

S2 Table. Functional annotation of *Rafflesia cantleyi* transcripts against Nr, Swiss-Prot, TAIR, PFAM, GO, COG and KEGG databases

Database	Number of annotated transcripts	Percentage (%)
Nr	31,444	35.0
Swiss-Prot	21,781	24.3
TAIR	25,970	28.9
PFAM-TMHMM	5,069	5.6
GO	28,922	32.3
COG	7,172	7.9
KEGG	5,850	6.5