

Supplemental Table S3. BUSCO analysis results using Firmicutes as the bacterial lineage on all the available *C. cateniformis* genome assemblies

Assembly	UC15921	ADKX01	AKCB01	BBDT01	QSCY01	QSVG01	QSVJ01
Complete BUSCO	221	219	221	183	221	221	221
Complete and single-copy BUSCOs	221	216	221	182	218	218	218
Complete and duplicated BUSCOs	0	3	0	1	3	3	3
Fragmented BUSCOs	0	0	0	29	0	0	0
Missing BUSCOs	11	13	11	20	11	11	11
Total BUSCO groups searched	232	232	232	232	232	232	232
% Complete	95.3%	94.4%	95.3%	78.9%	95.3%	95.3%	95.3%