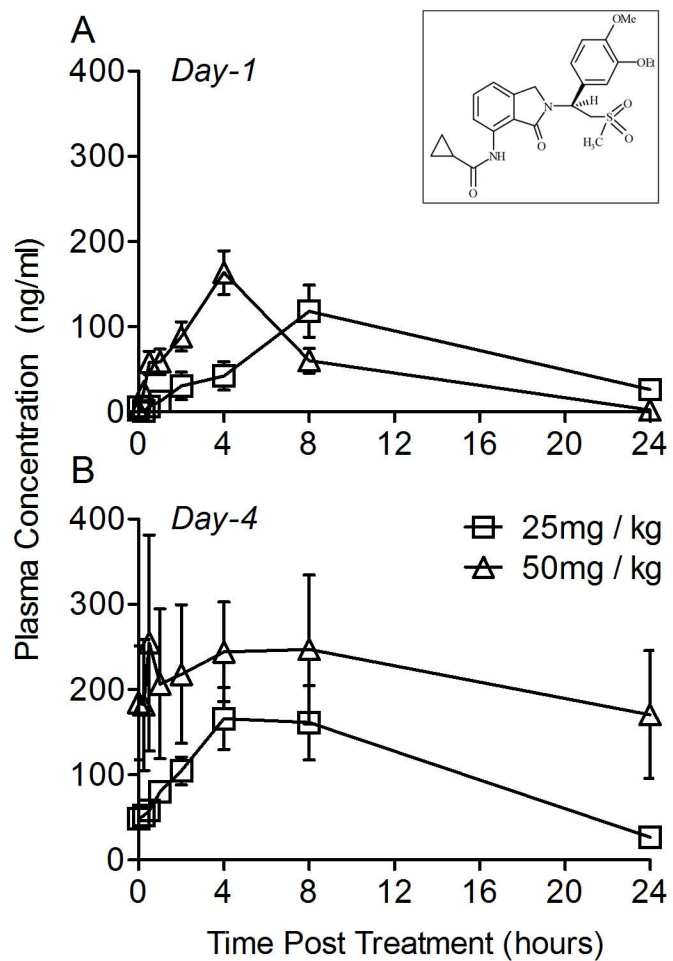
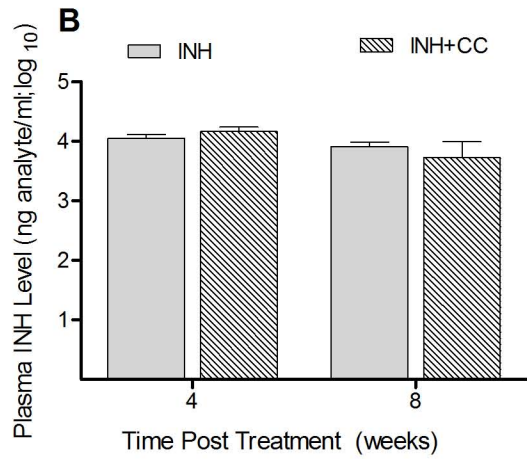
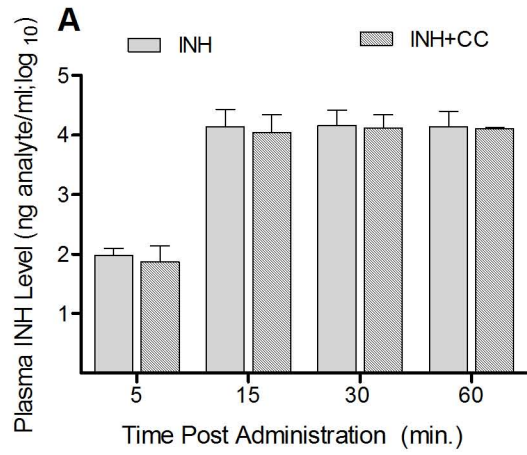


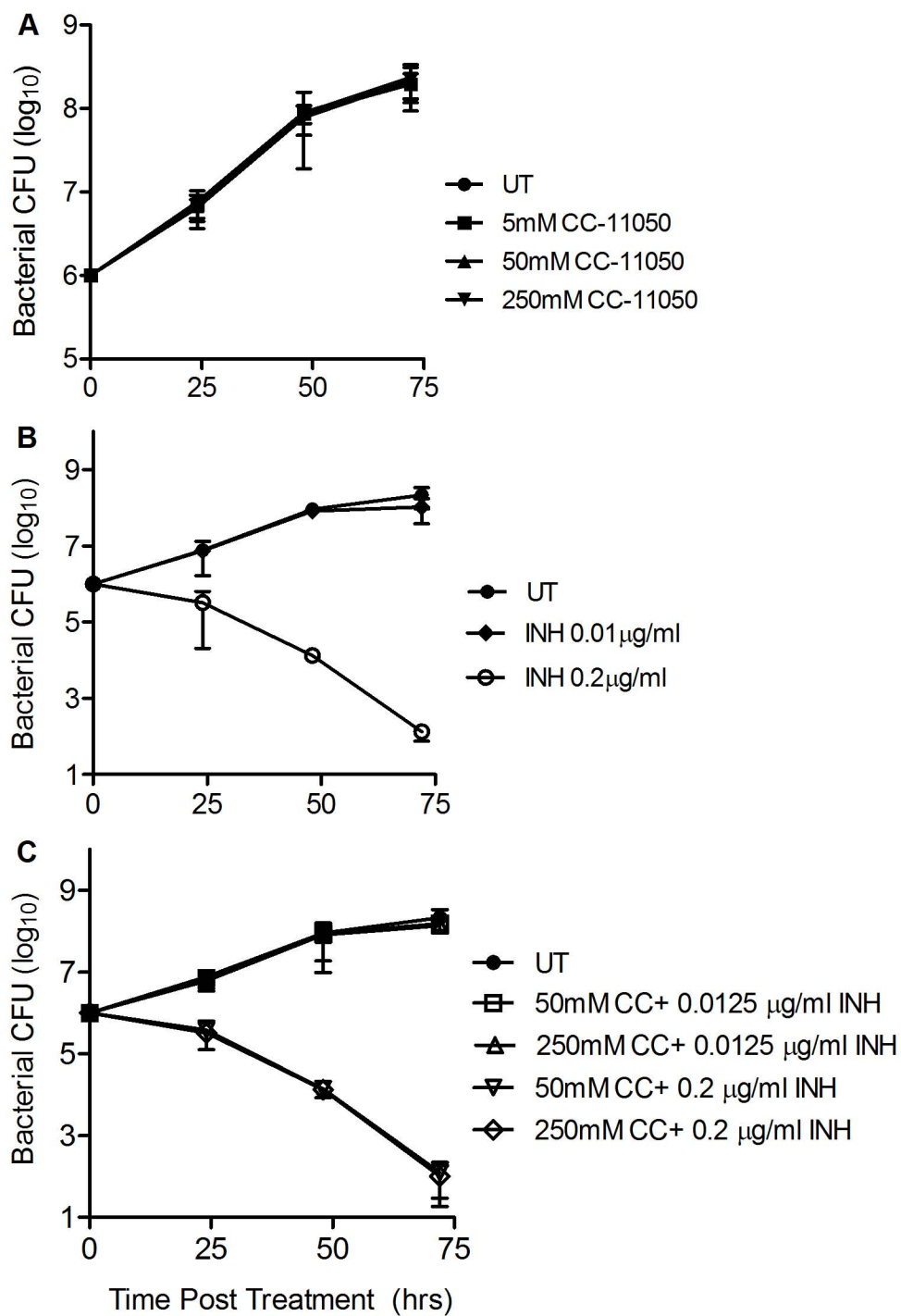
Supplementary Figure 1. Structure and pharmacokinetics of CC-11050



