_s_at	81618	ITM2C	integral membrane prote	585.42	318.44	534.58	194.83	1031.8	738.28	247.2	577.24	779.46	11437	542.99	323.12	33.527	83.342	50.067	43.392	129.57
221286_s_at	51237		hypothetical protein MGC	506.82	307.29	21.088	21.828	1374.2	1319.5	490.69	260.48	239.29	17479	16.949	17.981	29.661	37.977	33.723	31.289	40.291
206641_at	608	TNFRSF17	tumor necrosis factor rec	481.56	79.328	7.069	4.69	1187.8	365.71	263.56	626.98	442.39	14375	25.957	9.114	21.037	21.426	26.508	35.455	32.888
211641_x_at				669.73	445.68	99.639	82.5	1858	900.13	604.22	1247.4	2371.6	22774	146.08	117.63	154.76	164.45	135.49	142.59	353.57
224342_x_at				1159.5	1240.9	243.81	144.13	4950.9	3547.4	1665.6	1460.3	1353.7	44647	76.908	111.4	74.646	119.1	83.672	98.779	225.79
216412_x_at				132.76	183.85	15.044	25.193	757.95	436.2	527.22	512.07	467.82	6583.2	44.886	41.648	29.585	73.693	31.832	47.005	75.732
219118_at	51303	FKBP11	FK506 binding protein 11	1010.8	1771.9	1287	1715.5	833.49	736.14	618.66	596.84	431.99	13740	951.93	1748.7	153.69	225.52	152.68	102.36	170.24
234419_x_at				107.22	119.57	35.991	31.263	429.37	167.16	108.45	65.809	55.434	4308.9	28.589	30.469	25.767	36.928	23.977	25.044	43.659
217217_at				18.285	15.852	9.172	7.818	47.27	47.245	16.959	37.188	22.782	555.37	14.155	11.369	21.694	31.114	22.223	19.143	23.746
217179_x_at				589.65	458.26	148.22	85.462	3392.6	2245	842.29	1265.4	1225.6	26624	43.581	136.18	116.63	117.3	107.23	160.4	243.01
219003_s_at	79694	MANEA	mannosidase, endo-alpha	87.389	192.19	113.36	190.16	109.12	128.83	61.721	57.43	51.351	1510.6	85.751	124.28	111.99	82.277	196.15	182.86	5.879
217258_x_at				316.7	249.63	21.981	8.441	1613	1120.6	591.59	697.15	670.13	13019	31.014	20.293	21.717	30.131	21.158	20.484	70.638
212097_at	857	CAV1	caveolin 1, caveolae prot	45.356	22.639	15.996	72.569	96.76	64.758	65.577	46.166	24.508	1889.2	14.734	23.886	23.042	136.13	51.28	156.41	46.812

221383_at	10316	NMUR1	neuromedin U receptor 1	16.485	10.581	10.655	9.429	23.474	16.968	11.241	28.366	24.025	21.331	172.26	28.295	24.576	43.296	32.587	28.674	25.14
228948_at	2043	EPHA4	EPH receptor A4	232.57	88.394	232.39	42.03	171.22	92.377	36.349	140.79	81.496	54.073	2761.4	735.14	25.869	40.016	27.074	27.474	82.083
212062_at	10079	ATP9A	ATPase, Class II, type 9A	11.562	14.731	6.817	9.387	12.617	10.289	11.696	19.097	25.438	10.548	948.85	203.48	40.959	55.259	160.09	41.003	27.762
212070_at	9289	GPR56	G protein-coupled recept	450.23	46.85	680.21	378.36	273.85	146.6	17.949	47.686	42.818	31.702	5945.4	1929.6	50.639	140.34	46.329	72.247	180.88
206114_at	2043	EPHA4	EPH receptor A4	304.21	44.721	162.55	12.343	158.35	137.15	20.26	166.81	42.352	38.186	1388.8	396.3	25.82	30.08	30.872	25.258	53.499
217889_s_at	79901	CYBRD1	cytochrome b reductase	7.287	5.64	5.593	5.551	9.848	8.445	7.804	19.536	17.719	8.519	121.89	7.789	21.734	21.921	33.831	21.927	19.441
216050_at				56.285	18.537	102.77	15.889	50.579	25.757	25.26	112.58	61.378	25.283	535.44	147.87	18.017	23.31	13.838	26.439	28.242
228108_at				264.55	96.483	257.25	150.53	209.8	240.56	152.7	259.38	192.84	210.88	1220.4	304.67	157.85	96.387	413.32	116.43	274.48
223126_s_at	81563	C1orf21	chromosome 1 open reac	150.22	48.558	482.41	176.14	151.32	97.005	29.468	93.735	57.454	63.917	2150.6	903.11	84.932	390.11	31.397	23.844	159.08
204457_s_at	2619	GAS1	growth arrest-specific 1	12.055	10.149	9.647	9.038	20.012	12.016	8.361	24.734	25.129	12.794	178.7	35.304	17.643	34.257	42.159	26.652	44.204
226247_at	59338	PLEKHA1	pleckstrin homology dom	2609.8	1355.1	1304.4	665.37	1768.9	1304.9	762.61	848.99	955.81	356.19	7208.2	2925.4	617.2	652.31	909.71	1482.1	442.03
229374_at	2043	EPHA4	EPH receptor A4	900.99	422.48	455.53	254.47	417.9	429.13	273.55	717.74	602.07	596.28	2364.5	967.61	145.92	239.82	138.98	172.69	440
211209_x_at	4068	SH2D1A	SH2 domain protein 1A,	390.85	464.3	737.65	455.64	77.492	72.491	74.097	47.594	46.695	20.921	1693	746.65	51.221	51.22	39.309	41.259	66.209
227449_at	2043	EPHA4	EPH receptor A4	634.35	162.79	615.75	128.7	195.17	190.16	66.138	395.22	109.34	155.65	3974.9	1121.7	19.352	28.697	24.173	24.334	157.95
206355_at	2774	GL	guanine nucleotide bindi	85.932	63.9	41.075	28.466	74.428	43.235	33.021	90.869	85.197	79.187	193.61	90.085	39.564	54.834	62.992	63.343	84.722
243951_at				281.26	147.7	424.32	150.74	410.31	274.03	186.01	228.02	225.67	224.27	1488.5	773.91	78.088	55.655	48.71	46.583	298.63
226279_at	11098	PRSS23	protease, serine, 23	337.78	158.02	147.29	189.44	245.58	359.39	132.8	330.03	377.04	351.5	2740.9	775.99	92.684	108.22	107.02	101.46	407.85
229973_at	127254	C1orf173	chromosome 1 open reac	53.192	88.232	19.387	54.307	43.526	75.933	42.083	112.26	106.76	33.029	96.537	5221.1	20.035	16.235	31.973	19.899	76.112
230926_s_at	57489	ODF2L	outer dense fiber of sper	402.52	524.06	396.31	137.76	355.88	155.38	159.46	183.49	113.26	312.66	469.39	2852.3	29.954	88.016	51.95	64.323	111.68
228577_x_at	57489	ODF2L	outer dense fiber of sper	361.13	401.41	336.7	142.21	219.51	187.77	125.38	240.69	116.48	274.85	401.76	2054.4	33.763	61.725	39.819	107.49	86.458
