

## 附录2 全基因组复制后保留基因的GO功能富集结果

Appendix 2 Enriched GO terms for the retained genes following the whole genome duplication event of *Didymochlaena truncatula*

GO条目	GO term	本体 Ontology	描述 Description	数目 Number	p	错误发现率 FDR
GO:0051246	生物学过程 Biological process		蛋白质代谢过程调控	94*	8.70e-06*	0.02*
			Regulation of protein metabolic process	98#	2.6e-06#	0.0057#
GO:0006417	生物学过程 Biological process		翻译调控	90*	1.90e-05*	0.027*
			Regulation of translation	95#	3.2e-06#	0.0057#
GO:0010608	生物学过程 Biological process		基因表达的转录后调控	93*	1.70e-05*	0.027*
			Posttranscriptional regulation of gene expression	98#	3e-06#	0.0057#
GO:0032268	生物学过程 Biological process		细胞蛋白质代谢过程调控	94*	5.80e-06*	0.02*
			Regulation of cellular protein metabolic process	98#	1.7e-06#	0.0057#
GO:0016192	生物学过程 Biological process		囊泡介导转运	144*	2.70e-07*	0.0019*
			Vesicle-mediated transport	140#	8.5e-06#	0.012#
GO:0045182	分子功能 Molecular function		翻译调控活性	71*	1.90e-10*	5.30e-07*
			Translation regulator activity	183#	2.9e-11#	8.4e-08#
GO:0005515	分子功能 Molecular function		蛋白结合	1,126*	1.90e-09*	2.70e-06*
			Protein binding	1,170#	7.6e-11#	1.1e-07#
GO:0017025#	分子功能 Molecular function		TATA-binding蛋白结合 TATA-binding protein binding#	17#	5e-05#	0.048#

\*第一次全基因组复制特有的功能富集结果; #第二次全基因组复制特有的功能富集结果。

\* Represents the specific functional enrichment results of first whole genome duplication; # represents the specific functional enrichment results of the second whole genome duplication.