Figure S1: Cross-validation MRC1 Tajima's D Bin Size Lineage 2.2, Genome Coverage

Cross-validation tree of 1,000 MRC1 Tajima's D bin size lineage 2.2 genomes, using a 10-fold cross-validation. Genome coverage of 1,000 MRC1 Tajima's D bin size lineage 2.2 genomes. Tip colors indicate genome coverage, ranging from 100% (red) to 75% (blue).