

Table S2. Oligonucleotide Sequences Used in this Study

Bacterial Screening:	DNA Sequence (5'-3')
Universal27 forward	AGAGTTTGATCCTGGCTCAG
Universal1391 reverse	GACGGGCGGTGWGTRCA
<i>Myd88</i> Genotyping:	
Myd88 forward	TGGCATGCCTCCATCATAGTTA
Myd88 reverse	GTCAGAAACAACCACCACCAT
Myd88-lac reverse	ATCGCCTTCTATCGCCTTCTTG
Quantitative Real-Time PCR:	
L32 forward	CCTCTGGTGAAGCCCAAGATC
L32 reverse	TCTGGGTTTCCGCCAGTTT
SAA1/2 forward	GTAATTGGGGTCTTTGCC
SAA1/2 reverse	TTCTGCTCCCTGCTCCTG
SAA3 forward	TGCCATCATTCTTTGCATCTTGA
SAA3 reverse	CCGTGAACTTCTGAACAGCCT
TNF- α forward	CCAGACCCTCACACTCA
TNF- α reverse	CACTTGGTGGTTTGCTACGAC
TLR4 forward	ATGGCATGGCTTACACCACC
TLR4 reverse	GAGGCCAATTTTGTCTCCACA
Electromobility Shift Assays:	
NF- κ B biotinylated +	biotin-AGTTGAGGGGACTTTCCCAGGC
NF- κ B biotinylated -	biotin-GCCTGGGAAAGTCCCCTCAACT
NF- κ B unlabeled +	AGTTGAGGGGACTTTCCCAGGC
NF- κ B unlabeled -	GCCTGGGAAAGTCCCCTCAACT
NF- κ B mutated +	AGTTGAGCTCACTTTCCCAGGC
NF- κ B mutated -	GCCTGGGAAAGTGAGCTCAACT