

**Supplementary figure 1: *Nod2*- and *Rip2*-deficient mice showed enhanced susceptibility to DSS-induced colitis.** Single-housed wild-type (SH-WT; black circles, n=5), SH-*Nod2*<sup>-/-</sup> (gray circles, n=5) and SH-*Rip2*<sup>-/-</sup> (blue circles, n=4) mice received *ad libitum* 2% DSS for 7 days followed by 7 days of regular water. **(A)** Survival and **(B)** changes in body weight were evaluated daily and compared to that in SH-WT animals. **(C)** Representative colonoscopy and paraffin-embedded transversal sections of SH-WT, SH-*Nod2*<sup>-/-</sup> and SH-*Rip2*<sup>-/-</sup> mice colons at day 14. **(D)** Colon length and **(E)** histological scoring were evaluated at day 14.

**Supplementary Figure 2: Early enhanced IL-6-mediated response in DSS-treated *Nod2*-deficient mice.** **(A)** Protein levels of IL-6 were determined at day 9 and 14 by specific ELISA. **(B)** Immunohistochemical detection of IL-6 (brown) on colonic sections of DSS-treated WT, *Nod2*<sup>-/-</sup> and *Rip2*<sup>-/-</sup> mutant mice. IL-6 signal is detected in DC-like cells (arrowhead) and in epithelial cells. Scale bar, 50 µm. **(C)** Protein levels of Tnf-α were determined at day 9 and 14 by specific ELISA. **(D)** Relative colonic transcript levels of *Reg3β*, *Reg3γ*, *Spp1*, *Ptgs2*, *Il10*, *Mmp13*, *Tnfα* and *Il6* were determined by qRT-PCR analysis at day 0, 9 and 14. Mean ± SEM, are shown. Asterisk, significant difference P<0.05.

**Supplementary Figure 3: Enhanced IL-6 secretion in DSS-treated *Rip2*-deficient mice failed to co-localize with either CD11b<sup>+</sup>, CD4<sup>+</sup> or CD8a<sup>+</sup> cells.** Immunofluorescent stainings of IL-6 (green) and CD11b, CD4 or CD8a (red) on colonic sections of DSS-treated WT and *Rip2*<sup>-/-</sup> mutant mice. Arrows refer to dendritic-like cells that express IL-6. E, lu and m refer to epithelial cells, lumen and muscularis mucosae respectively. Scale bar, 50 µm.

**Supplementary Figure 4: Colonic response to fecal dysbiosis from *Nod2* deficient mice.** Wild-type animals were either single-housed (n=5) or co-housed at a 1:1 ratio with *Nod2*-deficient mice (n=5) on the same diet for a 4-week period. Microarray experiment was performed once on extracted RNAs from the colon of single-housed and co-housed WT animals **(A)** heat-map, **(B)** functional networking **(C)** and gene ontology of differentially expressed genes as determined by the Ingenuity pathway software. Significance is plotted as -log(p-value). Only biological terms with a significant enrichment are displayed as determined by a |z-score| < -2.

**Supplementary Figure 5: Nod2- and microbiota-independent IL-6 and Tnf- $\alpha$  secretion by the non-diseased colonic mucosa.** **(A)** Relative colonic transcript levels of *Tnf $\alpha$*  and *Il6* in both wild-type and *Nod2*-deficient mice bred under SPF and GF conditions were determined by qRT-PCR analysis. Mean  $\pm$  SEM, are shown. Asterisk, significant difference  $P<0.05$ .

**Supplementary Figure 6: Enhanced morbidity and mortality of *Nod2*- and *Rip2*-deficient mice in an experimental model of colitis-associated colorectal cancer.** Single-housed wild-type (SH-WT, black circles, n=10), SH-*Nod2*<sup>-/-</sup> (grey circles, n=11) and SH-*Rip2*<sup>-/-</sup> (blue circles, n=11) mice were treated with AOM (8 mg/kg). Five days later, mice received *ad libitum* 2% DSS in drinking water for 4 cycles of 5 days separated by 7-day period of regular water. **(A)** Disease activity index and **(B)** survival were monitored daily.

**Supplementary Figure 7: Enhanced IL-6, but not TNF- $\alpha$ , secretion in co-housed wild-type mice at early (but not at later) stage of intestinal tumorigenesis in an**

**experimental model of colitis-associated colorectal cancer.** Protein levels of Tnf- $\alpha$  and IL-6 were determined at day 9 and 34 by specific ELISA. Mean  $\pm$  SEM, are shown. Asterisk, significant difference  $P<0.05$ .

**Supplementary Figure 8: Gene-plus-dysbiosis interaction result in pathological IL-6-mediated response.** Five days after AOM administration (at 12 mg/kg), mice were exposed to 4 cycles of DSS, including 7 days separated by 10-day period of regular water. A combination of antibiotics (Atb) was given (n=4 wild-type mice and n=3 *Nod2*<sup>-/-</sup> mice) or not (n=3 wild-type mice and n=2 *Nod2*<sup>-/-</sup> mice) to animals. **(A)** Venn-diagram analysis. **(B)** Gene ontology. Significance is plotted as -log(p-value). Only top three biological terms with a significant enrichment are displayed as determined by Ingenuity pathway analysis.

**Supplementary Figure 9: Differential response to antibiotics treatment as a result of Nod2 deficiency.** Transcript levels of *Reg3 $\beta$* , *Reg3 $\gamma$* , *Spp1*, *Il6*, *Ptgs2*, *Tnf $\alpha$* , *Il17a*, *Il17f* and *Il22* were determined by qRT-PCR analysis in colons from *Nod2*<sup>+/+</sup> (black circles, n=4), *Nod2*<sup>-/-</sup> (gray circles, n=3), Atb-treated WT (black circles with green border, n=4) and Atb-treated *Nod2*<sup>-/-</sup> (gray circles with green border, n=3) at day 16. Mean  $\pm$  SEM, are shown.

**Supplementary Figure 10: Anti-IL6R treatment influences Nod2-driven dysbiosis in the DSS-AOM model.** High-throughput 16S rRNA pyrosequencing was performed on chronically inflamed **(A)** and tumoral **(B)** colonic specimens isolated from *Nod2*<sup>+/+</sup> and *Nod2*<sup>-/-</sup> animals (N=4). Clusters representing 20 OTUs which primarily contribute to the shift between *Nod2*<sup>+/+</sup> and *Nod2*<sup>-/-</sup> animals were depicted. Corresponding clusters (left and right) are created based on the same OTUs. Z-score normalized OTU abundances

are color-coded. Row dendograms display OTU abundance similarities while column dendograms display similarities between animals (dark green for *Nod2*<sup>+/+</sup>, light green for *Nod2*<sup>-/-</sup>). Column headers represent animal IDs, which are unique within a given group.

**Supplementary Figure 11: Fecal-associated dysbiosis in *Nod2*-deficient mice is mitigated by reciprocal fecal transplantation.** High-throughput 16S rRNA pyrosequencing was performed on fecal specimens isolated from all gnotobiotic animals upon injury (N=5). **(A)** Non parametric quantification of microbiota diversity. *P*-values were determined by the Mann-Whitney U-test. Mean  $\pm$  SEM, are shown. Asterisk, significant difference *P*<0.05. **(B)** Percentage abundance at the phyla level. **(C)** Clusters representing 20 OTUs which primarily contribute to the shift between the fecal-associated microbiota were depicted (N=5). Corresponding clusters (left and right) are created based on the same OTUs. Z-score normalized OTU abundances are color-coded. Row dendograms display OTU abundance similarities while column dendograms display similarities between animals (dark green for *Nod2*<sup>+/+</sup>, light green for *Nod2*<sup>-/-</sup>). Column headers represent animal IDs, which are unique within a given group.

**Supplementary Figure 12: Mucosal-associated dysbiosis in *Nod2*-deficient mice is mitigated by reciprocal fecal transplantation.** High-throughput 16S rRNA pyrosequencing was performed on colonic specimens isolated from all gnotobiotic animals upon injury (N=5). **(A)** Non parametric quantification of microbiota diversity. *P*-values were determined by the Mann-Whitney U-test. Mean  $\pm$  SEM, are shown. Asterisk, significant difference *P*<0.05. **(B)** Percentage abundance at the phyla level. **(C)** Clusters representing 20 OTUs which primarily contribute to the shift between the

mucosal-associated microbiota were depicted. Corresponding clusters (left and right) are created based on the same OTUs. Z-score normalized OTU abundances are color-coded. Row dendograms display OTU abundance similarities while column dendograms display similarities between animals (dark green for *Nod2*<sup>+/+</sup>, light green for *Nod2*<sup>-/-</sup>). Column headers represent animal IDs, which are unique within a given group.

**Supplementary Table 1 : Microarray analysis on co-housed wild-type animals.**

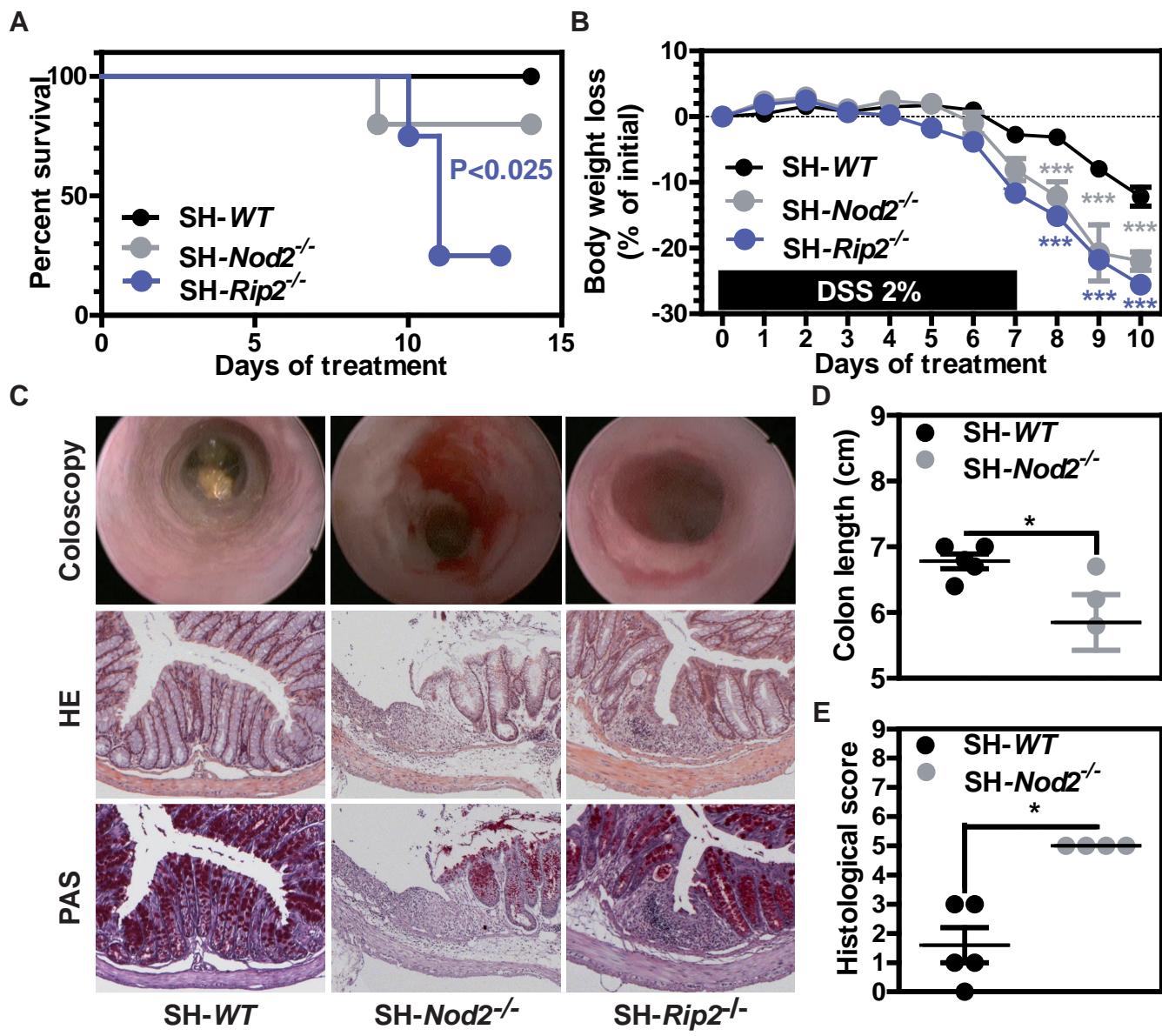
Differentially expressed genes with a *p*-value < 0.01 and an average fold change > 1 were listed.

**Supplementary Table 2 : Gene ontology analysis on differentially expressed genes upon co-housing.** Function annotation was performed by using the Ingenuity pathway software. Regulation z-score predict function activation state.

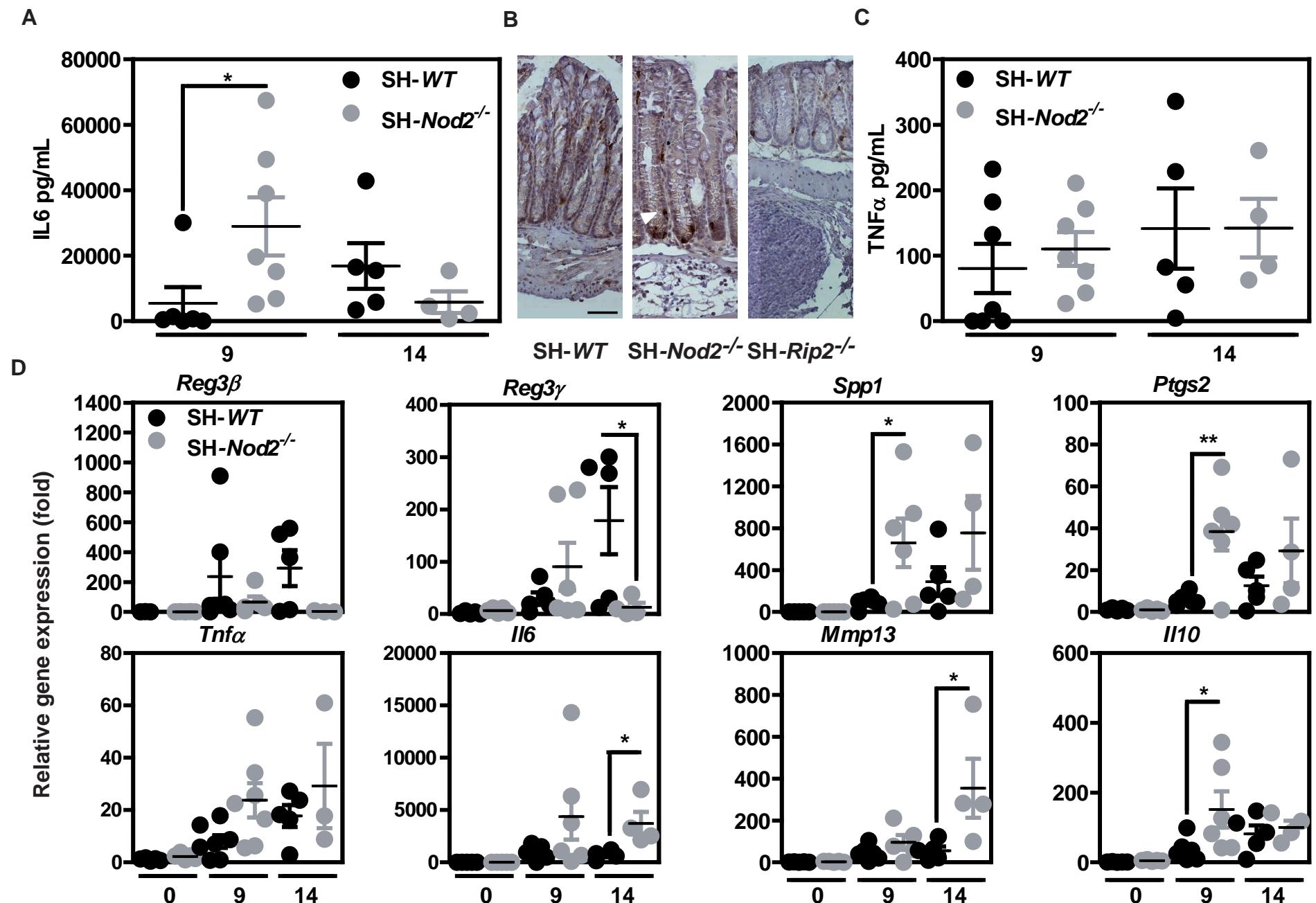
**Supplementary Table 3 : Microarray analysis in DSS/AOM-treated wild-type and *Nod2*-deficient mice according antibiotic treatment.** First and second group described differential responsiveness of wild-type and *Nod2*-deficient mice to antibiotic treatment when compared to untreated wild-type and *Nod2*-deficient animals respectively.

**Supplementary Table 4 : Gene ontology analysis on the differential *Nod2*-dependent response to antibiotics treatment.** Function annotation was performed by using the Ingenuity pathway software.

**Supplementary Table 5 : Primers sequences for qRT-PCR and qPCR analysis**

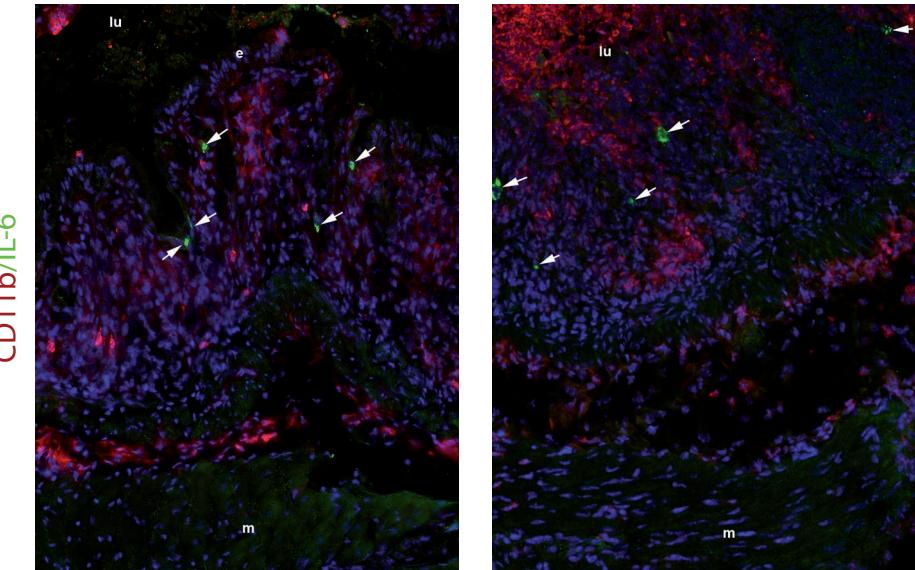


Supplementary figure 1

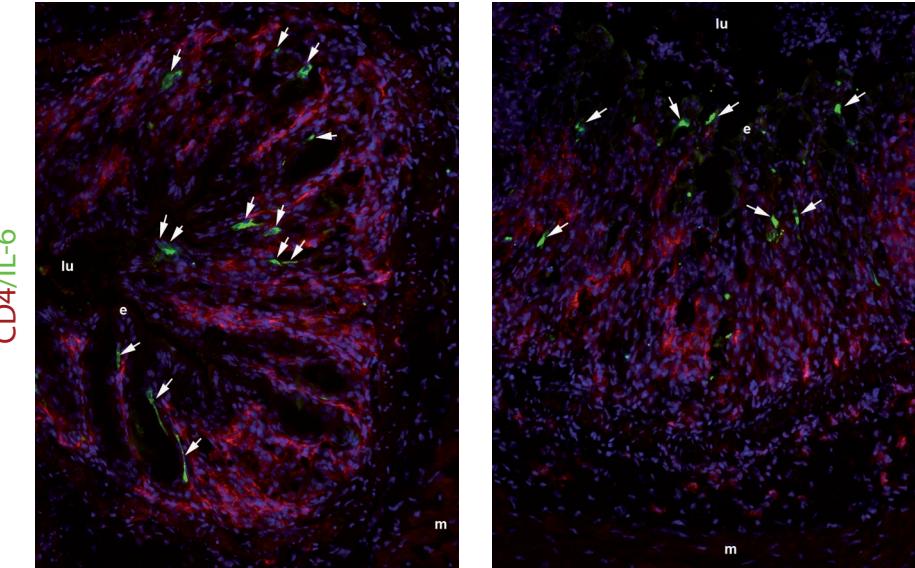


Supplementary figure 2

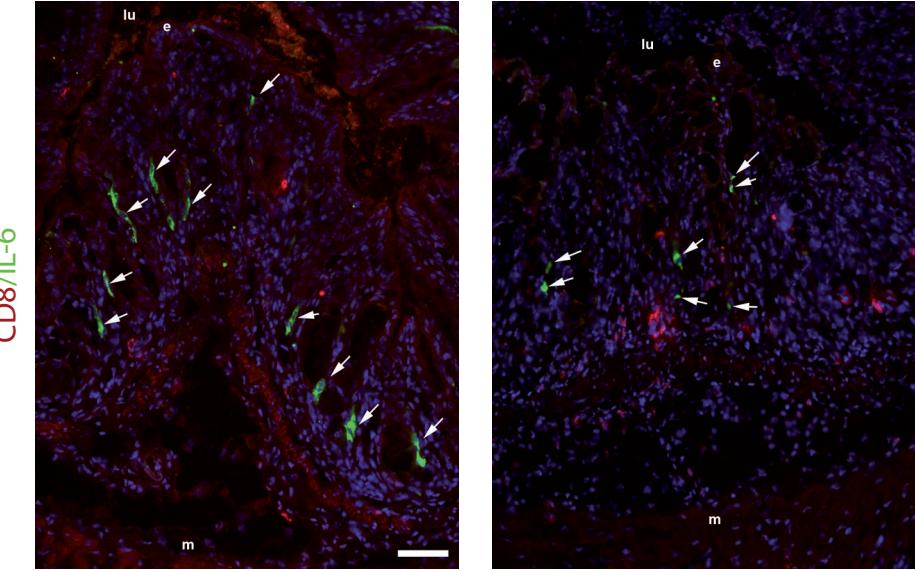
**A** *Rip2<sup>+/+</sup>* *Rip2<sup>-/-</sup>*



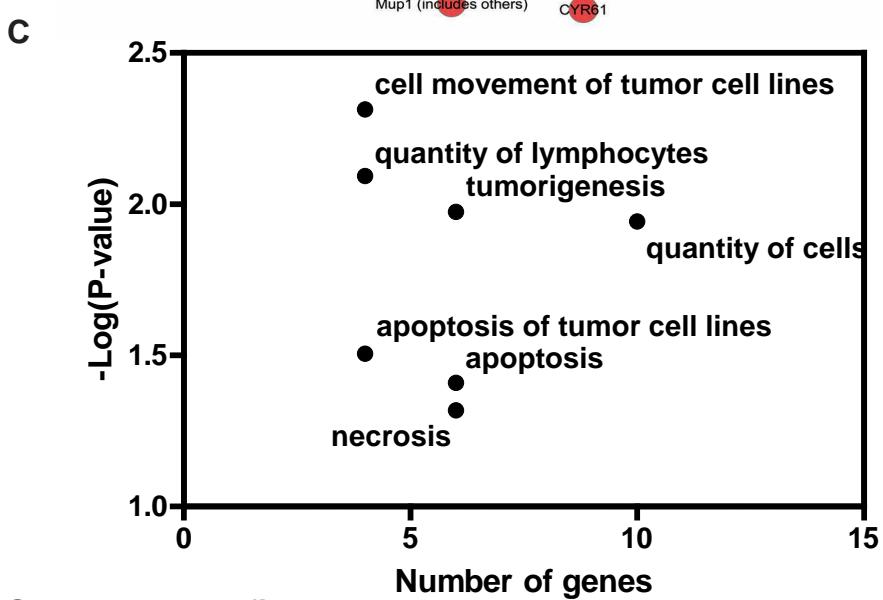
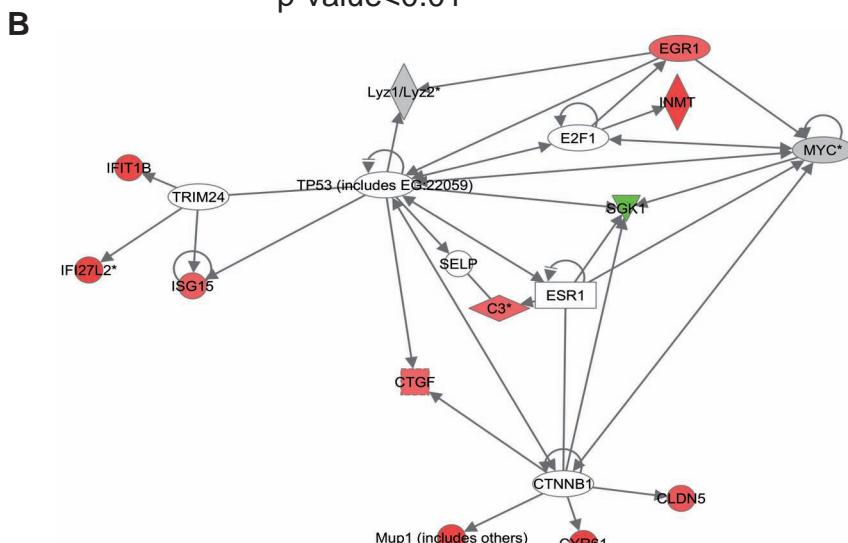
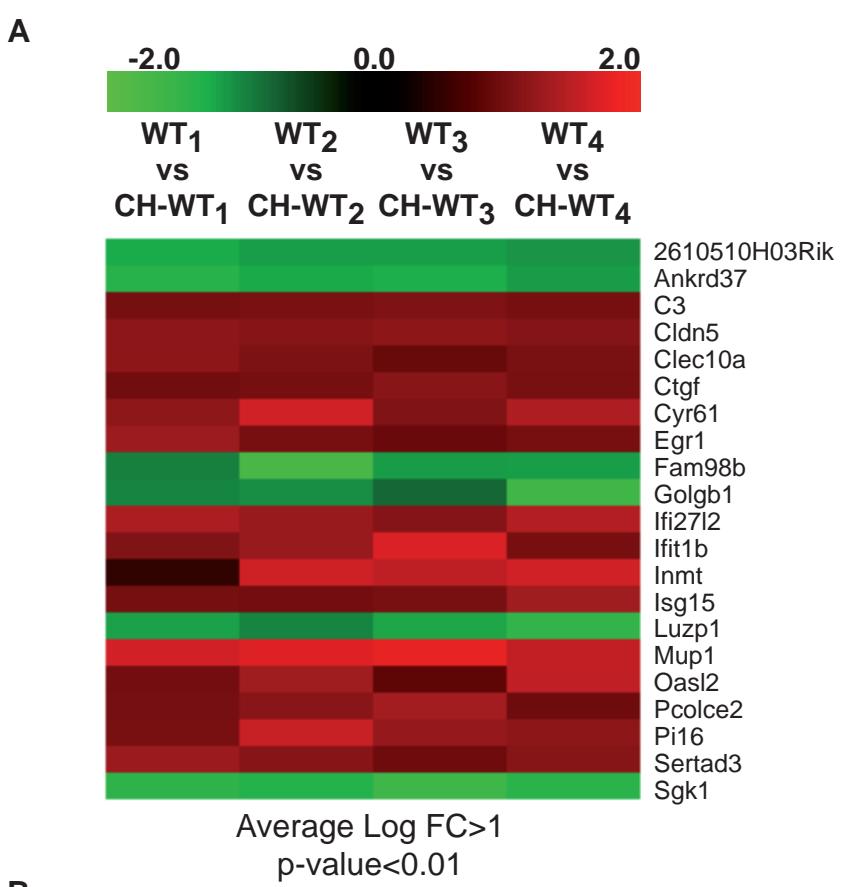
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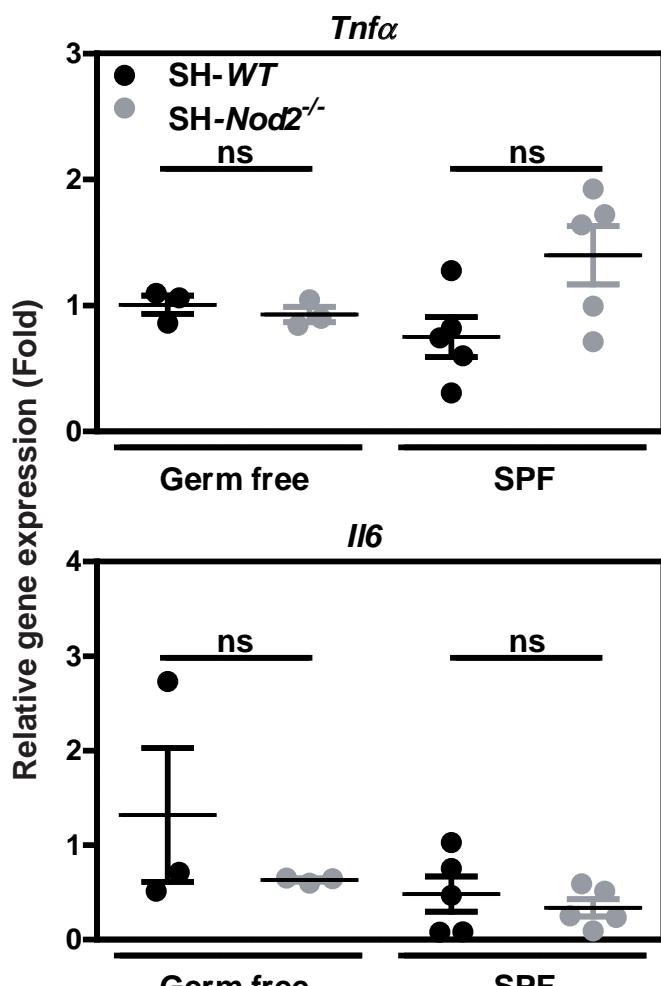
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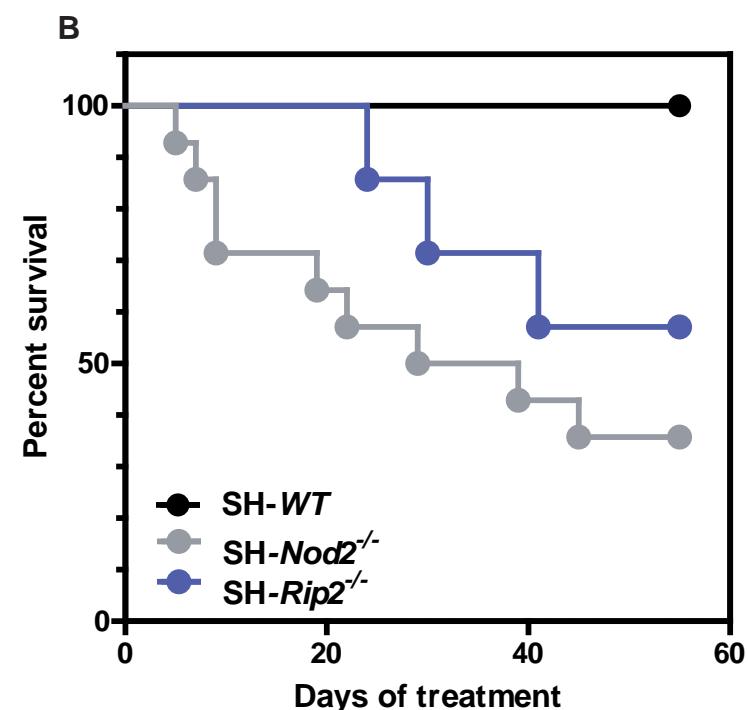
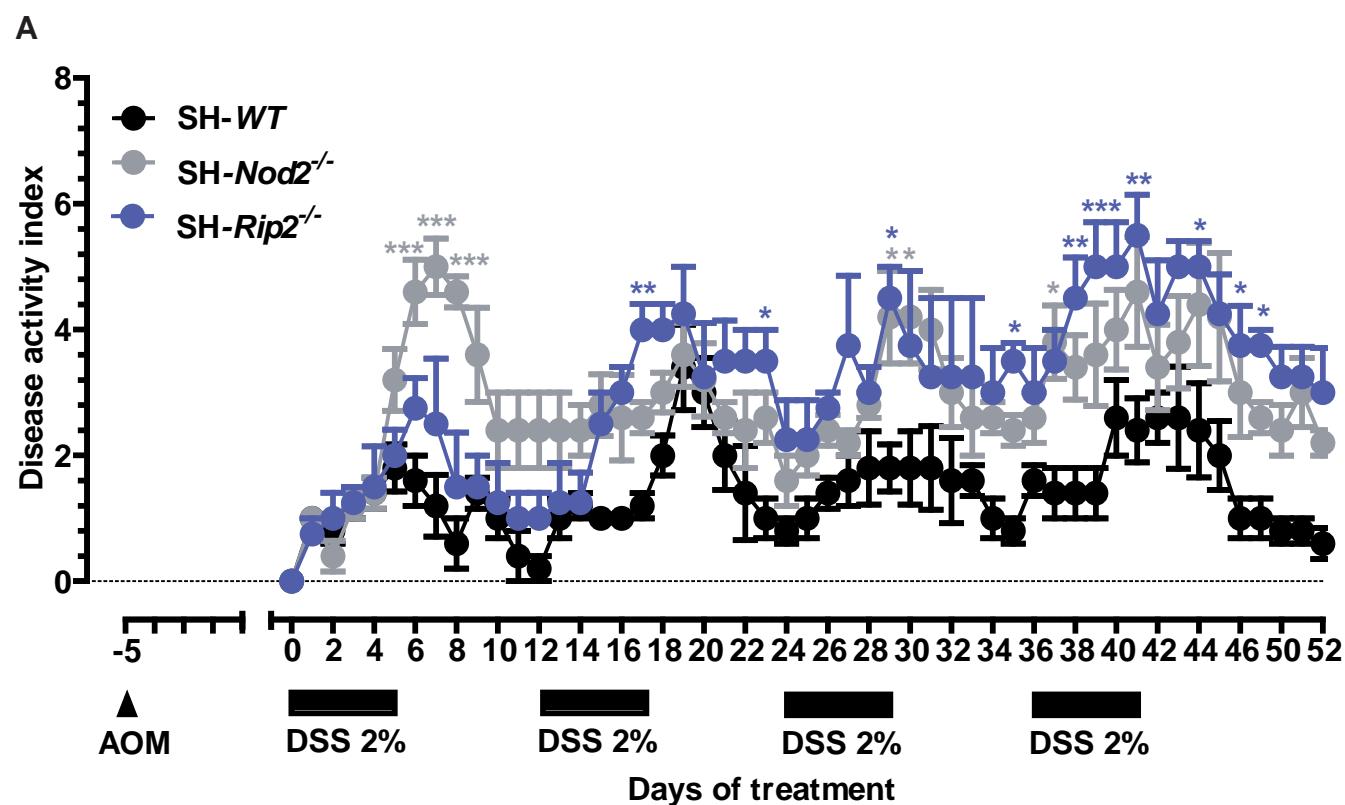
**Supplementary figure 3**



