

Supplementary Table S1

	Forward (5'-3' sequence)	Reverse (5'-3' sequence)
GAPDH	GGAGCGAGATCCCTCCAAAAT	GGCTGTTGTCATACTTCTCATGG
C5AR1	ATCTTTGCAGTCGTCTTCCTG	CGGCTACCGCCAAGTTGAG
IL10	ATGCCCCAAGCTGAGAACCAAGA	GCTGGGTCAGCTATCCCAGAG
PPP4C	CTGGATCAGATTCGGACAATCG	CACCGTCTCATTGAAGTGCCA