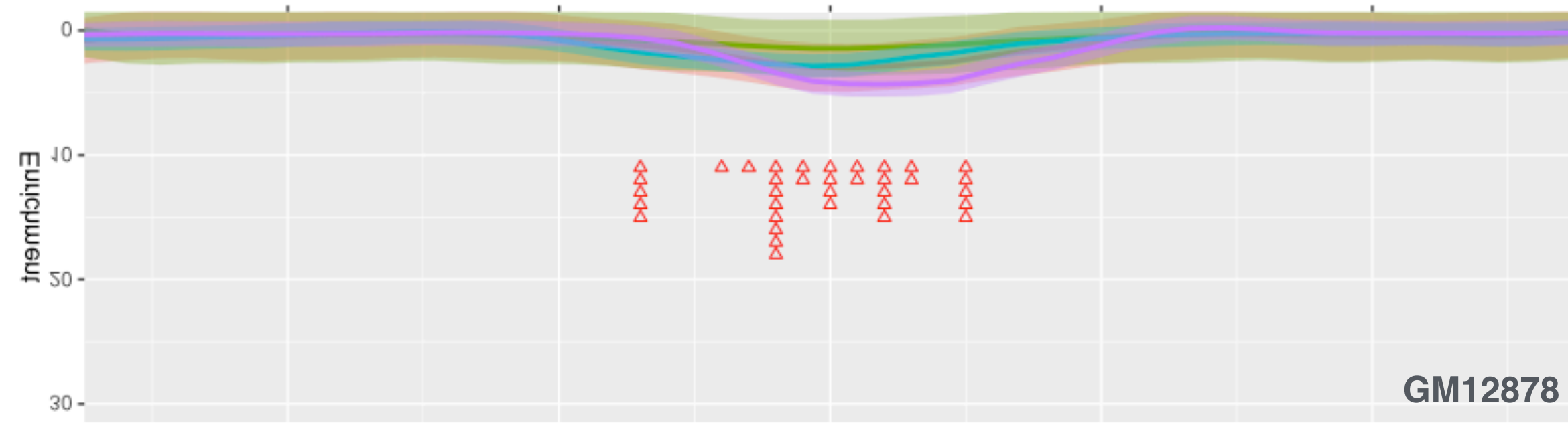
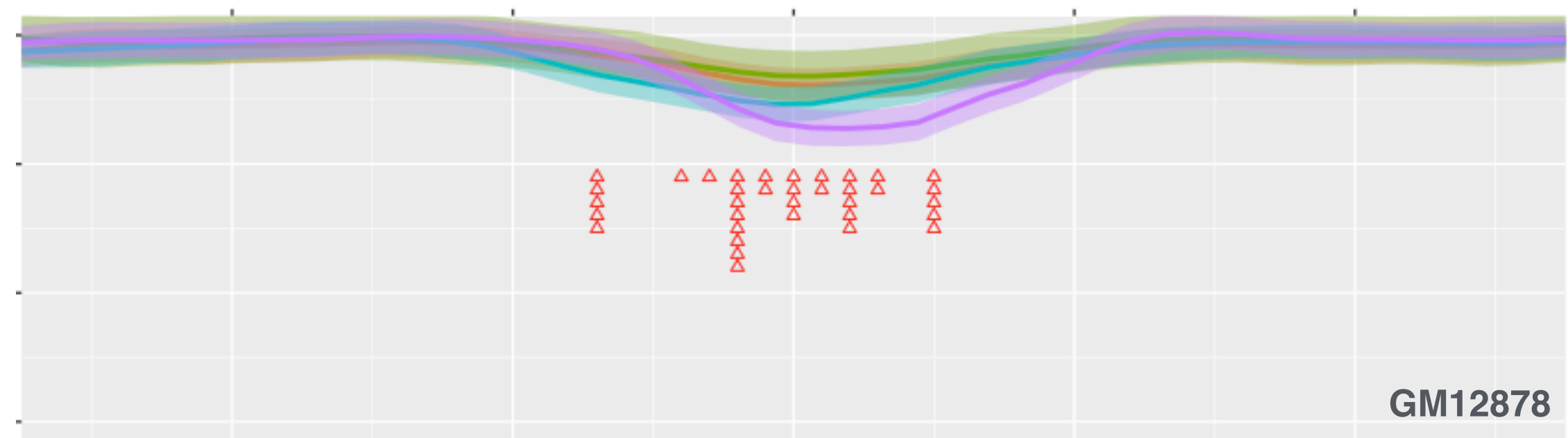


