Phaeobacter inhibens 1.000 denome traction 0.950 0.925 0.900 0.875 Shared 0.850 0.825 0.800 97.5 98.0 98.5 99.0 99.5 100.0 Average nucleotide identity (%) non-parametric KDE 2.0 Midpoint(s) 1.5 Density 1.0 0.5 0.0 98.0 98.5 99.5 97.5 99.0 100.0 97.0 Average nucleotide identity (%)