

Supplementary Table 2. Summary of benchmarking universal single-copy orthologs (BUSCO) analysis (lineage dataset was used as metazoa odb9; and number of BUSCOs was 978)

Type	Count	Ratio (%)
Complete BUSCOs	225	23
Complete and single-copy BUSCOs	223	22.8
Complete and duplicated BUSCOs	2	0.2
Fragmented BUSCOs	462	47.2
Missing BUSCOs	291	29.8