

hg38.108\_bin50000bp\_min2Mb\_genome\_state\_Phylo-HMRF\_contact50K\_norm\_KR\_CPhg38ToHg19\_LOwidth.min.bp30000\_kmer\_min2Mb\_consensusCp\_CFcontact50K\_NONE\_KR\_gaprangebins\_\_stateGrp4\_XEQUALScellingOFmean.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

$\log_{10}(\text{KR})$