track

- DNase
- H3K27ac
- H3K4me1
- H3K4me3

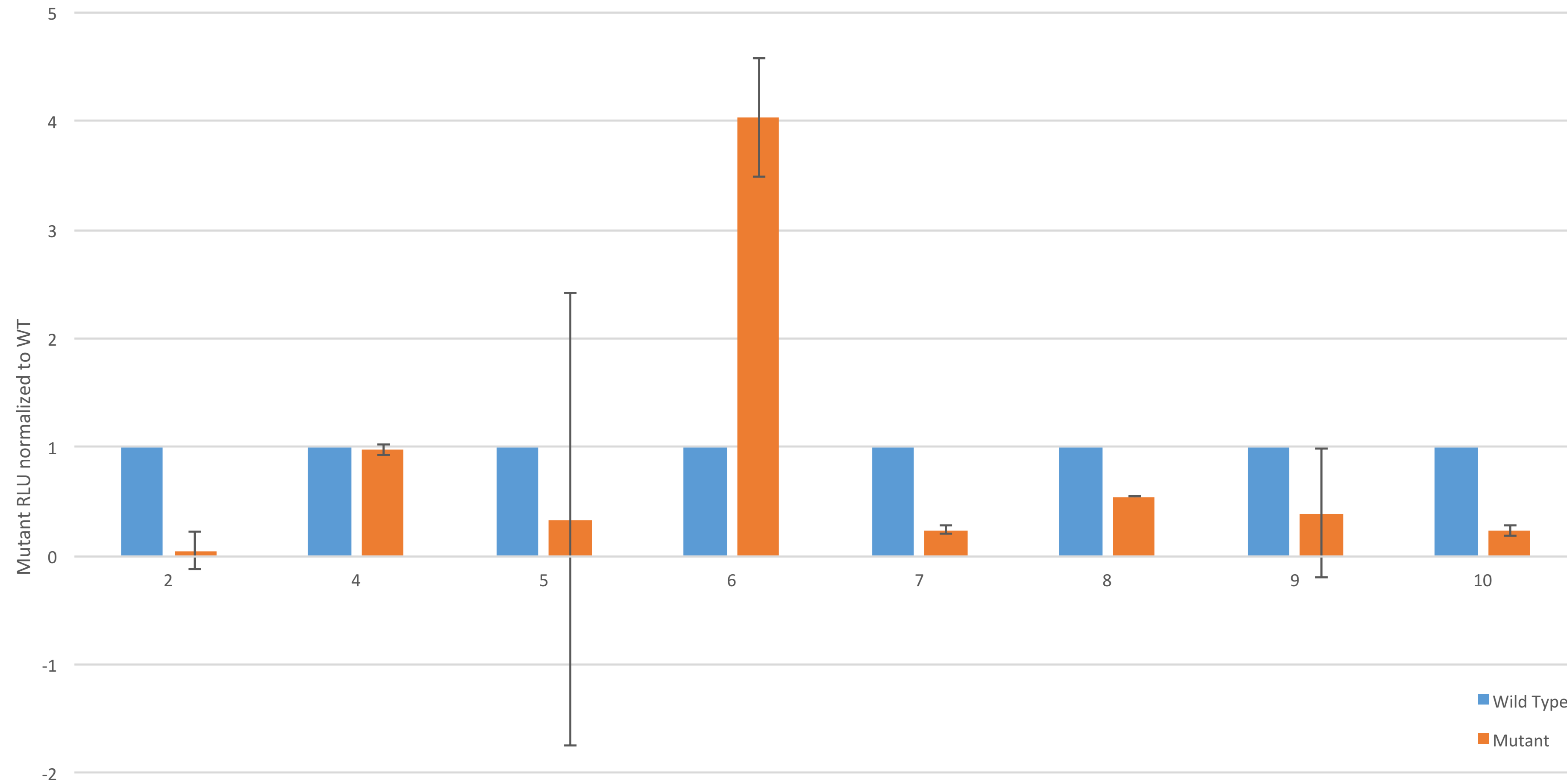
BCR

BCR



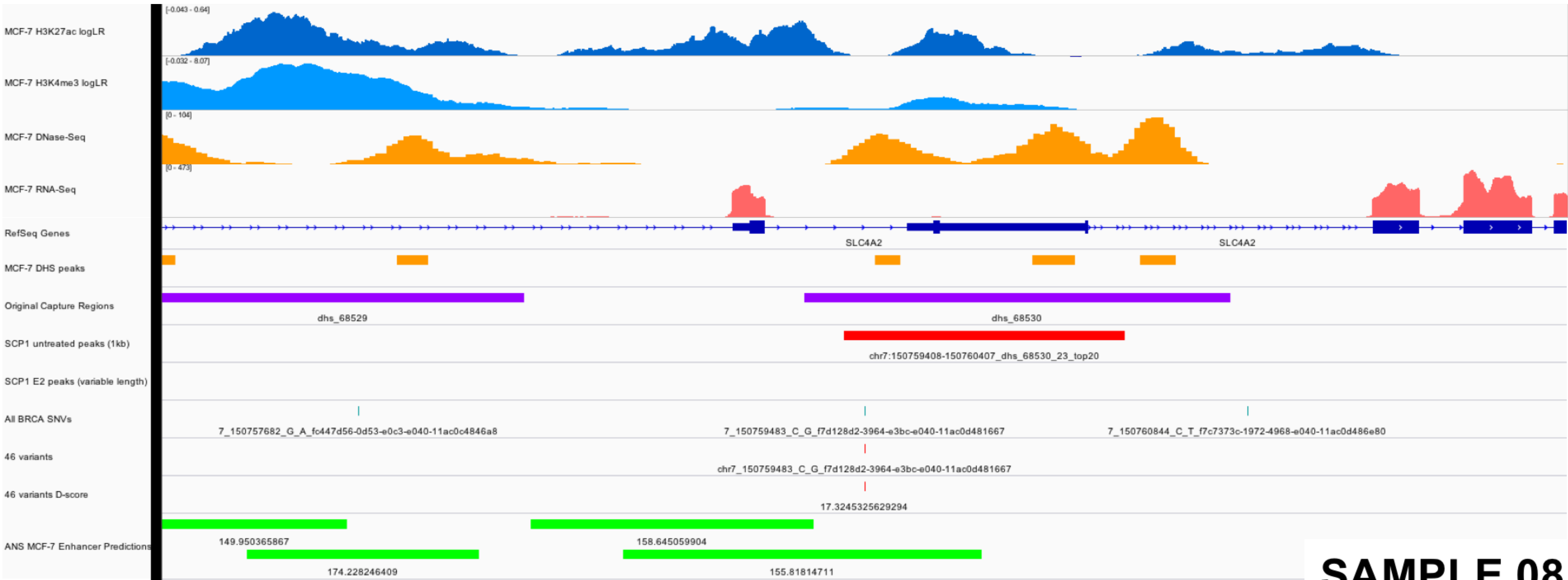
<fold change signal>

<p-val signal>

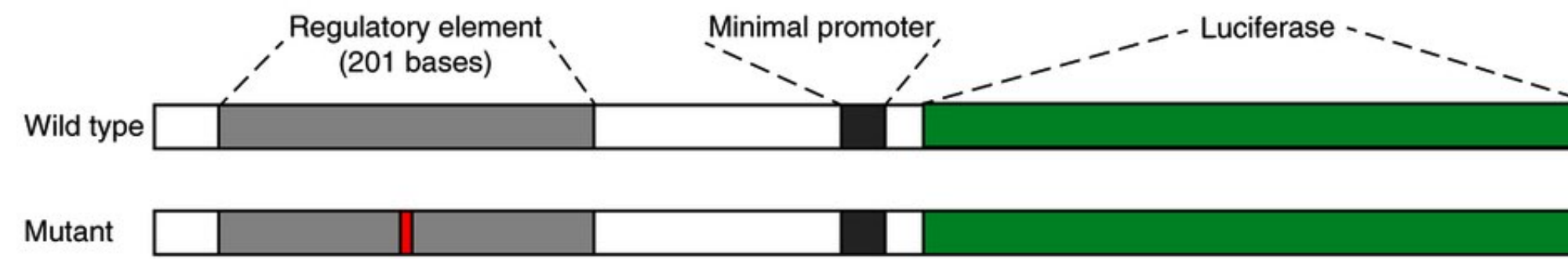