229893_at	257019	FRMD3	FERM domain containing	304.9	315.32	121.07	156.3	292.62	214.85	117.87	312.93	474.35	213.8	284.1	1797.8	111.72	76.125	133.01	202.21	294.68
203571_s_at	10974	C10orf116	chromosome 10 open rea	13.1	8.346	7.777	5.934	22.182	27.798	11.084	22.65	20.283	15.804	11.696	225.5	19.308	40.143	28.946	25.319	23.581
210218_s_at	6672	SP100	SP100 nuclear antigen	301.3	711.82	491.39	248.45	889.83	574.84	396.32	225.06	197.43	120.03	990.02	3884	572.42	755.13	276.02	661.01	630.04
219684_at	64108	RTP4	receptor (chemosensory)	156.03	1646.5	521.18	124.92	92.382	560.78	58.119	59.991	60.383	224.17	836.07	6115.6	346.26	438.55	577.13	1663.7	64.385
224973_at	55603	FAM46A	family with sequence sim	209.46	254.63	56.082	30.957	262.27	338.22	140.46	70.072	70.491	96.183	750.31	2775.4	406.81	129.88	401.53	597.47	211.98
228531_at	54809	SAMD9	sterile alpha motif domai	878.92	5053.9	3029.7	540.35	1463	1883.7	500	389.29	544.59	1078.5	4316.6	19829	471.56	584.64	274.6	2650.4	1593.1
230645_at	257019	FRMD3	FERM domain containing	290.98	504.45	214.86	121.57	355.76	267.45	226.52	846.65	811.22	387.72	223.91	3689.8	94.875	77.308	140.16	266.24	748.24
226103_at	91624	NEXN	nexilin (F actin binding p	67.656	627	67.589	118.05	87.618	101.62	150.77	82.89	38.41	30.593	115.11	2299.9	17.823	28.107	16.964	51.955	64.69
236156_at	3988	LIPA	lipase A, lysosomal acid,	34.715	451.51	34.551	44.167	32.135	29.07	32.918	52.246	56.176	76.157	21.641	2082.2	39.556	27.231	22.498	96.451	70.775
210705_s_at	85363	TRIM5	tripartite motif-containin	177.65	345.55	336.22	108.71	330.94	318.82	274.03	143.37	183.16	317.66	431.34	1212.8	328.47	236.23	326.44	417.5	213.38
218501_at	50650	ARHGEF3	Rho guanine nucleotide e	1839.8	2785.7	3143.2	928.31	802.1	999.06	266.06	374.46	472.91	440.33	4649.5	10693	808.73	521.37	1982.5	1260	265.32
228675_at				321.04	662.29	499.97	368.98	350.57	250.69	163.87	200.3	148.86	637.26	430.83	2257.5	109.55	72.205	85.077	413.35	267.85
236191_at				190.68	1376.2	626.48	604.37	248.31	303.43	158.5	49.547	60.247	882.36	259.52	4757.8	25.85	56.1	24.379	190.18	128.97
223434_at	2635	GBP3	guanylate binding protein	704.07	2327.8	2251.5	1177	507.74	597.15	260.27	260.49	282.58	680.29	3614.4	9689.2	235.89	408.95	477.37	1973.1	442.26
201185_at	5654	HTRA1	HtrA serine peptidase 1	298.66	211.29	129.67	158.13	244.18	325.94	217.11	254.78	217.39	378.02	182.02	100.41	4524.5	721.17	344.37	393.46	408.27
211165_x_at	2048	EPHB2	EPH receptor B2	18.128	14.747	13.441	10.721	22.435	19.212	13.813	41.378	38.509	19.184	36.082	33.919	750	37.251	66.137	111.51	40.488
211913_s_at	10461	MERTK	c-mer proto-oncogene ty	170.5	109.41	114.19	123.07	217.36	151.04	110.88	131.29	147.68	167.79	155.64	143.83	2057.5	444.14	347.38	196.23	221.36
208771_s_at	4048	LTA4H	leukotriene A4 hydrolase	2762.8	3065.3	1893.5	2088.7	5716.5	5135.1	4152	2095.9	2279.2	3178.3	2278	832.7	21704	2697.1	4330.1	1290.8	1902.3
209589_s_at	2048	EPHB2	EPH receptor B2	30.863	44.763	17.499	32.405	59.756	19.794	14.376	68.248	49.623	86.171	54.592	51.961	1348.1	30.996	332.32	326.9	110.81
206028_s_at	10461	MERTK	c-mer proto-oncogene ty	190.69	107.8	58.619	53.798	117.56	104.65	56.653	236.55	207.85	107.1	126.88	50.745	3131.1	768.85	422.35	132.67	247.23
227823_at	340526	RGAG4	retrotransposon gag dom	78.208	111.4	61.529	63.163	103.33	94.134	75.628	96.029	94.257	95.477	112.3	123.43	933.54	233.76	297.81	126.5	147.74
235751_s_at	284013	VMO1	vitelline membrane outer	34.258	39.726	11.115	11.268	16.153	14.451	19.847	23.683	20.328	23.967	362.26	11.41	2665.1	396.29	97.002	261.12	21.038
236646_at	120939	C12orf59	chromosome 12 open rea	47.859	43.294	34.943	30.558	36.092	85.694	45.534	69.956	54.285	134.15	72.249	101.61	821.29	191.39	106.19	115.88	143.71
218832_x_at	408	ARRB1	arrestin, beta 1	34.25	29.224	38.111	72.62	31.272	19.392	15.134	37.238	34.008	20.44	144	87.986	591.46	169.9	179.32	77.117	79.13
209588_at	2048	EPHB2	EPH receptor B2	197.37	142.57	136.13	141.82	204.37	159.93	109.4	240.71	226.39	202.71	117.7	98.776	1139.8	190.47	393.61	362.3	378.24
202838_at	2517	FUCA1	fucosidase, alpha-L- 1, ti	242.55	302.08	495.6	216.27	440.56	450.22	469.94	416.64	325.97	713.21	2269	1457.2	23621	1428.6	7804.2	2832.3	348.69
208862_s_at	51075	TXNDC14	thioredoxin domain conta	170.59	108.76	126.74	100.47	271.42	210.77	199.36	122.69	123.37	38.037	149.94	79.806	2508.8	1320.1	1363.3	601.22	135.64
212464_s_at	2335	FN1	fibronectin 1	26.448	27.259	17.197	26.945	30.48	25.495	42.183	81.327	103.35	16.885	366.11	53.143	16242	58.443	2883.5	2906	165.13
204787_at	11326	VSIG4	V-set and immunoglobuli	233	137.81	120.47	81.189	213.86	154.35	61.94	352.53	358.35	198.53	205.15	68.283	3604.2	370.12	157.55	209.64	455.5
210495_x_at	2335	FN1	fibronectin 1	70.789	23.169	29.882	44.304	59.421	56.337	17.899	52.106	56.731	16.433	698.9	104.41	19697	64.194	4322.9	4511	73.793
221565_s_at	51063	FAM26B	family with sequence sim	314.56	124.06	776.91	70.875	230.96	151.41	104.78	139.07	174.98	129.8	874.44	374.83	2336.4	451.57	899.63	218.93	379.34
