

hg38.108_bin50000bp_min2Mb_genome_state_Phylo-HMRF_contact50K_norm_KR_CPhg38ToHg19_LOwidth.min.bp30000_kmer_min2Mb_consensusCp_CFcontact50K_NONE_KR_gaprangebins__stateGrp4_XEQUALSceilingOFMODE.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

$\log_{10}(KR)$

