

Table SI. Primers for reverse transcription-quantitative PCR.

Gene	Sequence (5'→3')
GAPDH	F: GGCTCTCCAGAACATCATC R: TCTTCCTCTTGTGCTCTTG
TET2	F: TGATGCTGATGATGCTGATA R: GTAGTGGTGGTGTCTCTTA
P16 ^{INK4A}	F: ATATGCCTTCCCCCACTACC R: CGTGAGTGCTCACTCCAGAA
LCN2	F: CCACCTCAGACCTGATCCCA R: CCCCTGGAATTGGTTGTCTTG
S100A7	F: ACGTGATGACAAGATTGACAAGC R: GCGAGGTAATTTGTGCCCTTT
MMP9	F: AGACCTGGGCAGATTCCAAAC R: CGGCAAGTCTTCCGAGTAGT
CXCL1	F: GAAAGCTTGCCTCAATCCTG R: AACAGCCACCAGTGAGCTTC
IRF7	F: GCTGGACGTGACCATCATGTA R: GGGCCGTATAGGAACGTGC
IL-7R	F: CCCTCGTGGAGGTAAAGTGC R: CCTTCCCGATAGACGACACTC

F, forward; R, reverse; TET2, ten-eleven translocation-2; P16^{INK4A}, cyclin-dependent kinase inhibitor 2A; LCN2, lipocalin 2; S100A7, S100 calcium binding protein A7; MMP9, matrix metalloproteinase; CXCL1, C-X-C motif chemokine ligand 1; IRF7, interferon regulatory factor 7; IL-7R, interleukin-7 receptor.