Pseudomonas protegens 1.00 Shared genome fraction 0.98 0.96 0.94 0.92 0.90 0.88 0.86 97.5 98.5 97.0 98.0 99.0 99.5 100.0 Average nucleotide identity (%) non-parametric KDE 4 Midpoint(s) 3 Density 2 1 0 97.0 97.5 99.0 98.0 98.5 99.5 100.0 Average nucleotide identity (%)