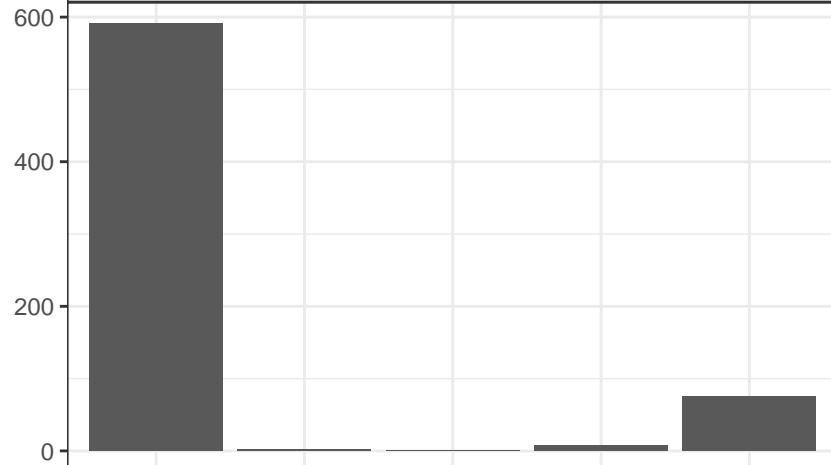


count

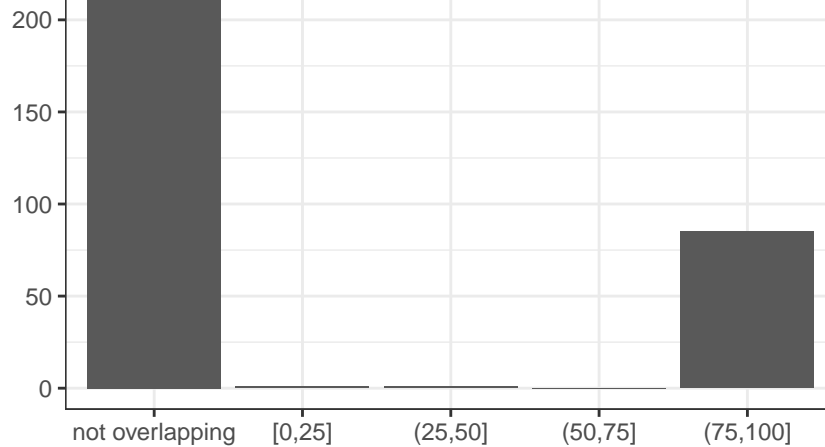
3' UTR



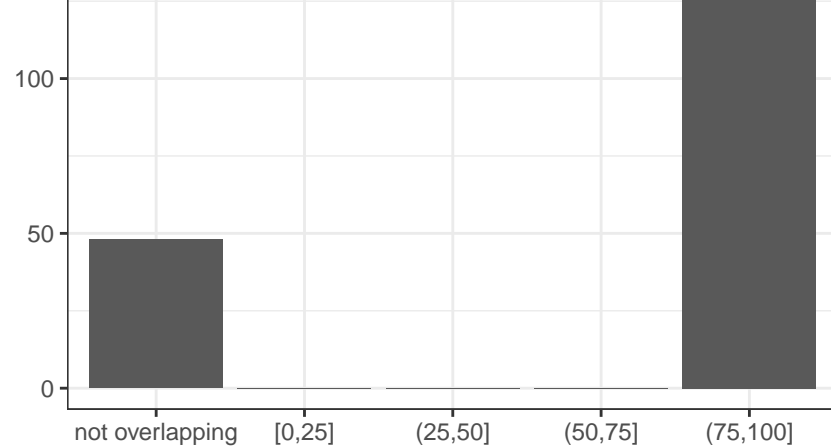
5' UTR



intergenic



internal



overlap with operon relative to prediction length [%]