



Additional file 12: Figure S10: ChIP-seq and DNase-seq data from mES cells can also be used as an input. (A) Browser view of the α globin locus. (B-D) ChIP-seq and DNase-seq data for mouse ES cells across the same region. Red lines show the pile-up of reads, and black points indicate the positions of binding sites identified by peak-calling. Data from the ENCODE project (58). (E) Browser view of the β globin locus. (F-H) ChIP-seq and DNase-seq data for mouse ES cells across the same region. Data from the ENCODE project (58).