

	+Klebsiella		
	WT	<i>Asc</i> ^{-/-}	<i>Nlrp3</i> ^{-/-}
GM-CSF	0	35	0
IL-1 α	6258	6859	4687
IL-1 β	5455	777	460
IL-2	2	20	13
IL-3	0	0	0
IL-4	0	1	0
IL-5	0	0	0
IL-6	13042	15840	7901
IL-9	9	25	0
IL-10	289	78	76
IL-13	0	0	1
IL-17 α	195	519	68
KC	36031	37240	38640
MCP-1	3709	5183	3679
M-CSF	1	54	0
RANTES	22593	27708	25863
TNF α	3582	4964	2256
VEGF	850	786	751

Table S1. Inflammatory cytokine and chemokine profiles in *Klebsiella* infected primary macrophage were quantified by RayBiotech custom Quantibody service.

	% MAC (\pm SEM)	% NEU (\pm SEM)	% EOS (\pm SEM)	% LYMP (\pm SEM)
+Klebsiella WT Mock	100.0 (\pm 0.0)	0.0 (\pm 0.0)	0.0 (\pm 0.0)	0.0 (\pm 0.0)
WT	19.0 (\pm 3.2)	79.3 (\pm 3.8)	0.5 (\pm 0.3)	1.2 (\pm 0.7)
<i>Asc</i> ^{-/-}	14.0 (\pm 3.6)	82.8 (\pm 4.1)	0.1 (\pm 0.1)	3.0 (\pm 0.9)
<i>Nlrp3</i> ^{-/-}	13.3 (\pm 2.0)	84.6 (\pm 2.8)	0.7 (\pm 0.4)	1.4 (\pm 0.8)
<i>Nlrc4</i> ^{-/-}	13.4 (\pm 1.8)	82.2 (\pm 1.8)	0.5 (\pm 0.2)	3.9 (\pm 1.2)

Table S2. Cellular composition of the BALF from Klebsiella infected mice. No significant differences were observed in the cellular composition of the BALF. BALF cellularity was assessed via morphology and differential staining of samples following cytopsin, red blood cell lysis, and total cell count assessments. Differentially stained cytopsin preparations were assessed by a trained reviewer (I.C.A.) at 20x and 40x magnification and the composition of each BALF sample was determined as the percent of macrophages/monocytes, neutrophils, eosinophils and lymphocyte present. No fewer than 200 cells per BALF sample were evaluated Mock, n=5; *Nlrp3*^{-/-}, n=9; *Nlrc4*^{-/-}, n=5; *Asc*^{-/-}, n=4; WT, n=12.