



gene_length.meta

18mo.2mo.log_ratio.mcg_genebody

9mo.2mo.activity.log_ratio.max.20kb-1Mb.enhancer_peak

18mo.9mo.contact.log_ratio.min.20kb-1Mb.enhancer_peak

count.DAR

18mo.2mo.log_ratio.std.atac 9mo.2mo.activity.log_ratio.count.0-10kb.enhancer_peak 18mo.2mo.T.median.loops

18mo.2mo.activity.log_ratio.std.20kb-1Mb.enhancer_peak 18mo.2mo.activity.log_ratio.count.20kb-1Mb.enhancer_DMR 18mo.9mo.activity.log_ratio.mean.20kb-1Mb.enhancer_peak 18mo.2mo.Q.min.loops

18mo.9mo.Q.median.loops

9mo.2mo.activity.log_ratio.std.20kb-1Mb.enhancer_peak 9mo.2mo.Q.std.loops

9mo.2mo.log ratio.std.atac

9mo.2mo.log ratio.DMR

9mo.2mo.log_ratio.mcg_genebody

 $18mo. 9mo. activity. log_ratio. max. 20kb-1 Mb. enhancer_peak$

Sum of 60 other features

