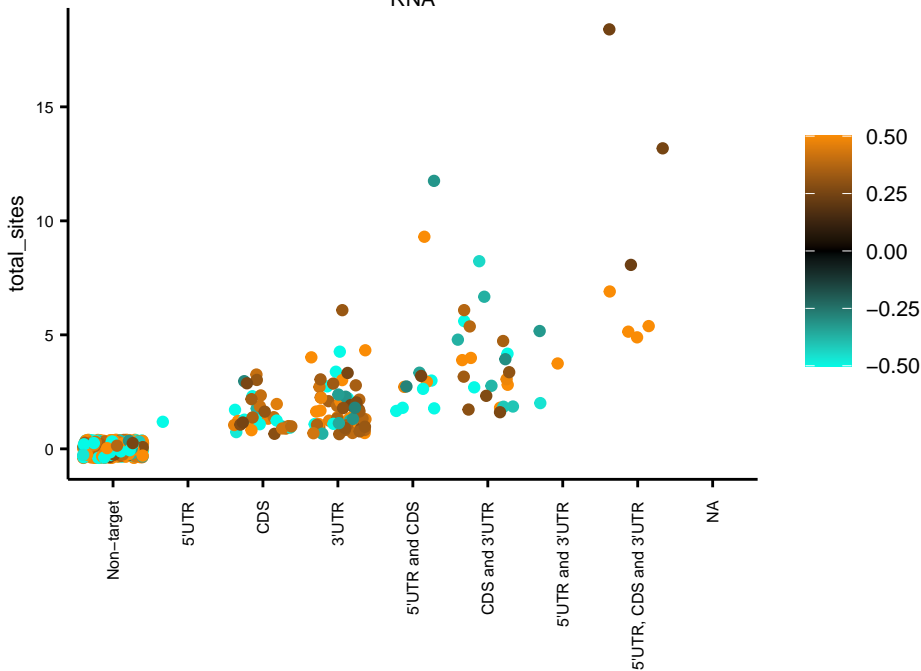


RNA



factor(bsregions, levels = c("Non-target", "5'UTR", "CDS", "3'UTR", ...