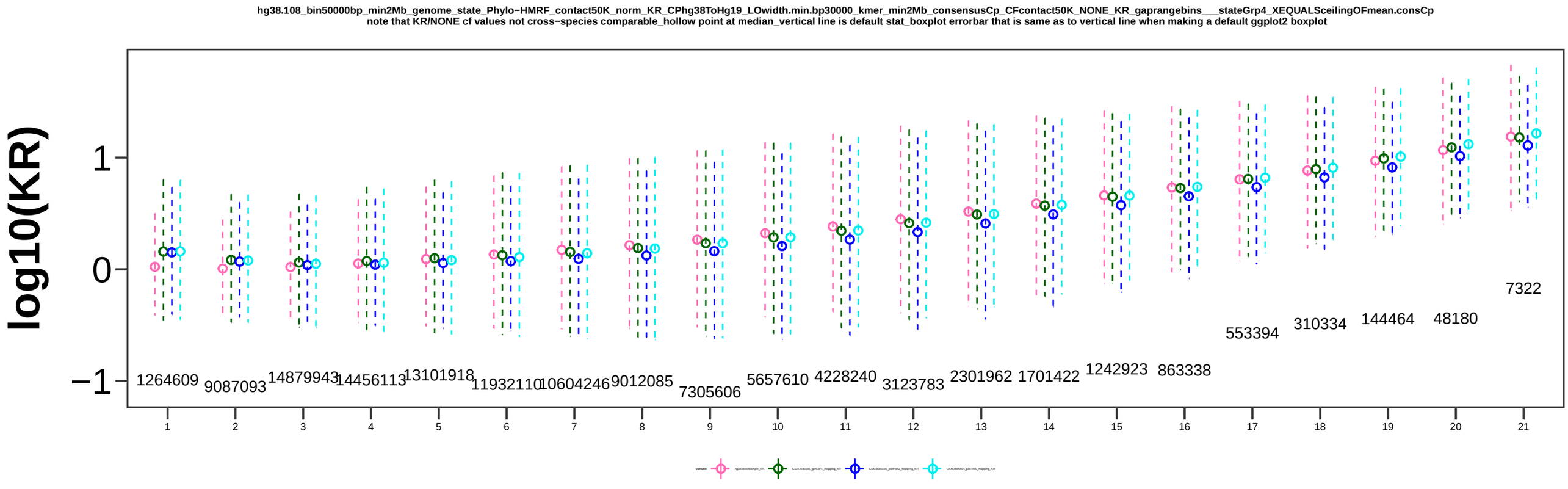
hg38.108_bin50000bp_min2Mb_genome_state_Phylo−HMRF_contact50K_norm_KR_CPhg38ToHg19_LOwidth.min.bp30000_kmer_min2Mb_consensusCp_CFcontact50K_NONE_KR_gaprangebins___stateGrp4_XEQUALSceilingOFmean.consCp
note that KR/NONE cf values not cross−species comparable_hollow point at median_vertical line is default stat_boxplot errorbar that is same as to vertical line when making a default ggplot2 boxplot
log10(KR)
1
0
−1
1264609
9087093
14879943
14456113
13101918
11932110
10604246
9012085
7305606
5657610
4228240
3123783
2301962
1701422
1242923
863338
553394
310334
144464
48180
7322
1
2
3
4
5
6
7
8
9
10
11
12
13
14
15
16
17
18
19
20
21
variable
hg38.downsample_KR
GSM3685956_gorGor4_mapping_KR
GSM3685955_panPan2_mapping_KR
GSM3685954_panTro5_mapping_KR