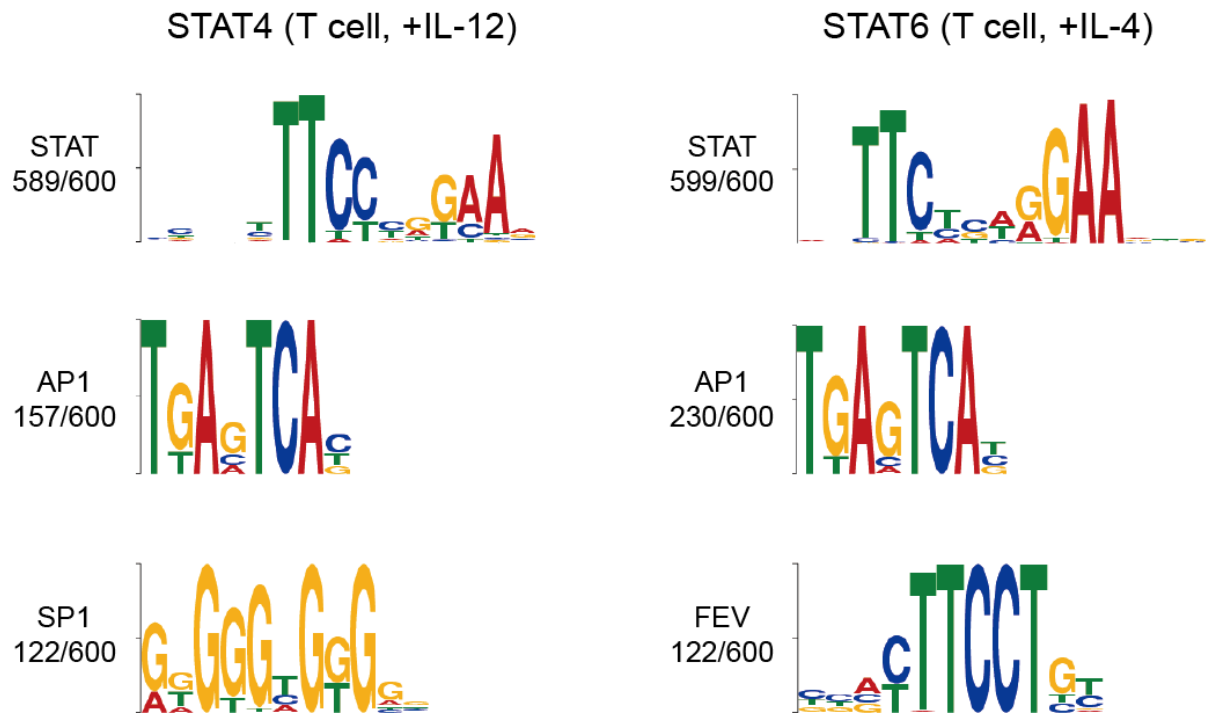




Additional file 5. Predicted motifs in top 600 STAT4 and STAT6 binding sites.
The same motif analysis (Fig. 4A) was performed with top 600 (ranked by peak height) STAT4 and STAT6 binding sites. The number of sites containing the predicted motifs is shown.