

18mo.2mo.log_ratio.mch_genebody
9mo.2mo.log_ratio.mch_genebody
18mo.2mo.log_ratio.max.atac
loop_size.log.mean.loops
18mo.9mo.Q.std.loops
gene_length.meta
18mo.2mo.log_ratio.mean.atac
count.atac

18mo.9mo.contact.log_ratio.max.20kb-5Mb.enhancer_peak
18mo.2mo.activity.log_ratio.min.20kb-5Mb.enhancer_peak
9mo.2mo.activity.log_ratio.median.20kb-5Mb.enhancer_peak
distance.log2.DMR

18mo.2mo.activity.log_ratio.median.20kb-5Mb.enhancer_peak 9mo.2mo.activity.log_ratio.mean.20kb-5Mb.enhancer_peak 18mo.2mo.contact.log_ratio.median.20kb-5Mb.enhancer_DMR 18mo.9mo.T.median.loops

9mo.2mo.contact.log_ratio.max.20kb-5Mb.enhancer_peak loop_size.log.min.loops

18mo.2mo.activity.log_ratio.max.20kb-5Mb.enhancer_peak
Sum of 41 other features



