

	Mean	SD	95% CI	P-value
Lyz				
WT	0.94	0.21	0.78-1.10	0.94
Nod2 ^{-/-}	0.93	0.22	0.78-1.09	
Reg3γ				
WT	0.97	0.42	0.65-1.30	0.74
Nod2 ^{-/-}	0.92	0.25	0.71-1.12	
Ang4				
WT	1.12	0.43	0.86-1.52	0.49
Nod2 ^{-/-}	1.35	0.53	0.96-1.72	
PanCrp				
WT	0.93	0.26	0.73-1.13	0.73
Nod2 ^{-/-}	0.88	0.30	0.67-1.10	
CRS1C				
WT	0.96	0.28	0.74-1.18	0.04
Nod2 ^{-/-}	0.73	0.12	0.64-0.82	
CRS4C				
WT	N.D.	-	-	-
Nod2 ^{-/-}	N.D.	-	-	-
Defa4				
WT	N.D.	-	-	-
Nod2 ^{-/-}	N.D.	-	-	-
Defa20				
WT	1.18	0.60	0.76-1.61	0.67
Nod2 ^{-/-}	1.07	0.52	0.67-1.47	
Defa3				
WT	1.14	0.45	0.76-1.52	0.53
Nod2 ^{-/-}	1.30	0.55	0.84-1.77	
Defa5				
WT	1.05	0.24	0.85-1.25	0.11
Nod2 ^{-/-}	0.89	0.16	0.77-1.01	

Supplemental Table 1. *WT and Nod2^{-/-} antimicrobial peptide mRNA expression levels.* This table displays the mean, standard deviation (SD), 95% confidence interval (CI), and p-value (based on Student's t-test) for all antimicrobial peptide transcript levels generated in this study (n = 8-10 mice/group for WT and *Nod2^{-/-}* mice).