The detailed results of ChIP-Seq analyses showing the enrichment of H3K27me3 in dE2f1 (A, corresponding to the peak in Fig. S1B), dCycE (B, corresponding to the peak in Fig. S2B) and stg (C, corresponding to the peak in Suppl. Fig. 3B) loci during the WPP stage. These results are obtained from the ChIP-Seq data sets provided by Dr. Kevin White laboratory from the modENCODE Project (http://modencode.oicr.on.ca/fgb2/gbrowse/fly/).