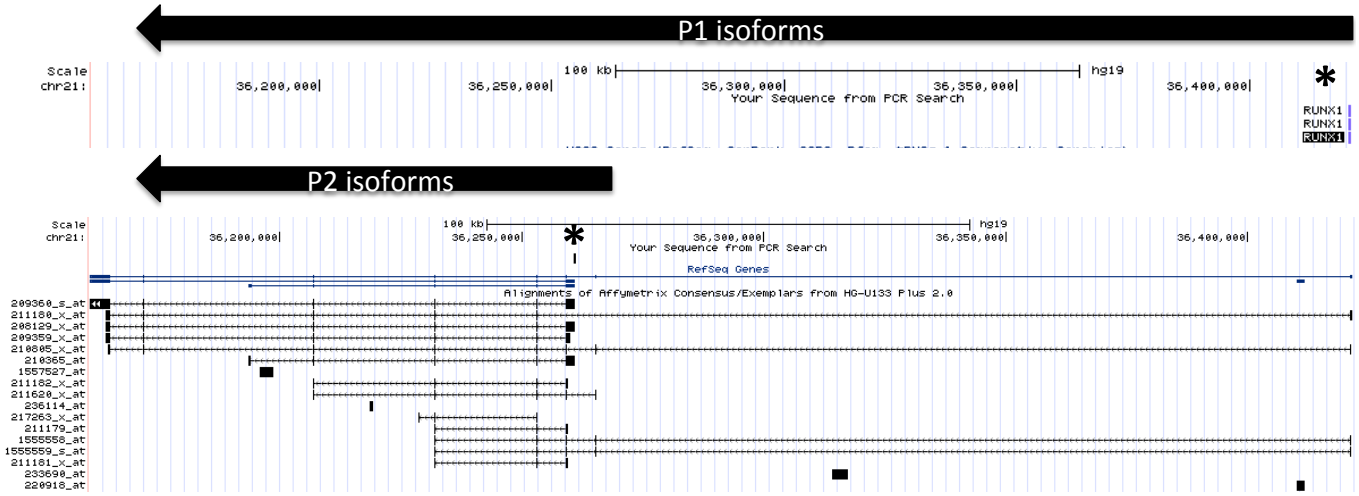

Figure S1 *RUNX1* gene with proximal (P1) and distal (P2) promoters, major isoforms, Affymetrix U133 plus 2.0 probe sets, and PCR primers (*) used to quantify P1 and P2 expression.

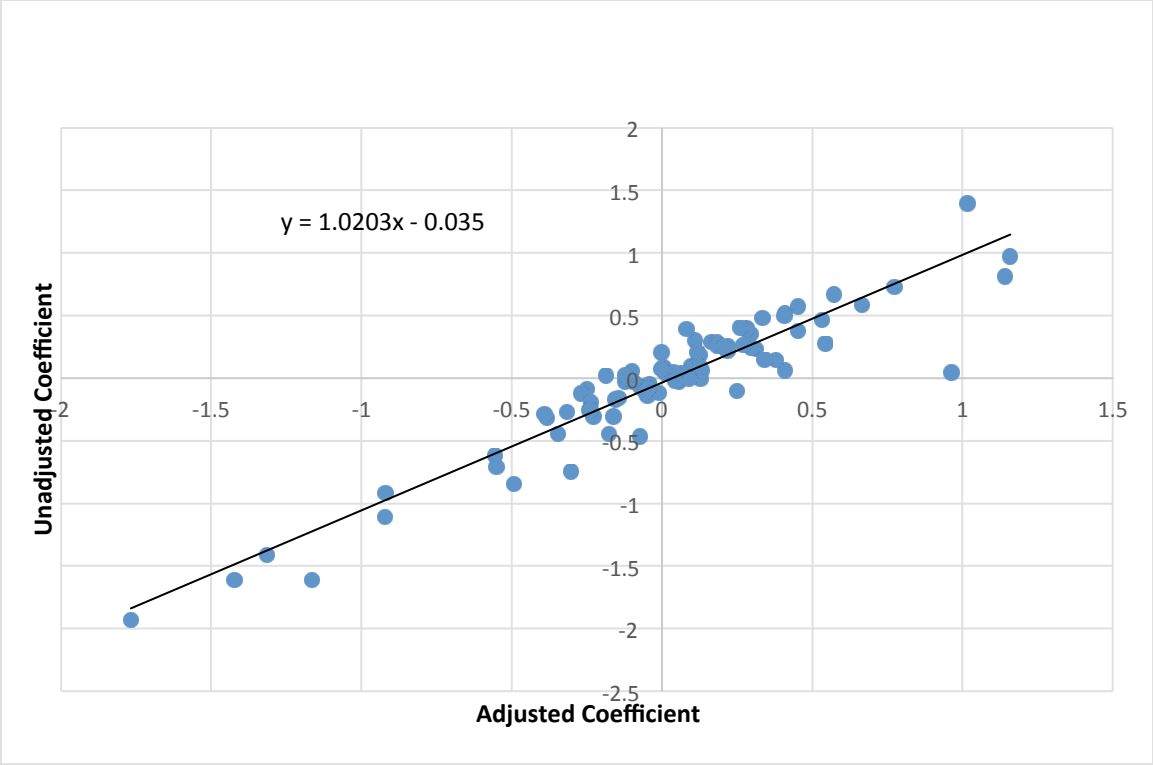

Figure S2 Comparison of coefficients for association of gene expression with cancer free survival in Jorissen et al dataset(Jorissen et al., 2009a) with (x-axis) vs. without (y-axis)adjustment for age, gender, adjuvant chemo- or radiation therapy. Each point represents a probe set mapping to either RUNX1 or an Aspirin Response Signature gene. The fitted line is the best linear model describing the points. A slope of 1 and intercept of 0 indicate that the unadjusted and adjusted coefficients are similar for most genes.

Supplemental Table 1 – Summary of *RUNX1* microarray probe set types and correlation with *RUNX1* P1 and P2 expression by PCR[1]					
Probe Set	Affymetrix probe set type[2]	*RUNX1* P2		*RUNX1* P1	
		Effect	P-value	Effect	P-value
233690_at	Unique	0.17	0.11	0.51	0.00
220918_at	Unique	0.03	0.48	0.28	0.00
209360_s_at	Identical	0.14	0.00	0.06	0.05
211620_x_at	Mixed	0.00	1.00	-0.17	0.10
209359_x_at	Mixed	0.06	0.47	-0.15	0.20
211180_x_at	Mixed	0.12	0.32	-0.15	0.35
208129_x_at	Mixed	0.06	1.00	-0.09	0.62
1557527_at	Unique	0.18	0.01	0.06	1.00
217263_x_at	Mixed	0.12	0.07	0.00	1.00
210365_at	Unique	0.20	0.07	-0.02	1.00
211181_x_at	Mixed	0.14	0.08	-0.07	1.00
211182_x_at	Mixed	0.10	0.29	-0.07	1.00
236114_at	Unique	0.09	0.54	0.01	1.00
210805_x_at	Mixed	0.00	1.00	-0.10	1.00
1555558_at	Unique	0.02	1.00	0.00	1.00
211179_at	Unique	0.04	1.00	0.01	1.00
1555559_s_at	Identical	-0.01	1.00	-0.03	1.00

[1] Shaded probe sets represent those that meet two criteria: 1) map uniquely to a single sequence and 2) are significantly ($p < 0.1$) associated with either *RUNX1* P1 or P2 expression by PCR.
[2]Unique probe sets contain probes that do not cross hybridize to other sequences; Identical probe sets contains probes that all cross-hybridize to sequences in different genes; Mixed probe sets contain at least one probe that cross-hybridizes to more than one sequence.

Additional Datasets

Gene expression data for two publicly available colon cancer datasets(Jorissen et al., 2009b, Smith et al., 2010) were downloaded (see Methods) and examined for differences by stage for in Aspirin Response Signature(Voora et al., 2013) probe set within each dataset (Smith and Jorrisen ANOVA sheets). Using meta-analysis we combined both datasets and performed pairwise tests between stage B-D vs. stage A for each ARS probe set (Meta BvA, CvA, and DvA sheets).

Gene expression data for two publicly available colon cancer datasets were downloaded (see Methods) and examined for in colon free survival for Aspirin Response Signature(Voora et al., 2013) probe sets within each dataset (Smith and Jorrisen Survival sheets sheets). Using meta-analysis we combined both datasets (Meta Survival sheet).

