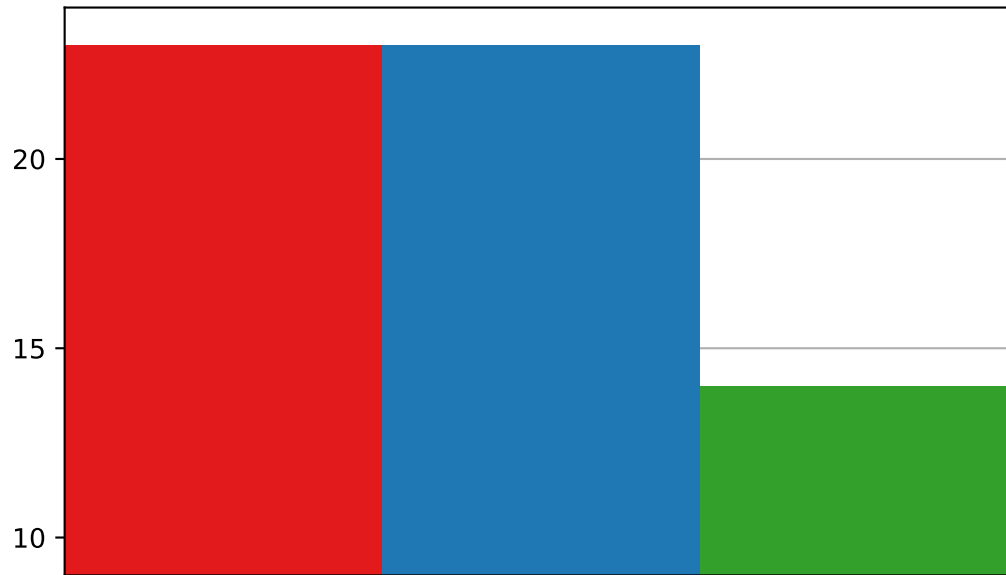


complete genomic features



SAMPLE3_SE.AF0.8.consensus.masked



SAMPLE1_PE.AF0.8.consensus.masked



SAMPLE2_PE.AF0.8.consensus.masked