

Figure S2

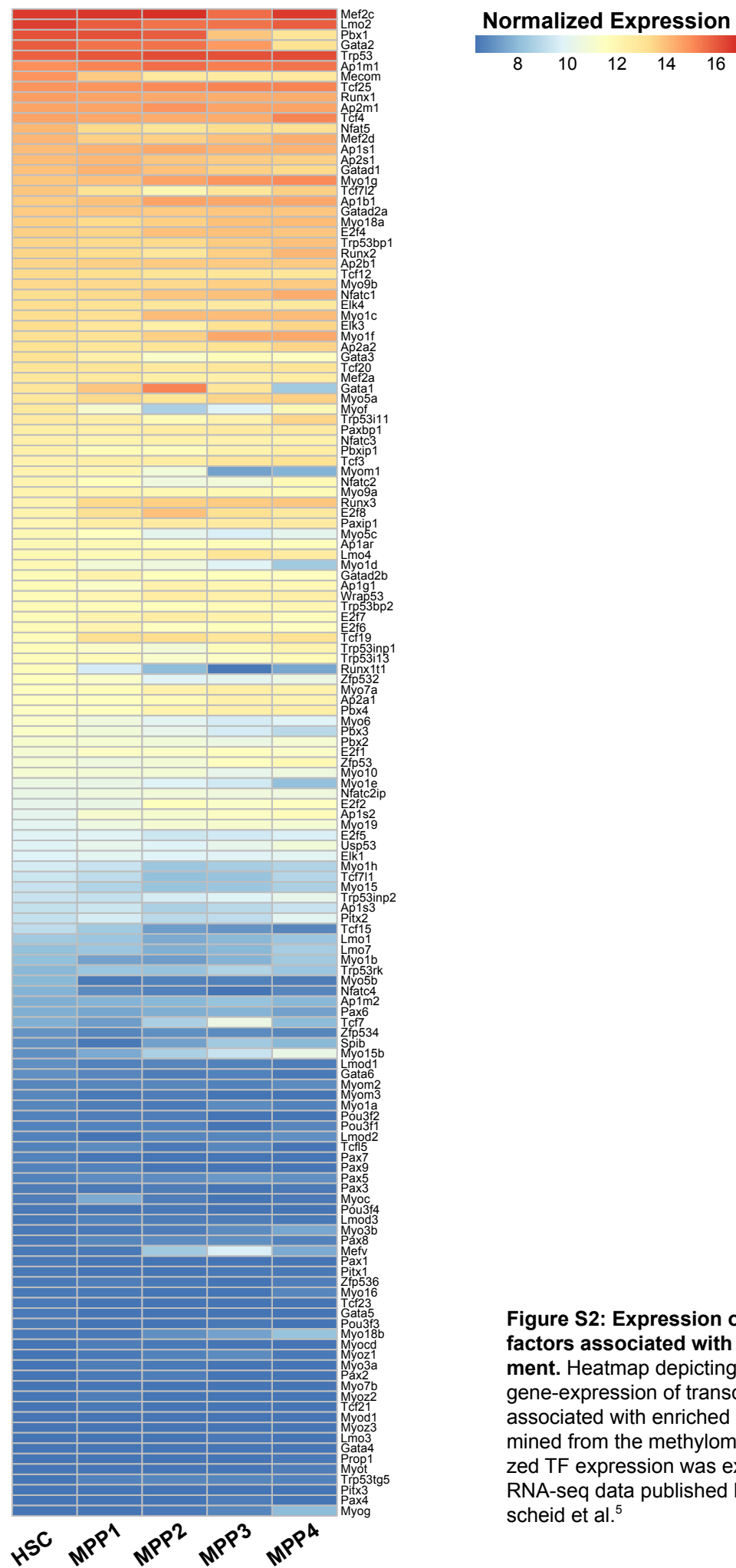


Figure S2: Expression of transcription factors associated with motif enrichment. Heatmap depicting normalized gene-expression of transcription factors associated with enriched motifs as determined from the methylome data. Normalized TF expression was extracted from the RNA-seq data published by Cabezas-Wallscheid et al.⁵