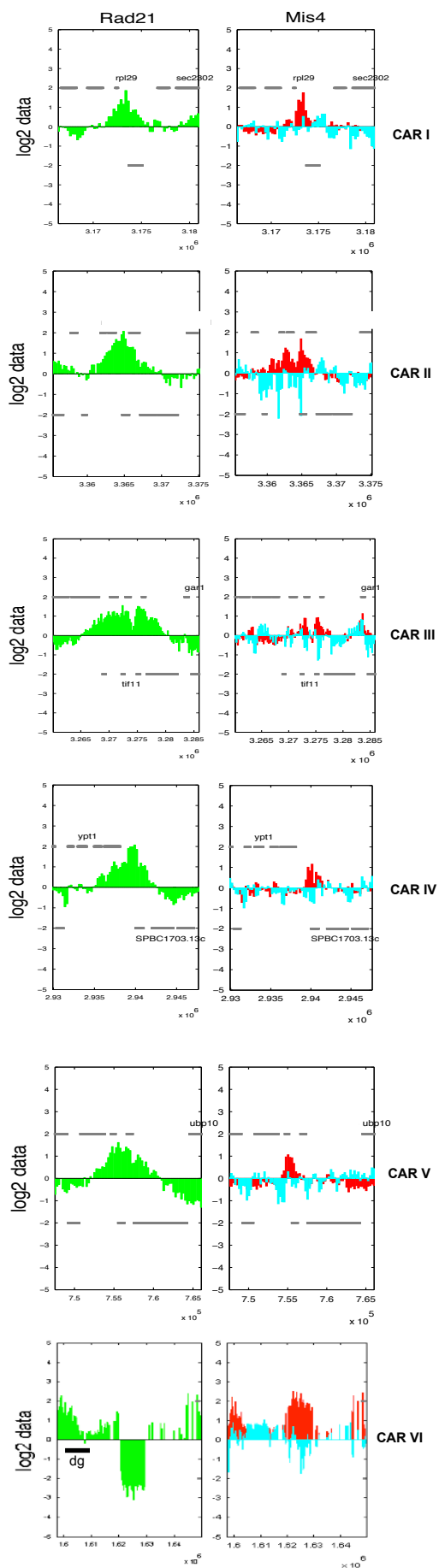
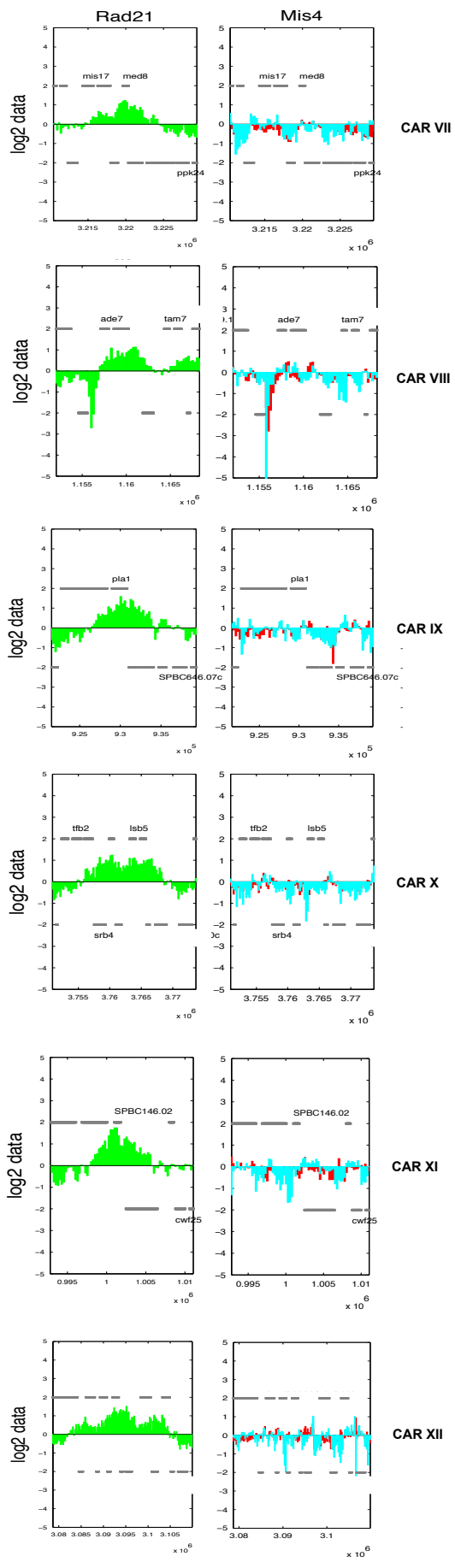


