|  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- |
| **Bacterial species** | **Isolate ID** | **Average coverage X** | **No. of contigs** | **Genome size (Mbps)** | **G+C content (%)** | **No. of CDS1** | **Ref.** | **Accession number** |
| *S. agnetis* | 59 | 9 | 53 | 2.40 | 35.7 | 2353 | This study | MRYT000000003 |
| 43 | 9 | 58 | 2.45 | 35.6 | 2384 | This study | MRYS000000003 |
| 6-4T, 2 | 8 | 58 | 2.48 | 35.6 | 2449 | This study | MRYU000000003 |
| 33 | 9 | 52 | 2.51 | 35.6 | 2457 | This study | MRYV000000003 |
| *S. chromogenes* | 46 | 11 | 29 | 2.21 | 36.7 | 2116 | This study | MRYW000000003 |
| 92 | 21 | 23 | 2.30 | 36.7 | 2231 | This study | MRYX000000003 |
| 101 | 27 | 26 | 2.31 | 36.7 | 2260 | This study | MRYY000000003 |
| 121 | 32 | 20 | 2.27 | 36.6 | 2196 | This study | MRYZ000000003 |
| 117 | 30 | 16 | 2.27 | 36.7 | 2192 | This study | MRZA000000004 |
| 38 | 20 | 34 | 2.34 | 36.8 | 2231 | This study | MRZB000000004 |
| 22 | 26 | 19 | 2.30 | 36.7 | 2221 | This study | MRZC000000003 |
| 72 | 17 | 24 | 2.32 | 36.6 | 2248 | This study | MRZD000000003 |
| *S. simulans* | 102 | 28 | 30 | 2.63 | 35.9 | 2453 | This study | MRZE000000004 |
| 97 | 17 | 34 | 2.70 | 35.8 | 2561 | This study | MRZF000000003 |
| 78 | 24 | 25 | 2.55 | 36.0 | 2368 | This study | MRZG000000004 |
| 113 | 19 | 32 | 2.59 | 36.0 | 2419 | This study | MRZH000000003 |
| 15 | 17 | 26 | 2.70 | 35.9 | 2552 | This study | MRZI000000004 |
| 116 | 19 | 28 | 2.60 | 37.3 | 2404 | This study | MRZJ000000003 |
| 52 | 21 | 31 | 2.63 | 35.9 | 2449 | This study | MRZK000000003 |
| 19 | 22 | 22 | 2.62 | 35.9 | 2451 | This study | MRZL000000004 |
| *S. aureus* | 110 | 28 | 25 | 2.66 | 32.6 | 2547 | (31) | JZIP00000000 |
| 112 | 20 | 29 | 2.65 | 32.6 | 2469 | (31) | JZIQ00000000 |
| 75 | 23 | 26 | 2.65 | 32.6 | 2544 | (31) | JZIO00000000 |
| 9 | 18 | 31 | 2.70 | 32.6 | 2478 | (31) | JZIN00000000 |