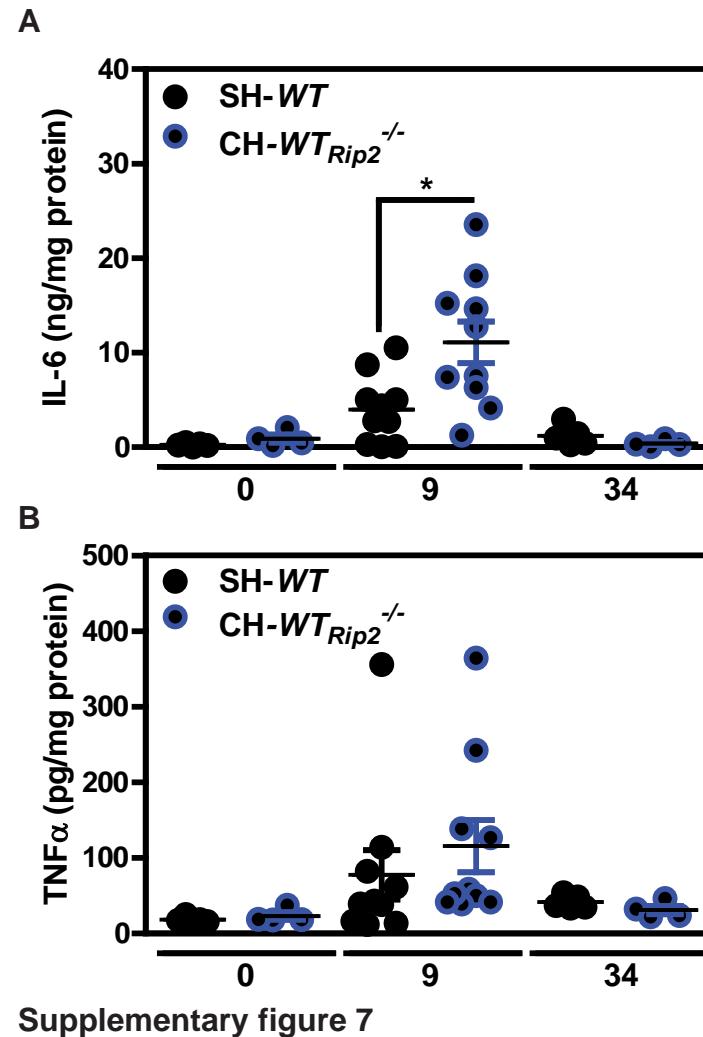
Supplementary figure 4

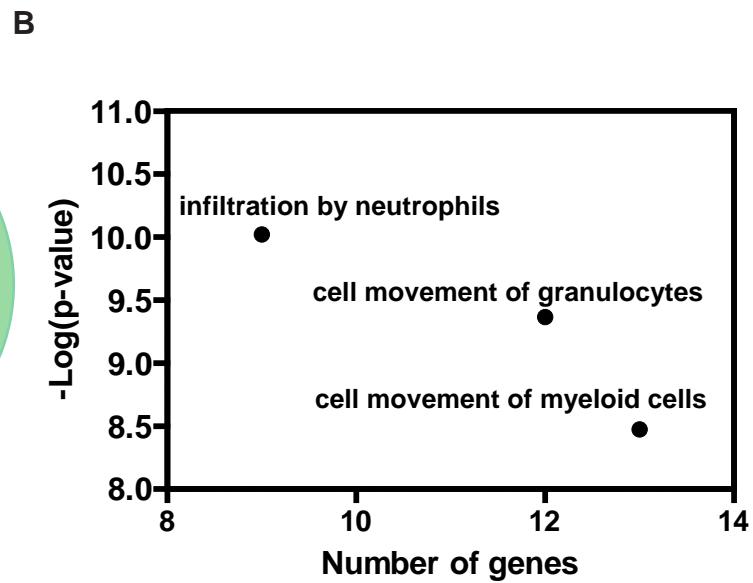


Supplementary figure 5

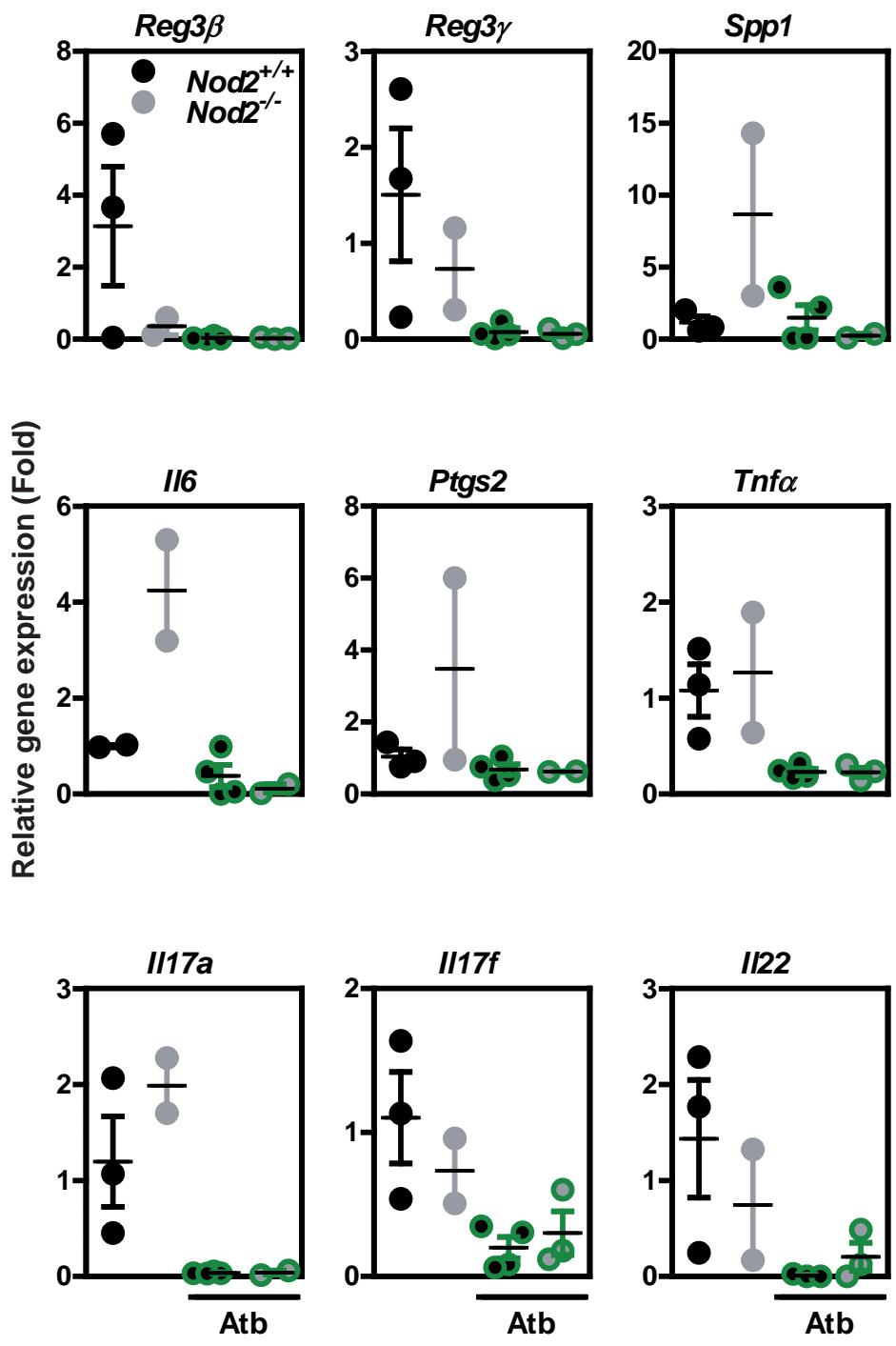


Supplementary figure 6

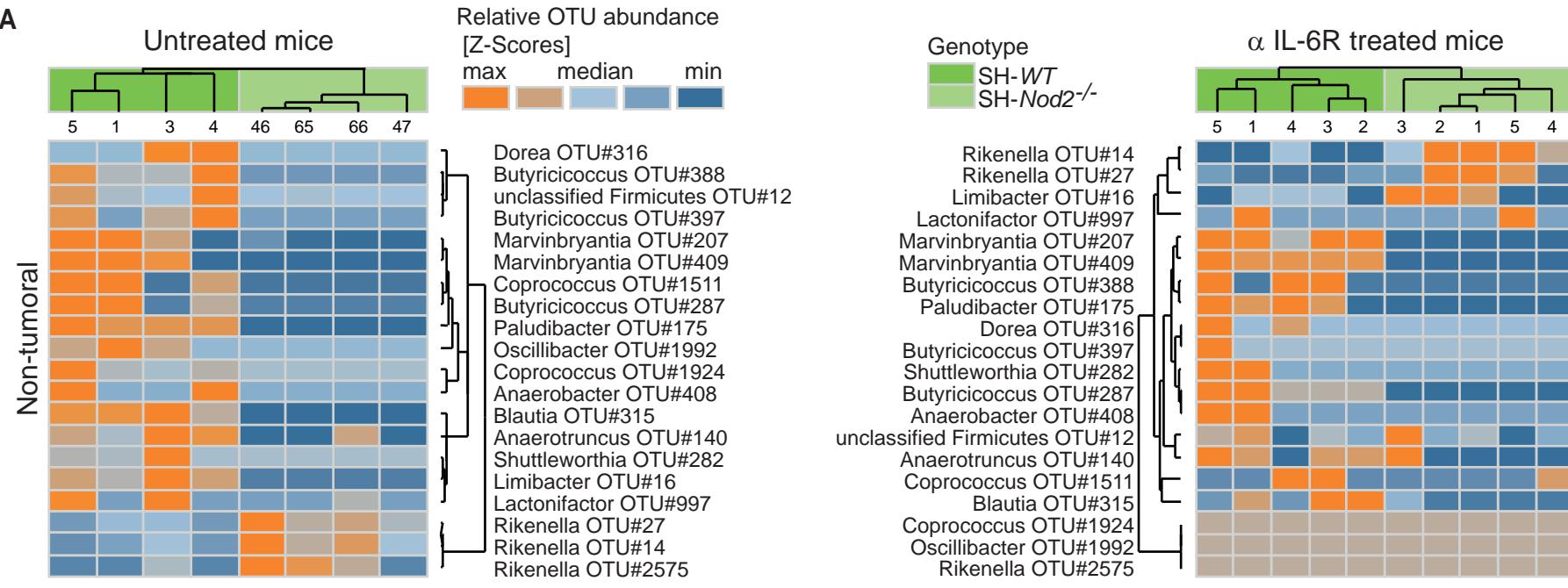
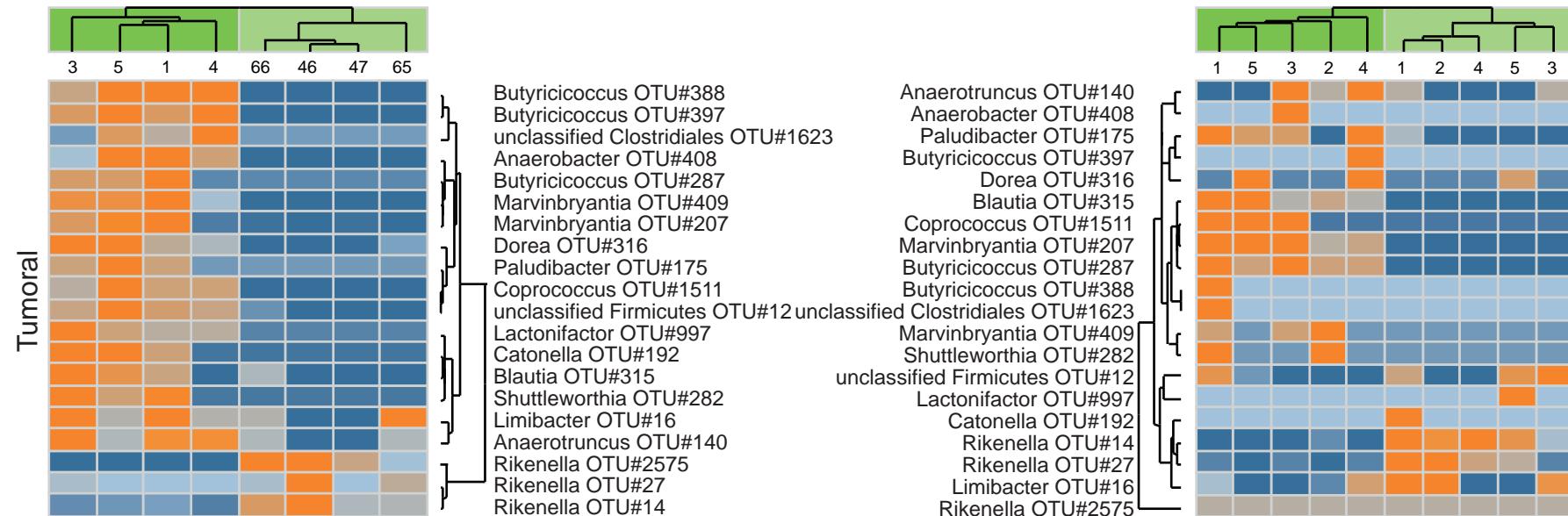


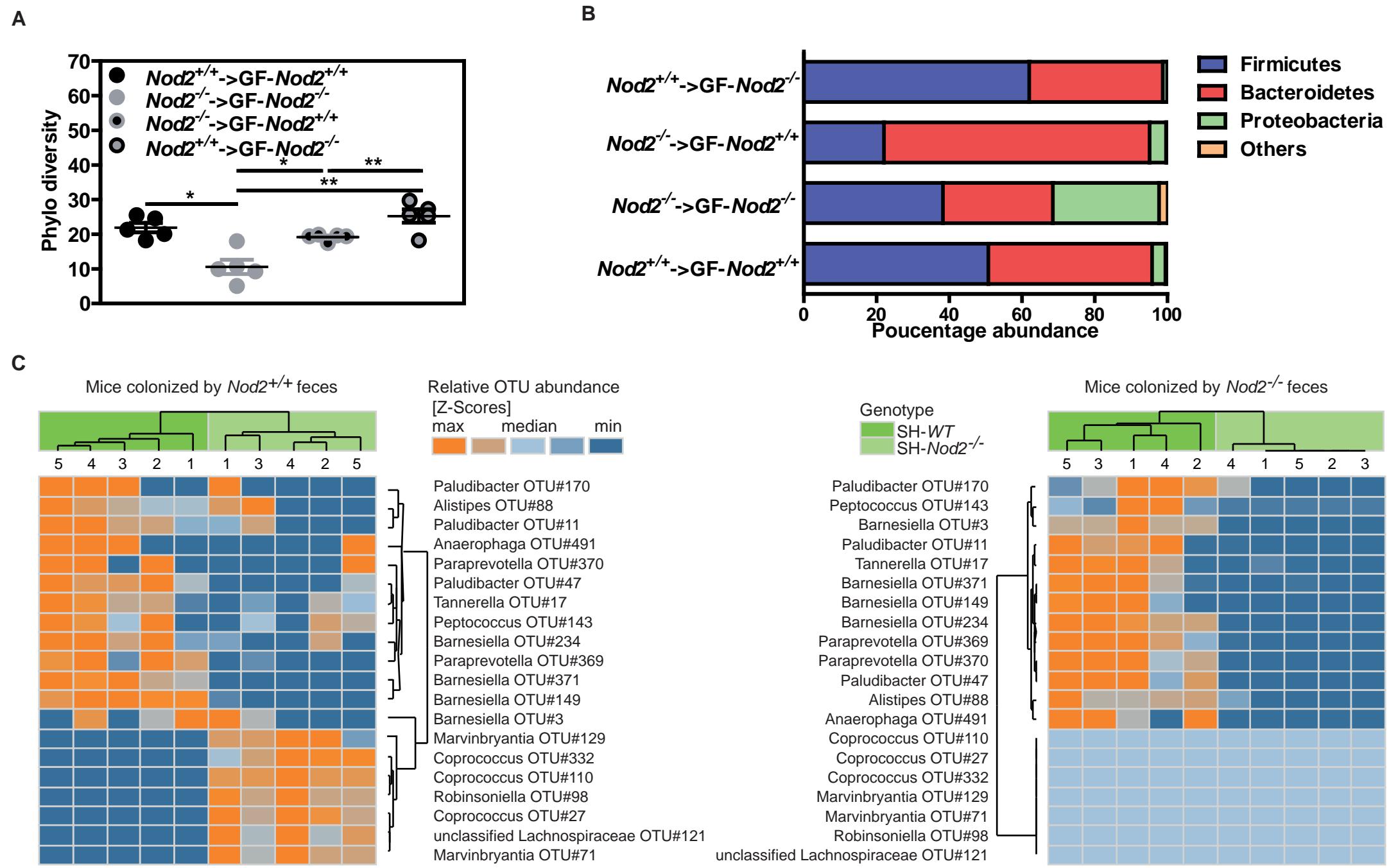


**Supplementary figure 8**



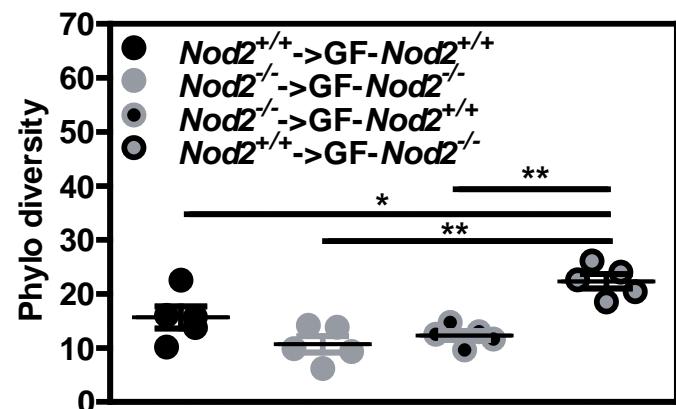
Supplementary Figure 9

**A****B****Supplementary figure 10**

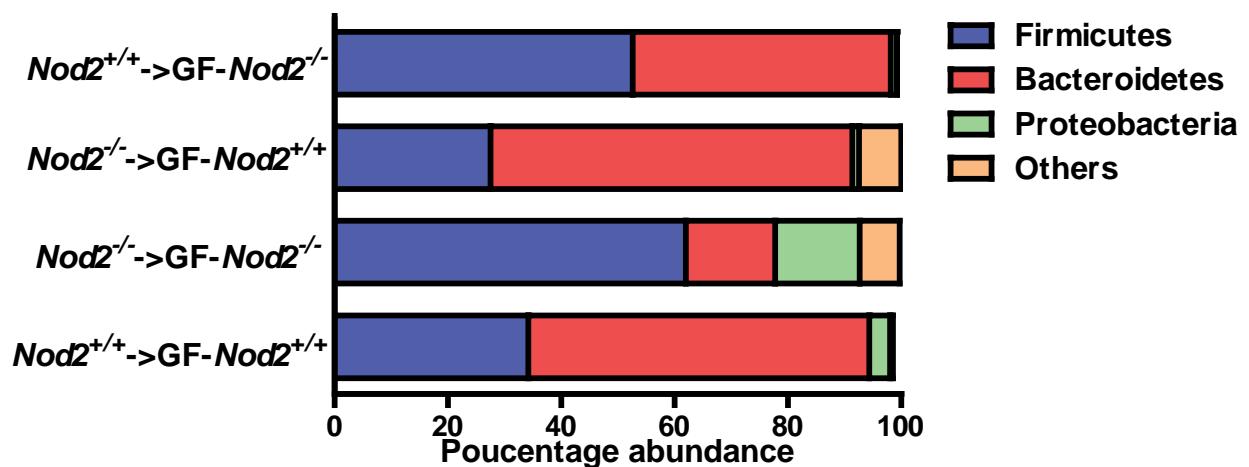


Supplementary figure 11

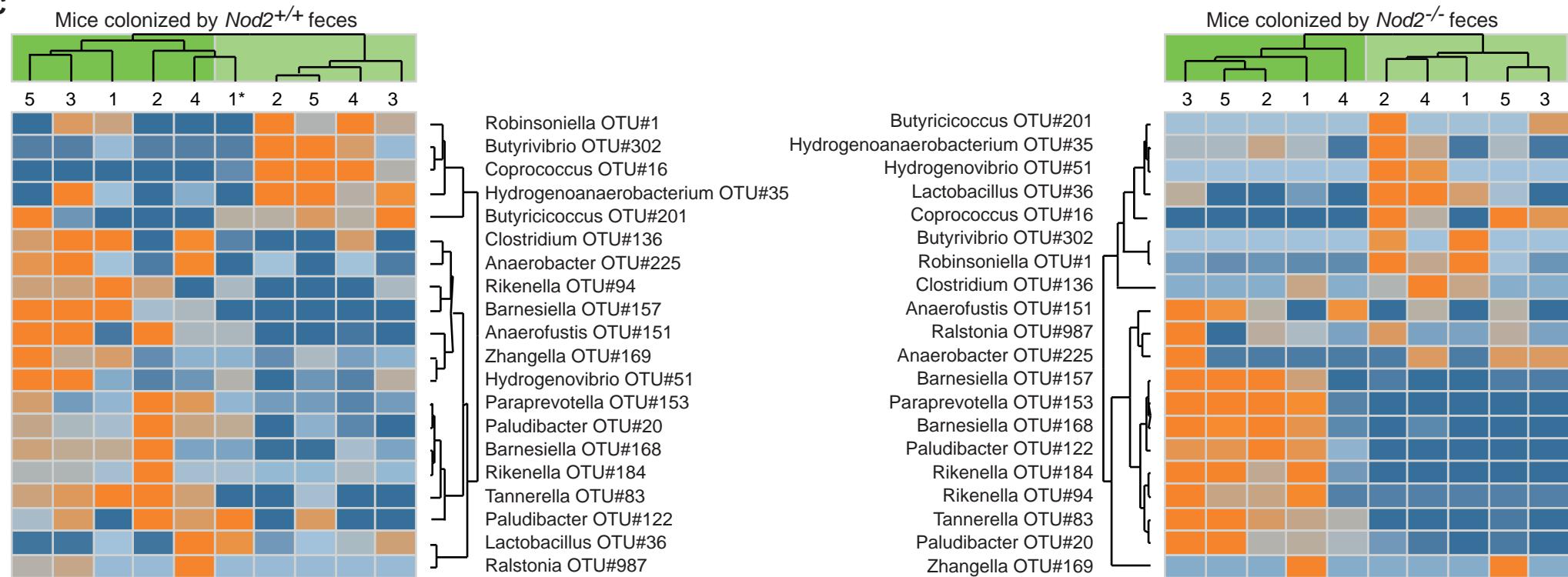
A



B



C



Supplementary figure 12

**Supplementary Table 1: Microarray analysis on co-housed wild-type animals.**

ID	Symbol	Entrez Gene Name	Log Ratio	p-value
A_51_P272066	2010109I03Rik	RIKEN cDNA 2010109I03 gene	-1,147	1,67E-06
A_52_P319438	ANKRD37	ankyrin repeat domain 37	-1,238	2,16E-06
A_51_P110301	C3	complement component 3	1,002	1,32E-06
A_51_P189814	CLDN5	claudin 5	1,131	1,53E-06
A_51_P141413	CLEC10A	C-type lectin domain family 10, member A	1,021	2,20E-04
A_51_P157042	CTGF	connective tissue growth factor	1,011	2,65E-06
A_51_P252859	CYR61	cysteine-rich, angiogenic inducer, 61	1,310	1,91E-05
A_51_P367866	EGR1	early growth response 1	1,040	2,55E-04
A_51_P202040	FAM98B	family with sequence similarity 98, member B	-1,225	1,94E-04
A_52_P675052	GOLGB1	golgin B1	-1,099	4,88E-04
A_52_P90363	IFI27L2	interferon, alpha-inducible protein 27-like 2	1,283	3,72E-06
A_51_P327751	IFIT1B	interferon-induced protein with tetratricopeptide repeats 1B	1,251	4,64E-04
A_51_P162162	INMT	indolethylamine N-methyltransferase	1,305	4,62E-03
A_52_P463936	ISG15	ISG15 ubiquitin-like modifier	1,052	2,55E-05
A_52_P359621	LUZP1	leucine zipper protein 1	-1,195	4,61E-06
A_51_P461703	Mup1	major urinary protein 1	1,663	5,24E-05
A_51_P387123	Oasl2	2'-5' oligoadenylate synthetase-like 2	1,143	1,00E-04
A_51_P245789	PCOLCE2	procollagen C-endopeptidase enhancer 2	1,085	1,50E-05
A_52_P413756	PI16	peptidase inhibitor 16	1,236	1,49E-04
A_52_P71105	SERTAD3	SERTA domain containing 3	1,096	4,79E-06
A_51_P118237	SGK1	serum/glucocorticoid regulated kinase 1	-1,414	1,12E-06

**Supplementary Table 2 : Gene ontology analysis on differentially expressed genes upon co-housing**

Category	Functions	Functions Annotation	p-Value	-log <sub>10</sub> p-Value	Predicted Activation State	Activation z-score	# Molecules
Cancer	tumorigenesis	tumorigenesis	1.14E-02	1.94309515	-1.414	C3,CLDN5,CLEC10A,CTGF,CYR61,EGR1,IFI27L2,IFIT1,ISG15,SGK1	10
Cell Death	apoptosis	apoptosis of tumor cell lines	3.12E-02	1.50584541	-0.482	CTGF,CYR61,EGR1,SGK1	4
Cell Death	necrosis	necrosis	3.90E-02	1.40893539	-0.283	20101090103Rik,C3,CTGF,CYR61,EGR1,SGK1	6
Cell Death	apoptosis	apoptosis	4.81E-02	1.31785492	0.065	20101090103Rik,C3,CTGF,CYR61,EGR1,SGK1	6
Tissue Morphology	quantity	quantity of cells	1.06E-02	1.97469413	0.280	C3,CLEC10A,CTGF,EGR1,ISG15,SGK1	6
Hematological System Development and Function	quantity	quantity of lymphocytes	8.08E-03	2.02958864	1.154	C3,CLEC10A,EGR1,ISG15	4
Tissue Morphology	quantity	quantity of lymphocytes	8.08E-03	2.02958864	1.154	C3,CLEC10A,EGR1,ISG15	4
Cellular Movement	cell movement	cell movement of tumor cell lines	4.86E-03	2.31336373	1.951	C3,CTGF,CYR61,EGR1	4
Cellular Assembly and Organization	quantity	quantity of focal complexes	6.38E-06	2.97E-05	CTGF,CYR61	2	
Cell Morphology	abnormal morphology	abnormal morphology of neutrophils	2.97E-05	CTGF	2		
Hematological System Development and Function	abnormal morphology	abnormal morphology of neutrophils	2.97E-05	CTGF	2		
Inflammatory Response	abnormal morphology	abnormal morphology of neutrophils	2.97E-05	CTGF	2		
Cancer	pituitary gland adenoma	pituitary gland adenoma	3.58E-04	3.38E-04	C3,EGR1,IFIT1B	3	
Cellular Compromise	injury	injury of cells	3.06E-04	3.06E-04	C3,EGR1	2	
Renal and Urological Disease	proteinuria	proteinuria	4.07E-04	3.07E-04	C3,CTGF,SGK1	3	
Cellular Growth and Proliferation	colony formation	colony formation of carcinoma cell lines	5.18E-04	2.91E-04	CYR61,EGR1	2	
Digestive System Development and Function	regeneration	regeneration of liver	5.51E-04	2.83E-04	C3,EGR1	2	
Hepatic System Development and Function	regeneration	regeneration of liver	5.51E-04	2.83E-04	C3,EGR1	2	
Organ Morphology	regeneration	regeneration of liver	5.51E-04	2.83E-04	C3,EGR1	2	
Cancer	benign tumor	benign tumor	1.01E-03	2.01E-03	C3,CTGF,CYR61,EGR1,IFIT1B	5	
Hepatic System Development and Function	adhesion	adhesion of hepatic stellate cells	1.07E-03	2.07E-03	CTGF	1	
Cell-To-Cell Signaling and Interaction	adhesion	adhesion of hepatic stellate cells	1.07E-03	2.07E-03	CTGF	1	
Connective Tissue Development and Function	adhesion	adhesion of hepatic stellate cells	1.07E-03	2.07E-03	CTGF	1	
Tissue Development	adhesion	adhesion of hepatic stellate cells	1.07E-03	2.07E-03	CTGF	1	
Cardiovascular System Development and Function	angiogenesis	angiogenesis of growth plate	1.07E-03	2.07E-03	CTGF	1	
Connective Tissue Development and Function	angiogenesis	angiogenesis of growth plate	1.07E-03	2.07E-03	CTGF	1	
Embryonic Development	angiogenesis	angiogenesis of growth plate	1.07E-03	2.07E-03	CTGF	1	
Organ Development	angiogenesis	angiogenesis of growth plate	1.07E-03	2.07E-03	CTGF	1	
Organismal Development	angiogenesis	angiogenesis of growth plate	1.07E-03	2.07E-03	CTGF	1	
Skeletal and Muscular System Development and Function	angiogenesis	angiogenesis of growth plate	1.07E-03	2.07E-03	CTGF	1	
Tissue Development	angiogenesis	angiogenesis of growth plate	1.07E-03	2.07E-03	CTGF	1	
Cellular Response to Therapeutics	antiestrogen resistance	antiestrogen resistance of breast cancer cell lines	1.07E-03	2.07E-03	CYR61	1	
Drug Metabolism	antiestrogen resistance	antiestrogen resistance of breast cancer cell lines	1.07E-03	2.07E-03	CYR61	1	
Cell Death	cytotoxicity	cytotoxicity of hepatoma cell lines	1.07E-03	2.07E-03	SGK1	1	
Cancer	hyperplasia	delay in hyperplasia of hepatocytes	1.07E-03	2.07E-03	C3	1	
Gastrointestinal Disease	hyperplasia	delay in hyperplasia of hepatocytes	1.07E-03	2.07E-03	C3	1	
Hepatic System Disease	hyperplasia	delay in hyperplasia of hepatocytes	1.07E-03	2.07E-03	C3	1	
Tissue Morphology	removal	delay in removal of parenchyma	1.07E-03	2.07E-03	C3	1	
Cellular Movement	dissemination	dissemination of hepatoma cell lines	1.07E-03	2.07E-03	CTGF	1	
Connective Tissue Development and Function	endochondral ossification	endochondral ossification of growth plate	1.07E-03	2.07E-03	CTGF	1	
Embryonic Development	endochondral ossification	endochondral ossification of growth plate	1.07E-03	2.07E-03	CTGF	1	
Organ Development	endochondral ossification	endochondral ossification of growth plate	1.07E-03	2.07E-03	CTGF	1	
Organismal Development	endochondral ossification	endochondral ossification of growth plate	1.07E-03	2.07E-03	CTGF	1	
Skeletal and Muscular System Development and Function	endochondral ossification	endochondral ossification of growth plate	1.07E-03	2.07E-03	CTGF	1	
Tissue Development	endochondral ossification	endochondral ossification of growth plate	1.07E-03	2.07E-03	CTGF	1	
Hematological System Development and Function	erythropoiesis	erythropoiesis of splenocytes	1.07E-03	2.07E-03	EGR1	1	
Cellular Development	erythropoiesis	erythropoiesis of splenocytes	1.07E-03	2.07E-03	EGR1	1	
Embryonic Development	erythropoiesis	erythropoiesis of splenocytes	1.07E-03	2.07E-03	EGR1	1	
Hematopoiesis	erythropoiesis	erythropoiesis of splenocytes	1.07E-03	2.07E-03	EGR1	1	
Lymphoid Tissue Structure and Development	erythropoiesis	erythropoiesis of splenocytes	1.07E-03	2.07E-03	EGR1	1	
Organ Development	erythropoiesis	erythropoiesis of splenocytes	1.07E-03	2.07E-03	EGR1	1	
Oncogen Development	erythropoiesis	erythropoiesis of splenocytes	1.07E-03	2.07E-03	EGR1	1	
Tissue Development	erythropoiesis	erythropoiesis of splenocytes	1.07E-03	2.07E-03	EGR1	1	
Infectious Disease	infection	infection by Chikungunya virus	1.07E-03	2.07E-03	ISG15	1	
Cell-To-Cell Signaling and Interaction	long-term potentiation	long-term potentiation of dentate gyrus	1.07E-03	2.07E-03	EGR1	1	
Nervous System Development and Function	long-term potentiation	long-term potentiation of dentate gyrus	1.07E-03	2.07E-03	EGR1	1	
Hepatic System Development and Function	metaphase/anaphase transition	metaphase/anaphase transition of liver cells	1.07E-03	2.07E-03	EGR1	1	
Cell Cycle	metaphase/anaphase transition	metaphase/anaphase transition of liver cells	1.07E-03	2.07E-03	EGR1	1	
Hematological System Development and Function	opsonization	opsonization of red blood cells	1.07E-03	2.07E-03	C3	1	
Cell Death	opsonization	opsonization of red blood cells	1.07E-03	2.07E-03	C3	1	
Cellular Growth and Proliferation	outgrowth	outgrowth of breast cancer cell lines	1.07E-03	2.07E-03	CYR61	1	
Cellular Development	outgrowth	outgrowth of breast cancer cell lines	1.07E-03	2.07E-03	CYR61	1	
Cardiovascular System Development and Function	structural integrity	structural integrity of vasculature	1.07E-03	2.07E-03	CYR61	1	
Cancer	tumorigenesis	tumorigenesis of endometrial cancer cell lines	1.07E-03	2.07E-03	CYR61	1	
Reproductive System Development and Function	function	function of reproductive system	1.07E-03	2.07E-03	C3,EGR1	2	
Connective Tissue Development and Function	quantity	quantity of connective tissue cells	1.21E-03	2.31E-03	C3,CTGF,EGR1	3	
Tissue Morphology	quantity	quantity of connective tissue cells	1.21E-03	2.31E-03	C3,CTGF,EGR1	3	
Reproductive System Disease	endometriosis	endometriosis	1.32E-03	2.42E-03	CTGF,CYR61,EGR1,SGK1	4	
Renal and Urological Disease	albuminuria	albuminuria	1.59E-03	2.69E-03	CTGF,SGK1	2	
Metabolic Disease	albuminuria	albuminuria	1.59E-03	2.69E-03	CTGF,SGK1	2	
Endocrine System Development and Function	ovulation	ovulation	1.82E-03	2.87E-03	CTGF,EGR1	2	
Connective Tissue Development and Function	opposition	opposition of femur	2.13E-03	3.13E-03	CTGF	1	
Skeletal and Muscular System Development and Function	opposition	opposition of femur	2.13E-03	3.13E-03	EGR1	1	
Cellular Compromises	atrophy	atrophy of Leydig cells	2.13E-03	3.13E-03	EGR1	1	
Organismal Injury and Abnormalities	atrophy	atrophy of Leydig cells	2.13E-03	3.13E-03	EGR1	1	
Lipid Metabolism	clearance	clearance of fatty acid	2.13E-03	3.13E-03	C3	1	
Molecular Transport	clearance	clearance of fatty acid	2.13E-03	3.13E-03	C3	1	
Small Molecule Biochemistry	clearance	clearance of fatty acid	2.13E-03	3.13E-03	C3	1	
Cellular Growth and Proliferation	colonization	colonization of endometrial cancer cell lines	2.13E-03	3.13E-03	CYR61	1	
Infectious Disease	infection	infection of central nervous system	2.13E-03	3.13E-03	C3	1	
Neurological Disease	infectivity	infection of central nervous system	2.13E-03	3.13E-03	C3	1	
Infectious Disease	localization	infectivity of Zaire ebolavirus	2.13E-03	3.13E-03	CLEC10A	1	
Hematological System Development and Function	localization	localization of neutrophils	2.13E-03	3.13E-03	C3	1	
Inflammatory Response	localization	localization of neutrophils	2.13E-03	3.13E-03	C3	1	
Cellular Movement	localization	localization of neutrophils	2.13E-03	3.13E-03	C3	1	
Cellular Function and Maintenance	localization	localization of neutrophils	2.13E-03	3.13E-03	C3	1	
Immune Cell Trafficking	localization	localization of neutrophils	2.13E-03	3.13E-03	C3	1	



