

# MSC1 vs MSC2

## Global Summary

%DE = 0.16  
 # genes with  $fdr < 0.2$  = 1846 ( 1002 + / 844 - )  
 # genes with  $fdr < 0.1$  = 1353 ( 812 + / 541 - )  
 # genes with  $fdr < 0.05$  = 1091 ( 692 + / 399 - )  
 # genes with  $fdr < 0.01$  = 694 ( 502 + / 192 - )  
 # genes in genesets = 14839

<FC> = 0  
 <shrinkage-t> = 0  
 <p-value> = 0.1  
 <fdr> = 0.84

## Global Genelist

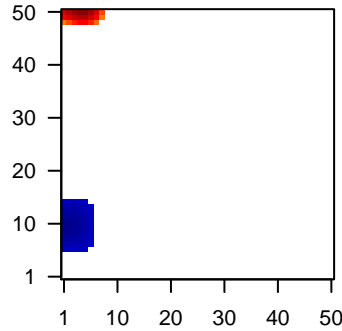
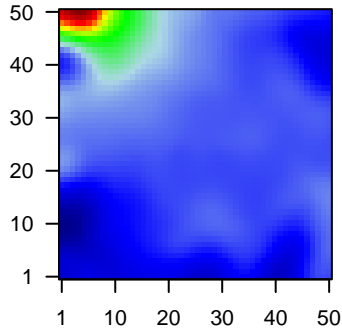
Rank	ID	log(FC)	fdr	p-value	Description
1	ACTL6A	1	2e-16	3e-14	4 x 50 actin-like 6A [Source:HGNC Symbol;Acc:HGNC:24124]
2	ANLN	1.04	2e-16	3e-14	5 x 50 anillin, actin binding protein [Source:HGNC Symbol;Acc:HGNC:24124]
3	ASF1B	1.5	2e-16	3e-14	4 x 50 anti-silencing function 1B histone chaperone [Source:HGNC Symbol;Acc:HGNC:24124]
4	ASPM	0.91	2e-16	3e-14	6 x 50 asp (abnormal spindle) homolog, microcephaly associated (D)
5	ATAD2	0.78	2e-16	3e-14	2 x 50 ATPase family, AAA domain containing 2 [Source:HGNC Symbol;Acc:HGNC:24124]
6	AURKB	1	2e-16	3e-14	6 x 50 aurora kinase B [Source:HGNC Symbol;Acc:HGNC:11390]
7	BARD1	0.81	2e-16	3e-14	2 x 50 BRCA1 associated RING domain 1 [Source:HGNC Symbol;Acc:HGNC:24124]
8	BIRC5	0.89	2e-16	3e-14	6 x 50 baculoviral IAP repeat containing 5 [Source:HGNC Symbol;Acc:HGNC:24124]
9	BLM	0.77	2e-16	3e-14	4 x 50 Bloom syndrome, RecQ helicase-like [Source:HGNC Symbol;Acc:HGNC:24124]
10	BLMH	0.83	2e-16	3e-14	11 x 50 bleomycin hydrolase [Source:HGNC Symbol;Acc:HGNC:1056]
11	BRC A2	0.72	2e-16	3e-14	2 x 50 breast cancer 2, early onset [Source:HGNC Symbol;Acc:HGNC:24124]
12	BUB1	0.96	2e-16	3e-14	6 x 50 BUB1 mitotic checkpoint serine/threonine kinase [Source:HGNC Symbol;Acc:HGNC:24124]
13	BUB1B	1.19	2e-16	3e-14	6 x 50 BUB1 mitotic checkpoint serine/threonine kinase B [Source:HGNC Symbol;Acc:HGNC:24124]
14	CASC5	0.82	2e-16	3e-14	6 x 50 cancer susceptibility candidate 5 [Source:HGNC Symbol;Acc:HGNC:24124]
15	CCDC150	0.83	2e-16	3e-14	5 x 50 coiled-coil domain containing 150 [Source:HGNC Symbol;Acc:HGNC:24124]
16	CCNA2	0.81	2e-16	3e-14	6 x 50 cyclin A2 [Source:HGNC Symbol;Acc:HGNC:1578]
17	CCNB2	0.96	2e-16	3e-14	6 x 50 cyclin B2 [Source:HGNC Symbol;Acc:HGNC:1580]
18	CCNE2	0.87	2e-16	3e-14	2 x 50 cyclin E2 [Source:HGNC Symbol;Acc:HGNC:1590]
19	CDC20	1.02	2e-16	3e-14	6 x 50 cell division cycle 20 [Source:HGNC Symbol;Acc:HGNC:1722]
20	CDC45	0.86	2e-16	3e-14	1 x 50 cell division cycle 45 [Source:HGNC Symbol;Acc:HGNC:1736]