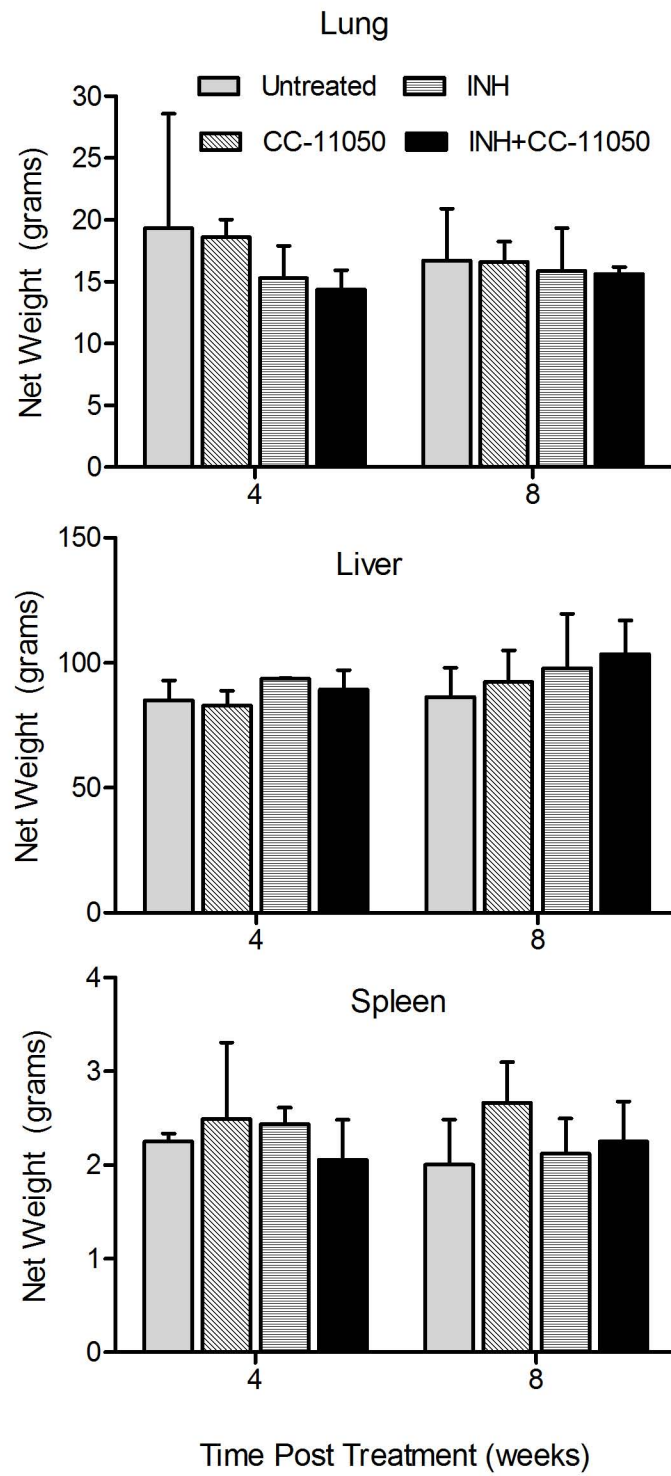
Supplementary Figure 2. Plasma pharmacokinetics of INH



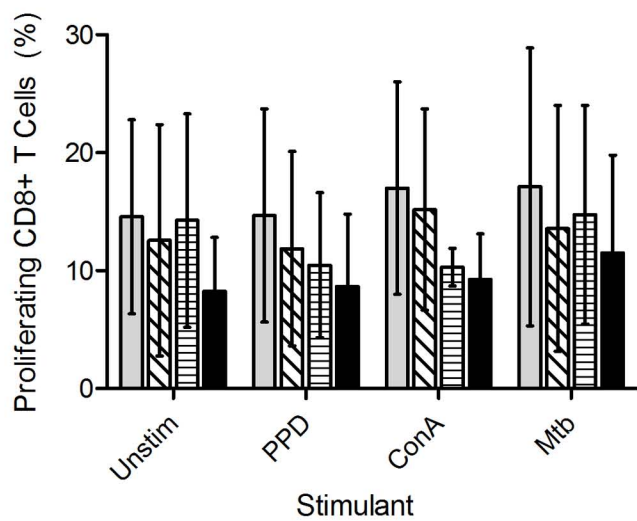
Supplementary Figure 3. In vitro susceptibility of Mtb to INH



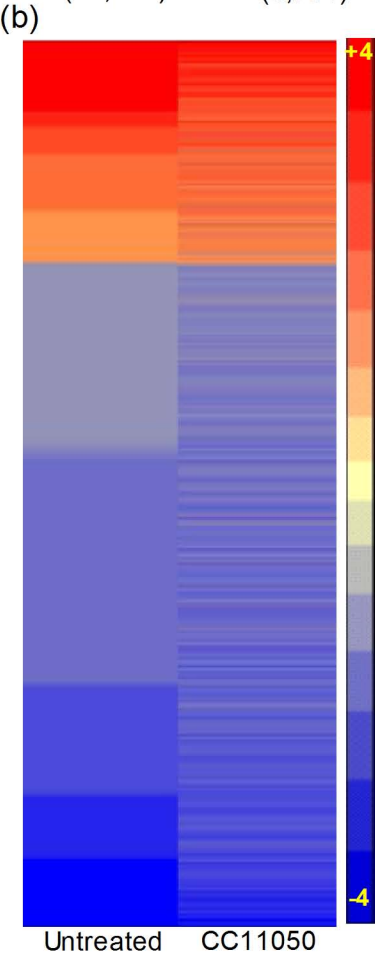
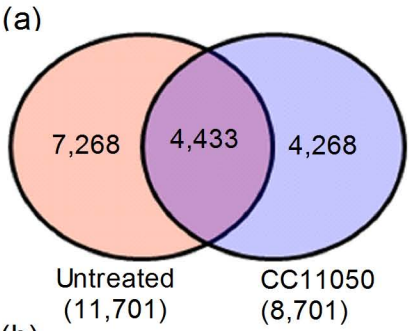
Supplementary Figure 4. Gross weight of the lungs, liver and spleen of Mtb-infected rabbits with or without CC-11050 treatment