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Smith ANOVA

probe set	GeneName	key.stage2	key.stage3	key.stage4	AveExpr	P.Value
226152_at	TTC7B	0.2923	0.4904	0.3385	7.1284	0.0015
201108_s_at	THBS1	0.3325	0.6837	0.353	9.804	0.0023
200665_s_at	SPARC	0.328	0.5019	0.2945	12.253	0.0029
212667_at	SPARC	0.3153	0.634	0.3808	10.877	0.0033
209651_at	TGFB1I1	0.2254	0.5343	0.3224	9.2159	0.0045
201058_s_at	MYL9	0.5333	0.7625	0.5152	9.8285	0.0062
1552773_at	CLEC4D	-0.018	-0.0485	-0.111	6.2424	0.008
212077_at	CALD1	0.1973	0.4362	0.3294	11.8559	0.0081
207414_s_at	PCSK6	-0.4293	-0.4399	-0.388	8.3864	0.0085
207156_at	HIST1H2AG	-0.2705	-0.0356	0.0807	5.4799	0.0115
204115_at	GNG11	0.1317	0.4188	0.1985	9.105	0.0128
212651_at	RHOBTB1	0.2225	0.3367	0.4575	8.3694	0.0161
216956_s_at	ITGA2B	-0.037	-0.0867	-0.1244	6.1998	0.0182
212813_at	JAM3	0.1495	0.3496	0.2439	8.8929	0.0209
208782_at	FSTL1	0.1227	0.3798	0.2061	10.7708	0.023
203817_at	GUCY1B3	0.2275	0.3665	0.2098	8.8591	0.0363
207808_s_at	PROS1	0.3209	0.5156	0.3246	9.0799	0.0381
224823_at	MYLK	0.3067	0.5018	0.478	10.8513	0.0416
201906_s_at	CTDSPL	0.0039	0.1421	-0.051	9.6744	0.0501
203680_at	PRKAR2B	-0.282	-0.2564	-0.6381	7.769	0.0552
227189_at	CPNE5	-0.1903	-0.168	-0.1652	7.0632	0.0631
227180_at	ELOVL7	0.3198	-0.0855	0.1455	8.8964	0.0734
206272_at	RAB4A	0.0904	0.1882	0.2676	8.8557	0.0768
225974_at	TMEM64	0.3555	0.5397	0.3158	9.1598	0.0785
57588_at	SLC24A3	0.1541	0.3498	0.2766	8.2488	0.0885
1555659_a_at	TREML1	-0.0068	-0.085	-0.0087	7.0007	0.094
219090_at	SLC24A3	0.1664	0.3596	0.2526	8.3653	0.0982
206655_s_at	GP1BB	-0.0125	-0.0267	-0.1395	7.0206	0.1118
37966_at	PARVB	0.3274	0.3677	0.1955	7.6395	0.1128
221556_at	CDC14B	-0.063	0.0192	0.1599	8.4855	0.1258
1553842_at	BEND2	0.0801	0.0985	0.0385	4.9597	0.132
206110_at	HIST1H3H	-0.2473	-0.2253	-0.0944	6.8185	0.1332
212573_at	ENDOD1	-0.0356	0.1342	-0.0226	9.027	0.1795
1560262_at	LRRC32	-0.0183	-0.0724	-0.0643	5.3142	0.1815
225354_s_at	SH3BGRL2	-0.1289	-0.0845	0.0623	10.1452	0.1867
206494_s_at	ITGA2B	-0.0435	-0.059	-0.0538	4.6583	0.1975
230690_at	TUBB1	0.016	-0.0398	0.0013	5.8163	0.2301
214974_x_at	CXCL5	0.6265	0.9969	0.7996	8.219	0.2343
203414_at	MMD	-0.0038	0.132	0.0764	9.7883	0.2633
206493_at	ITGA2B	-0.0032	-0.0636	-0.1066	7.5222	0.2786
206049_at	SELP	0.1399	0.0698	0.0527	7.6833	0.2954

Smith ANOVA

probe set	GeneName	key.stage2	key.stage3	key.stage4	AveExpr	P.Value
207815_at	PF4V1	0.0287	0.0595	-0.0289	4.0104	0.3048
226188_at	LGALSL	-0.079	-0.0367	0.0932	8.9842	0.3104
204628_s_at	ITGB3	0.0804	0.0535	0.073	7.4427	0.3313
228708_at	RAB27B	0.1182	-0.0354	-0.2794	9.0769	0.3514
208792_s_at	CLU	0.1836	0.3118	0.1689	8.5708	0.3528
222717_at	SDPR	0.1019	0.2236	0.2369	6.7815	0.3742
210387_at	HIST1H2BG	-0.3308	-0.1725	-0.1614	6.6259	0.3744
208601_s_at	TUBB1	0.0115	-0.0248	-0.0155	6.4864	0.39
205442_at	MFAP3L	-0.3517	-0.1781	-0.2555	6.8246	0.3902
212148_at	PBX1	-0.0403	0.1072	0.212	8.6328	0.392
220496_at	CLEC1B	-0.0045	-0.0418	-0.0312	7.4321	0.4158
207206_s_at	ALOX12	0.0022	-0.0398	-0.0119	6.8405	0.429
211026_s_at	MGLL	-0.0555	0.0689	0.1039	9.6804	0.4404
208791_at	CLU	0.1586	0.3047	0.1904	8.5151	0.4425
215779_s_at	HIST1H2BG	-0.1825	-0.1588	-0.0928	6.2462	0.4562
227451_s_at	C6ORF79	0.0645	0.0732	0.1737	8.8176	0.6051
203819_s_at	IGF2BP3	-0.067	0.2506	0.0809	6.3987	0.6466
212151_at	PBX1	0.0166	0.0765	0.1318	8.8926	0.6484
241133_at	PRSS1	-0.0293	-0.0443	-0.0384	5.2852	0.649
210986_s_at	TPM1	0.0733	0.0193	0.0946	12.1825	0.6579
227088_at	PDE5A	-0.0963	0.0685	-0.0401	8.1783	0.6617
230645_at	FRMD3	-0.0724	-0.0493	-0.1858	6.8784	0.6966
230942_at	CMTM5	-0.007	-0.0284	0.0122	7.282	0.7917
207550_at	MPL	0.0086	0.0226	0.0364	5.014	0.8011
206167_s_at	ARHGAP6	0.106	0.0787	-0.0643	7.0807	0.8495
214146_s_at	PPBP	0.3439	0.404	0.2773	7.3663	0.8594
201059_at	CTTN	0.0324	0.0627	0.035	10.0915	0.879
225166_at	ARHGAP18	-0.0786	-0.0453	-0.0853	6.8501	0.9219
235331_x_at	PCGF5	0.1088	0.0729	0.0383	7.0513	0.9229
229778_at	C12ORF39	0.0047	0.0062	0.0168	4.0643	0.9462
206390_x_at	PF4	0.001	-0.021	0.0139	7.8762	0.9885

Jorissen ANOVA

probe set	GeneName	key.stage2	key.stage3	key.stage4	AveExpr	P.Value
226152_at	TTC7B	0.3014	0.4639	0.3898	6.9515	0.0011
201058_s_at	MYL9	0.7127	0.7196	0.6899	9.2586	0.0013
227189_at	CPNE5	-0.2217	-0.1364	-0.2455	6.4551	0.0016
212667_at	SPARC	0.397	0.6172	0.4726	10.3232	0.0021
200665_s_at	SPARC	0.375	0.5161	0.3675	11.8067	0.0029
212077_at	CALD1	0.3262	0.4586	0.3739	11.5609	0.0033
201108_s_at	THBS1	0.3029	0.503	0.4438	9.2409	0.0058
207414_s_at	PCSK6	-0.3267	-0.344	-0.3072	7.9278	0.0078
203680_at	PRKAR2B	-0.2341	-0.348	-0.6581	7.5505	0.0092
209651_at	TGFB1I1	0.3577	0.4776	0.4332	8.6405	0.0098
220496_at	CLEC1B	0.0179	-0.0533	-0.0417	6.8405	0.0131
224823_at	MYLK	0.4785	0.5409	0.5361	10.3868	0.0169
212651_at	RHOBTB1	0.0973	0.2039	0.3708	7.5845	0.0226
219090_at	SLC24A3	0.2876	0.3322	0.4151	7.7354	0.0287
212813_at	JAM3	0.2007	0.2737	0.3131	8.6541	0.0307
203817_at	GUCY1B3	0.3059	0.3557	0.3262	8.3873	0.0345
204115_at	GNG11	0.2235	0.3861	0.2623	8.6923	0.0373
57588_at	SLC24A3	0.2025	0.2914	0.3776	7.6493	0.0386
228708_at	RAB27B	0.1345	0.0151	-0.3644	8.682	0.0477
1560262_at	LRRC32	-0.0646	-0.0912	-0.0623	4.8297	0.0572
208782_at	FSTL1	0.1624	0.3311	0.2514	10.3262	0.0603
212573_at	ENDOD1	0.0416	0.2132	0.0719	8.7469	0.0636
204628_s_at	ITGB3	0.1026	0.0268	0.0182	7.0451	0.0846
227180_at	ELOVL7	0.1566	-0.1437	-0.1045	8.4709	0.0899
214974_x_at	CXCL5	0.6081	0.9484	0.5997	7.9401	0.1015
212148_at	PBX1	0.0519	0.1092	0.3272	8.2904	0.1074
207815_at	PF4V1	0.0353	0.0515	-0.0308	3.6583	0.1136
1555659_a_at	TREML1	-0.0391	-0.0762	0.0013	6.6807	0.1244
207206_s_at	ALOX12	0.0432	0.005	-0.0302	6.301	0.1287

Jorissen ANOVA

probe set	GeneName	key.stage2	key.stage3	key.stage4	AveExpr	P.Value
225354_s_at	SH3BGRL2	-0.1191	-0.025	0.0638	9.7399	0.1336
1553842_at	BEND2	0.017	0.0479	-0.0199	4.64	0.1476
221556_at	CDC14B	-0.1091	-0.0057	0.0875	8.223	0.1577
1552773_at	CLEC4D	0.0513	0.0131	-0.0155	5.8716	0.1587
203414_at	MMD	0.0232	0.1177	0.13	9.5199	0.1792
222717_at	SDPR	0.2511	0.224	0.3861	6.3826	0.1897
207808_s_at	PROS1	0.0417	0.2183	0.2472	8.897	0.2113
212151_at	PBX1	0.0775	0.0402	0.2151	8.8109	0.2151
203819_s_at	IGF2BP3	0.3273	0.5313	0.3617	5.8796	0.2242
206049_at	SELP	0.0942	0.0052	-0.0065	7.3906	0.2422
235331_x_at	PCGF5	-0.1967	-0.1769	-0.2866	6.7696	0.2466
206167_s_at	ARHGAP6	0.3597	0.3128	0.3074	6.672	0.2901
208792_s_at	CLU	0.2599	0.3394	0.2847	8.3553	0.3048
226188_at	LGALSL	-0.0857	-0.0077	0.0614	8.7322	0.3089
210986_s_at	TPM1	0.0643	-0.0202	0.0894	11.9018	0.3536
225974_at	TMEM64	0.2306	0.2979	0.1912	8.7928	0.3628
225166_at	ARHGAP18	-0.097	-0.1978	-0.119	6.6369	0.3642
208791_at	CLU	0.2305	0.3174	0.2878	8.3694	0.366
230690_at	TUBB1	0.0193	-0.0202	0.015	5.4767	0.3663
210387_at	HIST1H2BG	0.0561	0.2282	0.206	6.363	0.3688
241133_at	PRSS1	-0.0368	-0.0625	-0.0584	5.1363	0.3967
207156_at	HIST1H2AG	-0.0735	0.0254	0.1151	5.349	0.4051
37966_at	PARVB	0.1461	0.1725	0.0548	7.3797	0.4273
206655_s_at	GP1BB	0.0106	0.0288	-0.0686	6.5017	0.4646
205442_at	MFAP3L	-0.1581	-0.1814	0.0208	6.5332	0.4706
206494_s_at	ITGA2B	-0.023	-0.0313	-0.0488	4.3658	0.4759
216956_s_at	ITGA2B	-0.0307	-0.0502	-0.0635	5.7151	0.48
206272_at	RAB4A	-0.0327	0.0201	0.0801	8.4287	0.5151
230942_at	CMTM5	-0.0521	-0.0609	-0.0287	6.8353	0.5293