219525_at	55244	FLJ10847	solute carrier family 47,	18.184	27.079	14.001	17.424	21.488	30.656	13.634	24.757	22.461	30.17	18.268	31.408	34.706	31.818	1294.5	62.187	37.893
203305_at	2162	F13A1	coagulation factor XIII, A	289.19	154.99	124.88	105.2	313.25	233.58	115.91	479.91	402.89	247.61	199.86	162.12	301.41	267.62	9213.4	463.63	663.21
206682_at	10462	CLEC10A	C-type lectin domain fam	117.66	77.553	54.795	58.318	113.97	116.72	20.433	119.71	111.38	104.15	242.18	51.339	666.23	247.07	10346	234.87	219.09
207328_at	246	ALOX15	arachidonate 15-lipoxyge	10.612	25.339	6.831	6.378	13.36	11.346	9.715	19.972	58.372	10.044	27.642	27.956	26.953	29.212	4539.9	72.346	182.96
204044_at	23475	QPRT	quinolinate phosphoribos	22.06	16.746	54.589	25.22	24.631	16.766	31.21	87.88	56.88	141.4	27.911	18.443	39.471	38.229	3854.3	108.41	77.585
237731_at				40.498	30.122	31.198	25.411	32.529	28.516	36.734	69.953	64.71	44.851	23.433	33.679	57.564	49.195	1113.4	122.32	43.176
203619_s_at	23017	FAIM2	Fas apoptotic inhibitory r	53.577	29.106	14.683	16.222	41.65	21.3	18.821	28.089	25.181	28.634	18.124	15.246	38.708	44.951	1986	47.632	35.496
220301_at	79839	CCDC102E	coiled-coil domain contai	13.112	8.355	4.523	5.864	5.023	8.302	1.015	37.086	9.202	8.124	16.601	9.295	16.732	22.858	224.25	24.244	10.698
201876_at	5445	PON2	paraoxonase 2	153.68	113.96	189.29	208.72	203.37	121.68	97.311	120.96	128.04	130.15	434.92	340.34	276.18	103.43	5310.8	766.1	141.11

207277_at	30835	CD209	CD209 molecule	220.38	169.13	83.777	91.853	207.13	163.27	99.57	313.17	251	187.27	118.57	188.32	365.69	835.46	7460.1	742.95	370.95
222240_s_at	51477	ISY1	myo-inositol 1-phosphate	285.49	212.26	179.91	183.44	292.53	198.33	85.288	65.867	135.68	52.564	172.47	121.04	178.53	181.87	2664.7	426.65	120.38
207278_s_at	30835	CD209	CD209 molecule	106.67	65.52	19.824	18.065	59.958	52.787	21.539	35.029	47.617	24.918	40.149	53.679	90.689	154.74	1321.4	166.3	140.49
201348_at	2878	GPX3	glutathione peroxidase 3	207.97	169.7	117.12	135.49	237.29	182.73	161.49	414.11	438.13	229.3	289.86	166.4	2820.8	773.89	17689	2021.5	666.64
214091_s_at	2878	GPX3	glutathione peroxidase 3	202.77	174.29	157.95	111.93	261.65	236.72	154.17	295.06	301.73	234.74	290.97	235.63	1813.8	724.23	10819	1542.4	612.69
210830_s_at	5445	PON2	paraoxonase 2	142.99	90.515	112.25	122	90.488	64.661	62.882	113.24	149.34	50.288	188.76	161.04	168.75	41.48	3833.2	579.46	69.345
223939_at	56670	SUCNR1	succinate receptor 1	181.08	728.19	164.8	113.29	340.37	277.13	142.74	423.79	260.43	305.45	166.02	133.08	158.29	517.6	4885.7	822.66	343.58
213230_at	30850	CDR2L	cerebellar degeneration-r	14.937	10.283	11.864	17.984	14.459	8.9	13.077	13.102	13.026	10.069	15.938	20.626	54.75	40.053	904.74	77.118	39.601
209183_s_at	11067	C10orf10	chromosome 10 open rea	37.582	61.905	18.124	17.372	28.084	23.813	25.516	43.537	39.406	29.808	24.176	45.143	33.926	63.693	95.817	2692.8	36.858
201564_s_at	6624	FSCN1	fascin homolog 1, actin-b	21.011	314.03	33.576	161.6	16.286	63.516	55.39	54.472	29.717	24.173	81.53	523.84	136.01	184.31	670.32	17366	98.237
218596_at	54662	TBC1D13	TBC1 domain family, mer	76.209	49.734	69.905	50.81	70.849	141.92	114.81	54.299	51.949	51.197	98.944	262.08	96.11	188.52	289.66	5087.5	56.742
237154_at				333.43	340.39	283.22	277.53	237.07	137.2	75.169	147.68	206.61	305.57	150.12	378.01	86.24	121.02	294.11	15782	284.28
205226_at	5157	PDGFRL	platelet-derived growth f	36.427	9.53	14.258	12.599	34.964	16.96	44.181	40.21	22.475	13.408	28.375	53.462	47.721	49.514	53.746	1387.7	20.595
202411_at	3429	IFI27	interferon, alpha-inducib	162.25	912.04	119.25	65.958	75.112	112.25	46.373	50.78	94.163	119.78	134.04	1488.9	210.18	325.75	138.05	17765	151.87
205890_s_at	10537	UBD	ubiquitin D	78.773	668.53	20.265	24.835	25.86	39.337	23.056	114.36	60.898	40.013	74.727	153.21	53.552	33.817	95.649	8820.3	107.24
227970_at	80045	GPR157	G protein-coupled recept	1127.6	936.33	442.99	373.89	792.28	765.69	864.41	672.41	417.83	1021.2	990.63	1117.5	941.2	500.12	390.29	9736.3	730.17
224399_at	80380	PDCD1LG2	programmed cell death 1	37.157	39.408	26.364	31.244	35.784	26.589	26.191	63.203	53.796	59.15	21.387	27.365	24.291	29.793	122.17	1974.3	53.788
209182_s_at	11067	C10orf10	chromosome 10 open rea	86.486	66.895	25.297	43.289	87.619	102.61	23.724	58.954	61.644	63.256	76.594	37.144	76.86	55.657	65.272	1186.5	89.081
211267_at	8820	HESX1	HESX homeobox 1	15.962	20.309	9.002	16.894	20.791	22.89	13.287	26.034	26.317	17.264	13.973	59.359	33.421	25.657	40.231	1691.5	22.801
210933_s_at	6624	FSCN1	fascin homolog 1, actin-b	20.332	566.4	33.336	168.56	71.603	102.41	67.491	10.366	6.506	7.758	126.4	620.29	171.77	344.98	1454.9	16899	52.809
208782_at	11167	FSTL1	follistatin-like 1	192.12	120.83	113.92	109.45	195.78	158.41	118.18	353.01	375.64	162.49	93.606	105.83	192.63	274.55	188.9	4875.7	381.91
44696_at	54662	TBC1D13	TBC1 domain family, mer	624.15	537.35	501.78	399.81	665.39	730.1	446.29	528.19	571.03	405.15	582	689.43	770.72	899.75	1147.7	7639.7	681.05
