Figure S7. Re-classification of ChIP-chip target intergenic regions (IGRs) according to both conservative and permissive E score thresholds. Light blue bars indicate the number of ChIP-chip "indirect" target IGRs that would be reclassified as "direct" targets according to the presence of a TF specific k-mer with an E Score > 0.45 from PBM data. Dark blue bars show this same reannotation at a more stringent minimum E Score (0.49). Light red bars indicate the number of ChIP-chip "direct" IGR targets that would be reclassified as "indirect" targets because they have no k-mers with an E Score > 0.45 from PBM data. Dark red bars impose a more stringent requirement (no kmers E score > 0.35) for reannotating an IGR from "direct" to "indirect."