Supplementary Figure 5. Proliferative capacity of CD8+ T cells obtained from Mtb-infected rabbits with or without CC-11050 treatment

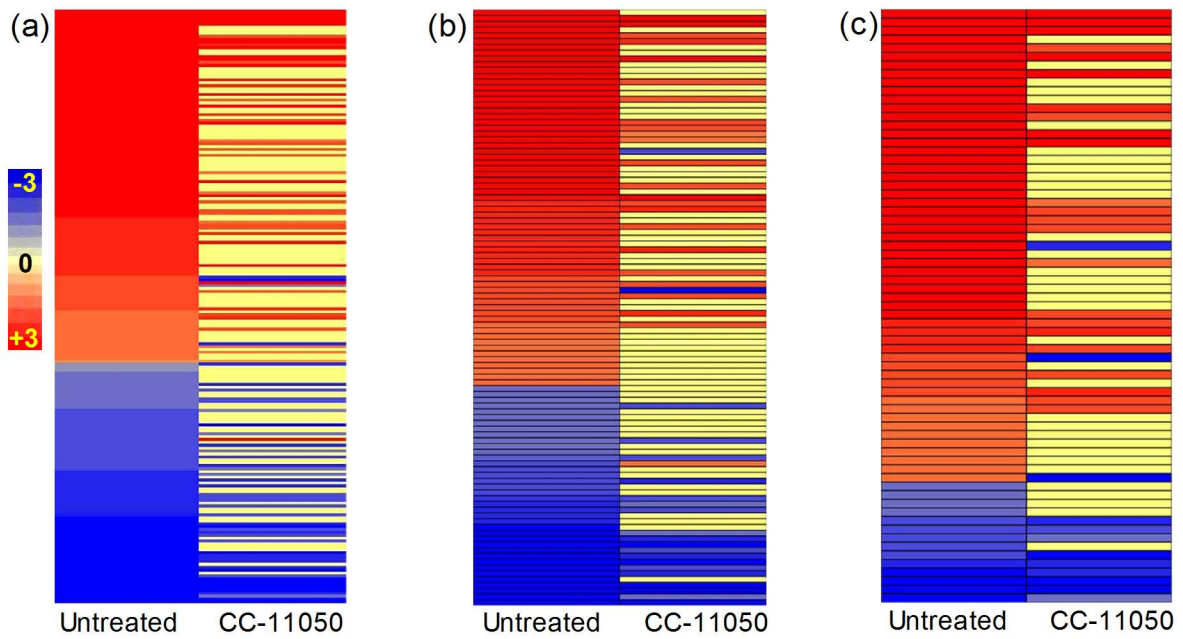


Supplementary Figure 6. Global gene expression analysis of Mtb-infected rabbit lungs with or without CC-11050 treatment

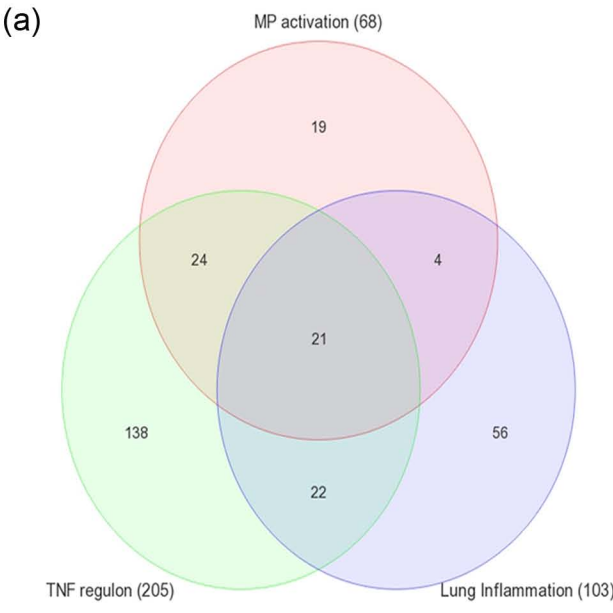


Supplementary Figure 7. Expression profile of genes associated with selected inflammatory response networks in the lungs of Mtb-infected rabbits with or without CC-11050 treatment

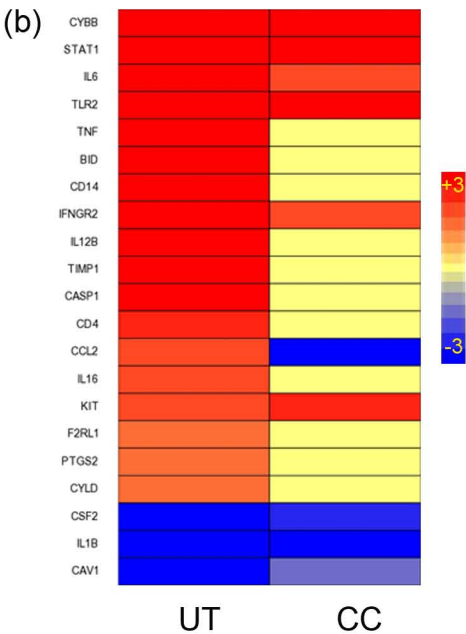
- (a). TNF- α regulon network
- (b). Macrophage activation network
- (c). Lung inflammation network



Supplementary Figure 8. Expression profile of genes common to networks associated with inflammatory response in Mtb-infected rabbit lungs with or without CC-11050 treatment

