

Additional data file 11. Temporal changes in levels of MED occupancy correlate with changes in DNaseI accessibility. 1 kb regions +/- 500 bp of the peak nucleotide of binding for each of the 400 regions most highly bound by MED at stages 5 (A and B), 10 (C) and 14 (D) were identified. (A) Plot of the ratio of ChIP-chip scores at stage 5 over those at 10 (x axis) versus the ratio of DNaseI-seq scores at stage 5 over those at 10 (y axis). (B) Plot of the ratio of ChIP-chip scores at stage 5 over those at 10 (x axis) versus the ratio of DNase-seq scores at stage 5 over those at 10 (y axis). (C) Plot of the ratio of ChIP-chip scores at stage 10 over those at 5 (x axis) versus the ratio of DNase-seq scores at stage 10 over those at 5 (y axis). (D) Plot of the ratio of ChIP-chip scores at stage 14 over those at 5 (x axis) versus the ratio of DNase-seq scores at stage 14 over those at 5 (y axis). The Pearson correlation coefficients (r) for each comparison are shown in the top right of each panel.

