

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

