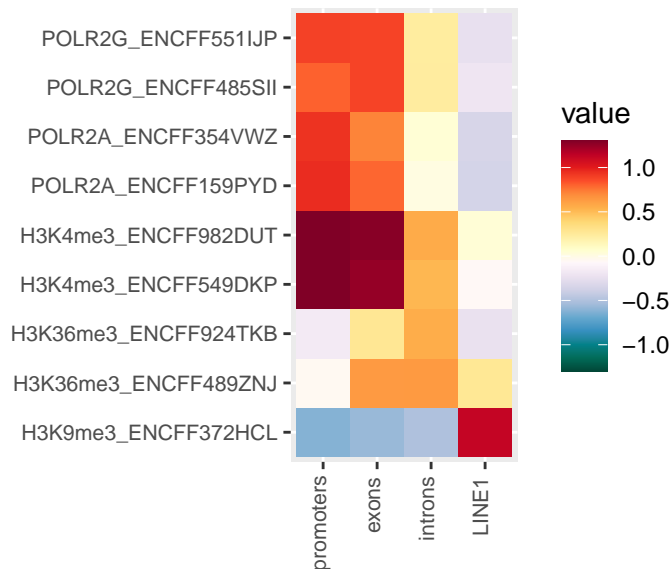


Association with genomic features

Normalized z-score matrix



resampleRegions