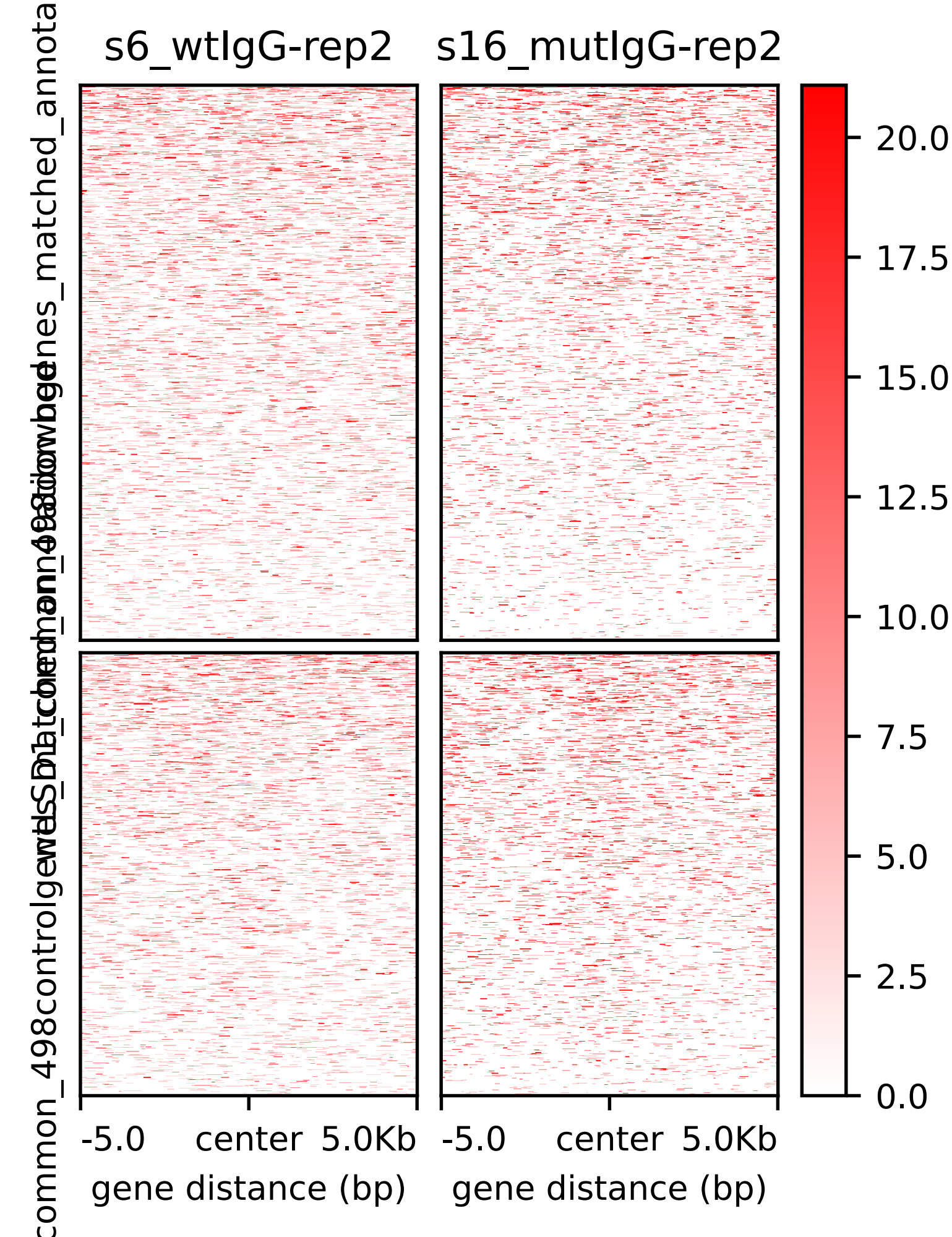
s6_wtIgG-rep2
s16_mutIgG-rep2
20.0
17.5
15.0
12.5
10.0
7.5
5.0
2.5
0.0
-5.0
center
5.0Kb
gene distance (bp)
-5.0
center
5.0Kb
gene distance (bp)
common_498controlgenes_matched_annotated.bed
wtSD1_common_498controlgenes_matched_annotated.bed