Tissue Morphology	quantity	quantity of blood platelets	6.07E-03	CLEC10A, EGR1
Inflammatory Response	inflammation	inflammation of liver	6.18E-03	C3, EGR1
Digestive System Development and Function	inflammation	inflammation of liver	6.18E-03	C3, EGR1
Hepatic System Development and Function	inflammation	inflammation of liver	6.18E-03	C3, EGR1
Organ Development	inflammation	inflammation of liver	6.18E-03	C3, EGR1
Cell Death	apoptosis	apoptosis of endometrial cancer cell lines	6.38E-03	CYR61
Cellular Development	maturatation	maturatation of plasmacytoid dendritic cells	6.38E-03	EGR1
Hematopoiesis	maturatation	maturatation of plasmacytoid dendritic cells	6.38E-03	EGR1
Cardiovascular System Development and Function	morphogenesis	morphogenesis of atrioventricular valve	6.38E-03	CYR61
Embryonic Development	morphogenesis	morphogenesis of atrioventricular valve	6.38E-03	CYR61
Organ Development	morphogenesis	morphogenesis of atrioventricular valve	6.38E-03	CYR61
Organismal Development	morphogenesis	morphogenesis of atrioventricular valve	6.38E-03	CYR61
Tissue Development	morphogenesis	morphogenesis of atrioventricular valve	6.38E-03	CYR61
Cellular Compromise	perturbation	perturbation of cells	6.38E-03	CYR61
Neurological Disease	Huntington's disease	Huntington's disease	6.50E-03	C3, CTGF, EGR1, SGK1
Skeletal and Muscular Disorders	Huntington's disease	Huntington's disease	6.50E-03	C3, CTGF, EGR1, SGK1
Heredity Disorders	Huntington's disease	Huntington's disease	6.50E-03	C3, CTGF, EGR1, SGK1
Connective Tissue Development and Function	development	cartilage development	6.51E-03	CTGF, CYR61
Skeletal and Muscular System Development and Function	development	cartilage development	6.51E-03	CTGF, CYR61
Tissue Development	development	cartilage development	6.51E-03	CTGF, CYR61
Organismal Survival	mortality	mortality	6.51E-03	C3, ISG15
Cellular Growth and Proliferation	proliferation	proliferation of brain cancer cell lines	7.18E-03	CYR61, EGR1
Cellular Development	proliferation	proliferation of brain cancer cell lines	7.18E-03	CYR61, EGR1
Tissue Morphology	size	size of infarct	7.18E-03	C3, EGR1
Cardiovascular Disease	size	size of infarct	7.18E-03	C3, EGR1
Embryonic Development	development	development of ovary	7.41E-03	CTGF, EGR1
Organ Development	development	development of ovary	7.41E-03	CTGF, EGR1
Organismal Development	development	development of ovary	7.41E-03	CTGF, EGR1
Tissue Development	development	development of ovary	7.41E-03	CTGF, EGR1
Reproductive System Development and Function	development	development of ovary	7.41E-03	CTGF, EGR1
Renal and Urological System Development and Function	infection	Pneumococcal infection	7.44E-03	C3
Infectious Disease	abnormal morphology	abnormal morphology of conducting airway	7.44E-03	CTGF
Respiratory System Development and Function	adhesion	adhesion of stomach cancer cell lines	7.44E-03	CYR61
Cell-To-Cell Signaling and Interaction	adhesion	adhesion of stomach cancer cell lines	7.44E-03	CYR61
Tissue Development	apoptosis	apoptosis of acinar gland cells	7.44E-03	C3
Cell Death	cytostasis	cytostasis of carcinoma cell lines	7.44E-03	CYR61
Cellular Growth and Proliferation	clastic heart failure	diastolic heart failure	7.44E-03	CTGF
Cardiovascular Disease	differentiation	differentiation of rhabdomyosarcoma cell lines	7.44E-03	CYR61
Cellule Development	morphogenesis	morphogenesis of atrial septum	7.44E-03	CTGF
Cardiovascular System Development and Function	morphogenesis	morphogenesis of atrial septum	7.44E-03	CYR61
Embryonic Development	morphogenesis	morphogenesis of atrial septum	7.44E-03	CTGF
Organ Development	morphogenesis	morphogenesis of atrial septum	7.44E-03	CYR61
Organismal Development	morphogenesis	morphogenesis of atrial septum	7.44E-03	CTGF
Tissue Development	morphogenesis	morphogenesis of atrial septum	7.44E-03	CYR61
Renal and Urological Disease	tubulo-interstitial fibrosis	tubulo-interstitial fibrosis of kidney	7.44E-03	C3
Organismal Injury and Abnormalities	tubulo-interstitial fibrosis	tubulo-interstitial fibrosis of kidney	7.44E-03	CTGF
Gastrointestinal Disease	damage	damage of liver	7.76E-03	C3, EGR1
Hepatic System Disease	damage	damage of liver	7.76E-03	C3, EGR1
Organismal Injury and Abnormalities	damage	damage of liver	7.76E-03	C3, EGR1
Cell Death	apoptosis	apoptosis of gonadal cells	8.49E-03	CTGF, EGR1
Skeletal and Muscular Disorders	K/BxN serum transfer arthritis	K/BxN serum transfer arthritis	8.50E-03	C3
Connective Tissue Disorders	K/BxN serum transfer arthritis	K/BxN serum transfer arthritis	8.50E-03	C3
Inflammatory Disease	K/BxN serum transfer arthritis	K/BxN serum transfer arthritis	8.50E-03	C3
Inflammatory Response	alternative complement pathway	alternative complement pathway	8.50E-03	C3
Antigen Presentation	alternative complement pathway	alternative complement pathway	8.50E-03	C3
Humoral Immune Response	alternative complement pathway	alternative complement pathway	8.50E-03	C3
Ophthalmic Disease	choroidal neovascularization	choroidal neovascularization of eye	8.50E-03	C3
Hematological System Development and Function	differentiation	differentiation of B-1 lymphocytes	8.50E-03	EGR1
Cellular Development	differentiation	differentiation of B-1 lymphocytes	8.50E-03	EGR1
Hematopoiesis	differentiation	differentiation of B-1 lymphocytes	8.50E-03	EGR1
Humoral Immune Response	differentiation	differentiation of B-1 lymphocytes	8.50E-03	EGR1
Cellular Function and Maintenance	differentiation	differentiation of B-1 lymphocytes	8.50E-03	CTGF
Cellular Assembly and Organization	function	function of chondrocytes	8.50E-03	Mup1 (includes others)
Cellular Function and Maintenance	morphogenesis	morphogenesis of mitochondria	8.50E-03	SGK1
Molecular Transport	reabsorption	reabsorption of Na+	8.50E-03	SGK1
Infectious Disease	reabsorption	reabsorption of Na+	8.50E-03	SGK1
Neurological Disease	scrapping	scrapping	8.50E-03	C3
Cancer	scrapping	scrapping	8.50E-03	C3
Cancer	tumorigenesis	tumorigenesis of cells	8.57E-03	C3, CYR61, EGR1
Haematological Disease	T-cell non-Hodgkin's disease	T-cell non-Hodgkin's disease	8.74E-03	CYR61, EGR1
Cellular Growth and Proliferation	T-cell non-Hodgkin's disease	T-cell non-Hodgkin's disease	8.74E-03	CYR61, EGR1
Cellular Development	proliferation	proliferation of epithelial cell lines	9.38E-03	CYR61, EGR1
Hair and Skin Development and Function	proliferation	proliferation of epithelial cell lines	9.38E-03	CYR61, EGR1
Tissue Morphology	abnormal morphology	abnormal morphology of nasal capsule	9.55E-03	CYR61, EGR1
Cell Death	cytolysis	cytolysis of leukemia cell lines	9.55E-03	CTGF
Cell Death	cytolysis	cytolysis of lymphoma cell lines	9.55E-03	ISG15
Cancer	development	development of T-cell non-Hodgkin lymphoma	9.55E-03	ISG15
Immunological Disease	development	development of T-cell non-Hodgkin lymphoma	9.55E-03	EGR1
Immunological Disease	hypersensitivity reaction type II	hypersensitivity reaction type II	9.55E-03	EGR1
Antimicrobial Response	immune protection	immune protection of organism	9.55E-03	C3
Cellular Movement	migration	migration of bone marrow cell lines	9.55E-03	C3
Inflammatory Response	pulmonary interstitial fibrosis	pulmonary interstitial fibrosis	9.55E-03	EGR1
Organismal Injury and Abnormalities	pulmonary interstitial fibrosis	pulmonary interstitial fibrosis	9.55E-03	EGR1
Connective Tissue Disorders	pulmonary interstitial fibrosis	pulmonary interstitial fibrosis	9.55E-03	EGR1
Inflammatory Disease	pulmonary interstitial fibrosis	pulmonary interstitial fibrosis	9.55E-03	EGR1
Respiratory Disease	pulmonary interstitial fibrosis	pulmonary interstitial fibrosis	9.55E-03	EGR1
Connective Tissue Development and Function	ossification	ossification of bone	1.02E-02	CTGF, ISG15
Embryonic Development	ossification	ossification of bone	1.02E-02	CTGF, ISG15
Organ Development	ossification	ossification of bone	1.02E-02	CTGF, ISG15







Organ Development	abnormal morphology	abnormal morphology of ulna	3.05E-02	CTGF
Organismal Development	abnormal morphology	abnormal morphology of ulna	3.05E-02	CTGF
Skeletal and Muscular System Development and Function	abnormal morphology	abnormal morphology of ulna	3.05E-02	CTGF
Tissue Development	adhesion	adhesion of embryonic cell lines	3.05E-02	C3
Cell-To-Cell Signaling and Interaction	adhesion	adhesion of embryonic cell lines	3.05E-02	C3
Embryonic Development	adhesion	adhesion of embryonic cell lines	3.05E-02	C3
Tissue Development	adhesion	adhesion of embryonic cell lines	3.05E-02	C3
Neurological Disease	age-related macular degeneration	age-related macular degeneration	3.05E-02	C3
Ophthalmic Disease	age-related macular degeneration	age-related macular degeneration	3.05E-02	C3
Infectious Disease	dengue hemorrhagic fever	dengue hemorrhagic fever	3.05E-02	PCOLCE2 (includes EG:26577)
Neurological Disease	febrile seizure	febrile seizure	3.05E-02	C3
Organismal Injury and Abnormalities	formation	formation of thrombus	3.05E-02	C3
Cardiovascular Disease	hypoplasia	hypoplasia of lung	3.05E-02	CTGF
Developmental Disorder	hypoplasia	hypoplasia of lung	3.05E-02	CTGF
Respiratory Disease	inflammation	inflammation of brain	3.05E-02	C3
Inflammatory Response	ossification	ossification of endochondral bone	3.05E-02	CTGF
Connective Tissue Development and Function	ossification	ossification of endochondral bone	3.05E-02	CTGF
Embryonic Development	ossification	ossification of endochondral bone	3.05E-02	CTGF
Organ Development	ossification	ossification of endochondral bone	3.05E-02	CTGF
Organismal Development	ossification	ossification of endochondral bone	3.05E-02	CTGF
Skeletal and Muscular System Development and Function	ossification	ossification of endochondral bone	3.05E-02	CTGF
Tissue Development	size	size of germinal center	3.05E-02	C3
Lymphatic Tissue Structure and Development	size	size of germinal center	3.05E-02	C3
Humoral Immune Response	synthesis	synthesis of proteoglycan	3.05E-02	CTGF
Small Molecule Biochemistry	synthesis	synthesis of proteoglycan	3.05E-02	CTGF
Carbohydrate Metabolism	vasculogenesis	vasculogenesis	3.09E-02	C3
Cardiovascular System Development and Function	vasculogenesis	vasculogenesis	3.09E-02	CTGF,CYR61
Organismal Development	concentration	concentration of triacylglycerol	3.12E-02	C3,CYR61
Lipid Metabolism	concentration	concentration of triacylglycerol	3.12E-02	EGR1
Molecular Transport	concentration	concentration of triacylglycerol	3.12E-02	C3,EGR1
Small Molecule Biochemistry	adhesion	adhesion of fibroblasts	3.15E-02	C3,EGR1
Cell-To-Cell Signaling and Interaction	adhesion	adhesion of fibroblasts	3.15E-02	CTGF
Connective Tissue Development and Function	adhesion	adhesion of fibroblasts	3.15E-02	CTGF
Tissue Development	adhesion	adhesion of fibroblasts	3.15E-02	CTGF
Organ Morphology	volume	volume of trabecular bone	3.15E-02	CTGF
Tissue Morphology	volume	volume of trabecular bone	3.15E-02	CTGF
uterine serous papillary cancer	uterine serous papillary cancer	3.23E-02	CLDN5,IG515	
uterine serous papillary cancer	uterine serous papillary cancer	3.23E-02	CLDN5,IG515	
Reproductive System Disease	abnormal morphology	abnormal morphology of long bone hypertrophic chondrocyte zone	3.25E-02	CTGF
Organ Morphology	abnormal morphology	abnormal morphology of long bone hypertrophic chondrocyte zone	3.25E-02	CTGF
Connective Tissue Development and Function	abnormal morphology	abnormal morphology of long bone hypertrophic chondrocyte zone	3.25E-02	CTGF
Embryonic Development	abnormal morphology	abnormal morphology of long bone hypertrophic chondrocyte zone	3.25E-02	CTGF
Organ Development	abnormal morphology	abnormal morphology of long bone hypertrophic chondrocyte zone	3.25E-02	CTGF
Organismal Development	abnormal morphology	abnormal morphology of long bone hypertrophic chondrocyte zone	3.25E-02	CTGF
Skeletal and Muscular System Development and Function	abnormal morphology	abnormal morphology of long bone hypertrophic chondrocyte zone	3.25E-02	CTGF
Tissue Development	abnormal morphology	abnormal morphology of long bone hypertrophic chondrocyte zone	3.25E-02	CTGF
Tissue Morphology	migration	migration of leukocyte cell lines	3.25E-02	C3
Hematological System Development and Function	migration	migration of leukocyte cell lines	3.25E-02	C3
Cellular Movement	migration	migration of leukocyte cell lines	3.25E-02	C3
Immune Cell Trafficking	tubulation	tubulation of vascular endothelial cells	3.25E-02	CYR61
Cell Biology	tubulation	tubulation of vascular endothelial cells	3.25E-02	CYR61
Cardiovascular System Development and Function	tubulation	tubulation of vascular endothelial cells	3.25E-02	CYR61
Cellular Development	tubulation	tubulation of vascular endothelial cells	3.25E-02	CYR61
Organismal Development	tubulation	tubulation of vascular endothelial cells	3.25E-02	CYR61
Tissue Development	tubulation	tubulation of vascular endothelial cells	3.25E-02	CYR61
Organ Morphology	abnormal morphology	abnormal morphology of cornea	3.36E-02	EGR1
Embryonic Development	abnormal morphology	abnormal morphology of cornea	3.36E-02	EGR1
Organ Development	abnormal morphology	abnormal morphology of cornea	3.36E-02	EGR1
Organismal Development	abnormal morphology	abnormal morphology of cornea	3.36E-02	EGR1
Tissue Development	abnormal morphology	abnormal morphology of cornea	3.36E-02	EGR1
Visual System Development and Function	abnormal morphology	abnormal morphology of cornea	3.36E-02	EGR1
Skeletal and Muscular System Development and Function	abnormal morphology	abnormal morphology of cornea	3.36E-02	EGR1
Cell-To-Cell Signaling and Interaction	abnormal morphology	abnormal morphology of rib cage	3.36E-02	CTGF
Tissue Development	adhesion	adhesion of epithelial cell lines	3.36E-02	C3
Hair and Skin Development and Function	adhesion	adhesion of epithelial cell lines	3.36E-02	C3
Organ Morphology	lack	lack of corpus luteum	3.36E-02	EGR1
Reproductive System Development and Function	lack	lack of corpus luteum	3.36E-02	EGR1
Cellular Growth and Proliferation	proliferation	proliferation of myeloma cell lines	3.36E-02	SGK1
Cellular Development	proliferation	proliferation of myeloma cell lines	3.36E-02	SGK1
Tissue Morphology	area	area of atherosclerotic lesion	3.46E-02	EGR1
Organismal Injury and Abnormalities	area	area of atherosclerotic lesion	3.46E-02	EGR1
Cardiovascular Disease	area	area of atherosclerotic lesion	3.46E-02	EGR1
Immunological Disease	area	area of atherosclerotic lesion	3.46E-02	EGR1
Hematological System Development and Function	lymphadenopathy	lymphadenopathy	3.46E-02	C3
Inflammatory Response	phagocytosis	phagocytosis of neutrophils	3.46E-02	C3
Cell-Cell Signaling and Interaction	phagocytosis	phagocytosis of neutrophils	3.46E-02	C3
Cellular Barriers and Maintenance	phagocytosis	phagocytosis of neutrophils	3.46E-02	C3
Antigen Presentation	phagocytosis	phagocytosis of neutrophils	3.46E-02	C3
Gastrointestinal Disease	phagocytosis	phagocytosis of neutrophils	3.46E-02	CTGF
Hepatic System Disease	primary sclerosing cholangitis	primary sclerosing cholangitis	3.46E-02	CTGF
Inflammatory Disease	primary sclerosing cholangitis	primary sclerosing cholangitis	3.46E-02	CTGF
Lipid Metabolism	primary sclerosing cholangitis	primary sclerosing cholangitis	3.46E-02	C3
Molecular Transport	uptake	uptake of fatty acid	3.46E-02	C3
Small Molecule Biochemistry	uptake	uptake of fatty acid	3.46E-02	C3
Hematological System Development and Function	adhesion	adhesion of blood platelets	3.56E-02	C3
Inflammatory Response	adhesion	adhesion of blood platelets	3.56E-02	C3
Cell-To-Cell Signaling and Interaction	adhesion	adhesion of blood platelets	3.56E-02	C3
Tissue Development	adhesion	adhesion of blood platelets	3.56E-02	C3
Endocrine System Development and Function	concentration	concentration of aldosterone	3.56E-02	SGK1
Lipid Metabolism	concentration	concentration of aldosterone	3.56E-02	SGK1





**Supplementary Table 3 : Microarray analysis in DSS/AOM-treated wild-type and *Nod2*-deficient mice according antibiotic treatment.**  
***Nod2*<sup>+/+</sup> Atb-treated vs *Nod2*<sup>+/+</sup>**

ID	Symbol	Entrez Gene Name	Gene Type	Chr	Start	End	Strand	RefSeq	Alt RefSeq	Protein ID	Protein Name	Log Ratio	p-value
A_52_P482781	HRNR	hornerin	protein	1	3,531	3,84E-06	-	A_51_P27128	IL6	A_52_P59190	claudin 8	-4,108	1,08E-08
A_51_P202501	KPRP	keratinocyte proline-rich protein	protein	1	3,234	3,19E-06	-	A_52_P59190	CLDN8	A_51_P401907	cystatin A (stefin A)	3,817	2,75E-08
A_51_P182876	Gm10229	predicted gene 10229	protein	1	2,913	1,70E-07	-	A_52_P64613	SULT1B1	A_51_P254855	sulfotransferase family, cytosolic, 1B, member 1	-3,664	1,41E-07
A_52_P460868	TCHH	trichohyalin	protein	1	2,828	1,88E-07	-	A_51_P254855	PTGS2	A_51_P481721	prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase)	3,556	5,38E-10
A_51_P424275	MT4	metallothionein 4	protein	1	2,827	3,17E-07	-	A_51_P51182	Sult1d1	A_51_P254855	sulfotransferase family 1D, member 1	-3,499	3,39E-09
A_51_P169671	REG3A	regenerating islet-derived 3 alpha	protein	1	2,755	1,16E-04	-	A_51_P20126	interferon activated gene 202B	A_51_P20126	interferon activated gene 202B	3,266	2,40E-06
A_52_P4244	LCE3D	late cornified envelope 3D	protein	1	2,668	3,39E-09	-	A_51_P230387	Sval1	A_52_P228281	seminal vesicle antigen-like	-3,227	3,75E-07
A_51_P183554	J2UFA1	POU class 2 associating factor 1	protein	1	2,630	2,73E-07	-	A_52_P230387	CAMK2L1	A_51_P20126	chemokine (C-X-C motif) ligand 1 (melanoma growth stimulating activity, alpha)	3,118	6,65E-08
A_52_P366773	Foxo3	soluble transcription factor 1	protein	1	2,612	4,06E-08	-	A_52_P230387	SLC7A11	A_51_P20126	RIGEN cDNA 1100001G20Rik	3,010	3,26E-05
A_52_P208521	Khd1b18	(includes others)	KH domain containing 1B	1	2,549	1,80E-06	-	A_51_P244560	Pbp2	A_52_P633446	phosphatidylethanolamine-binding protein 2	-2,920	3,14E-05
A_52_P62059	CYP2C19	cytochrome P450, family 2, subfamily C, polypeptide 19	protein	1	2,547	3,17E-07	-	A_51_P117422	HACO2	A_51_P456115	solute carrier family 7 (anionic amino acid transporter light chain, xc- system), member 11	-2,910	2,56E-05
A_52_P482251	GJB8	gap junction protein, beta 6, 30kDa	protein	1	2,537	2,64E-07	-	A_51_P456115	KIAA1199	A_51_P437889	hydroxycarboxyl oxidase 2 (long chain)	2,893	2,80E-05
A_51_P104681	KRTAP6-1	keratin associated protein 6-1	protein	1	2,528	4,08E-05	-	A_51_P325904	INHBB	A_51_P456115	KIAA1199	-2,844	1,43E-06
A_51_P104478	SMPX	small muscle protein, X-linked	protein	1	2,492	6,33E-08	-	A_52_P40244	LCE3D	A_51_P20126	anterior gradient 3 homolog (Xenopus laevis)	2,822	1,61E-07
A_51_P457187	TNFRSF13C	tumor necrosis factor receptor superfamily, member 13C	protein	1	2,448	3,30E-07	-	A_51_P230387	Vav1	A_52_P208521	inhibin, beta B	-2,819	2,58E-05
A_51_P351378	C5orf146	chromosome 5 open reading frame 46	protein	1	2,429	4,47E-07	-	A_51_P230387	CAMP	A_52_P208521	late cornified envelope 3D	2,814	4,69E-05
A_51_P336049	KRT33B	CAMP (includes EG:12796)	keratin 33B	1	2,417	5,72E-08	-	A_51_P385765	SPP1	(includes EG:100359743)	secreted phosphoprotein 1	-2,742	1,88E-07
A_51_P180423	CD161b	late cornified envelope 1B	protein	1	2,396	3,66E-08	-	A_52_P208521	Khd1b18	(includes others)	KH domain containing 1B	-2,727	4,77E-07
A_51_P125334	Lev1b	keratin 7	protein	1	2,391	1,36E-07	-	A_52_P652020	CYP2C19	A_51_P20126	cytochrome P450, family 2, subfamily C, polypeptide 19	2,725	6,19E-08
A_51_P125140	OP91	carbamoyl-phosphatase synthase 1, mitochondrial	protein	1	2,376	5,41E-07	-	A_51_P20126	E74-like factor 5	(includes EG:100359743)	E74-like factor 5 (zf-CAM domain transcription factor)	-2,700	1,09E-07
A_51_P149853	SEC14L4	SEC14-like 4 (S. cerevisiae)	protein	1	2,370	7,37E-06	-	A_51_P349415	AADC	A_52_P573872	WD repeat domain 72	-2,665	7,14E-05
A_51_P274796	ACE2	(includes EG:302668)	angiotensin I converting enzyme (peptidyl-dipeptidase A) 2	protein	2,350	8,60E-07	-	A_51_P404193	SP5	A_51_P20126	lysyl endopeptidase (esterase)	-2,645	2,16E-04
A_52_P677117	PTPRC	protein tyrosine phosphatase, receptor type, C	protein	1	2,343	5,72E-08	-	A_51_P20126	Sps	A_51_P20126	Sps transcription factor	-2,611	4,49E-05
A_51_P432641	CXCL10	chemokine (C-X-C motif) ligand 10	protein	1	2,290	4,25E-06	-	A_51_P20126	CAMP	(includes EG:12796)	cathelicidin antimicrobial peptide	-2,605	8,17E-06
A_51_P08649	SHISA2	shisa homolog 2 (Xenopus laevis)	protein	1	2,252	5,72E-08	-	A_51_P244523	VNN1	A_51_P20126	vanin 1	-2,599	1,27E-06
A_51_P349115	ADAC	arylacetamide deacetylase (esterase)	protein	1	2,246	2,33E-04	-	A_52_P560407	LILRB3	A_51_P20126	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3	-2,519	5,09E-05
A_51_P343789	AGR3	anterior gradient 3 homolog (Xenopus laevis)	protein	1	2,237	7,09E-07	-	A_51_P255699	MMP3	A_51_P20126	matrix metallopeptidase 3 (stromelysin 1, progelatinase A)	-2,518	4,33E-07
A_51_P231230	MMP8	matrix metalloproteinase 8 (neutrophil collagenase)	protein	1	2,234	5,94E-08	-	A_51_P677117	PTPRC	A_51_P443976	protein tyrosine phosphatase, receptor type, C	-2,514	1,19E-05
A_51_P414208	LCE1B	late cornified envelope 1B	protein	1	2,162	5,10E-05	-	A_51_P666200	TNFRSF11B	A_51_P20126	protein tyrosine phosphatase, receptor type, R	-2,507	9,66E-06
A_51_P33583	EFRA	ELL associated factor 2	protein	1	2,155	3,69E-07	-	A_51_P20126	TIMP1	A_51_P20126	tnur necrosis factor receptor superfamily, member 11b	-2,494	4,35E-05
A_51_P235015	EP0309G21Rik	RIKEN cDNA 201309G21 gene	protein	1	2,146	5,32E-06	-	A_51_P167242	ITGB3/Chi34	A_51_P20126	chitinase 3-like 4	-2,482	1,49E-06
A_51_P307150	ITGB3	hyaluronan and proteoglycan 2 (laminin chain)	protein	1	2,145	5,43E-08	-	A_51_P167242	OFL1	A_51_P20126	complement factor I	-2,462	7,17E-06
A_52_P307150	Kira15	indoleamine 2,3-dioxygenase 1	protein	1	2,129	5,10E-09	-	A_51_P174915	GJA1	A_51_P20126	vanin 1	-2,447	1,03E-07
A_52_P514407	Kira15	killer cell lectin-like receptor, subfamily A, member 15	protein	1	2,107	1,10E-05	-	A_51_P20126	TRPV3	A_51_P20126	gap junction protein, alpha 1, 43kDa	-2,442	8,67E-11
A_51_P273298	FAIM3	Fas apoptotic inhibitory molecule 3	protein	1	2,076	7,76E-07	-	A_51_P235702	RPN2	A_51_P20126	transient receptor potential cation channel, subfamily V, member 3	-2,410	1,83E-05
A_51_P211346	GPBP3	G protein-coupled receptor 83	protein	1	2,070	1,36E-06	-	A_52_P237764	HAS2	A_51_P20126	hyaluronan synthase 2	-2,398	1,69E-05
A_52_P151887	IGHA1	immunoglobulin heavy constant alpha 1	protein	1	2,069	7,25E-08	-	A_51_P213120	MMP8	A_51_P20126	matrix metallopeptidase 8 (neutrophil collagenase)	-2,396	1,88E-05
A_51_P390127	KLRC1	killer cell lectin-like receptor subfamily C, member 1	protein	1	2,052	4,98E-08	-	A_51_P135161	S100C	A_51_P20126	S100 calcium binding protein 2	-2,392	1,17E-05
A_51_P154485	FCRLA	Fc receptor-like A	protein	1	2,041	7,98E-07	-	A_51_P245561	Ilf1tm6	A_52_P59975	interferon induced transmembrane protein 6	-2,386	1,67E-05
A_51_P272685	UXO	urate oxidase, pseudogene	protein	1	2,041	2,38E-08	-	A_51_P20126	CXCR2	A_51_P20126	chemokine (C-X-C motif) receptor 2	-2,377	1,79E-05
A_51_P334785	IIl204	(includes others)	interferon activated gene 204	protein	2,025	3,08E-08	-	A_51_P488739	HCAR2	A_51_P20126	hydroxycarboxylic acid receptor 2	-2,369	1,07E-07
A_51_P310105	LTA	lymphotoxin alpha (TNF superfamily, member 1)	protein	1	2,022	2,81E-06	-	A_51_P207940	TMEM117	A_51_P20126	transmembrane protein 117	-2,366	8,28E-10
A_51_P461040	CRCT1	cytosteine-rich C-terminal 1	protein	1	1,972	2,09E-07	-	A_51_P185845	ANKK1	A_51_P20126	ankyrin repeat and kinase domain containing 1	-2,357	2,09E-05
A_52_P811324	E330019L11Rik	predicted gene E330019L11 gene	protein	1	1,962	6,44E-06	-	A_51_P233620	REBNX2	A_51_P20126	numerous E-box binding factor 2	-2,356	7,38E-05
A_51_P100907	SerpinB3a/SerpinB3c	serine (or cysteine) peptidase inhibitor, clade B (ovalbumin), member 3B	protein	1	1,955	1,66E-05	-	A_51_P265692	CEBPB	(includes EG:1051)	CCAAT/enrichment binding protein (C/EBP), beta	-2,355	4,45E-05
A_51_P271218	IL6	interleukin 6 (interferon, beta 2)	protein	1	1,943	1,70E-06	-	A_51_P118545	S100C	A_51_P20126	S100 calcium binding protein G	-2,346	4,17E-05
A_51_P442889	IGHM	immunoglobulin heavy constant mu	protein	1	1,927	3,68E-08	-	A_52_P441070	APOB	A_52_P877113	apolipoprotein B (including IgM antigen)	-2,345	6,92E-05
A_51_P272683	MSA1	membrane-spanning 4-domains, subfamily A, member 1	protein	1	1,925	1,41E-04	-	A_52_P5945	TIMP1	A_51_P20126	TIMP metallopeptidase inhibitor	-2,344	2,95E-05
A_52_P147466	Kira2	killer cell lectin-like receptor, subfamily A, member 2	protein	1	1,923	2,87E-05	-	A_51_P468558	TREML4	A_51_P20126	peroxisome proliferator-activated receptor gamma, coactivator 1 alpha	-2,313	2,09E-05
A_52_P413395	SLN	sarcoelin	protein	1	1,921	2,31E-06	-	A_52_P198898	SAMD5	A_52_P20126	triggering receptor expressed on myeloid cells-like 4	-2,305	2,77E-07
A_51_P401907	CSTA	cystatin A (stefin A)	protein	1	1,911	8,05E-05	-	A_52_P249926	HHIP	A_51_P20126	sterile alpha motif domain containing 5	-2,304	3,04E-05
A_51_P465640	TNFSF11	tnfrfncor factor (ligand) superfamily, member 11	protein	1	1,911	1,03E-05	-	A_52_P1084681	ADAM28	A_51_P20126	hedgehog interacting protein	-2,303	2,77E-07
A_51_P465646	CTSG	cathepsin G	protein	1	1,902	6,77E-08	-	A_52_P497235	C19orf59	A_51_P20126	ADAM metallopeptidase domain 28	-2,299	6,27E-07
A_51_P351381	FABP2	fatty acid binding protein 2, intestinal	protein	1	1,896	8,87E-08	-	A_51_P188200	TFAP2C	A_51_P20126	chromosome 19 open reading frame 59	-2,283	8,38E-07
A_51_P247534	MYH2	myosin, heavy chain 2, skeletal muscle, adult	protein	1	1,892	3,09E-06	-	A_51_P388491	TREM1	A_51_P20126	transcription factor AP-2 gamma (activating enhancer binding protein 2 gamma)	-2,280	1,15E-05
A_51_P424767	Igkv1	immunoglobulin kappa chain variable	protein	1	1,879	1,87E-05	-	A_51_P113169	Akr1c19	A_51_P20126	triggering receptor expressed on myeloid cells-1, member C19	-2,277	6,78E-05
A_51_P230837	Sival1	immunoglobulin kappa chain complex	protein	1	1,874	8,41E-05	-	A_51_P107315	SLO548	A_51_P20126	soluble carrier family 5 (oxide transporter), member 8	-2,260	2,62E-05
A_51_P167292	Chi3/Chi34	chitinas 3-like 2	protein	1	1,872	3,03E-04	-	A_51_P485778	HPGD	A_51_P20126	hydroxyprostaglandin dehydrogenase 5-(NAD)	-2,252	4,12E-05
A_52_P559575	CXCR2	chemokine (C-X-C motif) receptor 2	protein	1	1,868	1,60E-04	-	A_51_P211506	MUC20	A_51_P20126	mucin 20, cell surface associated	-2,238	1,83E-05
A_52_P227664	GPBP3	guanylate binding protein 5	protein	1	1,851	9,89E-08	-	A_51_P211436	GPR83	A_51_P20126	hedgehog interacting protein	-2,237	3,09E-07
A_52_P284177	Akr1c14	aldo-keto reductase family 1, member 1	protein	1	1,846	5,72E-08	-	A_51_P274746	AEC2	(includes EG:302668)	G protein-coupled receptor 83	-2,234	4,18E-05
A_51_P167294	ZGMA	granzyme A (granzyme 1, cytotoxic T-lymphocyte-associated serine esterase 3)	protein	1	1,836	1,51E-06	-	A_51_P202501	KPRP	A_51_P20126	keratinocyte proline-rich protein	-2,224	4,42E-05
A_51_P416617	DERL3	carbonic anhydrase III, muscle specific	protein	1	1,822	1,80E-08	-	A_51_P358012	Serpina3k	(includes others)	chromosome X open reading frame 57	-2,221	2,40E-05
A_51_P163578	UGT2B7	UDP glucuronosyltransferase 2 family, polypeptide B7	protein	1	1,806	2,34E-05	-	A_51_P273439	PRKG2	A_51_P20126	protein kinase, cGMP-dependent, type II	-2,198	2,14E-07
A_51_P371700	MARCO	macrophage receptor with collagenous structure	protein	1	1,799	2,86E-07	-	A_52_P307739	SOX2	A_51_P20126	SRY (sex determining region Y)-box 2	-2,195	1,79E-06
A_52_P104441	CCSP-B8B	SATB homeobox 1	protein	1	1,788	2,91E-05	-	A_52_P249572	CCL2	A_51_P20126	chemokine (C-C motif) ligand 2	-2,181	5,09E-05
A_52_P549180	GLDN8	coiled-coil domain containing 88B	protein	1	1,786	2,35E-05	-	A_51_P222141	OCDC88B	A_51_P20126	stem cell factor 1 like 2	-2,178	2,05E-05
A_51_P471715	ACSM3	acyl-CoA synthetase medium-chain family member 3	protein	1	1,784	1,16E-06	-	A_51_P297915	LRRK19	A_51_P20126	coiled-coil domain containing 88B	-2,148	2,72E-05
A_52_P111031	PCDH17	protocadherin 17	protein	1	1,782	3,39E-09	-	A_51_P233581	FABP2	A_51_P20126	leucine rich repeat containing 19	-2,140	1,07E-05
A_51_P221557	753kDa2B23Rik	protoactinin 17	protein	1	1,778	6,74E-08	-	A_51_P183881	SLFN12	A_51_P20126	fatty acid binding protein 2, intestinal	-2,135	1,37E-05
A_52_P485558	TMLM4	triggering receptor expressed on myeloid cells-4	protein	1	1,775	4,88E-07	-	A_51_P245090	AQP3	A_51_P20126	schafman family member 12	-2,120	3,64E-07
A_52_P524520	STAT1	signal transducer and activator of transcription 1, 91kDa	protein	1	1,770	2,06E-07	-	A_51_P279437	MFS2D2	A_51_P20126	major facilitator superfamily domain containing 2A	-2,114	8,04E-07
A_51_P742475	TLTL6	tubulin tyrosine ligase-like family, member 6	protein	1	1,768	5,18E-08	-	A_52_P247847	CLSTN2	A_51_P20126	calyminen 2	-2,113	1,77E-08
A_52_P428608	Ces1/Ces1b	carboxylesterase 1C	protein	1	1,765	2,40E-08	-	A_51_P224164	SLC26A4	A_51_P20126	solute carrier family 26, member 4	-2,111	1,53E-05
A_51_P366811	LOC100505002	uncharacterized LOC100505002	protein	1	1,764	3,16E-06	-	A_51_P383032	CLEC4D	A_51_P20126	C-type lectin domain family 4, member D	-2,097	8,83E-05

