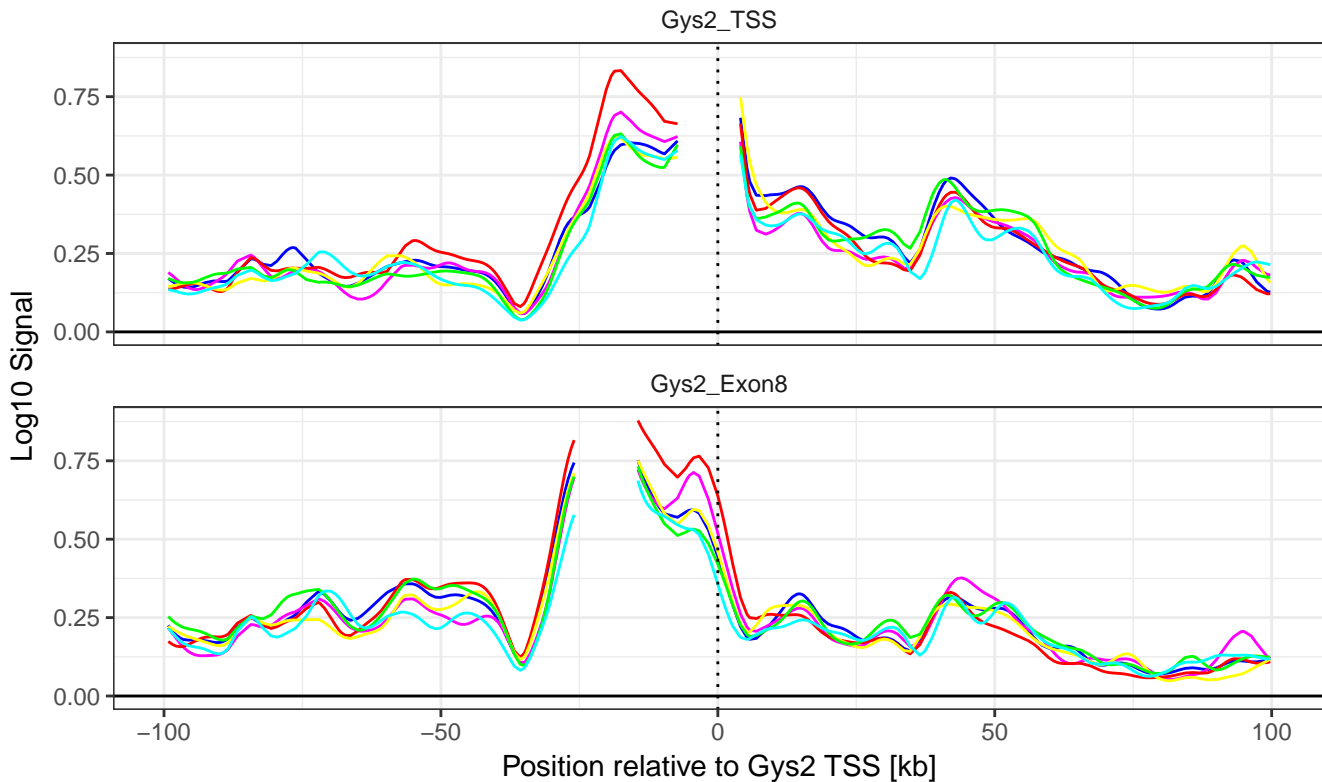


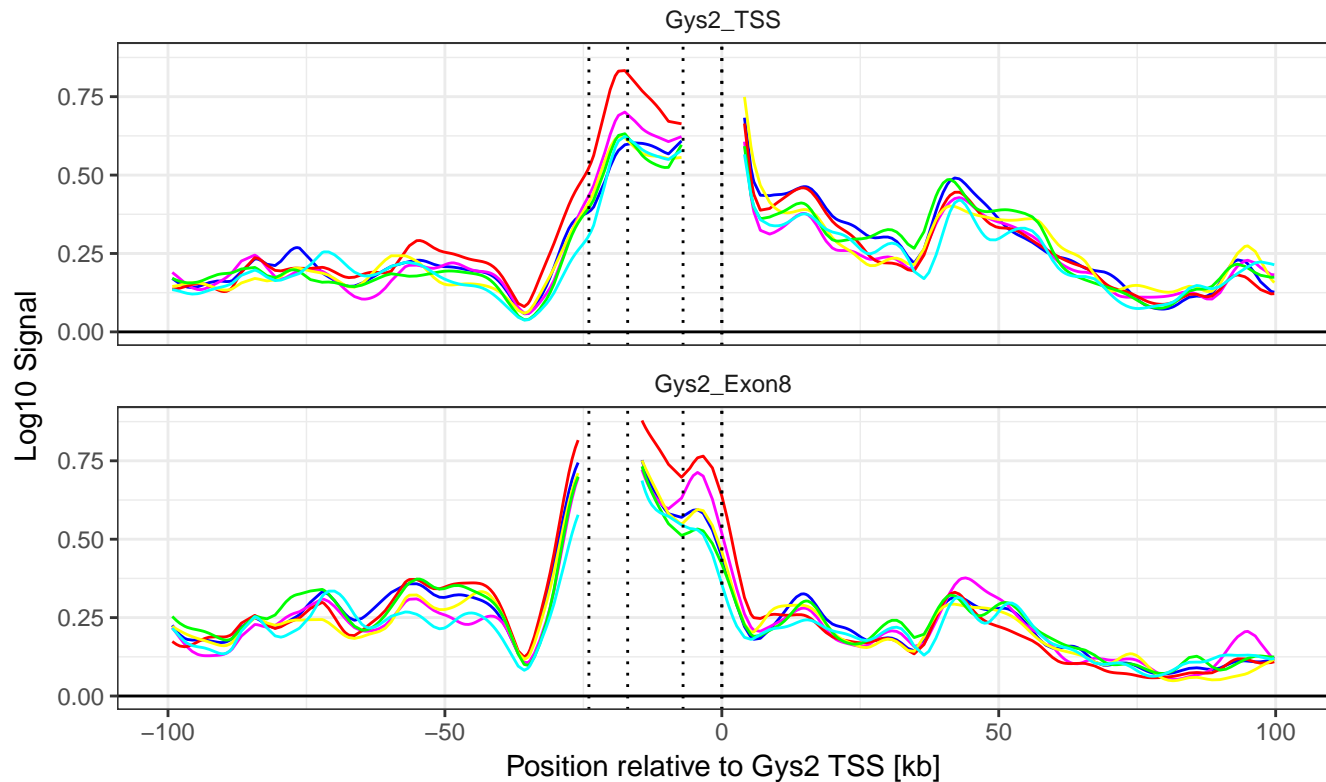
Gys2 merged



time

ZT00	ZT08	ZT16
ZT04	ZT12	ZT20

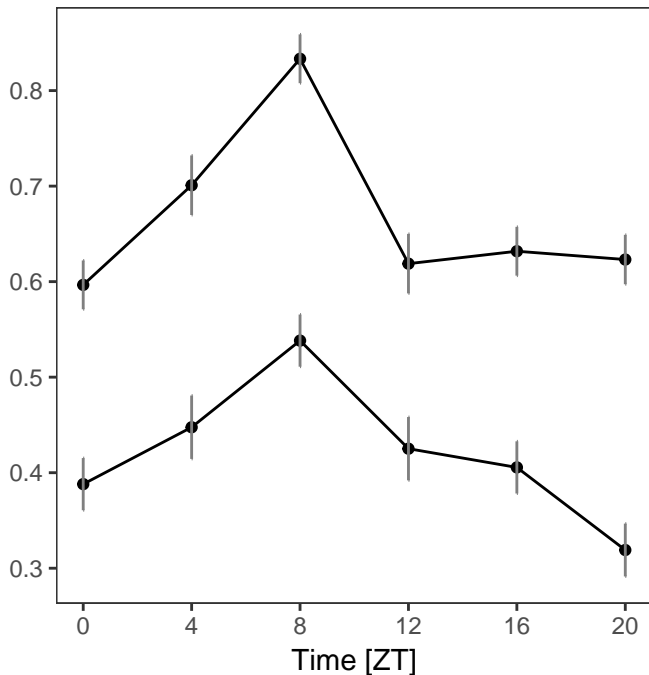
Gys2 merged



