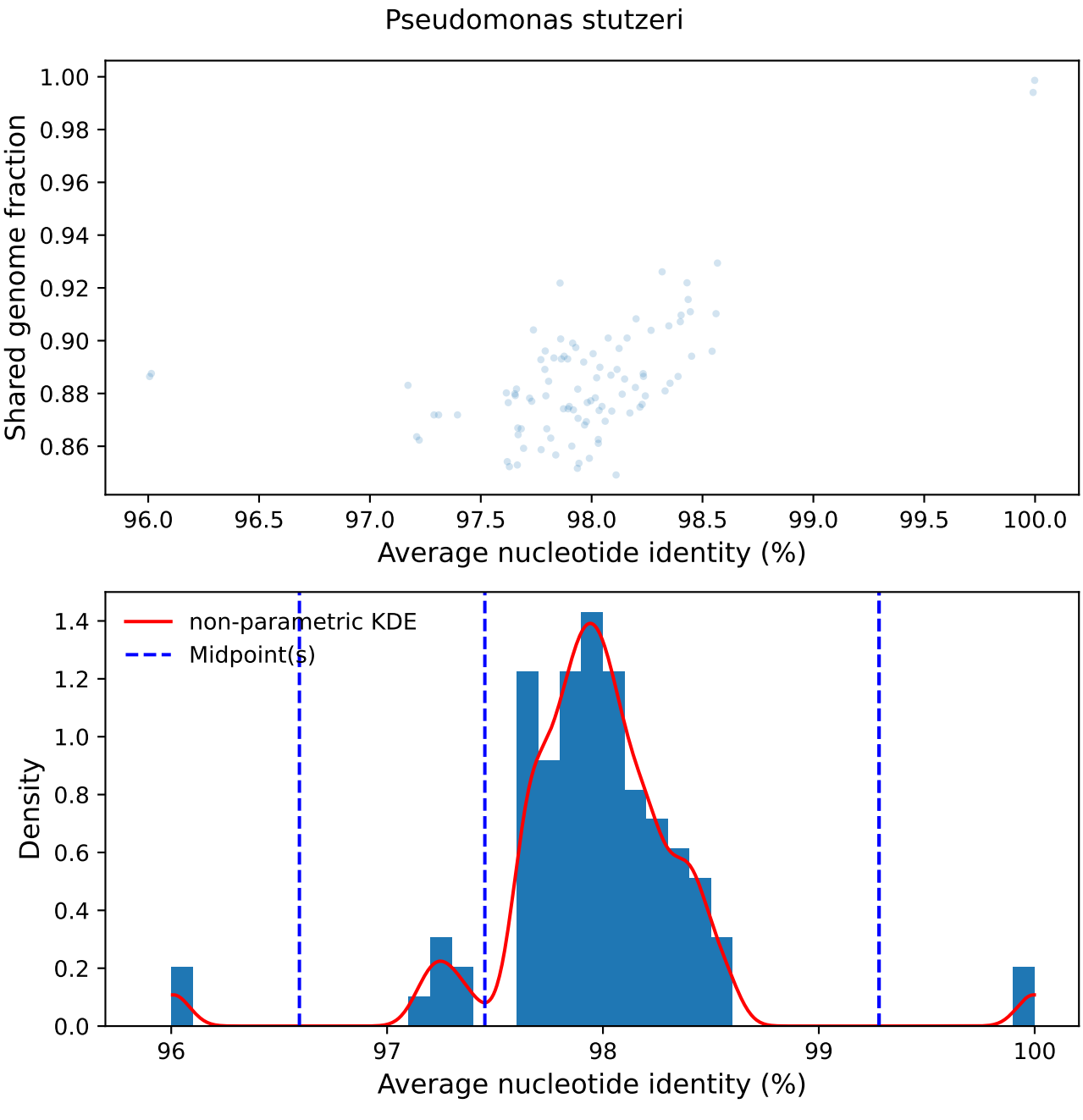
Pseudomonas stutzeri
Shared genome fraction
1.00
0.98
0.96
0.94
0.92
0.90
0.88
0.86
96.0
96.5
97.0
97.5
98.0
98.5
99.0
99.5
100.0
Average nucleotide identity (%)
non-parametric KDE
Midpoint(s)
Density
1.4
1.2
1.0
0.8
0.6
0.4
0.2
0.0
96
97
98
99
100
Average nucleotide identity (%)