Table 2 The ortholog candidate genes of *CCR2* in three species of Legume

Species	Gene name	ID No of gene	If the gene was Full-Length	Similarity	Evidence of EST
Medicago truncatula	*MtCCR2a*	AC134242_17	Yes	70.3%	F
	MtCCR2b	AC134242_43	Yes	70.6%	E
Lotus japonicas	*LjCRR2a*	chr3 CM0091 630 nd	Yes	64.0%	F
	LjCRR2b	chr1 CM0178 490 nc	Yes	66.8%	F
Glycine max	*GmCRR2a*	Glyma06g01470	Yes	68.6%	F
	GmCRR2b	Glyma11g12510	Yes	52.2%	F
	GmCRR2c	Glyma11g12490	Yes	55.8%	E
	GmCRR2d	Glyma11g12480	Yes	64.2%	F

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