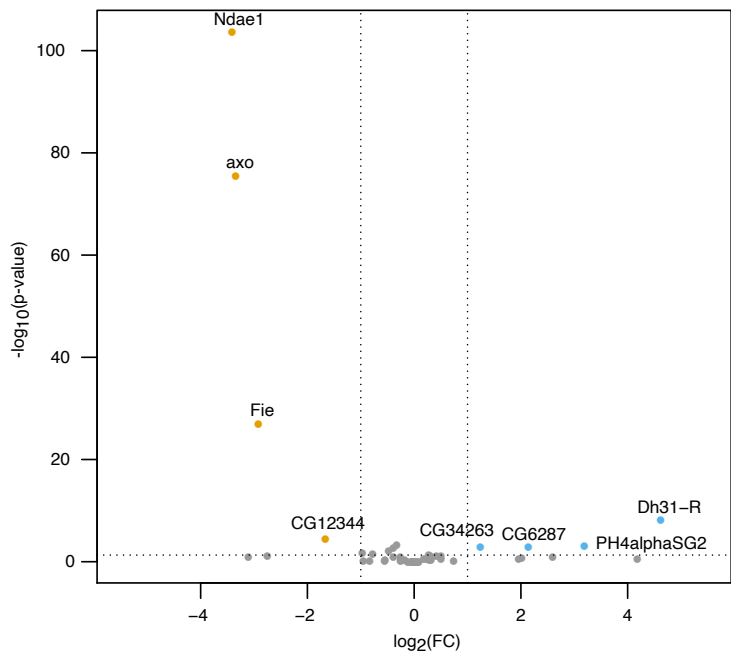
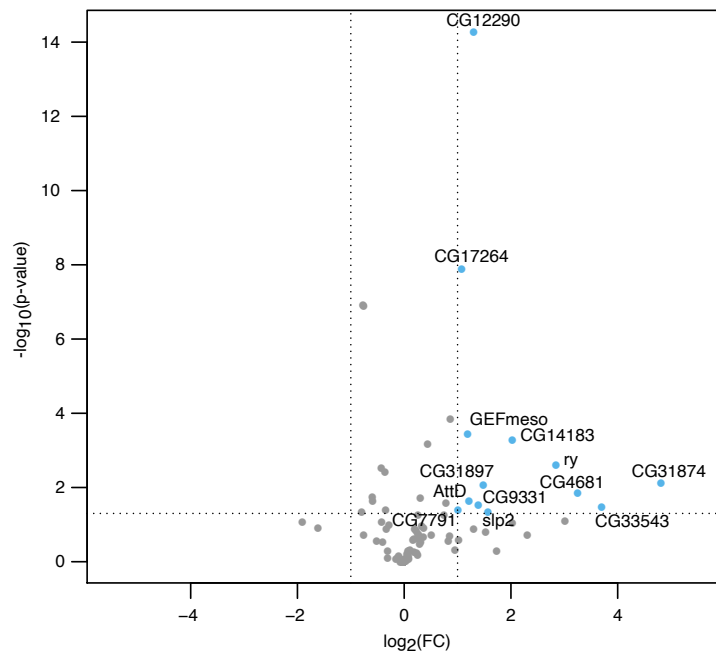


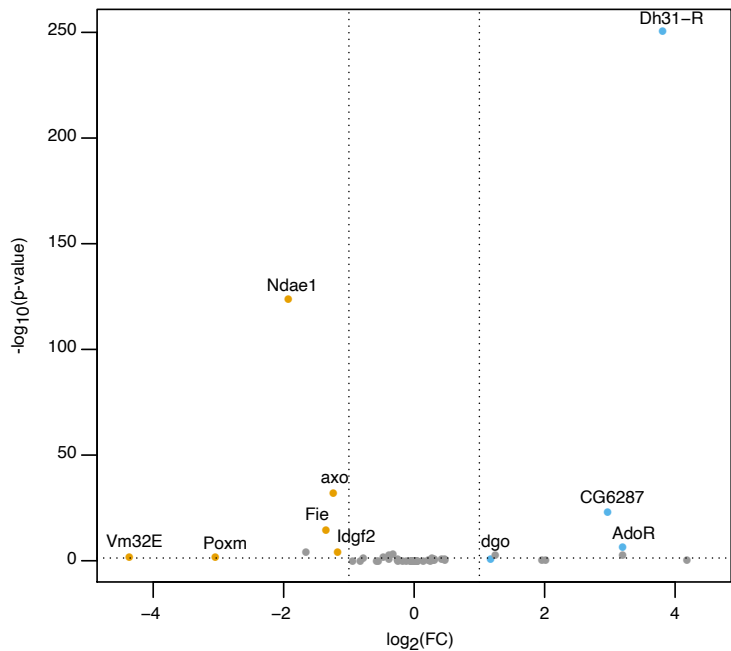
A

lost loops (BG3^{WT} vs BG3^{BEAF-32⁻})

B

maintained loops (BG3^{WT} vs BG3^{BEAF-32⁻})

C

lost loops (BG3^{WT} vs BG3^{Cp190⁻ Chro⁻})

D

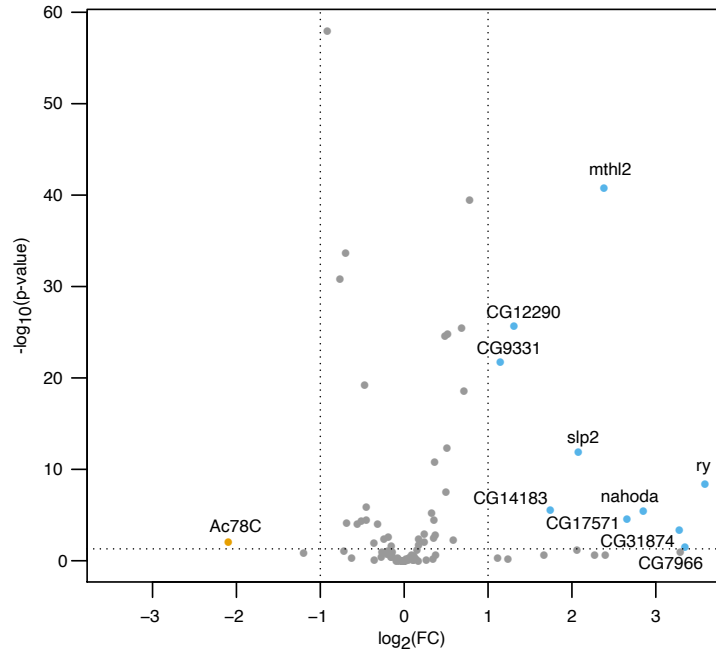
maintained loops (BG3^{WT} vs BG3^{Cp190⁻ Chro⁻})

Figure S12. *Differential expression of genes located at chromatin loops.* Volcano plots of the $-\log_{10}$ of the adjusted p-value as a function of \log_2 fold change. We considered all genes located at lost chromatin loops (A and C) and maintained chromatin loops (B and D) in BEAF-32 single knockdown (A and B) and Cp190 and Chro double knockdown (C and D).