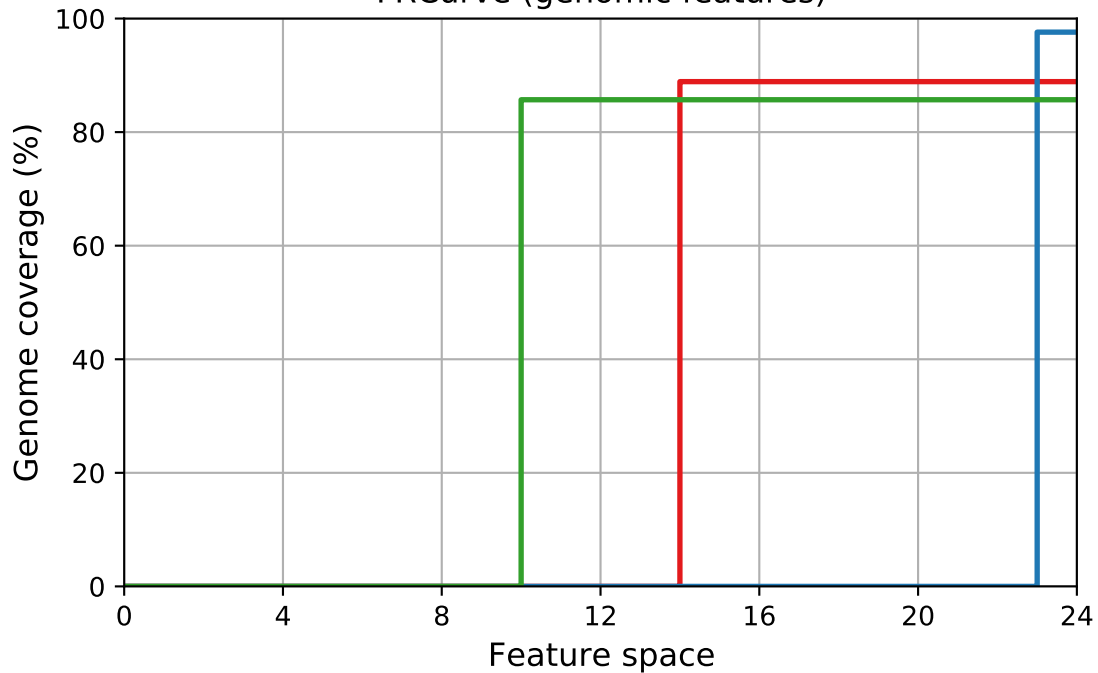


FRCurve (genomic features)



— SAMPLE3\_SE.AF0.75.consensus    — SAMPLE1\_PE.AF0.75.consensus    — SAMPLE2\_PE.AF0.75.consensus