

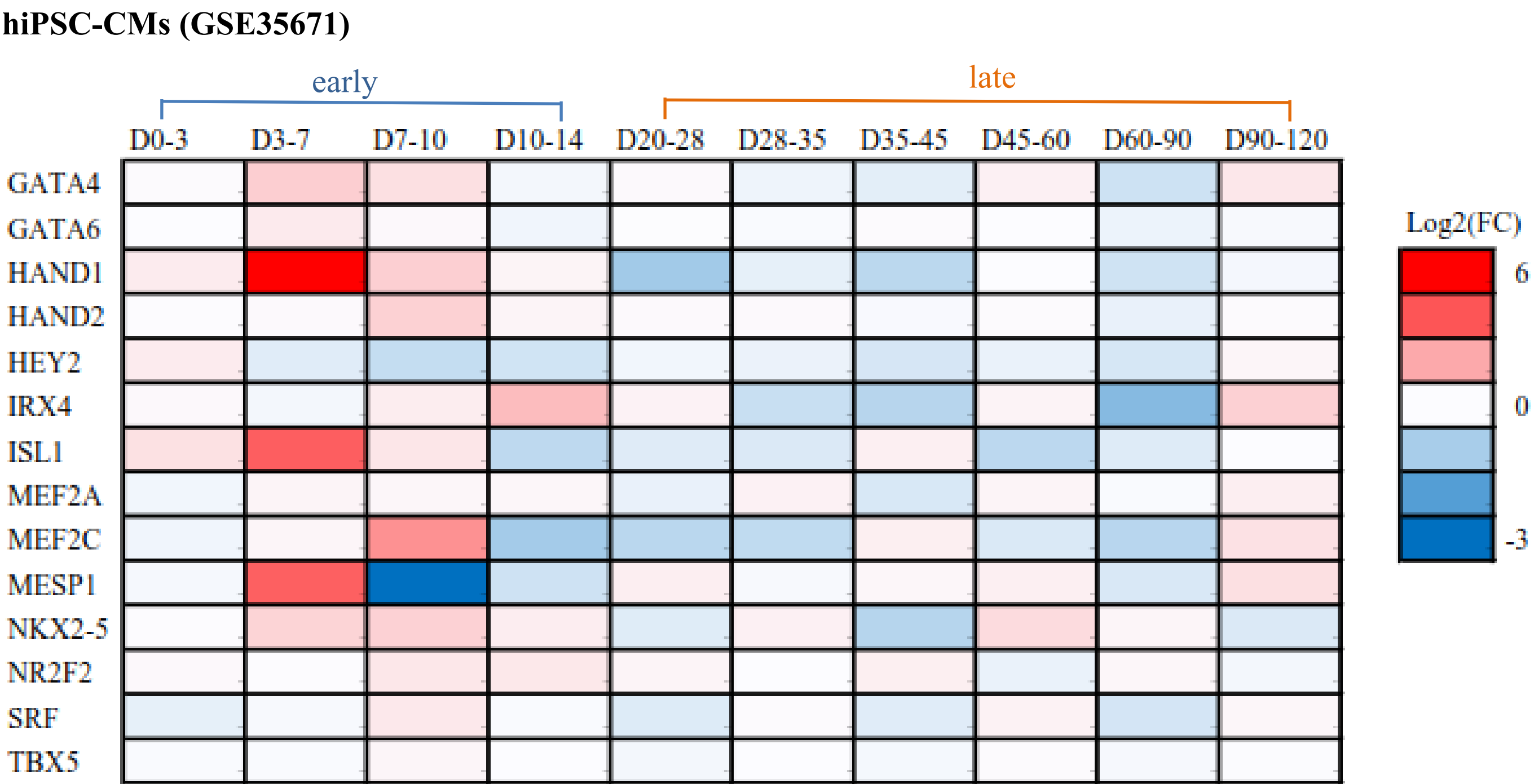
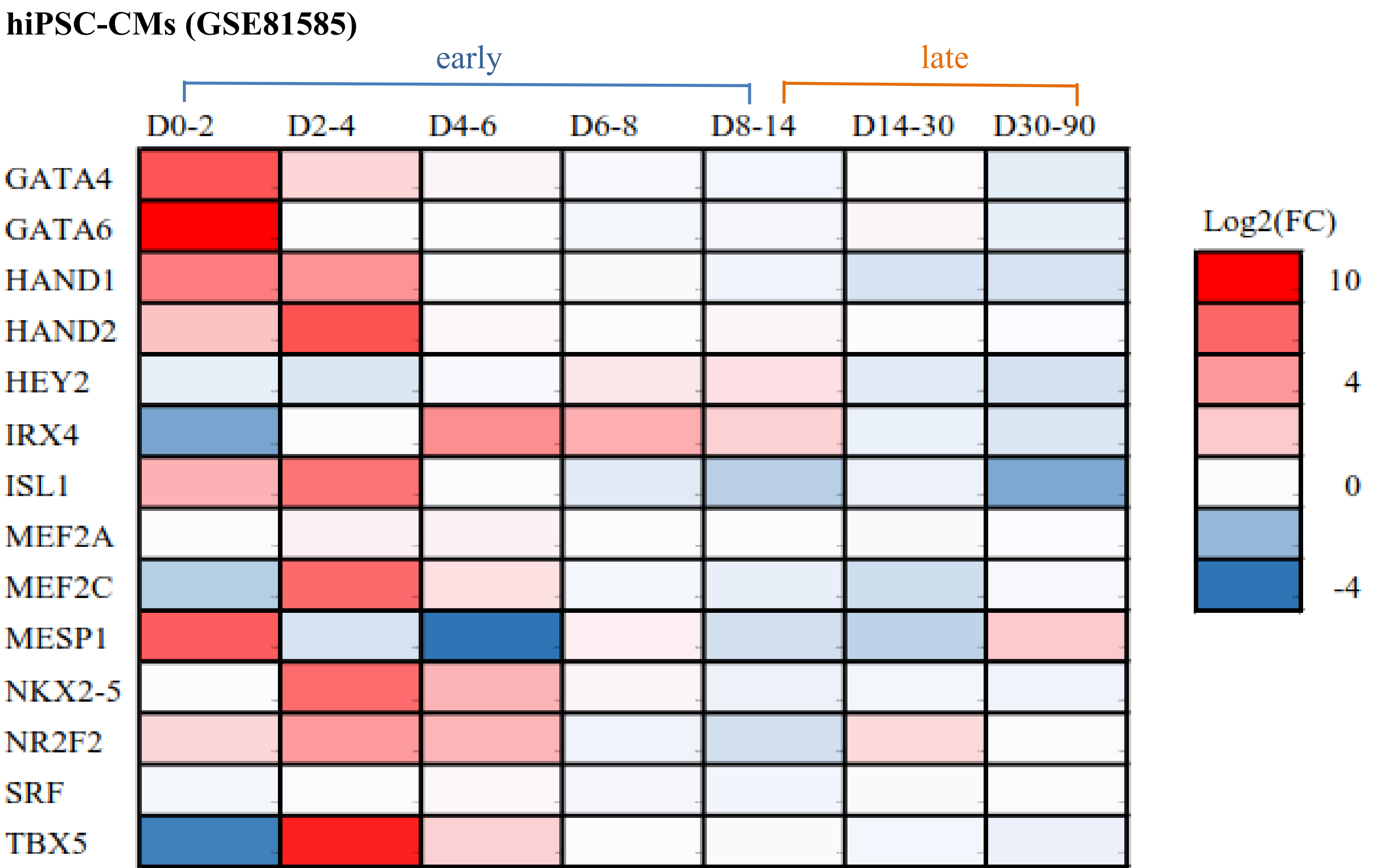