211083_s_at	9175	MAP3K13	mitogen-activated protei	6.504	4.804	3.886	3.113	5.906	5.464	4.814	8.781	9.781	11.148	9.844	6.678	11.997	17.73	14.728	173.95	11.757
206638_at	3357	HTR2B	5-hydroxytryptamine (se	34.96	28.74	11.051	3.945	26.466	10.004	15.287	25.989	17.253	29.89	44.103	23.668	20.289	20.805	23.311	815.72	23.146
223741_s_at	94015	TTYH2	tweety homolog 2 (Drosc	607.39	287.84	245.13	104.92	237.65	192.38	136.59	76.097	105.62	369.95	403.48	548.11	194.02	123.57	283.43	7052.4	203.15
220322_at	56300	IL1F9	interleukin 1 family, mem	91.241	91.419	44.993	41.394	72.339	78.498	53.339	171.73	175.77	85.443	75.673	95.566	193.5	13364	96.662	140.58	158.08
207316_at	3036	HAS1	hyaluronan synthase 1	83.836	49.02	39.758	51.693	82.304	62.179	22.661	135.41	153.81	107.37	52.122	23.743	85.531	6520.7	125.32	73.135	144.91
209278_s_at	7980	TFPI2	tissue factor pathway inh	15.299	25.449	6.514	7.562	15.212	135.04	88.061	22.203	21.938	12.314	14.302	264.75	228.32	10802	22.39	274.11	37.116
206569_at	11009	IL24	interleukin 24	85.352	83.076	38.318	38.162	251.23	72.213	118.18	67.58	126.55	22.751	89.089	27.311	98.807	7265	37.723	43.914	230.31
204475_at	4312	MMP1	matrix metallopeptidase	9.388	26.327	2.994	3.6	8.573	16.557	7.346	33.165	45.886	6.634	14.214	3.569	15.962	2464.7	22.158	26.477	38.896
209277_at	7980	TFPI2	tissue factor pathway inh	25.095	20.044	10.747	5.447	35.849	74.723	52.423	70.474	69.994	34.393	31.933	59.772	146.36	3886.6	28.623	142.85	22.053
207442_at	1440	CSF3	colony stimulating factor	18.01	24.088	13.039	11.862	29.41	19.209	19.475	26.511	26.774	20.875	47.925	24.496	27.293	2163.5	30.881	31.705	36.188
205207_at	3569	IL6	interleukin 6 (interferon,	207.34	440.66	94.669	75.292	320.72	1588	284.03	341.25	435.91	145.62	97.655	1336.6	1632.4	34971	204.43	2510.5	240.88
203510_at	4233	MET	met proto-oncogene (hep	11.918	12.065	5.665	56.352	11.912	10.115	35.396	24.648	16.854	32.194	15.715	16.009	360.15	4883.1	27.917	148.52	23.036
229435_at	169792	GLIS3	GLIS family zinc finger 3	51.725	42.231	9.847	76.931	13.202	16.12	14.462	83.659	111.02	64.578	13.486	37.029	185.27	1833	46.519	15.028	127.91
220014_at	51334	PRR16	proline rich 16	11.96	4.882	2.492	0.985	3.236	8.931	22.338	34.53	23.283	19.207	19.619	10.144	72.589	812.71	19.002	17.112	43.945
207852_at	6374	CXCL5	chemokine (C-X-C motif)	31.673	28.613	23.431	24.093	36.395	9.425	17.418	54.262	77.543	38.061	9.621	15.056	175.25	1677.9	40.346	43.681	99.159
207336_at	6660	SOX5	SRY (sex determining reg	9.836	7.861	7.661	6.475	29.154	8.14	11.336	19.539	23.041	16.023	9.393	10.435	26.932	296.21	30.578	32.724	14.187
220655_at	79931	TNIP3	TNFAIP3 interacting prot	30.483	496.06	87	200.55	28.186	19.561	27.081	64.546	62.118	37.863	13.75	34.176	660.08	8318.6	49.416	180.07	35.132
206421_s_at	8710	SERPINB7	serpin peptidase inhibitor	31.11	25.351	15.255	11.639	31.797	20.7	20.595	48.02	44.575	24.085	26.193	269.05	48.317	8770.5	32.96	803.26	45.827
204627_s_at	3690	ITGB3	integrin, beta 3 (platelet	77.937	52.599	25.61	12.752	87.344	100.16	79.679	27.204	28.815	13.054	82.961	102.15	50.385	2003.3	53.605	172.65	112.65
204748_at	5743	PTGS2	prostaglandin-endoperox	75.522	58.612	20.681	22.695	16.161	26.012	21.551	28.498	43.676	25.445	22.052	57.425	434.21	23846	79.633	293.33	2414
211163_s_at	8794	TNFRSF10	tumor necrosis factor rec	24.745	17.582	11.933	10.705	16.047	12.525	12.11	37.874	17.543	9.775	23.394	13.984	40.419	75.161	46.084	27.389	7494
220005_at	53829	P2RY13	purinergic receptor P2Y, (	20.207	15.818	7.568	6.407	29.631	76.257	13.312	70.916	33.945	22.424	15.617	12.05	22.463	29.8	106.12	24.156	7096
203435_s_at	4311	MME	membrane metallo-endop	52.68	35.537	27.734	32.273	47.653	44.56	32.565	106.81	102.32	68.168	45.207	25.599	56.171	108.62	66.398	51.768	5834.4
229967_at	146225	CMTM2	CKLF-like MARVEL transm	404.54	263.4	375.27	305.18	320.72	272.32	165.77	841.78	212.16	494.7	260.95	291.93	161.55	189.59	204.06	184.9	41610
234644_x_at				176.1	142.03	83.535	142.24	49.475	43.257	26.589	45.615	59.935	125.79	95.751	43.935	27.844	33.083	31.309	39.868	11321
206222_at	8794	TNFRSF10	tumor necrosis factor rec	122.04	82.36	80.849	72.071	101.24	94.444	66.289	143.19	149.05	94.772	69.516	66.842	124.52	98.004	132	127.39	7001.1
204351_at	6286	S100P	S100 calcium binding pro	341.03	96.34	29.987	26.748	230.68	299.56	120.18	69.519	41.807	30.985	35.679	40.597	196.07	287.74	71.192	50.918	21441
207094_at	3577	IL8RA	interleukin 8 receptor, alp	55.571	34.928	30.255	30.118	51.796	46.036	23.438	66.93	68.085	27.242	150.19	36.452	38.777	47.861	34.59	34.741	6702.9
232629_at	60675	PROK2	prokineticin 2	25.175	26.004	158.35	155.29	82.893	49.74	28.545	24.378	16.613	53.691	27.394	7.688	94.419	58.926	11.755	29.657	13698
220945_x_at	54682	MANSC1	MANSC domain containin	75.87	57.915	31.179	33.702	98.717	103.16	34.426	291.74	249.55	97.194	32.466	53.104	75.79	54.837	60.247	61.276	2613.4
229770_at	144423	GLT1D1	glycosyltransferase 1 dor	48.013	28.57	27.015	34.934	109.92	33.965	25.369	54.365	48.216	45.415	97.688	90.763	406.39	290.12	29.735	19.451	12243