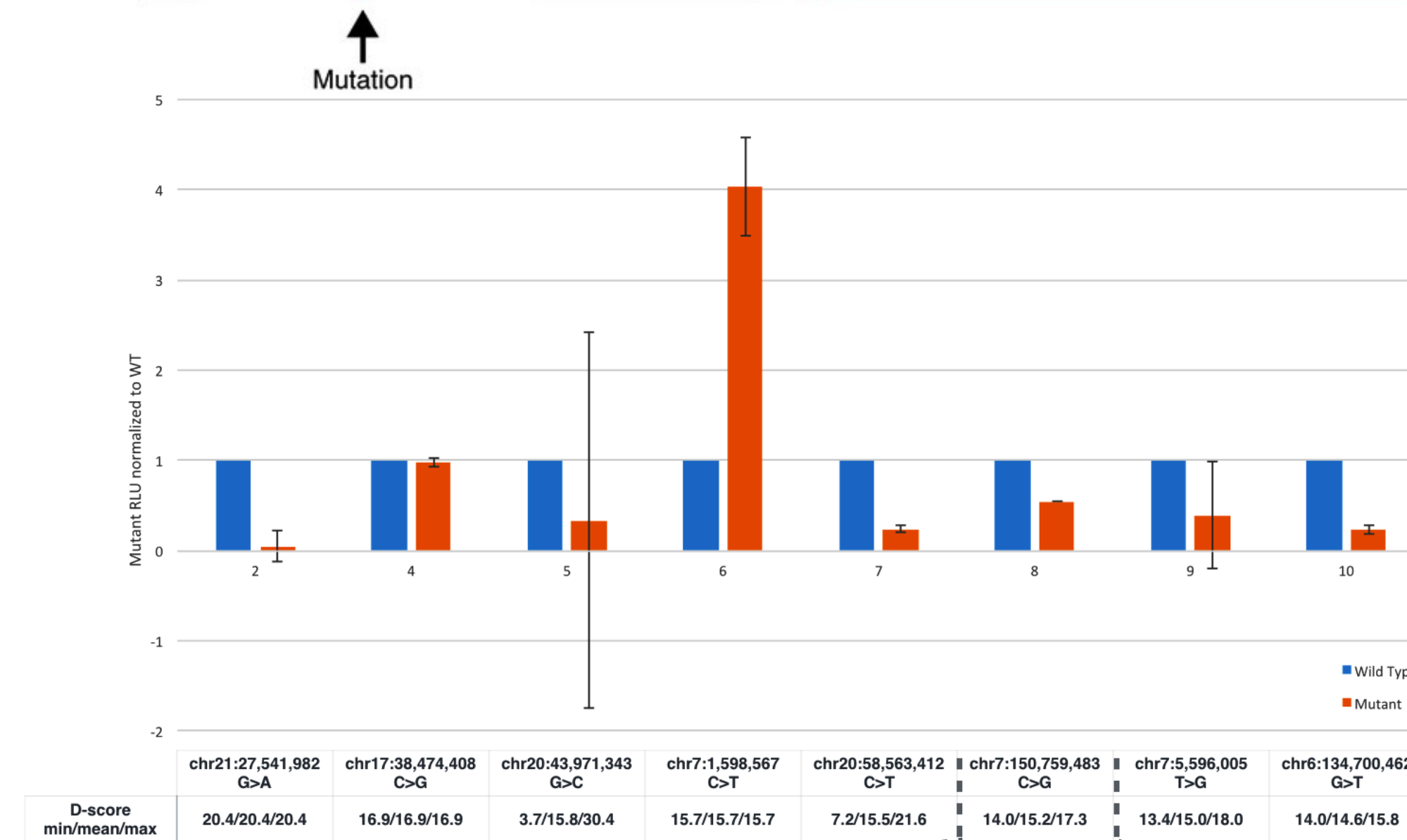


	chr21:27,541,982 G>A	chr17:38,474,408 C>G	chr20:43,971,343 G>C	chr7:1,598,567 C>T	chr20:58,563,412 C>T	chr7:150,759,483 C>G	chr7:5,596,005 T>G	chr6:134,700,462 G>T
<b>D-score min/mean/max</b>	20.4/20.4/20.4	16.9/16.9/16.9	3.7/15.8/30.4	15.7/15.7/15.7	7.2/15.5/21.6	14.0/15.2/17.3	13.4/15.0/18.0	14.0/14.6/15.8

*\*Sample 1 & 3 omitted due to issues with plasmid isolation*



**SAMPLE 08**

**a****b****c**