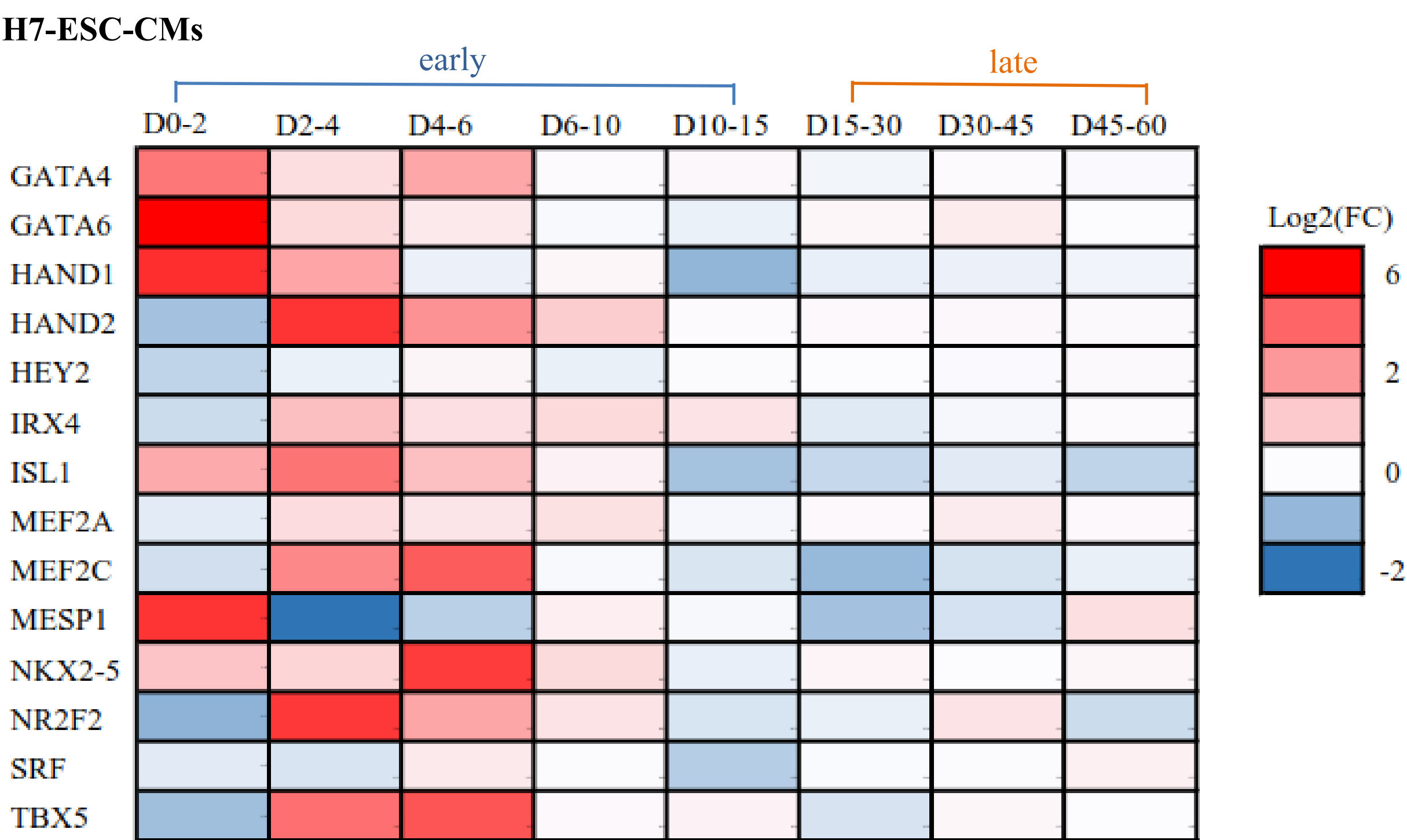


Figure S1. Heatmap of differential expression changes of the cardiac TFs during early and late stages of hESC-CM and hiPSC-CM differentiation.

