

gene\_length.meta
18mo.2mo.log ratio.mcg genebody

18mo.9mo.log\_ratio.std.atac

9mo.2mo.log\_ratio.std.atac

18mo.9mo.log ratio.mean.atac

9mo.2mo.log\_ratio.DAR

18mo.2mo.activity.log\_ratio.median.20kb-1Mb.enhancer\_peak

18mo.2mo.log\_ratio.mean.atac

9mo.2mo.activity.log ratio.min.10kb-20kb.enhancer DMR

count.atac

18mo.2mo.log ratio.DAR

18mo.2mo.contact.log ratio.std.20kb-1Mb.enhancer peak

18mo.9mo.T.max.loops

log2 distance.mean.atac

 $18mo.9mo.contact.log\_ratio.median.0-10kb.enhancer\_DMR$ 

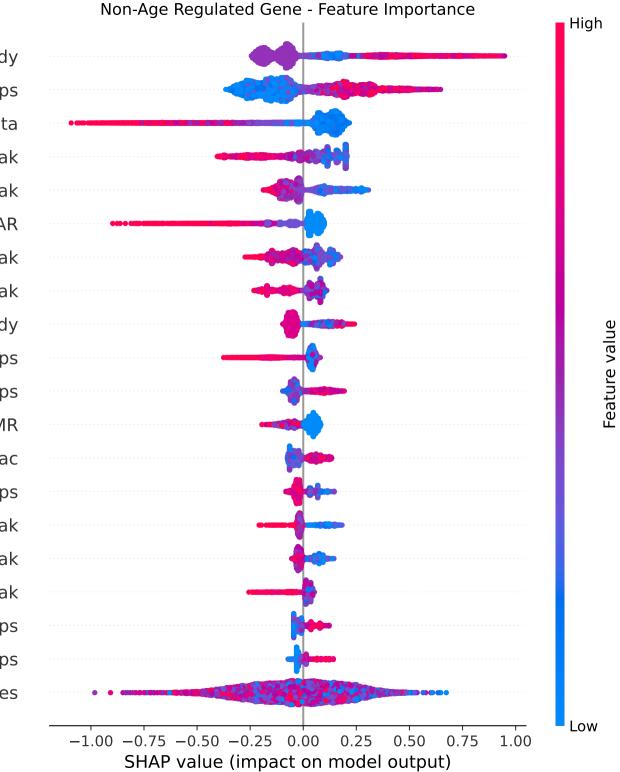
18mo.2mo.log ratio.mch genebody

 $18mo.9mo.activity.log\_ratio.max.0-10kb.enhancer\_DMR$ 

9mo.2mo.log\_ratio.mch\_genebody

9mo.2mo.Q.mean.loops

Sum of 234 other features



18mo.9mo.log ratio.mch genebody 9mo.2mo.Q.max.loops gene length.meta 18mo.2mo.activity.log ratio.mean.20kb-1Mb.enhancer peak 18mo.2mo.activity.log ratio.max.0-10kb.enhancer peak count.DAR 18mo.2mo.activity.log ratio.median.20kb-1Mb.enhancer peak

18mo.9mo.activity.log ratio.min.20kb-1Mb.enhancer peak 18mo.2mo.log ratio.mch genebody

Tanova.std.loops

9mo.2mo.Q.mean.loops

18mo.9mo.activity.log ratio.std.20kb-1Mb.enhancer DMR

18mo.9mo.log ratio.std.atac

18mo.9mo.Q.min.loops

18mo.9mo.activity.log ratio.mean.0-10kb.enhancer peak

18mo.9mo.activity.log ratio.mean.20kb-1Mb.enhancer peak

18mo.9mo.activity.log ratio.median.20kb-1Mb.enhancer peak

9mo.2mo.T.std.loops

18mo.2mo.T.std.loops

Sum of 234 other features

