



Supplemental Figure S1. qRT-PCR analysis of IFN β +TNF (I+T) upregulated DEGs positively regulated by STAT2 and IRF9. U3A cells were transfected with siCTRL (C), siSTAT2 (S2) or siIRF9 (F9). Cells were further left untreated or stimulated with I+T for 24h. Quantification of the mRNA corresponding to the indicated genes was performed by qRT-PCR and expressed as relative expression (DDCt) after normalization to the S9 mRNA levels. Mean \pm SEM, $n \geq 5$. Statistical comparison was conducted using Two-way ANOVA with tukey post-test. $p < 0.0001$ (****); ns: not significant.