

SUPPLEMENTARY TABLE S3. REVERSE TRANSCRIPTION-POLYMERASE CHAIN REACTION AND QUANTITATIVE REVERSE TRANSCRIPTION-POLYMERASE CHAIN REACTION PRIMERS

<i>Targets</i>	<i>Primer sequence (5'-3')</i>	<i>AT (°C)</i>	<i>PS (bp)</i>
hGAPDH	F- GGAGCGAGATCCCCTCCAAAAT R- GGCTGTTGTCATACTTCTCATGG	60	197
hIL-4	F- CGGCAACTTTGTCCACGGA R- TCTGTTACGGTCAACTCGGTG	60	111
hIL-13	F- CCTCATGGCGCTTTTGTGAC R- TCTGGTTCTGGGTGATGTTGA	60	134
mArg-1	F- TGTCCCTAATGACAGCTCCTT R- GCATCCACCCAAATGACACAT	60	204
mCD163	F- GGTGGACACAGAATGGTTCTTC R- CCAGGAGCGTTAGTGACAGC	60	128
mCD206	F- CTCTGTTTCAGCTATTGGACGC R- TGGCACTCCCAAACATAATTTGA	60	190
mGAPDH	F- GTGGCAAAGTGGAGATTGTTG R- CTCCTGGAAGATGGTGATGG	60	164
mIL-1 α	F- AGTATCAGCAACGTCAAGCAA R- TCCAGATCATGGGTATGGACTG	60	105
mIL-1 β	F-TGTAATGAAAGACGGCACACC R- TCTTCTTTGGGTATTGCTTGG	60	111
mIL-6	F- GATGGATGCTACCAAAGTGGAT R- CCAGGTAGCTATGGTACTCCAGA	60	85
mIL-10	F- GCTCTTACTGACTGGCATGAG R- CGCAGCTCTAGGAGCATGTG	60	105
miNOS	F- ATGTCCGAAGCAAACATCAC R- TAATGTCCAGGAAGTAGGTG	60	450
mTNF- α	F- GGGACAGTGACCTGGACTGT R- CTCCCTTTGCAGAACTCAGG	60	185

AT, annealing temperature; F, forward; PS, product size; R, reverse.