A_51_P461665	CXCL9	-1.715	5.83E-07	A_52_P430194	SMOX	-2.041	1.64E-08
A_51_P206019	Ighg3	-1.713	5.46E-05	A_52_P365615	Ces2e	2.038	1.34E-07
A_52_P5945	PPARGC1A	1.707	2.02E-07	A_51_P159565	NFE2	-2.036	2.58E-09
A_51_P305896	CFTR	1.706	3.17E-07	A_51_P248453	SEC14L4	2.036	7.41E-05
A_52_P300344	PM2D01	1.703	4.06E-06	A_51_P205209	CD244	-2.032	1.23E-09
A_52_P18746	IKZF3	1.702	6.66E-06	A_51_P346815	Arxes1/Arxes2	-2.023	6.91E-08
A_52_P12133	H2-Q8	1.696	5.72E-08	A_51_P406734	ST8SIA5	2.023	1.00E-08
A_51_P355702	TRPV3	1.694	7.82E-07	A_51_P163578	UGT2B7	2.014	1.19E-04
A_51_P470542	APOB	1.686	4.40E-06	A_51_P290826	CALCA	-2.005	3.29E-06
A_51_P452153	MZB1	1.685	6.96E-07	A_52_P619248	Pif2c2 (includes others)	-2.005	3.92E-07
A_51_P245855	PTGS2	1.680	2.07E-06	A_52_P456640	FGR	-1.999	3.56E-09
				A_51_P487177	ACSM2		
				A_51_P403477	DIO1	-1.997	1.06E-07
				A_51_P247161	WIF1	-1.997	6.80E-08
				A_51_P226234	SELL	-1.995	2.16E-07
				A_51_P183197	Corf1f69	-1.993	1.89E-07
				A_51_P108108	LRRK25	-1.983	2.13E-06
				A_51_P432607	Spink6	-1.964	4.45E-08
				A_51_P365516	SPINK1	-1.964	4.20E-09
				A_52_P402394	TMEM121	-1.959	7.63E-07
				A_51_P373142	A1854703	-1.954	8.39E-08
				A_52_P384264	Zfp185	-1.950	4.81E-09
				A_51_P109369	FBXO32	-1.948	7.81E-08
				A_52_P83906	SLC26A2	-1.946	3.61E-06
				A_51_P247097	IL1R2	-1.945	4.48E-06
				A_51_P203675	TREM3	-1.944	3.08E-07
				A_51_P362610	CXL10	-1.943	9.21E-06
				A_51_P144915	MAP1A	-1.929	3.88E-05
				A_51_P247098	GDA	-1.916	7.40E-03
				A_52_P274897	ARC2 (includes EG:11847)	-1.906	4.43E-07
				A_51_P274726	PTX3	-1.906	2.77E-07
				A_51_P407726	ZAN	-1.898	7.48E-04
				A_51_P270904	PLEKHS1	-1.894	5.57E-06
				A_52_P627816	TGM1	-1.894	2.40E-06
				A_51_P319460	OSMR	-1.893	1.55E-09
				A_51_P436652	CCL7	-1.883	6.57E-07
				A_52_P607060	Nebl	-1.883	8.78E-09
				A_51_P289796	ARL4C	-1.881	2.49E-09
				A_52_P518997	EPHA2	-1.877	5.69E-09
				A_51_P231557	753042BD23Rik	-1.874	1.33E-08
				A_52_P484194	IL1RL1	-1.872	7.53E-08
				A_51_P174633	MYL7	-1.867	3.17E-07
				A_51_P217463	LOC101928123	-1.866	5.14E-06
				A_51_P246653	CLECT7A	-1.863	2.58E-07
				A_51_P258690	SCRG1	-1.861	8.57E-08
				A_51_P263187	CXL2	-1.859	1.69E-07
				A_51_P252988	ISX	-1.859	3.06E-08
				A_51_P295034	KLK3	-1.859	2.62E-05
				A_52_P181116	CCL24	-1.858	4.61E-09
				A_51_P364600	FYB	-1.853	1.83E-09
				A_51_P326191	SerpinA3g (includes others)	-1.852	7.19E-06
				A_52_P811371	E330019L1Rik	-1.851	2.93E-05
				A_52_P300344	PM2D01	-1.848	9.43E-06
				A_52_P230181	PGYL	-1.845	2.96E-03
				A_52_P688951	IFNG (includes EG:15978)	-1.843	2.42E-08
				A_51_P422389	ODF3B	-1.842	6.35E-06
				A_52_P241818	HT	-1.838	1.98E-06
				A_52_P446364	LGII	-1.835	7.29E-06
				A_52_P253459	CD44	-1.834	1.47E-09
				A_51_P257457	LTPB2	-1.833	9.40E-06
				A_51_P210956	VCAM1	-1.828	7.08E-09
				A_51_P111192	CYP2D6	-1.827	5.77E-08
				A_52_P218058	CLEC5A	-1.825	2.18E-09
				A_51_P171616	WNT10A	-1.823	1.58E-08
				A_51_P238539	EN03	-1.823	3.84E-08
				A_51_P464394	KLB	-1.817	4.01E-05
				A_51_P390804	WISP2	-1.815	3.85E-07
				A_51_P464918	MEFV	-1.814	8.31E-07
				A_51_P393518	HMCN1	-1.814	2.31E-07
				A_52_P185162	SLC26A3	-1.811	9.66E-06
				A_51_P480337	TMEM190	-1.808	2.20E-08
				A_51_P180201	TMEM190	-1.805	9.16E-06
				A_51_P180201	TMEM190	-1.805	1.68E-07
				A_51_P417077	VILI	-1.805	7.85E-09
				A_51_P106000	KUA1211	-1.805	6.51E-05
				A_52_P483336	MSA1	-1.800	4.26E-10
				A_52_P106222	EMB	-1.798	1.03E-05
				A_51_P303791	9530076L18	-1.798	3.09E-07
				A_51_P190740	ADSSL1	-1.793	1.11E-06
				A_51_P245558	SCNN1G	-1.792	1.52E-08
				A_51_P184484	MMP13	-1.792	2.09E-06
				A_51_P496253	SLC6A4	-1.789	5.12E-08
				A_51_P516870	ITM2A	-1.785	7.36E-08
				A_51_P484020	CD300c	-1.785	2.39E-08
				A_51_P408343	II204 (includes others)	-1.785	3.89E-08
				A_51_P212426	CFB3	-1.784	3.89E-08
				A_51_P240837	DMNL2	-1.778	3.70E-08
				A_51_P408506	ICAM1	-1.778	1.18E-07
				A_51_P510891	AFP	-1.776	7.86E-05
				A_51_P236049	KRT33B	-1.770	1.99E-07
				A_52_P231729	HLA-B	-1.770	5.41E-08
				A_51_P231341	SULT1A3/SULT1A4	-1.769	4.01E-07
				A_52_P619192	RUND38	-1.768	2.24E-07
				A_51_P468482	ENTPD8	-1.766	4.62E-09

A_52_P100926	IL1A	interleukin 1, alpha	-1.760	4.00E-06
A_51_P161946	C17orf96	chromosome 17 open reading frame 96	-1.754	4.35E-07
A_52_P263680	ACER1	alkaline ceramidase 1	1.749	9.52E-05
A_51_P201804	ST3GAL1	ST3 beta-galactoside alpha-2,3-sialyltransferase 1	-1.748	5.64E-08
A_52_P266913	CNNM1	cyclin M1	1.748	1.62E-05
A_51_P173043	EGR2	early growth response 2	-1.747	4.90E-10
A_51_P488196	BMPER	BMP binding endothelial regulator	-1.744	4.69E-08
A_51_P268831	SLC2A6	solute carrier family 2 (facilitated glucose transporter), member 6	-1.734	2.55E-07
A_51_P122321	IL33	interleukin 33	-1.732	4.41E-06
A_51_P123630	IRG1	immunoresponsive 1 homolog (mouse)	-1.729	7.67E-03
A_52_P851352	IGFBP3	insulin-like growth factor binding protein 3	-1.728	2.43E-07
A_52_P242539	RETNLB	resistin like beta	-1.727	5.78E-03
A_52_P680827	ABAT	4-aminobutyrate aminotransferase	1.721	6.58E-07
A_51_P268834	DCLRE1A	claudin 1	-1.720	1.57E-07
A_52_P268833	DEFB103A/DEFB103B	defensin, beta 103B	-1.719	2.30E-06
A_51_P174961	F10	coagulation factor X	-1.712	2.98E-06
A_51_P461665	CXCL9	chemokine (C-X-C motif) ligand 9	-1.710	2.39E-07
A_51_P240857	CHPT1	choline phosphotransferase 1	1.710	7.59E-09
A_51_P464293	9130024f11Rik	RIKEN cDNA 9130024f11 gene	1.704	2.27E-07
A_51_P388260	TPPP	tubulin polymerization promoting protein	1.705	4.43E-07
A_51_P204486	EXOC3L4	exocyst complex component 3-like 4	-1.700	3.33E-04
A_51_P366448	GLODS	glyoxalase domain containing 5	1.699	7.33E-08
A_51_P511270	POU3F1	POU class 3 homeobox 1	-1.697	2.83E-06
A_51_P455326	SELE (includes EG-20339)	selectin E	-1.689	4.15E-08
A_52_P32453	1110020a21Rik	RIKEN cDNA 1110020a21 gene	1.689	3.50E-08
A_52_P504254	ADTRP	androgen-dependent TFP1-regulating protein	1.688	1.53E-05
A_51_P216524	TPEB	tumor protein p63	-1.687	3.92E-05
A_51_P420547	LGALS3	chloride intracellular channel 5	1.686	2.91E-07
A_51_P261955	DAPL1	2'-O-riboflavine synthetase 2, 69/71kDa	-1.684	6.40E-07
A_52_P161488	CLEC4E	death associated protein-like 1	-1.683	1.48E-09
A_51_P452768	CYP4F12	C-type lectin domain family 4, member E	-1.681	9.26E-05
A_52_P236664	UNC93A	cytochrome P450, family 4, subfamily F, polypeptide 12	1.680	4.29E-07
A_51_P461025	HSD11B2	unc-93 homolog A (C. elegans)	-1.673	1.06E-08
A_52_P485659	SLC38A4	hydroxysteroid (11-beta) dehydrogenase 2	1.673	1.17E-05
A_51_P304933	LOX	solute carrier family 38, member 4	-1.671	4.63E-05
A_51_P316902	SVOPL	lysyl oxidase	-1.666	3.88E-09
A_52_P578732	CCR5	SVOP-like	1.666	2.98E-06
A_51_P260312	CSGALNACT1	chemokine (C-C motif) receptor 5 (gene/pseudogene)	-1.662	4.92E-07
A_52_P63232	TFCP2L1	chondroitin sulfate N-acetylgalactosaminyltransferase 1	-1.657	1.06E-08
A_51_P436567	Ces2b/Ces2c	transcription factor CP2-like 1	1.651	1.07E-06
A_52_P423811	Ces2c	carboxylesterase 2C	1.650	4.05E-07
A_52_P261004	PRR15	cytochrome c oxidase, subunit VIIb	1.650	2.30E-07
A_51_P268673	2210016H18Rik	protein rich in proline	-1.648	2.40E-08
A_51_P468293	FAM84A	RIKEN cDNA 2210016H18 gene	1.644	4.07E-07
A_52_P247176	NAT8L	family with sequence similarity 84, member A	-1.643	1.06E-08
A_51_P132185	Sif1 (includes others)	N-acetyltransferase 8-like (GCNS-related, putative)	-1.642	9.65E-08
A_52_P35195	GPBP1L1	stefin A1	-1.642	7.49E-07
A_51_P438853	SERPINB11	GC-rich promoter binding protein 1-like 1	-1.639	2.42E-07
A_51_P117581	CABLES1	serpin peptidase inhibitor, clade B (ovalbumin), member 11 (gene/pseudogene)	-1.630	7.90E-07
A_51_P278368	DNER	Cdk5 and Abi enzyme substrate 1	1.630	2.26E-09
A_51_P340170	IL20RA	delta/notch-like EGF repeat containing	1.624	1.77E-07
A_51_P352875	LCP2	interleukin 20 receptor, alpha	-1.624	2.08E-08
A_51_P305896	CFTR	lymphocyte cytosolic protein 2 (SH2 domain containing leukocyte protein of 76kDa)	-1.623	8.28E-08
A_51_P482121	EDN2	cystic fibrosis transmembrane conductance regulator (ATP-binding cassette sub-family C, member 7)	1.622	2.27E-06
A_51_P282930	RORC	endothelin 2	1.611	2.72E-08
A_52_P261777	ESR1	RXR-RAR orphan receptor C	1.610	9.53E-07
A_51_P234938	KRT75	estrogen receptor 1	-1.612	1.47E-07
A_51_P256473	ORSK1	keratin 75	-1.612	7.47E-09
A_51_P263471	CLEC6A	olfactory receptor, family 5, subfamily K, member 1	1.609	1.34E-03
A_52_P507214	MMP9	C-type lectin domain family 6, member A	-1.604	1.06E-09
A_51_P309307	BCL2L15	matrix metalloproteinase 9 (gelatinase B, 92kDa gelatinase, 92kDa type IV collagenase)	-1.604	7.59E-07
A_51_P287925	ZC3H12A	BCL2-like 15	-1.603	1.62E-05
A_51_P255395	COL7A1	zinc-finger CCCH-type containing 12A	-1.602	2.13E-07
		collagen, type VII, alpha 1	-1.601	4.44E-08

Supplementary Table 4: Gene ontology analysis on the differential Nod2-dependent response to antibiotics treatment

Category	Functions	Functions Annotation	p-Value	-Log(P-value)	Molecules	# Molecules
Cellular Movement	infiltration	infiltration by neutrophils	9.54E-11	10.02045163	CFTR,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MMP8,PTGS2,SELL	9
Hematological System Development and Function	infiltration	infiltration by neutrophils	9.54E-11	10.02045163	CFTR,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MMP8,PTGS2,SELL	9
Immune Cell Trafficking	infiltration	infiltration by neutrophils	9.54E-11	10.02045163	CFTR,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MMP8,PTGS2,SELL	9
Inflammatory Response	infiltration	infiltration by neutrophils	9.54E-11	10.02045163	CFTR,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MMP8,PTGS2,SELL	9
Cellular Movement	cell movement	cell movement of granulocytes	4.30E-10	9.366531544	CAMP (includes EG:12796),CFTR,Ch3/3(Ch34,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MMP8,PTGS2,SELL	12
Hematological System Development and Function	cell movement	cell movement of granulocytes	4.30E-10	9.366531544	CAMP (includes EG:12796),CFTR,Ch3/3(Ch34,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MMP8,PTGS2,SELL	12
Immune Cell Trafficking	cell movement	cell movement of granulocytes	4.30E-10	9.366531544	CAMP (includes EG:12796),CFTR,Ch3/3(Ch34,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MMP8,PTGS2,SELL	12
Cellular Movement	cell movement	cell movement of myeloid cells	3.36E-09	8.473660723	CAMP (includes EG:12796),CFTR,Ch3/3(Ch34,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MMP8,PTGS2,PTPRC,SELL	13
Hematological System Development and Function	cell movement	cell movement of myeloid cells	3.36E-09	8.473660723	CAMP (includes EG:12796),CFTR,Ch3/3(Ch34,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MMP8,PTGS2,PTPRC,SELL	13
Immune Cell Trafficking	cell movement	cell movement of myeloid cells	3.36E-09	8.473660723	CAMP (includes EG:12796),CFTR,Ch3/3(Ch34,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MMP8,PTGS2,PTPRC,SELL	13
Cellular Movement	cell movement	cell movement of neutrophils	6.80E-09	8.167491087	CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MMP8,PTGS2,SELL	10
Hematological System Development and Function	cell movement	cell movement of neutrophils	6.80E-09	8.167491087	CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MMP8,PTGS2,SELL	10
Immune Cell Trafficking	cell movement	cell movement of neutrophils	6.80E-09	8.167491087	CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MMP8,PTGS2,SELL	10
Inflammatory Response	cell movement	cell movement of neutrophils	6.80E-09	8.167491087	CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MMP8,PTGS2,SELL	10
Infectious Disease	infection	infection of lung	1.09E-08		CAMP (Includes EG:12796),CXCL10,CXCL9,IFNG (includes EG:15978),IL6	5
Respiratory Disease	infection	infection of lung	1.09E-08		CAMP (Includes EG:12796),CXCL10,CXCL9,IFNG (includes EG:15978),IL6	5
Cellular Movement	recruitment	recruitment of cells	2.33E-08		APOB,CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,PTGS2,SELL	10
Cell-To-Cell Signaling and Interaction	recruitment	recruitment of cells	2.33E-08		APOB,CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,PTGS2,SELL	10
Antimicrobial Response	clearance	clearance of virus	2.52E-08		APOB,CAMP (Includes EG:12796),CFTR,Ch3/3(Ch34,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,PTPRC,SELL,ZC3H12A	5
Cellular Movement	migration	leukocyte migration	2.54E-08		APOB,CAMP (Includes EG:12796),CFTR,Ch3/3(Ch34,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,PTPRC,SELL,ZC3H12A	15
Immune Cell Trafficking	migration	leukocyte migration	2.54E-08		APOB,CAMP (Includes EG:12796),CFTR,Ch3/3(Ch34,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,PTPRC,SELL,ZC3H12A	15
Organismal Survival	survival	survival of organism	2.57E-08		CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MARCO,MMP8,PTGS2,PTPRC,ZC3H12A	13
Cellular Movement	cell movement	cell movement of leukocytes	3.28E-08		CAMP (Includes EG:12796),CFTR,Ch3/3(Ch34,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,PTPRC,SELL,ZC3H12A	14
Hematological System Development and Function	cell movement	cell movement of leukocytes	3.28E-08		CAMP (Includes EG:12796),CFTR,Ch3/3(Ch34,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,PTPRC,SELL,ZC3H12A	14
Immune Cell Trafficking	cell movement	cell movement of leukocytes	3.28E-08		CAMP (Includes EG:12796),CFTR,Ch3/3(Ch34,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,PTPRC,SELL,ZC3H12A	14
Inflammatory Response	inflammation	inflammation of lung	5.97E-08		CAMP (Includes EG:12796),CFTR,Ch3/3(Ch34,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MSA1,PTGS2,PTPRC,SELL	10
Respiratory Disease	inflammation	inflammation of lung	5.97E-08		CAMP (Includes EG:12796),CFTR,Ch3/3(Ch34,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MSA1,PTGS2,PTPRC,SELL	10
Cellular Movement	infiltration	infiltration of leukocytes	7.19E-08		CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,SELL,ZC3H12A	10
Hematological System Development and Function	infiltration	infiltration of leukocytes	7.19E-08		CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,SELL,ZC3H12A	10
Immune Cell Trafficking	infiltration	infiltration of leukocytes	7.19E-08		CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,SELL,ZC3H12A	10
Cellular Movement	recruitment	recruitment of phagocytes	1.19E-07		APOB,CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,SELL	8
Hematological System Development and Function	recruitment	recruitment of phagocytes	1.19E-07		APOB,CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,SELL	8
Immune Cell Trafficking	recruitment	recruitment of phagocytes	1.19E-07		APOB,CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,SELL	8
Inflammatory Response	recruitment	recruitment of phagocytes	1.19E-07		APOB,CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,SELL	8
Cell-To-Cell Signaling and Interaction	recruitment	recruitment of phagocytes	1.19E-07		APOB,CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,SELL	8
Cellular Movement	recruitment	recruitment of phagocytes	1.19E-07		APOB,CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,SELL	8
Hematological System Development and Function	recruitment	recruitment of phagocytes	1.19E-07		APOB,CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,SELL	8
Immune Cell Trafficking	recruitment	recruitment of phagocytes	1.19E-07		APOB,CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,SELL	8
Connective Tissue Disorders	ankylosing spondylitis	ankylosing spondylitis	1.90E-07		CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (Includes EG:15978),IL6,SELL	4
Inflammatory Disease	ankylosing spondylitis	ankylosing spondylitis	1.90E-07		CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (Includes EG:15978),IL6,SELL	4
Skeletal and Muscular Disorders	ankylosing spondylitis	ankylosing spondylitis	1.90E-07		CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (Includes EG:15978),IL6,SELL	4
Hematological Disease	toxemia	toxemia	2.07E-07		CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (Includes EG:15978),IL6,SELL	6
Infectious Disease	infection	Bacterial Infection	2.52E-07		CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,SELL	10
Energy Production	oxidation	oxidation of lipid	2.54E-07		AADAC,CYP2C19,FABP2,HAO2,HPGD,ILS,PPARGC1A,PTGS2	8
Lipid Metabolism	oxidation	oxidation of lipid	2.54E-07		AADAC,CYP2C19,FABP2,HAO2,HPGD,ILS,PPARGC1A,PTGS2	8
Small Molecule Biochemistry	oxidation	oxidation of lipid	2.54E-07		AADAC,CYP2C19,FABP2,HAO2,HPGD,ILS,PPARGC1A,PTGS2	8
Cellular Movement	cell movement	cell movement of phagocytes	4.01E-07		CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,SELL	11
Hematological System Development and Function	cell movement	cell movement of phagocytes	4.01E-07		CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,SELL	11
Immune Cell Trafficking	cell movement	cell movement of phagocytes	4.01E-07		CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,SELL	11
Inflammatory Response	cell movement	cell movement of phagocytes	4.01E-07		CAMP (Includes EG:12796),CFTR,CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,SELL	11
Cellular Movement	cell movement	cell movement of neutrophils	4.00E-07		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,SELL	6
Hematological System Development and Function	cell movement	cell movement of neutrophils	4.00E-07		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,SELL	6
Immune Cell Trafficking	cell movement	cell movement of neutrophils	4.00E-07		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,SELL	6
Hypersensitivity Response	cell movement	cell movement of neutrophils	4.00E-07		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,MMP8,PTGS2,SELL	6
Energy Production	oxidation	oxidation of fatty acid	4.80E-07		AADAC,FABP2,HAO2,HPGD,IL6,PPARGC1A,PTGS2	7
Lipid Metabolism	oxidation	oxidation of fatty acid	4.80E-07		AADAC,FABP2,HAO2,HPGD,IL6,PPARGC1A,PTGS2	7
Small Molecule Biochemistry	oxidation	oxidation of fatty acid	4.80E-07		AADAC,FABP2,HAO2,HPGD,IL6,PPARGC1A,PTGS2	7
Hematological System Development and Function	activation	activation of myeloid cells	5.48E-07		CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (Includes EG:15978),IL6,PTGS2,SELL	8
Immune Cell Trafficking	activation	activation of myeloid cells	5.48E-07		CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (Includes EG:15978),IL6,PTGS2,SELL	8
Inflammatory Response	activation	activation of myeloid cells	5.48E-07		CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (Includes EG:15978),IL6,PTGS2,SELL	8
Cell-To-Cell Signaling and Interaction	activation	activation of myeloid cells	5.48E-07		CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (Includes EG:15978),IL6,PTGS2,SELL	8
Cellular Movement	cell movement	cell movement of mononuclear leukocytes	6.53E-07		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,PTGS2,SELL	10
Hematological System Development and Function	cell movement	cell movement of mononuclear leukocytes	6.53E-07		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,PTGS2,SELL	10
Immune Cell Trafficking	cell movement	cell movement of mononuclear leukocytes	6.53E-07		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,PTGS2,SELL	10
Organismal Survival	binding	binding of phagocytes	7.87E-07		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,PTGS2,SELL	6
Inflammatory Response	binding	binding of phagocytes	7.87E-07		CAMP (Includes EG:12796),CXCL10,CXCR2,IFNG (Includes EG:15978),MARCO,PTPRC,SELL	6
Cell-To-Cell Signaling and Interaction	binding	binding of phagocytes	7.87E-07		CAMP (Includes EG:12796),CXCL10,CXCR2,IFNG (Includes EG:15978),MARCO,PTPRC,SELL	6
Cellular Movement	recruitment	recruitment of granulocytes	9.25E-07		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,SELL	7
Hematological System Development and Function	recruitment	recruitment of granulocytes	9.25E-07		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,SELL	7
Immune Cell Trafficking	recruitment	recruitment of granulocytes	9.25E-07		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,SELL	7
Cell-To-Cell Signaling and Interaction	recruitment	recruitment of granulocytes	9.25E-07		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,SELL	7
Organismal Survival	survival	survival of Bacillus anthracis	9.25E-07		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,SELL	3
Cell Death and Survival	survival	survival of Bacillus anthracis	9.25E-07		CAMP (Includes EG:12796),CXCL10,CXCR2,IFNG (Includes EG:15978),MARCO,PTPRC,SELL	6
Organismal Survival	mortality	mortality	9.49E-07		CAMP (Includes EG:12796),CXCL10,CXCR2,IFNG (Includes EG:15978),MARCO,PTPRC,SELL	7
Cellular Movement	cell movement	cell movement of T lymphocytes	1.02E-06		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,SELL	7
Hematological System Development and Function	cell movement	cell movement of T lymphocytes	1.02E-06		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,SELL	7
Immune Cell Trafficking	cell movement	cell movement of T lymphocytes	1.02E-06		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,SELL	7
Cell-mediated Immune Response	cell movement	cell movement of T lymphocytes	1.02E-06		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),IL6,SELL	7
Cellular Movement	chemotaxis	chemotaxis of granulocytes	1.02E-06		CAMP (Includes EG:12796),Ch3/3(Ch34,CXCL10,CXCR2,IFNG (Includes EG:15978),SELL	7

Hematological System Development and Function	chemotaxis	1.02E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,_CXCL9,_CXCR2,IFNG (includes EG:15978),SELL
Immune Cell Trafficking	chemotaxis	1.02E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,_CXCL9,_CXCR2,IFNG (includes EG:15978),SELL
Inflammatory Response	chemotaxis	1.02E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,_CXCL9,_CXCR2,IFNG (includes EG:15978),SELL
Cellular Movement	cell movement	1.03E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,_CXCL9,IFNG (includes EG:15978),IL6,PTGS2,SELL,ZC3H12A
Hematological System Development and Function	cell movement	1.03E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,_CXCL9,IFNG (includes EG:15978),IL6,PTGS2,SELL,ZC3H12A
Immune Cell Trafficking	cell movement	1.03E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,_CXCL9,IFNG (includes EG:15978),IL6,PTGS2,SELL,ZC3H12A
Hematological System Development and Function	binding	1.05E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,_CXCL9,CXCR2,IFNG (includes EG:15978),MARCO,PTPRC,SELL
Cell-To-Cell Signaling and Interaction	binding	1.05E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,_CXCL9,CXCR2,IFNG (includes EG:15978),MARCO,PTPRC,SELL
Inflammatory Response	inflammation	1.06E-06	CXCL10,IFNG (includes EG:15978),PTGS2,PTPRC
Dermatological Diseases and Conditions	psoriasis	1.13E-06	CAMP (includes EG:12796),CSTA,CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,LCE3D,PM20D1,PTGS2,PTPRC
Cell-To-Cell Signaling and Interaction	binding	1.23E-06	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Inflammatory Response	inflammatory response	1.26E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MARCO,MP8,PTGS2,PTPRC,SELL
Cardiovascular System Development and Function	systolic pressure	1.38E-06	ACE2 (includes EG:302688),IL6,PTGS2
Cellular Movement	chemotaxis	1.43E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),PTPRC,SELL
Hematological System Development and Function	chemotaxis	1.43E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),PTPRC,SELL
Immune Cell Trafficking	chemotaxis	1.43E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),PTPRC,SELL
Inflammatory Response	chemotaxis	1.43E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),PTPRC,SELL
Lipid Metabolism	conversion	1.48E-06	Akr1c14,CFTR,CYP2C19,HPGD,IFNG (includes EG:15978),IL6,PTGS2
Small Molecule Biochemistry	conversion	1.48E-06	Akr1c14,CFTR,CYP2C19,HPGD,IFNG (includes EG:15978),IL6,PTGS2
Humoral Immune Response	quantity	1.64E-06	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,PTGS2
Infectious Disease	quantity	1.64E-06	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,PTGS2
Cellular Movement	tuberculosis	1.88E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,CXCL9,IFNG (includes EG:15978),IL6
Hematological System Development and Function	chemotaxis	1.91E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,PTGS2,PTPRC,SELL
Immune Cell Trafficking	activation	2.35E-06	CAMP (includes EG:12796),CXCL10,CXCL9,FOXA2,IFNG (includes EG:15978),IL6,PTGS2,SELL
Inflammatory Response	activation	2.35E-06	CAMP (includes EG:12796),CXCL10,CXCL9,FOXA2,IFNG (includes EG:15978),IL6,PTGS2,SELL
Cell-To-Cell Signaling and Interaction	activation	2.35E-06	CAMP (includes EG:12796),CXCL10,CXCL9,FOXA2,IFNG (includes EG:15978),IL6,PTGS2,SELL
Inflammation	inflammation	2.68E-06	CAMP (includes EG:12796),CFTR,CYP2C19,HPGD,IFNG (includes EG:15978),IL6,MS4A1,PTGS2,PTPRC,SELL
Nutrition Disease	body mass index	2.69E-06	ACSM3,APOB,CFTR,CYP2C19,HPGD,IFNG (includes EG:15978),IL6,SELL
Cellular Movement	chemotaxis	3.12E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,CXCL9
Hematological System Development and Function	chemotaxis	3.12E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,CXCL9
Immune Cell Trafficking	chemotaxis	3.12E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,CXCL9
Inflammatory Response	chemotaxis	3.12E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,CXCL9
Hypersensitivity Response	chemotaxis	3.12E-06	CAMP (includes EG:12796),Ch3(3)Ch3(4)_CXCL10,CXCL9
Connective Tissue Development and Function	formation	3.61E-06	IFNG (includes EG:15978),IL6,SELL
Tissue Development	formation	3.61E-06	IFNG (includes EG:15978),IL6,SELL
Hematological System Development and Function	function	3.65E-06	CXCL10,CXCR2,IFNG (includes EG:15978),IL6,MARCO,PTGS2,SELL
Cellular Function and Maintenance	function	3.65E-06	CXCL10,CXCR2,IFNG (includes EG:15978),IL6,MARCO,PTGS2,SELL
Cell Death and Survival	cell death	3.82E-06	CAMP (includes EG:12796),IFNG (includes EG:15978),PTPRC,ZC3H12A
Cellular Movement	recruitment	4.46E-06	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Hematological System Development and Function	recruitment	4.46E-06	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Immune Cell Trafficking	recruitment	4.46E-06	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Inflammatory Response	recruitment	4.46E-06	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Cell-To-Cell Signaling and Interaction	recruitment	4.46E-06	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Organismal Functions	healing	4.62E-06	CAMP (includes EG:12796),CXCL10,CXCL9,IFNG (includes EG:15978),IL6,SELL
Tissue Morphology	healing	4.62E-06	CXCR2,IFNG (includes EG:15978),IL6,PTGS2,SCNN1G,SELL
Organismal Injury and Abnormalities	wound	4.78E-06	CAMP (includes EG:12796),CXCR2,IFNG (includes EG:15978),IL6,PTGS2,SELL
Hematological System Development and Function	accumulation	4.92E-06	CXCL10,CXCL9,IFNG (includes EG:15978),MMP8,SELL
Immune Cell Trafficking	accumulation	4.92E-06	CXCL10,CXCL9,IFNG (includes EG:15978),MMP8,SELL
Inflammatory Response	accumulation	4.92E-06	CXCL10,CXCL9,IFNG (includes EG:15978),MMP8,SELL
Tissue Development	accumulation	4.92E-06	CXCL10,CXCL9,IFNG (includes EG:15978),MMP8,SELL
Cellular Movement	migration	5.10E-06	CXCL10,CXCL9,IFNG (includes EG:15978),SELL
Hematological System Development and Function	migration	5.10E-06	CXCL10,CXCL9,IFNG (includes EG:15978),SELL
Immune Cell Trafficking	migration	5.10E-06	CXCL10,CXCL9,IFNG (includes EG:15978),SELL
Innate Immune Response	migration	5.10E-06	CXCL10,CXCL9,IFNG (includes EG:15978),SELL
Immunologic Disease	hypersensitive reaction	5.10E-06	CXCL10,CXCL9,IFNG (includes EG:15978),SELL
Lipid Metabolism	concentration	5.53E-06	CAMP (includes EG:12796),CFTR,CYCLICAMP (includes EG:12796),CFTR,FABP2,FOXA2,HPGD,IFNG (includes EG:15978),IL6,PPARGC1A,PTGS2
Small Molecule Biochemistry	concentration	5.53E-06	AADAC,ACER1,APOB,CAMP (includes EG:12796),CFTR,FABP2,FOXA2,HPGD,IFNG (includes EG:15978),IL6,PPARGC1A,PTGS2
Molecular Transport	concentration	5.53E-06	AADAC,ACER1,APOB,CAMP (includes EG:12796),CFTR,FABP2,FOXA2,HPGD,IFNG (includes EG:15978),IL6,PPARGC1A,PTGS2
Cellular Movement	cell movement	5.65E-06	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Hematological System Development and Function	cell movement	5.65E-06	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Immune Cell Trafficking	cell movement	5.65E-06	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Inflammatory Response	cell movement	5.65E-06	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Infectious Disease	infection	5.67E-06	CAMP (includes EG:12796),CFTR,CYCLICAMP (includes EG:15978),IL6,SELL
Infectious Disease	anthrax	5.95E-06	CAMP (includes EG:12796),CFTR,CYCLICAMP (includes EG:15978),IL6,SELL
Dermatological Diseases and Conditions	chronic large plaque psoriasis	5.95E-06	CXCL10,CXCL9,IFNG (includes EG:15978)
Dermatological Diseases and Conditions	chronic small plaque psoriasis	5.95E-06	CXCL10,CXCL9,IFNG (includes EG:15978)
Organismal Injury and Abnormalities	formation	5.95E-06	CXCL10,CXCL9,IFNG (includes EG:15978)
Cardiovascular Disease	formation	5.95E-06	CXCL10,CXCL9,IFNG (includes EG:15978)
Cellular Movement	transmigration	6.66E-06	CAMP (includes EG:12796),CFTR,CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MARCO,MMP8
Hematological System Development and Function	transmigration	6.66E-06	CAMP (includes EG:12796),CFTR,CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MARCO,MMP8
Gene Expression	transmigration	6.66E-06	CAMP (includes EG:12796),CFTR,CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MARCO,MMP8
Hematological System Development and Function	function	6.69E-06	IFNG (includes EG:15978),IL6
Cellular Function and Maintenance	function	6.87E-06	CXCL10,IFNG (includes EG:15978),IL6,MARCO,PTGS2,SELL
Humoral Immune Response	quantity	6.96E-06	CXCL10,IFNG (includes EG:15978),IL6,PTGS2,ZC3H12A
Protein Synthesis	quantity	6.96E-06	CXCL10,IFNG (includes EG:15978),IL6,PTGS2,ZC3H12A
Cellular Movement	cell movement	7.25E-06	CAMP (includes EG:12796),CFTR,CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978)
Hematological System Development and Function	cell movement	7.25E-06	CAMP (includes EG:12796),CFTR,CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978)
Immune Cell Trafficking	cell movement	7.25E-06	CAMP (includes EG:12796),CFTR,CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978)
Inflammatory Response	cell movement	7.25E-06	CAMP (includes EG:12796),CFTR,CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978)
Hypersensitivity Response	cell movement	7.25E-06	CAMP (includes EG:12796),CFTR,CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978)
Cellular Movement	recruitment	8.01E-06	CAMP (includes EG:12796),CFTR,CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978)
Hematological System Development and Function	recruitment	8.01E-06	CAMP (includes EG:12796),CFTR,CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978)
Immune Cell Trafficking	recruitment	8.01E-06	CAMP (includes EG:12796),CFTR,CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978)
Cell-To-Cell Signaling and Interaction	recruitment	8.01E-06	CAMP (includes EG:12796),CFTR,CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978)
Hematological System Development and Function	adhesion	8.85E-06	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MARCO,PTPRC,SELL
Immune Cell Trafficking	adhesion	8.85E-06	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MARCO,PTPRC,SELL
Cell-To-Cell Signaling and Interaction	adhesion	8.85E-06	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MARCO,PTPRC,SELL

