





gene_length.meta
18mo.2mo.activity.log_ratio.min.20kb-1Mb.enhancer_peak
9mo.2mo.activity.log_ratio.count.20kb-1Mb.enhancer_DMR
18mo.2mo.log_ratio.std.atac
18mo.9mo.Q.std.loops
count.DAR

Qanova.min.loops

9mo.2mo.activity.log_ratio.mean.20kb-1Mb.enhancer_peak 9mo.2mo.activity.log_ratio.min.20kb-1Mb.enhancer_peak 18mo.9mo.log_ratio.std.atac

18mo.2mo.activity.log_ratio.median.20kb-1Mb.enhancer_peak 18mo.2mo.contact.log_ratio.std.20kb-1Mb.enhancer_peak

log2_distance.std.atac

9mo.2mo.log_ratio.DMR

9mo.2mo.log_ratio.mean.atac

18mo.9mo.log_ratio.mch_genebody

18mo.9mo.T.mean.loops

Sum of 102 other features