A

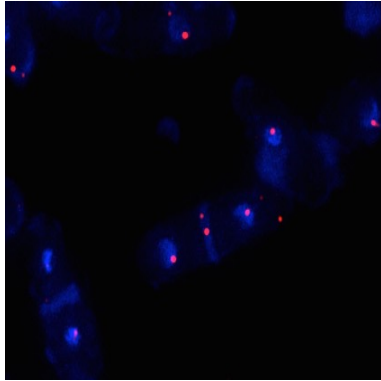
Mis4<sup>+</sup>/Rad21<sup>+</sup>

B

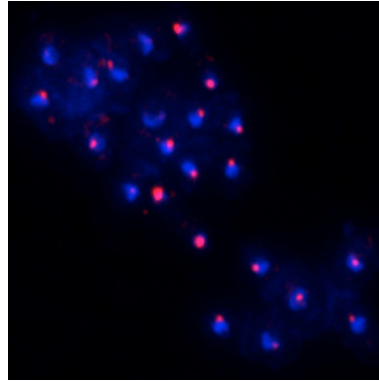
Rad21<sup>+</sup>

Mis4+/Rad21+

A



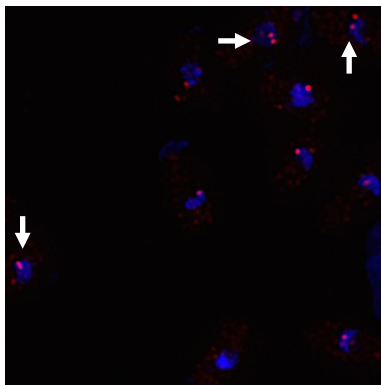
