

# Supplementary Materials

**Table S1.** Genome statistics.

Attribute	Value
Genome size (bp)	3,343,585
DNA coding (bp)	2,951,289
DNA G + C (bp)	2,223,709
DNA scaffolds	7
Total genes	3,275
Protein coding genes	3,185
RNA genes	58
Pseudo genes	32
Genes in SEED subsystems	1391
CRISPR repeats	0

**Table S2.** Project information.

Property	Term
Finishing quality	Draft
Libraries used	Paired End
Sequencing platforms	Illumina MiSeq
Fold coverage	70x
Assemblers	A5 pipeline, SOAPdenovo v2
Gene calling method	Prodigal
Genbank ID	SPVH00000000.1
Genebank Date of Release	3-April-2019
BIOPROJECT	PRJNA505530

**Table S3.** BUSCO results: The number of single copy conserved orthologs found in the genome.

Parameter	Value
Complete BUSCO	430 (99.5%)
Single-copy BUSCO	429 (99.3%)
Duplicated BUSCO	1 (0.2%)
Fragmented BUSCO	2 (0.5%)
Missing BUSCO	0 (0.0%)
Total BUSCO genes	432 (100.0%)