time

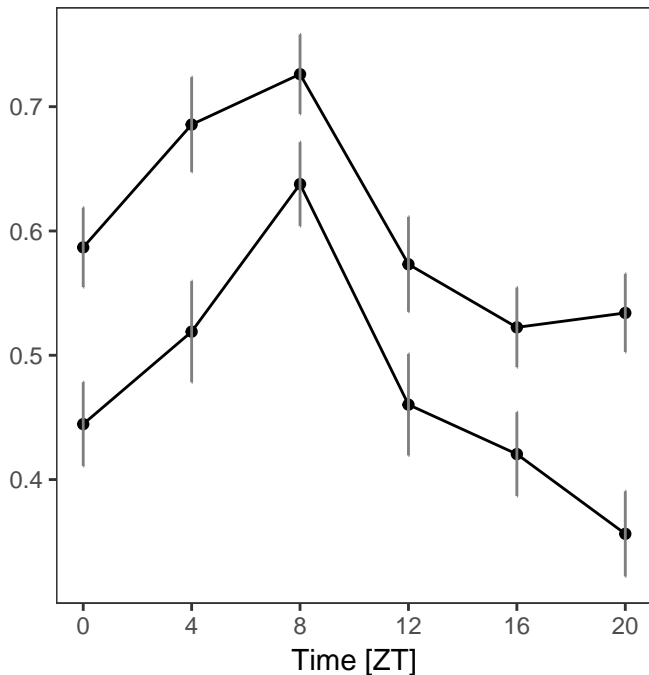
ZT00	ZT08	ZT16
ZT04	ZT12	ZT20

Pval=0.011, Phase=7.7, Log2FC=0.57



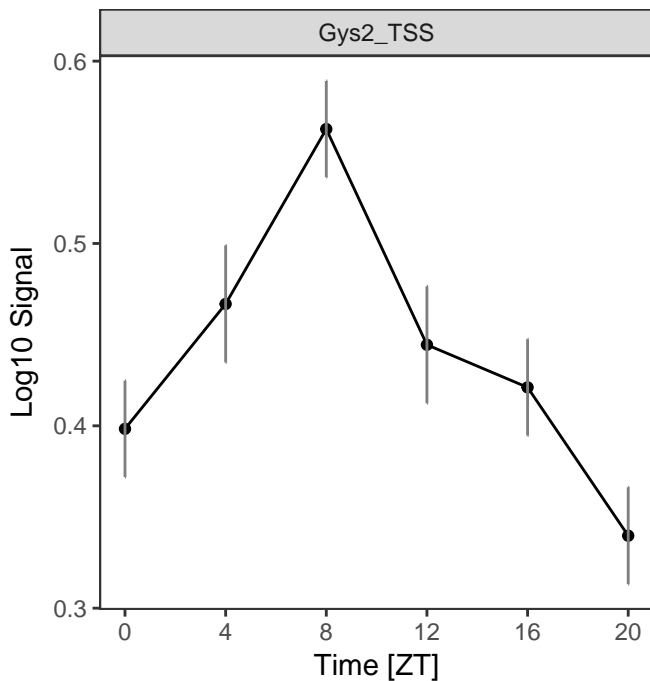
genotype — Gys2_TSS

