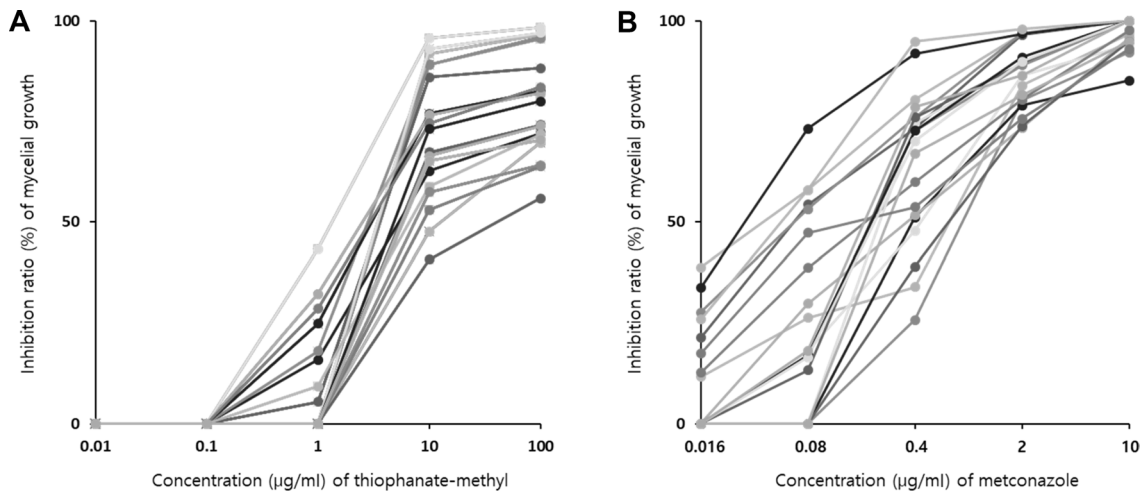


**Supplementary Fig. 1.** Phylogenetic tree based on internal transcribed spacer (ITS) region and translation elongation factor 1- $\alpha$  (*tef1*),  $\beta$ -tubulin (*tub2*) sequences. Sequences were compared using multiple alignment analyses with the Neighbor Joining method in MEGA X. The reference sequences were obtained from National Center for Biotechnology Information GenBank.



**Supplementary Fig. 2.** Inhibition ratio (%) of selected 20 less sensitive and sensitive *Botryosphaeria* spp. isolates against thiophanate-methyl WP (A) and metconazole SC (B).