CAR IV



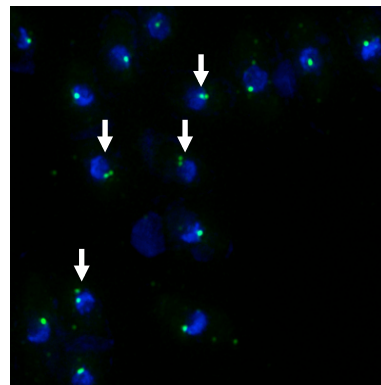
CAR VI

Rad21+

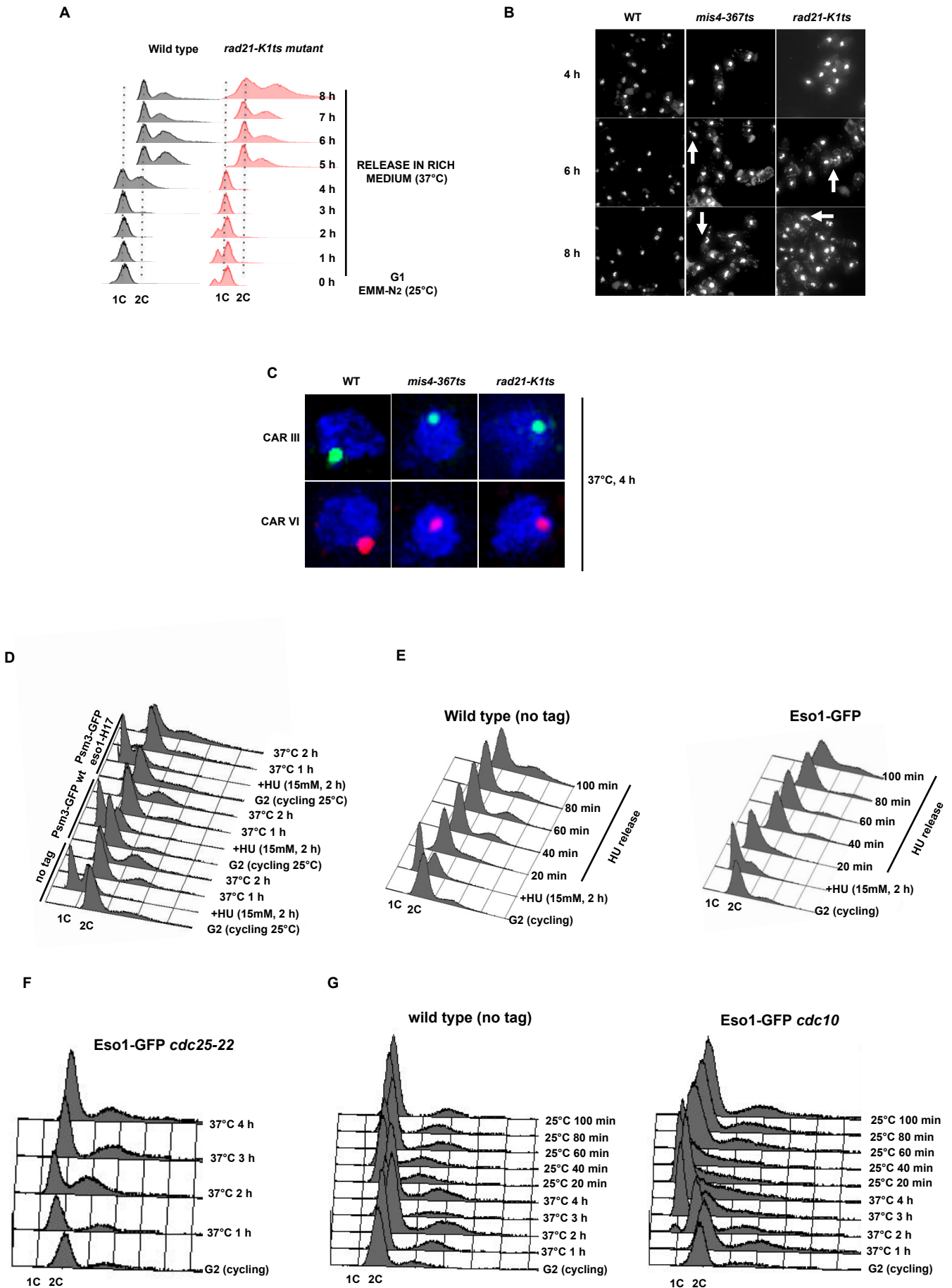
B



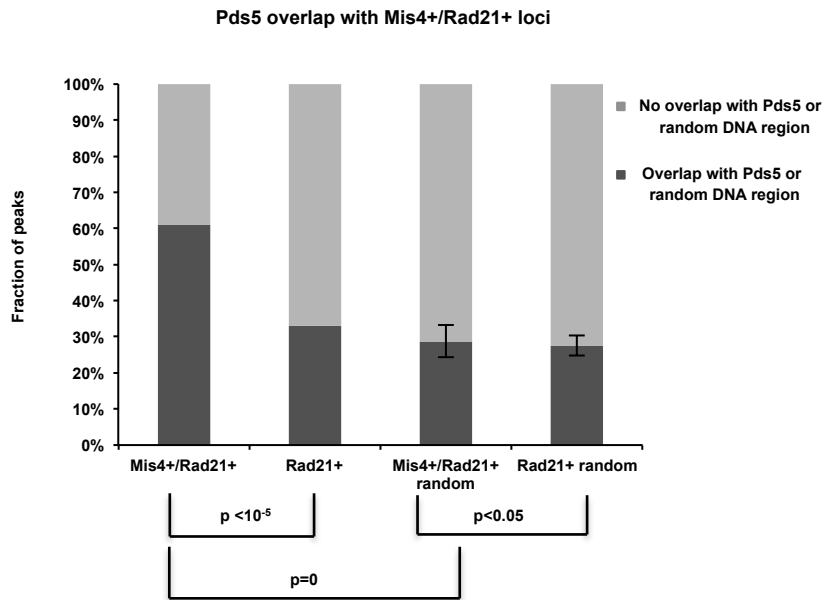
CAR VII



