

Supplementary Table S3. KEGG pathway annotation of *Dacrydium pectinatum* unigenes.

Pathway	Pathway ID	Number of unigenes	Percentage (%)
Ribosome	ko03010	1244	8.95
Carbon metabolism	ko01200	696	5.00
Biosynthesis of amino acids	ko01230	499	3.59
Plant-pathogen interaction	ko04626	483	3.47
Protein processing in endoplasmic reticulum	ko04141	398	2.86
Spliceosome	ko03040	366	2.63
Oxidative phosphorylation	ko00190	346	2.49
Glycolysis / Gluconeogenesis	ko00010	326	2.34
RNA transport	ko03013	311	2.24
Endocytosis	ko04144	286	2.06
Glyoxylate and dicarboxylate metabolism	ko00630	251	1.80
Phenylpropanoid biosynthesis	ko00940	233	1.68
Starch and sucrose metabolism	ko00500	223	1.60
Plant hormone signal transduction	ko04075	219	1.57
Carbon fixation in photosynthetic organisms	ko00710	216	1.55
Pyruvate metabolism	ko00620	212	1.52
Citrate cycle (TCA cycle)	ko00020	205	1.47
Peroxisome	ko04146	200	1.44
Purine metabolism	ko00230	197	1.42
Amino sugar and nucleotide sugar metabolism	ko00520	189	1.36