Yersinia ruckeri 0.98 Shared genome fraction 0.96 0.94 0.92 0.90 0.88 0.86 98.0 98.5 99.0 97.5 99.5 100.0 Average nucleotide identity (%) 3.5 non-parametric KDE Midpoint(s) 3.0 2.5 Density 2.0 1.5 1.0 0.5 0.0 98.0 98.5 99.0 99.5 100.0 97.5 Average nucleotide identity (%)