

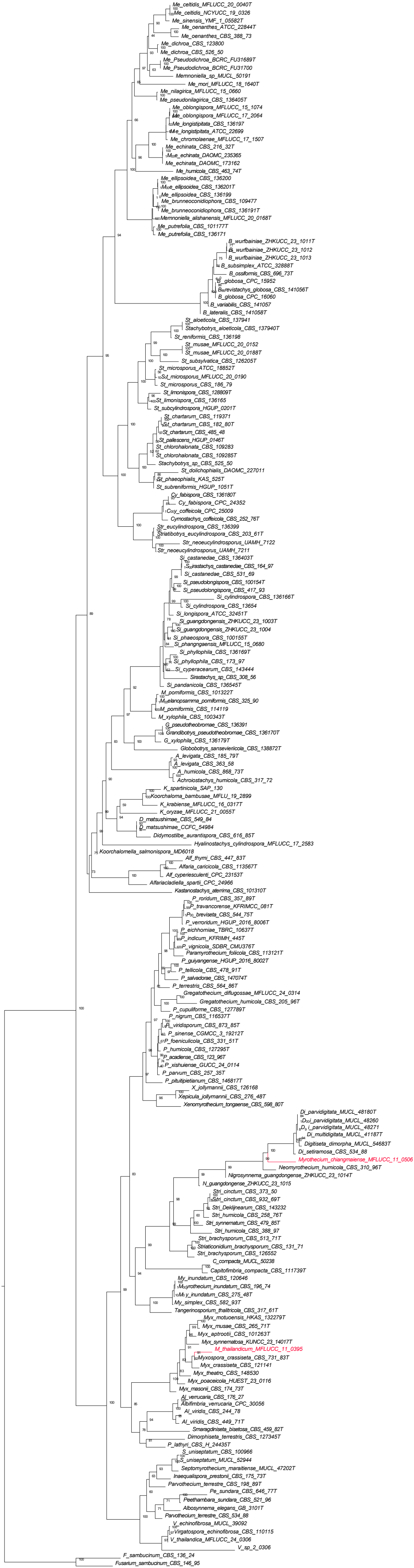


Fig. S1 Maximum likelihood tree of *Myrothecium* and related genera in Stachybotryaceae based on the concatenated ITS, LSU, rpb2, tef, tub2 and cal sequence data. The tree was rooted with *Fusarium sambucinum* (CBS 146.95 and CBS 136.24). ML support values over 70% are indicated. Type strains are in bold and the new combinations are in red. Scale bar indicates the number of substitutions per site.