

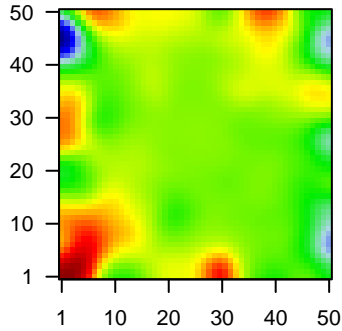
MPI-246

Global Summary

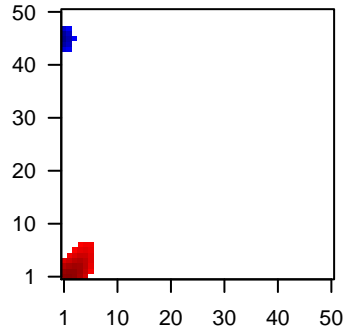
%DE = 0.06
 # genes with $fdr < 0.2$ = 863 (411 + / 452 -)
 # genes with $fdr < 0.1$ = 565 (256 + / 309 -)
 # genes with $fdr < 0.05$ = 540 (244 + / 296 -)
 # genes with $fdr < 0.01$ = 302 (123 + / 179 -)
 # genes in genesets = 13152

<FC> = 0
 <t-score> = 0.06
 <p-value> = 0.22
 <fdr> = 0.94

Portrait



Regulated Metagenes



Global Genelist

Rank	ID	log(FC)	fdr	p-value	Description
1	204591_at	1.97	2e-16	3e-13	17 x 7 cell adhesion molecule L1 like [Source:HGNC Symbol;Acc:HGNC:12691]
2	208621_s_at	-1.61	2e-16	3e-13	0 x 48 ezrin [Source:HGNC Symbol;Acc:HGNC:12691]
3	208930_s_at	-1.65	2e-16	3e-13	1 x 45 interleukin enhancer binding factor 3 [Source:HGNC Symbol;Acc:HGNC:12691]
4	209038_s_at	-1.3	2e-16	3e-13	0 x 42 EH domain containing 1 [Source:HGNC Symbol;Acc:HGNC:12691]
5	209173_at	2.12	2e-16	3e-13	17 x 13 anterior gradient 2, protein disulphide isomerase family memt
6	209754_s_at	-1.51	2e-16	3e-13	2 x 46 thymopoietin [Source:HGNC Symbol;Acc:HGNC:11875]
7	210258_at	-2.47	2e-16	3e-13	49 x 40 regulator of G protein signaling 13 [Source:HGNC Symbol;Acc:HGNC:12691]
8	210763_x_at	2.31	2e-16	3e-13	29 x 0 natural cytotoxicity triggering receptor 3 [Source:HGNC Synt
9	211583_x_at	2.9	2e-16	3e-13	20 x 3 natural cytotoxicity triggering receptor 3 [Source:HGNC Synt
10	212016_s_at	-1.33	2e-16	3e-13	0 x 44 polypyrimidine tract binding protein 1 [Source:HGNC Symbol;Acc:HGNC:12691]
11	212589_at	-1.75	2e-16	3e-13	0 x 7 RAS related 2 [Source:HGNC Symbol;Acc:HGNC:17271]
12	213606_s_at	-1.83	2e-16	3e-13	0 x 45 Rho GDP dissociation inhibitor alpha [Source:HGNC Symbol;Acc:HGNC:12691]
13	215780_s_at	-1.04	2e-16	3e-13	1 x 43 SET pseudogene 4 [Source:HGNC Symbol;Acc:HGNC:4292]
14	218302_at	-1.35	2e-16	3e-13	1 x 42 presenilin enhancer gamma-secretase subunit [Source:HGNC
15	219517_at	-1.63	2e-16	3e-13	49 x 41
16	219518_s_at	-1.92	2e-16	3e-13	49 x 41
17	221234_s_at	-1.89	2e-16	3e-13	45 x 49 BTB domain and CNC homolog 2 [Source:HGNC Symbol;Acc:HGNC:12691]
18	221807_s_at	-1.6	2e-16	3e-13	49 x 48 TraB domain containing [Source:HGNC Symbol;Acc:HGNC:2
19	201102_s_at	-1	4e-16	5e-12	2 x 39 phosphofructokinase, liver type [Source:HGNC Symbol;Acc:HGNC:12691]
20	203795_s_at	-1.37	7e-16	5e-12	42 x 49 BCL tumor suppressor 7A [Source:HGNC Symbol;Acc:HGNC:12691]

Global Geneset Analysis

Rank	GSZ	p-value	#all	Geneset
<i>Overexpressed</i>				
1	12.35	NULL	700	Chr Chr 12
2	8.9	NULL	589	Colon Cancer
3	8.86	NULL	447	Gloma
4	8.18	NULL	72	Reference
5	8.03	NULL	261	GSEA C2
6	7.85	NULL	14	Cancer
7	7.69	NULL	115	Gloma
8	7.64	NULL	231	Gloma
9	7.47	NULL	166	HM
10	7.36	NULL	386	GSEA C2
11	7.32	NULL	1052	GSEA C2
12	7.27	NULL	236	GSEA C2
13	7.25	NULL	317	Cancer
14	7.19	NULL	269	Gloma
15	7.17	NULL	439	GSEA C2
16	7	NULL	431	BP
17	6.99	NULL	265	GSEA C2
18	6.98	NULL	76	HM
19	6.91	NULL	431	GSEA C2
20	6.91	NULL	319	Melanom
<i>Underexpressed</i>				
1	-7.06	NULL	6368	Colon Cancer
2	-6.95	NULL	848	Colon Cancer
3	-6.83	NULL	106	Reference
4	-5.92	NULL	1174	Colon Cancer
5	-5.75	NULL	3150	TF
6	-5.7	NULL	17	Lymphom
7	-5.67	NULL	1764	BP
8	-5.63	NULL	1655	BP
9	-5.43	NULL	3451	TF
10	-5.26	NULL	186	GSEA C2
11	-5.25	NULL	275	GSEA C2
12	-5.21	NULL	18	Lymphom
13	-5.19	NULL	70	CC
14	-5.17	NULL	42	GSEA C2
15	-5.05	NULL	600	GSEA C2
16	-5.04	NULL	3796	TF
17	-5.03	NULL	2946	Chromatin
18	-5.01	NULL	226	GSEA C2
19	-4.88	NULL	227	Lymphom
20	-4.88	NULL	4138	Chromatin

p-values

