

## **Draft genome sequence data of *Streptomyces sp. FH025***

### **ABSTRACT**

The genome data of *Streptomyces sp. FH025* comprised of 8,381,474 bp with a high GC content of 72.51%. The genome contains 7035 coding sequences spanning 1261 contigs. *Streptomyces sp. FH025* contains 57 secondary metabolite gene clusters including polyketide synthase, nonribosomal polyketide synthase and other biosynthetic pathways such as amglyccycl, butyrolactone, terpenes, siderophores, lanthipeptide-class-iv, and ladderane. 16S rRNA analysis of *Streptomyces sp. FH025* is similar to the *Streptomyces genus*. This whole genome project has been deposited at NCBI under the accession JAFNG000000000.