## Global Geneset Analysis

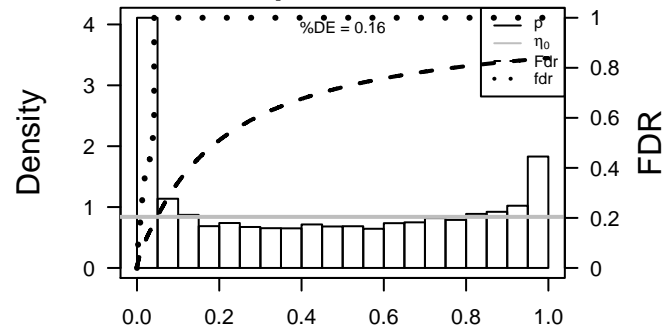
Rank	GSZ	p-value	#all	Geneset
<i>Overexpressed</i>				
1	48.04	0	142	Glio WILLSCHER_GBM_Verhaak-CL_up ( C )
2	46.18	0	550	GSEA C2SOBERT_OLIGODENDROCYTE_DIFFERENTIATION_UP
3	45.47	0	305	GSEA C2DUTERTRE ESTRADIOL_RESPONSE_24HR_UP
4	44.46	0	242	GSEA C2KOBAYASHI_EGFR_SIGNALING_24HR_DN
5	42.64	0	139	GSEA C2ROSTY_CERVICAL_CANCER_PROLIFERATION_CLUSTER
6	38.78	0	145	GSEA C2CHANG_CYCLING_GENES
7	37.47	0	16	Cancer SOTIRIOU_BREAST_CANCER_GRADE_1_VS_3_UP
8	37.26	0	197	HM HALLMARK_E2F_TARGETS
9	36.31	0	700	GSEA C2MARSON_BOUND_BY_E2F4_UNSTIMULATED
10	36.05	0	267	GSEA C2ZHANG_TLX_TARGETS_60HR_DN
11	35.42	0	1192	GSEA C2KINSEY_TARGETS_OF_EWSR1_FLII_FUSION_UP
12	35.28	0	93	GSEA C2KONG_E2F3_TARGETS
13	35.26	0	96	GSEA C2CROONQUIST_IL6_DEPRIVATION_DN
14	35	0	81	GSEA C2GRAHAM_NORMAL QUIESCENT_VS_NORMAL_DIVIDING_DN
15	34.48	0	124	GSEA C2ZHOU_CELL_CYCLE_GENES_IN_IR_RESPONSE_24HR
16	34.26	0	99	GSEA C2BURTON_ADIPOGENESIS_3
17	34.23	0	54	GSEA C2KANG_DOXORUBICIN_RESISTANCE_UP
18	34.07	0	436	GSEA C2SHEDDEN_LUNG_CANCER_POOR_SURVIVAL_A6
19	33.62	0	110	GSEA C2WHITEFORD_PEDIATRIC_CANCER_MARKERS
20	33.62	0	155	GSEA C2HOFFMANN_LARGE_TO_SMALL_PRE_BII_LYMPHOCYTE_UP
<i>Underexpressed</i>				
1	-7.26	2e-04	2984	CC integral component of membrane
2	-6.53	4e-04	3396	Lymphoid OPP_Repressed
3	-6.39	5e-04	2798	Colon Ca56inhG1_Colon
4	-6.1	5e-04	3081	Brain Mid_Frontal_Lobe_ZNF
5	-5.7	9e-04	5155	Colon Ca56inhWk1_Colon
6	-5.48	1e-03	1949	Brain Fetal_TssF
7	-5.38	1e-03	302	GSEA C2KINSEY_TARGETS_OF_EWSR1_FLII_FUSION_DN
8	-5.34	1e-03	2185	Brain Fetal_TssA
9	-5.29	1e-03	730	GSEA C2RODRIGUES_THYROID_CARCINOMA_POORLY_DIFFERENTIAL
10	-5.27	1e-03	616	CC endosome
11	-5.2	1e-03	2142	Colon Ca56inhPC_Colon
12	-5.12	1e-03	1889	Colon Ca56inhPCWk_Colon
13	-5.11	1e-03	9653	Colon Ca56inh_Colon
14	-5.07	2e-03	695	MF transmembrane transporter activity
15	-5.05	2e-03	8	GSEA C2TURJANSKI_MAPK7_TARGETS
16	-5.02	2e-03	472	GSEA C2DUTERTRE ESTRADIOL_RESPONSE_24HR_DN
17	-4.97	2e-03	1907	Brain Fetal_TxTrans
18	-4.95	2e-03	3088	CC plasma membrane
19	-4.92	2e-03	2159	Colon Ca56inhPC_Colon
20	-4.84	2e-03	2698	Colon Ca56inh_Colon