207008_at	3579	IL8RB	interleukin 8 receptor, be	48.606	7.934	13.365	4.607	34.921	15.821	19.504	13.723	15.532	19.566	392.19	100.84	111.39	20.749	453.51	83.791	12028
218963_s_at	25984	KRT23	keratin 23 (histone deace	21.721	25.964	13.516	29.433	77.001	62.235	52.891	151.99	138.74	106.63	36.835	49.514	68.24	67.635	90.373	68.849	3796.8
204007_at	2214	FCGR3A	Fc fragment of IgG, low a	420.69	197.38	238.39	130.46	284.33	180.1	91.841	248.53	239.36	162.95	901.02	693.72	1388	330.29	466.68	231.55	32749
203434_s_at	4311	MME	membrane metallo-endop	21.906	20.104	7.349	5.374	33.004	64.683	12.237	32.656	30.344	18.396	18.525	13.907	21.308	93.096	27.296	25.716	2580
220421_at	79908	BTNL8	butyrophilin-like 8	18.42	11.019	9.26	19.771	28.565	22.44	13.582	25.459	26.706	26.866	19.269	15.886	38.278	55.531	50.442	42.432	1195.4
210484_s_at	254896		hypothetical protein MGC	86.434	63.098	71.649	52.289	87.488	50.563	44.979	99.362	61.429	110.7	56.534	90.265	162.63	125.69	139.7	144.21	2915.3
240347_at				1854.5	61.454	48.362	44.29	999.91	57.159	47.999	62.948	64.1	66.571	43.711	36.304	28.181	36.962	26.687	31.636	101.33

227614_at	80201	HKDC1	hexokinase domain conta	442.35	35.677	47.464	21.902	45.899	34.716	31.414	65.151	73.108	71.596	26.961	45.619	23.442	29.542	32.629	24.274	58.015
204760_s_at	9572	NR1D1	nuclear receptor subfami	297.62	19.749	6.99	17.411	66.318	13.655	10.1	11.711	9.278	13.786	10.559	20.107	16.677	28.563	25.355	25.463	33.51
234967_at	3572	IL6ST	interleukin 6 signal trans	500.94	54.356	55.673	34.638	18.455	33.371	45.084	127.48	253.1	93.452	18.648	75.83	17.295	19.91	12.723	20.567	66.204
226342_at	6711	SPTBN1	spectrin, beta, non-eryth	1419.8	51.589	46.397	44.595	332.99	37.121	54.203	108.92	253.63	122.46	40.456	30.066	29.822	33.681	25.029	28.475	88.178
226388_at	6920	TCEA3	transcription elongation f	1084.5	225.21	238.7	62.339	68.575	50.839	46.818	75.369	171.03	152.37	116.74	24.173	42.689	80.724	58.11	61.04	198.9
215262_at				1279.9	295.26	403.78	149.61	92.396	109.06	137.22	125.98	246.18	43.598	125.04	63.024	38.942	29.201	84.617	132.45	136.94
221790_s_at	26119	LDLRAP1	low density lipoprotein re	2504.4	344.2	835.09	62.217	197.35	98.984	173	135.23	159.29	284.72	512.32	232.72	316.47	283.63	570.71	174.09	140.7
230304_at				2596.8	466.59	585.78	390.5	474.34	448.8	270.5	136.42	258	408.3	1183.6	722.03	174.49	97.182	94.599	65.5	229.94
207906_at	3562	IL3	interleukin 3 (colony-stin	11.796	382.21	7.019	134.51	10.679	10.166	9.046	36.347	16.231	9.837	9.531	7.443	16.465	22.998	20.151	21.977	18.31
222974_at	50616	IL22	interleukin 22	51.201	3661.6	26.909	880.39	42.27	53.862	16.571	102.85	70.52	146.28	86.691	40.447	29.994	57.904	25.763	21.935	91.888
227984_at				1219.7	1270.8	138.68	191.49	111.82	123.14	198.43	173.84	193.14	182.04	146.1	153.11	42.787	41.659	142.33	46.938	114.22
232584_at				1125	2341.9	195.23	381.74	356.46	283.78	174.75	205.37	190.32	240.26	149.76	123.63	72.559	162.63	116.99	71.606	208.72
207892_at	959	CD40LG	CD40 ligand (TNF superfa	492.38	838.31	67.853	167.51	104.76	46.611	20.655	64.876	58.402	42.182	40.464	49.649	34.919	30.912	38.454	28.851	88.189
224211_at	50943	FOXP3	forkhead box P3	322.53	1418.7	167.46	635.89	89.541	65.338	48.38	220.77	206.11	146.06	127.15	81.559	32.344	77.866	46.594	46.038	122.99
224801_at				562.06	2564.1	486.91	2284.5	335.08	441.29	350.3	661.08	704.09	614.13	370.94	370.3	183.16	259.95	348.39	267.85	711.33
236341_at	1493	CTLA4	cytotoxic T-lymphocyte-a	1240.2	4434.9	489.94	1618.8	53.729	52.941	16.366	134.12	85.046	151.54	9.094	48.237	37.288	50.691	101.61	64.781	132.91
220485_s_at	55423	SIRPG	signal-regulatory protein	1279.7	344.58	1870.4	377.07	176.03	122.11	36.387	183.64	157.75	97.431	105.48	80.092	107.77	138.53	127.52	151.82	220.38
206804_at	917	CD3G	CD3g molecule, gamma (	2246.6	916.04	3622.2	876.21	201.71	87.324	19.905	53.472	55.177	32.644	477.55	288.48	44.127	84.826	45.794	46.494	83.39
206974_at	10663	CXCR6	chemokine (C-X-C motif)	147.04	969.58	1662.6	350.63	135.37	191.12	73.848	262.77	265.17	136.31	406.47	375.57	152.71	177.96	163.36	154.32	292.65
230218_at	3090	HIC1	hypermethylated in canc	37.26	320.67	1359.6	1251.4	43.644	62.872	53.679	62.066	56.536	43.042	190.21	119.43	47.963	42.613	46.522	177.53	73.215
205484_at	27240	SIT1	signaling threshold regul	823.62	686.41	2063	786.65	1305.4	447.22	556.99	658.06	732.74	394.33	199.48	132.05	101.21	123.34	407.85	119.45	174.4
233857_s_at	51676	ASB2	ankyrin repeat and SOCS	380.33	402.91	2482.4	1135.9	242.59	289.52	203.62	242.59	200.29	612.31	274.66	272.28	135.17	143.8	94.268	172.17	376.9
227903_x_at	91978	C19orf20	chromosome 19 open rea	147.3	83.667	652.05	427.54	76.829	56.022	175.17	42.183	46.635	136.02	53.805	54.227	136.66	121.66	142.86	168.85	43.394
209670_at				9535.5	4889.7	11371	4324.1	989.81	734	297.26	509.97	406.85	471.19	786.27	699.43	305.97	565.94	236.82	296.6	844.63
204890_s_at	3932	LCK	lymphocyte-specific prote	4328.6	1960.6	5769.7	2096.1	582.59	401.34	265.26	313.1	297.33	85.982	2360.8	993.94	268.64	389.46	154.42	148.7	415.79
