Supplemental Fig. S9 – Peak calling consistency across laboratories by gcapc and SPP on three transcription factors. (A) CAT plot shows the improvements of consistency by removing GC effects for *YY1*. (B) GC content of called peaks by our approach and SPP for *YY1*. (C) Motif enrichment in called peaks for *YY1*. (D) as (A) but for transcription factor *EP300*. (E) as (B) but for transcription factor *EP300*. (F) as (A) but for transcription factor *GATA2*. (G) as (B) but for transcription factor *GATA2*. (H) as (C) but for transcription factor *GATA2*. The motif information was downloaded from JASPAR database 2016. Motif analysis for *EP300* was omitted as we didn't find good source of motif information in public databases.