| Isolate | NCBI submission name | Sanger sequence assignment | GTDBtk assignment | 16s copy number | Genome size (bp) | Number of contigs | GC content | Total coding sequences | Completeness | Contamination |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 1 | *Achromobacter veterisilvae* AB1 | Achromobacter | *Achromobacter veterisilvae* | 4 | 6,549,044 | 1 | 0.67 | 5,991 | 100.00 | 0.98 |
| 2 | *Ochrobactrum teleogrylli* AB1 | Ochrobactrum | *Ochrobactrum\_B teleogrylli* | 1 | 5,064,753 | 3 | 0.57 | 4,714 | 99.97 | 1.41 |
| 3 | *Pseudomonas fluorescens* AB1 | Pseudomonas | *Pseudomonas\_E fluorescens\_BH* | 2 | 6,759,727 | 1 | 0.60 | 6,053 | 100.00 | 0.15 |
| 4 | *Stenotrophomonas* sp. AB1 (2024) | Variovorax | *Variovorax sp003019815* | 1 | 6,611,336 | 2 | 0.68 | 6,041 | 99.99 | 0.37 |
| 5 | *Variovorax* sp. AB1 (2024) | Stenotrophomonas | *Stenotrophomonas spp.* | 1 | 3,926,254 | 2 | 0.66 | 3,384 | 100.00 | 0.04 |