Pval=4e-04, Phase=6.6, Log2FC=0.72



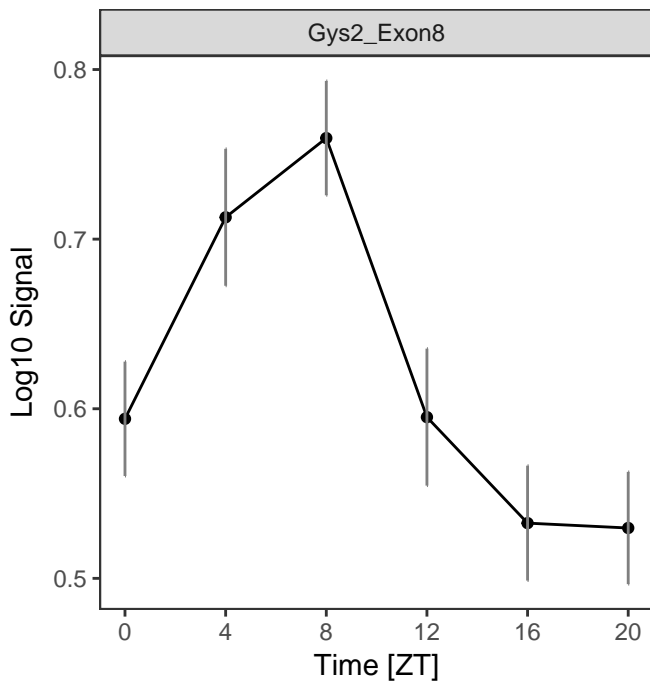
genotype — Gys2_Exon8

Fragment range: -24000 to -17000
Best frag: -23137.5 Pval=0.051, Phas

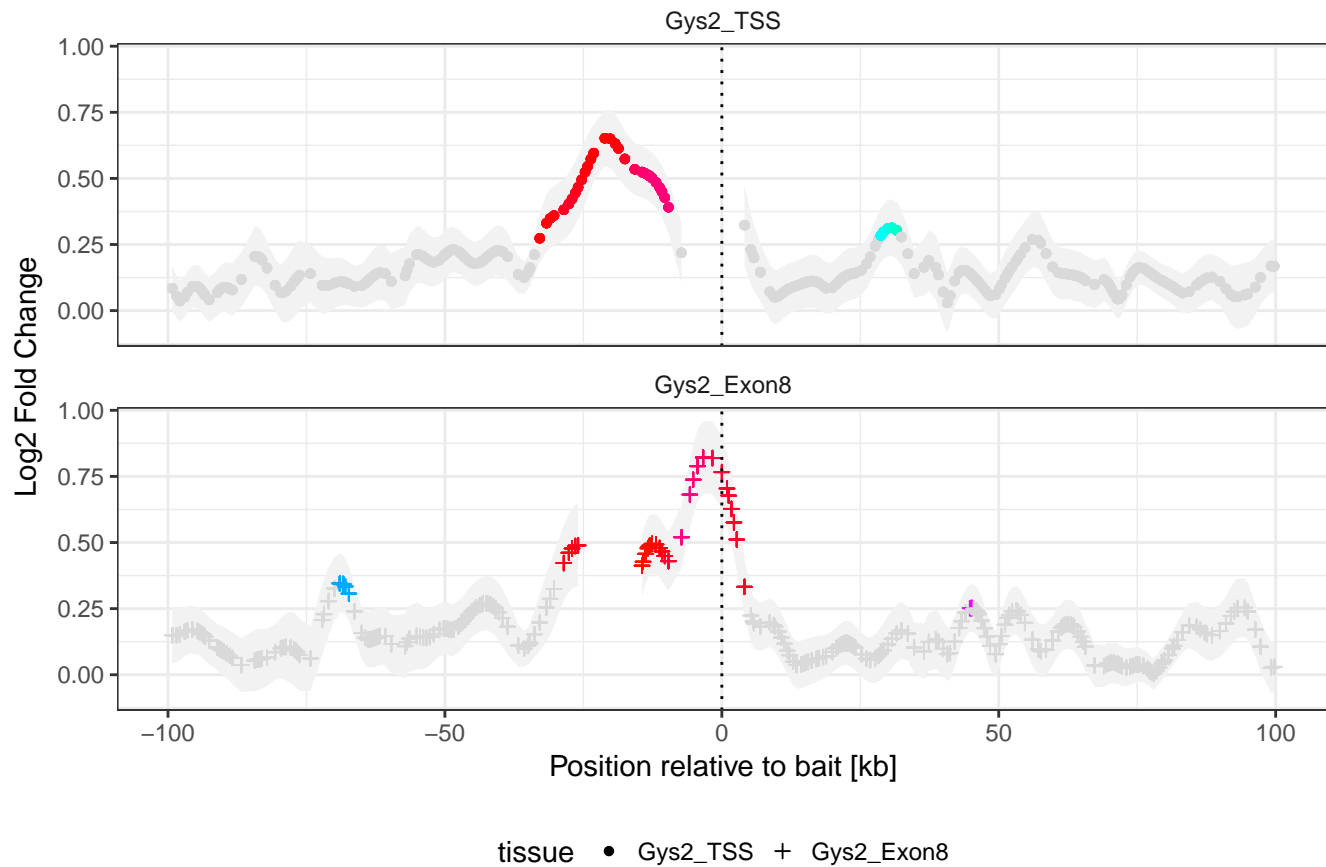