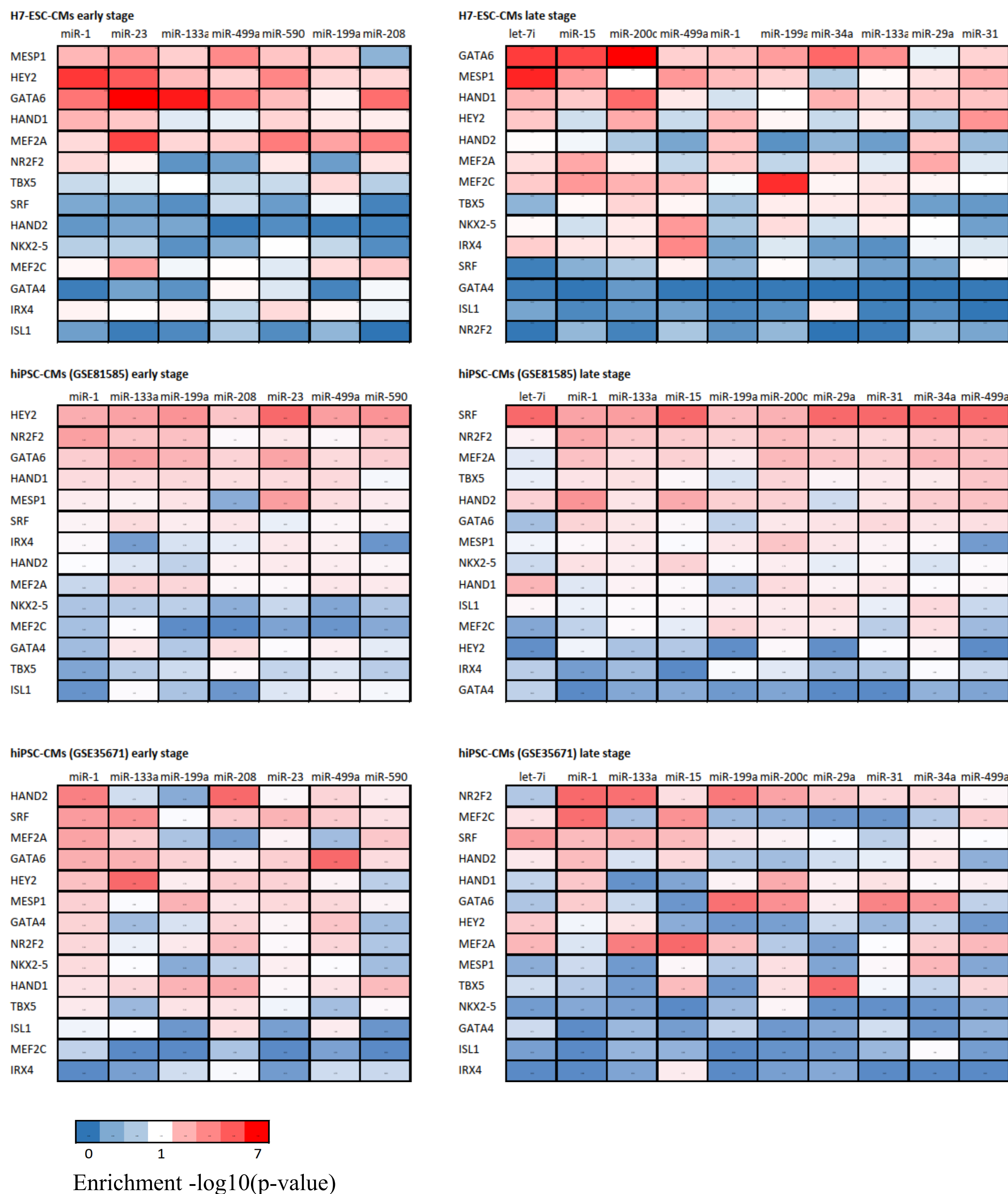


Figure S2. Enrichment analysis of overlaps between the target genes of the cardiac TFs and of miRNAs during early and late stages of hESC-CM/hiPSC-CM differentiation. The p-values account for the significance of the enrichment analysis based on hypergeometric distribution.