Jorissen ANOVA

probe set	GeneName	key.stage2	key.stage3	key.stage4	AveExpr	P.Value
207550_at	MPL	-0.0063	-0.0278	0.0233	4.7068	0.5339
214146_s_at	PPBP	0.4085	0.5172	0.4298	6.4749	0.5765
208601_s_at	TUBB1	-0.027	-0.0364	-0.0409	5.6026	0.6164
229778_at	C12ORF39	0.0071	-0.0182	0.0027	3.7252	0.6626
211026_s_at	MGLL	0.031	0.1102	0.0946	9.4465	0.6747
215779_s_at	HIST1H2BG	0.0445	0.1159	0.1094	5.9326	0.6874
201906_s_at	CTDSPL	-0.0644	-0.0156	-0.068	9.312	0.709
230645_at	FRMD3	0.079	0.0603	-0.0638	6.8273	0.7653
206390_x_at	PF4	-0.1064	-0.0703	-0.0567	7.4299	0.7913
227451_s_at	C6ORF79	-0.0321	-0.0391	0.0435	8.4604	0.7986
201059_at	CTTN	-0.0273	0.0139	-0.0399	9.8714	0.8205
206110_at	HIST1H3H	0.0864	0.0939	0.0467	6.3834	0.8246
206493_at	ITGA2B	0.0128	-0.0245	-0.0319	6.9668	0.8433
227088_at	PDE5A	0.0689	0.0028	0.0498	7.8206	0.9334

Meta BvA

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
201058_s_at	MYL9	0.6302	0.1457	4.3263	0
207414_s_at	PCSK6	-0.3652	0.084	-4.3468	0
227189_at	CPNE5	-0.2078	0.0485	-4.2831	0
200665_s_at	SPARC	0.351	0.0956	3.6708	0.0002
226152_at	TTC7B	0.2972	0.0853	3.4855	0.0005
212667_at	SPARC	0.3599	0.1186	3.0356	0.0024
224823_at	MYLK	0.3958	0.1306	3.0295	0.0024
203817_at	GUCY1B3	0.2668	0.0896	2.978	0.0029
212077_at	CALD1	0.2677	0.0917	2.9182	0.0035
209651_at	TGFB1I1	0.2968	0.1068	2.7783	0.0055
204628_s_at	ITGB3	0.0907	0.0333	2.7233	0.0065
201108_s_at	THBS1	0.3143	0.1159	2.7125	0.0067
219090_at	SLC24A3	0.2353	0.1001	2.3498	0.0188
212813_at	JAM3	0.1787	0.0792	2.2569	0.024
206049_at	SELP	0.1132	0.0513	2.2068	0.0273
37966_at	PARVB	0.2096	0.0967	2.1668	0.0302
225974_at	TMEM64	0.2798	0.1312	2.1336	0.0329
214974_x_at	CXCL5	0.615	0.3006	2.046	0.0408
57588_at	SLC24A3	0.1824	0.0969	1.8823	0.0598
204115_at	GNG11	0.1814	0.0979	1.8528	0.0639
207156_at	HIST1H2AG	-0.1658	0.091	-1.8227	0.0684
1560262_at	LRRC32	-0.047	0.026	-1.8049	0.0711
225354_s_at	SH3BGRL2	-0.123	0.069	-1.7824	0.0747
203680_at	PRKAR2B	-0.2534	0.1448	-1.75	0.0801
205442_at	MFAP3L	-0.2348	0.1342	-1.7498	0.0802
208792_s_at	CLU	0.2209	0.1268	1.7417	0.0816
227180_at	ELOVL7	0.2162	0.1248	1.7326	0.0832
206167_s_at	ARHGAP6	0.2616	0.1516	1.7261	0.0843
206494_s_at	ITGA2B	-0.0336	0.0204	-1.6513	0.0987

Meta BvA

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
212651_at	RHOBTB1	0.1514	0.0923	1.6397	0.1011
208782_at	FSTL1	0.1443	0.0937	1.5391	0.1238
222717_at	SDPR	0.1725	0.1131	1.5247	0.1273
208791_at	CLU	0.1961	0.1333	1.4714	0.1412
1553842_at	BEND2	0.0387	0.0273	1.4168	0.1565
207808_s_at	PROS1	0.1569	0.1147	1.3688	0.1711
214146_s_at	PPBP	0.384	0.2954	1.2999	0.1936
241133_at	PRSS1	-0.0327	0.0263	-1.2419	0.2143
221556_at	CDC14B	-0.0888	0.073	-1.2155	0.2242
226188_at	LGALSL	-0.0831	0.0688	-1.2071	0.2274
210986_s_at	TPM1	0.0681	0.0592	1.1499	0.2502
216956_s_at	ITGA2B	-0.0337	0.0293	-1.1492	0.2505
225166_at	ARHGAP18	-0.0892	0.0866	-1.0305	0.3028
207815_at	PF4V1	0.0331	0.0327	1.0111	0.312
230942_at	CMTM5	-0.0316	0.0329	-0.9623	0.3359
207206_s_at	ALOX12	0.0228	0.0249	0.9189	0.3581
203819_s_at	IGF2BP3	0.1815	0.2017	0.8996	0.3683
1555659_a_at	TREML1	-0.0252	0.031	-0.8131	0.4162
235331_x_at	PCGF5	-0.0851	0.1049	-0.8112	0.4173
228708_at	RAB27B	0.1284	0.1587	0.8088	0.4186
210387_at	HIST1H2BG	-0.1002	0.1255	-0.7986	0.4245
230690_at	TUBB1	0.018	0.023	0.7841	0.433
206110_at	HIST1H3H	-0.0615	0.082	-0.7501	0.4532
206390_x_at	PF4	-0.0595	0.0799	-0.7452	0.4561
215779_s_at	HIST1H2BG	-0.0588	0.0832	-0.7067	0.4797
1552773_at	CLEC4D	0.0175	0.025	0.7013	0.4831
212151_at	PBX1	0.051	0.0806	0.6325	0.5271
201906_s_at	CTDSPL	-0.0359	0.0568	-0.6315	0.5277
220496_at	CLEC1B	0.0083	0.0222	0.3713	0.7104

Meta BvA

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
229778_at	C12ORF39	0.006	0.0196	0.3087	0.7576
208601_s_at	TUBB1	-0.0063	0.021	-0.2993	0.7647
203414_at	MMD	0.0129	0.0597	0.2164	0.8287
206272_at	RAB4A	0.0136	0.0665	0.2046	0.8379
212148_at	PBX1	0.0171	0.1123	0.1522	0.879
212573_at	ENDOD1	0.0085	0.0718	0.1188	0.9055
206493_at	ITGA2B	0.0055	0.0474	0.1158	0.9078
211026_s_at	MGLL	-0.0052	0.0825	-0.0628	0.9499
227451_s_at	C6ORF79	0.004	0.0788	0.0514	0.959
230645_at	FRMD3	0.0048	0.115	0.0415	0.9669
207550_at	MPL	0.0008	0.0279	0.0301	0.976
206655_s_at	GP1BB	-0.0012	0.0489	-0.0237	0.9811
201059_at	CTTN	-0.0011	0.0529	-0.0206	0.9836
227088_at	PDE5A	0.0013	0.1112	0.0121	0.9904

Meta CvA

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
200665_s_at	SPARC	0.5088	0.0959	5.3072	0
201058_s_at	MYL9	0.7394	0.1461	5.0619	0
201108_s_at	THBS1	0.5727	0.1162	4.9274	0
204115_at	GNG11	0.4012	0.0982	4.0847	0
207414_s_at	PCSK6	-0.3802	0.0843	-4.5112	0
209651_at	TGFB1I1	0.5039	0.1071	4.704	0
212077_at	CALD1	0.4484	0.092	4.8743	0
212667_at	SPARC	0.6249	0.1189	5.2549	0
226152_at	TTC7B	0.4764	0.0855	5.5717	0
203817_at	GUCY1B3	0.3611	0.0898	4.0194	0.0001
212813_at	JAM3	0.3066	0.0794	3.8617	0.0001
224823_at	MYLK	0.522	0.131	3.9849	0.0001
208782_at	FSTL1	0.3534	0.094	3.7594	0.0002
219090_at	SLC24A3	0.3441	0.1004	3.4258	0.0006
57588_at	SLC24A3	0.3158	0.0972	3.2498	0.0012
1560262_at	LRRC32	-0.084	0.0261	-3.2157	0.0013
214974_x_at	CXCL5	0.9665	0.3016	3.2049	0.0014
227189_at	CPNE5	-0.1505	0.0487	-3.092	0.002
225974_at	TMEM64	0.3938	0.1316	2.9932	0.0028
207808_s_at	PROS1	0.3417	0.115	2.9714	0.003
212651_at	RHOBTB1	0.2616	0.0926	2.8255	0.0047
1555659_a_at	TREML1	-0.08	0.0311	-2.5735	0.0101
208792_s_at	CLU	0.3252	0.1272	2.5574	0.0105
212573_at	ENDOD1	0.1792	0.072	2.4891	0.0128
37966_at	PARVB	0.2414	0.0971	2.4862	0.0129
1553842_at	BEND2	0.0654	0.0274	2.3865	0.017
208791_at	CLU	0.3113	0.1337	2.3287	0.0199
216956_s_at	ITGA2B	-0.0677	0.0294	-2.3018	0.0213
206494_s_at	ITGA2B	-0.0457	0.0204	-2.2389	0.0252