Tissue Development	adhesion	adhesion of immune cells	8.85E-06	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MARCO,PTPRC,SELL
Cellular Movement	chemotaxis	chemotaxis of endothelial cell lines	9.11E-06	CXCL10,CXCL9,CXCR2
Cardiovascular System Development and Function	chemotaxis	chemotaxis of endothelial cell lines	9.11E-06	CAMP (Includes EG:12796),IL6,PTGS2
Lipid Metabolism	induction	induction of prostaglandin	9.11E-06	CAMP (Includes EG:12796),IL6,PTGS2
Small Molecule Biochemistry	induction	induction of prostaglandin	9.11E-06	CAMP (Includes EG:12796),IL6,PTGS2
Inflammatory Response	inflammation	inflammation of peritoneum	9.11E-06	CFTR,PTGS2,SELL
Hematological Disease	chronic B-cell leukemia	chronic B-cell leukemia	9.18E-06	CXCL9,IFNG (includes EG:15978),IL6,MSA1,SELL
Immunological Disease	chronic B-cell leukemia	chronic B-cell leukemia	9.18E-06	CXCL9,IFNG (includes EG:15978),IL6,MSA1,SELL
Cancer	chronic B-cell leukemia	chronic B-cell leukemia	9.18E-06	CXCL9,IFNG (includes EG:15978),IL6,MSA1,SELL
Cellular Movement	migration	migration of neutrophils	9.18E-06	CAMP (Includes EG:12796),CXCR2,IFNG (includes EG:15978),IL6,SELL
Hematological System Development and Function	migration	migration of neutrophils	9.18E-06	CAMP (Includes EG:12796),CXCR2,IFNG (includes EG:15978),IL6,SELL
Immune Cell Trafficking	migration	migration of neutrophils	9.18E-06	CAMP (Includes EG:12796),CXCR2,IFNG (includes EG:15978),IL6,SELL
Inflammatory Response	migration	migration of neutrophils	9.18E-06	CAMP (Includes EG:12796),CXCR2,IFNG (includes EG:15978),IL6,SELL
Cellular Movement	migration	migration of neutrophils	1.03E-05	CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Hematological System Development and Function	migration	migration of neutrophils	1.03E-05	CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Immune Cell Trafficking	migration	migration of neutrophils	1.03E-05	CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Inflammatory Response	migration	migration of neutrophils	1.03E-05	CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Lipid Metabolism	metabolism	fatty acid metabolism	1.03E-05	ACSM3,APOB,CFTR,FABP2,HPG6,IFNG (includes EG:15978),IL6,MSA1,PPARGC1A,PTGS2
Small Molecule Biochemistry	metabolism	fatty acid metabolism	1.03E-05	ACSM3,APOB,CFTR,FABP2,HPG6,IFNG (includes EG:15978),IL6,MSA1,PPARGC1A,PTGS2
Cellular Movement	homing	homing of T lymphocytes	1.05E-05	CAMP (Includes EG:12796),Ch3/Ch34,CXCL10,CXCL9,SELL
Hematological System Development and Function	homing	homing of T lymphocytes	1.05E-05	CAMP (Includes EG:12796),Ch3/Ch34,CXCL10,CXCL9,SELL
Immune Cell Trafficking	homing	homing of T lymphocytes	1.05E-05	CAMP (Includes EG:12796),Ch3/Ch34,CXCL10,CXCL9,SELL
Cell-mediated Immune Response	homing	homing of T lymphocytes	1.05E-05	CAMP (Includes EG:12796),Ch3/Ch34,CXCL10,CXCL9,SELL
Cardiovascular Disease	abdominal aortic aneurysm	abdominal aortic aneurysm	1.10E-05	CXCL10,IFNG (includes EG:15978),PTGS2
Hematological Disease	follicular non-Hodgkin's disease	follicular non-Hodgkin's disease	1.10E-05	IFNG (Includes EG:15978),IL6,MSA1,SELL
Cancer	follicular non-Hodgkin's disease	follicular non-Hodgkin's disease	1.10E-05	IFNG (Includes EG:15978),IL6,MSA1,SELL
Inflammatory Response	immune response	immune response of phagocytes	1.12E-05	CAMP (Includes EG:12796),CXCL10,IFNG (includes EG:15978),IL6,MARCO,PTPRC
Cell-To-Cell Signaling and Interaction	immune response	immune response of phagocytes	1.12E-05	CAMP (Includes EG:12796),CXCL10,IFNG (includes EG:15978),IL6,MARCO,PTPRC
Lymphatic Tissue Structure and Development	morphology	morphology of spleen	1.20E-05	CXCR2,IFNG (includes EG:15978),MARCO,PTGS2,PTPRC,SELL,ZC3H12A
Organ Morphology	morphology	morphology of spleen	1.20E-05	CXCR2,IFNG (includes EG:15978),MARCO,PTGS2,PTPRC,SELL,ZC3H12A
Inflammatory Disease	dermatitis	Dermatitis	1.23E-05	CAMP (Includes EG:12796),CXCL9,IFNG (includes EG:15978),LCE3D,PPARGC1A,PTGS2,SELL,TRPV3
Dermatological Diseases and Conditions	dermatitis	Dermatitis	1.23E-05	CAMP (Includes EG:12796),CXCL9,IFNG (includes EG:15978),LCE3D,PPARGC1A,PTGS2,SELL,TRPV3
Cell-To-Cell Signaling and Interaction	response	response of myeloid cells	1.25E-05	CAMP (Includes EG:12796),CXCL9,IFNG (includes EG:15978),IL6,MARCO,PTPRC
Hematological System Development and Function	activation	activation of granulocytes	1.30E-05	CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Immune Cell Trafficking	activation	activation of granulocytes	1.30E-05	CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Inflammatory Response	activation	activation of granulocytes	1.30E-05	CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Cell-To-Cell Signaling and Interaction	activation	activation of granulocytes	1.30E-05	CXCL9,CXCR2,IFNG (includes EG:15978),IL6,SELL
Cellular Movement	attraction	attraction of dendritic cells	1.32E-05	CAMP (Includes EG:12796),CXCL10,CXCL9
Immune Cell Trafficking	attraction	attraction of dendritic cells	1.32E-05	CAMP (Includes EG:12796),CXCL10,CXCL9
Cell-To-Cell Signaling and Interaction	attraction	attraction of dendritic cells	1.32E-05	CAMP (Includes EG:12796),CXCL10,CXCL9
Cardiovascular System Development and Function	differentiation	differentiation of vascular endothelial cells	1.32E-05	CXCL10,IFNG (includes EG:15978),PTGS2
Cellular Development	differentiation	differentiation of vascular endothelial cells	1.32E-05	CXCL10,IFNG (includes EG:15978),PTGS2
Cellular Movement	transmigration	transmigration of leukocytes	1.41E-05	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6
Hematological System Development and Function	transmigration	transmigration of leukocytes	1.41E-05	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6
Immune Cell Trafficking	transmigration	transmigration of leukocytes	1.41E-05	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6
Inflammatory Response	transmigration	transmigration of leukocytes	1.41E-05	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6
Cell-To-Cell Signaling and Interaction	transmigration	transmigration of leukocytes	1.41E-05	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6
Tissue Morphology	quantity	quantity of leukocytes	1.43E-05	CAMP (Includes EG:12796),CXCL10,CXCR2,GPR83,IFNG (includes EG:15978),IL6,MMP8,MSA1,PTGS2,PTPRC,SELL,ZC3H12A
Hematological System Development and Function	quantity	quantity of leukocytes	1.43E-05	CAMP (Includes EG:12796),CXCL10,CXCR2,GPR83,IFNG (includes EG:15978),IL6,MMP8,MSA1,PTGS2,PTPRC,SELL,ZC3H12A
Inflammatory Response	phagocytosis	phagocytosis of myeloid cells	1.47E-05	CAMP (Includes EG:12796),CXCL10,IFNG (includes EG:15978),MARCO,PTPRC
Cell-To-Cell Signaling and Interaction	phagocytosis	phagocytosis of myeloid cells	1.47E-05	CAMP (Includes EG:12796),CXCL10,IFNG (includes EG:15978),MARCO,PTPRC
Cellular Function and Maintenance	phagocytosis	phagocytosis of myeloid cells	1.47E-05	CAMP (Includes EG:12796),CXCL10,IFNG (includes EG:15978),MARCO,PTPRC
Inflammatory Response	phagocytosis	phagocytosis of myeloid cells	1.47E-05	CAMP (Includes EG:12796),CXCL10,IFNG (includes EG:15978),MARCO,PTPRC
Cell-To-Cell Signaling and Interaction	phagocytosis	phagocytosis of myeloid cells	1.53E-05	CAMP (Includes EG:12796),CXCL10,IFNG (includes EG:15978),MARCO,PTPRC
Cellular Function and Maintenance	phagocytosis	phagocytosis of myeloid cells	1.53E-05	CAMP (Includes EG:12796),CXCL10,IFNG (includes EG:15978),MARCO,PTPRC
Cell Death and Survival	internalization	internalization of cells	1.53E-05	CAMP (Includes EG:12796),CXCL10,IFNG (includes EG:15978),MARCO,PTPRC
Inflammatory Response	cell death	cell death of lung cancer cell lines	1.66E-05	CAMP (Includes EG:12796),CFTR,IFNG (Includes EG:15978),MARCO,PTPRC
Cell-To-Cell Signaling and Interaction	immunoreponse	immunoreponse of macrophages	1.75E-05	CAMP (Includes EG:12796),FOXA2,IFNG (Includes EG:15978),IL6,MARCO,PTPRC
Cell Death and Survival	immunoreponse	immunoreponse of macrophages	1.88E-05	CAMP (Includes EG:12796),IFNG (Includes EG:15978),IL6,MARCO,PTPRC
Cell Death and Survival	apoptosis	apoptosis of neutrophils	1.90E-05	CAMP (Includes EG:12796),IFNG (Includes EG:15978),IL6,MMP8
Immunological Disease	Graves' ophthalmopathy	Graves' ophthalmopathy	2.00E-05	IL6,MSA1
Endocrine System Disorders	Graves' ophthalmopathy	Graves' ophthalmopathy	2.00E-05	IL6,MSA1
Hereditary Disorder	Graves' ophthalmopathy	Graves' ophthalmopathy	2.00E-05	IL6,MSA1
Metabolic Disease	Graves' ophthalmopathy	Graves' ophthalmopathy	2.00E-05	IL6,MSA1
Ophthalmic Disease	Graves' ophthalmopathy	Graves' ophthalmopathy	2.00E-05	IL6,MSA1
Hematological System Development and Function	aggregation	aggregation of hematopoietic progenitor cells	2.00E-05	CXCL10,CXCL9
Tissue Development	aggregation	aggregation of hematopoietic progenitor cells	2.00E-05	CXCL10,CXCL9
Cellular Development	aggregation	aggregation of hematopoietic progenitor cells	2.00E-05	CXCL10,CXCL9
Cellular Movement	attraction	attraction of Th1 cells	2.00E-05	CXCL10,CXCL9
Hematological System Development and Function	attraction	attraction of Th1 cells	2.00E-05	CXCL10,CXCL9
Immune Cell Trafficking	attraction	attraction of Th1 cells	2.00E-05	CXCL10,CXCL9
Cell-To-Cell Signaling and Interaction	attraction	attraction of Th1 cells	2.00E-05	CXCL10,CXCL9
Cellular Movement	attraction	attraction of plasmacytoid dendritic cells	2.00E-05	CXCL10,CXCL9
Hematopoiesis	attraction	attraction of plasmacytoid dendritic cells	2.00E-05	CXCL10,CXCL9
Cellular Movement	attraction	attraction of plasmacytoid dendritic cells	2.00E-05	CXCL10,CXCL9
Cell Morphology	autophagy	autophagy of monocytes	2.00E-05	CAMP (Includes EG:12796),IFNG (Includes EG:15978)
Cell Morphology	autophagy	autophagy of monocytes	2.00E-05	CAMP (Includes EG:12796),IFNG (Includes EG:15978)
Lipid Metabolism	conversion	conversion of eicosapentaenoic acid	2.00E-05	CFTR,PTGS2
Small Molecule Biochemistry	conversion	conversion of eicosapentaenoic acid	2.00E-05	CFTR,PTGS2
Hematological System Development and Function	polarization	polarization of hematopoietic progenitor cells	2.00E-05	CXCL10,CXCL9
Cell Morphology	polarization	polarization of hematopoietic progenitor cells	2.00E-05	CXCL10,CXCL9
Hematopoiesis	polarization	polarization of hematopoietic progenitor cells	2.00E-05	CXCL10,CXCL9
Cellular Movement	influx	influx of leukocytes	2.02E-05	CXCR2,IFNG (Includes EG:15978),IL6,MARCO
Hematological System Development and Function	influx	influx of leukocytes	2.02E-05	CXCR2,IFNG (Includes EG:15978),IL6,MARCO
Immune Cell Trafficking	influx	influx of leukocytes	2.02E-05	CXCR2,IFNG (Includes EG:15978),IL6,MARCO
Cellular Movement	influx	influx of leukocytes	2.03E-05	CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (Includes EG:15978),IL6,MMP8,MSA1,PTGS2,PTPRC,SELL,SLC7A11
Immunological Disease	systemic autoimmune syndrome	systemic autoimmune syndrome	2.08E-05	CXCR2,IFNG (Includes EG:15978),IL6,MARCO,PTGS2
Renal Disease	damage	damage to lung	2.08E-05	CXCR2,IFNG (Includes EG:15978),IL6,MARCO,PTGS2
Organismal Injury and Abnormalities	damage	damage of lung	2.08E-05	CXCR2,IFNG (Includes EG:15978),IL6,MARCO,PTGS2
Cellular Movement	chemotaxis	chemotaxis of neutroglia	2.15E-05	CXCL10,CXCL9,IL6
Infectious Disease	quantity	quantity of virus	2.16E-05	CXCL9,IFNG (Includes EG:15978),IL6,PTGS2

Cellular Function and Maintenance	function	function of phagocytes	2.30E-05	CXCL10.CXCR2.IFNG (includes EG:15978),IL6, MARCO.PTGS2,SELL	7
Hematological System Development and Function	activation	activation of leukocytes	2.30E-05	CAMP (includes EG:12796),CXCL10.CXCL9.CXCR2,FOXA2,IFNG (includes EG:15978),IL6,PTPS2,PTPRC,SELL	10
Immune Cell Trafficking	activation	activation of leukocytes	2.30E-05	CAMP (includes EG:12796),CXCL10.CXCL9.CXCR2,FOXA2,IFNG (includes EG:15978),IL6,PTGS2,PTPRC,SELL	10
Inflammatory Response	activation	activation of leukocytes	2.39E-05	CAMP (includes EG:12796),CXCL10.CXCL9.CXCR2,FOXA2,IFNG (includes EG:15978),IL6,PTGS2,PTPRC,SELL	10
Cell-To-Cell Signaling and Interaction	activation	activation of leukocytes	2.39E-05	CAMP (includes EG:12796),CXCL10.CXCL9.CXCR2,FOXA2,IFNG (includes EG:15978),IL6,PTGS2,PTPRC,SELL	10
Cell-To-Cell Signaling and Interaction	binding	binding of leukemia cell lines	2.44E-05	CXCR2,IFNG (includes EG:15978),IL6,SELL	4
Cellular Movement	chemotaxis	chemotaxis of embryonic cell lines	2.48E-05	CAMP (includes EG:12796),CXCL10.CXCR2	3
Embryonic Development	chemotaxis	chemotaxis of embryonic cell lines	2.48E-05	CAMP (includes EG:12796),CXCL10.CXCR2	3
Cellular Movement	chemotaxis	chemotaxis of epithelial cell lines	2.48E-05	CAMP (includes EG:12796),CXCL10.CXCR2	3
Hair and Skin Development and Function	chemotaxis	chemotaxis of epithelial cell lines	2.48E-05	CAMP (includes EG:12796),CXCL10.CXCR2	3
Cellular Movement	trafficking	trafficking of T lymphocytes	2.48E-05	CXCL10,IFNG (includes EG:15978),SELL	3
Hematological System Development and Function	trafficking	trafficking of T lymphocytes	2.48E-05	CXCL10,IFNG (includes EG:15978),SELL	3
Immune Cell Trafficking	trafficking	trafficking of T lymphocytes	2.48E-05	CXCL10,IFNG (includes EG:15978),SELL	3
Cell-mediated Immune Response	trafficking	trafficking of T lymphocytes	2.48E-05	CXCL10,IFNG (includes EG:15978),SELL	3
Cell-To-Cell Signaling and Interaction	binding	binding of cells	2.64E-05	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MARCO.PTPRC,SELL	9
Tissue Morphology	abnormal morphology	abnormal morphology of lymph node	2.78E-05	CXCR2,IFNG (includes EG:15978),PTGS2,SELL,ZC3H12A	5
Lymphoid Tissue Structure and Development	abnormal morphology	abnormal morphology of lymph node	2.78E-05	CXCR2,IFNG (includes EG:15978),PTGS2,SELL,ZC3H12A	5
Organ Morphology	abnormal morphology	abnormal morphology	2.78E-05	CXCR2,IFNG (includes EG:15978),PTGS2,SELL,ZC3H12A	5
Cellular Movement	cell rolling	cell rolling of mononuclear leukocytes	2.85E-05	CXCR2,IFNG (includes EG:15978),SELL	3
Hematological System Development and Function	cell rolling	cell rolling of mononuclear leukocytes	2.85E-05	CXCR2,IFNG (includes EG:15978),SELL	3
Immune Cell Trafficking	cell rolling	cell rolling of mononuclear leukocytes	2.85E-05	CXCR2,IFNG (includes EG:15978),SELL	3
Cellular Movement	recruitment	recruitment of inflammatory leukocytes	2.85E-05	CXCL10,IFNG (includes EG:15978),IL6	3
Hematological System Development and Function	recruitment	recruitment of inflammatory leukocytes	2.85E-05	CXCL10,IFNG (includes EG:15978),IL6	3
Immune Cell Trafficking	recruitment	recruitment of inflammatory leukocytes	2.85E-05	CXCL10,IFNG (includes EG:15978),IL6	3
Cell-To-Cell Signaling and Interaction	recruitment	recruitment of inflammatory leukocytes	2.85E-05	CXCL10,IFNG (includes EG:15978),IL6	3
Molecular Transport	quantity	quantity of metal ion	2.92E-05	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,PTGS2,PTPRC,SCNN1G,SELL	9
Cellular Function and Maintenance	function	function of leukocytes	3.03E-05	CXCL10,CXCR2,IFNG (includes EG:15978),IL6,MARCO.MS4A1,PTGS2,PTPRC,SELL	9
Hematological Disease	lymphocytic leukemia	lymphocytic leukemia	3.05E-05	CXCL10,IFNG (includes EG:15978),IL6,MS4A1,PTPRC,SELL	6
Cancer	lymphocytic leukemia	lymphocytic leukemia	3.20E-05	CXCL9,IFNG (includes EG:15978),IL6,MS4A1,PTPRC,SELL	6
Cell-To-Cell Signaling and Interaction	activation	activation of cells	3.23E-05	CAMP (includes EG:12796),CXCL10,IFNG (includes EG:15978),TRPV3	12
Cell-To-Cell Signaling and Interaction	activation	activation of epithelial cells	3.25E-05	CAMP (includes EG:12796),IFNG (includes EG:15978),IL6,PTGS2	3
Organismal Injury and Abnormalities	formation	formation of lesion	3.27E-05	APOB.CTR,CXCL10,IFNG (includes EG:15978),IL6,PTGS2	6
Hematological System Development and Function	adhesion	adhesion of phagocytes	3.42E-05	CXCR2,IFNG (includes EG:15978),IL6,MARCO,SELL	5
Immune Cell Trafficking	adhesion	adhesion of phagocytes	3.42E-05	CXCR2,IFNG (includes EG:15978),IL6,MARCO,SELL	5
Inflammatory Response	adhesion	adhesion of phagocytes	3.42E-05	CXCR2,IFNG (includes EG:15978),IL6,MARCO,SELL	5
Cell-To-Cell Signaling and Interaction	adhesion	adhesion of phagocytes	3.42E-05	CXCR2,IFNG (includes EG:15978),IL6,MARCO,SELL	5
Tissue Development	adhesion	adhesion of phagocytes	3.42E-05	CXCR2,IFNG (includes EG:15978),IL6,MARCO,SELL	5
Hematological System Development and Function	binding	binding of myeloid cells	3.47E-05	CXCL10,CXCR2,IFNG (includes EG:15978),SELL	4
Cell-To-Cell Signaling and Interaction	binding	binding of myeloid cells	3.47E-05	CXCL10,CXCR2,IFNG (includes EG:15978),SELL	4
Hematological System Development and Function	maturity	maturity of T lymphocytes	3.47E-05	IFNG (includes EG:15978),IL6,PTGS2,PTPRC	4
Cellular Development	maturity	maturity of T lymphocytes	3.47E-05	IFNG (includes EG:15978),IL6,PTGS2,PTPRC	4
Hematopoiesis	maturity	maturity of T lymphocytes	3.47E-05	IFNG (includes EG:15978),IL6,PTGS2,PTPRC	4
Cellular Movement	cell movement	cell movement of antigen presenting cells	3.63E-05	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),PTGS2,SELL	7
Hematological System Development and Function	cell movement	cell movement of antigen presenting cells	3.63E-05	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),PTGS2,SELL	7
Immune Cell Trafficking	cell movement	cell movement of antigen presenting cells	3.63E-05	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),PTGS2,SELL	7
Cardiovascular Disease	heart failure	heart failure	3.68E-05	ACE2 (includes EG:302668),HPGD,IFNG (includes EG:15978),IL6,PPARGC1A,PTGS2	6
Humoral Immune Response	quantity	quantity of IgG	3.76E-05	CXCL10,CXCL9,IFNG (includes EG:15978),IL6,PTGS2,ZC3H12A	6
Protein Response	quantity	quantity of IgG	3.76E-05	CXCL10,CXCL9,IFNG (includes EG:15978),IL6,PTGS2,ZC3H12A	6
Hematological System Development and Function	quantity	quantity of lymphocytes	3.92E-05	CAMP (includes EG:12796),CXCL10,CXCR2,GPR83,IFNG (includes EG:15978),IL6,MS4A1,PTPRC,SELL,ZC3H12A	10
Tissue Morphology	quantity	quantity of lymphocytes	3.92E-05	CAMP (includes EG:12796),CXCL10,CXCR2,GPR83,IFNG (includes EG:15978),IL6,MS4A1,PTPRC,SELL,ZC3H12A	10
Respiratory Disease	growth	growth of lung carcinoma	4.00E-05	CXCR2,PTGS2	2
Cancer	growth	growth of lung carcinoma	4.00E-05	CXCR2,PTGS2	2
Cellular Movement	migration	migration of cytotoxic T cells	4.00E-05	CXCL10,CXCL9	2
Hematological System Development and Function	migration	migration of cytotoxic T cells	4.00E-05	CXCL10,CXCL9	2
Immune Cell Trafficking	migration	migration of cytotoxic T cells	4.00E-05	CXCL10,CXCL9	2
Cell-mediated Immune Response	migration	migration of cytotoxic T cells	4.00E-05	CXCL10,CXCL9	2
Cell-mediated Immune Response	migration	migration of cytotoxic T cells	4.00E-05	HPGD,PTGS2	2
Energy Production	oxidation	oxidation of prostaglandin	4.00E-05	HPGD,PTGS2	2
Lipid Metabolism	oxidation	oxidation of prostaglandin	4.00E-05	HPGD,PTGS2	2
Small Molecule Biochemistry	oxidation	oxidation of prostaglandin	4.00E-05	HPGD,PTGS2	2
Cellular Movement	recruitment	recruitment of Th1 cells	4.00E-05	CXCL10,CXCL9	2
Hematological System Development and Function	recruitment	recruitment of Th1 cells	4.00E-05	CXCL10,CXCL9	2
Immune Cell Trafficking	recruitment	recruitment of Th1 cells	4.00E-05	CXCL10,CXCL9	2
Cell-To-Cell Signaling and Interaction	recruitment	recruitment of Th1 cells	4.00E-05	CXCL10,CXCL9	2
Cell-mediated Immune Response	recruitment	recruitment of Th1 cells	4.00E-05	CXCL10,CXCL9	2
Hematological System Development and Function	sensitization	sensitization of macrophages	4.04E-05	IFNG (includes EG:15978),IL6	2
Inflammatory Response	sensitization	sensitization of macrophages	4.04E-05	IFNG (includes EG:15978),IL6	2
Cell-To-Cell Signaling and Interaction	sensitization	sensitization of macrophages	4.04E-05	IFNG (includes EG:15978),IL6	2
Cell Morphology	whole-cell conductance	whole-cell conductance of oocytes	4.04E-05	CFTR,SCNN1G	2
Reproductive System Development and Function	whole-cell conductance	whole-cell conductance of oocytes	4.04E-05	CFTR,SCNN1G	2
Hematological System Development and Function	activation	activation of antigen presenting cells	4.03E-05	CAMP (includes EG:12796),CXCL10,FOXA2,IFNG (includes EG:15978),IL6,PTGS2	6
Immune Cell Trafficking	activation	activation of antigen presenting cells	4.03E-05	CAMP (includes EG:12796),CXCL10,FOXA2,IFNG (includes EG:15978),IL6,PTGS2	6
Inflammatory Response	activation	activation of antigen presenting cells	4.03E-05	CAMP (includes EG:12796),CXCL10,FOXA2,IFNG (includes EG:15978),IL6,PTGS2	6
Cell-To-Cell Signaling and Interaction	activation	activation of antigen presenting cells	4.03E-05	CAMP (includes EG:12796),CXCL10,FOXA2,IFNG (includes EG:15978),IL6,PTGS2	6
Lipid Metabolism	conversion	conversion of eicosanoid	4.16E-05	CFTR,HPGD,PTGS2	3
Small Molecule Biochemistry	conversion	conversion of eicosanoid	4.16E-05	CFTR,HPGD,PTGS2	3
Cellular Movement	recruitment	recruitment of lymphocytes	4.53E-05	CXCL10,CXCL9,IFNG (includes EG:15978),SELL	4
Hematological System Development and Function	recruitment	recruitment of lymphocytes	4.53E-05	CXCL10,CXCL9,IFNG (includes EG:15978),SELL	4
Immune Cell Trafficking	recruitment	recruitment of lymphocytes	4.53E-05	CXCL10,CXCL9,IFNG (includes EG:15978),SELL	4
Immunological Disease	immediate hypersensitivity	immediate hypersensitivity	4.55E-05	CAMP (includes EG:12796),CXCL9,IFNG (includes EG:15978),LCE3D,PPARGC1A,PTGS2,PTPRC	7
Hematological System Development and Function	quantity	quantity of B lymphocytes	4.63E-05	CAMP (includes EG:12796),CXCR2,IFNG (includes EG:15978),IL6,MS4A1,PTPRC,ZC3H12A	7
Humoral Immune Response	quantity	quantity of B lymphocytes	4.63E-05	CAMP (includes EG:12796),CXCR2,IFNG (includes EG:15978),IL6,MS4A1,PTPRC,ZC3H12A	7
Tissue Morphology	quantity	quantity of B lymphocytes	4.63E-05	CAMP (includes EG:12796),CXCR2,IFNG (includes EG:15978),IL6,MS4A1,PTPRC,ZC3H12A	7
Cellular Movement	chemoattraction	chemoattraction of myeloid cells	4.67E-05	CAMP (includes EG:12796),CXCL10,CXCL9	3
Hematological System Development and Function	chemoattraction	chemoattraction of myeloid cells	4.67E-05	CAMP (includes EG:12796),CXCL10,CXCL9	3
Immune Cell Trafficking	chemoattraction	chemoattraction of myeloid cells	4.67E-05	CAMP (includes EG:12796),CXCL10,CXCL9	3
Inflammatory Response	chemoattraction	chemoattraction of myeloid cells	4.67E-05	CAMP (includes EG:12796),CXCL10,CXCL9	3
Cell-To-Cell Signaling and Interaction	chemoattraction	chemoattraction of myeloid cells	4.67E-05	CAMP (includes EG:12796),CXCL10,CXCL9	3

Cellular Movement	chemotaxis	chemotaxis of antigen presenting cells	4.72E-05	CAMP (includes EG:12796) CXCL10,CXCL9,CXCR2,SELL
Hematological System Development and Function	chemotaxis	chemotaxis of antigen presenting cells	4.72E-05	CAMP (includes EG:12796) CXCL10,CXCL9,CXCR2,SELL
Immune Cell Trafficking	chemotaxis	chemotaxis of antigen presenting cells	4.72E-05	CAMP (includes EG:12796) CXCL10,CXCL9,CXCR2,SELL
Inflammatory Response	chemotaxis	chemotaxis of antigen presenting cells	4.72E-05	CAMP (includes EG:12796) CXCL10,CXCL9,CXCR2,SELL
Lipid Metabolism	synthesis	synthesis of fatty acid	4.87E-05	ACSM3,CFTR,HPGD,IFNG (includes EG:15978) IL6,PPARGC1A,PTGS2
Small Molecule Biochemistry	synthesis	synthesis of fatty acid	4.87E-05	ACSM3,CFTR,HPGD,IFNG (includes EG:15978) IL6,PPARGC1A,PTGS2
Hematological System Development and Function	accumulation	accumulation of leukocytes	4.93E-05	CXCL10,CXCL9,IFNG (includes EG:15978) IL6,MMP8,SELL
Immune Cell Trafficking	accumulation	accumulation of leukocytes	4.93E-05	CXCL10,CXCL9,IFNG (includes EG:15978) IL6,MMP8,SELL
Inflammatory Response	accumulation	accumulation of leukocytes	4.93E-05	CXCL10,CXCL9,IFNG (includes EG:15978) IL6,MMP8,SELL
Tissue Development	accumulation	accumulation of leukocytes	4.93E-05	CXCL10,CXCL9,IFNG (includes EG:15978) IL6,MMP8,SELL
Hematological System Development and Function	function	function of lymphocytes	5.21E-05	CXCL10,CXCR2,IFNG (includes EG:15978) IL6,MS4A1,PTPRC,SELL
Cellular Function and Maintenance	function	function of lymphocytes	5.21E-05	CXCL10,CXCR2,IFNG (includes EG:15978) IL6,MS4A1,PTPRC,SELL
Immunological Disease	autoimmune thyroid disease	autoimmune thyroid disease	5.22E-05	CXCL10,IL6,MS4A1
Endocrine System Disorders	autoimmune thyroid disease	autoimmune thyroid disease	5.22E-05	CXCL10,IL6,MS4A1
Cellular Movement	chemotaxis	chemotaxis of kidney cell lines	5.22E-05	CAMP (includes EG:12796) CXCL10,CXCR2
Renal and Urological System Development and Function	chemotaxis	chemotaxis of kidney cell lines	5.22E-05	CAMP (includes EG:12796) CXCL10,CXCR2
Cellular Movement	cell movement	cell movement of macrophages	5.26E-05	CAMP (includes EG:12796) CXCL10,CXCL9,IFNG (includes EG:15978) PTGS2,SELL
Hematological System Development and Function	cell movement	cell movement of macrophages	5.26E-05	CAMP (includes EG:12796) CXCL10,CXCL9,IFNG (includes EG:15978) PTGS2,SELL
Immune Cell Trafficking	cell movement	cell movement of macrophages	5.26E-05	CAMP (includes EG:12796) CXCL10,CXCL9,IFNG (includes EG:15978) PTGS2,SELL
Inflammatory Response	cell movement	cell movement of macrophages	5.26E-05	CAMP (includes EG:12796) CXCL10,CXCL9,IFNG (includes EG:15978) PTGS2,SELL
Cell-To-Cell Signaling and Interaction	phagocytosis	phagocytosis of cells	5.37E-05	CAMP (includes EG:12796) CXCL10,IFNG (includes EG:15978) IL6,MARCO,PTPRC
Cellular Function and Maintenance	phagocytosis	phagocytosis of cells	5.37E-05	CAMP (includes EG:12796) CXCL10,IFNG (includes EG:15978) IL6,MARCO,PTPRC
Cellular Function and Maintenance	flux	flux of ion	5.85E-05	CAMP (includes EG:12796) CFTR,CXCL10,IFNG (includes EG:15978) IL6,MS4A1,SCNN1G
Molecular Transport	flux	flux of ion	5.85E-05	CAMP (includes EG:12796) CFTR,CXCL10,IFNG (includes EG:15978) IL6,MS4A1,SCNN1G
Hematological System Development and Function	activation	activation of macrophages	6.20E-05	CAMP (includes EG:12796) CXCL10,IPRC (includes EG:15978) IL6,PTGS2
Immune Cell Trafficking	activation	activation of macrophages	6.20E-05	CAMP (includes EG:12796) CXCL10,IPRC (includes EG:15978) IL6,PTGS2
Inflammatory Response	activation	activation of macrophages	6.20E-05	CAMP (includes EG:12796) CXCL10,IPRC (includes EG:15978) IL6,PTGS2
Cell-To-Cell Signaling and Interaction	activation	activation of macrophages	6.20E-05	CAMP (includes EG:12796) CXCL10,IPRC (includes EG:15978) IL6,PTGS2
Lipid Metabolism	conversion	conversion of docosahexaenoic acid	6.65E-05	CFTR,PTGS2
Small Molecule Biochemistry	conversion	conversion of docosahexaenoic acid	6.65E-05	CFTR,PTGS2
Organismal Development	formation	formation of bile duct	6.65E-05	FOXA2,IL6
Infectious Disease	hantavirus pulmonary syndrome	hantavirus pulmonary syndrome	6.65E-05	IFNG (includes EG:15978),IL6
Respiratory Disease	hantavirus pulmonary syndrome	hantavirus pulmonary syndrome	6.65E-05	IFNG (includes EG:15978),IL6
Connective Tissue Disorders	hyperplasia	hyperplasia of synovial cells	6.65E-05	IFNG (includes EG:15978),IL6
Cancer	hyperplasia	hyperplasia of synovial cells	6.65E-05	IFNG (includes EG:15978),IL6
Cancer	growth	growth of tumor	6.98E-05	CXCL9,CXCR2,IFNG (includes EG:15978) IL6,MS4A1,PTGS2,SELL
Humoral Immune Response	class switching	class switching	7.04E-05	IFNG (includes EG:15978),IL6,PTGS2,ZC3H12A
Infectious Disease	sexually transmitted disease	sexually transmitted disease	7.04E-05	IFNG (includes EG:15978),IL6,PTGS2,PTPRC
Lymphoid Tissue Structure and Development	abnormal morphology	abnormal morphology of spleen	7.05E-05	CXCR2,IFNG (includes EG:15978),PTGS2,PTPRC,SELL,ZC3H12A
Organ Pathology	abnormal morphology	abnormal morphology of spleen	7.05E-05	CXCR2,IFNG (includes EG:15978),PTGS2,PTPRC,SELL,ZC3H12A
Hematological System Development and Function	proliferation	proliferation of myeloid progenitor cells	7.12E-05	CXCL10,IFNG (includes EG:15978),IL6
Lymphoid Tissue Structure and Development	proliferation	proliferation of myeloid progenitor cells	7.12E-05	CXCL10,IFNG (includes EG:15978),IL6
Cellule Development	proliferation	proliferation of myeloid progenitor cells	7.12E-05	CXCL10,IFNG (includes EG:15978),IL6
Hematopoiesis	proliferation	proliferation of myeloid progenitor cells	7.12E-05	CXCL10,IFNG (includes EG:15978),IL6
Cellular Growth and Proliferation	proliferation	proliferation of myeloid progenitor cells	7.12E-05	CXCL10,IFNG (includes EG:15978),IL6
Nutritional Disease	cachexia	cachexia	7.36E-05	CFTR,IFNG (includes EG:15978),IL6,PPARGC1A
Organismal Injury and Abnormalities	cachexia	cachexia	7.36E-05	CFTR,IFNG (includes EG:15978),IL6,PPARGC1A
Cellular Movement	cell movement	cell movement of epithelial cell lines	7.70E-05	CAMP (includes EG:12796) CXCL10,CXCR2,PTGS2
Hair and Skin Development and Function	cell movement	cell movement of epithelial cell lines	7.70E-05	CAMP (includes EG:12796) CXCL10,CXCR2,PTGS2
Cellular Movement	chemotaxis	chemotaxis of neutrophils	7.79E-05	CAMP (includes EG:12796) CXCL10,CXCR2,IFNG (includes EG:15978),SELL
Hematological System Development and Function	chemotaxis	chemotaxis of neutrophils	7.79E-05	CAMP (includes EG:12796) CXCL10,CXCR2,IFNG (includes EG:15978),SELL
Immune Cell Trafficking	chemotaxis	chemotaxis of neutrophils	7.79E-05	CAMP (includes EG:12796) CXCL10,CXCR2,IFNG (includes EG:15978),SELL
Inflammatory Response	chemotaxis	chemotaxis of neutrophils	7.79E-05	CAMP (includes EG:12796) CXCL10,CXCR2,IFNG (includes EG:15978),SELL
Cell Death and Survival	apoptosis	apoptosis of oligodendrocytes	8.06E-05	CAMP (includes EG:12796) CXCL10,CXCR2,IFNG (includes EG:15978),PTGS2
Cellular Development	cell movement	cell movement of colon cancer cell lines	8.06E-05	CXCL10,CXOL9,IFNG (includes EG:15978),IL6
Cellular Movement	cell movement	cell movement of colon cancer cell lines	8.06E-05	CXCL10,IFNG (includes EG:15978),IL6,PTPRC
Hematological System Development and Function	proliferation	proliferation of myeloid cells	8.06E-05	CXCL10,IFNG (includes EG:15978),IL6,PTPRC
Cellular Development	proliferation	proliferation of myeloid cells	8.06E-05	CXCL10,IFNG (includes EG:15978),IL6,PTPRC
Cellular Growth and Proliferation	proliferation	proliferation of myeloid cells	8.06E-05	APOB,CXCL10,CXCR2,IFNG (includes EG:15978)
Cellular Movement	recruitment	recruitment of macrophages	8.06E-05	APOB,CXCL10,CXCR2,IFNG (includes EG:15978)
Hematological System Development and Function	recruitment	recruitment of macrophages	8.06E-05	APOB,CXCL10,CXCR2,IFNG (includes EG:15978)
Immune Cell Trafficking	recruitment	recruitment of macrophages	8.06E-05	APOB,CXCL10,CXCR2,IFNG (includes EG:15978)
Inflammatory Response	recruitment	recruitment of macrophages	8.06E-05	APOB,CXCL10,CXCR2,IFNG (includes EG:15978)
Cell-To-Cell Signaling and Interaction	recruitment	recruitment of macrophages	8.06E-05	APOB,CXCL10,CXCR2,IFNG (includes EG:15978)
Organismal Survival	organismal death	organismal death	8.20E-05	ACE2 (includes EG:302668),APOB,CAMP (includes EG:12796),CFTR,CXCL10,CXCL9,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MARCO,PPARGC1A,PTGS2,PTPRC,ZC3H12A
Hematological System Development and Function	accumulation	accumulation of eosinophils	8.61E-05	CXCL10,CXCL9,SELL
Immune Cell Trafficking	accumulation	accumulation of eosinophils	8.61E-05	CXCL10,CXCL9,SELL
Inflammatory Response	accumulation	accumulation of eosinophils	8.61E-05	CXCL10,CXCL9,SELL
Hypersensitivity Response	accumulation	accumulation of eosinophils	8.61E-05	CXCL10,CXCL9,SELL
Tissue Development	accumulation	accumulation of eosinophils	8.61E-05	CXCL10,CXCL9,SELL
Lipid Metabolism	synthesis	synthesis of steroid	9.12E-05	APOB,CFTR,CXCL10,IFNG (includes EG:15978),IL6,PPARGC1A
Small Molecule Biochemistry	synthesis	synthesis of steroid	9.12E-05	APOB,CFTR,CXCL10,IFNG (includes EG:15978),IL6,PPARGC1A
Vitamin and Mineral Metabolism	synthesis	synthesis of steroid	9.12E-05	APOB,CFTR,CXCL10,IFNG (includes EG:15978),IL6,PPARGC1A
Infectious Disease	infection	infection by DNA virus	9.18E-05	CAMP (includes EG:12796),IFNG (includes EG:15978),MARCO,PTPRC
Hematological System Development and Function	phagocytosis	phagocytosis by macrophages	9.18E-05	CAMP (includes EG:12796),IFNG (includes EG:15978),MARCO,PTPRC
Inflammatory Response	phagocytosis	phagocytosis by macrophages	9.18E-05	CAMP (includes EG:12796),IFNG (includes EG:15978),MARCO,PTPRC
Cell-To-Cell Signaling and Interaction	phagocytosis	phagocytosis by macrophages	9.18E-05	CAMP (includes EG:12796),IFNG (includes EG:15978),MARCO,PTPRC
Cellular Function and Maintenance	phagocytosis	phagocytosis by macrophages	9.18E-05	CAMP (includes EG:12796),IFNG (includes EG:15978),MARCO,PTPRC
Molecular Transport	mobilization	mobilization of Ca2+	9.92E-05	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),PTPRC,SELL
Vitamin and Mineral Metabolism	mobilization	mobilization of Ca2+	9.92E-05	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),PTPRC,SELL
Cell Signaling	mobilization	mobilization of Ca2+	9.92E-05	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),PTPRC,SELL
Cell-To-Cell Signaling and Interaction	activation	activation of keratinocytes	9.96E-05	IFNG (includes EG:15978),TRPV3
Hair and Skin Development and Function	activation	activation of keratinocytes	9.96E-05	IFNG (includes EG:15978),TRPV3
Tissue Development	involution	delay in involution of mammary gland	9.96E-05	IL6,PTGS2
Organ Morphology	involution	delay in involution of mammary gland	9.96E-05	IL6,PTGS2
Reproductive System Development and Function	involution	delay in involution of mammary gland	9.96E-05	IL6,PTGS2
Connective Tissue Disorders	experimentally-induced arthritis	experimentally-induced arthritis of ankle joint	9.96E-05	IFNG (includes EG:15978),IL6
Inflammatory Disease	experimentally-induced arthritis	experimentally-induced arthritis of ankle joint	9.96E-05	IFNG (includes EG:15978),IL6

