

Expanded View Figures

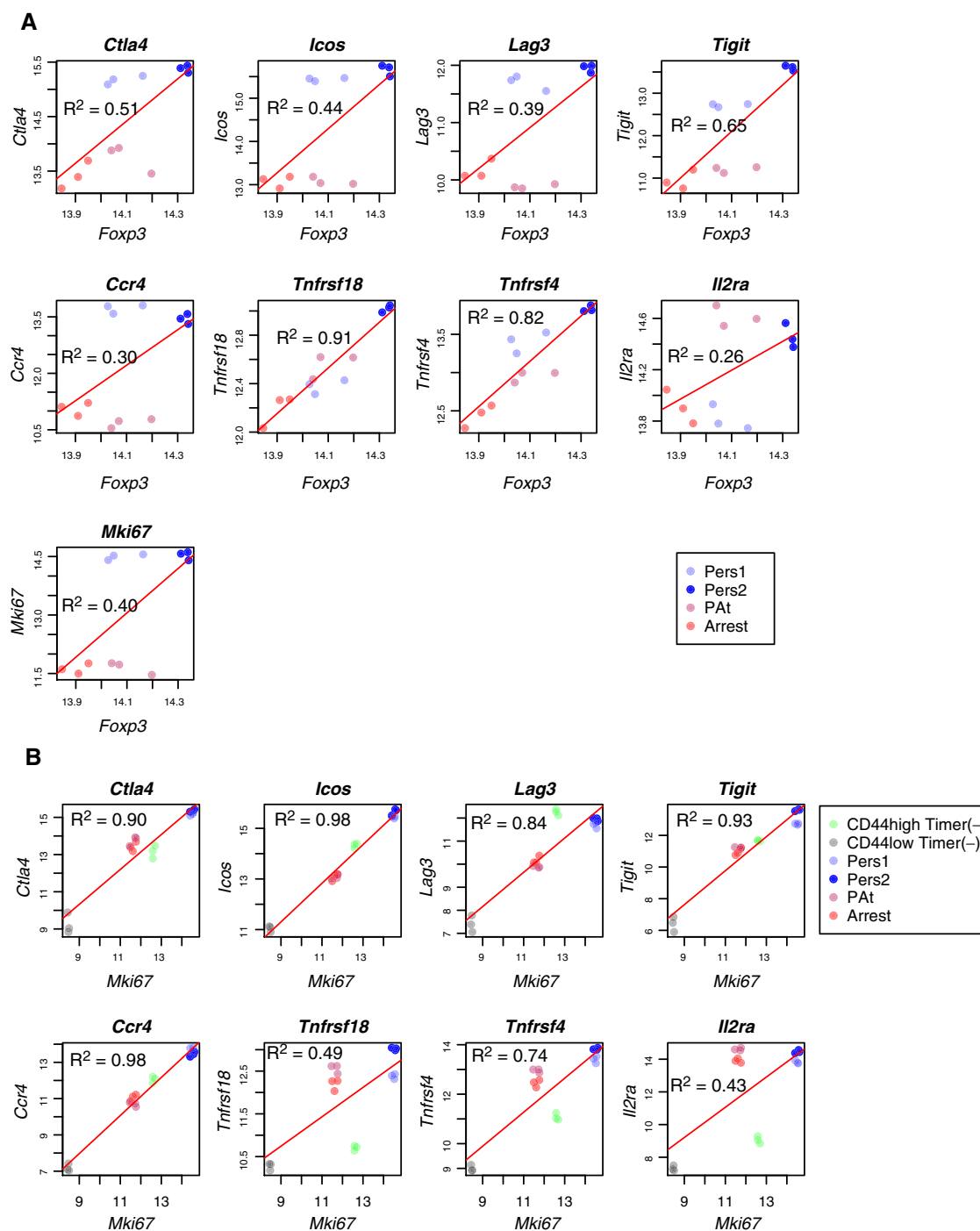


Figure EV1. Analysis of correlations between *Foxp3* and *Mki67* in RNA-seq data from Fig 5.

A, B Scatter plots showing the correlations between the indicated transcripts (normalised log2 of read counts) and *Foxp3* transcripts and *Mki67* transcripts. Note that *Foxp3*⁺ cells only are shown in (A). Red lines indicate linear regression line, and R-square values are shown. Group designations are shown by colours (see legend), *n* = 3.