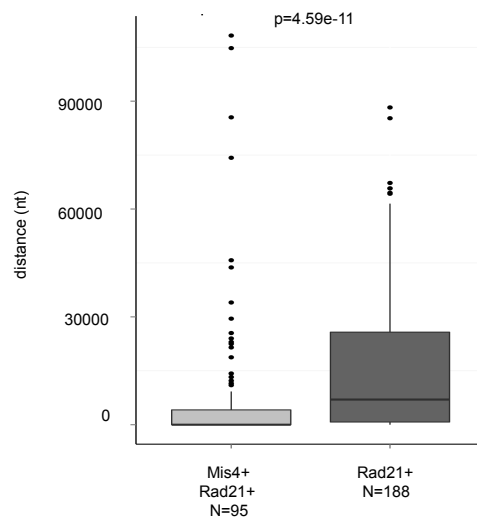
CAR VIII



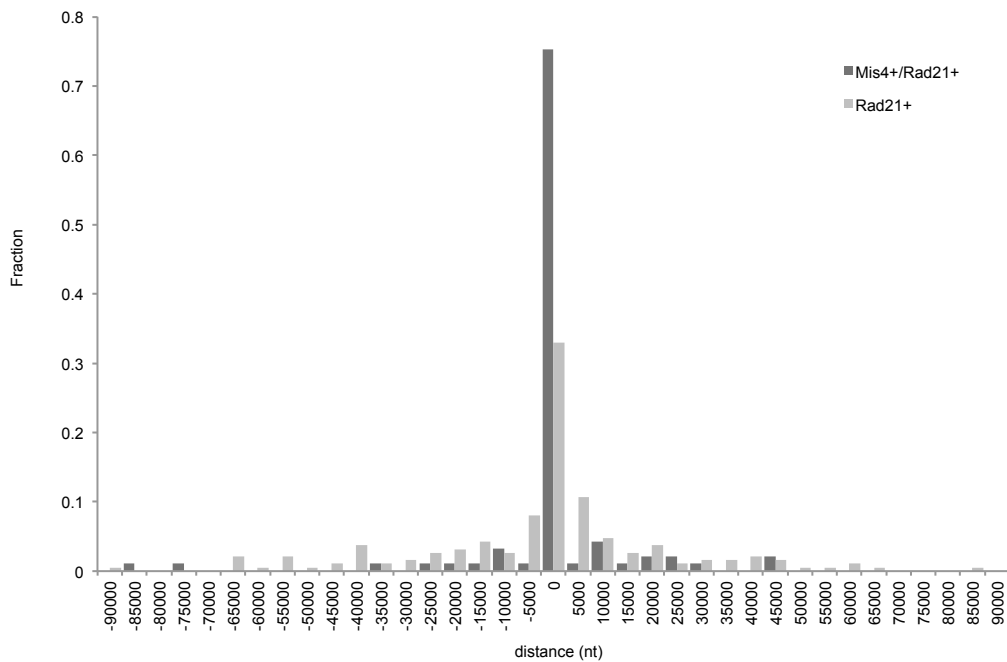
A

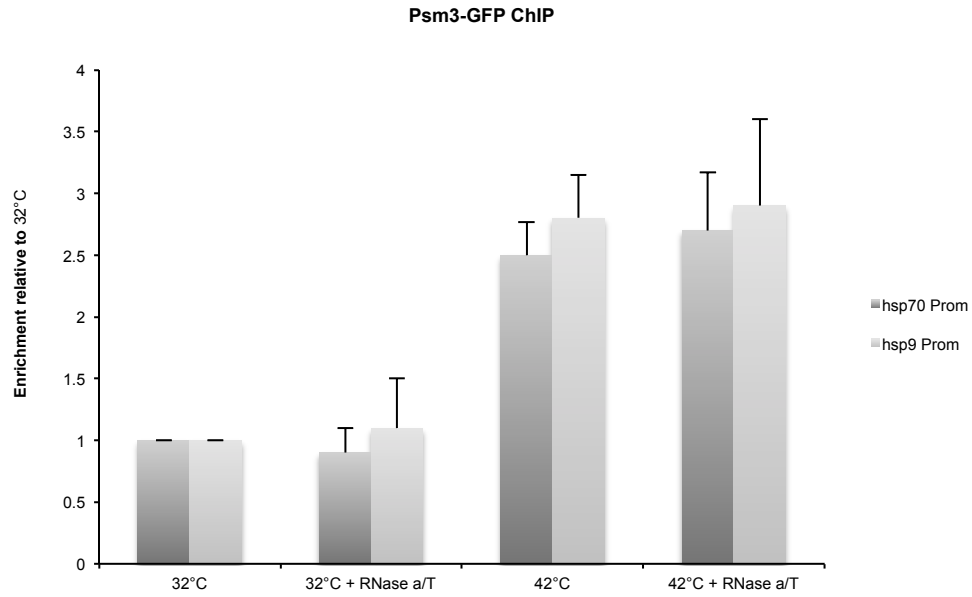
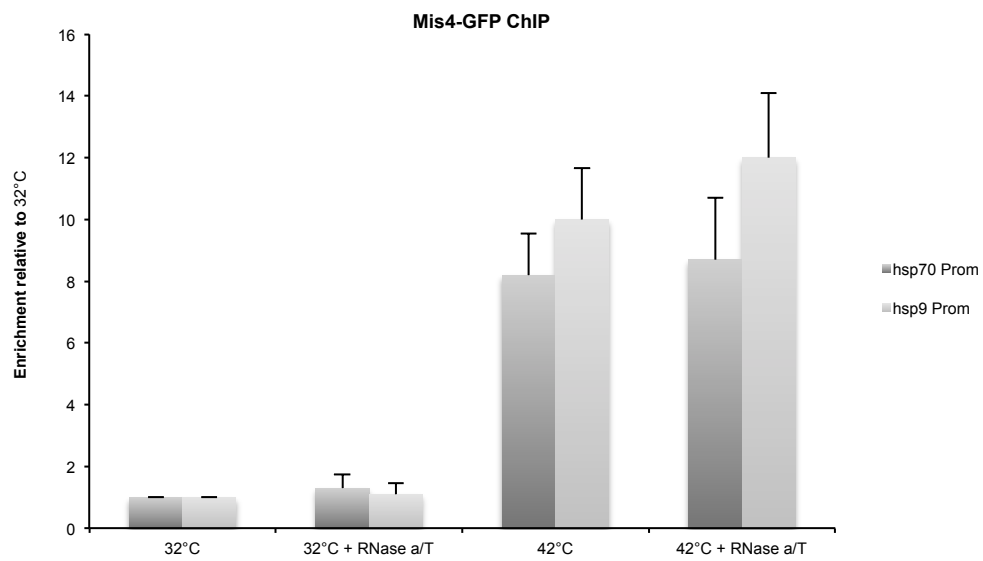