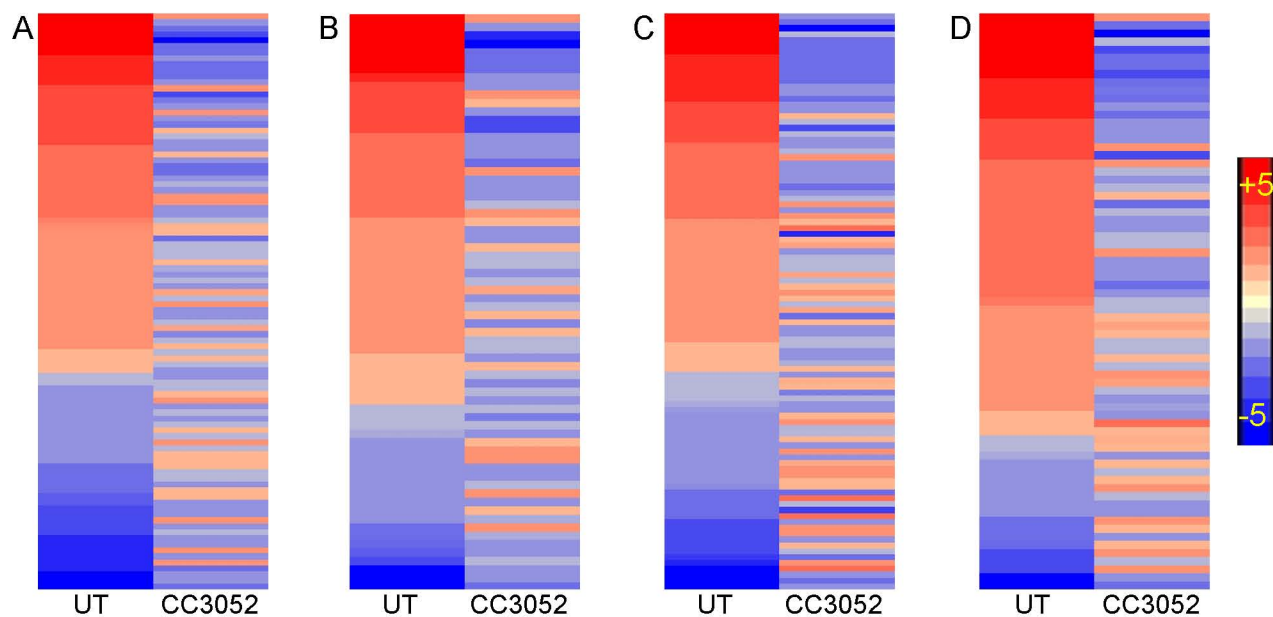


Intensity plot of common genes



Supplementary Figure 9. Expression profile of genes associated with selected inflammatory response and fibrosis networks in the lungs of Mtb-infected rabbits with or without CC-3052 treatment

- A. TNF- α regulon network
- B. Macrophage activation network
- C. Lung inflammation network
- D. Lung fibrosis network



Supplementary Table 1. Description of gene specific primers used in this study

Genes	Forward primer	Reverse primer	Description	Gene ID
<i>1L6</i>	ACTGGCGGAAGTCAATCTGC	CCTGAAC TTGGCCTGAAGGT	interleukin 6	100008733
<i>CCL3</i>	TGTGACCTCCACAGCTAGGT	TGATGACAGCCACTCGGTTG	chemokine ligand 3	6348
<i>CCR2</i>	CCTGTATCTCCGCCTTCACT	CTTACCCGCTCTCGTTGGTA	chemokine (CC) receptor 2	729230
<i>CRP</i>	GCCAGAGGCAAGCATTATTC	CGTGTACTTCACCACGTA CT	C-reactive protein	NM_001082265.1
<i>CXCR2</i>	GGTGGTCTCGCTTGTGAAGG	GTGGCGCTTCTGGATCATCG	chemokine (CXC) receptor 2	100328627
<i>IL12A</i>	TTCCAGTGCCTTAACCACTC	TTCACCGTGCTGGTCTTATC	interleukin 12A	100349007
<i>MCP1</i>	TTCTGTGCCTGCTGCTCATA	GGACACTTGGTGCTGTTGAT	macrophage cationic peptide 1	100009115
<i>S100A9</i>	CACGAGGAGATGCACAAGAA	TTATTTTCCCCACCTGCAAC	calcium binding protein A9	100008704
<i>SPP1</i>	TCTCCTAACACCGCAGAATG	TCTGTAAGCCACACTGTCAC	secreted phosphoprotein 1	100008982
<i>MMP2</i>	CCGATGTCCAGCGAGTAGAC	CCATCTTCTTCTTCACCTCA	matrix metalloproteinase,2	NM_001082209.1
<i>COL1A</i>	GATGGATTACATGGACCAG	TCTTGGGAAGGCTCCTGTCT	collagen, type I	1277
<i>MMP1</i>	AATGGCTAAGGAAGGCCAAG	ATCAGGATGATGCGAGTGAC	matrix metalloproteinase,1	100009110
<i>MMP3</i>	TCTACAACGCCTTTCACAGAC	CCATAGGCACTCCAGAGTTA	matrix metalloproteinase,3	NM_001082280.1
<i>MMP9</i>	CGCCAGCTACGACAAGGACA	AAGTGGTGGCACACCAGAGG	matrix metalloproteinase,9	NM_001082203.1
<i>MMP12</i>	CCA ACTGGCTGTGACCACAA	AGCAGCCTCAATGCCTGAAG	matrix metalloproteinase,12	NM_001082771.1
<i>MMP13</i>	TCTTGAGCTGGACTCATTGC	GCAGGATTCAGAGGATGGTA	matrix metalloproteinase,13	NM_001082037.1
<i>MMP14</i>	CCACAAGATGCCTCCTCAAC	GTAGCCGTCCATCACTTGGT	matrix metalloproteinase,14	NM_001082793.1
<i>DCN</i>	CAAGGTCTTCCTCCTTCTCT	ACCACCAGGTA CTCCGATAA	decorin	100009171
<i>COL3A1</i>	TAGGAGGACTCGCAGGCTAT	GCAGGACCAGATGGACCTAT	collagen, type III, alpha 1	1281
<i>MMP8</i>	GCATTCAGGCCATCTATGGA	TGTAGCTGAGGATGCCTTCT	matrix metalloproteinase,8	4317
<i>GAPDH</i>	GGCGTGAACCACGAGAAGTA	TCCACAATGCCGAAGTGGTC	glyceraldehyde 3-phosphate dehydrogenase	100009074

Supplementary Table 2a. Pharmacokinetics of CC-11050 in rabbits

Dose (mg/kg)	Day	C _{max} (ng/ml) at T _{max}	AUC ₂₄ (ng*h/ml)	Accumulation Ratio*
25	1	118 +/- 52.8	1370 +/- 371	NC
25	4	214 +/- 33.7	2260 +/- 551	1.77 +/- 0.71
50	1	163 +/- 44.8	2380 +/- 702	NC
50	4	280 +/- 126	5130 +/- 3240	2.37 +/- 1.86

*Day 4 AUC_{24h}/Day 1 AUC_{24h}; NC-not calculated.

Supplementary Table 2b. Individual plasma concentrations for CC-11050 and INH in rabbits after two weeks of treatment

	Concentration (ng/ml; Mean (SD))
Treatment Group	CC-11050
INH only	BLOQ
INH+CC-11050	118.02
Untreated	BLOQ
CC-11050 only	178.94

BLOQ: below the level of quantification.

