

### Legends for Supplementary Figures

Figure S1. Maximum-likelihood phylogenetic tree of *Helvella lacunosa* clade inferred from combined Hsp90 dataset. Bootstrap values  $\geq 70\%$  (left) are indicated at nodes. Asterisk denotes 100% bootstrap.

Figure S2. Maximum-likelihood phylogenetic tree of *Helvella lacunosa* clade inferred from combined ITS dataset. Bootstrap values  $\geq 70\%$  (left) are indicated at nodes. Asterisk denotes 100% bootstrap.

Figure S3. Maximum-likelihood phylogenetic tree of *Helvella lacunosa* clade inferred from combined LSU dataset. Bootstrap values  $\geq 70\%$  (left) are indicated at nodes. Asterisk denotes 100% bootstrap.

Figure S4. Maximum-likelihood phylogenetic tree of *Helvella lacunosa* clade inferred from combined TEF1 dataset. Bootstrap values  $\geq 70\%$  (left) are indicated at nodes. Asterisk denotes 100% bootstrap.