Profile

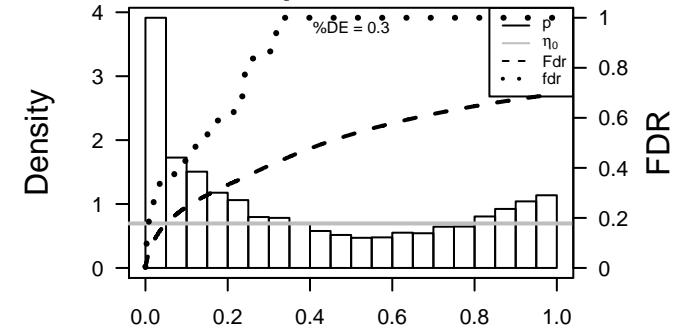
Regulated Spots



p-values



p-values



# MSC1 vs MSC2

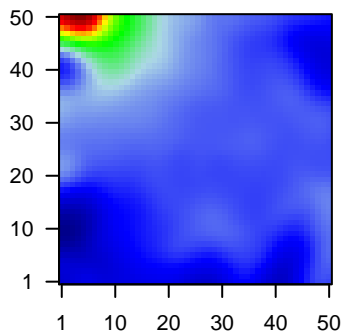
## Local Summary

%DE = 1  
 # metagenes = 23  
 # genes = 352  
 # genes in genesets = 351  
  
 # genes with  $fdr < 0.1 = 350$  ( 350 + / 0 -)  
 # genes with  $fdr < 0.05 = 350$  ( 350 + / 0 -)  
 # genes with  $fdr < 0.01 = 347$  ( 347 + / 0 -)

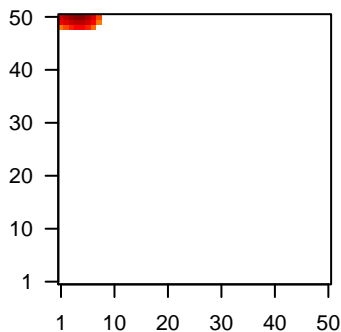
$\langle r \rangle$  metagenes = 0.99  
 $\langle r \rangle$  genes = 0.87

$\langle FC \rangle = 0.63$   
 $\langle \text{shrinkage-t} \rangle = 3.27$   
 $\langle p\text{-value} \rangle = 0$   
 $\langle fdr \rangle = 0.02$

Profile



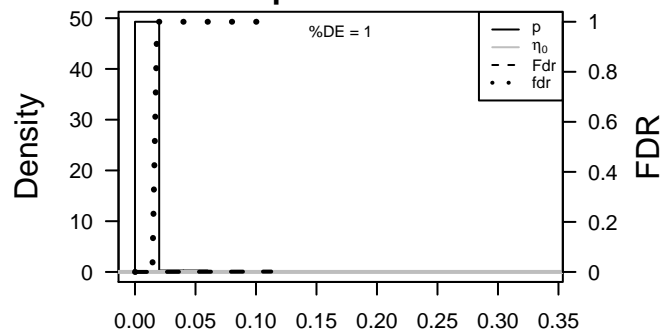
Spot



## Local Genelist

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3	ASF1B	1.5	2e-16	3e-18	4 x 50 anti-silencing function 1B histone chaperone [Source:HGNC Symbol;Acc:HGNC:24124]
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8	BIRC5	0.89	2e-16	3e-18	6 x 50 baculoviral IAP repeat containing 5 [Source:HGNC Symbol;Acc:HGNC:11390]
9	BLM	0.77	2e-16	3e-18	4 x 50 Bloom syndrome, RecQ helicase-like [Source:HGNC Symbol;Acc:HGNC:11390]
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14	CCDC150	0.83	2e-16	3e-18	5 x 50 coiled-coil domain containing 150 [Source:HGNC Symbol;Acc:HGNC:11390]
15	CCNA2	0.81	2e-16	3e-18	6 x 50 cyclin A2 [Source:HGNC Symbol;Acc:HGNC:1578]
16	CCNB2	0.96	2e-16	3e-18	6 x 50 cyclin B2 [Source:HGNC Symbol;Acc:HGNC:1580]
17	CCNE2	0.87	2e-16	3e-18	2 x 50 cyclin E2 [Source:HGNC Symbol;Acc:HGNC:1590]
18	CDC20	1.02	2e-16	3e-18	6 x 50 cell division cycle 20 [Source:HGNC Symbol;Acc:HGNC:1723]
19	CDC45	0.86	2e-16	3e-18	1 x 50 cell division cycle 45 [Source:HGNC Symbol;Acc:HGNC:1735]
20	CDC6	1.19	2e-16	3e-18	1 x 50 cell division cycle 6 [Source:HGNC Symbol;Acc:HGNC:1744]