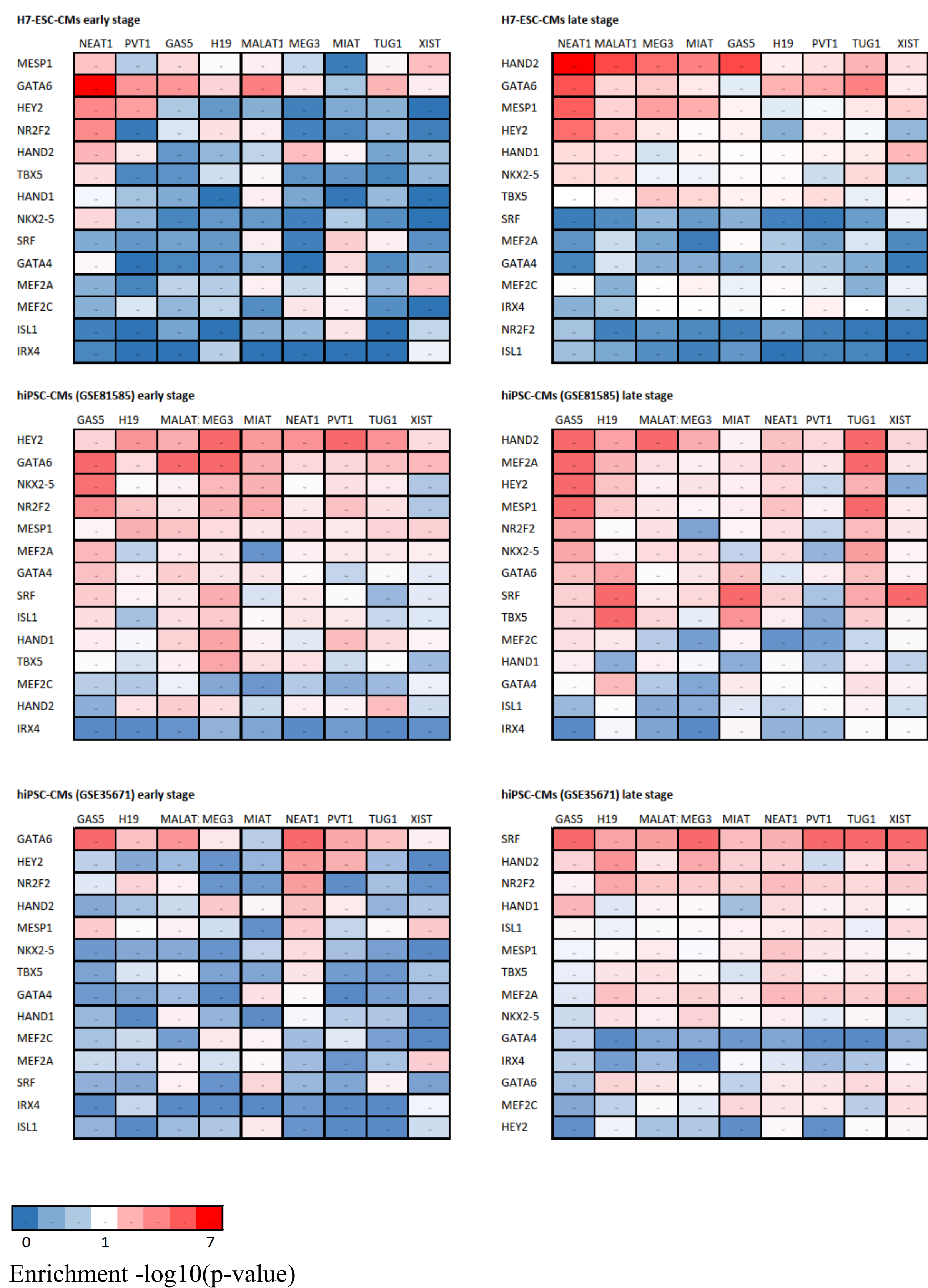


Figure S3. Enrichment analysis of overlaps between the target genes of the cardiac TFs and of lncRNAs during early and late stages of hESC-CM/hiPSC-CM differentiation. The p-values account for the significance of the enrichment analysis based on hypergeometric distribution.