Trueperella pyogenes 1.00 Shared genome fraction 86.0 96.0 06.0 06.0 06.0 0.98 0.96 88.0 99.5 98.0 98.5 99.0 100.0 Average nucleotide identity (%) non-parametric KDE 2.0 Midpoint(s) 1.5 Density 1.0 0.5 0.0 98.0 98.5 99.0 99.5 100.0

Average nucleotide identity (%)