|  |  |
| --- | --- |
| **Draft genome** | 11,949 scaffolds with 213,981 gaps; N-content: 80,946,504 bp |
| **Long reads** | 104 Pacific Biosciences RS II reads of insert reported by SSPACE-LongRead to create the scaffolds (min read length 229, max read length 9726, total sequence 281,827 bp) |
| **FGAP output** | 11,949 scaffolds with 124,799 gaps; N-content: 35,400,095 bp |