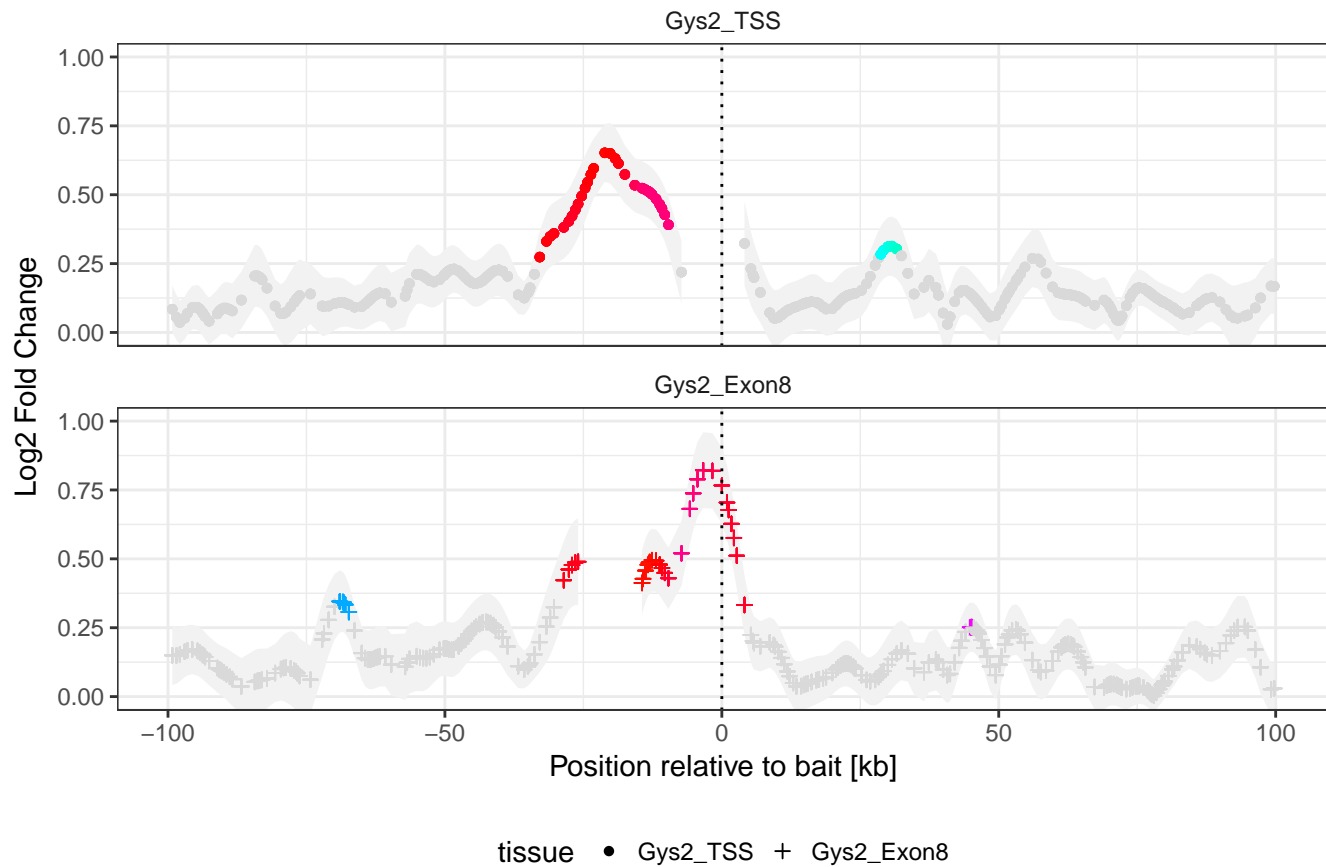
genotype — Gys2_TSS

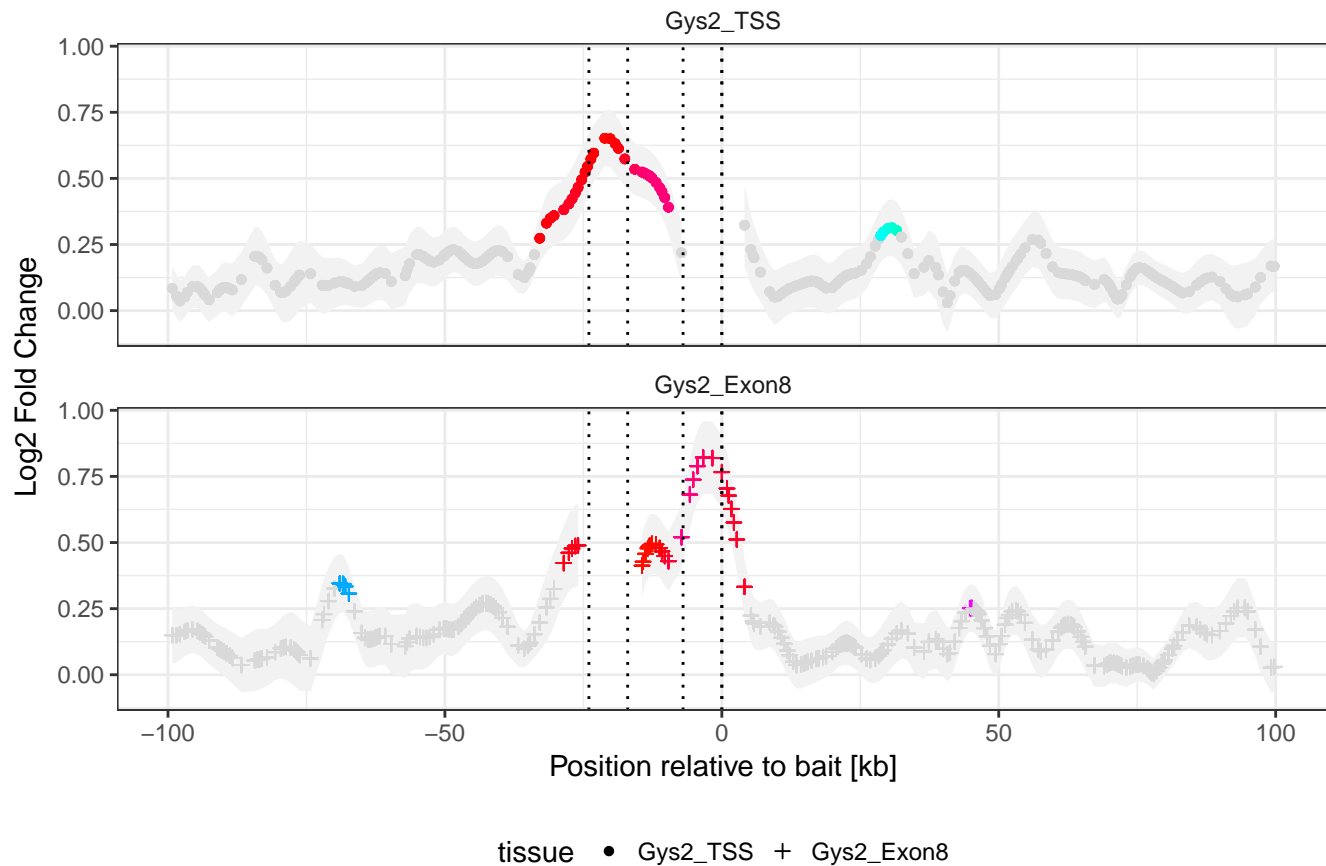
Fragment range: -7000 to 0
Best frag: -4383 Pval=0.016, Phase=6