219836_at	79413	ZBED2	zinc finger, BED-type con	153.9	3485.5	158.99	5721.5	244.85	328.8	549.38	314.44	244.52	123.05	130.91	87.84	77.241	139.93	117.45	87.879	257.46
204695_at	993	CDC25A	cell division cycle 25 hom	13.246	24.781	88.757	868.1	16.454	12.278	11.699	27.212	27.494	47.288	12.808	8.661	21.27	35.54	28.316	18.804	23.371
241682_at	151230	KLHL23	kelch-like 23 (Drosophila	60.364	94.954	220.96	1774.3	72.694	58.921	60.413	79.842	48.502	36.977	86.999	91.376	27.185	36.326	19.515	22.162	51.824
225655_at	29128	UHRF1	ubiquitin-like, containing	236.33	1119.1	3816.9	11233	321.11	227.75	796.15	223.23	171.58	1438.7	184.73	95.148	68.223	51.556	36.754	33.832	223.99
242890_at				233.92	290.59	813.35	3551	308.18	279.31	439.72	282.41	256.85	363.93	137.58	95.521	32.602	48.19	34.163	57.902	99.755
223229_at	29089	UBE2T	ubiquitin-conjugating enz	174.69	798.98	1705.3	6581	117.92	255.28	393.67	100.27	84.534	842.79	137.04	110.9	41.216	64.275	62.825	50.933	53.295
220085_at	3070	HELLS	helicase, lymphoid-specil	47.141	87.792	454.04	1303.1	93.846	106.47	136.72	41.992	43.732	61.13	6.926	6.673	16.983	29.67	25.008	24.039	28.993
204558_at	8438	RAD54L	RAD54-like (S. cerevisiae	37.236	27.188	174	529.88	29.891	30.037	23.372	35.105	31.035	17.64	55.063	35.177	56.263	47.656	48.391	46.296	57.866
227350_at	3070	HELLS	helicase, lymphoid-specil	284.84	423.31	1327.9	3798.5	346.27	421.1	591.66	309.23	325.53	424.17	176.76	154	34.591	65.708	34.752	50.621	298.73
205242_at	10563	CXCL13	chemokine (C-X-C motif)	19.513	1354.7	29.476	1874.5	18.396	16.105	190.19	60.868	41.46	23.525	28.792	94.868	25.984	158.1	27.05	242.1	41.323
219000_s_at	79075	DCC1	defective in sister chroma	41.096	98.875	266.43	1054.4	34.983	45.943	45.53	38.164	54.297	115.2	61.206	23.793	101.91	136.67	98.441	95.428	61.939
206975_at	4049	LTA	lymphotoxin alpha (TNF s	36.869	2186	19.704	2896.6	23.214	305.93	19.847	22.388	21.473	14.365	18.138	118.74	20.557	29.577	27.044	28.376	32.132
222504_s_at	10328	COX4NB	COX4 neighbor	27.346	73.761	32.207	695.77	25.677	34.441	63.591	60.775	60.12	98.738	29.093	23.251	36.363	39.324	34.294	22.123	37.046
235401_s_at	84824	FCRLA	Fc receptor-like A	130.31	41.849	42.774	32.087	3782.4	1105.1	2016.2	1131.8	1993.1	118.29	45.306	35.399	33.046	52.991	28.414	16.273	85.514
230877_at	3495		immunoglobulin heavy co	668.07	294.22	50.476	56.156	21387	11410	6419.6	332.54	1278.4	329.1	45.323	42.148	14.306	59.328	16.55	23.564	65.166
227198_at				533.25	92.106	343.14	118.34	9753.7	6992	4487.8	1347.4	2163	17.917	278.65	183.3	30.892	227.31	28.426	22.836	36.823
219667_s_at	55024	BANK1	B-cell scaffold protein wi	131.67	62.309	17.332	13.043	5063.1	3369.7	1609	4849.6	4389	269.4	87.887	93.53	35.155	98.414	81.002	42.046	96.465
217418_x_at	931	MS4A1	membrane-spanning 4-d	756	462.64	129.79	55.3	13482	12327	7914.7	12484	14351	330.25	161.02	229.51	115.45	406.1	130.5	190.71	468.78
243780_at				743.39	815.94	246.9	284.77	15139	11615	7550.3	4127.6	4829.9	1153.1	188.29	83.886	44.602	109.46	38.954	42.475	172.67
210356_x_at	931	MS4A1	membrane-spanning 4-d	909.27	610.61	159.55	92.926	15292	14082	8618.6	14009	16289	294.53	225.68	328.16	265.38	525.55	198.48	283.11	571.83
230983_at	199786	BCNP1	family with sequence sim	1149.9	781.16	583.52	548.37	15403	7850.1	4753.7	3450	6123.6	822.34	702.73	706.12	256.69	301.58	217.31	334.95	1098.2
243968_x_at	115350	FCRL1	Fc receptor-like 1	416.95	132.19	110.29	114.67	10113	3156.3	1094.1	1986.9	2423.5	875.88	108.17	132.07	64.362	69.389	74.8	93.924	289.88
232286_at				415.85	176.56	284.27	153.54	7089.1	3593	3206.7	1259.2	1392.3	207.42	191.43	156.86	97.122	121.18	138.05	110.89	567.87
228599_at	931	MS4A1	membrane-spanning 4-d	291.55	183.21	166.4	186.99	8517.3	10792	5230.7	3721.6	4253.6	312.53	80.829	114.15	101.59	113.37	101.97	103.15	253.83
228592_at	931	MS4A1	membrane-spanning 4-d	1316.4	1018.6	104.4	58.217	33100	35052	20987	23579	26239	395.68	326.13	440.54	128.31	417.2	81.803	158.62	326.6
222891_s_at	53335	BCL11A	B-cell CLL/lymphoma 11/	338.58	370.66	69.794	59.023	13141	14169	6339.9	3436.7	3985.5	528.8	278.77	344.03	188.3	1126.3	142.9	935.66	524.55
205933_at	26040	SETBP1	SET binding protein 1	164.27	180.62	200.7	219.58	1157	3083.3	786.02	716.76	599.74	526.88	182.44	205.55	75.283	105.32	347.1	146.98	127.52
219498_s_at	53335	BCL11A	B-cell CLL/lymphoma 11/	138.73	133.94	25.675	41.093	2742.6	3364.6	1258.2	1038.8	1261.6	187.64	51.388	86.743	146.78	603.73	84.55	594.52	212.77
217084_at				41.785	67.193	9.726	5.421	163.95	47.628	32.218	45.183	41.792	1278.4	6.244	13.577	15.288	17.852	17.773	16.565	14.048
211642_at				13.19	15.156	7.21	7.64	141.74	45.987	36.33	21.507	51.597	827.93	8.998	9.784	20.519	27.983	23.596	15.285	39.095
217378_x_at				968.69	815.12	23.298	23.718	5042.3	2888.9	2118.1	2838.8	3291.2	34965	22.09	30.102	44.255	58.244	47.213	46.657	53.954
214777_at				443.22	466.97	17.373	12.526	4208.7	2671.2	1577.3	1877	2561.6	34431	25.025	27.54	26.512	58.57	29.03	25.412	43.521