Meta CvA

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
220496_at	CLEC1B	-0.0483	0.0223	-2.1673	0.0302
203680_at	PRKAR2B	-0.3109	0.1452	-2.1409	0.0323
203819_s_at	IGF2BP3	0.4268	0.2024	2.1086	0.035
203414_at	MMD	0.1232	0.0599	2.0567	0.0397
241133_at	PRSS1	-0.0526	0.0264	-1.9919	0.0464
222717_at	SDPR	0.2238	0.1134	1.9738	0.0484
207815_at	PF4V1	0.0542	0.0328	1.6502	0.0989
214146_s_at	PPBP	0.4741	0.2964	1.5994	0.1097
225166_at	ARHGAP18	-0.1329	0.0868	-1.5302	0.126
206167_s_at	ARHGAP6	0.2218	0.152	1.4585	0.1447
208601_s_at	TUBB1	-0.0301	0.0211	-1.4294	0.1529
230942_at	CMTM5	-0.0461	0.0329	-1.398	0.1621
205442_at	MFAP3L	-0.1801	0.1346	-1.3378	0.181
206272_at	RAB4A	0.0838	0.0667	1.2556	0.2093
204628_s_at	ITGB3	0.0411	0.0334	1.2307	0.2184
230690_at	TUBB1	-0.0281	0.023	-1.2192	0.2228
211026_s_at	MGLL	0.0928	0.0828	1.1213	0.2622
227180_at	ELOVL7	-0.1224	0.1252	-0.9774	0.3284
212148_at	PBX1	0.1085	0.1126	0.9629	0.3356
206493_at	ITGA2B	-0.0425	0.0475	-0.8948	0.3709
201906_s_at	CTDSPL	0.0506	0.057	0.8878	0.3746
235331_x_at	PCGF5	-0.085	0.1053	-0.808	0.4191
225354_s_at	SH3BGRL2	-0.0488	0.0692	-0.7048	0.481
207206_s_at	ALOX12	-0.0173	0.0249	-0.6957	0.4866
212151_at	PBX1	0.0561	0.0808	0.6941	0.4876
1552773_at	CLEC4D	-0.0171	0.0251	-0.6813	0.4957
201059_at	CTTN	0.0354	0.0531	0.6681	0.5041
206049_at	SELP	0.0323	0.0515	0.6273	0.5305
206390_x_at	PF4	-0.0487	0.0801	-0.6077	0.5434

Meta CvA

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
206110_at	HIST1H3H	-0.0484	0.0822	-0.589	0.5558
210387_at	HIST1H2BG	0.0653	0.1259	0.5188	0.6039
229778_at	C12ORF39	-0.0071	0.0197	-0.3622	0.7172
226188_at	LGALSL	-0.0192	0.069	-0.2776	0.7813
227088_at	PDE5A	0.0298	0.1116	0.2671	0.7894
207550_at	MPL	-0.0036	0.028	-0.128	0.8981
215779_s_at	HIST1H2BG	-0.0098	0.0835	-0.1177	0.9063
221556_at	CDC14B	0.0053	0.0732	0.0728	0.9419
210986_s_at	TPM1	-0.0034	0.0594	-0.0567	0.9548
230645_at	FRMD3	0.0063	0.1153	0.0548	0.9563
227451_s_at	C6ORF79	0.0032	0.079	0.0403	0.9679
207156_at	HIST1H2AG	-0.0034	0.0912	-0.0372	0.9703
228708_at	RAB27B	-0.0039	0.1592	-0.0245	0.9805
206655_s_at	GP1BB	0.0004	0.049	0.0085	0.9932

Meta DvA

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
203680_at	PRKAR2B	-0.6499	0.1558	-4.1718	0
212651_at	RHOBTB1	0.4089	0.0993	4.1185	0
201058_s_at	MYL9	0.6082	0.1566	3.8835	0.0001
226152_at	TTC7B	0.3655	0.0917	3.9878	0.0001
227189_at	CPNE5	-0.2094	0.0522	-4.0127	0.0001
207414_s_at	PCSK6	-0.3381	0.0904	-3.7384	0.0002
212077_at	CALD1	0.3534	0.0986	3.5831	0.0003
224823_at	MYLK	0.5077	0.1404	3.616	0.0003
212667_at	SPARC	0.4303	0.1275	3.3752	0.0007
209651_at	TGFB1I1	0.3813	0.1148	3.3208	0.0009
212813_at	JAM3	0.2827	0.0851	3.3205	0.0009
201108_s_at	THBS1	0.4083	0.1247	3.274	0.0011
200665_s_at	SPARC	0.3298	0.1027	3.21	0.0013
57588_at	SLC24A3	0.3349	0.1042	3.2126	0.0013
219090_at	SLC24A3	0.3437	0.1077	3.1918	0.0014
216956_s_at	ITGA2B	-0.093	0.0315	-2.9512	0.0032
203817_at	GUCY1B3	0.2673	0.0963	2.7761	0.0055
222717_at	SDPR	0.3063	0.1215	2.522	0.0117
206494_s_at	ITGA2B	-0.0514	0.0219	-2.3496	0.0188
212148_at	PBX1	0.2828	0.1209	2.34	0.0193
1552773_at	CLEC4D	-0.0628	0.0268	-2.3372	0.0194
208782_at	FSTL1	0.2304	0.1008	2.2856	0.0223
207808_s_at	PROS1	0.2797	0.1234	2.2679	0.0233
1560262_at	LRRC32	-0.0631	0.028	-2.2505	0.0244
204115_at	GNG11	0.2326	0.1053	2.2091	0.0272
206272_at	RAB4A	0.152	0.0716	2.1229	0.0338
214974_x_at	CXCL5	0.6752	0.3236	2.0867	0.0369
212151_at	PBX1	0.1782	0.0867	2.0558	0.0398
206655_s_at	GP1BB	-0.1053	0.0525	-2.0031	0.0452

Meta DvA

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
228708_at	RAB27B	-0.332	0.1709	-1.9429	0.052
225974_at	TMEM64	0.2412	0.1411	1.709	0.0874
203414_at	MMD	0.1091	0.0643	1.698	0.0895
208791_at	CLU	0.2405	0.1433	1.6789	0.0932
241133_at	PRSS1	-0.0474	0.0283	-1.6765	0.0936
208792_s_at	CLU	0.2247	0.1363	1.6489	0.0992
220496_at	CLEC1B	-0.0371	0.0239	-1.5526	0.1205
221556_at	CDC14B	0.1201	0.0785	1.5291	0.1262
235331_x_at	PCGF5	-0.1656	0.1129	-1.4659	0.1427
210986_s_at	TPM1	0.0916	0.0637	1.4382	0.1504
204628_s_at	ITGB3	0.0479	0.0358	1.3395	0.1804
206493_at	ITGA2B	-0.0668	0.0509	-1.3106	0.19
208601_s_at	TUBB1	-0.027	0.0226	-1.1993	0.2304
203819_s_at	IGF2BP3	0.2558	0.2172	1.178	0.2388
214146_s_at	PPBP	0.3709	0.318	1.1662	0.2435
225166_at	ARHGAP18	-0.1044	0.0931	-1.1215	0.2621
211026_s_at	MGLL	0.0985	0.0888	1.1095	0.2672
227451_s_at	C6ORF79	0.0931	0.0848	1.0982	0.2721
207156_at	HIST1H2AG	0.0987	0.0978	1.0095	0.3127
37966_at	PARVB	0.1051	0.1042	1.0088	0.3131
230645_at	FRMD3	-0.1245	0.1235	-1.008	0.3135
226188_at	LGALSL	0.0741	0.0741	1.0009	0.3169
201906_s_at	CTDSPL	-0.0608	0.0611	-0.9946	0.3199
207550_at	MPL	0.0296	0.03	0.9879	0.3232
206167_s_at	ARHGAP6	0.161	0.1631	0.9868	0.3237
207815_at	PF4V1	-0.0301	0.0352	-0.8557	0.3922
225354_s_at	SH3BGRL2	0.0631	0.0743	0.8504	0.3951
207206_s_at	ALOX12	-0.021	0.0267	-0.7848	0.4326
205442_at	MFAP3L	-0.0908	0.1444	-0.6284	0.5297

Meta DvA

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
229778_at	C12ORF39	0.0092	0.0211	0.4355	0.6632
210387_at	HIST1H2BG	0.0548	0.1351	0.4057	0.685
212573_at	ENDOD1	0.0307	0.0772	0.3976	0.691
230690_at	TUBB1	0.0094	0.0247	0.3814	0.7029
206049_at	SELP	0.0186	0.0552	0.3366	0.7364
206390_x_at	PF4	-0.0253	0.0859	-0.2949	0.7681
230942_at	CMTM5	-0.0098	0.0353	-0.2782	0.7809
206110_at	HIST1H3H	-0.0169	0.0881	-0.1917	0.848
215779_s_at	HIST1H2BG	0.0158	0.0895	0.1769	0.8596
201059_at	CTTN	-0.0064	0.0569	-0.1131	0.9099
227088_at	PDE5A	0.0124	0.1197	0.1033	0.9177
1555659_a_at	TREML1	-0.0031	0.0333	-0.092	0.9267
227180_at	ELOVL7	-0.0115	0.1343	-0.0853	0.932
1553842_at	BEND2	0.0006	0.0294	0.0213	0.983

