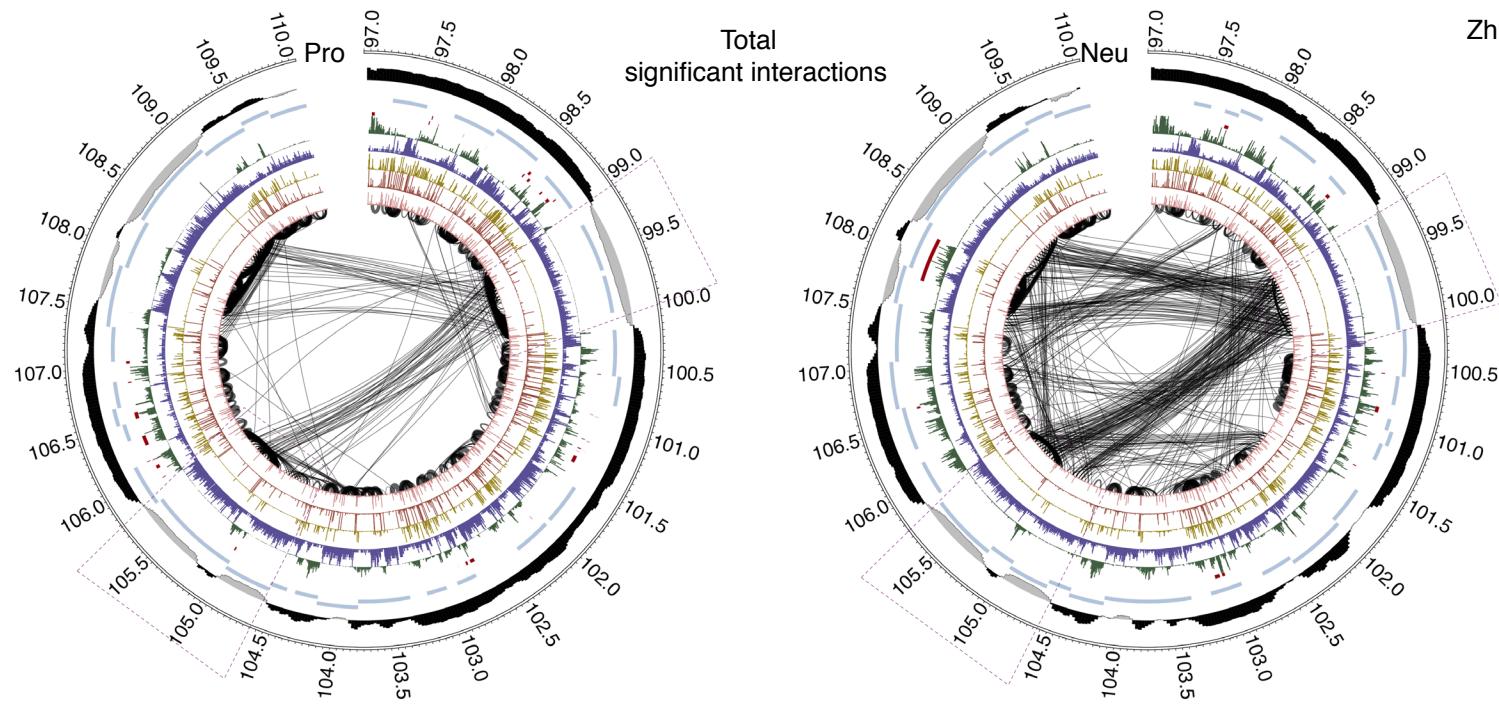
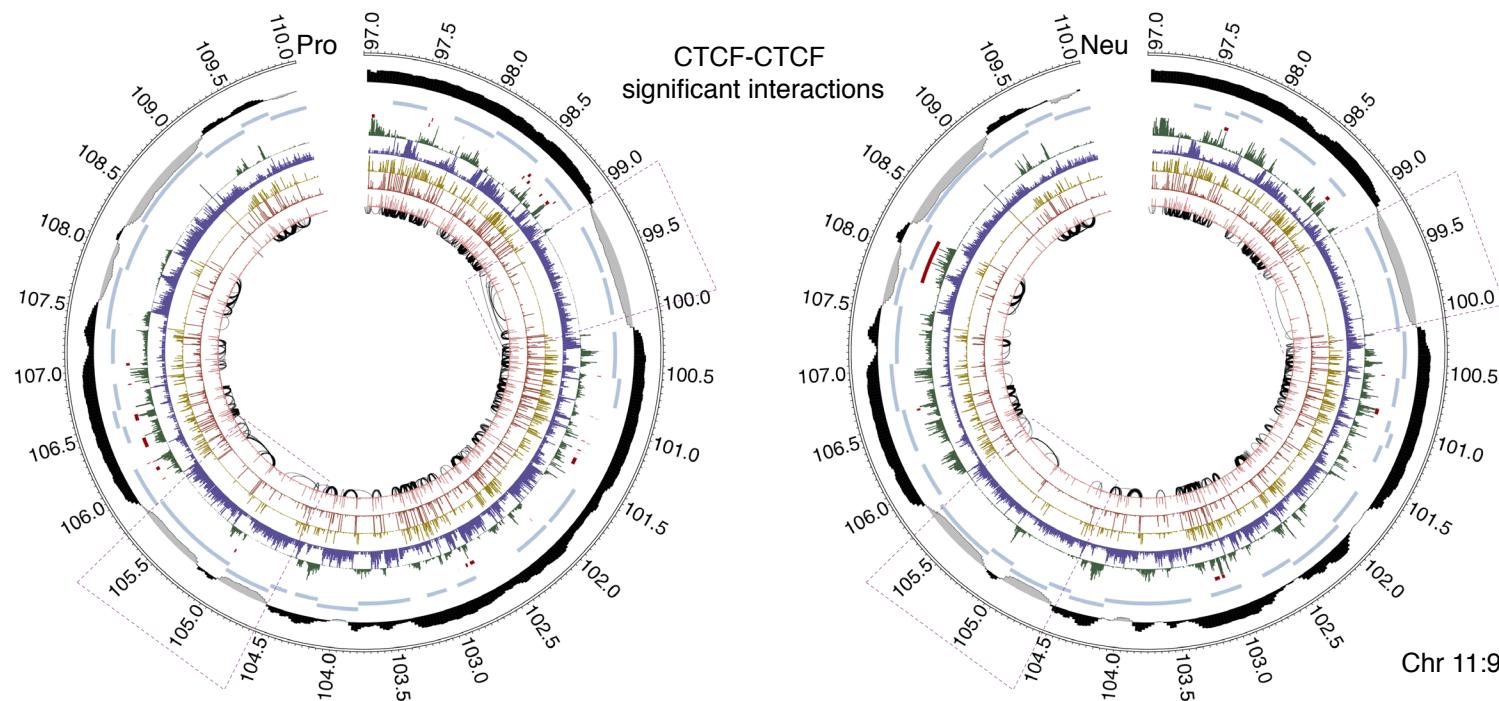


**A****B**

outer →  
inner  
 Less RAD21 peaks  
**PC1 TAD**  
 Diff. genes  
 GRO-Seq  
 H3K27me3  
 H3K4me2  
 RAD21  
 CTCF  
 Interactions

Chr 11:97,000,000-110,000,000