

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

count