Meta BvA

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
201058_s_at	MYL9	0.6302	0.1457	4.3263	0
207414_s_at	PCSK6	-0.3652	0.084	-4.3468	0
227189_at	CPNE5	-0.2078	0.0485	-4.2831	0
200665_s_at	SPARC	0.351	0.0956	3.6708	0.0002
226152_at	TTC7B	0.2972	0.0853	3.4855	0.0005
212667_at	SPARC	0.3599	0.1186	3.0356	0.0024
224823_at	MYLK	0.3958	0.1306	3.0295	0.0024
203817_at	GUCY1B3	0.2668	0.0896	2.978	0.0029
212077_at	CALD1	0.2677	0.0917	2.9182	0.0035
209651_at	TGFB1I1	0.2968	0.1068	2.7783	0.0055
204628_s_at	ITGB3	0.0907	0.0333	2.7233	0.0065
201108_s_at	THBS1	0.3143	0.1159	2.7125	0.0067
219090_at	SLC24A3	0.2353	0.1001	2.3498	0.0188
212813_at	JAM3	0.1787	0.0792	2.2569	0.024
206049_at	SELP	0.1132	0.0513	2.2068	0.0273
37966_at	PARVB	0.2096	0.0967	2.1668	0.0302
225974_at	TMEM64	0.2798	0.1312	2.1336	0.0329
214974_x_at	CXCL5	0.615	0.3006	2.046	0.0408
57588_at	SLC24A3	0.1824	0.0969	1.8823	0.0598
204115_at	GNG11	0.1814	0.0979	1.8528	0.0639
207156_at	HIST1H2AG	-0.1658	0.091	-1.8227	0.0684
1560262_at	LRRC32	-0.047	0.026	-1.8049	0.0711
225354_s_at	SH3BGRL2	-0.123	0.069	-1.7824	0.0747
203680_at	PRKAR2B	-0.2534	0.1448	-1.75	0.0801
205442_at	MFAP3L	-0.2348	0.1342	-1.7498	0.0802
208792_s_at	CLU	0.2209	0.1268	1.7417	0.0816
227180_at	ELOVL7	0.2162	0.1248	1.7326	0.0832
206167_s_at	ARHGAP6	0.2616	0.1516	1.7261	0.0843
206494_s_at	ITGA2B	-0.0336	0.0204	-1.6513	0.0987
212651_at	RHOBTB1	0.1514	0.0923	1.6397	0.1011
208782_at	FSTL1	0.1443	0.0937	1.5391	0.1238
222717_at	SDPR	0.1725	0.1131	1.5247	0.1273
208791_at	CLU	0.1961	0.1333	1.4714	0.1412
1553842_at	BEND2	0.0387	0.0273	1.4168	0.1565
207808_s_at	PROS1	0.1569	0.1147	1.3688	0.1711
214146_s_at	PPBP	0.384	0.2954	1.2999	0.1936
241133_at	PRSS1	-0.0327	0.0263	-1.2419	0.2143
221556_at	CDC14B	-0.0888	0.073	-1.2155	0.2242
226188_at	LGALSL	-0.0831	0.0688	-1.2071	0.2274
210986_s_at	TPM1	0.0681	0.0592	1.1499	0.2502
216956_s_at	ITGA2B	-0.0337	0.0293	-1.1492	0.2505

Meta BvA

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
225166_at	ARHGAP18	-0.0892	0.0866	-1.0305	0.3028
207815_at	PF4V1	0.0331	0.0327	1.0111	0.312
230942_at	CMTM5	-0.0316	0.0329	-0.9623	0.3359
207206_s_at	ALOX12	0.0228	0.0249	0.9189	0.3581
203819_s_at	IGF2BP3	0.1815	0.2017	0.8996	0.3683
1555659_a_at	TREML1	-0.0252	0.031	-0.8131	0.4162
235331_x_at	PCGF5	-0.0851	0.1049	-0.8112	0.4173
228708_at	RAB27B	0.1284	0.1587	0.8088	0.4186
210387_at	HIST1H2BG	-0.1002	0.1255	-0.7986	0.4245
230690_at	TUBB1	0.018	0.023	0.7841	0.433
206110_at	HIST1H3H	-0.0615	0.082	-0.7501	0.4532
206390_x_at	PF4	-0.0595	0.0799	-0.7452	0.4561
215779_s_at	HIST1H2BG	-0.0588	0.0832	-0.7067	0.4797
1552773_at	CLEC4D	0.0175	0.025	0.7013	0.4831
212151_at	PBX1	0.051	0.0806	0.6325	0.5271
201906_s_at	CTDSPL	-0.0359	0.0568	-0.6315	0.5277
220496_at	CLEC1B	0.0083	0.0222	0.3713	0.7104
229778_at	C12ORF39	0.006	0.0196	0.3087	0.7576
208601_s_at	TUBB1	-0.0063	0.021	-0.2993	0.7647
203414_at	MMD	0.0129	0.0597	0.2164	0.8287
206272_at	RAB4A	0.0136	0.0665	0.2046	0.8379
212148_at	PBX1	0.0171	0.1123	0.1522	0.879
212573_at	ENDOD1	0.0085	0.0718	0.1188	0.9055
206493_at	ITGA2B	0.0055	0.0474	0.1158	0.9078
211026_s_at	MGLL	-0.0052	0.0825	-0.0628	0.9499
227451_s_at	C6ORF79	0.004	0.0788	0.0514	0.959
230645_at	FRMD3	0.0048	0.115	0.0415	0.9669
207550_at	MPL	0.0008	0.0279	0.0301	0.976
206655_s_at	GP1BB	-0.0012	0.0489	-0.0237	0.9811
201059_at	CTTN	-0.0011	0.0529	-0.0206	0.9836
227088_at	PDE5A	0.0013	0.1112	0.0121	0.9904

Meta CvA

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
200665_s_at	SPARC	0.5088	0.0959	5.3072	0
201058_s_at	MYL9	0.7394	0.1461	5.0619	0
201108_s_at	THBS1	0.5727	0.1162	4.9274	0
204115_at	GNG11	0.4012	0.0982	4.0847	0
207414_s_at	PCSK6	-0.3802	0.0843	-4.5112	0
209651_at	TGFB1I1	0.5039	0.1071	4.704	0
212077_at	CALD1	0.4484	0.092	4.8743	0
212667_at	SPARC	0.6249	0.1189	5.2549	0
226152_at	TTC7B	0.4764	0.0855	5.5717	0
203817_at	GUCY1B3	0.3611	0.0898	4.0194	0.0001
212813_at	JAM3	0.3066	0.0794	3.8617	0.0001
224823_at	MYLK	0.522	0.131	3.9849	0.0001
208782_at	FSTL1	0.3534	0.094	3.7594	0.0002
219090_at	SLC24A3	0.3441	0.1004	3.4258	0.0006
57588_at	SLC24A3	0.3158	0.0972	3.2498	0.0012
1560262_at	LRRC32	-0.084	0.0261	-3.2157	0.0013
214974_x_at	CXCL5	0.9665	0.3016	3.2049	0.0014
227189_at	CPNE5	-0.1505	0.0487	-3.092	0.002
225974_at	TMEM64	0.3938	0.1316	2.9932	0.0028
207808_s_at	PROS1	0.3417	0.115	2.9714	0.003
212651_at	RHOBTB1	0.2616	0.0926	2.8255	0.0047
1555659_a_at	TREML1	-0.08	0.0311	-2.5735	0.0101
208792_s_at	CLU	0.3252	0.1272	2.5574	0.0105
212573_at	ENDOD1	0.1792	0.072	2.4891	0.0128
37966_at	PARVB	0.2414	0.0971	2.4862	0.0129
1553842_at	BEND2	0.0654	0.0274	2.3865	0.017
208791_at	CLU	0.3113	0.1337	2.3287	0.0199
216956_s_at	ITGA2B	-0.0677	0.0294	-2.3018	0.0213
206494_s_at	ITGA2B	-0.0457	0.0204	-2.2389	0.0252
220496_at	CLEC1B	-0.0483	0.0223	-2.1673	0.0302
203680_at	PRKAR2B	-0.3109	0.1452	-2.1409	0.0323
203819_s_at	IGF2BP3	0.4268	0.2024	2.1086	0.035
203414_at	MMD	0.1232	0.0599	2.0567	0.0397
241133_at	PRSS1	-0.0526	0.0264	-1.9919	0.0464
222717_at	SDPR	0.2238	0.1134	1.9738	0.0484
207815_at	PF4V1	0.0542	0.0328	1.6502	0.0989
214146_s_at	PPBP	0.4741	0.2964	1.5994	0.1097
225166_at	ARHGAP18	-0.1329	0.0868	-1.5302	0.126
206167_s_at	ARHGAP6	0.2218	0.152	1.4585	0.1447
208601_s_at	TUBB1	-0.0301	0.0211	-1.4294	0.1529
230942_at	CMTM5	-0.0461	0.0329	-1.398	0.1621