203065_s_at	857	CAV1	caveolin 1, caveolae prot	78.802	62.756	35.597	78.088	88.197	45.911	78.45	119.82	124.39	1721.2	32.745	47.461	43.329	83.562	51.181	114.21	120.86
216557_x_at				677.66	476.06	157.64	134.86	2782.3	1358.7	976.07	737.17	1560.2	21772	156.03	222.68	145.78	228.12	167.6	153.98	378.45
232478_at				21.964	34.023	28.871	23.483	28.206	39.402	43.26	60.684	63.493	39.157	1357.6	1006	18.416	25.441	20.915	20.658	43.551

221417_x_at				15.586	12.486	8.636	8.962	16.528	11.045	13.645	18.64	16.741	10.684	636.77	472.14	16.048	20.789	17.434	18.811	19.978
233743_x_at				67.846	44.061	34.084	28.856	38.968	48.293	33.393	82.363	80.263	80.665	1050.1	856.24	18.334	27.73	21.672	26.124	41.053
230464_at				717.93	282.15	666.48	162.36	414.42	319.18	33.136	75.413	64.085	83.253	10394	9089.6	32.645	56.232	50.011	38.707	78.204
201681_s_at	9231	DLG5	discs, large homolog 5 (D	48.54	24.378	25.433	19.693	44.627	16.9	23.514	39.08	38.174	37.877	593.45	203.75	22.159	47.671	29.067	22.446	35.15
225688_s_at	257068	PLCXD2	phosphatidylinositol-spec	357.59	187.77	433.56	256.29	505.27	522.21	174.05	345.23	277.7	248.14	3883.5	2824.8	57.803	75.66	44.914	112.46	259.01
219383_at	79899	FLJ14213	hypothetical protein FLJ1	180.15	197.03	170.88	145.01	166.91	153.61	34.386	135.85	178.07	56.217	1486.1	742.55	54.253	57.897	48.31	112.77	211.31
204731_at	7049	TGFBR3	transforming growth fact	533.63	343.64	541.31	190.35	192.1	131.58	96.953	183.11	213.56	161.31	4462.7	3660.1	148.41	166.84	82.078	112.41	324.45
205898_at	1524	CX3CR1	chemokine (C-X3-C motif	1329.3	195.08	634.97	42.747	441.76	79.278	23.436	156.29	157.04	89.718	11626	7080.3	846.43	55.954	46.831	65.088	703.38
220646_s_at	51348	KLRF1	killer cell lectin-like recep	728.3	114.8	131.7	24.566	751.36	240.92	38.515	203.31	195.7	100.14	7620.2	5981.3	75.985	74.24	100.25	97.755	204.09
205171_at	5775	PTPN4	protein tyrosine phospha	960.31	792.99	1209.5	415.43	695.24	526.92	334.2	378.6	407.28	491.97	6757.7	6224.2	130.69	179.31	485.75	147.81	541.49
236935_at				424.84	365.3	571.27	223.34	289.89	221.9	158.53	250.08	392.4	343.3	3242.9	2902.9	90.022	88.882	109.14	78.403	328.46
226625_at	7049	TGFBR3	transforming growth fact	2940.3	2267.2	2912	1216.9	588.96	522.78	231.16	741.98	630.24	599.46	14704	12273	167.22	280.79	134	140.35	1264.7
223836_at	83888	KSP37	fibroblast growth factor b	2087.8	79.084	8123.5	2358.6	435.28	230.64	175.08	88.643	54.91	54.863	37240	11588	30.563	76.403	31.688	27.257	521.22
223464_at	114879	OSBPL5	oxysterol binding protein	301	127.76	248.21	171.33	202.29	110.69	69.018	132.79	113.05	114.13	2031.1	2266.5	101.74	59.833	139.28	77.974	127.23
235643_at	219285	SAMD9L	sterile alpha motif domai	768.39	9098.1	1581.8	440.99	1694.1	2391.7	104.07	157.76	419.91	2176.5	2468.3	25077	389.99	535.76	573.04	4217.1	849.73
222816_s_at	54877	ZCCHC2	zinc finger, CCHC domain	2470.2	5750.5	1562.2	2493.9	1912.9	2185.9	1046.7	922.62	816.44	599.11	2777.4	13364	2181.8	2110.6	3070.1	2447.8	1979.3
210865_at	356	FASLG	Fas ligand (TNF superfam	272.33	789.32	739.24	912.05	263.67	181.14	81.324	227.78	206.89	174.67	1767.8	4539.7	170.68	255.05	115.16	146.5	379.82
204070_at	5920	RARRES3	retinoic acid receptor res	2719.8	2664.6	3060.1	235.33	849.11	278.5	127.62	432.41	367.02	1801.9	7967.4	15536	142.96	272	114.99	681.38	388.82
226549_at	388228	SBK1	SH3-binding domain kina	913.96	523.96	775.1	283.02	343.93	279.65	196.79	279.46	191.22	270.74	1362.5	3717.5	45.33	75.229	66.294	84.376	443.53
230036_at	219285	SAMD9L	sterile alpha motif domai	1329.8	9480.5	1860.7	712.66	1880.3	2386.9	353.87	846.09	942.06	2698.5	2070.9	15246	608.57	541.2	934.09	3897.5	1137.3
205237_at	2219	FCN1	ficolin (collagen/fibrinoge	467.99	60.392	78.933	58.575	284.26	339.72	113.46	134.87	123.98	180.89	671.24	314.21	12909	835.39	177.32	267.64	5507.7
215049_x_at	9332	CD163	CD163 molecule	101.82	119.76	79.024	61.356	118.4	74.272	44.964	221.24	226.95	210.04	263.89	77.313	9682.4	8620.1	827.99	239.35	265.16
219890_at	23601	CLEC5A	C-type lectin domain fam	119.11	95.737	92.389	84.429	228.91	150.82	73.627	226.83	219.5	159.9	1117.4	316.18	19857	11344	2299.7	896.82	347.84
203645_s_at	9332	CD163	CD163 molecule	71.858	85.886	38.251	24.845	68.466	53.077	31.555	174.52	279.05	140.63	241.85	36.597	8883.9	7051.5	908.66	212.78	175.95
223204_at	51313	C4orf18	chromosome 4 open reac	156.34	93.035	118.44	123.52	130.26	140.87	68.801	294.18	302.38	218.98	437.88	202.42	4948.7	822.28	4023.9	569.82	309.26
204150_at	23166	STAB1	stabilin 1	128.66	107.3	93.651	72.915	83.267	94.68	113.24	128.71	106.57	64	600.3	293.96	10839	1936.2	8595.5	1317.1	174.1
223567_at	10501	SEMA6B	sema domain, transmem	33.318	21.347	24.505	33.317	76.208	44.16	22.369	63.408	40.779	118.86	107.53	73.612	1094.7	910.27	52.127	143.82	30.585
201506_at	7045	TGFBI	transforming growth fact	314.89	110.94	91.258	36.355	79.571	101.04	43.168	62.25	132.31	102.34	3343	793.16	31162	3217	25407	5132.9	103.22
204392_at	8536	CAMK1	calcium/calmodulin-depe	231.77	200.59	264.45	485.72	194.55	167.43	176.85	157.92	185.87	191.37	336.18	168.74	4099.8	349.29	3162	432.58	573.02
