

Table S2: List of primers. AttB sites for GateWay cloning are highlined.

<i>promStIT1</i>	F	<u>GGGGACAAGTTTGTACAAAAAAGCAGGCT</u> GATCTGCCACATAACATAACAAATG	Y1H
<i>promStIT1</i>	R	<u>GGGGACCACTTTGTACAAGAAAGCTGGGT</u> TGCTCATATAAAACTTGGGATCAC	Y1H
<i>promStBEL5 region_1</i>	F	<u>GGGGACAAGTTTGTACAAAAAAGCAGGCT</u> AGATGAAACCAATAAAGATTAGCA	Y1H
<i>promStBEL5 region_1</i>	R	<u>GGGGACCACTTTGTACAAGAAAGCTGGGT</u> CCAAGAAATGAAAGTACACTGC	Y1H
<i>promStBEL5 region_2</i>	F	<u>GGGGACAAGTTTGTACAAAAAAGCAGGCT</u> CATTGTACCATTGCGTCACG	Y1H
<i>promStBEL5 region_2</i>	R	<u>GGGGACCACTTTGTACAAGAAAGCTGGGT</u> ACTTGAAAGATGAAAGGTGCGT	Y1H
<i>StNLP5 cds</i>	F	<u>GGGGACAAGTTTGTACAAAAAAGCAGGCT</u> ATGAATTTCATTTTCCGTTCAAAGCCT	Y1H
<i>StNLP5 cds</i>	R	<u>GGGGACCACTTTGTACAAGAAAGCTGGGT</u> TCATTCACCTGAGCTCTCACAGG	Y1H
<i>StNLP3 cds</i>	F	<u>GGGGACAAGTTTGTACAAAAAAGCAGGCT</u> ATGTCGGAACCGGAAGGAGG	Y1H
<i>StNLP3 cds</i>	R	<u>GGGGACCACTTTGTACAAGAAAGCTGGGT</u> TCATTTACCTGAGCTCTCACAGGAG	Y1H
<i>promStCLE4 region_1</i>	F	<u>GGGGACAAGTTTGTACAAAAAAGCAGGCT</u> TAACACTCCTATAAAGCAAGAACTC	Y1H
<i>promStCLE4 region_1</i>	R	<u>GGGGACCACTTTGTACAAGAAAGCTGGGT</u> TAATTCTTGAATAATTAATGATATATATCTAAGGTT	Y1H and GUS
<i>promStCLE4 region_2</i>	F	<u>GGGGACAAGTTTGTACAAAAAAGCAGGCT</u> GCAGCCGACATCAAATAGAAG	Y1H
<i>promStCLE4 region_2</i>	R	<u>GGGGACCACTTTGTACAAGAAAGCTGGGT</u> GGACCTTAGTTTAGTTAAAATGTG	Y1H
<i>StCLE4 cds</i>	F	<u>GGGGACAAGTTTGTACAAAAAAGCAGGCT</u> ATGGCTAGTGCTTCTAGGTTTTTGT	Overexpr.
<i>StCLE4 cds</i>	R	<u>GGGGACCACTTTGTACAAGAAAGCTGGGTC</u> TAGTGATGTCGAGGATCAGGTCC	Overexpr.
<i>StCLE4 G6T cds</i>	R	<u>GGGGACCACTTTGTACAAGAAAGCTGGGTC</u> CTAGTGATGTCGAGGATCAGGTGTGTCAGGAC	Overexpr.
<i>promStCLE4</i>	F	<u>GGGGACAAGTTTGTACAAAAAAGCAGGCT</u> CTCATAAATCCCTAACATAGGTAAATAAT	GUS