Figure S2. Sequence logos for all de novo discovered motifs and their corresponding TFs. There are 79 motifs in total, and they are ordered alphabetically by motif name. UA1–UA12 are 12 motifs that were not annotated in TRANSFAC or JASPAR. UA12 was identified in the subset of HDAC2-associated GATA1 peaks in K562. The remaining 78 motifs were identified in the top 500 ChIP-seq peaks using our motif discovery pipeline.