p-values



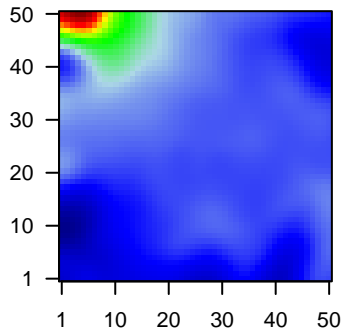
# MSC1 vs MSC2

## Local Summary

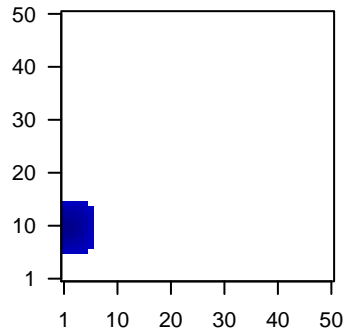
%DE = 0.74  
 # metagenes = 58  
 # genes = 571  
 # genes in genesets = 567  
  
 # genes with  $fdr < 0.1$  = 269 ( 4 + / 265 -)  
 # genes with  $fdr < 0.05$  = 222 ( 1 + / 221 -)  
 # genes with  $fdr < 0.01$  = 137 ( 1 + / 136 -)

<r> metagenes = 1  
 <r> genes = 0.7  
  
 <FC> = -0.17  
 <shrinkage-t> = -0.92  
 <p-value> = 0.01  
 <fdr> = 0.59

Profile



Spot



## Local Genelist

Rank	ID	log(FC)	p-value	fdr	Description
1	KLHL24	-0.72	2e-16	3e-14	1 x 11 kelch-like family member 24 [Source:HGNC Symbol;Acc:HGI
2	TMEM177	-0.67	5e-14	2e-09	3 x 10 transmembrane protein 177 [Source:HGNC Symbol;Acc:HGN
3	SS18L1	-0.56	2e-11	2e-09	1 x 10 synovial sarcoma translocation gene on chromosome 18-like
4	FYCO1	-0.57	3e-11	7e-09	4 x 13 FYVE and coiled-coil domain containing 1 [Source:HGNC Sy
5	MEF2C	-0.52	8e-11	8e-09	5 x 15 myocyte enhancer factor 2C [Source:HGNC Symbol;Acc:HGT
6	MCF2	-0.59	1e-10	8e-09	5 x 10 MCF.2 cell line derived transforming sequence [Source:HGNC
7	MCF2L	-0.57	2e-10	3e-08	1 x 10 MCF.2 cell line derived transforming sequence-like [Source:HGNC
8	KRTAP19-1	-0.59	4e-10	8e-08	1 x 8 keratin associated protein 19-1 [Source:HGNC Symbol;Acc:HGNC:1
9	POPDC2	-0.58	1e-09	8e-08	1 x 9 popeye domain containing 2 [Source:HGNC Symbol;Acc:HGT
10	GREB1	-0.52	1e-09	2e-07	1 x 9 growth regulation by estrogen in breast cancer 1 [Source:HGI
11	CCDC171	-0.43	3e-09	4e-07	1 x 11 coiled-coil domain containing 171 [Source:HGNC Symbol;Acc:HGNC:1
12	MYC	-0.48	5e-09	4e-06	5 x 15 v-myc avian myelocytomatosis viral oncogene homolog [Sou
13	SLC9A1	-0.48	5e-08	4e-06	3 x 10 solute carrier family 9, subfamily A (NHE1, cation proton anti
14	ZNF559	-0.52	7e-08	4e-06	3 x 15 zinc finger protein 559 [Source:HGNC Symbol;Acc:HGNC:28
15	MVB12A	-0.46	9e-08	5e-06	1 x 6 multivesicular body subunit 12A [Source:HGNC Symbol;Acc:HGNC:1
16	SSPN	-0.47	1e-07	5e-06	4 x 13 sarcospan [Source:HGNC Symbol;Acc:HGNC:11322]
17	SNAI2	-0.43	2e-07	6e-06	1 x 11 snail family zinc finger 2 [Source:HGNC Symbol;Acc:HGNC:1
18	PDE3A	-0.45	2e-07	6e-06	2 x 12 phosphodiesterase 3A, cGMP-inhibited [Source:HGNC Synt
19	FOS	-0.37	2e-07	6e-06	4 x 7 FBJ murine osteosarcoma viral oncogene homolog [Source:HGNC
20	HSPB8	-0.39	3e-07	6e-06	1 x 11 heat shock 22kDa protein 8 [Source:HGNC Symbol;Acc:HGNC

p-values

