

Table S4. Primers used for EMSA analysis

Gene	Primer	Sequence (5'-3')	Amplified region (+1 as the TSS)
<i>CmNRT</i>			
#1	NRT-110bp-F6	GTGACAACGGAGTGCTGGCG	-510 to -401
	NRT-110bp-R6	TTTTTTAGAAATTGTGGAGTTTGCTAGTTGT	
#2	NRT-110bp-F7	TTCTAAAAAACCTCCAAACGCTTGCAACT	-410 to -301
	NRT-110bp-R7	TCCTAGTTCGCGTAAAAGGCAAACG	
#3	NRT-110bp-F8	CGAACTAGGAAACACGTTTCGCGAT	-310 to -201
	NRT-110bp-R8	ATACGACGCCGCTACGCTCC	
#4	NRT-110bp-F9	GGCGTCGATTTTCGGGATCG	-210 to -101
	NRT-110bp-R9	TGCAACCGAATCGTTGACCGTTGA	
#5	NRT-110bp-F10	TTCGGTTGCAGCGGCGCGCG	-110 to +2
	NRT-110bp-R10	GCCAAACACGAATACAACGTGTGGC	
<i>CmNIR</i>			
#1	SirB-GS-500bp-F2	ACCGTGCTACGACCGCACCG	-506 to -397
	SirB-110bp-R1	TTACCTCTTCTTGATATACTAGCCGC	
#2	SirB-110bp-F2	GAAGAGGTAAGGGTTATGGAACCTTGG	-406 to -297
	SirB-110bp-R2	GGACAAGTCTTGACACATGCCAGTG	
#3	SirB-110bp-F3	AGACTTGTCCACTTAAAACAATTAACATGA	-306 to -197
	SirB-110bp-R3	CCGGCCTCATAGCGGACAC	
#4	SirB-110bp-F4	ATGAGGCCGGATCGGCATCG	-206 to -97
	SirB-110bp-R4	CTGTCCAGTTAAGTCTATAGTTGGCTCG	
#5	SirB-110bp-F5	AACTGGACAGGTCTGCGATGCTC	-106 to +4
	SirB-GS-500bp-R2	AAACGAAGTAAGGCAAAAGCCATCCTGAAT	
<i>CmGS</i>			
#1	GS-110bp-F1	GTAGGTCGATCTGCGTAGTTTGCAG	-510 to -401
	GS-110bp-R1	CTCTGCTTTGCTATATCACTGAGGAC	
#2	GS-110bp-F2	CAAAGCAGAGCTGTTCCCGCG	-410 to -301
	GS-110bp-R2	CGATAGAACGTATGAAGCAATCTGAATAC	
#3	GS-110bp-F3	ATCTGAATACCCGAGATTTTCGTACCTTTC	-310 to -201
	GS-110bp-R3	GACGATGGACACCAGATTGCAG	
#4	GS-110bp-F4	GTCCATCGTCCACAGAAACGTC	-210 to -101
	GS-110bp-R4	TGGACGGCAGCGCACGCT	
#5	GS-110bp-F5	CTGCCGTCCATTCGTCAGCG	-110 to +2
	GS-110bp-R5	AAGCGCTGACACGTCAAGCTCG	