Skeletal and Muscular Disorders	experimentally-induced arthritis	experimentally-induced arthritis of ankle joint	9.9E-05	IFNG (includes EG:15978),IL6	2
Hematological System Development and Function	inhibition	inhibition of Th17 cells	9.9E-05	IFNG (includes EG:15978),PTGS2	2
Cellular Growth and Proliferation	inhibition	inhibition of Th17 cells	9.9E-05	IFNC (includes EG:15978),PTGS2	3
Hematological System Development and Function	activation	activation of neutrophils	9.9E-05	CXCR2,IFNG (includes EG:15978),IL6,SELL	4
Immune Cell Trafficking	activation	activation of neutrophils	9.9E-05	CXCR2,IFNG (includes EG:15978),IL6,SELL	4
Inflammatory Response	activation	activation of neutrophils	9.9E-05	CXCR2,IFNG (includes EG:15978),IL6,SELL	4
Cell-To-Cell Signaling and Interaction	activation	activation of neutrophils	9.9E-05	CXCR2,IFNG (includes EG:15978),IL6,SELL	4
Cancer	growth	growth of malignant tumor	9.9E-05	CXCL9,CXCR2,IL6,PTGS	4
Hematological System Development and Function	function	function of T lymphocytes	1.02E-04	CXCL10,CXCR2,IFNG (includes EG:15978),IL6,PTPRC,SELL	6
Cellular Function and Maintenance	function	function of T lymphocytes	1.02E-04	CXCL10,CXCR2,IFNG (includes EG:15978),IL6,PTPRC,SELL	6
Cell Death and Survival	apoptosis	apoptosis of lung cell lines	1.03E-04	CAMP (Includes EG:12796),IFNG (includes EG:15978),PTGS2	3
Cellular Movement	chemoattraction	chemoattraction of phagocytes	1.03E-04	CAMP (Includes EG:12796),CXCL10,CXCL9	3
Hematological System Development and Function	chemoattraction	chemoattraction of phagocytes	1.03E-04	CAMP (Includes EG:12796),CXCL10,CXCL9	3
Immune Cell Trafficking	chemoattraction	chemoattraction of phagocytes	1.03E-04	CAMP (Includes EG:12796),CXCL10,CXCL9	3
Inflammatory Response	chemoattraction	chemoattraction of phagocytes	1.03E-04	CAMP (Includes EG:12796),CXCL10,CXCL9	3
Cell-To-Cell Signaling and Interaction	chemoattraction	chemoattraction of phagocytes	1.03E-04	CAMP (Includes EG:12796),CXCL10,CXCL9	3
Connective Tissue Development and Function	formation	formation of osteoclasts	1.04E-04	Chi3l3/Ch34,IFNG (includes EG:15978),IL6,PTGS2	4
Cellular Growth and Proliferation	formation	formation of osteoclasts	1.04E-04	Chi3l3/Ch34,IFNG (includes EG:15978),IL6,PTGS2	4
Skeletal and Muscular System Development and Function	formation	formation of osteoclasts	1.04E-04	Chi3l3/Ch34,IFN (includes EG:15978),IL6,PTGS2	4
Cancer	incidence	incidence of tumor	1.06E-04	IFNG (includes EG:15978),IL6,MMP8,PPARGC1A,PTGS2,SELL	6
Lipid Metabolism	metabolism	steroid metabolism	1.06E-04	Akr1c14,APOB,CFTR,CYP2C19,IFNG (includes EG:15978),IL6	6
Small Molecule Biochemistry	metabolism	steroid metabolism	1.06E-04	Akr1c14,APOB,CFTR,CYP2C19,IFNG (includes EG:15978),IL6	6
Vitamin and Mineral Metabolism	metabolism	steroid metabolism	1.06E-04	Akr1c14,APOB,CFTR,CYP2C19,IFNG (includes EG:15978),IL6	6
Inflammatory Disease	hepatitis	Hepatitis	1.07E-04	APOB,CXL10,IFNG (includes EG:15978),IL6,PTPRC	5
Gastrointestinal Disease	hepatitis	Hepatitis	1.07E-04	APOB,CXL10,IFNG (includes EG:15978),IL6,PTPRC	5
Hepatic System Disease	hepatitis	Hepatitis	1.07E-04	APOB,CXL10,IFNG (Includes EG:12796),IL6,PTPRC	5
Cellular Movement	chemotaxis	chemotaxis of T lymphocytes	1.09E-04	CAMP (Includes EG:12796),Chi3l3/Ch34,CXCL10,CXCL9	4
Hematological System Development and Function	chemotaxis	chemotaxis of T lymphocytes	1.09E-04	CAMP (Includes EG:12796),Chi3l3/Ch34,CXCL10,CXCL9	4
Immune Cell Trafficking	chemotaxis	chemotaxis of T lymphocytes	1.09E-04	CAMP (Includes EG:12796),Chi3l3/Ch34,CXCL10,CXCL9	4
Inflammatory Response	chemotaxis	chemotaxis of T lymphocytes	1.09E-04	CAMP (Includes EG:12796),Chi3l3/Ch34,CXCL10,CXCL9	4
Cell-mediated Immune Response	chemotaxis	chemotaxis of T lymphocytes	1.09E-04	CAMP (Includes EG:12796),Chi3l3/Ch34,CXCL10,CXCL9	4
Infectious Disease	endotoxemia	endotoxemia	1.09E-04	CAMP (Includes EG:12796),IFNG (includes EG:15978),IL6,PTGS2	4
Hematological Disease	endotoxemia	endotoxemia	1.09E-04	CAMP (Includes EG:12796),IFNG (includes EG:15978),IL6,PTGS2	4
Inflammatory Response	immune response	immune response of T lymphocytes	1.09E-04	CAMP (Includes EG:12796),IFNG (includes EG:15978),IL6,PTPRC	4
Cell-To-Cell Signaling and Interaction	immune response	immune response of T lymphocytes	1.09E-04	CAMP (Includes EG:12796),IFNG (includes EG:15978),IL6,PTPRC	4
Molecular Transport	quantity	quantity of Ca2+	1.09E-04	CXCL10,CXCL9,CXCR2,IFN (includes EG:15978),IL6,PTGS2,PTPRC,SELL	8
Vitamin and Mineral Metabolism	quantity	quantity of Ca2+	1.09E-04	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,PTGS2,PTPRC,SELL	8
Cell Signaling	quantity	quantity of Ca2+	1.09E-04	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,PTGS2,PTPRC,SELL	8
Respiratory Disease	fibrosis	fibrosis of lung	1.10E-04	CFTR,IFNG (includes EG:15978),IL6,PTGS2,SELL	5
Inflammatory Disease	fibrosis	fibrosis of lung	1.10E-04	CFTR,IFNG (includes EG:15978),IL6,PTGS2,SELL	5
Organismal Injury and Abnormalities	fibrosis	fibrosis of lung	1.10E-04	CFTR,IFNG (includes EG:15978),IL6,PTGS2,SELL	5
Protein Synthesis	quantity	quantity of protein in blood	1.16E-04	APOB,CXR2,FABP2,IFNG (includes EG:15978),IL6,PPARGC1A,PTGS2,SELL	8
Hematological System Development and Function	binding	binding of lymphocytes	1.22E-04	CXCL10,CXCL9,IFNG (includes EG:15978),SELL	4
Cell-To-Cell Signaling and Interaction	binding	binding of lymphocytes	1.22E-04	CXCL10,CXCL9,IFNG (includes EG:15978),SELL	4
Cellular Movement	chemotaxis	chemotaxis of macrophages	1.22E-04	CAMP (Includes EG:12796),CXCL10,CXCL9,SELL	4
Hematological System Development and Function	chemotaxis	chemotaxis of macrophages	1.22E-04	CAMP (Includes EG:12796),CXCL10,CXCL9,SELL	4
Immune Cell Trafficking	chemotaxis	chemotaxis of macrophages	1.22E-04	CAMP (Includes EG:12796),CXCL10,CXCL9,SELL	4
Inflammatory Response	chemotaxis	chemotaxis of macrophages	1.22E-04	CAMP (Includes EG:12796),CXCL10,CXCL9,SELL	4
Tissue Development	development	lung development	1.23E-04	CFTR,FOXA2,IFNG (includes EG:15978),IL6,PPARGC1A,SELL	6
Embryonic Development	development	lung development	1.23E-04	CFTR,FOXA2,IFNG (includes EG:15978),IL6,PPARGC1A,SELL	6
Organismal Development	development	lung development	1.23E-04	CFTR,FOXA2,IFNG (includes EG:15978),IL6,PPARGC1A,SELL	6
Organ Development	development	lung development	1.23E-04	CFTR,FOXA2,IFNG (includes EG:15978),IL6,PPARGC1A,SELL	6
Respiratory System Development and Function	development	lung development	1.23E-04	CFTR,FOXA2,IFNG (includes EG:15978),IL6,PPARGC1A,SELL	6
Hematological System Development and Function	accumulation	accumulation of lymphocytes	1.27E-04	CXCL10,IFNG (includes EG:15978),IL6,SELL	4
Immune Cell Trafficking	accumulation	accumulation of lymphocytes	1.27E-04	CXCL10,IFNG (includes EG:15978),IL6,SELL	4
Inflammatory Response	accumulation	accumulation of lymphocytes	1.27E-04	CXCL10,IFNG (includes EG:15978),IL6,SELL	4
Tissue Development	accumulation	accumulation of lymphocytes	1.27E-04	CXCL10,IFNG (includes EG:15978),IL6,SELL	4
Cell-To-Cell Signaling and Interaction	immunoreaction	immunoreaction of tumor cell lines	1.27E-04	IFNG (Includes EG:12796),IL6,MARCO,MS4A1	4
Cell Morphology	immunoreaction	immunoreaction of tumor cell lines	1.27E-04	IFNG (Includes EG:15978),IL6,MARCO,MS4A1	4
Organismal Injury and Abnormalities	polarization	polarization of blood cells	1.28E-04	CXCL10,CXCL9,IFNC (includes EG:15978),IL6	4
Gastrointestinal Disease	damage	damage of liver	1.28E-04	CXCL10,CXCR2,IFNG (includes EG:15978),IL6,PTGS2	5
Hepatic System Disease	damage	damage of liver	1.28E-04	CXCL10,CXCR2,IFNG (includes EG:15978),IL6,PTGS2	5
Nutritional Disease	obesity	obesity	1.29E-04	APOB,CXL10,FABP2,FOXA2,IL6,PPARGC1A,PTGS2	7
Free Radical Scavenging	production	production of reactive oxygen species	1.29E-04	CAMP (Includes EG:12796),CXCL10,CXCL9,IFNG (includes EG:15978),IL6,MS4A1,PPARGC1A,PTGS2	7
Hematological System Development and Function	quantity	quantity of T lymphocytes	1.29E-04	CAMP (Includes EG:12796),CXCL10,GP83,IFNG (includes EG:15978),IL6,PTPRC,SELL,ZC3H12A	8
Tissue Morphology	quantity	quantity of T lymphocytes	1.29E-04	CAMP (Includes EG:12796),CXCL10,GP83,IFNG (includes EG:15978),IL6,PTPRC,SELL,ZC3H12A	8
Tissue Morphology	quantity	quantity of cells	1.30E-04	APOB,CAMP (Includes EG:12796),CFTR,CXL10,CXCR2,GPR83,IFNG (includes EG:15978),IL6,MS4A1,PPARGC1A,PTGS2,PTPRC,SELL,ZC3H12A	15
Free Radical Scavenging	synthesis	synthesis of reactive oxygen species	1.30E-04	CAMP (Includes EG:12796),CXL10,GPR83,IFNG (Includes EG:15978),IL6,MS4A1,PPARGC1A,PTGS2,ZC3H12A	8
Hematological System Development and Function	quantity	quantity of helper T lymphocytes	1.32E-04	CXCL10,GPR83,IL6,PTPRC,SELL	5
Tissue Morphology	quantity	quantity of helper T lymphocytes	1.32E-04	CXCL10,GPR83,IL6,PTPRC,SELL	5
Cellular Compromise	depletion	depletion of lymphocytes	1.32E-04	IFNG (Includes EG:15978),MS4A1,SELL	3
Cell Death and Survival	cell death	cell death of carcinoma cell lines	1.35E-04	CAMP (Includes EG:12796),IFNG (Includes EG:15978),IL6,PTGS2,SLC7A11	5
Hematological System Development and Function	accumulation	accumulation of TREG cells	1.39E-04	IFNG (Includes EG:15978),IL6	2
Immune Cell Trafficking	accumulation	accumulation of TREG cells	1.39E-04	IFNG (Includes EG:15978),IL6	2
Inflammatory Response	accumulation	accumulation of TREG cells	1.39E-04	IFNG (Includes EG:15978),IL6	2
Tissue Development	accumulation	accumulation of TREG cells	1.39E-04	IFNG (Includes EG:15978),IL6	2
Endocrine System Development and Function	activation	activation of hypothalamus-pituitary-adrenal axis	1.39E-04	IFNG (Includes EG:15978),IL6	2
Hematological System Development and Function	activation	activation of monocyte-derived macrophages	1.39E-04	CAMP (Includes EG:12796),IFNG (Includes EG:15978)	2
Immune Cell Trafficking	activation	activation of monocyte-derived macrophages	1.39E-04	CAMP (Includes EG:12796),IFNG (Includes EG:15978)	2
Inflammatory Response	activation	activation of monocyte-derived macrophages	1.39E-04	CAMP (Includes EG:12796),IFNG (Includes EG:15978)	2
Cell-To-Cell Signaling and Interaction	activation	activation of monocyte-derived macrophages	1.39E-04	CAMP (Includes EG:12796),IFNG (Includes EG:15978)	2
Cellular Movement	attraction	attraction of natural killer cells	1.39E-04	CXCL10,CXCL9	2
Immune Cell Trafficking	attraction	attraction of natural killer cells	1.39E-04	CXCL10,CXCL9	2
Cell-To-Cell Signaling and Interaction	attraction	attraction of natural killer cells	1.39E-04	CXCL10,CXCL9	2
Small Molecule Biochemistry	incorporation	incorporation of guanosine 5'-O-(3-thiotriphosphate)	1.39E-04	CXCL10,LOXL9	2
Cell Signaling	incorporation	incorporation of guanosine 5'-O-(3-thiotriphosphate)	1.39E-04	CXCL10,CXCL9	2
Nucleic Acid Metabolism	incorporation	incorporation of guanosine 5'-O-(3-thiotriphosphate)	1.39E-04	CXCL10,CXCL9	2
Cell-To-Cell Signaling and Interaction	stimulation	stimulation of hepatoma cell lines	1.39E-04	IFNG (Includes EG:15978),IL6	2

Cellular Growth and Proliferation	stimulation	stimulation of hepatoma cell lines	1.39E-04	IFNG (includes EG:15978),IL6	2
Cardiovascular System Development and Function	vascularization	vascularization of hindlimb	1.39E-04	CAMP (Includes EG:12796),PTGS2	2
Organismal Development	vascularization	vascularization of hindlimb	1.39E-04	CAMP (Includes EG:12796),PTGS2	2
Lipid Metabolism	phospholipid flip-flop	phospholipid flip-flop of phosphatidyserine	1.43E-04	CAMP (Includes EG:12796),IFNG (includes EG:15978),PTGS2	3
Small Molecule Biochemistry	phospholipid flip-flop	phospholipid flip-flop of phosphatidyserine	1.43E-04	CAMP (Includes EG:12796),IFNG (includes EG:15978),PTGS2	3
Hematological System Development and Function	adhesion	adhesion of granulocytes	1.43E-04	CXCR2,IFNG (includes EG:15978),IL6,SELL	4
Immune Cell Trafficking	adhesion	adhesion of granulocytes	1.43E-04	CXCR2,IFNG (includes EG:15978),IL6,SELL	4
Cell-To-Cell Signaling and Interaction	adhesion	adhesion of granulocytes	1.43E-04	CXCR2,IFNG (includes EG:15978),IL6,SELL	4
Tissue Development	adhesion	adhesion of granulocytes	1.43E-04	CXCR2,IFNG (includes EG:15978),IL6,SELL	4
Lipid Metabolism	quantity	quantity of sphingolipid	1.43E-04	ACER1,CFTR,IFNG (includes EG:15978),IL6	4
Small Molecule Biochemistry	quantity	quantity of sphingolipid	1.43E-04	ACER1,CFTR,IFNG (includes EG:15978),IL6	4
Molecular Transport	quantity	quantity of sphingolipid	1.43E-04	ACER1,CFTR,IFNG (includes EG:15978),IL6	4
Hematological System Development and Function	activation	activation of dendritic cells	1.48E-04	CAMP (Includes EG:12796),FOXA2,IFNG (includes EG:15978),IL6	4
Immune Cell Trafficking	activation	activation of dendritic cells	1.48E-04	CAMP (Includes EG:12796),FOXA2,IFNG (includes EG:15978),IL6	4
Inflammatory Response	activation	activation of dendritic cells	1.48E-04	CAMP (Includes EG:12796),FOXA2,IFNG (includes EG:15978),IL6	4
Cell-To-Cell Signaling and Interaction	activation	activation of dendritic cells	1.48E-04	CAMP (Includes EG:12796),FOXA2,IFNG (includes EG:15978),IL6	4
Inflammatory Response	fever	fever	1.66E-04	IFNG (Includes EG:15978),IL6,PTGS2	3
Organismal Functions	fever	fever	1.66E-04	IFNG (Includes EG:15978),IL6,PTGS2	3
Infectious Disease	mycosis	mycosis	1.66E-04	IFNG (Includes EG:15978),IL6,PTGS2	3
Cardiovascular Disease	congestive heart failure	congestive heart failure	1.66E-04	CXCR2,IFNG (Includes EG:15978),IL6	3
Neurological Disease	demyelination	demyelination	1.66E-04	ACE2 (Includes EG:302668),HPGD,IL6,PPARGC1A	4
Cell-To-Cell Signaling and Interaction	activation	activation of astrocytes	1.78E-04	CXCL10,IFNG (Includes EG:15978),IL6,PTGS2	3
Nervous System Development and Function	activation	activation of astrocytes	1.78E-04	IFNG (Includes EG:15978),IL6,PTGS2	3
Cellular Movement	cell movement	cell movement of bone marrow cells	1.78E-04	Ch313/Ch34,IFNG (Includes EG:15978),IL6	3
Immune Cell Trafficking	cell movement	cell movement of bone marrow cells	1.78E-04	Ch313/Ch34,IFNG (Includes EG:15978),IL6	3
Inflammatory Response	cell movement	cell movement of bone marrow cells	1.78E-04	Ch313/Ch34,IFNG (Includes EG:15978),IL6	3
Hematopoiesis	cell movement	cell movement of bone marrow cells	1.78E-04	Ch313/Ch34,IFNG (Includes EG:15978),IL6	3
Lipid Metabolism	incorporation	incorporation of lipid	1.78E-04	ACE2,CFTR,FAFB2	3
Small Molecule Biochemistry	incorporation	incorporation of lipid	1.78E-04	ACER1,CFTR,ABP2	3
Hematological System Development and Function	phagocytosis	phagocytosis of neutrophils	1.78E-04	CAMP (Includes EG:12796),CXCL10,IFNG (Includes EG:15978)	3
Inflammatory Response	phagocytosis	phagocytosis of neutrophils	1.78E-04	CAMP (Includes EG:12796),CXCL10,IFNG (Includes EG:15978)	3
Cell-To-Cell Signaling and Interaction	phagocytosis	phagocytosis of neutrophils	1.78E-04	CAMP (Includes EG:12796),CXCL10,IFNG (Includes EG:15978)	3
Cellular Function and Maintenance	phagocytosis	phagocytosis of neutrophils	1.78E-04	CAMP (Includes EG:12796),CXCL10,IFNG (Includes EG:15978)	3
Lipid Metabolism	concentration	concentration of phospholipid	1.79E-04	CFTR,FOX2,IFNG (Includes EG:15978),IL6,PTGS2	5
Small Molecule Biochemistry	concentration	concentration of phospholipid	1.79E-04	CFTR,FOX2,IFNG (Includes EG:15978),IL6,PTGS2	5
Molecular Transport	concentration	concentration of phospholipid	1.79E-04	CFTR,FOX2,IFNG (Includes EG:15978),IL6,PTGS2	5
Cardiovascular System Development and Function	development	development of cardiovascular system	1.81E-04	ACE2 (Includes EG:302668),APOB,CAMP (Includes EG:12796),CXCL10,CXCR2,FOXA2,IFNG (Includes EG:15978),IL6,PPARGC1A,PTGS2,SELL	11
Cellular Movement	attraction	attraction of memory T lymphocytes	1.85E-04	CXCL10,CXCL9	2
Hematological System Development and Function	attraction	attraction of memory T lymphocytes	1.85E-04	CXCL10,CXCL9	2
Immune Cell Trafficking	attraction	attraction of memory T lymphocytes	1.85E-04	CXCL10,CXCL9	2
Cell-To-Cell Signaling and Interaction	attraction	attraction of memory T lymphocytes	1.85E-04	CXCL10,CXCL9	2
Cell Death and Survival	cell death	cell death of dermal fibroblasts	1.85E-04	CAMP (Includes EG:12796),IFNG (Includes EG:15978)	2
Cell Death and Survival	cell viability	cell viability of peripheral blood leukocytes	1.85E-04	APOB,IL6	2
Lipid Metabolism	concentration	concentration of palmitic acid	1.85E-04	PPARGC1A,PTGS2	2
Small Molecule Biochemistry	concentration	concentration of palmitic acid	1.85E-04	PPARGC1A,PTGS2	2
Molecular Transport	concentration	concentration of palmitic acid	1.85E-04	PPARGC1A,PTGS2	2
Cellular Growth and Proliferation	cytostasis	cytostasis of melanoma cell lines	1.85E-04	IFNG (Includes EG:15978),IL6	2
Tissue Development	fibrogenesis	fibrogenesis of liver	1.85E-04	IFNG (Includes EG:15978),IL6	2
Embryonic Development	fibrogenesis	fibrogenesis of liver	1.85E-04	IFNG (Includes EG:15978),IL6	2
Organismal Development	fibrogenesis	fibrogenesis of liver	1.85E-04	IFNG (Includes EG:15978),IL6	2
Organ Development	fibrogenesis	fibrogenesis of liver	1.85E-04	IFNG (Includes EG:15978),IL6	2
Cellular Assembly and Organization	fibrogenesis	fibrogenesis of liver	1.85E-04	IFNG (Includes EG:15978),IL6	2
Digestive System Development and Function	fibrogenesis	fibrogenesis of liver	1.85E-04	IFNG (Includes EG:15978),IL6	2
Hepatic System Development and Function	fibrogenesis	fibrogenesis of liver	1.85E-04	IFNG (Includes EG:15978),IL6	2
Cellular Movement	invasion	invasion of macrophages	1.85E-04	CXCL10,IFNG (Includes EG:15978)	2
Immune Cell Trafficking	invasion	invasion of macrophages	1.85E-04	CXCL10,IFNG (Includes EG:15978)	2
Inflammatory Response	invasion	invasion of macrophages	1.85E-04	CXCL10,IFNG (Includes EG:15978)	2
Cellular Movement	localization	localization of T lymphocytes	1.85E-04	CXCL10,IFNG (Includes EG:15978)	2
Hematological System Development and Function	localization	localization of T lymphocytes	1.85E-04	IFNG (Includes EG:15978),IL6	2
Immune Cell Trafficking	localization	localization of T lymphocytes	1.85E-04	IFNG (Includes EG:15978),IL6	2
Inflammatory Response	localization	localization of T lymphocytes	1.85E-04	IFNG (Includes EG:15978),IL6	2
Cellular Function and Maintenance	localization	localization of T lymphocytes	1.85E-04	IFNG (Includes EG:15978),IL6	2
Cell-To-Cell Signaling and Interaction	permeability	permeability of tight junctions	1.85E-04	IFNG (Includes EG:15978),IL6	2
Cell Morphology	permeability	permeability of tight junctions	1.85E-04	IFNG (Includes EG:15978),IL6	2
Cellular Assembly and Organization	permeability	permeability of tight junctions	1.85E-04	IFNG (Includes EG:15978),IL6	2
Hematological System Development and Function	stimulation	stimulation of monocytes	1.85E-04	IFNG (Includes EG:15978),IL6	2
Inflammatory Response	stimulation	stimulation of monocytes	1.85E-04	IFNG (Includes EG:15978),IL6	2
Cell-To-Cell Signaling and Interaction	stimulation	stimulation of monocytes	1.85E-04	IFNG (Includes EG:15978),IL6	2
Cellular Growth and Proliferation	stimulation	stimulation of monocytes	1.85E-04	IFNG (Includes EG:15978),IL6	2
Cell-To-Cell Signaling and Interaction	stimulation	stimulation of smooth muscle cells	1.85E-04	IFNG (Includes EG:15978),IL6	2
Cellular Movement	stimulation	stimulation of smooth muscle cells	1.85E-04	IFNG (Includes EG:15978),IL6	2
Hematological System Development and Function	chemotaxis	chemotaxis of phagocytes	1.90E-04	CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (Includes EG:15978),SELL	6
Immune Cell Trafficking	chemotaxis	chemotaxis of phagocytes	1.90E-04	CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (Includes EG:15978),SELL	6
Inflammatory Response	chemotaxis	chemotaxis of phagocytes	1.90E-04	CAMP (Includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (Includes EG:15978),SELL	6
Lipid Metabolism	concentration	concentration of prostaglandin E2	1.91E-04	HPGD,IFNG (Includes EG:15978),PTGS2	3
Small Molecule Biochemistry	concentration	concentration of prostaglandin E2	1.91E-04	HPGD,IFNG (Includes EG:15978),PTGS2	3
Molecular Transport	concentration	concentration of prostaglandin E2	1.91E-04	HPGD,IFNG (Includes EG:15978),PTGS2	3
Drug Metabolism	concentration	concentration of prostaglandin E2	1.91E-04	HPGD,IFNG (Includes EG:15978),PTGS2	3
Hematological Disease	neutrophilia	neutrophilia	1.91E-04	IFNG (Includes EG:15978),IL6,SELL	3
Organismal Injury and Abnormalities	neutrophilia	neutrophilia	1.91E-04	IFNG (Includes EG:15978),IL6,SELL	3
Immunological Disease	neutrophilia	neutrophilia	1.91E-04	IFNG (Includes EG:15978),IL6,SELL	3
Respiratory Disease	injury	injury of lung	1.91E-04	CXCR2,IL6,MARCO,PTGS2	4
Organismal Injury and Abnormalities	injury	injury of lung	1.91E-04	CXCR2,IL6,MARCO,PTGS2	4
Hematological Disease	non-Hodgkin's disease	non-Hodgkin's disease	1.95E-04	CXCL10,CXCL9,IFNG (Includes EG:15978),IL6,M34A1,SELL	6
Cancer	non-Hodgkin's disease	non-Hodgkin's disease	1.95E-04	CXCL10,CXCL9,IFNG (Includes EG:15978),IL6,MS4A1,SELL	6
Lipid Metabolism	concentration	concentration of fatty acid	2.00E-04	APOB,CFTR,HPGD,IFNG (Includes EG:15978),PPARGC1A,PTGS2	6
Small Molecule Biochemistry	concentration	concentration of fatty acid	2.00E-04	APOB,CFTR,HPGD,IFNG (Includes EG:15978),PPARGC1A,PTGS2	6

Molecular Transport	concentration	concentration of fatty acid	2.00E-04	APOB,CFTR,HPGDIIFNG (includes EG:15978),PPARGC1A,PTGS2	6
Hematological System Development and Function	binding	binding of neutrophils	2.19E-04	CXCL10,CXCR2,SELL	3
Immune Cell Trafficking	binding	binding of neutrophils	2.19E-04	CXCL10,CXCR2,SELL	3
Inflammatory Response	binding	binding of neutrophils	2.19E-04	CXCL10,CXCR2,SELL	3
Cell-To-Cell Signaling and Interaction	binding	binding of neutrophils	2.19E-04	CXCL10,CXCR2,SELL	3
Tissue Development	binding	binding of neutrophils	2.19E-04	CXCL10,CXCR2,SELL	3
Hematological System Development and Function	function	function of Th1 cells	2.19E-04	CXCR2,IFNG (includes EG:15978),IL6	3
Cellular Function and Maintenance	function	function of Th1 cells	2.19E-04	CXCR2,IFNG (includes EG:15978),IL6	3
Cellular Movement	influx	influx of neutrophils	2.19E-04	CXCR2,IL6,MARCO	3
Hematological System Development and Function	influx	influx of neutrophils	2.19E-04	CXCR2,IL6,MARCO	3
Immune Cell Trafficking	influx	influx of neutrophils	2.19E-04	CXCR2,IL6,MARCO	3
Inflammatory Response	influx	influx of neutrophils	2.19E-04	CXCR2,IL6,MARCO	3
Organismal Injury and Abnormalities	injury	injury of mice	2.19E-04	IFNG (includes EG:15978),IL6,PTGS2	3
Connective Tissue Development and Function	proliferation	proliferation of hepatic stellate cells	2.19E-04	IFNG (includes EG:15978),IL6,PTGS2	3
Cellular Development	proliferation	proliferation of hepatic stellate cells	2.19E-04	IFNG (includes EG:15978),IL6,PTGS2	3
Cellular Growth and Proliferation	proliferation	proliferation of hepatic stellate cells	2.19E-04	IFNG (includes EG:15978),IL6,PTGS2	3
Hepatic System Development and Function	proliferation	proliferation of hepatic stellate cells	2.19E-04	IFNG (includes EG:15978),IL6,PTGS2	3
Hematological System Development and Function	adhesion	adhesion of lymphocytes	2.19E-04	CXCL9,IFNG (includes EG:15978),IL6,SELL	4
Immune Cell Trafficking	adhesion	adhesion of lymphocytes	2.19E-04	CXCL9,IFNG (includes EG:15978),IL6,SELL	4
Cell-To-Cell Signaling and Interaction	adhesion	adhesion of lymphocytes	2.19E-04	CXCL9,IFNG (includes EG:15978),IL6,SELL	4
Tissue Development	adhesion	adhesion of lymphocytes	2.19E-04	CXCL9,IFNG (includes EG:15978),IL6,SELL	4
Dermatological Diseases and Conditions	plaque psoriasis	plaque psoriasis	2.27E-04	CSTA,CXCL10,CXCL9,IFNG (includes EG:15978)	4
Cellular Movement	chemotaxis	chemotaxis of endothelial cells	2.34E-04	CAMP (includes EG:12796),CXCR2,PTGS2	3
Cardiovascular System Development and Function	chemotaxis	chemotaxis of endothelial cells	2.34E-04	CAMP (includes EG:12796),CXCR2,PTGS2	3
Tissue Development	involution	involution	2.34E-04	CFTR,IL6,PTGS2	3
Organ Morphology	involution	involution	2.34E-04	CFTR,IL6,PTGS2	3
Cellular Development	proliferation	proliferation of mesangial cells	2.34E-04	CXCL10,CXCL9,IL6	3
Renal and Urological System Development and Function	proliferation	proliferation of mesangial cells	2.34E-04	CXCL10,CXCL9,IL6	3
Cellular Growth and Proliferation	proliferation	proliferation of mesangial cells	2.34E-04	CXCL10,CXCL9,IL6	3
Cellular Movement	infiltration	infiltration by lymphocytes	2.35E-04	CXCL10,IFNG (includes EG:15978),IL6,ZC3H12A	4
Hematological System Development and Function	infiltration	infiltration by lymphocytes	2.35E-04	CXCL10,IFNG (includes EG:15978),IL6,ZC3H12A	4
Immune Cell Trafficking	infiltration	infiltration by lymphocytes	2.35E-04	CXCL10,IFNG (includes EG:15978),IL6,ZC3H12A	4
Cell Death and Survival	apoptosis	apoptosis of cholangiocarcinoma cell lines	2.38E-04	IFNG (includes EG:15978),IL6	2
Cell Death and Survival	apoptosis	apoptosis of chronic lymphocytic leukemia B cells	2.38E-04	IL6,MSA1	2
Tumor Morphology	apoptosis	apoptosis of chronic lymphocytic leukemia B cells	2.38E-04	IL6,MSA1	2
Cell Death and Survival	apoptosis	apoptosis of trophoblast cells	2.38E-04	IFNG (includes EG:15978),IL6	2
Embryonic Development	apoptosis	apoptosis of trophoblast cells	2.38E-04	IFNG (includes EG:15978),IL6	2
Cell Death and Survival	cell death	cell death of bronchial epithelial cells	2.38E-04	CAMP (Includes EG:12796),IFNG (includes EG:15978)	2
Cellular Movement	chemoattraction	chemoattraction of monocytes	2.38E-04	CXCL10,CXCL9	2
Hematological System Development and Function	chemoattraction	chemoattraction of monocytes	2.38E-04	CXCL10,CXCL9	2
Immune Cell Trafficking	chemoattraction	chemoattraction of monocytes	2.38E-04	CXCL10,CXCL9	2
Inflammatory Response	chemoattraction	chemoattraction of monocytes	2.38E-04	CXCL10,CXCL9	2
Cell-To-Cell Signaling and Interaction	chemoattraction	chemoattraction of monocytes	2.38E-04	CXCL10,CXCL9	2
Cellular Movement	chemotaxis	chemotaxis of hematopoietic progenitor cells	2.38E-04	CXCL10,CXCL9	2
Hematological System Development and Function	chemotaxis	chemotaxis of hematopoietic progenitor cells	2.38E-04	CXCL10,CXCL9	2
Hematopoiesis	chemotaxis	chemotaxis of hematopoietic progenitor cells	2.38E-04	IFNG (includes EG:15978),IL6	2
Hematological System Development and Function	colony formation	colony formation of granulocyte-macrophage progenitor cells	2.38E-04	IFNG (includes EG:15978),IL6	2
Hematopoiesis	colony formation	colony formation of granulocyte-macrophage progenitor cells	2.38E-04	IFNG (includes EG:15978),IL6	2
Cellular Growth and Proliferation	colony formation	colony formation of granulocyte-macrophage progenitor cells	2.38E-04	IFNG (includes EG:15978),IL6	2
Organismal Functions	healing	delay in healing	2.38E-04	IL6,PTGS2	2
Hematological System Development and Function	generation	generation of naïve T lymphocytes	2.38E-04	IFNG (includes EG:15978),IL6	2
Tissue Development	generation	generation of naïve T lymphocytes	2.38E-04	IFNG (includes EG:15978),IL6	2
Cellular Growth and Proliferation	generation	generation of naïve T lymphocytes	2.38E-04	IFNG (includes EG:15978),IL6	2
Infectious Disease	infection	infection by Human Cytomegalovirus	2.38E-04	IFNG (includes EG:15978),IL6	2
Inflammatory Response	infection	infection by Human Herpesvirus 4	2.38E-04	CXCL10,CXCL9	2
Respiratory Disease	pulmonary interstitial fibrosis	pulmonary interstitial fibrosis	2.38E-04	CXCL10,CXCL9	2
Connective Tissue Disorders	pulmonary interstitial fibrosis	pulmonary interstitial fibrosis	2.38E-04	CFTR,PTGS2	2
Inflammatory Disease	pulmonary interstitial fibrosis	pulmonary interstitial fibrosis	2.38E-04	CFTR,PTGS2	2
Organismal Injury and Abnormalities	pulmonary interstitial fibrosis	pulmonary interstitial fibrosis	2.38E-04	CFTR,PTGS2	2
Tissue Morphology	quantity	quantity of endocrine cell lines	2.38E-04	CFTR,PTGS2	2
Endocrine System Development and Function	quantity	quantity of endocrine cell lines	2.38E-04	IFNC (includes EG:15978),IL6	2
Hematological System Development and Function	regulation	regulation of antigen presenting cells	2.38E-04	IFNG (includes EG:15978),IL6	2
Cellular Function and Maintenance	regulation	regulation of antigen presenting cells	2.38E-04	IFNG (includes EG:15978),IL6	2
Cell-To-Cell Signaling and Interaction	communication	communication of cells	2.38E-04	IFNG (includes EG:15978),IL6	2
Hematological System Development and Function	quantity	quantity of neutrophils	2.39E-04	CAMP (Includes EG:12796),CFTR,CXCL10,CXCL9,IFNG (includes EG:15978),IL6	6
Inflammatory Response	quantity	quantity of neutrophils	2.39E-04	CXCR2,IFNG (includes EG:15978),IL6,MMP8,SELL	5
Tissue Morphology	quantity	quantity of neutrophils	2.39E-04	CXCR2,IFNG (includes EG:15978),IL6,MMP8,SELL	5
Hematological Disease	lymphomagenesis	lymphomagenesis	2.41E-04	CXCR2,IFNG (includes EG:15978),IL6,MMP8,SELL	5
Cancer	lymphomagenesis	lymphomagenesis	2.41E-04	CXCL10,CXCL9,IFNG (includes EG:15978),IL6,MSA1,PTPRC,SELL	7
Cellular Movement	localization	localization of cells	2.49E-04	CAMP (Includes EG:12796),IFNG (includes EG:15978),IL6	3
Cellular Function and Maintenance	localization	localization of cells	2.49E-04	CAMP (Includes EG:12796),IFNG (includes EG:15978),IL6	3
Cell-To-Cell Signaling and Interaction	binding	binding of endothelial cells	2.50E-04	CXCR2,IFNG (includes EG:15978),IL6,SELL	4
Cardiovascular System Development and Function	binding	binding of endothelial cells	2.50E-04	CXCR2,IFNG (includes EG:15978),IL6,SELL	4
Cellular Development	proliferation	proliferation of neuroglia	2.50E-04	CXCR2,IFNG (includes EG:15978),IL6,PTPRC	4
Cellular Growth and Proliferation	proliferation	proliferation of neuroglia	2.50E-04	CXCR2,IFNG (includes EG:15978),IL6,PTPRC	4
Nervous System Development and Function	proliferation	proliferation of neuroglia	2.50E-04	CXCR2,IFNG (includes EG:15978),IL6,PTPRC	4
Cell Morphology	shape change	shape change of connective tissue cells	2.66E-04	CXCR2,IFNG (includes EG:15978),IL6	3
Respiratory Disease	interstitial lung disease	interstitial lung disease	2.67E-04	CFTR,IFNG (includes EG:15978),MSA1,PTGS2	4
Connective Tissue Disorders	interstitial lung disease	interstitial lung disease	2.67E-04	CFTR,IFNG (includes EG:15978),MSA1,PTGS2	4
Cancer	hyperplasia	hyperplasia	2.73E-04	CFTR,FOXA2,GPR83,IFNG (includes EG:15978),IL6,PTGS2,PTPRC,SELL	8
Connective Tissue Disorders	rheumatoid arthritis	rheumatoid arthritis	2.79E-04	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MSA1,PTGS2,PTPRC,SLC7A11	9
Inflammatory Disease	rheumatoid arthritis	rheumatoid arthritis	2.79E-04	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MSA1,PTGS2,PTPRC,SLC7A11	9
Skeletal and Muscular Disorders	rheumatoid arthritis	rheumatoid arthritis	2.79E-04	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MSA1,PTGS2,PTPRC,SLC7A11	9
Immunological Disease	rheumatoid arthritis	rheumatoid arthritis	2.79E-04	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MSA1,PTGS2,PTPRC,SLC7A11	9
Gene Expression	induction	induction of RNA	2.82E-04	IFNG (Includes EG:15978),IL6,PTPRC	3
Hematological System Development and Function	inhibition	inhibition of T lymphocytes	2.82E-04	IFNG (Includes EG:15978),IL6,PTGS2	3
Cellular Growth and Proliferation	inhibition	inhibition of T lymphocytes	2.82E-04	IFNG (Includes EG:15978),IL6,PTGS2	3
Inflammatory Response	phagocytosis	phagocytosis of tumor cell lines	2.82E-04	IFNG (Includes EG:15978),IL6,MARCO	3