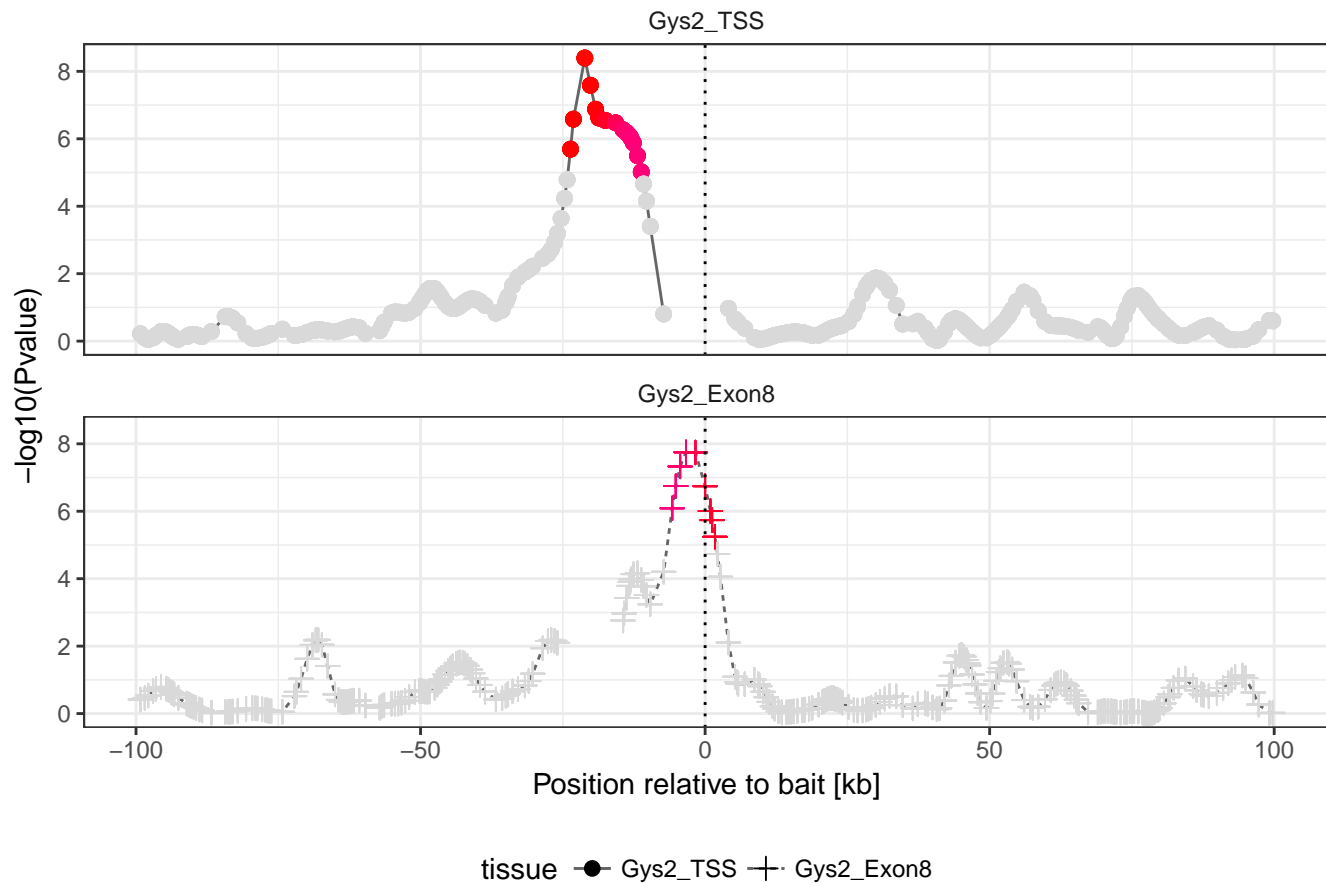


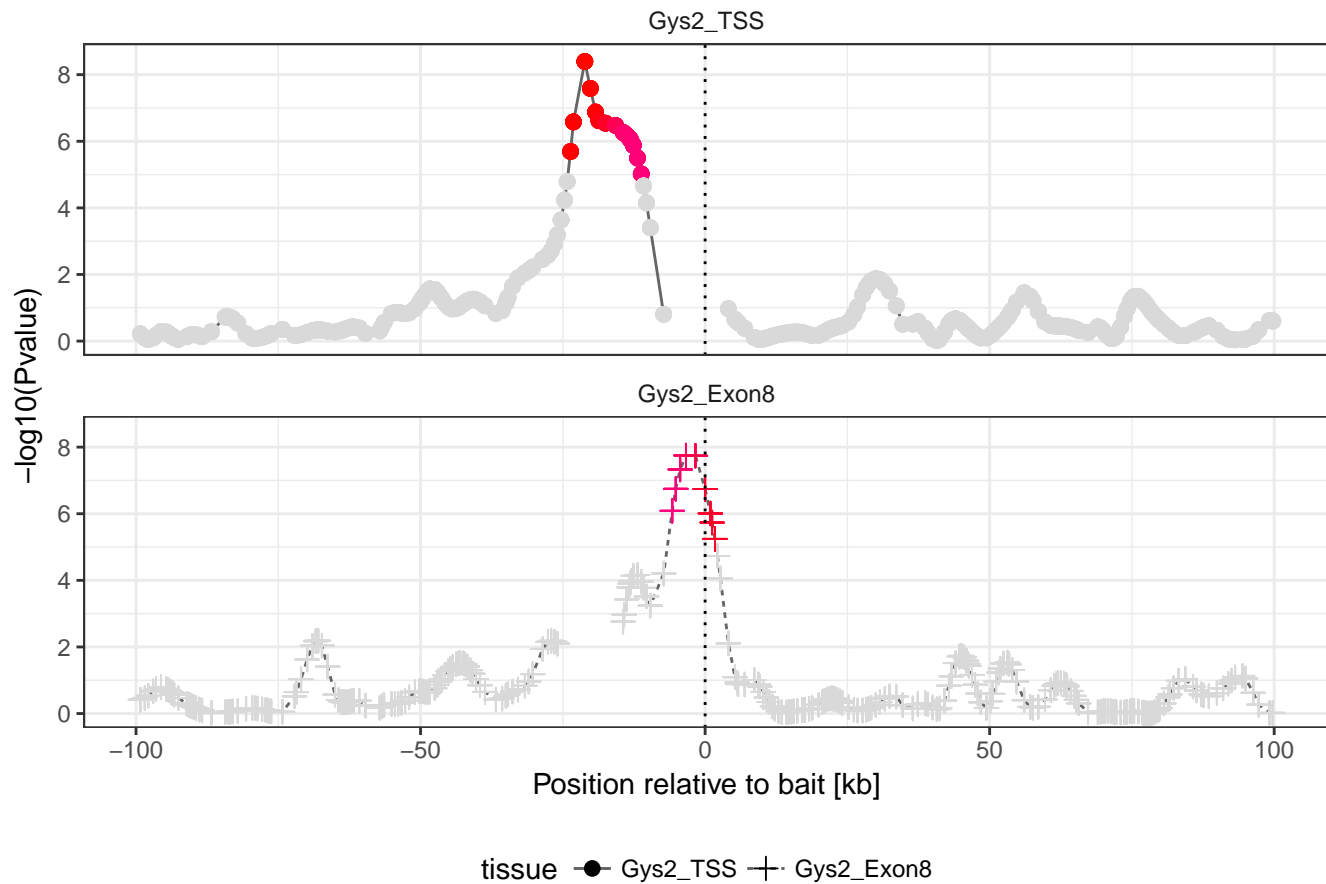
genotype — Gys2_Exon8

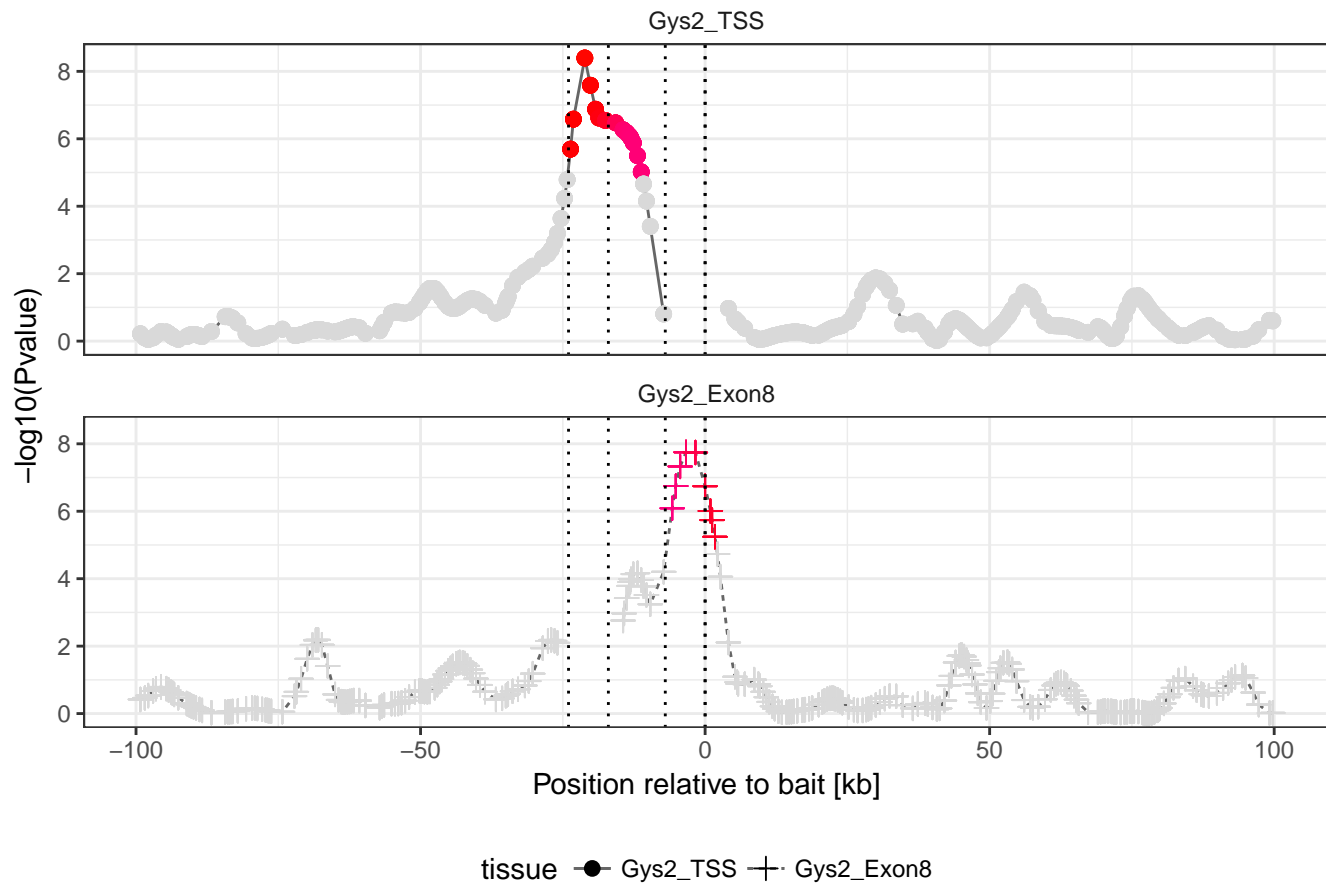


