



Figure S2. DNA sequence alignment of *C. gloeosporioides* s.l. mitogenomes at the short *cox3* gene coding region (76-218 bp) where ZH03 (*C. siamense* s.l.) showed strong lineage-specific sequence divergence. In the alignment, amino acid codons are shaded in cyan or yellow. SNP nucleotides in ZH03 are highlighted in red, whereas ones in other isolates are highlighted in blue. ‘*’ marks nonsynonymous codon and the corresponding amino acid changes are shown in the bottom.