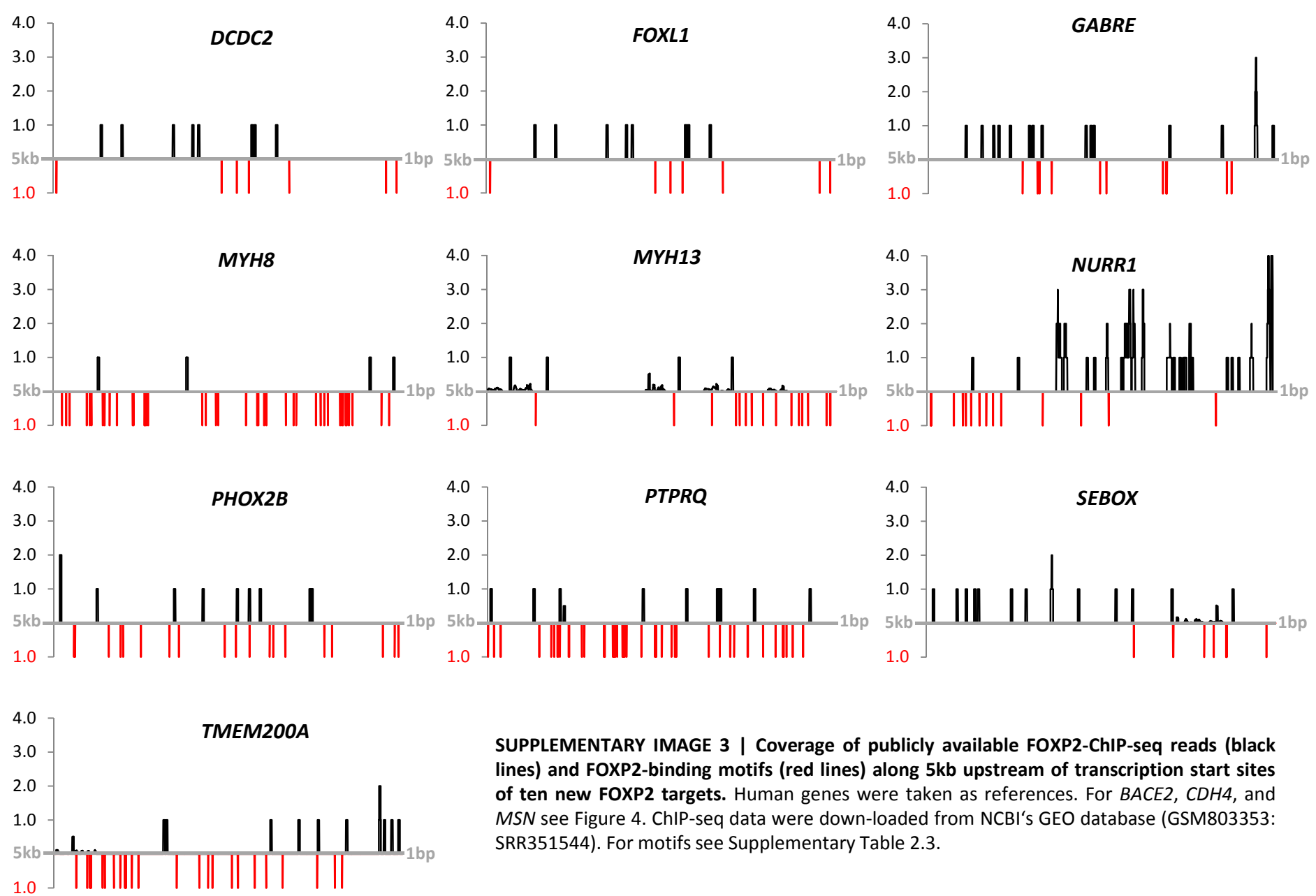


**SUPPLEMENTARY IMAGE 3**



**SUPPLEMENTARY IMAGE 3 | Coverage of publicly available FOXP2-ChIP-seq reads (black lines) and FOXP2-binding motifs (red lines) along 5kb upstream of transcription start sites of ten new FOXP2 targets.** Human genes were taken as references. For *BACE2*, *CDH4*, and *MSN* see Figure 4. ChIP-seq data were down-loaded from NCBI's GEO database (GSM803353: SRR351544). For motifs see Supplementary Table 2.3.