

Supplementary Table 18. GO enrichment of unique gene families of the *C. goeringii* genome.

GO_ID	GO_Term	GO_Class	Pvalue	AdjustedPv	x1	x2	n	N	GOlevl
GO:0000956	nuclear-transcribed mRNA catabolic process	BP	1.94E-06	0.002	10	14	4870	34243	8
GO:0000184	nuclear-transcribed mRNA catabolic process, nonsense-mediated decay	BP	8.25E-06	0.009	6	6	4870	34243	9
GO:0044270	cellular nitrogen compound catabolic process	BP	3.92E-05	0.044	16	39	4870	34243	5