Supplement Table 3. Mean CFU (log10) from individual rabbits

Time (wks)	Untreated									
0	2.9	2.8	3.18	3.12	3.48	3.72	3.77	3.5	3.4	3.5
4	7.37	7.7	7.22	6.09	7.12	6				
8	6.48	7.66	7.63	5.66	5.4	6.8				
12	5.43	6	6.8	6.59	8.5	7.84	6.02			
Time (wks)	CC-11050									
0	2.9	2.8	3.18	3.12	3.48	3.72	3.77	3.5	3.4	3.5
4	7.37	7.7	7.22	6.09	7.12	6				
8	7.12	7.02	6.16	6.9	6.97	5				
12	7.1	5.47	5.26	6	5.86	5	5.71	4.52		
Time (wks)	INH									
0	2.9	2.8	3.18	3.12	3.48	3.72	3.77	3.5	3.4	3.5
4	7.37	7.7	7.22	6.09	7.12	6				
8	6.3	5.92	7.41	6.41	6	5.61				
12	5.14	4.6	6	5.61	5.99	5.3	5.21	4.7		
Time (wks)	INH+CC-11050									
0	2.9	2.8	3.18	3.12	3.48	3.72	3.77	3.5	3.4	3.5
4	7.37	7.7	7.22	6.09	7.12	6				
8	5.68	6.09	6.03	5	5.91	4.3				
12	5.16	3.17	3.92	3.81	3.62	5.2	4.2	4.9		

Supplementary Table 4. qPCR validation of microarray gene expression

Genes	qPCR (se)	Array
<i>1L6</i>	1.071 (0.153)	1.85
<i>CCL3</i>	1.812 (0.395)	2.1
<i>CCR2</i>	1.584 (0.289)	1.32
<i>CRP</i>	5.804 (0.649)	1.76
<i>CXCR2</i>	1.595 (0.309)	1.51
<i>IL12A</i>	1.264 (0.043)	2.28
<i>MCP1</i>	-1.130 (0.044)	-3.02
<i>S100A9</i>	1.1638 (0.259)	2.2
<i>SPP1</i>	-2.08 (0.829)	-2.16

Supplementary Table 5: Top ten SDEG in Mtb infected untreated or CC-11050 treated rabbit lungs

Untreated

<u>Up regulated</u>	<u>Exp. Value</u>
GZMA	711.230
MMP1	109.323
CXCL9	88.635
CLEC4A	61.612
FAM26F	43.008
ATP12A	36.495
ATP1A4	34.821
SLAMF7	34.075
GZMB	32.580
CXCL13	27.808

<u>Down regulated</u>	<u>Exp. Value</u>
CREG2	-24.422
NTS	-23.653
SLC16A12	-16.857
GPM6A	-16.848
UPK1B	-15.987
ITLN1	-12.634
SLC26A3	-12.441
SCG5	-11.666
SRL	-11.495
SEMA3E	-11.303

CC-11050 treated

<u>Up regulated</u>	<u>Exp. Value</u>
GZMA	77.496
CYLC2	27.298
USP12	25.658
SLC17A6	23.049
CXCL9	16.566
PNLIP	16.061
SLC44A5	11.997
MMP1	11.329
MDGA2	10.783
CLEC4A	10.433

<u>Down regulated</u>	<u>Exp. Value</u>
ITLN1	-17.702
ALDH1A2	-16.497
NTS	-9.257

<i>ME3</i>	-8.198
<i>PHYH</i>	-8.090
<i>SPINK2</i>	-7.898
<i>MYOT</i>	-7.119
<i>CPVL</i>	-6.866
<i>UPK1B</i>	-6.400
<i>CXCL8</i>	-6.328

Supplementary Table 6. List of genes and their expression levels in selected networks

<i>SDEG involved in TNFa regulon network</i>		
<i>Symbol</i>	<i>Untreated</i>	<i>CC-11050 treated</i>
<i>CXCL13</i>	27.8077	5.37504
<i>CLEC4E</i>	18.9198	4.2693
<i>CYBB</i>	14.4262	3.38956
<i>STAT1</i>	14.0654	5.23444
<i>OAS2</i>	13.9082	5.04546
<i>KYNU</i>	12.7519	2.74291
<i>CCL5</i>	12.358	0
<i>CFB</i>	12.2153	0
<i>FPR1</i>	11.3865	0
<i>IL6</i>	11.1379	1.84644
<i>TNFSF13B</i>	10.8439	4.70939
<i>CD5</i>	10.8011	3.02577
<i>NCF1</i>	10.7237	2.39023
<i>CXCL10</i>	10.0354	3.58632
<i>PLAU</i>	9.75851	0
<i>SPP1</i>	9.57236	0
<i>TLR2</i>	9.34011	3.80716
<i>RGS1</i>	9.16372	3.54841
<i>TF</i>	7.67308	1.85165
<i>CTSB</i>	7.32619	2.52052
<i>GUSB</i>	7.27346	0
<i>SPHK1</i>	7.27082	0

<i>TNF</i>	7.24064	0
<i>CD86</i>	6.10309	0
<i>PSMB9</i>	5.93947	2.90161
<i>RUNX2</i>	5.87558	0
<i>IL18BP</i>	5.67449	2.28553
<i>BCL2A1</i>	5.63305	0
<i>TAP1</i>	5.5465	0
<i>CD274</i>	5.3788	2.54051
<i>CD3E</i>	5.15547	0
<i>CD40LG</i>	4.76094	1.87891
<i>BID</i>	4.73392	0
<i>CD69</i>	4.71329	8.93939
<i>CTSS</i>	4.69755	0
<i>IL21R</i>	4.50233	0
<i>HLA-DRA</i>	4.48618	2.49508
<i>CD14</i>	4.38048	0
<i>PSMB10</i>	4.26662	1.88834
<i>GRIA1</i>	4.02756	4.20764
<i>IRF5</i>	3.8695	0
<i>BIRC3</i>	3.83273	0
<i>ITGB2</i>	3.83038	0
<i>S100A8</i>	3.79368	0
<i>LTB</i>	3.72229	0
<i>CD28</i>	3.62874	1.47712
<i>IRF8</i>	3.52603	2.00602
<i>IRF1</i>	3.47754	0
<i>IFNGR2</i>	3.43932	1.8924
<i>CSF1R</i>	3.43681	0