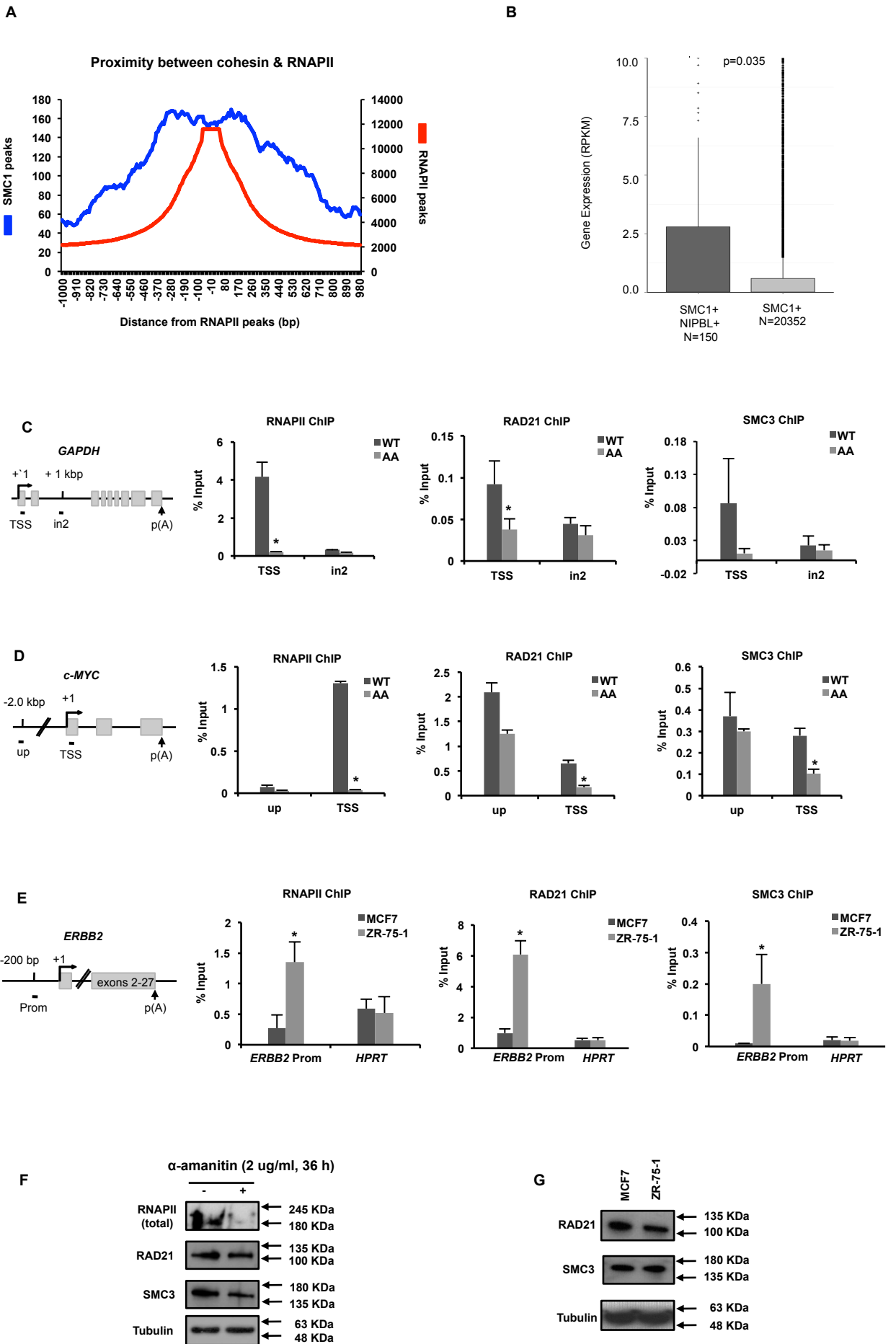
B



C

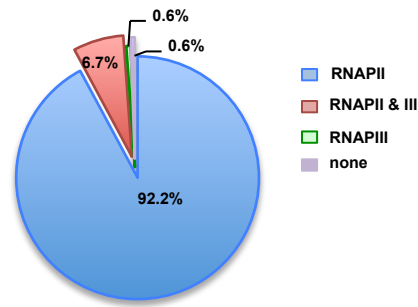


**A****B**

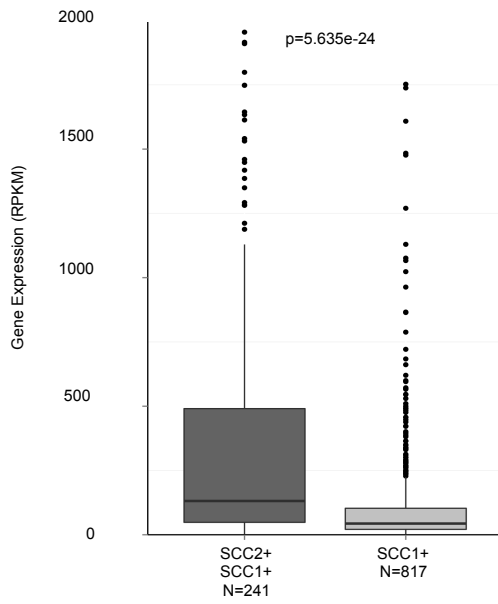


A

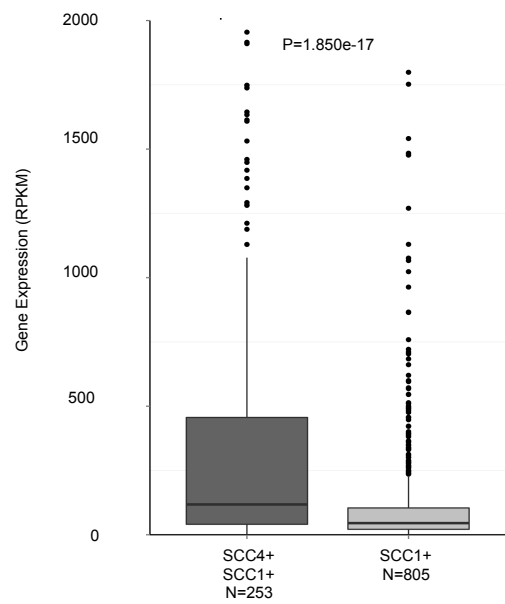
Cohesin (1058 peaks) distribution in *S. cerevisiae*, relative to RNAPII & III



