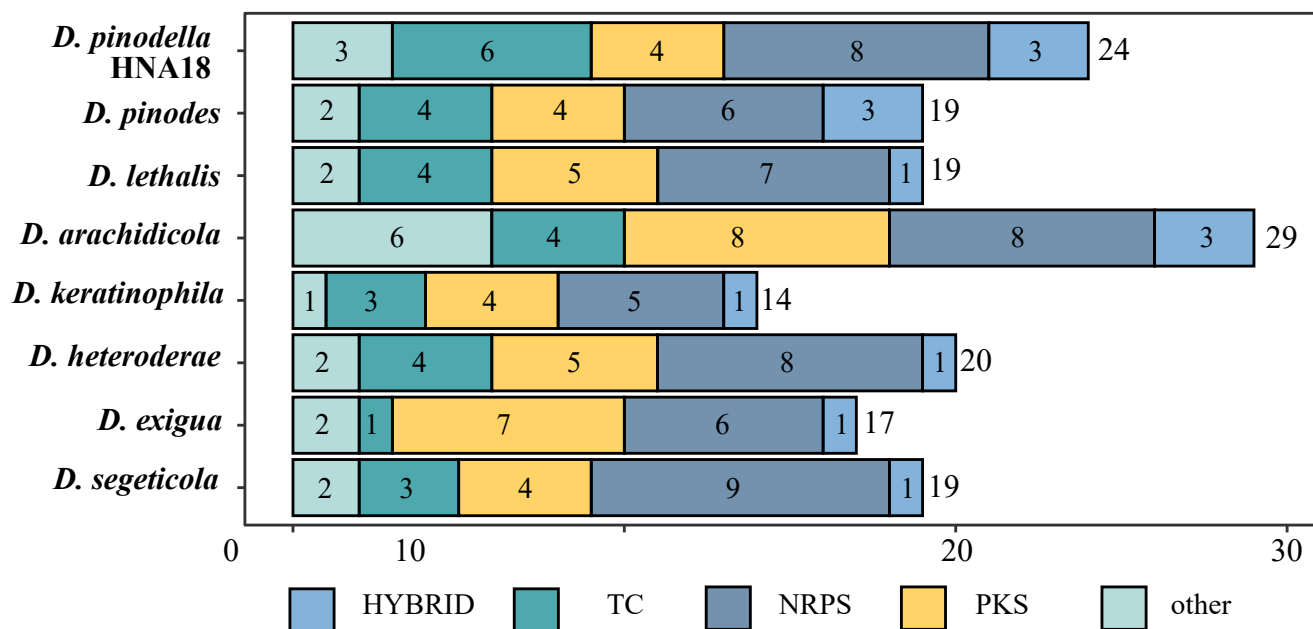




Supplementary Figure 1. The composition of secondary metabolic gene clusters (SMGCs) in eight *Didymella* species. Predicted secondary metabolite genes for each species were divided by the backbone enzyme. NRPS: non-ribosomal peptide synthetase, PKS: polyketide synthase, HYBRID: a backbone gene containing domains from NRPS and PKS backbones, NRPS-like: non-ribosomal peptide synthetase like, TC: terpene cyclase.