Supplement 4: Transcript levels of four selected genes by qPCR in the three study groups

STAT1

SLE>Controls: \( p < 0.001 \), SLE>SSc: \( p = 0.009 \), SSc>Controls: \( p < 0.001 \)

IFIT3

SLE>Controls: \( p < 0.001 \), SLE>SSc: \( p = 0.008 \), SSc>Controls: \( p < 0.001 \)

TLR5

SLE<->Controls: \( p = 0.426 \), SLE<->SSc: \( p = 0.811 \), SSc<->Controls: \( p = 0.384 \)