Supplementary Figure 3. Phylogenetic sources and abundance of KS-domains from PKS predicted gene models for Zoopagomycota and Mucoromycota species. The heatmap (right) represents the abundances KS-domains predicted for each domain clustered within each clade. Glom.: Glomeromycotina. Mort.: Morteriellomycotina. Ent.: Entomophtoromycotina. Kickx.: Kickxellomycotina. Zoo.: Zoopagomycotina