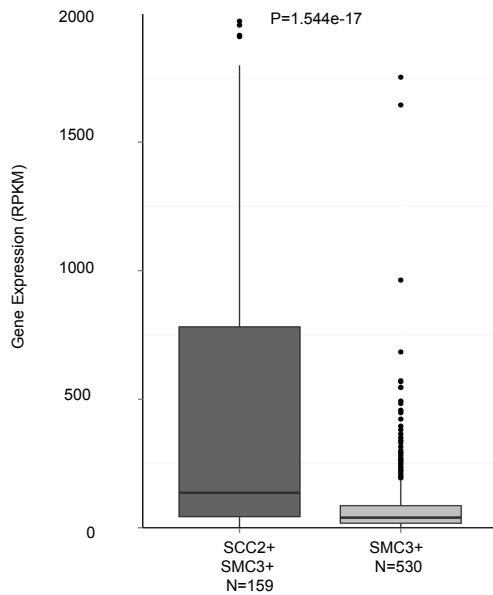
B



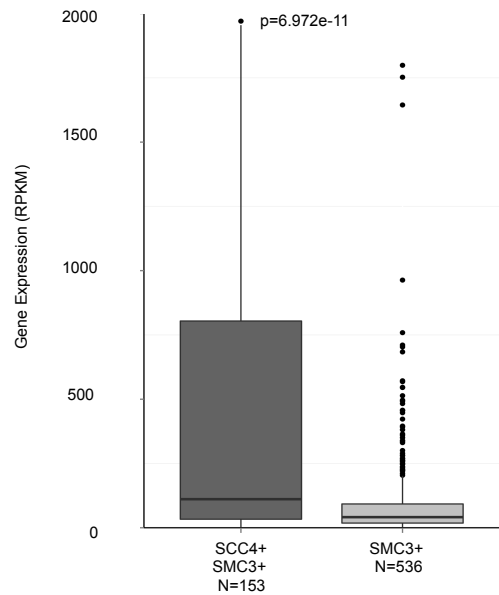
C



D

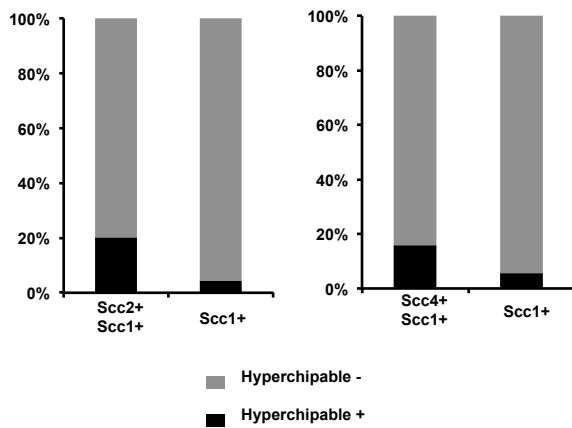


E



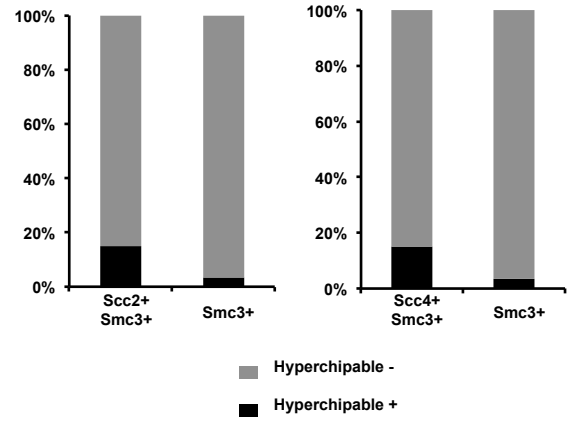
A

“Hyperchipable regions” plotted relative to *Scs2+*/*Scs1+* & *Scs1+* or *Scs4+*/*Scs1+* & *Scs1+* sites in *S. cerevisiae*

