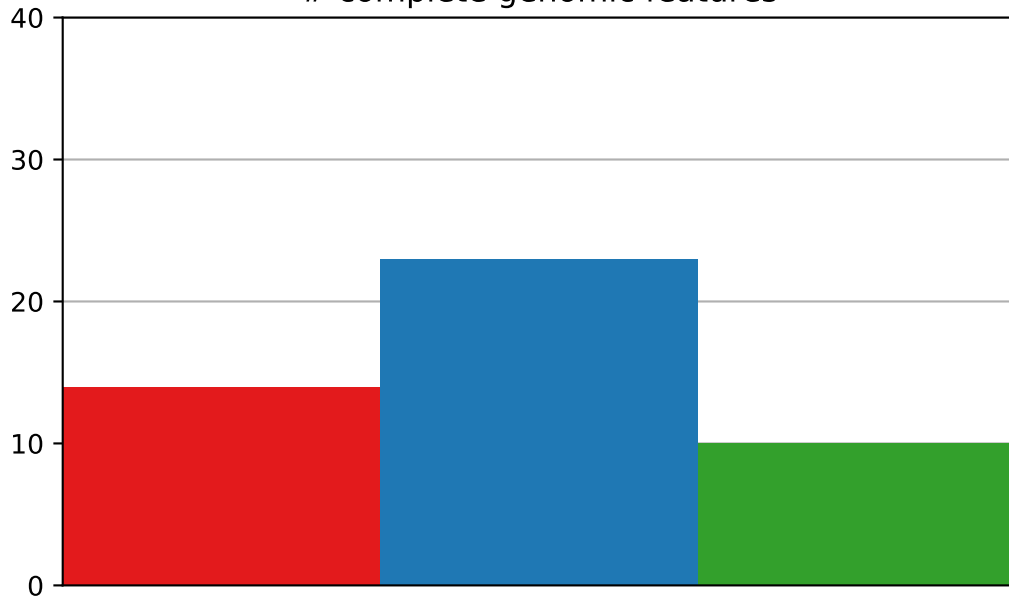


# complete genomic features



SAMPLE3\_SE.AF0.8.consensus



SAMPLE1\_PE.AF0.8.consensus



SAMPLE2\_PE.AF0.8.consensus