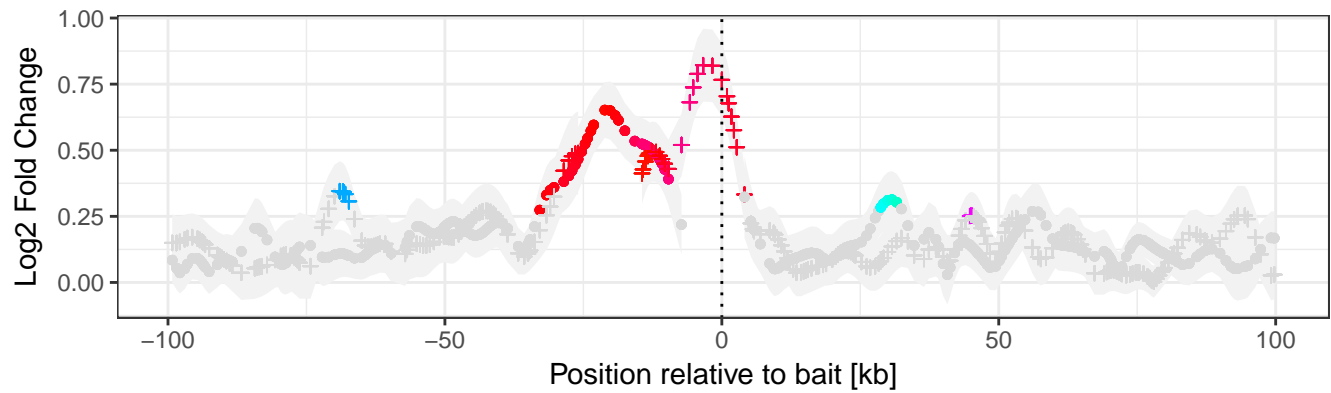




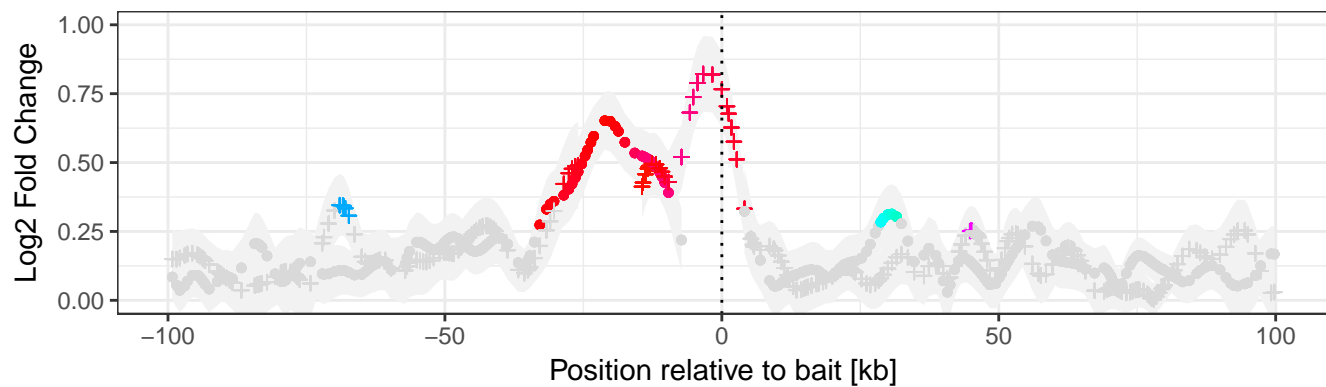




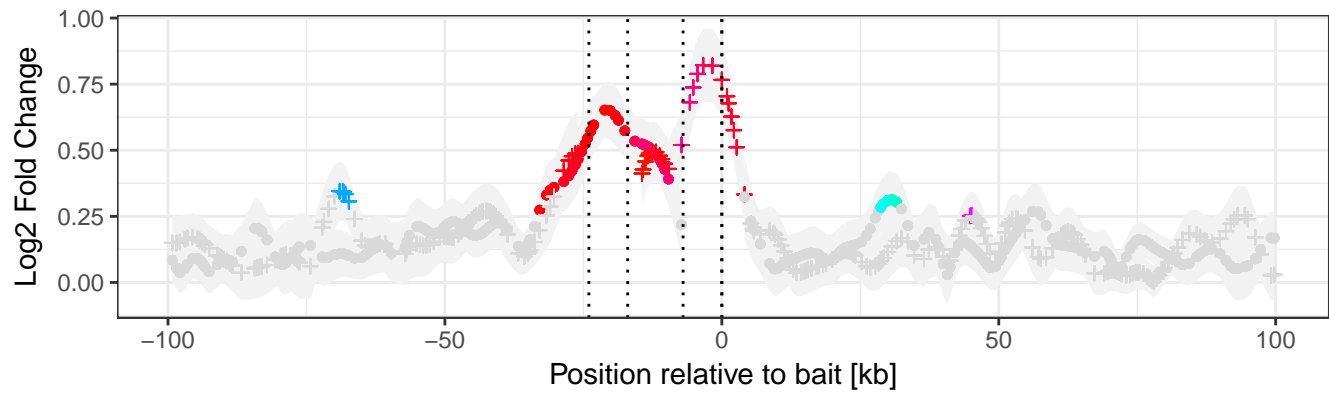