<i>B2M</i>	3.3902	1.81578
<i>INHBA</i>	3.37736	0
<i>ACSL1</i>	3.30325	0
<i>NFKBIE</i>	3.26748	0
<i>HLA-C</i>	3.24267	0
<i>IL12B</i>	3.1419	0
<i>ITGAL</i>	3.13904	1.49532
<i>TIMP1</i>	2.98939	0
<i>PSMB8</i>	2.9789	0
<i>NPPA</i>	2.9505	3.16043
<i>C11orf82</i>	2.92906	0
<i>GRN</i>	2.90628	0
<i>CD40</i>	2.88273	0
<i>FBXO32</i>	2.87271	1.50951
<i>AK2</i>	2.85238	3.89079
<i>CIITA</i>	2.80008	0
<i>SLC7A8</i>	2.77249	1.96136
<i>ITGB7</i>	2.75781	0
<i>CASP1</i>	2.73833	0
<i>RRM2</i>	2.69883	1.99582
<i>IL2RA</i>	2.57588	1.60276
<i>HEXB</i>	2.55882	0
<i>NCF2</i>	2.46975	0
<i>PSME2</i>	2.44397	1.50133
<i>TNFSF15</i>	2.37462	1.58851
<i>RFTN1</i>	2.33682	1.92651
<i>SERPINB9</i>	2.33163	0
<i>IRF7</i>	2.30504	2.19894

<i>NFKB2</i>	2.29707	0
<i>CTSZ</i>	2.25932	0
<i>LRRC8C</i>	2.2593	2.7555
<i>PDGFRA</i>	2.23633	0
<i>IDO1</i>	2.18388	0
<i>TIMP2</i>	2.17808	0
<i>CD4</i>	2.16589	0
<i>IL4R</i>	2.14877	0
<i>PCK2</i>	2.12973	0
<i>CTLA4</i>	2.10674	0
<i>COL1A2</i>	2.08269	3.17903
<i>LITAF</i>	2.07957	0
<i>TNFRSF9</i>	2.06428	0
<i>WISP1</i>	2.04103	0
<i>MCL1</i>	1.952	-2.27535
<i>CCL2</i>	1.94896	-3.0179
<i>EDN1</i>	1.90454	2.75582
<i>SMPDL3A</i>	1.90401	-1.46037
<i>IL16</i>	1.79424	0
<i>PIGR</i>	1.77355	1.58886
<i>PSME1</i>	1.75066	0
<i>TBX21</i>	1.71805	0
<i>TAPBP</i>	1.66906	0
<i>ITGAM</i>	1.65944	0
<i>ADRB3</i>	1.60324	0
<i>KIT</i>	1.59277	2.18036
<i>GSTM2</i>	1.54875	0
<i>BGN</i>	1.51776	1.89442

<i>ACE</i>	1.51531	2.12671
<i>CCL1</i>	1.48391	0
<i>ITGA4</i>	1.46196	0
<i>CASP7</i>	1.44851	0
<i>STMN1</i>	1.41627	1.66908
<i>CD83</i>	1.39628	0
<i>F2RL1</i>	1.38788	0
<i>TLR8</i>	1.38513	0
<i>FCGRT</i>	1.36696	0
<i>LOX</i>	1.34615	-2.15553
<i>ACOX1</i>	1.33438	1.39234
<i>PTGS2</i>	1.32614	0
<i>PPARGC1A</i>	1.32498	1.47731
<i>CYLD</i>	1.29265	0
<i>EFNA1</i>	1.28524	0
<i>BIRC2</i>	1.14046	1.13197
<i>NCAM1</i>	-1.02438	-2.08653
<i>CDKN2C</i>	-1.0849	0
<i>DBT</i>	-1.11953	0
<i>PLA2G16</i>	-1.27425	0
<i>TRAF6</i>	-1.29496	0
<i>TNFSF10</i>	-1.32895	0
<i>ECH1</i>	-1.3302	0
<i>DBI</i>	-1.35529	-2.33195
<i>CD82</i>	-1.35712	0
<i>BCKDHB</i>	-1.37294	-1.78782
<i>HERC1</i>	-1.39794	0
<i>TXNIP</i>	-1.42883	0

<i>ARF4</i>	-1.45088	-1.74103
<i>CD247</i>	-1.46004	-1.97349
<i>LCN2</i>	-1.50754	0
<i>CIB2</i>	-1.52438	0
<i>TPT1</i>	-1.57261	-1.15256
<i>HES1</i>	-1.5843	0
<i>CYP1A1</i>	-1.61685	0
<i>VCAM1</i>	-1.63183	0
<i>FGF2</i>	-1.63355	0
<i>PSMA2</i>	-1.64394	-2.59416
<i>VIPR2</i>	-1.67679	0
<i>ROBO1</i>	-1.68618	0
<i>IGFBP5</i>	-1.68912	-1.35629
<i>RGS2</i>	-1.71889	0
<i>THY1</i>	-1.73614	2.51446
<i>HSD11B1</i>	-1.8109	0
<i>ACADM</i>	-1.8637	-2.0408
<i>RGS7</i>	-1.89144	0
<i>RARRES2</i>	-1.89277	-1.48071
<i>IL18</i>	-1.90189	0
<i>PLA2G4A</i>	-1.90356	-2.40989
<i>ADORA2A</i>	-1.90787	0
<i>CH25H</i>	-1.92944	0
<i>RNF149</i>	-1.92968	-2.15522
<i>CTNNB1</i>	-1.93057	-1.51553
<i>WLS</i>	-2.09422	0
<i>IGFBP6</i>	-2.09696	-1.26441
<i>CCL20</i>	-2.11032	0

<i>ANGPT2</i>	-2.11739	-2.0785
<i>GPD1</i>	-2.12202	0
<i>SCNN1B</i>	-2.15696	-2.42077
<i>KITLG</i>	-2.26762	0
<i>ALDH2</i>	-2.33202	0
<i>RNASE4</i>	-2.34052	-1.96516
<i>ALAD</i>	-2.42816	-1.68142
<i>RND3</i>	-2.44171	-1.72475
<i>IL17RD</i>	-2.45162	0
<i>TDRD7</i>	-2.47031	-1.59923
<i>LXN</i>	-2.48021	0
<i>PDZD2</i>	-2.49178	0
<i>KRT23</i>	-2.49771	-1.93479
<i>ITGB6</i>	-2.52538	0
<i>OCLN</i>	-2.59521	0
<i>CSF2</i>	-2.70658	-2.02953
<i>IL1B</i>	-2.71143	-4.15226
<i>TFPI</i>	-2.73683	-1.77196
<i>SOD1</i>	-2.78391	-2.13858
<i>ADAMTS5</i>	-2.80094	-1.82814
<i>PCSK6</i>	-2.91087	0
<i>SEMA3C</i>	-2.92051	-1.92201
<i>OGN</i>	-2.95231	0
<i>CPT1A</i>	-3.10994	0
<i>EMP1</i>	-3.17009	0
<i>DCN</i>	-3.17218	-3.62229
<i>LPL</i>	-3.21357	-2.33424
<i>PPARA</i>	-3.23528	-2.22701

