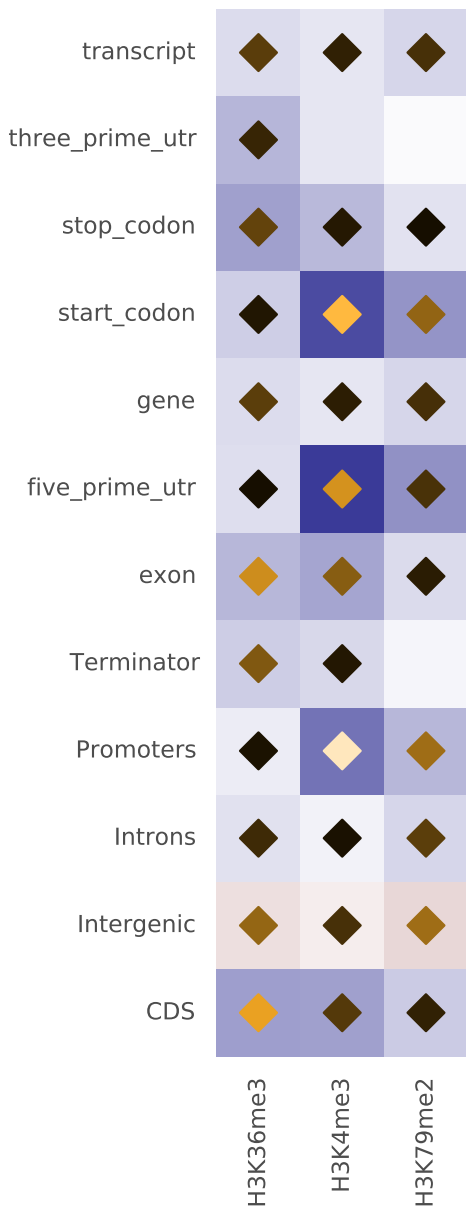
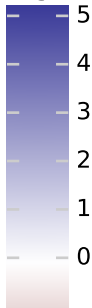


Feature



log2(fold change) for summed bp overlaps



-log10(p-value)