Cell-To-Cell Signaling and Interaction	phagocytosis	phagocytosis of tumor cell lines	2.82E-04	IFNG (includes EG:15978),IL6,MARCO	3
Cellular Function and Maintenance	phagocytosis	phagocytosis of tumor cell lines	2.82E-04	IFNG (includes EG:15978),IL6,MARCO	3
Organismal Injury and Abnormalities	injury	injury of liver	2.94E-04	CXCL10,CXCR2,IFNG (includes EG:15978),IL6	4
Gastrointestinal Disease	injury	injury of liver	2.94E-04	CXCL10,CXCR2,IFNG (includes EG:15978),IL6	4
Hepatic System Disease	injury	injury of liver	2.94E-04	CXCL10,CXCR2,IFNG (includes EG:15978),IL6	4
Tissue Morphology	abnormal morphology	abnormal morphology of cervical lymph node	2.97E-04	CXCR2,SELL	2
Lymphoid Tissue Structure and Development	abnormal morphology	abnormal morphology of cervical lymph node	2.97E-04	CXCR2,SELL	2
Organ Morphology	abnormal morphology	abnormal morphology of cervical lymph node	2.97E-04	CXCR2,SELL	2
Small Molecule Biochemistry	binding	binding of proteoglycan	2.97E-04	CXCL10,IFNG (includes EG:15978)	2
Carbohydrate Metabolism	binding	binding of proteoglycan	2.97E-04	CXCL10,IFNG (includes EG:15978)	2
Antimicrobial Response	clearance	clearance of influenza virus	2.97E-04	IFNG (includes EG:15978),IL6	2
Hematological System Development and Function	detachment	detachment of leukocytes	2.97E-04	PTPRC,SELL	2
Immune Cell Trafficking	detachment	detachment of leukocytes	2.97E-04	PTPRC,SELL	2
Cell-To-Cell Signaling and Interaction	detachment	detachment of leukocytes	2.97E-04	PTPRC,SELL	2
Tissue Development	detachment	detachment of leukocytes	2.97E-04	PTPRC,SELL	2
Hematological System Development and Function	erythropoiesis	erythropoiesis of erythroid progenitor cells	2.97E-04	IFNG (includes EG:15978),IL6	2
Cardiovascular System Development and Function	erythropoiesis	erythropoiesis of erythroid progenitor cells	2.97E-04	IFNG (includes EG:15978),IL6	2
Tissue Development	erythropoiesis	erythropoiesis of erythroid progenitor cells	2.97E-04	IFNG (includes EG:15978),IL6	2
Cellular Development	erythropoiesis	erythropoiesis of erythroid progenitor cells	2.97E-04	IFNG (includes EG:15978),IL6	2
Hematopoiesis	erythropoiesis	erythropoiesis of erythroid progenitor cells	2.97E-04	IFNG (includes EG:15978),IL6	2
Organismal Development	erythropoiesis	erythropoiesis of erythroid progenitor cells	2.97E-04	IFNG (includes EG:15978),PPARGC1A	2
Carbohydrate Metabolism	excretion	excretion of carbohydrate	2.97E-04	IFNG (includes EG:15978),PPARGC1A	2
Hematological System Development and Function	generation	generation of Th1 cells	2.97E-04	IFNG (includes EG:15978),IL6	2
Tissue Development	generation	generation of Th1 cells	2.97E-04	IFNG (includes EG:15978),IL6	2
Cellular Growth and Proliferation	generation	generation of Th1 cells	2.97E-04	IFNG (includes EG:15978),IL6	2
Cellular Movement	migration	migration of bone marrow cell lines	2.97E-04	CXCL10,CXCL9	2
Cellular Development	osteoclastogenesis	osteoclastogenesis of macrophage cancer cell lines	2.97E-04	IFNG (includes EG:15978),IL6	2
Skeletal and Muscular System Development and Function	osteoclastogenesis	osteoclastogenesis of macrophage cancer cell lines	2.97E-04	IFNG (includes EG:15978),IL6	2
Cell Movement	transmigration	transmigration of eosinophils	2.97E-04	CXCL10,CXCL9	2
Hematological System Development and Function	transmigration	transmigration of eosinophils	2.97E-04	CXCL10,CXCL9	2
Immune Cell Trafficking	transmigration	transmigration of eosinophils	2.97E-04	CXCL10,CXCL9	2
Hypersensitivity Response	transmigration	transmigration of eosinophils	2.97E-04	CXCL10,CXCL9	2
Hematological System Development and Function	binding	binding of macrophages	3.00E-04	IFNG (includes EG:15978),MARCO,PTPRC	3
Immune Cell Trafficking	binding	binding of macrophages	3.00E-04	IFNG (includes EG:15978),MARCO,PTPRC	3
Inflammatory Response	binding	binding of macrophages	3.00E-04	IFNG (includes EG:15978),MARCO,PTPRC	3
Cell-To-Cell Signaling and Interaction	binding	binding of macrophages	3.00E-04	IFNG (includes EG:15978),MARCO,PTPRC	3
Tissue Development	binding	binding of macrophages	3.00E-04	IFNG (includes EG:15978),MARCO,PTPRC	3
Cell-To-Cell Signaling and Interaction	response	response of mononuclear leukocytes	3.01E-04	CAMP (includes EG:12796),CXCL9,IFNG (includes EG:15978),IL6,PTPRC	5
Cellular Development	differentiation	differentiation of keratinocytes	3.03E-04	CS1A,IFNG (includes EG:15978),IL6,PTGS2	4
Hair and Skin Development and Function	differentiation	differentiation of keratinocytes	3.03E-04	CS1A,IFNG (includes EG:15978),IL6,PTGS2	4
Hematological Disease	eosinophilia	eosinophilia	3.03E-04	Ch313/Ch314,CXCL9,IFNG (includes EG:15978),PPARGC1A	4
Immunological Disease	eosinophilia	eosinophilia	3.03E-04	Ch313/Ch314,CXCL9,IFNG (includes EG:15978),PPARGC1A	4
Lipid Metabolism	steroidogenesis	steroidogenesis	3.33E-04	APOB,CFTR,CXCL10,IFNG (includes EG:15978),IL6	5
Small Molecule Biochemistry	steroidogenesis	steroidogenesis	3.33E-04	APOB,CFTR,CXCL10,IFNG (includes EG:15978),IL6	5
Cellular Movement	cell movement	cell movement of epithelial cells	3.42E-04	CXCL10,IFNG (includes EG:15978),IL6,SELL	4
Cellular Movement	infiltration	infiltration by macrophages	3.52E-04	CXCL10,IFNG (includes EG:15978),PTGS2,SELL	4
Hematological System Development and Function	infiltration	infiltration by macrophages	3.52E-04	CXCL10,IFNG (includes EG:15978),PTGS2,SELL	4
Immune Cell Trafficking	infiltration	infiltration by macrophages	3.52E-04	CXCL10,IFNG (includes EG:15978),PTGS2,SELL	4
Inflammatory Response	infiltration	infiltration by macrophages	3.52E-04	CXCL10,IFNG (includes EG:15978),PTGS2,SELL	4
Endocrine System Disorders	diabetes mellitus	diabetes mellitus	3.58E-04	APOB,FABP2,FOXA2,IFNG (includes EG:15978),IL6,MSA41,PTGS2,SELL,SLC7A11	9
Metabolic Disease	diabetes mellitus	diabetes mellitus	3.58E-04	APOB,FABP2,FOXA2,IFNG (includes EG:15978),IL6,MSA41,PTGS2,SELL,SLC7A11	9
Gastrointestinal Disease	diabetes mellitus	diabetes mellitus	3.58E-04	APOB,FABP2,FOXA2,IFNG (includes EG:15978),IL6,MSA41,PTGS2,SELL,SLC7A11	9
Small Molecule Biochemistry	synthesis	synthesis of nitric oxide	3.59E-04	ACE2 (includes EG:302668),CAMP (includes EG:12796),CFTR,IFNG (includes EG:15978),IL6,PTGS2	6
Cell Signaling	synthesis	synthesis of nitric oxide	3.59E-04	ACE2 (includes EG:302668),CAMP (includes EG:12796),CFTR,IFNG (includes EG:15978),IL6,PTGS2	6
Tissue Development	branching morphogenesis	branching morphogenesis of lung	3.62E-04	FOX4,IL6	2
Organ Morphology	branching morphogenesis	branching morphogenesis of lung	3.62E-04	FOX4,IL6	2
Embryonic Development	branching morphogenesis	branching morphogenesis of lung	3.62E-04	FOX4,IL6	2
Organism Development	branching morphogenesis	branching morphogenesis of lung	3.62E-04	FOX4,IL6	2
Organ Development	branching morphogenesis	branching morphogenesis of lung	3.62E-04	FOX4,IL6	2
Respiratory System Development and Function	branching morphogenesis	branching morphogenesis of lung	3.62E-04	FOX4,IL6	2
Small Molecule Biochemistry	catabolism	catabolism of L-tryptophan	3.62E-04	IFNC (includes EG:15978),IL6	2
Amino Acid Metabolism	catabolism	catabolism of L-tryptophan	3.62E-04	IFNG (includes EG:15978),IL6	2
Neurological Disease	demyelination	demyelination of cells	3.62E-04	IFNG (includes EG:15978),IL6	2
Cellular Compromise	depletion	depletion of B lymphocytes	3.62E-04	IFNG (includes EG:15978),MSA41	2
Hematological System Development and Function	expansion	expansion of Th17 cells	3.62E-04	IFNG (includes EG:15978),IL6	2
Cellular Development	expansion	expansion of Th17 cells	3.62E-04	IFNG (includes EG:15978),IL6	2
Cellular Growth and Proliferation	expansion	expansion of Th17 cells	3.62E-04	IFNG (includes EG:15978),IL6	2
Cellular Movement	infiltration	infiltration by mast cells	3.62E-04	CFTR,SELL	2
Hematological System Development and Function	infiltration	infiltration by mast cells	3.62E-04	CFTR,SELL	2
Immune Cell Trafficking	infiltration	infiltration by mast cells	3.62E-04	CFTR,SELL	2
Inflammatory Response	infiltration	infiltration by mast cells	3.62E-04	CFTR,SELL	2
Infectious Disease	replication	replication of Coxsackievirus B3	3.62E-04	IFNG (includes EG:15978),PTPRC	2
Hematological System Development and Function	proliferation	proliferation of hematopoietic progenitor cells	3.67E-04	CXCL10,IFNG (includes EG:15978),IL6,PTGS2,PTPRC	5
Cellular Development	proliferation	proliferation of hematopoietic progenitor cells	3.67E-04	CXCL10,IFNG (includes EG:15978),IL6,PTGS2,PTPRC	5
Hematopoiesis	proliferation	proliferation of hematopoietic progenitor cells	3.67E-04	CXCL10,IFNG (includes EG:15978),IL6,PTGS2,PTPRC	5
Cellular Growth and Proliferation	proliferation	proliferation of hematopoietic progenitor cells	3.67E-04	CXCL10,IFNG (includes EG:15978),IL6,PTGS2,PTPRC	5
Tissue Development	aggregation	aggregation of cells	3.70E-04	CXCL10,CXCL9,IL6,PTGS2,PTPRC,SELL	6
Cancer	growth	growth of carcinoma	3.77E-04	CXCR2,IL6,PTGS2	3
Tumor Morphology	regression	regression of tumor	3.77E-04	CXCL10,CXCL9,IFNG (includes EG:15978)	3
Cell Death and Survival	cytolysis	cytolysis	3.77E-04	CXCL10,CXCL9,IFNG (includes EG:15978)	3
Inflammatory Disease	experimental autoimmune encephalomyelitis	experimental autoimmune encephalomyelitis	3.89E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),IL6,MSA41,PTPRC	5
Neurological Disease	experimental autoimmune encephalomyelitis	experimental autoimmune encephalomyelitis	3.89E-04	CXCL10,IFNG (includes EG:15978),IL6,PTPRC,SELL	5
Cell Death and Survival	cell death	cell death of B lymphocytes	3.89E-04	CXCL10,IFNG (includes EG:15978),IL6,PTPRC,SELL	5
Lipid Metabolism	quantity	quantity of polyunsaturated fatty acids	3.95E-04	CFTR,HPGD,IFNG (includes EG:15978),PTGS2	4
Small Molecule Biochemistry	quantity	quantity of polyunsaturated fatty acids	3.95E-04	CFTR,HPGD,IFNG (includes EG:15978),PTGS2	4
Molecular Transport	quantity	quantity of polyunsaturated fatty acids	3.95E-04	CFTR,HPGD,IFNG (includes EG:15978),PTGS2	4

Infectious Disease	hepatitis C	hepatitis C	3.98E-04	IFNG (includes EG:15978),IL6,PTPRC	3
Inflammatory Disease	hepatitis C	hepatitis C	3.98E-04	IFNG (includes EG:15978),IL6,PTPRC	3
Gastrointestinal Disease	hepatitis C	hepatitis C	3.98E-04	IFNG (includes EG:15978),IL6,PTPRC	3
Hepatic System Disease	hepatitis C	hepatitis C	3.98E-04	IFNG (includes EG:15978),IL6,PTPRC	3
Small Molecule Biochemistry	synthesis	synthesis of nitrite	3.98E-04	IFNG (includes EG:15978),IL6,ZC3H12A	3
Lipid Metabolism	synthesis	synthesis of lipid	4.12E-04	ACSM3,APOB,CFTR,CXCL10,HPGD,IFNG (includes EG:15978),IL6,PPARGC1A,PTGS2	9
Small Molecule Biochemistry	synthesis	synthesis of lipid	4.12E-04	ACSM3,APOB,CFTR,CXCL10,HPGD,IFNG (includes EG:15978),IL6,PPARGC1A,PTGS2	9
Cell-To-Cell Signaling and Interaction	contact growth inhibition	contact growth inhibition of leukocytes	4.34E-04	IFNG (includes EG:15978),PTPRC	2
Cellular Growth and Proliferation	contact growth inhibition	contact growth inhibition of leukocytes	4.34E-04	IFNG (includes EG:15978),PTPRC	2
Cell Cycle	contact growth inhibition	contact growth inhibition of leukocytes	4.34E-04	IFNG (includes EG:15978),PTPRC	2
Lipid Metabolism	conversion	conversion of arachidonic acid	4.34E-04	CFTR,PTGS2	2
Small Molecule Biochemistry	conversion	conversion of arachidonic acid	4.34E-04	CFTR,PTGS2	2
Cell-To-Cell Signaling and Interaction	induction	induction of tumor cell lines	4.34E-04	IFNG (includes EG:15978),IL6	2
Cellular Growth and Proliferation	induction	induction of tumor cell lines	4.34E-04	IFNG (includes EG:15978),IL6	2
Skeletal and Muscular System Development and Function	muscularization	muscularization	4.34E-04	AEC2 (includes EG:302668),IL6	2
Cell-To-Cell Signaling and Interaction	stimulation	stimulation of cancer cells	4.34E-04	IFNG (includes EG:15978),IL6	2
Cancer	stimulation	stimulation of cancer cells	4.34E-04	IFNG (includes EG:15978),IL6	2
Cellular Growth and Proliferation	stimulation	stimulation of cancer cells	4.34E-04	IFNG (includes EG:15978),IL6	2
Tumor Morphology	stimulation	stimulation of cancer cells	4.34E-04	IFNG (includes EG:15978),IL6	2
Cell-To-Cell Signaling and Interaction	stimulation	stimulation of leukemia cell lines	4.34E-04	IFNG (includes EG:15978),IL6	2
Cellular Growth and Proliferation	stimulation	stimulation of leukemia cell lines	4.34E-04	IFNG (includes EG:15978),IL6	2
Cellular Movement	cell movement	cell movement of peripheral blood leukocytes	4.42E-04	CAMP (includes EG:12796),CXCL9,SELL	3
Hematological System Development and Function	cell movement	cell movement of peripheral blood leukocytes	4.42E-04	CAMP (includes EG:12796),CXCL9,SELL	3
Immune Cell Trafficking	cell movement	cell movement of peripheral blood leukocytes	4.42E-04	CAMP (includes EG:12796),CXCL9,SELL	3
Cardiovascular System Development and Function	development	development of blood vessel	4.62E-04	AEC2 (includes EG:302668),APOB,CAMP (includes EG:12796),CXCL10,CXCR2,IFNG (includes EG:15978),IL6,PTGS2,SELL	9
Organismal Development	development	development of blood vessel	4.62E-04	AEC2 (includes EG:302668),APOB,CAMP (includes EG:12796),CXCL10,CXCR2,IFNG (includes EG:15978),IL6,PTGS2,SELL	9
Cell Death and Survival	killing	killing of bacteria	4.65E-04	CAMP (includes EG:12796),CXCL10,IFNG (includes EG:15978)	3
Cellular Movement	recruitment	recruitment of T lymphocytes	4.65E-04	CXCL10,CXCL9,IFNG (includes EG:15978)	3
Hematological System Development and Function	recruitment	recruitment of T lymphocytes	4.65E-04	CXCL10,CXCL9,IFNG (includes EG:15978)	3
Immune Cell Trafficking	recruitment	recruitment of T lymphocytes	4.65E-04	CXCL10,CXCL9,IFNG (includes EG:15978)	3
Cell-To-Cell Signaling and Interaction	recruitment	recruitment of T lymphocytes	4.65E-04	CXCL10,CXCL9,IFNG (includes EG:15978)	3
Cell-mediated Immune Response	recruitment	recruitment of T lymphocytes	4.65E-04	CXCL10,CXCL9,IFNG (includes EG:15978)	3
Connective Tissue Disorders	rheumatic disease	Rheumatic Disease	4.67E-04	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MMP8,MS4A1,PTGS2,PTPRC,SLC7A11	11
Inflammatory Disease	rheumatic disease	Rheumatic Disease	4.67E-04	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MMP8,MS4A1,PTGS2,PTPRC,SLC7A11	11
Skeletal and Muscular Disorders	rheumatic disease	Rheumatic Disease	4.67E-04	CAMP (includes EG:12796),CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,MMP8,MS4A1,PTGS2,PTPRC,SLC7A11	11
Cellular Growth and Proliferation	colony formation	colony formation of tumor cell lines	4.68E-04	FOXA2,HPGD,IFNG (includes EG:15978),IL6,PTGS2	5
Cellular Development	maturity	maturity of leukocytes	4.68E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),IL6,PTGS2,PTPRC	5
Hematopoiesis	maturity	maturity of leukocytes	4.68E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),IL6,PTGS2,PTPRC	5
Inflammatory Response	pneumonia	pneumonia	4.80E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),MS4A1,PTGS2	4
Respiratory Disease	pneumonia	pneumonia	4.80E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),MS4A1,PTGS2	4
Hematological System Development and Function	stimulation	stimulation of phagocytes	4.89E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),IL6	3
Inflammatory Response	stimulation	stimulation of phagocytes	4.89E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),IL6	3
Cell-To-Cell Signaling and Interaction	stimulation	stimulation of phagocytes	4.89E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),IL6	3
Cellular Growth and Proliferation	stimulation	stimulation of phagocytes	4.89E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),IL6	3
Lipid Metabolism	synthesis	synthesis of eicosanoid	5.03E-04	CFTR,HPGD,IFNG (includes EG:15978),IL6,PTGS2	5
Small Molecule Biochemistry	synthesis	synthesis of eicosanoid	5.03E-04	CFTR,HPGD,IFNG (includes EG:15978),IL6,PTGS2	5
Humoral Immune Response	quantity	quantity of IgM	5.06E-04	IFNG (includes EG:15978),MARCO,PTGS2,ZC3H12A	4
Protein Synthesis	quantity	quantity of IgM	5.06E-04	IFNG (includes EG:15978),MARCO,PTGS2,ZC3H12A	4
Neurological Disease	damage	damage of axons	5.12E-04	IFNG (includes EG:15978),IL6	2
Hematological System Development and Function	differentiation	differentiation of bone-marrow-derived monocyte/macrophage precursor cells	5.12E-04	IFNG (includes EG:15978),IL6	2
Cellular Development	differentiation	differentiation of bone-marrow-derived monocyte/macrophage precursor cells	5.12E-04	IFNG (includes EG:15978),IL6	2
Hematopoiesis	differentiation	differentiation of bone-marrow-derived monocyte/macrophage precursor cells	5.12E-04	IFNG (includes EG:15978),IL6	2
Organismal Functions	healing	healing of bone	5.12E-04	IL6,PTGS2	2
Respiratory Disease	hyperreactivity	hyperreactivity of airway	5.12E-04	CXCL10,CXCL9	2
Organismal Injury and Abnormalities	hyperreactivity	hyperreactivity of airway	5.12E-04	CXCL10,CXCL9	2
Inflammatory Response	immune response	immune response of B lymphocytes	5.12E-04	IFNG (includes EG:15978),PTPRC	2
Cell-To-Cell Signaling and Interaction	immune response	immune response of B lymphocytes	5.12E-04	IFNG (includes EG:15978),PTPRC	2
Humoral Immune Response	immune response	immune response of D lymphocytes	5.12E-04	IFNG (includes EG:15978),PTPRC	2
Cellular Movement	migration	migration of mesangial cells	5.12E-04	CXCL10,CXCL9	2
Renal and Urological System Development and Function	migration	migration of mesangial cells	5.12E-04	CXCL10,CXCL9	2
Cellular Movement	migration	migration of plasmacytoid dendritic cells	5.12E-04	CXCL10,CXCL9	2
Hematological System Development and Function	migration	migration of plasmacytoid dendritic cells	5.12E-04	CXCL10,CXCL9	2
Immune Cell Trafficking	migration	migration of plasmacytoid dendritic cells	5.12E-04	CXCL10,CXCL9	2
Inflammatory Response	migration	migration of plasmacytoid dendritic cells	5.12E-04	CXCL10,CXCL9	2
Hematological System Development and Function	quantity	quantity of peripheral T lymphocyte	5.12E-04	IL6,PTPRC	2
Tissue Morphology	quantity	quantity of peripheral T lymphocyte	5.12E-04	IL6,PTPRC	2
Lipid Metabolism	quantity	quantity of ceramide	5.14E-04	CFTR,IFNG (includes EG:15978),IL6	3
Small Molecule Biochemistry	quantity	quantity of ceramide	5.14E-04	CFTR,IFNG (includes EG:15978),IL6	3
Molecular Transport	quantity	quantity of ceramide	5.14E-04	CFTR,IFNG (includes EG:15978),IL6	3
Hematological System Development and Function	lymphopoiesis	lymphopoiesis	5.40E-04	IFNG (includes EG:15978),IL6,PTPRC	3
Cellular Development	lymphopoiesis	lymphopoiesis	5.40E-04	IFNG (includes EG:15978),IL6,PTPRC	3
Hematopoiesis	lymphopoiesis	lymphopoiesis	5.40E-04	IFNG (includes EG:15978),IL6,PTPRC	3
Protein Synthesis	quantity	quantity of cytokine	5.40E-04	IFNG (includes EG:15978),IL6,PTPRC	3
Hematological System Development and Function	aggregation	aggregation of blood cells	5.50E-04	CXCR2,FABP2,IFNG (includes EG:15978),IL6,PTGS2	5
Tissue Development	aggregation	aggregation of blood cells	5.50E-04	CXCL10,CXCL9,IL6,PTGS2,SELL	5
Lipid Metabolism	synthesis	synthesis of prostaglandin E2	5.62E-04	HPGD,IFNG (includes EG:15978),IL6,PTGS2	4
Small Molecule Biochemistry	synthesis	synthesis of prostaglandin E2	5.62E-04	HPGD,IFNG (includes EG:15978),IL6,PTGS2	4
Drug Metabolism	synthesis	synthesis of prostaglandin E2	5.62E-04	HPGD,IFNG (includes EG:15978),IL6,PTGS2	4
Inflammatory Response	inflammation	inflammation of intestine	5.79E-04	HPGD,IFNG (includes EG:15978),IL6,PTGS2	5
Infectious Disease	infection	Viral Infection	5.91E-04	ACE2 (includes EG:302668),APOB,CAMP (includes EG:12796),CFTR,CXCL10,CXCL9,CXCR2,HPGD,IFNG (includes EG:15978),IL6,PM20D1,PPARGC1A,PTGS2,PTPRC	14
Cell-To-Cell Signaling and Interaction	activation	activation of endothelial cells	5.93E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),IL6	3
Cardiovascular System Development and Function	activation	activation of endothelial cells	5.93E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),IL6	3
Metabolic Disease	albuminuria	albuminuria	5.93E-04	IFNG (includes EG:15978),IL6,PTGS2	3
Renal and Urological Disease	albuminuria	albuminuria	5.93E-04	IFNG (includes EG:15978),IL6,PTGS2	3
Cell Cycle	interphase	arrest in interphase of melanoma cell lines	5.96E-04	IFNG (includes EG:15978),IL6	2
Antimicrobial Response	clearance	clearance of bacteria	5.96E-04	IFNG (includes EG:15978),IL6,MARCO	2
Cell Death and Survival	kill	killing of macrophages	5.96E-04	CAMP (includes EG:12796),IFNG (includes EG:15978)	2
Hematological System Development and Function	regulation	regulation of phagocytes	5.96E-04	IFNG (includes EG:15978),IL6	2
Inflammatory Response	regulation	regulation of phagocytes	5.96E-04	IFNG (includes EG:15978),IL6	2

Cellular Function and Maintenance	regulation	regulation of phagocytes	5.96E-04	IFNG (includes EG:15978),IL6	2
Cell-To-Cell Signaling and Interaction	response	response of macrophage cancer cell lines	5.96E-04	IFNG (includes EG:15978),MARCO	2
Metabolic Disease	glucose metabolism disorder	glucose metabolism disorder	6.05E-04	APOB,FABP2,FOXA2,IFNG (includes EG:15978),IL6,MSA1,PPARCC1A,PTGS2,SELL,SLC7A11	10
Connective Tissue Disorders	systemic lupus erythematosus	systemic lupus erythematosus	6.07E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),MMMP8,PTPRC	4
Inflammatory Disease	systemic lupus erythematosus	systemic lupus erythematosus	6.07E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),MMMP8,PTPRC	4
Skeletal and Muscular Disorders	systemic lupus erythematosus	systemic lupus erythematosus	6.07E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),MMMP8,PTPRC	4
Immunological Disease	systemic lupus erythematosus	systemic lupus erythematosus	6.07E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),MMMP8,PTPRC	4
Cardiovascular System Development and Function	angiogenesis	angiogenesis	6.08E-04	APOB,CAMP (includes EG:12796),CXCL10,CXCL9,IFNG (includes EG:15978),IL6,PTGS2,SELL	8
Inflammatory Response	TH1 immune response	TH1 immune response	6.21E-04	CXCL10,IFNG (includes EG:15978),IL6	3
Cardiovascular System Development and Function	abnormal morphology	abnormal morphology	6.21E-04	ACE2 (includes EG:302668),IL6,PPARGC1A	3
Organ Morphology	abnormal morphology	abnormal morphology	6.21E-04	ACE2 (includes EG:302668),IL6,PPARGC1A	3
Cell-To-Cell Signaling and Interaction	binding	binding	6.21E-04	CXCR2,IFNG (includes EG:15978),SELL	3
Cardiovascular System Development and Function	binding	binding	6.21E-04	CXCR2,IFNG (includes EG:15978),SELL	3
Cellular Movement	migration	migration	6.21E-04	CAMP (includes EG:12796),CXCL10,IFNG (includes EG:15978)	3
Inflammatory Response	ulcerative colitis	ulcerative colitis	6.50E-04	HPGD,MSA1,PTGS2	3
Inflammatory Disease	ulcerative colitis	ulcerative colitis	6.50E-04	HPGD,MSA1,PTGS2	3
Gastrointestinal Disease	ulcerative colitis	ulcerative colitis	6.50E-04	HPGD,MSA1,PTGS2	3
Cellular Growth and Proliferation	inhibition	inhibition	6.54E-04	CXCL10,IFNG (includes EG:15978),IL6,PTGS2	4
Inflammatory Disease	pancreatitis	Pancreatitis	6.80E-04	CFTR,IFNG (includes EG:15978),PTGS2	3
Endocrine System Disorders	pancreatitis	Pancreatitis	6.80E-04	CFTR,IFNG (includes EG:15978),PTGS2	3
Gastrointestinal Disease	pancreatitis	Pancreatitis	6.80E-04	CAMP (includes EG:12796),FABP2,FOXA2,IFNG (includes EG:15978),IL6,PPARGC1A,PTGS2	7
Carbohydrate Metabolism	quantity	quantity	6.83E-04	IFNG (includes EG:15978),IL6	2
Molecular Transport	accumulation	accumulation	6.87E-04	CAMP (includes EG:15978),IFNG (includes EG:15978)	2
Cell Death and Survival	apoptosis	apoptosis	6.87E-04	CAMP (includes EG:12796),SELL	2
Cellular Movement	cell movement	cell movement	6.87E-04	CAMP (includes EG:12796),SELL	2
Hematological System Development and Function	cell movement	cell movement	6.87E-04	CAMP (includes EG:12796),SELL	2
Immune Cell Trafficking	cell movement	cell movement	6.87E-04	CAMP (includes EG:12796),SELL	2
Inflammatory Response	cell movement	cell movement	6.87E-04	CAMP (includes EG:12796),SELL	2
Cellular Movement	cell rolling	cell rolling	6.97E-04	IFNG (includes EG:15978),SELL	2
Hematological System Development and Function	cell rolling	cell rolling	6.97E-04	IFNG (includes EG:15978),SELL	2
Immune Cell Trafficking	cell rolling	cell rolling	6.97E-04	IFNG (includes EG:15978),SELL	2
Cell-mediated Immune Response	cell rolling	cell rolling	6.97E-04	IFNG (includes EG:15978),SELL	2
Cellular Movement	extravasation	extravasation	6.97E-04	IFNG (includes EG:15978),SELL	2
Hematological System Development and Function	extravasation	extravasation	6.97E-04	Ch319/Ch34,SELL	2
Small Molecule Biochemistry	formation	formation	6.97E-04	Ch319/Ch34,SELL	2
Free Radical Scavenging	formation	formation	6.97E-04	IFNG (includes EG:15978),IL6	2
Cell Death and Survival	killing	killing	6.97E-04	IFNG (includes EG:15978),IL6	2
Organismal Injury and Abnormalities	atherogenesis	atherogenesis	7.11E-04	CAMP (includes EG:12796),IFNG (includes EG:15978)	3
Cardiovascular Disease	atherogenesis	atherogenesis	7.11E-04	APOB,CXCL10,IFNG (includes EG:15978)	3
Carbohydrate Metabolism	binding	binding	7.11E-04	APOB,CXCL10,IFNG (includes EG:15978)	3
Cellular Movement	cell movement	cell movement	7.11E-04	CAMP (includes EG:12796),CXCL10,IFNG (includes EG:15978)	3
Hematological System Development and Function	cell movement	cell movement	7.11E-04	CXCL10,CXCL9,IFNG (includes EG:15978)	3
Cellular Movement	migration	migration	7.11E-04	CXCL10,CXCL9,IFNG (includes EG:15978)	3
Hair and Skin Development and Function	migration	migration	7.11E-04	migration of keratinocytes	3
Cellular Movement	mobilization	mobilization	7.11E-04	mobilization of blood cells	3
Hematological System Development and Function	mobilization	mobilization	7.11E-04	mobilization of blood cells	3
Immunological Disease	allergy	allergy	7.21E-04	CFK,IFNG (includes EG:15978),SELL	4
Cellular Movement	cell movement	cell movement	7.21E-04	CFK,IFNG (includes EG:15978),SELL	4
Hematological System Development and Function	cell movement	cell movement	7.21E-04	CFK,IFNG (includes EG:15978),SELL	4
Cellular Movement	cell movement	cell movement	7.21E-04	CFK,IFNG (includes EG:15978),SELL	4
Tumor Morphology	apoptosis	apoptosis	7.42E-04	IFNG (includes EG:15978),IL6,MSA1	3
Cell Death and Survival	apoptosis	apoptosis	7.42E-04	IFNG (includes EG:15978),IL6,MSA1	3
Organismal Injury and Abnormalities	cell death	cell death	7.56E-04	IFNG (includes EG:15978),IL6,PTGS2,SELL	4
Respiratory Disease	fibrosis	fibrosis	7.71E-04	ACE2 (includes EG:302668),IL6,PTGS2,SLC7A11	7
Inflammatory Disease	asthma	asthma	7.75E-04	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978),IL6,PTGS2,PTPRC	4
Cell Death and Survival	asthma	asthma	7.75E-04	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978)	4
Hematological System Development and Function	activation	activation	7.84E-04	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978)	4
Immune Cell Trafficking	activation	activation	7.84E-04	CXCL10,CXCL9,CXCR2,IFNG (includes EG:15978)	4
Inflammatory Response	activation	activation	7.84E-04	IFNG (includes EG:15978),IL6	2
Cell-to-Cell Signaling and Interaction	activation	activation	7.84E-04	IFNG (includes EG:15978),IL6	2
Cellular Movement	chemotaxis	chemotaxis	7.84E-04	IFNG (includes EG:15978),IL6	2
Hematological System Development and Function	chemotaxis	chemotaxis	7.84E-04	IFNG (includes EG:15978),IL6	2
Immune Cell Trafficking	chemotaxis	chemotaxis	7.84E-04	IFNG (includes EG:15978),IL6	2
Inflammatory Response	chemotaxis	chemotaxis	7.84E-04	IFNG (includes EG:15978),IL6	2
Cellular Movement	chemotaxis	chemotaxis	7.84E-04	IFNG (includes EG:15978),IL6	2
Hematological System Development and Function	colon formation	colon formation	7.84E-04	IFNG (includes EG:15978),IL6	2
Hematopoiesis	colon formation	colon formation	7.84E-04	IFNG (includes EG:15978),IL6	2
Cellular Growth and Proliferation	colon formation	colon formation	7.84E-04	IFNG (includes EG:15978),IL6	2
Cancer	hyperplasia	hyperplasia	7.84E-04	FOXA2,IFNG (includes EG:15978)	2
Hematological System Development and Function	induction	induction	7.84E-04	IFNG (includes EG:15978),IL6	2
Cell-To-Cell Signaling and Interaction	induction	induction	7.84E-04	IFNG (includes EG:15978),IL6	2
Cellular Growth and Proliferation	induction	induction	7.84E-04	IFNG (includes EG:15978),IL6	2
Connective Tissue Disorders	damage	damage	7.84E-04	IFNG (includes EG:15978),IL6	2
Skeletal and Muscular Disorders	damage	damage	7.84E-04	IFNG (includes EG:15978),IL6	2
Tissue Morphology	damage	damage	7.84E-04	IFNG (includes EG:15978),IL6	2
Organismal Injury and Abnormalities	damage	damage	7.84E-04	IFNG (includes EG:15978),IL6,PTGS2	3
Infectious Disease	infection	infection	7.84E-04	IFNG (includes EG:15978),IL6,PTGS2	3
Cell Death and Survival	necrosis	necrosis	7.84E-04	IFNG (includes EG:15978),IL6,PTGS2	5
Gastrointestinal Disease	digestive organ tumor	digestive organ tumor	8.29E-04	AADAC,AGR3,CFTR,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MSA1,PTGS2,PTPRC,SLC7A11,UGT2B7	13
Infectious Disease	digestive organ tumor	digestive organ tumor	8.29E-04	AADAC,AGR3,CFTR,CXCL10,CXCR2,FOXA2,IFNG (includes EG:15978),IL6,MSA1,PTGS2,PTPRC,SLC7A11,UGT2B7	13
Hematological System Development and Function	infection	infection	8.31E-04	CXCL10,IFNG (includes EG:15978),IL6,PTGS2	4
Immune Cell Trafficking	accumulation	accumulation	8.42E-04	CXCL10,IFNG (includes EG:15978),IL6,PTGS2	3
Inflammatory Response	accumulation	accumulation	8.42E-04	CXCL10,IFNG (includes EG:15978),IL6,PTGS2	3
Tissue Development	accumulation	accumulation	8.42E-04	CXCL10,IFNG (includes EG:15978),IL6	3
Cellular Movement	infiltration	infiltration	8.42E-04	CXCL10,IFNG (includes EG:15978),IL6	3
Hematological System Development and Function	infiltration	infiltration	8.42E-04	CXCL10,IFNG (includes EG:15978),IL6	3

