

Table S1: Related to STAR Methods**Primer sequences for genotyping and qRT-PCR studies**

Target	Note	Sequence
<i>Ccl2</i>	Forward	TGGCTCAGCCAGATGCAGT
	Reverse	TTGGGATCATCTTGCTGGTG
<i>Cxcl10</i>	Forward	GCCGTCATTTTCTGCCTCA
	Reverse	CGTCCTTGCGAGAGGGATC
<i>Ccr2</i>	Forward	GTTACCTCAGTTCATCCA
	Reverse	CAAGGCTCACCATCATCGTAGTC
<i>Cxcr3</i>	Forward	AGTTTTGGCCCGCAACTGT
	Reverse	GGATTGAGGCAGCAGTGCAT
<i>Il1b</i>	Forward	ACCTGTCCTGTGTAATGAAAGACG
	Reverse	TGGGTATTGCTTGGGATCCA
<i>Tnfa</i>	Forward	GCACAGAAAGCATGATCCG
	Reverse	GCCCCCATCTTTTGGG
<i>Gapdh</i>	Forward	GGCAAATTCAACGGCACAGT
	Reverse	AGATGGTGATGGGCTTCCC
WNV-E	Forward	TCAGCGATCTCTCCACCAAAG
	Reverse	GGGTCAGCCGTTTGTGATTG
RIPK3-2xFV (intact transgene)	Forward	CCAGAGGCCACTTGTGTAGCG
	Reverse	CGCTTTAGAAGCCTTCAGGTTGAC
RIPK3-2xFV (deleted stop cassette)	Forward	AGCACTTGCTCTCCCAAAGTC
	Reverse	TCACCTCCCAGGATATCTTCTTCG
Rosa26 locus	Forward	AGCACTTGCTCTCCCAAAGTC
	Reverse	CCGACAAAACCGAAAATCTGTGGG
Mox2-Cre	Common	GGGACCACCTTCTTTTGGCTTC
	WT	AAGATGTGGAGAGTTCCGGGGTAG
	Mutant	CCAGATCCTCCTCAGAAATCAGC