



Supplemental Figure S13. Mutations in CPA machinery result in increased H3K9me enrichment at constitutive heterochromatic loci. Replicate data for H3K9me ChIP-seq for *pfs2-11* (Figure 5B-D). Telomeres from chromosomes I and II and centromeres from chromosomes I, II and II depicted. Parent (blue), *pfs2-11* mutant (orange).