Meta CvA

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
205442_at	MFAP3L	-0.1801	0.1346	-1.3378	0.181
206272_at	RAB4A	0.0838	0.0667	1.2556	0.2093
204628_s_at	ITGB3	0.0411	0.0334	1.2307	0.2184
230690_at	TUBB1	-0.0281	0.023	-1.2192	0.2228
211026_s_at	MGLL	0.0928	0.0828	1.1213	0.2622
227180_at	ELOVL7	-0.1224	0.1252	-0.9774	0.3284
212148_at	PBX1	0.1085	0.1126	0.9629	0.3356
206493_at	ITGA2B	-0.0425	0.0475	-0.8948	0.3709
201906_s_at	CTDSPL	0.0506	0.057	0.8878	0.3746
235331_x_at	PCGF5	-0.085	0.1053	-0.808	0.4191
225354_s_at	SH3BGRL2	-0.0488	0.0692	-0.7048	0.481
207206_s_at	ALOX12	-0.0173	0.0249	-0.6957	0.4866
212151_at	PBX1	0.0561	0.0808	0.6941	0.4876
1552773_at	CLEC4D	-0.0171	0.0251	-0.6813	0.4957
201059_at	CTTN	0.0354	0.0531	0.6681	0.5041
206049_at	SELP	0.0323	0.0515	0.6273	0.5305
206390_x_at	PF4	-0.0487	0.0801	-0.6077	0.5434
206110_at	HIST1H3H	-0.0484	0.0822	-0.589	0.5558
210387_at	HIST1H2BG	0.0653	0.1259	0.5188	0.6039
229778_at	C12ORF39	-0.0071	0.0197	-0.3622	0.7172
226188_at	LGALSL	-0.0192	0.069	-0.2776	0.7813
227088_at	PDE5A	0.0298	0.1116	0.2671	0.7894
207550_at	MPL	-0.0036	0.028	-0.128	0.8981
215779_s_at	HIST1H2BG	-0.0098	0.0835	-0.1177	0.9063
221556_at	CDC14B	0.0053	0.0732	0.0728	0.9419
210986_s_at	TPM1	-0.0034	0.0594	-0.0567	0.9548
230645_at	FRMD3	0.0063	0.1153	0.0548	0.9563
227451_s_at	C6ORF79	0.0032	0.079	0.0403	0.9679
207156_at	HIST1H2AG	-0.0034	0.0912	-0.0372	0.9703
228708_at	RAB27B	-0.0039	0.1592	-0.0245	0.9805
206655_s_at	GP1BB	0.0004	0.049	0.0085	0.9932

Meta DvA

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
203680_at	PRKAR2B	-0.6499	0.1558	-4.1718	0
212651_at	RHOBTB1	0.4089	0.0993	4.1185	0
201058_s_at	MYL9	0.6082	0.1566	3.8835	0.0001
226152_at	TTC7B	0.3655	0.0917	3.9878	0.0001
227189_at	CPNE5	-0.2094	0.0522	-4.0127	0.0001
207414_s_at	PCSK6	-0.3381	0.0904	-3.7384	0.0002
212077_at	CALD1	0.3534	0.0986	3.5831	0.0003
224823_at	MYLK	0.5077	0.1404	3.616	0.0003
212667_at	SPARC	0.4303	0.1275	3.3752	0.0007
209651_at	TGFB1I1	0.3813	0.1148	3.3208	0.0009
212813_at	JAM3	0.2827	0.0851	3.3205	0.0009
201108_s_at	THBS1	0.4083	0.1247	3.274	0.0011
200665_s_at	SPARC	0.3298	0.1027	3.21	0.0013
57588_at	SLC24A3	0.3349	0.1042	3.2126	0.0013
219090_at	SLC24A3	0.3437	0.1077	3.1918	0.0014
216956_s_at	ITGA2B	-0.093	0.0315	-2.9512	0.0032
203817_at	GUCY1B3	0.2673	0.0963	2.7761	0.0055
222717_at	SDPR	0.3063	0.1215	2.522	0.0117
206494_s_at	ITGA2B	-0.0514	0.0219	-2.3496	0.0188
212148_at	PBX1	0.2828	0.1209	2.34	0.0193
1552773_at	CLEC4D	-0.0628	0.0268	-2.3372	0.0194
208782_at	FSTL1	0.2304	0.1008	2.2856	0.0223
207808_s_at	PROS1	0.2797	0.1234	2.2679	0.0233
1560262_at	LRRC32	-0.0631	0.028	-2.2505	0.0244
204115_at	GNG11	0.2326	0.1053	2.2091	0.0272
206272_at	RAB4A	0.152	0.0716	2.1229	0.0338
214974_x_at	CXCL5	0.6752	0.3236	2.0867	0.0369
212151_at	PBX1	0.1782	0.0867	2.0558	0.0398
206655_s_at	GP1BB	-0.1053	0.0525	-2.0031	0.0452
228708_at	RAB27B	-0.332	0.1709	-1.9429	0.052
225974_at	TMEM64	0.2412	0.1411	1.709	0.0874
203414_at	MMD	0.1091	0.0643	1.698	0.0895
208791_at	CLU	0.2405	0.1433	1.6789	0.0932
241133_at	PRSS1	-0.0474	0.0283	-1.6765	0.0936
208792_s_at	CLU	0.2247	0.1363	1.6489	0.0992
220496_at	CLEC1B	-0.0371	0.0239	-1.5526	0.1205
221556_at	CDC14B	0.1201	0.0785	1.5291	0.1262
235331_x_at	PCGF5	-0.1656	0.1129	-1.4659	0.1427
210986_s_at	TPM1	0.0916	0.0637	1.4382	0.1504
204628_s_at	ITGB3	0.0479	0.0358	1.3395	0.1804
206493_at	ITGA2B	-0.0668	0.0509	-1.3106	0.19

Meta DvA

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
208601_s_at	TUBB1	-0.027	0.0226	-1.1993	0.2304
203819_s_at	IGF2BP3	0.2558	0.2172	1.178	0.2388
214146_s_at	PPBP	0.3709	0.318	1.1662	0.2435
225166_at	ARHGAP18	-0.1044	0.0931	-1.1215	0.2621
211026_s_at	MGLL	0.0985	0.0888	1.1095	0.2672
227451_s_at	C6ORF79	0.0931	0.0848	1.0982	0.2721
207156_at	HIST1H2AG	0.0987	0.0978	1.0095	0.3127
37966_at	PARVB	0.1051	0.1042	1.0088	0.3131
230645_at	FRMD3	-0.1245	0.1235	-1.008	0.3135
226188_at	LGALSL	0.0741	0.0741	1.0009	0.3169
201906_s_at	CTDSPL	-0.0608	0.0611	-0.9946	0.3199
207550_at	MPL	0.0296	0.03	0.9879	0.3232
206167_s_at	ARHGAP6	0.161	0.1631	0.9868	0.3237
207815_at	PF4V1	-0.0301	0.0352	-0.8557	0.3922
225354_s_at	SH3BGRL2	0.0631	0.0743	0.8504	0.3951
207206_s_at	ALOX12	-0.021	0.0267	-0.7848	0.4326
205442_at	MFAP3L	-0.0908	0.1444	-0.6284	0.5297
229778_at	C12ORF39	0.0092	0.0211	0.4355	0.6632
210387_at	HIST1H2BG	0.0548	0.1351	0.4057	0.685
212573_at	ENDOD1	0.0307	0.0772	0.3976	0.691
230690_at	TUBB1	0.0094	0.0247	0.3814	0.7029
206049_at	SELP	0.0186	0.0552	0.3366	0.7364
206390_x_at	PF4	-0.0253	0.0859	-0.2949	0.7681
230942_at	CMTM5	-0.0098	0.0353	-0.2782	0.7809
206110_at	HIST1H3H	-0.0169	0.0881	-0.1917	0.848
215779_s_at	HIST1H2BG	0.0158	0.0895	0.1769	0.8596
201059_at	CTTN	-0.0064	0.0569	-0.1131	0.9099
227088_at	PDE5A	0.0124	0.1197	0.1033	0.9177
1555659_a_at	TREML1	-0.0031	0.0333	-0.092	0.9267
227180_at	ELOVL7	-0.0115	0.1343	-0.0853	0.932
1553842_at	BEND2	0.0006	0.0294	0.0213	0.983

probe set	GeneName	p-value	coef	se(coef)
200665_s_at	SPARC	0.0007	1.16	0.3656
226152_at	TTC7B	0.0068	1.0027	0.3797
201059_at	CTTN	0.0075	1.77	0.708
212667_at	SPARC	0.01	0.6115	0.2392
206494_s_at	ITGA2B	0.0124	-4.2229	1.7721
209651_at	TGFB1I1	0.015	0.6664	0.2838
207808_s_at	PROS1	0.0158	0.6132	0.2557
206167_s_at	ARHGAP6	0.0232	0.3732	0.1564
212077_at	CALD1	0.025	0.7635	0.3588
225974_at	TMEM64	0.0353	0.4378	0.2198
212573_at	ENDOD1	0.0404	0.8399	0.4159
241133_at	PRSS1	0.0502	-2.4021	1.2565
208782_at	FSTL1	0.0517	0.5833	0.2949
207414_s_at	PCSK6	0.062	-0.5893	0.311
201108_s_at	THBS1	0.0669	0.3965	0.2163
212813_at	JAM3	0.0776	0.5943	0.325
201058_s_at	MYL9	0.1035	0.3212	0.1993
228708_at	RAB27B	0.1179	0.2355	0.1473
210986_s_at	TPM1	0.125	0.7216	0.4965
216956_s_at	ITGA2B	0.135	-1.6176	1.1078
201906_s_at	CTDSPL	0.1811	-0.6141	0.4608
1555659_a_at	TREML1	0.1919	-1.1708	0.9216
230942_at	CMTM5	0.1939	-1.1224	0.875
206390_x_at	PF4	0.2164	-0.4466	0.3727
221556_at	CDC14B	0.2219	0.4561	0.3736
204115_at	GNG11	0.2335	0.3405	0.2805
207156_at	HIST1H2AG	0.2611	-0.3583	0.3344
206110_at	HIST1H3H	0.2786	-0.3734	0.3557
222717_at	SDPR	0.2947	-0.403	0.3962

