Pseudomonas sp 1.00 Shared genome fraction 0.95 0.90 0.85 0.80 97 95 98 99 96 100 Average nucleotide identity (%) non-parametric KDE 1.0 Midpoint(s) 8.0 Density 0.6 0.4 0.2 0.0 97 98 95 96 99 100 Average nucleotide identity (%)