

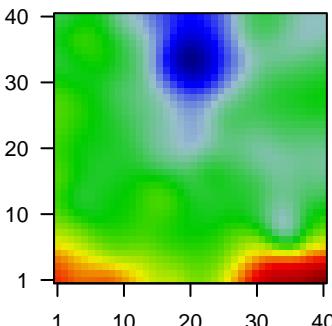
# 3021M

## Global Summary

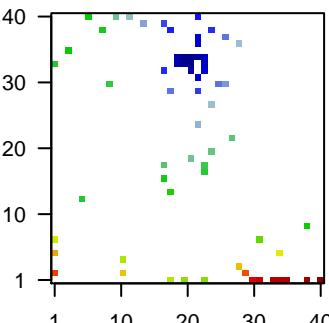
%DE = 0.07  
 # genes with fdr < 0.2 = 2124 ( 1215 + / 909 - )  
 # genes with fdr < 0.1 = 1602 ( 934 + / 668 - )  
 # genes with fdr < 0.05 = 1225 ( 719 + / 506 - )  
 # genes with fdr < 0.01 = 681 ( 388 + / 293 - )  
 # genes in genesets = 16360

$\langle FC \rangle = 0$   
 $\langle t\text{-score} \rangle = 0.1$   
 $\langle p\text{-value} \rangle = 0.23$   
 $\langle fdr \rangle = 0.93$

## Portrait



## Top 100 DE genes



## Global Genelist

Rank	ID	log(FC)	fdr p-value	Metagene	Description		
<b>Overexpressed</b>							
1	1567628_at	-1.46	2e-16	5e-13	20 x 33 CD74 molecule [Source:HGNC Symbol;Acc:HGNC:1697]		
2	200660_at	-1.3	2e-16	5e-13	20 x 34 S100 calcium binding protein A11 [Source:HGNC Symbol;Acc:		
3	201137_s_at	-1.14	2e-16	5e-13	19 x 34 major histocompatibility complex, class II, DP beta 1 [Source:		
4	202295_s_at	-1.31	2e-16	5e-13	23 x 32 cathepsin H [Source:HGNC Symbol;Acc:HGNC:2535]		
5	202376_at	-1.58	2e-16	5e-13	19 x 34 serpin family A member 3 [Source:HGNC Symbol;Acc:HGNC		
6	203240_at	-1.5	2e-16	5e-13	20 x 34 Fc fragment of IgG binding protein [Source:HGNC Symbol;Ac		
7	203973_s_at	-1.02	2e-16	5e-13	22 x 37 CCAAT enhancer binding protein delta [Source:HGNC Symb		
8	204670_x_at	-1.53	2e-16	5e-13	19 x 33 major histocompatibility complex, class II, DR beta 1 [Source:		
9	204990_s_at	-1.25	2e-16	5e-13	22 x 36 integrin subunit beta 4 [Source:HGNC Symbol;Acc:HGNC:61		
10	208306_x_at	-1.16	2e-16	5e-13	19 x 33 major histocompatibility complex, class II, DR beta 1 [Source:		
11	208894_at	-1.04	2e-16	5e-13	20 x 33 major histocompatibility complex, class II, DR alpha [Source:t		
12	209101_at	-1.12	2e-16	5e-13	11 x 4 cellular communication network factor 2 [Source:HGNC Symt		
13	209312_x_at	-1.16	2e-16	5e-13	19 x 33 major histocompatibility complex, class II, DR beta 1 [Source:		
14	209619_at	-0.92	2e-16	5e-13	20 x 34 CD74 molecule [Source:HGNC Symbol;Acc:HGNC:1697]		
15	209703_x_at	-0.99	2e-16	5e-13	27 x 22 methyltransferase like 7A [Source:HGNC Symbol;Acc:HGNC:		
16	210982_s_at	-1.22	2e-16	5e-13	20 x 33 major histocompatibility complex, class II, DR alpha [Source:t		
17	211990_at	-1.14	2e-16	5e-13	19 x 34 major histocompatibility complex, class II, DP alpha 1 [Source:t		
18	214317_x_at	-0.84	2e-16	5e-13	9 x 30 ribosomal protein S9 [Source:HGNC Symbol;Acc:HGNC:104		
19	215193_x_at	-1.36	2e-16	5e-13	19 x 33 major histocompatibility complex, class II, DR beta 3 [Source:		
20	219269_at	-1.04	2e-16	5e-13	14 x 39 homeobox containing 1 [Source:HGNC Symbol;Acc:HGNC:21		
<b>Underexpressed</b>							
1		-18.9	NULL		17	BP	antigen processing and presentation of peptide or polysaccharide a
2		-16.12	NULL		43	BP	antigen processing and presentation
3		-15.97	NULL		564	BP	immune system process
4		-13.87	NULL		388	BP	immune response
5		-10.88	NULL		417	BP	innate immune response
6		-10.58	NULL		460	BP	neutrophil degranulation
7		-9.53	NULL		364	BP	inflammatory response
8		-7.81	NULL		155	BP	regulation of immune response
9		-7.41	NULL		184	BP	defense response to virus
10		-7.12	NULL		289	BP	cytokine-mediated signaling pathway
11		-7.04	NULL		222	BP	adaptive immune response
12		-6.89	NULL		16	BP	negative regulation of growth
13		-6.74	NULL		59	BP	positive regulation of T cell proliferation
14		-6.51	NULL		93	BP	antigen processing and presentation of exogenous peptide antigen
15		-6.46	NULL		31	BP	cellular response to cadmium ion
16		-6.13	NULL		41	BP	positive regulation of interferon-gamma production
17		-6.12	NULL		47	BP	complement activation
18		-6.09	NULL		160	BP	T cell receptor signaling pathway
19		-6.04	NULL		17	BP	cellular response to zinc ion
20		-5.99	NULL		151	BP	defense response to bacterium

## Global Geneset Analysis