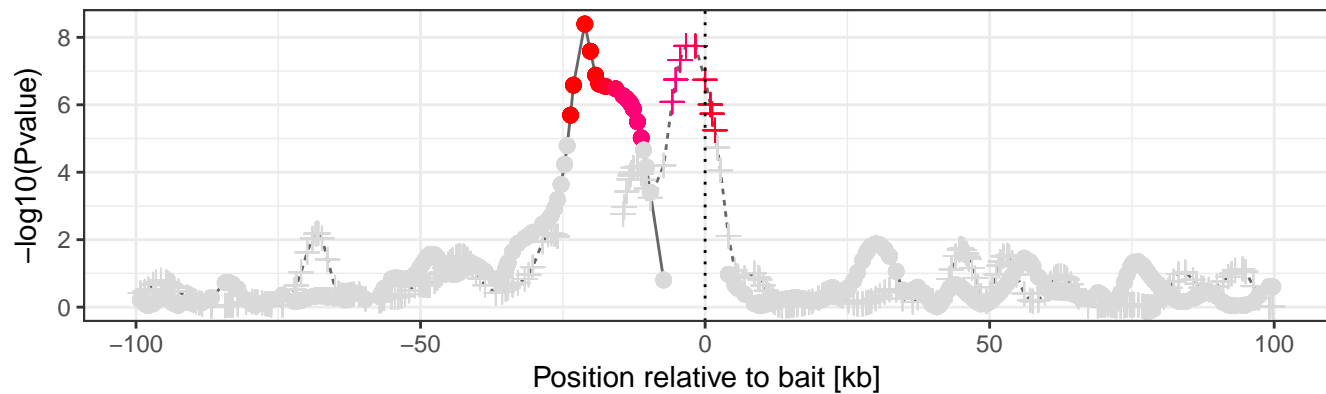
tissue • Gys2_TSS + Gys2_Exon8



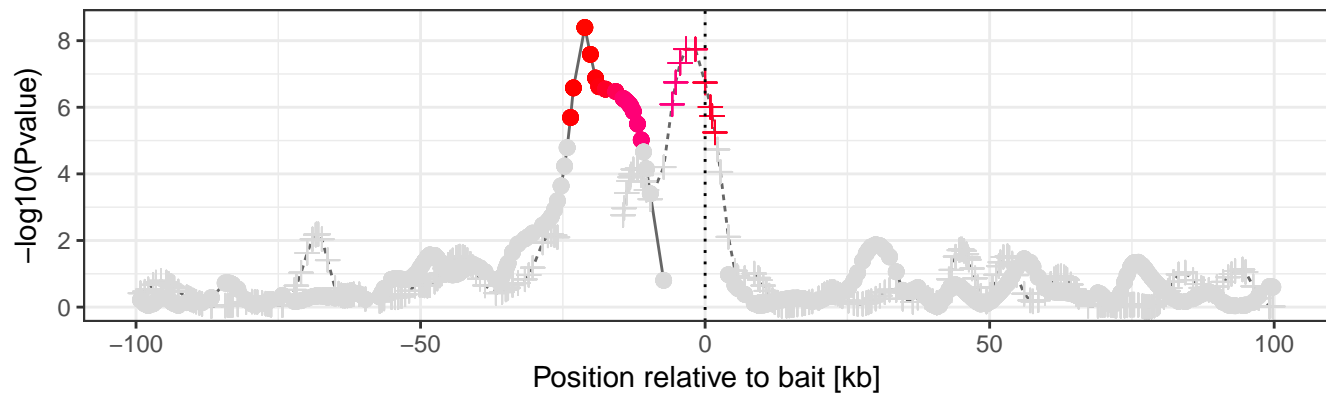
tissue • Gys2_TSS + Gys2_Exon8



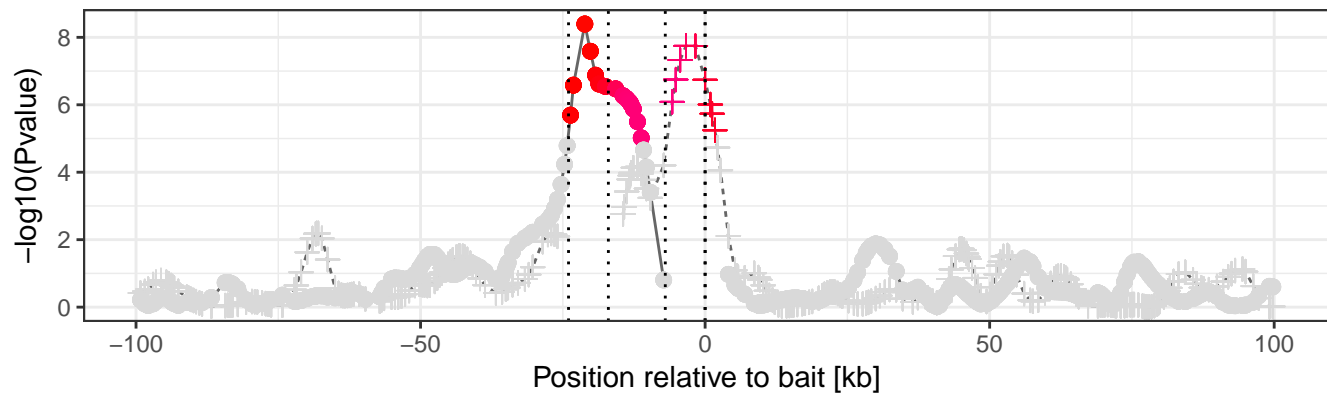
tissue • Gys2_TSS + Gys2_Exon8



tissue ● Gys2_TSS ✚ Gys2_Exon8



tissue ● Gys2_TSS ✚ Gys2_Exon8



tissue ● Gys2_TSS + Gys2_Exon8