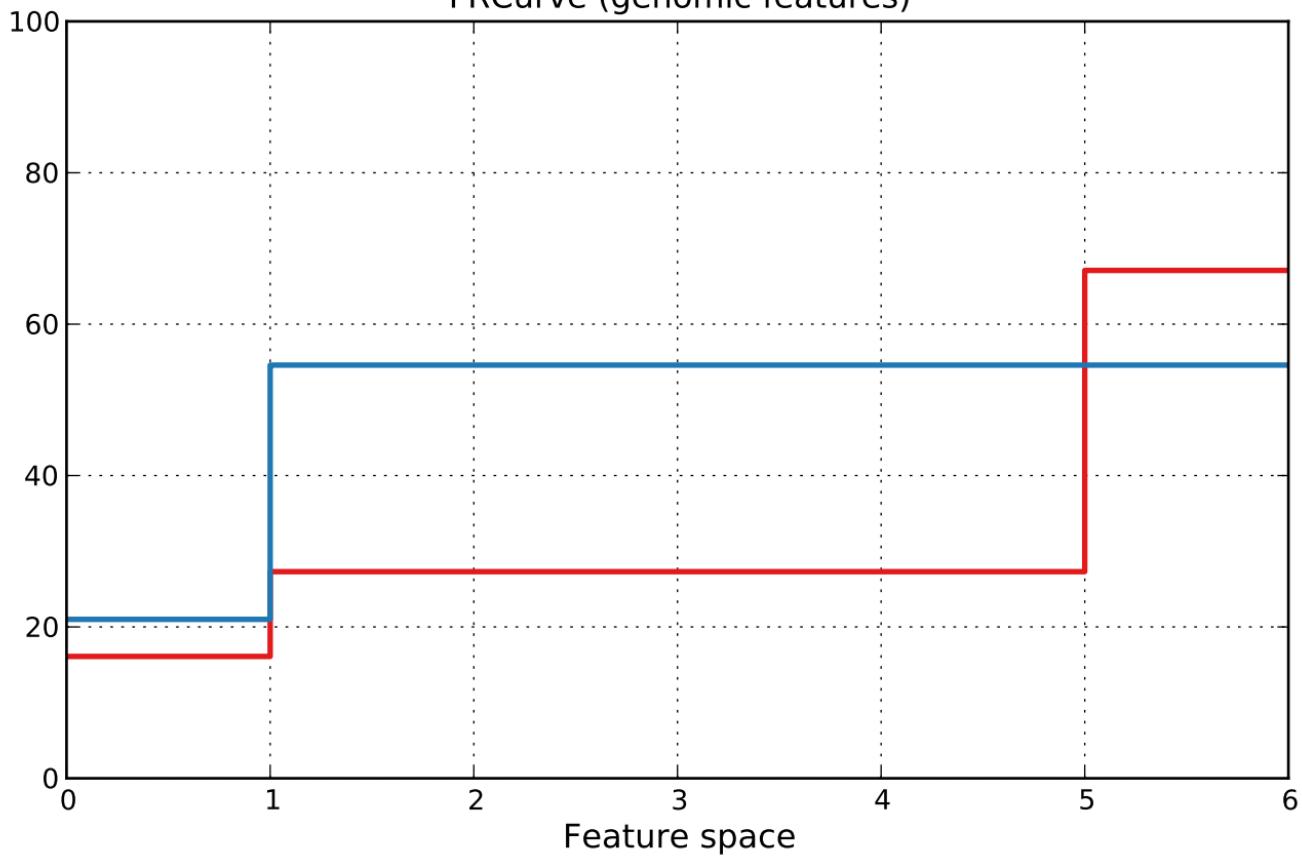


FRCurve (genomic features)

Genome coverage (%)



— contigs_1 — contigs_2