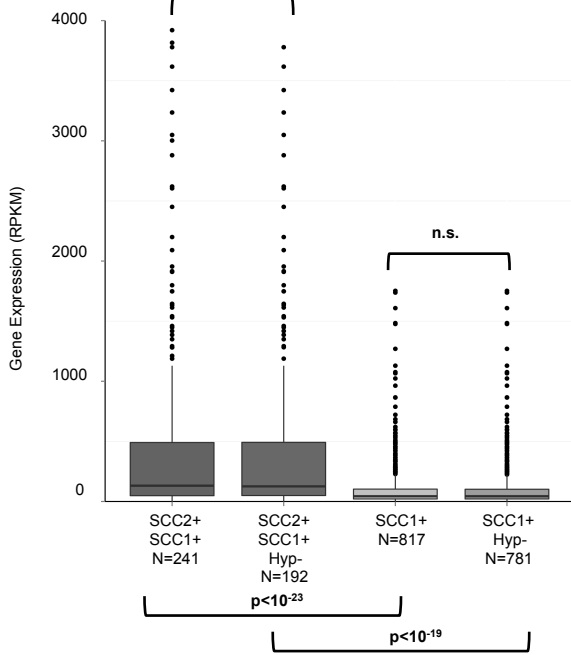


B

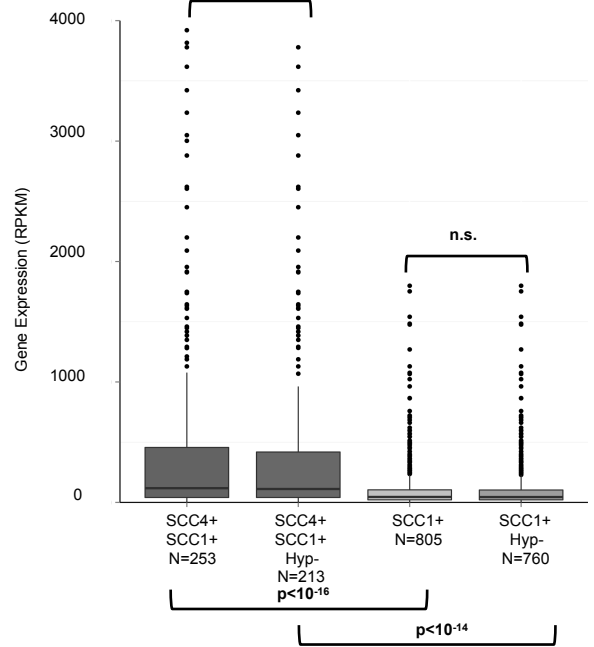
“Hyperchipable regions” plotted relative to *Scs2+*/*Scs3+* & *Scs3+* or *Scs4+*/*Scs3+* & *Scs3+* sites in *S. cerevisiae*



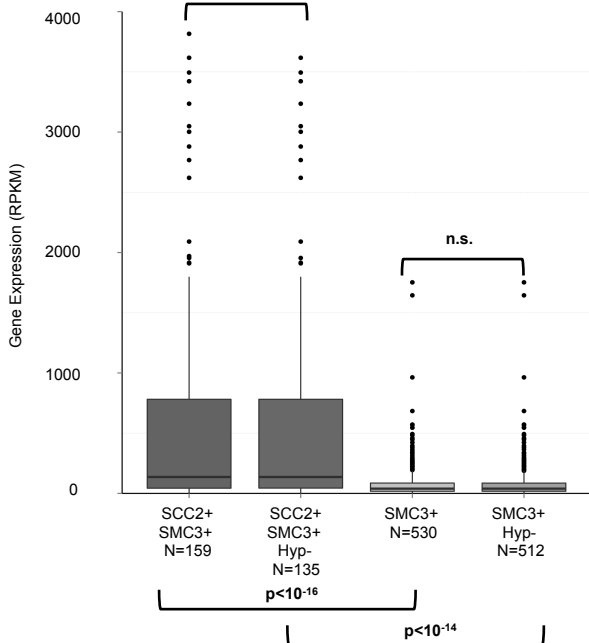
C



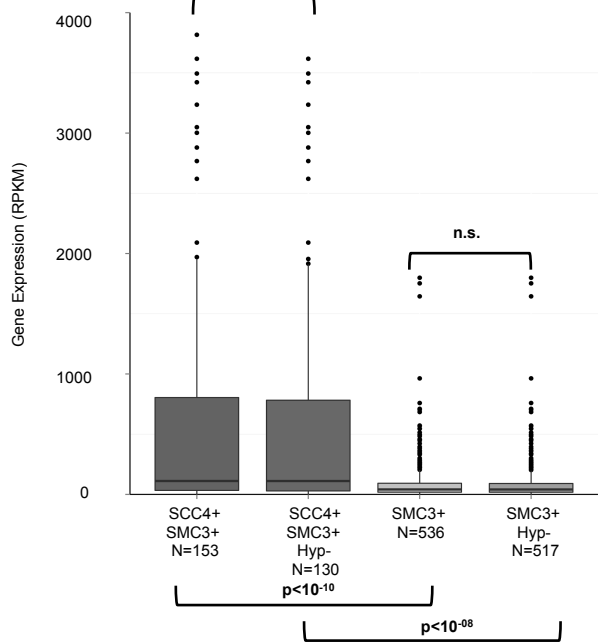
D



E



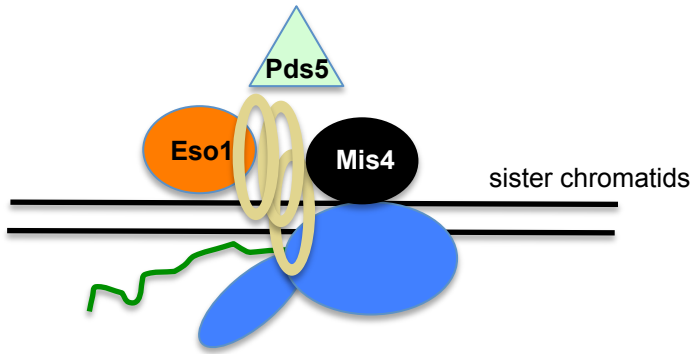
F





Mis4+/Rad21+

high transcription  
stable cohesion



Rad21+

low transcription  
unstable cohesion

