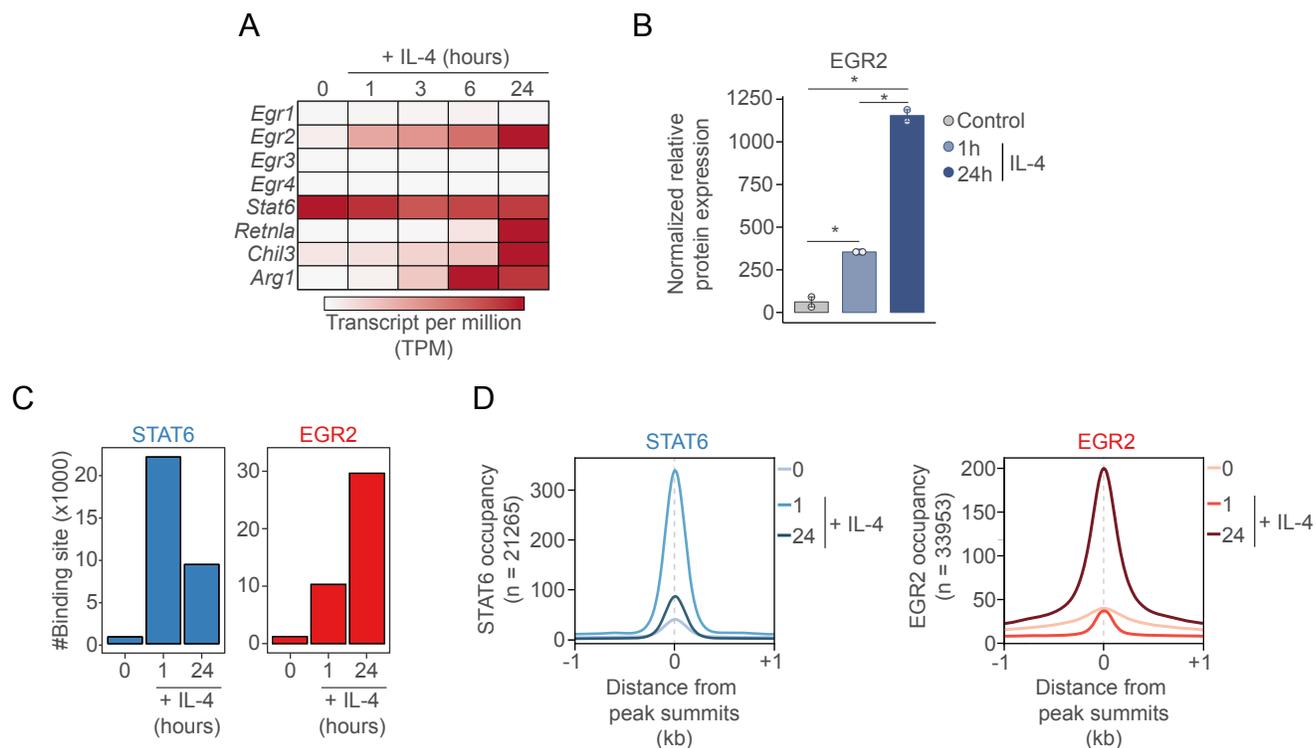


Supplemental Fig. S2



Supplemental Fig. S2. *Egr2* is selectively induced by IL-4 and populates the macrophage genome. Related to Fig. 2.

A, Heatmap representation of the expression of *Egr* transcription factor family member genes and alternative polarization marker genes over the indicated time course determined by RNA-seq.

B, Quantification of western blot results presented on Fig. 2D by densitometry. Experiments were repeated four times, and significant changes between groups were calculated by two-way analysis of variance (ANOVA).

C, Bar plot visualization of the overall number of binding sites for STAT6 and EGR2 from ChIP-seq experiments at the indicated time points.

D, Histograms of average STAT6 and EGR2 binding signals at two different time points following IL-4 polarization.