Immune Cell Trafficking	infiltration	infiltration by T lymphocytes	8.42E-04	CXCL10,IFNG (includes EG:15978),IL6	3
Cell-mediated Immune Response	infiltration	infiltration by T lymphocytes	8.42E-04	CXCL10,IFNG (includes EG:15978),IL6	3
Hematological System Development and Function	quantity	quantity of phagocytes	8.49E-04	CXCL10,CXCR2,IFNG (includes EG:15978),IL6,MMP8,SELL	6
Inflammatory Response	quantity	quantity of phagocytes	8.49E-04	CXCL10,CXCR2,IFNG (includes EG:15978),IL6,MMP8,SELL	6
Tissue Morphology	quantity	quantity of phagocytes	8.49E-04	CXCL10,CXCR2,IFNG (includes EG:15978),IL6,MMP8,SELL	6
Hematological System Development and Function	binding	binding of T lymphocytes	8.77E-04	CXCL10,CXCL9,IFNG (includes EG:15978)	3
Cell-To-Cell Signaling and Interaction	binding	binding of T lymphocytes	8.77E-04	CXCL10,CXCL9,IFNG (includes EG:15978)	3
Tissue Development	binding	binding of T lymphocytes	8.77E-04	CXCL10,CXCL9,IFNG (includes EG:15978)	3
Energy Production	oxidation	oxidation of long chain fatty acid	8.77E-04	FABP2,IL6,PPARGC1A	3
Lipid Metabolism	oxidation	oxidation of long chain fatty acid	8.77E-04	FABP2,IL6,PPARGC1A	3
Small Molecule Biochemistry	oxidation	oxidation of long chain fatty acid	8.77E-04	FABP2,IL6,PPARGC1A	3
Cell-To-Cell Signaling and Interaction	binding	binding of hepatoma cell lines	8.87E-04	CXCL10,IL6	2
Cellular Movement	chemoattraction	chemoattraction of T lymphocytes	8.87E-04	CXCL10,CXCL9	2
Hematological System Development and Function	chemoattraction	chemoattraction of T lymphocytes	8.87E-04	CXCL10,CXCL9	2
Immune Cell Trafficking	chemoattraction	chemoattraction of T lymphocytes	8.87E-04	CXCL10,CXCL9	2
Inflammatory Response	chemoattraction	chemoattraction of T lymphocytes	8.87E-04	CXCL10,CXCL9	2
Cell-To-Cell Signaling and Interaction	chemoattraction	chemoattraction of T lymphocytes	8.87E-04	CXCL10,CXCL9	2
Cell-mediated Immune Response	chemoattraction	chemoattraction of T lymphocytes	8.87E-04	CXCL10,CXCL9	2
Cellular Movement	chemotaxis	chemotaxis of vascular endothelial cells	8.87E-04	CXCR2,PTGS2	2
Cardiovascular System Development and Function	chemotaxis	chemotaxis of vascular endothelial cells	8.87E-04	CXCR2,PTGS2	2
Hematological System Development and Function	differentiation	differentiation of memory B cells	8.87E-04	IL6,PTGS2	2
Humoral Immune Response	differentiation	differentiation of memory B cells	8.87E-04	IL6,PTGS2	2
Cellular Development	differentiation	differentiation of memory B cells	8.87E-04	IL6,PTGS2	2
Hematopoiesis	differentiation	differentiation of memory B cells	8.87E-04	IL6,PTGS2	2
Infectious Disease	quantity	quantity of influenza A virus	8.87E-04	IFNG (includes EG:15978),PTGS2	2
Lipid Metabolism	quantity	quantity of fat	8.87E-04	APOB,ASP2	2
Small Molecule Biochemistry	quantity	quantity of fat	8.87E-04	APOB,FABP2	2
Molecular Transport	quantity	quantity of fat	8.87E-04	APOB,FABP2	2
Organismal Functions	thermoregulation	thermoregulation	9.07E-04	IFNG (includes EG:15978),IL6,PPARGC1A,PTGS2	4
Hematological System Development and Function	adhesion	adhesion of neutrophils	9.13E-04	CXCR2,IL6,SELL	3
Immune Cell Trafficking	adhesion	adhesion of neutrophils	9.13E-04	CXCR2,IL6,SELL	3
Inflammatory Response	adhesion	adhesion of neutrophils	9.13E-04	CXCR2,IL6,SELL	3
Cell-To-Cell Signaling and Interaction	adhesion	adhesion of neutrophils	9.13E-04	CXCR2,IL6,SELL	3
Tissue Development	adhesion	adhesion of neutrophils	9.13E-04	CXCR2,IL6,SELL	3
Neurological Disease	demyelination	demyelination of nervous tissue	9.13E-04	CXCL10,IFNG (includes EG:15978),IL6	3
Cardiovascular Disease	hyperlipidemia	hyperlipidemia	9.13E-04	APOB,IL6,PTGS2	3
Metabolic Disease	hyperlipidemia	hyperlipidemia	9.13E-04	APOB,IL6,PTGS2	3
Cellular Development	differentiation	differentiation of epithelial cells	9.30E-04	CSTA,FOXA2,IFNG (includes EG:15978),IL6,PTGS2	5
Inflammatory Response	antibody response	antibody response	9.30E-04	CAMP (includes EG:12796),IFNG (includes EG:15978),IL6,PTPRC	4
Nervous System Development and Function	myelination	myelination	9.30E-04	CXCL10,CXCR2,IFNG (includes EG:15978),PTPRC	4
Cardiovascular System Development and Function	vasculogenesis	vasculogenesis	9.45E-04	ACE2 (includes EG:302668),APOB,CAMP (includes EG:12796),CXCL10,CXCR2,IFNG (includes EG:15978),IL6,PTGS2	8
Organismal Development	vasculogenesis	vasculogenesis	9.45E-04	ACE2 (includes EG:302668),APOB,CAMP (includes EG:12796),CXCL10,CXCR2,IFNG (includes EG:15978),IL6,PTGS2	8
Cell Death and Survival	cytotoxicity	cytotoxicity of T lymphocytes	9.49E-04	IFNG (includes EG:15978),IL6,PTPRC	3
Vitamin and Mineral Metabolism	quantity	quantity of vitamin	9.49E-04	IFNG (includes EG:15978),IL6,PTGS2	3
Gastrointestinal Disease	diarrhea	diarrhea	9.87E-04	CFTR,PTGS2,PTPRC	3
Cell Death and Survival	apoptosis	apoptosis of macrophage cancer cell lines	9.96E-04	IFNG (includes EG:15978),ZC3H12A	2
Cellular Development	growth	arrest in growth of leukemia cell lines	9.96E-04	IFNG (includes EG:15978),IL6	2
Cellular Growth and Proliferation	growth	arrest in growth of leukemia cell lines	9.96E-04	IFNG (includes EG:15978),IL6	2
Gene Expression	binding	binding of Sis-inducible element	9.96E-04	IFNG (includes EG:15978),IL6	2
Cell-To-Cell Signaling and Interaction	binding	binding of neuroglia	9.96E-04	CXCR2,IFNG (includes EG:15978)	2
Hematological System Development and Function	development	development of Th17 cells	9.96E-04	IFNG (includes EG:15978),IL6	2
Cell-mediated Immune Response	development	development of Th17 cells	9.96E-04	IFNG (includes EG:15978),IL6	2
Cellular Function and Maintenance	development	development of Th17 cells	9.96E-04	IFNG (includes EG:15978),IL6	2
Lymphoid Tissue Structure and Development	development	development of Th17 cells	9.96E-04	IFNG (includes EG:15978),IL6	2
Cellular Development	development	development of Th17 cells	9.96E-04	IFNG (includes EG:15978),IL6	2
Hematopoiesis	development	development of Th17 cells	9.96E-04	IFNG (includes EG:15978),IL6	2
Organismal Injury and Abnormalities	edema	edema of retina	9.96E-04	IFNG (includes EG:15978),IL6	2
Cell-To-Cell Signaling and Interaction	gap junctional intercellular communication	gap junctional intercellular communication of cells	9.96E-04	CTFR,IL6	2
Infectious Disease	infection	infection by hepatitis C virus	9.96E-04	CXCL10,CXCL9	2
Small Molecule Biochemistry	quantity	quantity of nitrite	9.96E-04	IFNC (includes EG:15978),IL6	2
Molecular Transport	quantity	quantity of nitrite	9.96E-04	IFNC (includes EG:15978),IL6	2
Infectious Disease	replication	replication of flavivirus	9.96E-04	CXCL10,IFNG (includes EG:15978)	2
Dermatological Diseases and Conditions	atopic dermatitis	atopic dermatitis	1.00E-03	CAMP (includes EG:12796),CXCL9,CE3D,PPARGC1A,PTGS2	5
Immunological Disease	atopic dermatitis	atopic dermatitis	1.00E-03	CAMP (includes EG:12796),CXCL9,CE3D,PPARGC1A,PTGS2	5
Hematological System Development and Function	differentiation	differentiation of erythroid progenitor cells	1.03E-03	IFNG (includes EG:15978),IL6,PTPRC	3
Cellular Development	differentiation	differentiation of erythroid progenitor cells	1.03E-03	IFNG (includes EG:15978),IL6,PTPRC	3
Hematopoiesis	differentiation	differentiation of erythroid progenitor cells	1.03E-03	IFNG (includes EG:15978),IL6,PTPRC	3
Small Molecule Biochemistry	quantity	quantity of nitric oxide	1.03E-03	ACE2 (includes EG:302668),IFNG (includes EG:15978),IL6	3
Cell Signaling	quantity	quantity of nitric oxide	1.03E-03	ACE2 (includes EG:302668),IFNG (includes EG:15978),IL6	3
Small Molecule Biochemistry	metabolism	metabolism of D-glucose	1.07E-03	FOXA2,IL6,PPARGC1A	3
Carbohydrate Metabolism	metabolism	metabolism of D-glucose	1.07E-03	FOXA2,IL6,PPARGC1A	3
Lymphoid Tissue Structure and Development	morphology	morphology of spleen component	1.07E-03	CXCR2,IFNG (includes EG:15978),MARCO	3
Organ Morphology	morphology	morphology of spleen component	1.07E-03	CXCR2,IFNG (includes EG:15978),MARCO	3
Hematological System Development and Function	development	development of myeloid cells	1.11E-03	IFNG (includes EG:15978),IL6,PTGS2	3
Lymphoid Tissue Structure and Development	development	development of myeloid cells	1.11E-03	IFNG (includes EG:15978),IL6,PTGS2	3
Cellular Development	development	development of myeloid cells	1.11E-03	IFNG (includes EG:15978),IL6,PTGS2	3
Hematopoiesis	development	development of myeloid cells	1.11E-03	IFNG (includes EG:15978),IL6,PTGS2	3
Reproductive System Development and Function	gestation	gestation	1.11E-03	HPGD,IL6,PTGS2	3
Humoral Immune Response	quantity	quantity of IgA	1.11E-03	IFNG (includes EG:15978),IL6,ZC3H12A	3
Protein Synthesis	quantity	quantity of IgA	1.11E-03	IFNG (includes EG:15978),IL6,ZC3H12A	3
Cell Death and Survival	apoptosis	apoptosis of liver cells	1.11E-03	IFNG (includes EG:15978),IL6,PTGS2,PTPRC	4
Cell-To-Cell Signaling and Interaction	attachment	attachment of blood cells	1.11E-03	ACE2 (includes EG:302668),SELL	2
Tissue Development	attachment	attachment of blood cells	1.11E-03	ACE2 (includes EG:302668),SELL	2
Cell Cycle	cell cycle progression	cell cycle progression of liver cells	1.11E-03	IFNG (includes EG:15978),IL6	2
Cell Death and Survival	cell viability	cell viability of sympathetic neuron	1.11E-03	CXCL9,IL6	2
Nervous System Development and Function	cell viability	cell viability of sympathetic neuron	1.11E-03	CXCL9,IL6	2
Neurological Disease	demyelination	demyelination of spinal cord	1.11E-03	CXCL10,IFNG (includes EG:15978),PTGS2	2
Carbohydrate Metabolism	efflux	efflux of carbohydrate	1.11E-03	IFNG (includes EG:15978),PTGS2	2

Cellular Function and Maintenance	flux	flux of chloride	1.11E-03	CFTR,IFNG (includes EG:15978)	2
Molecular Transport	flux	flux of chloride	1.11E-03	CFTR,IFNG (includes EG:15978)	2
Hematological System Development and Function	proliferation	proliferation of microglia	1.11E-03	IL6,PTPRC	2
Cellular Development	proliferation	proliferation of microglia	1.11E-03	IL6,PTPRC	2
Cellular Growth and Proliferation	proliferation	proliferation of microglia	1.11E-03	IL6,PTPRC	2
Nervous System Development and Function	multiple sclerosis	multiple sclerosis	1.13E-03	IFNG (includes EG:15978),MSA1,PTGS2,PTPRC	2
Inflammatory Disease	multiple sclerosis	multiple sclerosis	1.13E-03	IFNG (includes EG:15978),MSA1,PTGS2,PTPRC	2
Skeletal and Muscular Disorders	multiple sclerosis	multiple sclerosis	1.13E-03	IFNG (includes EG:15978),MSA1,PTGS2,PTPRC	2
Neurological Disease	multiple sclerosis	multiple sclerosis	1.13E-03	IFNG (includes EG:15978),MSA1,PTGS2,PTPRC	2
Hematological System Development and Function	stimulation	stimulation of leukocytes	1.13E-03	CAMP (includes EG:12796),IFNG (includes EG:15978),IL6,PTPRC	2
Cell-To-Cell Signaling and Interaction	stimulation	stimulation of leukocytes	1.13E-03	CAMP (includes EG:12796),IFNG (includes EG:15978),IL6,PTPRC	2
Cellular Growth and Proliferation	stimulation	stimulation of leukocytes	1.13E-03	CAMP (includes EG:12796),IFNG (includes EG:15978),IL6,PTPRC	2
Small Molecule Biochemistry	concentration	concentration of hormone	1.17E-03	FABP2,HPGD,IFNG (includes EG:15978),IL6,PPARGC1A,PTGS2	6
Molecular Transport	concentration	concentration of hormone	1.17E-03	FABP2,HPGD,IFNG (includes EG:15978),IL6,PPARGC1A,PTGS2	6
Endocrine System Development and Function	concentration	concentration of hormone	1.17E-03	FABP2,HPGD,IFNG (includes EG:15978),IL6,PPARGC1A,PTGS2	6
Cellular Movement	cell movement	cell movement of muscle cells	1.18E-03	CXCL10,IFNG (includes EG:15978),IL6,PTGS2	4
Lipid Metabolism	homeostasis	homeostasis of lipid	1.18E-03	APOB,FABP2,PPARGC1A,PTGS2	4
Small Molecule Biochemistry	homeostasis	homeostasis of lipid	1.18E-03	APOB,FABP2,PPARGC1A,PTGS2	4
Skeletal and Muscular Disorders	damage	damage of bone	1.19E-03	IFNG (includes EG:15978),IL6,PTGS2	3
Tissue Morphology	damage	damage of bone	1.19E-03	IFNG (includes EG:15978),IL6,PTGS2	3
Organismal Injury and Abnormalities	size	size of tumor	1.19E-03	IL6,PPARGC1A,PTGS2	3
Tumor Morphology	size	size of tumor	1.19E-03	IL6,PPARGC1A,PTGS2	3
Tissue Development	morphology	morphology of lung	1.23E-03	CFTR,FOXA2,PPARGC1A,SELL	4
Organ Morphology	morphology	morphology of lung	1.23E-03	CFTR,FOXA2,PPARGC1A,SELL	4
Embryonic Development	morphology	morphology of lung	1.23E-03	CFTR,FOXA2,PPARGC1A,SELL	4
Organismal Development	morphology	morphology of lung	1.23E-03	CFTR,FOXA2,PPARGC1A,SELL	4
Organ Development	morphology	morphology of lung	1.23E-03	CFTR,FOXA2,PPARGC1A,SELL	4
Regulated System Development and Function	morphology	morphology of lung	1.23E-03	CFTR,FOXA2,PPARGC1A,SELL	4
Cellular Movement	attraction	attraction of neutrophils	1.23E-03	CAMP (includes EG:12796),CXCL10	2
Hematological System Development and Function	attraction	attraction of neutrophils	1.23E-03	CAMP (includes EG:12796),CXCL10	2
Immune Cell Trafficking	attraction	attraction of neutrophils	1.23E-03	CAMP (includes EG:12796),CXCL10	2
Cell-To-Cell Signaling and Interaction	attraction	attraction of neutrophils	1.23E-03	CAMP (includes EG:12796),CXCL10	2
Cellular Function and Maintenance	function	function of keratinocytes	1.23E-03	SELL,TRPV3	2
Hematological System Development and Function	generation	generation of Th17 cells	1.23E-03	IFNG (includes EG:15978),IL6	2
Tissue Development	generation	generation of Th17 cells	1.23E-03	IFNG (includes EG:15978),IL6	2
Cellular Growth and Proliferation	generation	generation of Th17 cells	1.23E-03	IFNG (includes EG:15978),IL6	2
Organismal Injury and Abnormalities	pathogenesis	pathogenesis	1.23E-03	IFNG (includes EG:15978),MSA1	2
Small Molecule Biochemistry	secretion	secretion of nitric oxide	1.23E-03	IFNG (includes EG:15978),PTPRC	2
Molecular Transport	secretion	secretion of nitric oxide	1.23E-03	IFNG (includes EG:15978),PTPRC	2
Cell Signaling	secretion	secretion of nitric oxide	1.23E-03	IFNG (includes EG:15978),PTPRC	2
Humoral Immune Response	quantity	quantity of IgG3	1.23E-03	IFNG (includes EG:15978),PTGS2,ZC3H12A	3
Protein Synthesis	quantity	quantity of IgG3	1.23E-03	IFNG (includes EG:15978),PTGS2,ZC3H12A	3
Cell-To-Cell Signaling and Interaction	signaling	signaling of cells	1.25E-03	CAMP (includes EG:12796),CXCL10,CXCL9,IFNG (includes EG:15978),IL6	5
Cell Death and Survival	apoptosis	apoptosis of lung cancer cell lines	1.25E-03	CAMP (includes EG:12796),FOXA2,IL6,PTGS2	4
Cardiovascular System Development and Function	proliferation	proliferation of endothelial cells	1.27E-03	CAMP (includes EG:12796),CXCL10,IFNG (includes EG:15978),IL6,PTGS2	5
Tissue Development	proliferation	proliferation of endothelial cells	1.27E-03	CAMP (includes EG:12796),CXCL10,IFNG (includes EG:15978),IL6,PTGS2	5
Cellular Development	proliferation	proliferation of endothelial cells	1.27E-03	CAMP (includes EG:12796),CXCL10,IFNG (includes EG:15978),IL6,PTGS2	5
Organismal Development	proliferation	proliferation of endothelial cells	1.27E-03	CAMP (includes EG:12796),CXCL10,IFNG (includes EG:15978),IL6,PTGS2	5
Cellular Growth and Proliferation	proliferation	proliferation of endothelial cells	1.27E-03	CAMP (includes EG:12796),CXCL10,IFNG (includes EG:15978),IL6,PTGS2	5
Lipid Metabolism	synthesis	synthesis of sterol	1.28E-03	APOB,CFTR,IFNG (includes EG:15978)	3
Small Molecule Biochemistry	synthesis	synthesis of sterol	1.28E-03	APOB,CFTR,IFNG (includes EG:15978)	3
Vitamin and Mineral Metabolism	synthesis	synthesis of sterol	1.28E-03	APOB,CFTR,IFNG (includes EG:15978)	3
Cellular Movement	transmigration	transmigration of phagocytes	1.28E-03	CXCR2,IFNG (includes EG:15978),IL6	3
Hematological System Development and Function	transmigration	transmigration of phagocytes	1.28E-03	CXCR2,IFNG (includes EG:15978),IL6	3
Immune Cell Trafficking	transmigration	transmigration of phagocytes	1.28E-03	CXCR2,IFNG (includes EG:15978),IL6	3
Inflammatory Response	transmigration	transmigration of phagocytes	1.28E-03	CXCR2,IFNG (includes EG:15978),IL6	3
Inflammatory Disease	transmigration	transmigration of phagocytes	1.28E-03	CXCR2,IFNG (includes EG:15978),IL6	3
Gastrointestinal Disease	Crrohn's disease	Crrohn's disease	1.28E-03	CAMP (includes EG:12796) HPGD,IL6,PTGS2	4
Lipid Metabolism	concentration	concentration of cholesterol	1.30E-03	APOB,FABP2,IFNG (includes EG:15978),IL6,PTGS2	5
Small Molecule Biochemistry	concentration	concentration of cholesterol	1.30E-03	APOB,FABP2,IFNG (includes EG:15978),IL6,PTGS2	5
Cellular Movement	concentration	concentration of cholesterol	1.30E-03	APOB,FABP2,IFNG (includes EG:15978),IL6,PTGS2	5
Transmembrane Transport	concentration	concentration of cholesterol	1.30E-03	APOB,FABP2,IFNG (includes EG:15978),IL6,PTGS2	5
Hematological System Development and Function	non-insulin-dependent diabetes mellitus	non-insulin-dependent diabetes mellitus	1.30E-03	APOB,FABP2,FOXA2,IL6,PTGS2	5
Metabolic Disease	non-insulin-dependent diabetes mellitus	non-insulin-dependent diabetes mellitus	1.30E-03	APOB,FABP2,FOXA2,IL6,PTGS2	5
Gastrointestinal Disease	non-insulin-dependent diabetes mellitus	non-insulin-dependent diabetes mellitus	1.30E-03	APOB,FABP2,FOXA2,IL6,PTGS2	5
Nutritional Disease	weight gain	weight gain	1.32E-03	IFNG (includes EG:15978),IL6,PPARGC1A,PTGS2,SELL	5
Inflammatory Response	cell-mediated response	cell-mediated response	1.33E-03	CXCL10,IFNG (includes EG:15978),IL6,SELL	4
Cell Morphology	morphology	morphology of tumor cell lines	1.33E-03	FABP2,IFNG (includes EG:15978),IL6,MARCO	4
Hematological System Development and Function	adhesion	adhesion of hematopoietic progenitor cells	1.36E-03	CXCL10,CXCL9	2
Cell-To-Cell Signaling and Interaction	adhesion	adhesion of hematopoietic progenitor cells	1.36E-03	CXCL10,CXCL9	2
Tissue Development	adhesion	adhesion of hematopoietic progenitor cells	1.36E-03	CXCL10,CXCL9	2
Hematopoiesis	adhesion	adhesion of hematopoietic progenitor cells	1.36E-03	CXCL10,CXCL9	2
Cell Death and Survival	apoptosis	apoptosis of hematopoietic progenitor cells	1.36E-03	IFNG (includes EG:15978),IL6	2
Tumor Morphology	apoptosis	apoptosis of hematopoietic progenitor cells	1.36E-03	IFNG (includes EG:15978),IL6	2
Lipid Metabolism	concentration	concentration of myeloid leukemia cells	1.36E-03	IFNG (includes EG:15978),PTGS2	2
Small Molecule Biochemistry	concentration	concentration of myeloid leukemia cells	1.36E-03	IFNG (includes EG:15978),PTGS2	2
Molecular Transport	concentration	concentration of myeloid leukemia cells	1.36E-03	IFNG (includes EG:15978),PTGS2	2
Vitamin and Mineral Metabolism	concentration	concentration of myeloid leukemia cells	1.36E-03	IFNG (includes EG:15978),PTGS2	2
Drug Metabolism	concentration	concentration of myeloid leukemia cells	1.36E-03	IFNG (includes EG:15978),PTGS2	2
Hematological System Development and Function	differentiation	differentiation of effector T lymphocytes	1.36E-03	IFNG (includes EG:15978),IL6	2
Cell-mediated Immune Response	differentiation	differentiation of effector T lymphocytes	1.36E-03	IFNG (includes EG:15978),IL6	2
Cellular Function and Maintenance	differentiation	differentiation of effector T lymphocytes	1.36E-03	IFNG (includes EG:15978),IL6	2
Lymphoid Tissue Structure and Development	differentiation	differentiation of effector T lymphocytes	1.36E-03	IFNG (includes EG:15978),IL6	2
Cellular Development	differentiation	differentiation of effector T lymphocytes	1.36E-03	IFNG (includes EG:15978),IL6	2
Hematopoiesis	differentiation	differentiation of effector T lymphocytes	1.36E-03	IFNG (includes EG:15978),IL6	2
Respiratory Disease	eosinophilia	eosinophilia of airway	1.36E-03	CXCL9,IFNG (includes EG:15978)	2
Hematological Disease	eosinophilia	eosinophilia of airway	1.36E-03	CXCL9,IFNG (includes EG:15978)	2
Immunological Disease	eosinophilia	eosinophilia of airway	1.36E-03	CXCL9,IFNG (includes EG:15978)	2
Hematological System Development and Function	generation	generation of regulatory T lymphocytes	1.36E-03	IFNG (includes EG:15978),IL6	2

Tissue Development	generation	generation of regulatory T lymphocytes	1.36E-03	IFNG (includes EG:15978),IL6	2
Cellular Growth and Proliferation	generation	generation of regulatory T lymphocytes	1.36E-03	IFNC (includes EG:15978),IL6	2
Lipid Metabolism	metabolism	metabolism of arachidonic acid	1.36E-03	CETP,PTGS2	2
Small Molecule Biochemistry	metabolism	metabolism of arachidonic acid	1.36E-03	CETP,PTGS2	2
Cellular Growth and Proliferation	proliferation	proliferation of duct cells	1.36E-03	IFNG (includes EG:15978),IL6	2
Infectious Disease	replication	replication of hepatitis B virus	1.36E-03	IFNG (includes EG:15978),PPARGC1A	2
Cellular Movement	translocation	translocation of cells	1.36E-03	CXCL9,IFNG (includes EG:15978)	2
Cell-To-Cell Signaling and Interaction	detachment	detachment of cells	1.37E-03	CXCL10,PTPRC,SELL	3
Tissue Development	detachment	detachment of cells	1.37E-03	CXCL10,PTPRC,SELL	3
Cardiovascular Disease	hypertrophy	hypertrophy of heart ventricle	1.37E-03	IL6,PPARGC1A,PTGS2	3
Developmental Disorder	hypertrophy	hypertrophy of heart ventricle	1.37E-03	IL6,PPARGC1A,PTGS2	3
Cell Death and Survival	cell viability	cell viability of mononuclear leukocytes	1.39E-03	APOB,IFNG (includes EG:15978),IL6,PTPRC	4
Infectious Disease	endotoxin shock response	endotoxin shock response	1.42E-03	IFNG (includes EG:15978),IL6,PTGS2	3
Hematological Disease	endotoxin shock response	endotoxin shock response	1.42E-03	IFNG (includes EG:15978),IL6,PTGS2	3
Organismal Injury and Abnormalities	endotoxin shock response	endotoxin shock response	1.42E-03	IFNG (includes EG:15978),IL6,PTGS2	3
Hematological System Development and Function	proliferation	proliferation of antigen presenting cells	1.42E-03	IFNG (includes EG:15978),IL6,PTPRC	3
Cellular Development	proliferation	proliferation of antigen presenting cells	1.42E-03	IFNG (includes EG:15978),IL6,PTPRC	3
Cellular Growth and Proliferation	proliferation	proliferation of antigen presenting cells	1.42E-03	IFNG (includes EG:15978),IL6,PTPRC	3
Cell Death and Survival	cell death	cell death of dermal cells	1.47E-03	CAMP (includes EG:12796),IFNG (includes EG:15978),PTGS2	3
Cellular Movement	migration	migration of vascular smooth muscle cells	1.47E-03	IFNG (includes EG:15978),IL6,PTGS2	3
Skeletal and Muscular System Development and Function	migration	migration of vascular smooth muscle cells	1.47E-03	IFNG (includes EG:15978),IL6,PTGS2	3
Lymphoid Tissue Structure and Development	proliferation	proliferation of splenocytes	1.47E-03	CXCL10,IFNG (includes EG:15978),IL6	3
Cellular Development	proliferation	proliferation of splenocytes	1.47E-03	CXCL10,IFNG (includes EG:15978),IL6	3
Cellular Growth and Proliferation	proliferation	proliferation of splenocytes	1.47E-03	CXCL10,IFNG (includes EG:15978),IL6	3
Cellular Development	proliferation	proliferation of liver cells	1.47E-03	CXCR2,IFNG (includes EG:15978),IL6,PTGS2	4
Cellular Growth and Proliferation	proliferation	proliferation of liver cells	1.47E-03	CXCR2,IFNG (includes EG:15978),IL6,PTGS2	4
Hepatic System Development and Function	proliferation	proliferation of liver cells	1.47E-03	CXCR2,IFNG (includes EG:15978),IL6,PTGS2	4
Reproductive System Disease	endometriosis	endometriosis	1.47E-03	CXCL9,CXCL9,EGFR2,IL6,PTGS2,UGT2B7	6
Inflammatory Response	allergic contact dermatitis	allergic contact dermatitis	1.49E-03	CAMP (includes EG:12796),IFNG (Includes EG:15978)	2
Dermatological Diseases and Conditions	allergic contact dermatitis	allergic contact dermatitis	1.49E-03	CAMP (includes EG:12796),IFNG (Includes EG:15978)	2
Immunological Disease	allergic contact dermatitis	allergic contact dermatitis	1.49E-03	CAMP (includes EG:12796),IFNG (Includes EG:15978)	2
Cell-To-Cell Signaling and Interaction	binding	binding of endothelial cell lines	1.49E-03	IL6,SELL	2
Cardiovascular System Development and Function	binding	binding of endothelial cell lines	1.49E-03	IL6,SELL	2
Tissue Development	generation	generation of dendritic cells	1.49E-03	IFNG (includes EG:15978),IL6	2
Cellular Growth and Proliferation	generation	generation of dendritic cells	1.49E-03	IFNG (includes EG:15978),IL6	2
Inflammatory Response	inflammation	inflammation of stomach	1.49E-03	IFNG (includes EG:15978),PTGS2	2
Hematological System Development and Function	maturation	maturation of helper T lymphocytes	1.49E-03	IFNG (includes EG:15978),PTGS2	2
Cellular Development	maturation	maturation of helper T lymphocytes	1.49E-03	IFNG (includes EG:15978),PTGS2	2
Hematopoiesis	maturation	maturation of helper T lymphocytes	1.49E-03	IFNG (includes EG:15978),PTGS2	2

**Supplementary Table 5 : Primers sequences for qRT-PCR and qPCR analysis**

Mouse gene name	Forward primer (5'-3')	Reverse primer (5'-3')
Actb	GAATGGGTCAAGGACTCCTATG	CCATGTCGTCCCAGTTGGTAA
Cldn5	GGGCCGTGGGTGAGCATTC	TCGTCGCGCCGTACGATG
Il6	ACACATGTTCTCTGGAAATCGT	AAGTCATCATCGTTGTTCATACA
Il10	ATACTGCTAACCGACTCCTTAATG	ACCCAGGAAATTCAAATGCTC
Il17a	ACCGCAATGAAGACCCTGATA	GCACTGAGCTTCCCAGAT
Il17f	ATTCCAGAACCGCTCCAGTTC	GGTCTCGAGTGATGTTGTAATCC
Il22	CATCGTCAACCGCACCTT	GCCGGACGCTGTGTTGTTA
Mmp13	TTTATTGTTGCTGCCATGA	TTTTGGGATGCTTAGGGTTG
Ptgs2	GGGCCCTTCCTCCCGTAGCA	CCATGGCCCAGTCCTCGGGT
Reg3b	ATGCTGCTCCCTGCCTGATG	CTAATGCGTGCAGGAGGGTATATTTC
Reg3g	CTGTGGTACCCCTGTCAAGAGC	GGCCTGAATTGCAAGACAT
Spp1	TCCCCAACGGCCGAGGTGAT	GCTCAGAAGCTGGGCAACAGGG
Tnfa	GACCCTCACACTCAGATCATCTTCT	CCACTTGCTGGTTGGTACGA
Bacterial name	Forward primer (5'-3')	Reverse primer (5'-3')
Bacteroides	GGTTCTGAGAGGAAGGTCCC	GCTGCCTCCCGTAGGAGT