Pantoea ananatis 0.94 Shared genome fraction 0.92 0.90 88.0 0.86 0.84 0.82 0.80 96.5 97.0 97.5 99.0 98.0 98.5 Average nucleotide identity (%) 3.0 non-parametric KDE Midpoint(s) 2.5 2.0 Density 1.5 1.0 0.5 0.0 99.0 96.5 97.0 98.0 97.5 98.5 Average nucleotide identity (%)