38487_at	23166	STAB1	stabilin 1	161.92	111.78	150.14	170.8	190.98	219.39	144.43	301.11	260.52	81.386	705.38	383.12	13007	3057.7	9961.8	2083.4	329.11
213119_at	206358	SLC36A1	solute carrier family 36 (	69.352	112.18	167.49	162.58	136.7	242.86	134.87	81.805	73.633	100.04	249.87	161.79	2534.6	715.27	1603.8	653.16	330.92
215784_at	913	CD1E	CD1e molecule	23.446	31.112	24.51	19.626	10.493	18.039	17.494	38.018	79.575	21.279	40.886	10.775	97.286	121.26	16075	5709.7	69.112
202953_at	713	C1QB	complement component	57.155	68.223	38.545	24.608	93.249	62.42	33.698	50.778	36.277	18.398	77.011	102.19	182.48	146.48	10592	7856.7	41.177
218232_at	712	C1QA	complement component	62.558	24.101	23.867	20.628	32.379	49.396	11.998	83.199	44.593	74.948	50.142	86.203	184.27	108.09	7282.4	2928.5	76.151
206749_at	910	CD1B	CD1b molecule	116.44	64.685	43.037	23.273	75.038	76.53	60.891	184.23	121.28	202.85	106.28	55.634	90.609	383.87	17114	4264.6	170.68
208592_s_at	913	CD1E	CD1e molecule	76.574	73.689	34.628	34.141	101.33	67.955	48.996	74.338	72.827	86.424	115.69	80.099	81.505	94.16	6752.7	1417.6	118.17
225353_s_at	714	C1QC	complement component	110.65	140.74	27.702	61.741	34.068	71.675	33.642	118.86	56.562	125.48	139.21	130.21	495.99	88.076	12507	11043	156.99
204518_s_at	5480	PPIC	peptidylprolyl isomerase	47.172	48.238	37.09	16.31	46.384	27.873	72.802	76.562	100.23	28.991	26.824	12.981	32.336	55.873	707.73	310.34	39.174
206120_at	945	CD33	CD33 molecule	23.86	26.068	11.41	9.258	25.716	20.375	14.645	37.029	35.662	17.374	19.578	19.477	679.76	121.04	1593.4	133.38	60.619
210325_at	909	CD1A	CD1a molecule	334.06	215.3	239.04	192.86	595.37	330.27	283.03	765.57	729.57	334.33	236.08	255.17	374.32	400.57	17758	3408.6	715.59
217757_at	2	A2M	alpha-2-macroglobulin	56.154	90.243	91.964	68.09	88.213	61.413	70.048	110.28	217.55	116.79	212.65	114.82	533.47	194.51	14937	15010	110.11
206407_s_at	6357	CCL13	chemokine (C-C motif) li	103.8	102.11	70.412	68.673	165.5	111.1	53.946	134.15	164.72	100.83	157.64	289.58	242.42	455.08	6434.5	11487	174.82
213415_at	1193	CLIC2	chloride intracellular chai	19.734	13.509	4.991	7.239	27.693	27.378	16.056	17.633	17.894	9.958	115.54	209.36	190.72	234.49	3590.7	8375.9	22.283
203680_at	5577	PRKAR2B	protein kinase, cAMP-dep	109.78	78.126	61.402	92.489	117.09	64.795	78.53	96.114	111.03	96.123	108.68	84.149	166.4	144.73	669.88	3171.9	167.26
219519_s_at	6614	SIGLEC1	sialic acid binding Ig-like	23.821	16.503	13.074	11.874	18.704	14.555	17.02	27.273	27.555	13.05	29.371	89.235	275.86	294.87	710.24	6005.1	49.332
220049_s_at	80380	PDCD1LG2	programmed cell death 1	16.213	14.334	24.966	7.891	14.958	12.758	13.968	20.981	22.152	13.788	33.651	23.536	50.538	60.569	174.05	1520.5	26.35
205686_s_at	942	CD86	CD86 molecule	183.62	162.09	106.74	90.275	293.85	267.82	183.33	618.49	400.78	281.29	137.07	318.48	630.29	242.14	2070.3	8560	203.25
202357_s_at	629	CFB	complement factor B	136.58	132.94	46.834	56.07	131.35	83.048	43.087	129.28	79.264	59.938	110.84	127.86	139.77	1128.5	252.42	4969.9	187.81
204614_at	5055	SERPINB2	serpin peptidase inhibitor	104.82	108.06	56.517	32.963	71.147	96.217	41.219	113.71	115.17	71.027	83.1	95.581	4287.5	24461	98.613	352.89	181.87
219093_at	55022	FLJ20701	phosphotyrosine interact	68.932	25.883	27.311	36.14	27.062	43.915	33.14	85.735	108.6	45.643	154.69	54.538	7487.2	11205	34.221	105.68	78.269
207850_at	2921	CXCL3	chemokine (C-X-C motif)	24.788	25.074	21.049	26.677	53.966	18.063	38.137	56.646	50.331	87.434	63.616	22.828	3914.7	27356	131.59	522.63	73.882
215101_s_at	6374	CXCL5	chemokine (C-X-C motif)	32.322	5.255	14.219	3.561	33.529	12.421	21.868	49.028	37.239	15.912	20.307	12.939	4063.1	33167	20.269	445.88	102.86
204470_at	2919	CXCL1	chemokine (C-X-C motif)	102.29	72.235	49.323	38.146	109.13	94.808	78.084	156.84	130.59	91.55	86.887	118.64	6214.2	35585	100.46	1191.3	1368.5
210118_s_at	3552	IL1A	interleukin 1, alpha	88.623	145.24	90.422	35.318	155.24	63.8	81.281	287.8	220.03	111.54	112.61	77.232	2599.8	22595	213.69	660.82	323.06
217767_at	718	C3	complement component	59.883	26.757	34.401	17.188	28.973	26.25	17.639	27.042	25.763	39.066	62.908	45.944	5647.8	8096.8	246.51	282.92	49.628
221463_at	6369	CCL24	chemokine (C-C motif) li	44.054	21.628	24.646	14.316	57.382	49.296	26.492	25.721	26.113	19.318	235.72	198.21	9587.6	13839	113.99	365.61	68.75
210511_s_at	3624	INHBA	inhibin, beta A	16.237	135.48	7.304	7.813	26.306	14.216	20.578	20.333	18.133	8.488	20.287	104.69	536.09	11449	171.22	4927.5	19.921
205922_at	8875	VNN2	vanin 2	269.9	77.17	513.82	12.6	355.24	189.21	167.27	287.16	314.52	168.83	499.61	320.66	223.25	4735.8	44.71	72.248	19069
203591_s_at	1441	CSF3R	colony stimulating factor	106.53	44.825	45.942	38.679	176.05	82.909	41.943	74.399	38.012	29.283	24.289	44.759	819.72	100.75	190.43	109.83	10132
223809_at	64407	RGS18	regulator of G-protein sig	348.74	282.09	132.72	112.99	257.38	151.54	93.752	264.46	229.04	186.15	315.04	200.03	374.78	94.765	1421.8	103.53	10444