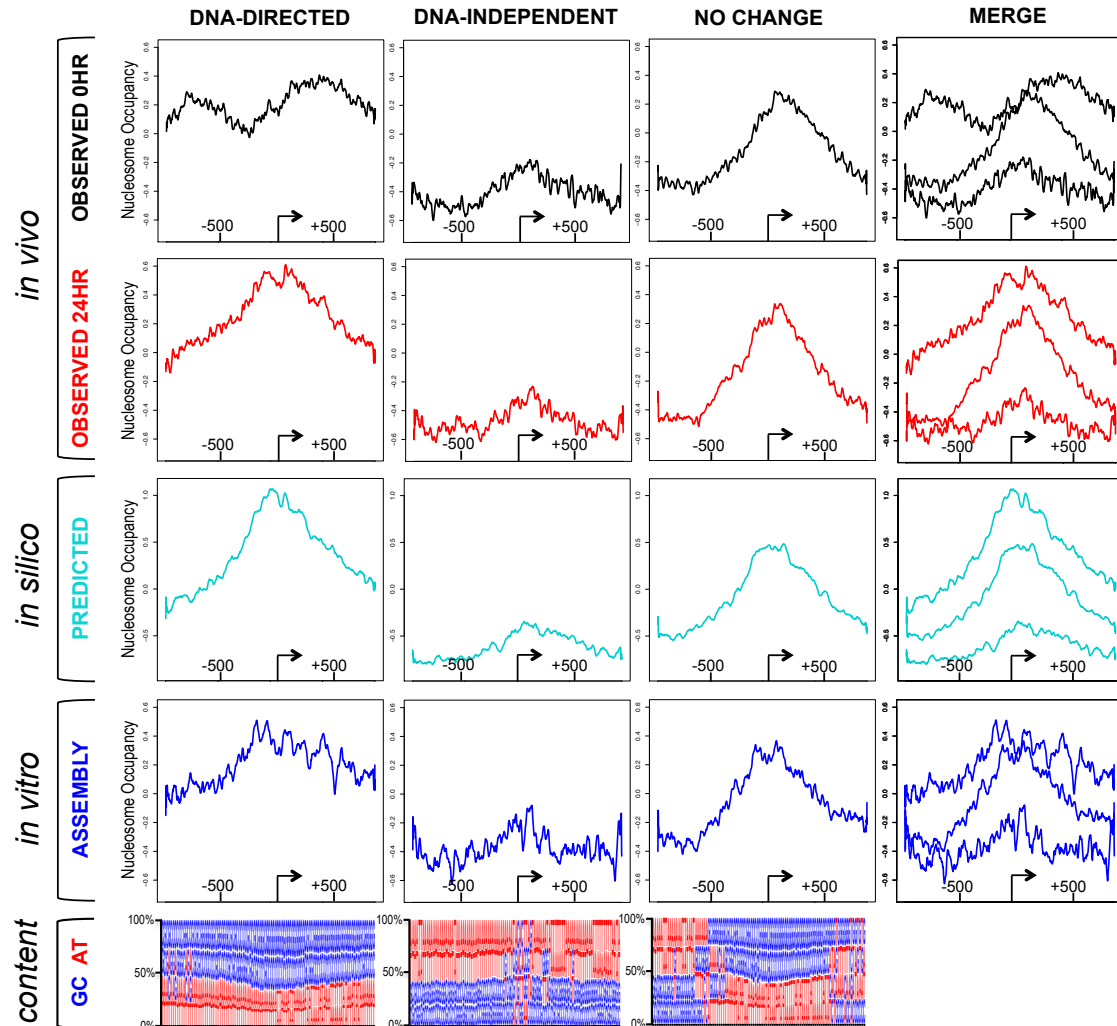


Sexton et al. _ Supplementary Figure 17



Supp. Fig. 17. Average values for all genes identified as DNA directed (left column), DNA independent (second column), and no change (third column), calculated by alignment of loci to the TSS for each of: *in vivo* observed 0 hours (black), *in vivo* observed 24 hours (red), *in silico* A375 prediction (10) (cyan), and *in vitro* assembly (blue). The last column is the merge of DNA directed, DNA independent, and no change for each row. Average GC content at each position is represented as a sequence logo below each column; GC, blue; AT, red.