probe set	GeneName	p-value	coef	se(coef)
226188_at	LGALSL	0.2975	-0.3809	0.3643
1553842_at	BEND2	0.324	-0.8626	0.8985
207815_at	PF4V1	0.3257	-0.8234	0.925
206272_at	RAB4A	0.3294	-0.3819	0.393
203680_at	PRKAR2B	0.3633	-0.1792	0.1967
207206_s_at	ALOX12	0.4231	-1.0305	1.3097
203819_s_at	IGF2BP3	0.4346	0.0962	0.121
214146_s_at	PPBP	0.4367	0.0595	0.0743
37966_at	PARVB	0.4385	0.1988	0.259
206493_at	ITGA2B	0.4508	-0.4913	0.6512
210387_at	HIST1H2BG	0.4719	0.1475	0.2022
229778_at	C12ORF39	0.4795	0.671	0.8788
204628_s_at	ITGB3	0.4917	0.5964	0.8558
208601_s_at	TUBB1	0.5099	-1.0219	1.5701
214974_x_at	CXCL5	0.5212	0.0544	0.0843
207550_at	MPL	0.5222	0.6329	0.978
219090_at	SLC24A3	0.5349	0.178	0.2873
227189_at	CPNE5	0.5686	0.3442	0.5912
220496_at	CLEC1B	0.5744	-0.7464	1.3398
225354_s_at	SH3BGRL2	0.583	0.1957	0.3607
1552773_at	CLEC4D	0.5937	-0.6349	1.1939
230690_at	TUBB1	0.6484	0.628	1.3766
1560262_at	LRRC32	0.6493	-0.4511	0.9989
212651_at	RHOBTB1	0.6608	-0.1288	0.2925
212151_at	PBX1	0.673	-0.1516	0.3599
203414_at	MMD	0.677	0.185	0.4474
57588_at	SLC24A3	0.6775	0.1189	0.2862
211026_s_at	MGLL	0.7065	-0.1168	0.3082
227180_at	ELOVL7	0.7329	0.0668	0.1962

probe set	GeneName	p-value	coef	se(coef)
215779_s_at	HIST1H2BG	0.7456	0.112	0.3413
212148_at	PBX1	0.7633	-0.0689	0.2277
224823_at	MYLK	0.7851	0.0609	0.2239
205442_at	MFAP3L	0.8084	0.0464	0.1905
227451_s_at	C6ORF79	0.8179	0.0711	0.3084
206049_at	SELP	0.8328	-0.1165	0.552
225166_at	ARHGAP18	0.8335	0.0678	0.3233
227088_at	PDE5A	0.8734	-0.0406	0.2545
203817_at	GUCY1B3	0.9191	-0.0357	0.3508
208791_at	CLU	0.9398	0.0199	0.2633
230645_at	FRMD3	0.9476	0.0165	0.2509
208792_s_at	CLU	0.958	-0.0155	0.2937
235331_x_at	PCGF5	0.9848	0.0039	0.2065
206655_s_at	GP1BB	0.999	-0.0008	0.6498

Meta Survival

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
200665_s_at	SPARC	0.7364	0.1931	3.8134	0.0001
212077_at	CALD1	0.7018	0.2081	3.3726	0.0007
212667_at	SPARC	0.4668	0.141	3.3097	0.0009
226152_at	TTC7B	0.6533	0.2014	3.2433	0.0012
209651_at	TGFB1I1	0.4801	0.1561	3.0765	0.0021
212573_at	ENDOD1	0.6567	0.2234	2.9399	0.0033
208782_at	FSTL1	0.516	0.1766	2.9224	0.0035
206167_s_at	ARHGAP6	0.2821	0.0998	2.8268	0.0047
201059_at	CTTN	1.0095	0.3676	2.7465	0.006
206494_s_at	ITGA2B	-2.2302	0.8628	-2.5847	0.0097
207414_s_at	PCSK6	-0.5022	0.1959	-2.5635	0.0104
201058_s_at	MYL9	0.2788	0.1092	2.5539	0.0107
228708_at	RAB27B	0.2425	0.0962	2.521	0.0117
207808_s_at	PROS1	0.386	0.1547	2.4958	0.0126
201108_s_at	THBS1	0.3303	0.1337	2.4702	0.0135
212813_at	JAM3	0.4729	0.1953	2.4208	0.0155
1555659_a_at	TREML1	-1.3264	0.5507	-2.4084	0.016
225974_at	TMEM64	0.3188	0.1335	2.3877	0.017
207550_at	MPL	1.1338	0.5744	1.974	0.0484
241133_at	PRSS1	-1.3065	0.6851	-1.907	0.0565
204115_at	GNG11	0.3039	0.1641	1.8526	0.0639
210986_s_at	TPM1	0.4896	0.2826	1.7327	0.0831
216956_s_at	ITGA2B	-0.9809	0.605	-1.6215	0.1049
206390_x_at	PF4	-0.3201	0.2155	-1.4851	0.1375
207815_at	PF4V1	-0.8771	0.5949	-1.4744	0.1404
203680_at	PRKAR2B	-0.172	0.1172	-1.4678	0.1422
230942_at	CMTM5	-0.6783	0.5001	-1.3563	0.175
226188_at	LGALSL	-0.304	0.2289	-1.3285	0.184
206272_at	RAB4A	-0.3239	0.2447	-1.3238	0.1856

Meta Survival

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
224823_at	MYLK	0.1472	0.1245	1.183	0.2368
221556_at	CDC14B	0.2488	0.2148	1.1582	0.2468
212151_at	PBX1	-0.251	0.2174	-1.155	0.2481
207156_at	HIST1H2AG	-0.2145	0.1925	-1.114	0.2653
1553842_at	BEND2	-0.6083	0.5646	-1.0774	0.2813
1552773_at	CLEC4D	-0.712	0.6731	-1.0577	0.2902
230690_at	TUBB1	0.8528	0.8113	1.0511	0.2932
214146_s_at	PPBP	0.0504	0.049	1.028	0.304
206110_at	HIST1H3H	-0.196	0.197	-0.9951	0.3197
206493_at	ITGA2B	-0.3381	0.3563	-0.949	0.3426
229778_at	C12ORF39	0.5116	0.5699	0.8977	0.3693
214974_x_at	CXCL5	0.0476	0.0533	0.8916	0.3726
1560262_at	LRRC32	-0.5451	0.6605	-0.8253	0.4092
212148_at	PBX1	-0.1213	0.1472	-0.8236	0.4101
203819_s_at	IGF2BP3	0.0623	0.0758	0.8217	0.4112
37966_at	PARVB	0.1255	0.1677	0.7483	0.4543
203817_at	GUCY1B3	0.1328	0.1922	0.691	0.4896
210387_at	HIST1H2BG	0.078	0.1188	0.6569	0.5112
208791_at	CLU	0.0766	0.1305	0.5873	0.557
225354_s_at	SH3BGRL2	0.1346	0.2311	0.5823	0.5604
219090_at	SLC24A3	0.0984	0.1698	0.5792	0.5624
222717_at	SDPR	-0.0931	0.1647	-0.5656	0.5717
201906_s_at	CTDSPL	-0.1523	0.2745	-0.5549	0.5789
206049_at	SELP	-0.1659	0.3339	-0.497	0.6192
204628_s_at	ITGB3	0.2112	0.4554	0.4638	0.6428
225166_at	ARHGAP18	0.0889	0.2003	0.4437	0.6572
205442_at	MFAP3L	-0.0524	0.124	-0.4227	0.6725
235331_x_at	PCGF5	-0.0587	0.1406	-0.4173	0.6764
215779_s_at	HIST1H2BG	0.0741	0.1798	0.412	0.6803

Meta Survival

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
227088_at	PDE5A	-0.0586	0.1532	-0.3826	0.702
220496_at	CLEC1B	0.2904	0.775	0.3747	0.7079
212651_at	RHOBTB1	-0.0618	0.1722	-0.3589	0.7197
57588_at	SLC24A3	0.0595	0.173	0.3439	0.7309
211026_s_at	MGLL	-0.0582	0.1828	-0.3183	0.7502
208601_s_at	TUBB1	-0.2562	0.8365	-0.3063	0.7594
227189_at	CPNE5	0.0779	0.3372	0.2311	0.8173
227180_at	ELOVL7	0.0257	0.127	0.2026	0.8394
208792_s_at	CLU	0.0219	0.1363	0.1605	0.8725
227451_s_at	C6ORF79	-0.0188	0.1968	-0.0955	0.9239
230645_at	FRMD3	-0.0117	0.1308	-0.0897	0.9285
206655_s_at	GP1BB	0.0153	0.3192	0.048	0.9617
203414_at	MMD	0.0125	0.2727	0.0458	0.9634
207206_s_at	ALOX12	0.0106	0.5903	0.018	0.9857

