

8272M

Global Summary

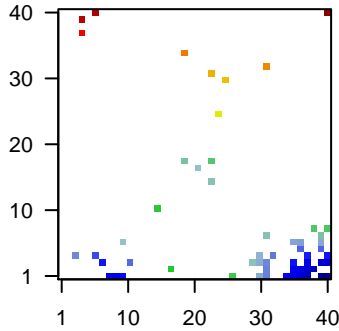
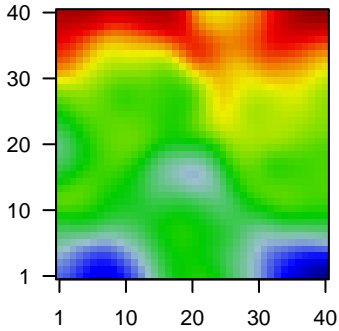
%DE = 0.06
 # genes with $fdr < 0.2$ = 990 (239 + / 751 -)
 # genes with $fdr < 0.1$ = 571 (99 + / 472 -)
 # genes with $fdr < 0.05$ = 384 (55 + / 329 -)
 # genes with $fdr < 0.01$ = 176 (19 + / 157 -)

 # genes in genesets = 16360

<FC> = 0
 <t-score> = -0.18
 <p-value> = 0.28
 <fdr> = 0.94

Portrait

Top 100 DE genes



Global Genelist

Rank	ID	log(FC)	fdr	p-value	Description
1	203797_at	-1.71	2e-16	1e-12	40 x 1 visinin like 1 [Source:HGNC Symbol;Acc:HGNC:12722]
2	203889_at	-2.31	2e-16	1e-12	32 x 4 secretogranin V [Source:HGNC Symbol;Acc:HGNC:10816]
3	203999_at	-2.05	2e-16	1e-12	40 x 1 synaptotagmin 1 [Source:HGNC Symbol;Acc:HGNC:11509]
4	204035_at	-1.85	2e-16	1e-12	31 x 2 secretogranin II [Source:HGNC Symbol;Acc:HGNC:10575]
5	204684_at	-2.16	2e-16	1e-12	40 x 1 neuronal pentraxin 1 [Source:NCBI gene;Acc:4884]
6	205691_at	-2.15	2e-16	1e-12	39 x 1 synaptogyrin 3 [Source:HGNC Symbol;Acc:HGNC:11501]
7	206018_at	-2.31	2e-16	1e-12	30 x 3 forkhead box G1 [Source:HGNC Symbol;Acc:HGNC:3811]
8	218901_at	-1.65	2e-16	1e-12	24 x 25 phospholipid scramblase 4 [Source:HGNC Symbol;Acc:HGNC:11509]
9	221805_at	-2.62	2e-16	1e-12	40 x 1 neurofilament light [Source:HGNC Symbol;Acc:HGNC:7739]
10	229057_at	-1.76	2e-16	1e-12	35 x 1 sodium voltage-gated channel alpha subunit 2 [Source:HGNC Symbol;Acc:HGNC:11509]
11	239671_at	-2.35	3e-15	1e-10	36 x 1 synaptotagmin 16 [Source:HGNC Symbol;Acc:HGNC:23142]
12	205029_s_at	-1.75	4e-15	1e-09	23 x 18 fatty acid binding protein 7 [Source:HGNC Symbol;Acc:HGNC:11509]
13	205030_at	-1.41	2e-14	2e-09	23 x 18 fatty acid binding protein 7 [Source:HGNC Symbol;Acc:HGNC:11509]
14	227189_at	-1.91	7e-14	2e-09	35 x 2 copine 5 [Source:HGNC Symbol;Acc:HGNC:2318]
15	213714_at	-2.19	8e-14	5e-09	39 x 7 calcium voltage-gated channel auxiliary subunit beta 2 [Source:HGNC Symbol;Acc:HGNC:11509]
16	203998_s_at	-2.03	2e-13	5e-09	40 x 1 synaptotagmin 1 [Source:HGNC Symbol;Acc:HGNC:11509]
17	213245_at	-1.86	3e-13	2e-08	40 x 4 adenylate cyclase 1 [Source:HGNC Symbol;Acc:HGNC:232]
18	1558009_at	-1.74	7e-13	2e-08	9 x 1 solute carrier family 1 member 2 [Source:HGNC Symbol;Acc:HGNC:11509]
19	213609_s_at	-1.38	1e-12	2e-08	31 x 1 seizure related 6 homolog like [Source:HGNC Symbol;Acc:HGNC:11509]
20	219619_at	-1.4	1e-12	3e-07	37 x 4 DIRAS family GTPase 2 [Source:HGNC Symbol;Acc:HGNC:11509]

Global Geneset Analysis

Rank	GSZ	p-value	#all	Geneset
<i>Overexpressed</i>				
1	8.98	NULL	564	BP immune system process
2	8.19	NULL	417	BP innate immune response
3	7.85	NULL	388	BP immune response
4	6.61	NULL	364	BP inflammatory response
5	6	NULL	184	BP defense response to virus
6	5.68	NULL	151	BP defense response to bacterium
7	5.44	NULL	148	BP chemotaxis
8	5.14	NULL	74	BP neutrophil chemotaxis
9	4.96	NULL	121	BP defense response
10	4.85	NULL	460	BP neutrophil degranulation
11	4.82	NULL	65	BP chemokine-mediated signaling pathway
12	4.8	NULL	41	BP negative regulation of viral genome replication
13	4.65	NULL	109	BP response to virus
14	4.61	NULL	222	BP adaptive immune response
15	4.59	NULL	51	BP antimicrobial humoral response
16	4.57	NULL	16	BP positive regulation of macrophage activation
17	4.49	NULL	155	BP regulation of immune response
18	4.45	NULL	115	BP keratinization
19	4.08	NULL	30	BP lipopolysaccharide binding
20	4.02	NULL	23	BP hydrogen peroxide catabolic process
<i>Underexpressed</i>				
1	-13.66	NULL	7387	BP membrane
2	-13.56	NULL	574	BP synapse
3	-10.72	NULL	6202	BP cytoplasm
4	-9.85	NULL	236	BP chemical synaptic transmission
5	-9.47	NULL	240	BP postsynaptic membrane
6	-9.29	NULL	4740	BP cytosol
7	-8.98	NULL	4278	BP plasma membrane
8	-8.44	NULL	505	BP nervous system development
9	-6.99	NULL	1435	BP mitochondrion
10	-6.71	NULL	27	BP glutamate secretion
11	-6.56	NULL	146	BP homophilic cell adhesion via plasma membrane adhesion molecules
12	-6.38	NULL	657	BP calcium ion binding
13	-6.24	NULL	12	BP regulation of postsynaptic neurotransmitter receptor activity
14	-6.16	NULL	16	BP positive regulation of calcium ion-dependent exocytosis
15	-6.15	NULL	28	BP synaptic vesicle exocytosis
16	-6.11	NULL	1242	BP Golgi apparatus
17	-6.03	NULL	33	BP regulation of exocytosis
18	-5.94	NULL	545	BP protein ubiquitination
19	-5.91	NULL	31	BP regulation of NMDA receptor activity
20	-5.89	NULL	51	BP neurotransmitter secretion

