

**Table 3.** General features of the *F. acidarmanus* fer1 genome.

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Genome size	1,935,211 bp
Coding region, %	88.8
GC content %	37.56
Predicted ORFs	1963
Functional predictions	1363 (69.4%)
Conserved hypotheticals	301 (15.3%)
Hypothetical protein	299 (15.2%)
ORFs involved in transport	168
Transposase-related	94
rRNAs	1 5S; 1 16S; 1 23S
tRNAs	44

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