<i>LAMP3</i>	-3.29693	-2.05295
<i>SMURF2</i>	-3.31064	0
<i>DAG1</i>	-3.3143	-2.07702
<i>SMURF1</i>	-3.329	-2.90496
<i>RASD1</i>	-3.34956	0
<i>AGTR1</i>	-3.36595	-1.76157
<i>NEDD9</i>	-3.59528	-2.5899
<i>CDO1</i>	-3.60602	-3.50987
<i>EGFR</i>	-3.68555	-3.0543
<i>DPP4</i>	-3.89939	-4.52489
<i>GHR</i>	-4.83601	-2.70707
<i>WNT7A</i>	-5.27012	-2.26628
<i>CAV1</i>	-5.47809	-1.43909
<i>PENK</i>	-5.62031	-3.81191
<i>ZNF365</i>	-7.19321	-3.59045
<i>SDEG involved in macrophage activation network</i>		
<i>Symbol</i>	Untreated	CC-11050 treated
<i>CXCL13</i>	27.8077	5.37504
<i>NFKBID</i>	4.81049	5.30073
<i>STAT1</i>	14.0654	5.23444
<i>TLR2</i>	9.34011	3.80716
<i>CXCL10</i>	10.0354	3.58632
<i>CYBB</i>	14.4262	3.38956
<i>CD274</i>	5.3788	2.54051
<i>IRF7</i>	2.30504	2.19894
<i>IGJ</i>	6.25816	2.19384

<i>KIT</i>	1.59277	2.18036
<i>IRF8</i>	3.52603	2.00602
<i>LY86</i>	5.85646	1.89766
<i>IFNGR2</i>	3.43932	1.8924
<i>IL6</i>	11.1379	1.84644
<i>B2M</i>	3.3902	1.81578
<i>IL6ST</i>	1.50018	1.81162
<i>GABRA3</i>	2.13962	1.78236
<i>VWF</i>	1.43487	1.71411
<i>TRPV2</i>	2.41192	1.69712
<i>NOD1</i>	2.57257	1.69222
<i>PIGR</i>	1.77355	1.58886
<i>ITGAL</i>	3.13904	1.49532
<i>CD28</i>	3.62874	1.47712
<i>FPR1</i>	11.3865	0
<i>SPP1 (includes EG:20750)</i>	9.57236	0
<i>BPIFA1</i>	8.24569	0
<i>SPHK1</i>	7.27082	0
<i>TNF</i>	7.24064	0
<i>TAP1</i>	5.5465	0
<i>BID</i>	4.73392	0
<i>CD14</i>	4.38048	0
<i>IRF5</i>	3.8695	0
<i>BIRC3</i>	3.83273	0
<i>ITGB2</i>	3.83038	0
<i>LTB</i>	3.72229	0
<i>RAC2</i>	3.32531	0

<i>IL12B</i>	3.1419	0
<i>TIMP1</i>	2.98939	0
<i>GRN</i>	2.90628	0
<i>CD40</i>	2.88273	0
<i>CASP1</i>	2.73833	0
<i>TOP2A</i>	2.64811	0
<i>CD4</i>	2.16589	0
<i>IL16</i>	1.79424	0
<i>ITGAM (includes EG:16409)</i>	1.65944	0
<i>IKBKB</i>	1.40326	0
<i>CD83</i>	1.39628	0
<i>F2RL1</i>	1.38788	0
<i>TLR8</i>	1.38513	0
<i>PTGS2</i>	1.32614	0
<i>CYLD</i>	1.29265	0
<i>LTA</i>	1.27459	0
<i>TNFSF10</i>	-1.32895	0
<i>C1QBP</i>	-1.44234	0
<i>HNRNPD</i>	-1.47079	0
<i>LCN2</i>	-1.50754	0
<i>CAV1</i>	-5.47809	-1.43909
<i>RARRES2</i>	-1.89277	-1.48071
<i>LUM</i>	-1.86244	-1.89639
<i>CSF2</i>	-2.70658	-2.02953
<i>LCP1</i>	3.18949	-2.04624
<i>CBY1</i>	-1.75393	-2.11757
<i>DCLRE1A</i>	-2.48055	-2.38638

<i>TYMS</i>	-1.94008	-2.75204
<i>STAT3</i>	1.25057	-2.75316
<i>CCL2</i>	1.94896	-3.0179
<i>DCN</i>	-3.17218	-3.62229
<i>IL1B</i>	-2.71143	-4.15226
<i>SDEG involved in lung inflammation network</i>		
<i>Symbol</i>	Untreated	CC-11050 treated
<i>ARG1</i>	18.1586	0
<i>CYBB</i>	14.4262	3.38956
<i>STAT1</i>	14.0654	5.23444
<i>CCL5</i>	12.358	0
<i>IL6</i>	11.1379	1.84644
<i>NCF1</i>	10.7237	2.39023
<i>PLAU</i>	9.75851	0
<i>SPP1</i>	9.57236	0
<i>TLR2</i>	9.34011	3.80716
<i>TNF</i>	7.24064	0
<i>CD86</i>	6.10309	0
<i>LGALS3</i>	5.74066	0
<i>CD40LG</i>	4.76094	1.87891
<i>BID</i>	4.73392	0
<i>CTSS</i>	4.69755	0
<i>PTPRC</i>	4.67186	1.56613
<i>MS4A1</i>	4.65916	0
<i>LCP2</i>	4.46924	0
<i>CD14</i>	4.38048	0

<i>IFNAR1</i>	3.57649	1.70404
<i>IFNGR2</i>	3.43932	1.8924
<i>CLCA1</i>	3.39839	1.15025
<i>SELP</i>	3.2436	1.52051
<i>IL12B</i>	3.1419	0
<i>LIPA</i>	3.0335	-2.00638
<i>TIMP1</i>	2.98939	0
<i>ENTPD1</i>	2.9056	1.5757
<i>CIITA</i>	2.80008	0
<i>CASP1</i>	2.73833	0
<i>TOP2A</i>	2.64811	0
<i>NOD1</i>	2.57257	1.69222
<i>TNFAIP8L2</i>	2.52363	0
<i>GPC3</i>	2.50629	3.426
<i>TNFSF15</i>	2.37462	1.58851
<i>TGFB1</i>	2.34997	2.09126
<i>PDGFRA</i>	2.23633	0
<i>TPI1</i>	2.19757	0
<i>MTOR</i>	2.18113	1.96763
<i>CD4</i>	2.16589	0
<i>IL4R</i>	2.14877	0
<i>CTLA4</i>	2.10674	0
<i>MAP2K1</i>	2.10336	2.1419
<i>NBR1</i>	2.09089	0
<i>TNFRSF9</i>	2.06428	0
<i>CCR1</i>	2.06133	0
<i>CASP3</i>	2.04918	1.68297
<i>GIT2</i>	1.98891	0

