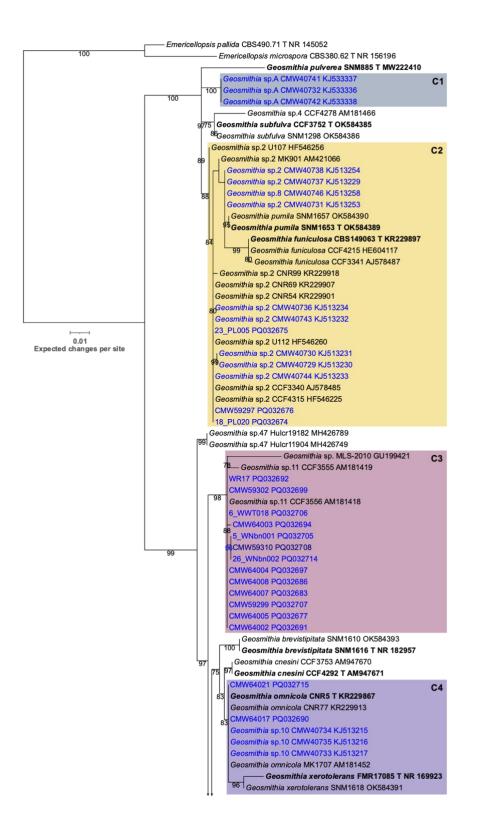
SUPPLEMENTARY FIGURE

Figure S1A – ITS phylogeny



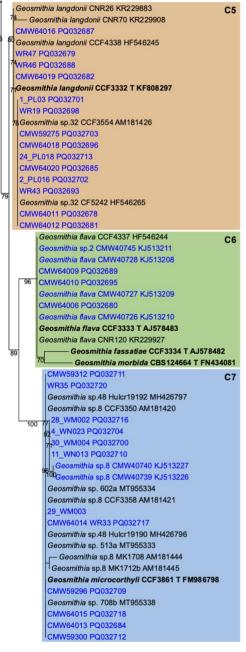


Figure S1. Maximum Likelihood phylogeny of *Geosmithia* (A) ITS, (B) *BT*, (C) *EF1a* exon, (D) *RPB2* and (E) *EF1a* intron sequences, displaying Transfer Bootstrap Expectation (TBE) support values. Clades containing the South African isolates investigated in this study (shown in blue) are highlighted and labelled C1-C7. Isolates of Machingambi et al., (2014) are labelled according to their original identifications. Type strains (denoted with T) of previously described *Geosmithia* species are shown in bold.

Figure S1B – BT phylogeny

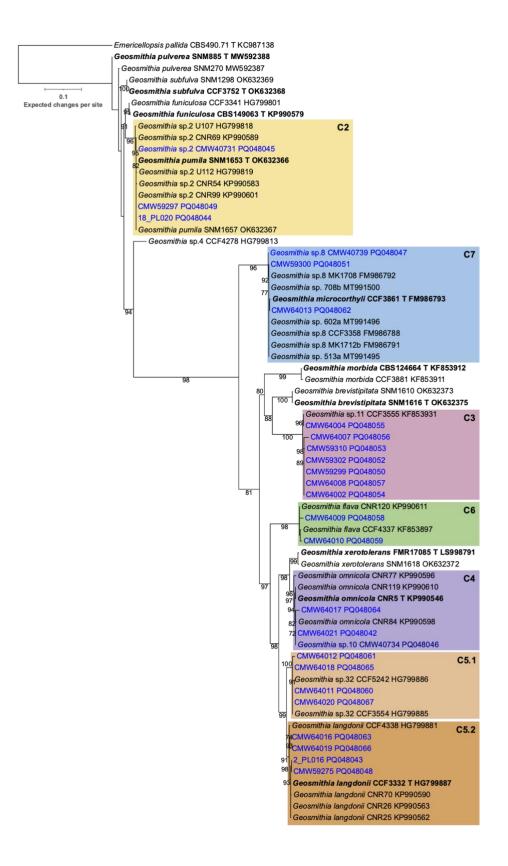


Figure S1C – EF1a exon phylogeny

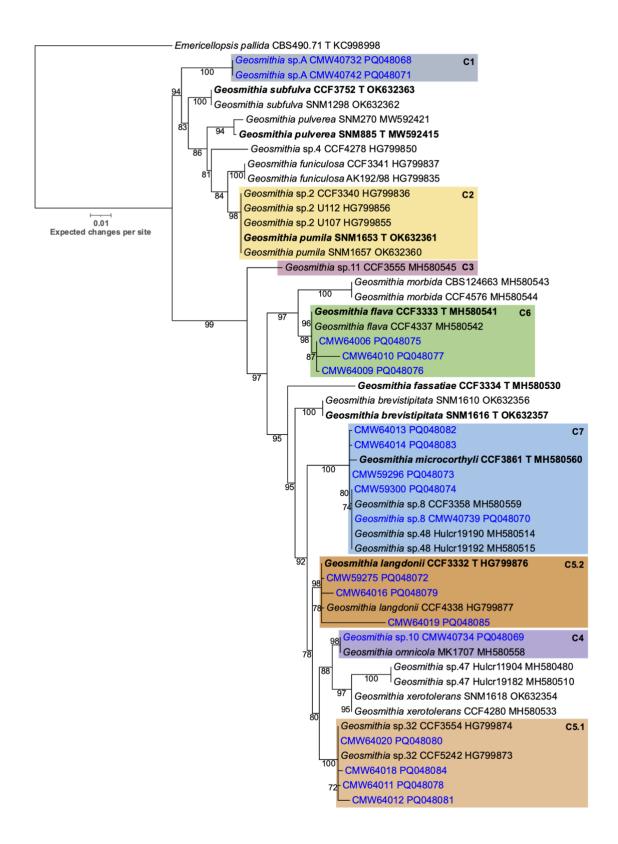


Figure S1D – *RPB2* phylogeny

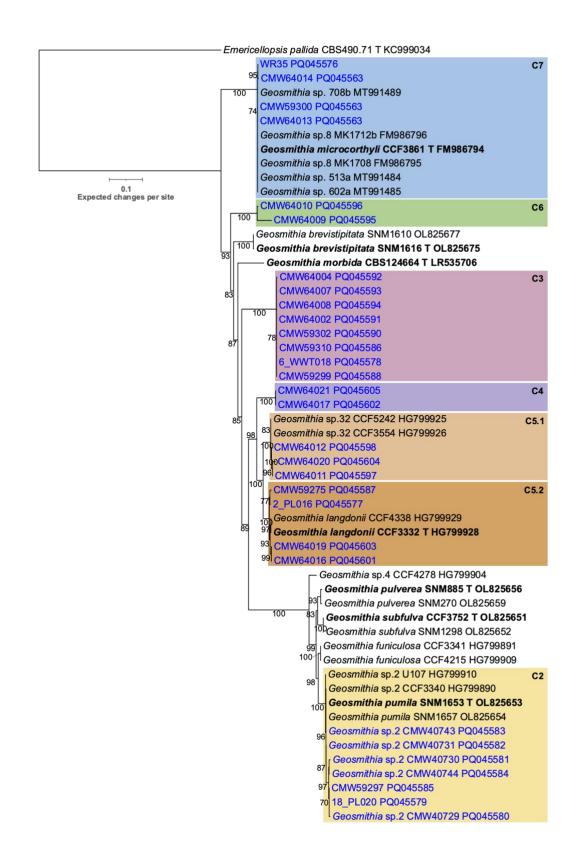


Figure S1E – EF1a intron phylogeny

