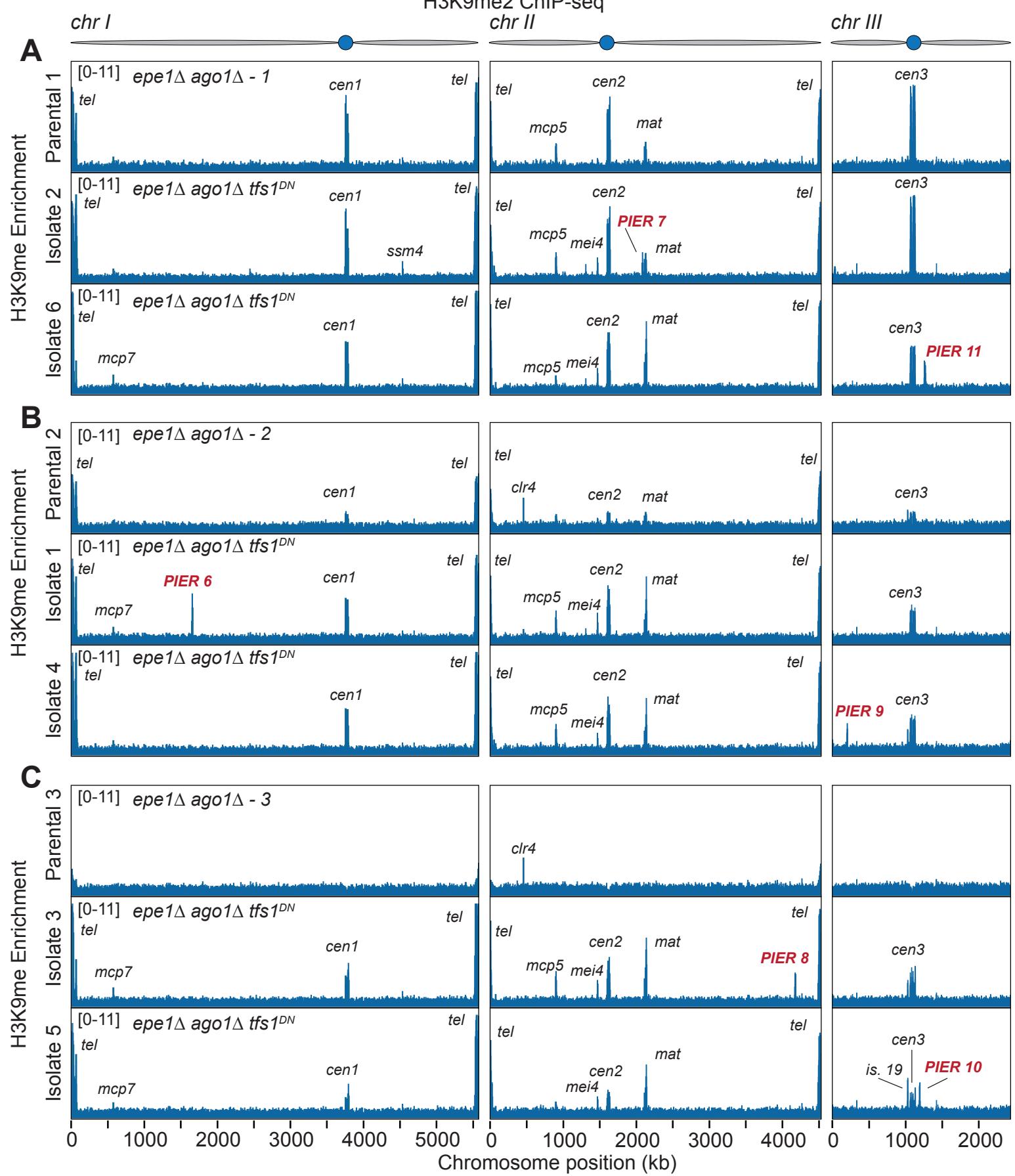


Supplemental Figure S11.

Parsa\_Supplemental\_FigS11



**Supplemental Figure S11. PIER formation is RNAi-independent and expression of TFIIS<sup>DN</sup> can rescue the adaptive silencing of *clr4*<sup>+</sup> observed in *epe1*<sup>Δ</sup> *ago1*<sup>Δ</sup> strains. A-C)** Genome wide plots of H3K9me2 enrichment. **A)** Parental (*epe1*<sup>Δ</sup> *ago1*<sup>Δ-1</sup>) and two independent isolates of *epe1*<sup>Δ</sup> *ago1*<sup>Δ</sup> *tfis1*<sup>DN</sup> derived from this parental are plotted depicting PIERs 7 and 11. **B)** Parental (*epe1*<sup>Δ</sup> *ago1*<sup>Δ-2</sup>), initially harboring H3K9me2 at *clr4*<sup>+</sup>, and two independent isolates of *epe1*<sup>Δ</sup> *ago1*<sup>Δ</sup> *tfis1*<sup>DN</sup> derived from this parental are plotted depicting PIERs 6 and 9. **C)** Parental (*epe1*<sup>Δ</sup> *ago1*<sup>Δ-3</sup>), initially harboring H3K9me2 at *clr4*<sup>+</sup>, and two independent isolates of *epe1*<sup>Δ</sup> *ago1*<sup>Δ</sup> *tfis1*<sup>DN</sup> derived from this parental are plotted depicting PIERs 8 and 10.