<i>IGF1</i>	1.96856	1.99985
<i>CCL2</i>	1.94896	-3.0179
<i>CRP</i>	1.90823	1.77819
<i>IL16</i>	1.79424	0
<i>TANK</i>	1.77772	0
<i>KIT</i>	1.59277	2.18036
<i>CBL</i>	1.56745	0
<i>SLC6A2</i>	1.55024	1.65
<i>ITK</i>	1.53474	0
<i>RORA</i>	1.47585	0
<i>ADA</i>	1.45241	0
<i>IKBKB</i>	1.40326	0
<i>F2RL1</i>	1.38788	0
<i>NFE2L2</i>	1.38277	0
<i>FCGRT</i>	1.36696	0
<i>PTGS2</i>	1.32614	0
<i>CYLD</i>	1.29265	0
<i>LTA</i>	1.27459	0
<i>PPP3CB</i>	-1.23811	0
<i>LUC7L3</i>	-1.23856	0
<i>FBN1</i>	-1.27844	0
<i>INSIG1</i>	-1.28992	-1.92175
<i>TRAF6</i>	-1.29496	0
<i>TNFRSF1A</i>	-1.3672	0
<i>FOXA2</i>	-1.41504	0
<i>FOXO3</i>	-1.47902	0
<i>TRAF3IP2</i>	-1.49289	0
<i>SELL</i>	-1.49725	-1.84906

<i>NEO1</i>	-1.52892	0
<i>BCR</i>	-1.54105	0
<i>PPP3CA</i>	-1.5581	-1.91243
<i>NR3C1</i>	-1.60672	1.25902
<i>DUSP1</i>	-1.86747	0
<i>IL18</i>	-1.90189	0
<i>PLA2G4A</i>	-1.90356	-2.40989
<i>ADORA2A</i>	-1.90787	0
<i>IL12A</i>	-1.91473	0
<i>SELENBP1</i>	-2.02197	-1.98049
<i>LDLR</i>	-2.0712	-1.40861
<i>ANXA5</i>	-2.1427	-1.90766
<i>F2</i>	-2.28177	0
<i>ALDH2</i>	-2.33202	0
<i>ITGB6</i>	-2.52538	0
<i>VEGFA</i>	-2.59968	-1.27549
<i>CSF2</i>	-2.70658	-2.02953
<i>IL1B</i>	-2.71143	-4.15226
<i>TFPI</i>	-2.73683	-1.77196
<i>IL4</i>	-2.77449	-2.23974
<i>SCGB1A1</i>	-3.03608	-2.70236
<i>AGTR1</i>	-3.36595	-1.76157
<i>PRDX6</i>	-3.44223	-2.49206
<i>ALB</i>	-3.50069	0
<i>ANXA3</i>	-4.09888	-2.94295
<i>TAC1</i>	-4.62156	-3.50564
<i>CAV1</i>	-5.47809	-1.43909
<i>CCL11</i>	-5.99061	-2.5123

Common SDEG involved in TNF-α, macrophage activation and lung inflammation networks		
<i>Symbol</i>	Untreated	CC-11050 treated
<i>CYBB</i>	14.4262	3.38956
<i>STAT1</i>	14.0654	5.23444
<i>IL6</i>	11.1379	1.84644
<i>TLR2</i>	9.34011	3.80716
<i>TNF</i>	7.24064	0
<i>BID</i>	4.73392	0
<i>CD14</i>	4.38048	0
<i>IFNGR2</i>	3.43932	1.8924
<i>IL12B</i>	3.1419	0
<i>TIMP1</i>	2.98939	0
<i>CASP1</i>	2.73833	0
<i>CD4</i>	2.16589	0
<i>CCL2</i>	1.94896	-3.0179
<i>IL16</i>	1.79424	0
<i>KIT</i>	1.59277	2.18036
<i>F2RL1</i>	1.38788	0
<i>PTGS2</i>	1.32614	0
<i>CYLD</i>	1.29265	0
<i>CSF2</i>	-2.70658	-2.02953
<i>IL1B</i>	-2.71143	-4.15226
<i>CAV1</i>	-5.47809	-1.43909

SDEG involved in lung fibrosis network in rabbits at 12 weeks post infection		
Symbol	Untreated	CC11050
ARG1	18.16	0.00
STAT1	14.07	5.23
IL6	11.14	1.85
NCF1	10.72	2.39
PLAU	9.76	0.00
SPP1	9.57	0.00
CTSK	8.08	0.00
SPHK1	7.27	0.00
TNF	7.24	0.00
LGALS3	5.74	0.00
HCK	4.97	0.00
CD40LG	4.76	1.88
BID	4.73	0.00
HLA-DRA	4.49	2.50
IFNGR2	3.44	0.00
SELP	3.24	0.00
IL12B	3.14	0.00
IL22	3.11	0.00
ENTPD1	2.91	0.00
SERPINE1	2.63	3.61
IL7	2.42	0.00
COL3A1	2.42	2.99
TGFB1	2.35	2.09

<i>PDGFRA</i>	2.24	0.00
<i>CXCL12</i>	2.14	0.00
<i>IGF1</i>	1.97	0.00
<i>KIT</i>	1.59	2.18
<i>ITGA5</i>	1.57	0.00
<i>TSTA3</i>	1.50	0.00
<i>ADA</i>	1.45	0.00
<i>IKBKB</i>	1.40	0.00
<i>TOP1</i>	1.40	1.31
<i>LOX</i>	1.35	0.00
<i>PTGS2</i>	1.33	0.00
<i>HMGA2</i>	1.32	0.00
<i>FOXM1</i>	1.27	0.00
<i>PDE5A</i>	0.00	1.60
<i>IL1RN</i>	0.00	-1.58
<i>RET</i>	-1.08	0.00
<i>TNFRSF1A</i>	-1.37	0.00
<i>PIK3CB</i>	-1.45	-1.62
<i>SELL</i>	-1.50	0.00
<i>BCR</i>	-1.54	0.00
<i>NR3C1</i>	-1.61	0.00
<i>THY1</i>	-1.74	0.00
<i>SCGB3A2</i>	-1.89	0.00
<i>TGFA</i>	-1.90	0.00
<i>PLA2G4A</i>	-1.90	-2.41
<i>ADORA2A</i>	-1.91	0.00
<i>IL12A</i>	-1.91	0.00
<i>ARG2</i>	-1.99	0.00

<i>IL23A</i>	-2.18	0.00
<i>FGFR2</i>	-2.49	0.00
<i>CSF2</i>	-2.71	-2.03
<i>IL1B</i>	-2.71	-4.15
<i>IL4</i>	-2.77	-2.24
<i>TGFB2</i>	-3.34	-3.23
<i>AGTR1</i>	-3.37	0.00
<i>CAV1</i>	-5.48	0.00

