表 7 ZIK4 在豆科 3 个物种中的直系同源候选基因

Table 7 The orthologous candidate genes of ZIK4 in three species of Legume

物种	基因名 Gene name	基因号 ID No of gene	是否全长			相似性	EST 证据
Species			If t	he	gene	wasSimilarity	Evidence of EST
			Full-Length				
蒺藜苜蓿	MtZIK4a	AC171498_31	Yes			56.7%	E
Medicago truncatula	MtZIK4b	AC135565_12	Yes			56.9%	E
	MtZIK4c	AC202374_9	Yes			52.4%	E
百脉根	LjZIK4a	LjB12P21_90_nc	Yes			55.3%	E
Lotus japonicas	<i>LjZIK4b</i>	chr5_CM0357_880_nc	Yes			50.6%	E
大豆	GmZIK4a	Glyma20g37180	Yes			61.1%	E
Glycine max	GmZIK4b	Glyma19g43210	Yes			57.6%	F
	GmZIK4c	Glyma03g40550	Yes			43.7%	F
	GmZIK4d	Glyma10g30210	Yes			53.5%	F
	GmZIK4e	Glyma10g39390	Yes			55.8%	E

注: F 代表整个候选基因有 PUT 序列覆盖,或者两端有 PUT 序列覆盖并且整个基因 80%的序列被 PUT 序列覆盖; E 则表示该基因的某个部分或者几个部分有符合要求的 PUT 序列覆盖,但是还没有达到 F 的标准

Note: F: the whole candidate gene was covered by PUT sequence, or both ends were covered by PUT sequence and 80% of the whole gene sequences were covered by PUT sequence; E: a certain part or several parts of the gene were covered by suitable PUT sequence, but had not reached the standard of F