Paenibacillus polymyxa 1.00 Shared genome fraction 0.95 0.90 0.85 0.80 97 96 98 99 95 100 Average nucleotide identity (%) non-parametric KDE 1.4 Midpoint(s) 1.2 1.0 Density 8.0 0.6 0.4 0.2 0.0 96 97 98 99 95 100

Average nucleotide identity (%)