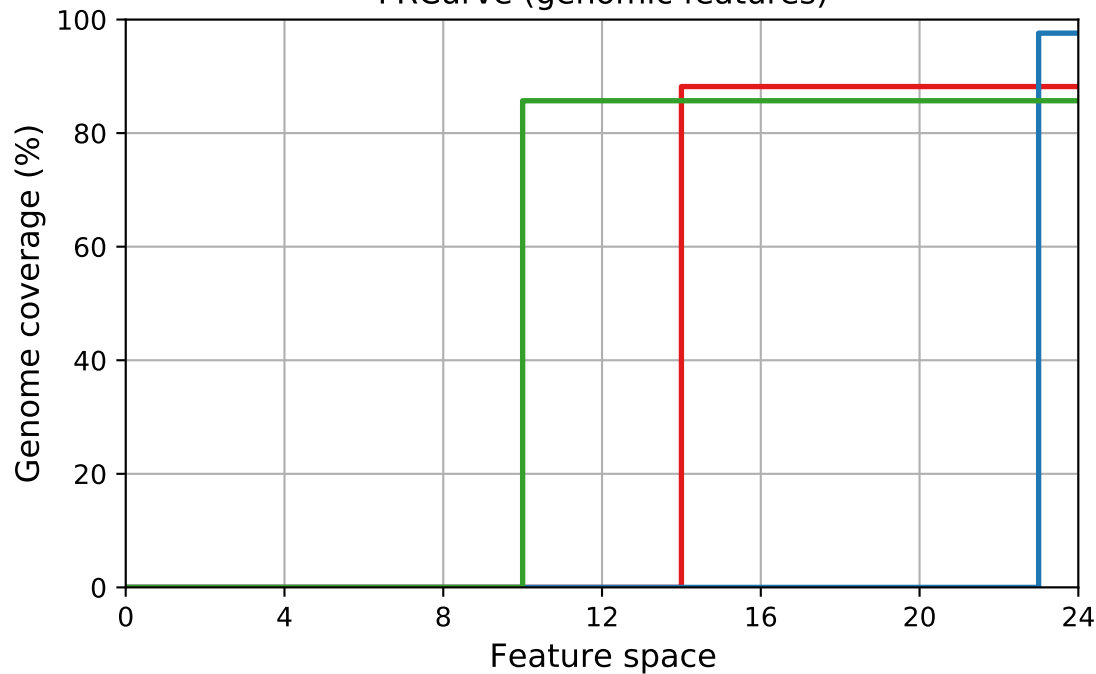


FRCurve (genomic features)



— SAMPLE3_SE.AF0.8.consensus — SAMPLE1_PE.AF0.8.consensus — SAMPLE2_PE.AF0.8.consensus