Jorissen Survival

probe set	GeneName	Pvalue	coef	se(coef)
212077_at	CALD1	0.0066	0.6705	0.2554
200665_s_at	SPARC	0.0084	0.5725	0.2274
212667_at	SPARC	0.0233	0.3896	0.1746
208782_at	FSTL1	0.0296	0.4784	0.2204
209651_at	TGFB1I1	0.0302	0.3993	0.1868
226152_at	TTC7B	0.0302	0.5165	0.2376
212573_at	ENDOD1	0.0304	0.5824	0.2648
1555659_a_at	TREML1	0.0345	-1.4128	0.6869
201058_s_at	MYL9	0.0476	0.2607	0.1305
207550_at	MPL	0.0516	1.3975	0.7096
228708_at	RAB27B	0.0539	0.2477	0.127
207414_s_at	PCSK6	0.0793	-0.4449	0.2523
201059_at	CTTN	0.0846	0.7288	0.4301
201108_s_at	THBS1	0.0902	0.2893	0.1701
206494_s_at	ITGA2B	0.0925	-1.611	0.9879
206167_s_at	ARHGAP6	0.101	0.2195	0.1296
212813_at	JAM3	0.1075	0.4042	0.2444
225974_at	TMEM64	0.1251	0.2493	0.1681
204115_at	GNG11	0.1669	0.285	0.2023
207808_s_at	PROS1	0.1871	0.2549	0.1942
207815_at	PF4V1	0.1947	-0.915	0.7769
224823_at	MYLK	0.2161	0.1859	0.1497
203680_at	PRKAR2B	0.2459	-0.1681	0.1459
212151_at	PBX1	0.257	-0.3082	0.2727
210986_s_at	TPM1	0.2625	0.3785	0.3436
241133_at	PRSS1	0.2909	-0.843	0.8173
206390_x_at	PF4	0.3192	-0.2565	0.2642
216956_s_at	ITGA2B	0.3222	-0.7104	0.7221
230690_at	TUBB1	0.342	0.9725	1.0043
1552773_at	CLEC4D	0.3508	-0.7479	0.8151
206272_at	RAB4A	0.3584	-0.2872	0.3127
203817_at	GUCY1B3	0.3698	0.2051	0.2298
226188_at	LGALSL	0.3901	-0.254	0.2942
220496_at	CLEC1B	0.3923	0.8118	0.9501
212148_at	PBX1	0.4161	-0.1589	0.193
205442_at	MFAP3L	0.439	-0.1251	0.1634
230942_at	CMTM5	0.4431	-0.4628	0.6094
1560262_at	LRRC32	0.478	-0.6182	0.8804
214146_s_at	PPBP	0.5167	0.0434	0.0653
206493_at	ITGA2B	0.5229	-0.2727	0.4257
208791_at	CLU	0.5314	0.0951	0.1502

Jorissen Survival

probe set	GeneName	Pvalue	coef	se(coef)
214974_x_at	CXCL5	0.5338	0.043	0.0689
207156_at	HIST1H2AG	0.5364	-0.1431	0.2355
1553842_at	BEND2	0.5367	-0.4423	0.7259
235331_x_at	PCGF5	0.5579	-0.1128	0.192
221556_at	CDC14B	0.577	0.1465	0.2625
229778_at	C12ORF39	0.6112	0.3959	0.7486
206110_at	HIST1H3H	0.6132	-0.1176	0.2366
206049_at	SELP	0.6409	-0.1945	0.4193
203819_s_at	IGF2BP3	0.6816	0.0404	0.0973
207206_s_at	ALOX12	0.6823	0.2761	0.6613
225166_at	ARHGAP18	0.6885	0.102	0.2552
227088_at	PDE5A	0.7191	-0.0688	0.1919
37966_at	PARVB	0.7416	0.0725	0.2201
227451_s_at	C6ORF79	0.7525	-0.0806	0.2557
225354_s_at	SH3BGRL2	0.7589	0.092	0.3011
201906_s_at	CTDSPL	0.7655	0.1017	0.3418
210387_at	HIST1H2BG	0.779	0.0414	0.1467
215779_s_at	HIST1H2BG	0.7802	0.0595	0.2115
219090_at	SLC24A3	0.7915	0.0556	0.2106
203414_at	MMD	0.7946	-0.0895	0.344
208792_s_at	CLU	0.8355	0.0321	0.1538
222717_at	SDPR	0.8743	-0.0284	0.1811
230645_at	FRMD3	0.884	-0.0223	0.1533
227189_at	CPNE5	0.9019	-0.0505	0.4105
212651_at	RHOBTB1	0.902	-0.0263	0.213
57588_at	SLC24A3	0.9072	0.0253	0.2173
211026_s_at	MGLL	0.9075	-0.0264	0.227
204628_s_at	ITGB3	0.9127	0.0591	0.5379
206655_s_at	GP1BB	0.9554	0.0205	0.3664
208601_s_at	TUBB1	0.9619	0.0473	0.9885
227180_at	ELOVL7	0.9815	-0.0039	0.1665

Meta Survival

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
200665_s_at	SPARC	0.7364	0.1931	3.8134	0.0001
212077_at	CALD1	0.7018	0.2081	3.3726	0.0007
212667_at	SPARC	0.4668	0.141	3.3097	0.0009
226152_at	TTC7B	0.6533	0.2014	3.2433	0.0012
209651_at	TGFB1I1	0.4801	0.1561	3.0765	0.0021
212573_at	ENDOD1	0.6567	0.2234	2.9399	0.0033
208782_at	FSTL1	0.516	0.1766	2.9224	0.0035
206167_s_at	ARHGAP6	0.2821	0.0998	2.8268	0.0047
201059_at	CTTN	1.0095	0.3676	2.7465	0.006
206494_s_at	ITGA2B	-2.2302	0.8628	-2.5847	0.0097
207414_s_at	PCSK6	-0.5022	0.1959	-2.5635	0.0104
201058_s_at	MYL9	0.2788	0.1092	2.5539	0.0107
228708_at	RAB27B	0.2425	0.0962	2.521	0.0117
207808_s_at	PROS1	0.386	0.1547	2.4958	0.0126
201108_s_at	THBS1	0.3303	0.1337	2.4702	0.0135
212813_at	JAM3	0.4729	0.1953	2.4208	0.0155
1555659_a_at	TREML1	-1.3264	0.5507	-2.4084	0.016
225974_at	TMEM64	0.3188	0.1335	2.3877	0.017
207550_at	MPL	1.1338	0.5744	1.974	0.0484
241133_at	PRSS1	-1.3065	0.6851	-1.907	0.0565
204115_at	GNG11	0.3039	0.1641	1.8526	0.0639
210986_s_at	TPM1	0.4896	0.2826	1.7327	0.0831
216956_s_at	ITGA2B	-0.9809	0.605	-1.6215	0.1049
206390_x_at	PF4	-0.3201	0.2155	-1.4851	0.1375
207815_at	PF4V1	-0.8771	0.5949	-1.4744	0.1404
203680_at	PRKAR2B	-0.172	0.1172	-1.4678	0.1422
230942_at	CMTM5	-0.6783	0.5001	-1.3563	0.175
226188_at	LGALSL	-0.304	0.2289	-1.3285	0.184
206272_at	RAB4A	-0.3239	0.2447	-1.3238	0.1856
224823_at	MYLK	0.1472	0.1245	1.183	0.2368
221556_at	CDC14B	0.2488	0.2148	1.1582	0.2468
212151_at	PBX1	-0.251	0.2174	-1.155	0.2481
207156_at	HIST1H2AG	-0.2145	0.1925	-1.114	0.2653
1553842_at	BEND2	-0.6083	0.5646	-1.0774	0.2813
1552773_at	CLEC4D	-0.712	0.6731	-1.0577	0.2902
230690_at	TUBB1	0.8528	0.8113	1.0511	0.2932
214146_s_at	PPBP	0.0504	0.049	1.028	0.304
206110_at	HIST1H3H	-0.196	0.197	-0.9951	0.3197
206493_at	ITGA2B	-0.3381	0.3563	-0.949	0.3426
229778_at	C12ORF39	0.5116	0.5699	0.8977	0.3693
214974_x_at	CXCL5	0.0476	0.0533	0.8916	0.3726

Meta Survival

probe set	GeneName	Effect.fixed	SE.fixed	Z.fixed	Pvalue.fixed
1560262_at	LRRC32	-0.5451	0.6605	-0.8253	0.4092
212148_at	PBX1	-0.1213	0.1472	-0.8236	0.4101
203819_s_at	IGF2BP3	0.0623	0.0758	0.8217	0.4112
37966_at	PARVB	0.1255	0.1677	0.7483	0.4543
203817_at	GUCY1B3	0.1328	0.1922	0.691	0.4896
210387_at	HIST1H2BG	0.078	0.1188	0.6569	0.5112
208791_at	CLU	0.0766	0.1305	0.5873	0.557
225354_s_at	SH3BGRL2	0.1346	0.2311	0.5823	0.5604
219090_at	SLC24A3	0.0984	0.1698	0.5792	0.5624
222717_at	SDPR	-0.0931	0.1647	-0.5656	0.5717
201906_s_at	CTDSPL	-0.1523	0.2745	-0.5549	0.5789
206049_at	SELP	-0.1659	0.3339	-0.497	0.6192
204628_s_at	ITGB3	0.2112	0.4554	0.4638	0.6428
225166_at	ARHGAP18	0.0889	0.2003	0.4437	0.6572
205442_at	MFAP3L	-0.0524	0.124	-0.4227	0.6725
235331_x_at	PCGF5	-0.0587	0.1406	-0.4173	0.6764
215779_s_at	HIST1H2BG	0.0741	0.1798	0.412	0.6803
227088_at	PDE5A	-0.0586	0.1532	-0.3826	0.702
220496_at	CLEC1B	0.2904	0.775	0.3747	0.7079
212651_at	RHOBTB1	-0.0618	0.1722	-0.3589	0.7197
57588_at	SLC24A3	0.0595	0.173	0.3439	0.7309
211026_s_at	MGLL	-0.0582	0.1828	-0.3183	0.7502
208601_s_at	TUBB1	-0.2562	0.8365	-0.3063	0.7594
227189_at	CPNE5	0.0779	0.3372	0.2311	0.8173
227180_at	ELOVL7	0.0257	0.127	0.2026	0.8394
208792_s_at	CLU	0.0219	0.1363	0.1605	0.8725
227451_s_at	C6ORF79	-0.0188	0.1968	-0.0955	0.9239
230645_at	FRMD3	-0.0117	0.1308	-0.0897	0.9285
206655_s_at	GP1BB	0.0153	0.3192	0.048	0.9617
203414_at	MMD	0.0125	0.2727	0.0458	0.9634
207206_s_at	ALOX12	0.0106	0.5903	0.018	0.9857