

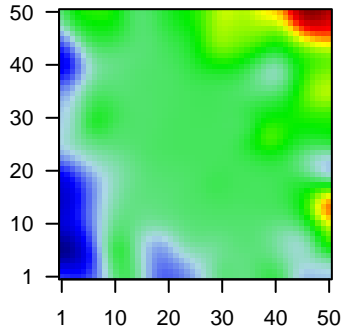
# MPI-095

## Global Summary

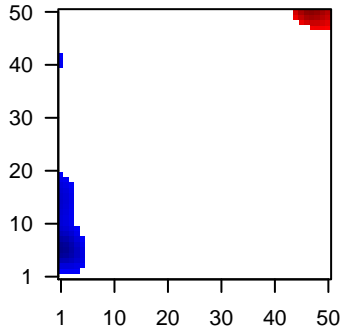
%DE = 0.05  
 # genes with  $fdr < 0.2$  = 548 ( 285 + / 263 - )  
 # genes with  $fdr < 0.1$  = 400 ( 207 + / 193 - )  
 # genes with  $fdr < 0.05$  = 224 ( 110 + / 114 - )  
 # genes with  $fdr < 0.01$  = 158 ( 86 + / 72 - )  
  
 # genes in genesets = 13152

<FC> = 0  
 <t-score> = -0.04  
 <p-value> = 0.27  
 <fdr> = 0.95

### Portrait



### Regulated Metagenes



## Global Genelist

Rank	ID	log(FC)	fdr	p-value	Description
1	212288_at	-1.42	2e-16	2e-12	0 x 5 formin binding protein 1 [Source:HGNC Symbol;Acc:HGNC:1
2	216412_x_at	2.57	2e-16	2e-12	11 x 5
3	204882_at	-1.31	4e-16	4e-11	0 x 6 Rho GTPase activating protein 25 [Source:HGNC Symbol;Acc
4	217022_s_at	-1.57	2e-15	6e-09	0 x 2 immunoglobulin heavy constant alpha 2 (A2m marker) [Sourc
5	217258_x_at	2.16	3e-13	2e-08	11 x 5
6	201946_s_at	0.96	2e-12	2e-08	7 x 48 chaperonin containing TCP1 subunit 2 [Source:HGNC Symbc
7	211813_x_at	-1.7	2e-12	3e-08	8 x 0 decorin [Source:HGNC Symbol;Acc:HGNC:2705]
8	AFFX-r2-Hs1	2.05	4e-12	3e-08	49 x 47
9	216430_x_at	2.03	6e-12	3e-08	29 x 17
10	212660_at	-1.28	6e-12	2e-07	0 x 14 jade family PHD finger 2 [Source:HGNC Symbol;Acc:HGNC:2
11	220423_at	-1.25	2e-11	2e-07	2 x 3 phospholipase A2 group IID [Source:HGNC Symbol;Acc:HGNC
12	215379_x_at	0.89	2e-11	2e-07	41 x 42 immunoglobulin lambda constant 2 [Source:HGNC Symbol;Acc
13	204914_s_at	1.95	4e-11	2e-07	45 x 49 SRY-box 11 [Source:HGNC Symbol;Acc:HGNC:11191]
14	209598_at	1.94	5e-11	2e-07	44 x 46 PNMA family member 2 [Source:HGNC Symbol;Acc:HGNC:9
15	211663_x_at	-0.99	6e-11	3e-07	2 x 3
16	217373_x_at	1.92	7e-11	3e-07	45 x 40 MDM2 proto-oncogene [Source:HGNC Symbol;Acc:HGNC:6
17	219628_at	-1.55	9e-11	2e-06	0 x 10 zinc finger matrix-type 3 [Source:HGNC Symbol;Acc:HGNC:1
18	215121_x_at	0.72	2e-10	2e-06	41 x 42 immunoglobulin lambda constant 2 [Source:HGNC Symbol;Acc
19	219914_at	1.85	4e-10	2e-06	23 x 13 endothelin converting enzyme like 1 [Source:HGNC Symbol;A
20	217227_x_at	1.85	4e-10	2e-06	10 x 5

## Global Geneset Analysis

Rank	GSZ	p-value	#all	Geneset
<i>Overexpressed</i>				
1	15.58	NULL	319	Melanoma
2	14.78	NULL	669	GSEA C2JOHNSTONE_PARVB_TARGETS_3_DN
3	14.22	NULL	726	GSEA C2PUJANA_CHEK2_PCC_NETWORK
4	14.02	NULL	99	Lymphoma
5	13.43	NULL	966	GSEA C2KINSEY_TARGETS_OF_EWSR1_FLII_FUSION_UP
6	13.34	NULL	42	GSEA C2HUMMEL_BURKITT'S_LYMPHOMA_UP
7	12.8	NULL	526	GSEA C2MARSON_BOUND_BY_E2F4_UNSTIMULATED
8	12.63	NULL	192	Lymphoma
9	12.5	NULL	1052	GSEA C2DODD_NASOPHARYNGEAL_CARCINOMA_DN
10	12.11	NULL	439	GSEA C2SHEDDEN_LUNG_CANCER_POOR_SURVIVAL_A6
11	11.87	NULL	431	GSEA C2SOBERT_OLIGODENDROCYTE_DIFFERENTIATION_UP
12	11.82	NULL	400	GSEA C2PUJANA_BRCA2_PCC_NETWORK
13	11.73	NULL	575	GSEA C2CAIRO_HEPATOBLASTOMA_CLASSES_UP
14	11.66	NULL	1527	GSEA C2PUJANA_BRCA1_PCC_NETWORK
15	11.66	NULL	294	GSEA C2WONG_EMBRYONIC_STEM_CELL_CORE
16	11.44	NULL	321	GSEA C2BLUM_RESPONSE_TO_SALIRASIB_DN
17	11.26	NULL	347	GSEA C2REACTOME_CELL_CYCLE
18	11.26	NULL	703	GSEA C2LEE_BMP2_TARGETS_DN
19	11.22	NULL	244	GSEA C2KOBAYASHI_EGFR_SIGNALING_24HR_DN
20	11	NULL	509	GSEA C2RODRIGUES_THYROID_CARCINOMA_POORLY_DIFFERENTIATED
<i>Underexpressed</i>				
1	-16.93	NULL	317	Cancer SPANG_BCL6-index2
2	-15.2	NULL	85	Lymphoma
3	-13.03	NULL	3223	Chromatin
4	-11.27	NULL	2380	Chromatin
5	-11.11	NULL	447	Glioma
6	-10.95	NULL	3682	Chromatin
7	-10.87	NULL	5404	Lymphoma
8	-10.69	NULL	90	GSEA C2BASSO_CD40_SIGNALING_UP
9	-10.64	NULL	186	Cancer SPANG_LPS-index2
10	-10.64	NULL	3767	Chromatin
11	-10.5	NULL	2203	Chromatin
12	-10.23	NULL	6791	Chromatin
13	-10.2	NULL	14	GSEA C2HUMMEL_BURKITT'S_LYMPHOMA_DN
14	-10.15	NULL	1930	Chromatin
15	-10.11	NULL	4079	Chromatin
16	-10	NULL	432	Chromatin
17	-9.97	NULL	7078	Chromatin
18	-9.92	NULL	3838	Chromatin
19	-9.57	NULL	12	Lymphoma
20	-9.38	NULL	3270	CC

### p-values

