

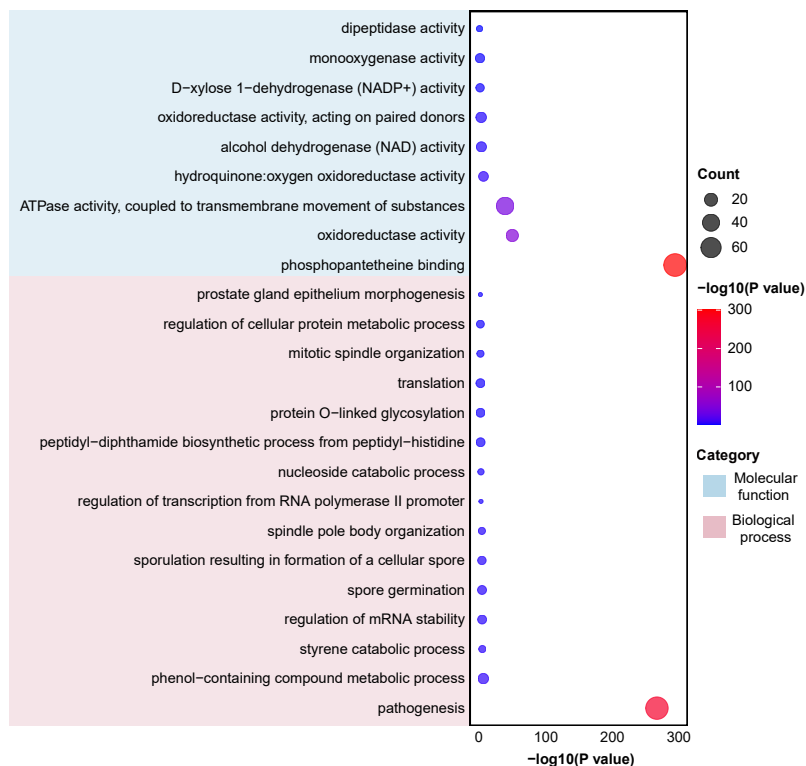
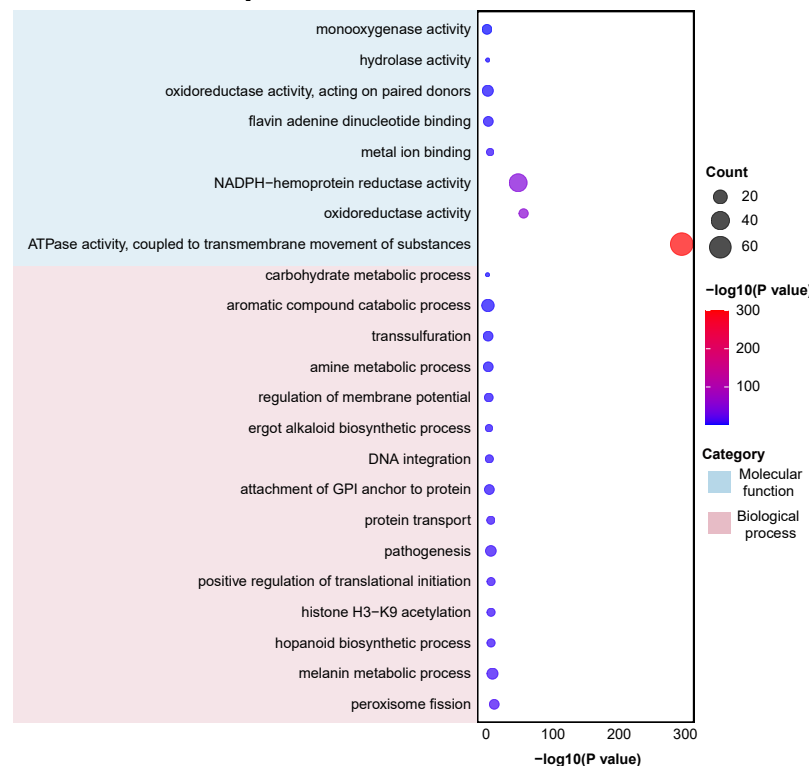
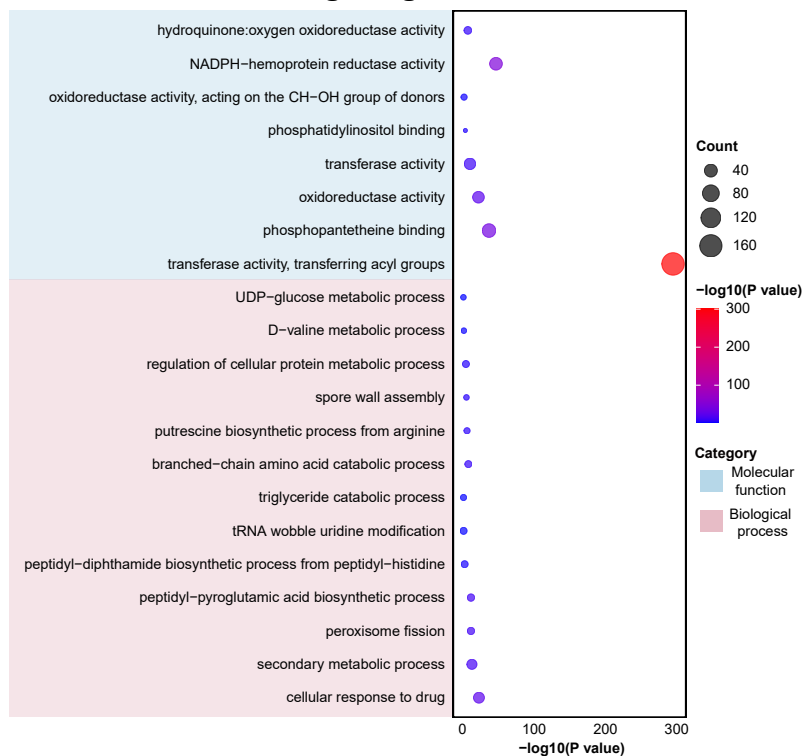
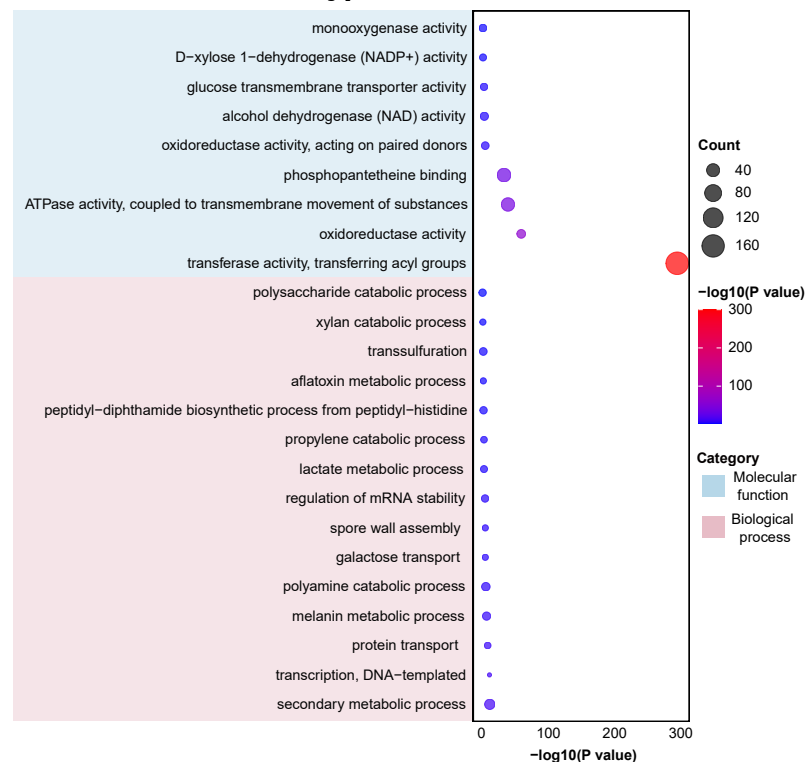
A***C. hawksworthii*****B*****C. pauciramosa*****C*****C. hongkongensis*****D*****C. eucalypti***

Fig. S2. GO enrichment analysis for the species exhibit significant gene family expansions. **(A)** GO enrichment analysis for the 70 expanded gene families in *C. hawksworthii*. **(B)** GO enrichment analysis for the 67 expanded gene families in *C. pauciramosa*. **(C)** GO enrichment analysis for the 65 expanded gene families in *C. hongkongensis*. **(D)** GO enrichment analysis for the 60 expanded gene families in *C. eucalypti*.