



Figure S3 Cap-H2 ChIP-seq peak regions validated by ChIP-qPCR. (A-D) Integrated Genome Viewer (IGV) screenshot of ChIP-seq peaks for Cap-H2 and Mrg15 in regions near *Esc* (A), *squ* (B), *Orc2* (C), and *RpL27* (D) validated by ChIP-qPCR in Figure 2B. Grey arrows indicate the region that was amplified by qPCR.