Supplemental Table 1. Primers used to generate the protein fusion constructs.

Name	Primer sequence (5' to 3')	Restriction enzyme
p98EF1	TATGGTCAATGAGCTCATTGTCCTTCGGCA	Sac I
p98ER1	TGTTTTGGACCGGTGAAAAAAAAAACTAAT	Age I
98Fegad	GCGGCCGAATTCCCCGGGATGGAGAATTTCGTCGACG	Eco RI
98Regad	TCCGGTGGATCCAAGCTTGAGTCCATGAACAAAAG	Bam HI
DD2F	GATTACGCCAAGCTTCTCTGTTTCTTATCAGA	Hind III
DD2R	GCTCACCATGGATCCTTTAATATCACTAATACTGC	Bam HI
DD4F	TGATTACGCCAAGCTTTTTCATACTTAAAAATAGTTTGG	Hind III
DD4R	GCTCACCATGGATCCAAAATCAAAATAGTAAAAGCA	Bam HI
DD11F	GATTACGCCTGCAGTTTTCCTTAACGGCGAACGA	Pst I
DD11R	TGCTCACCATGGATCCAGTTTCAAAAGCGAGAGGAGCAC	Bam HI
DD12F	GATTACGCCTGCAGGTGCTTTTTTCCCCATAAAAAACC	Sbf I
DD12R	TGCTCACCATGGATCCCCCACCGCCTCCACCGCC ACCAGAATATGGTGAAAC	Bam HI
DD32F	TGATTACGCCAAGCTTCAGACGACCCATGTAAGAGT	Hind III
DD32R	TGCTCACCATGGATCCATCCCAAGAATAGCAATGA	Bam HI
5941-98F	TATGGTGAATTCACTCATTGTCCTTCGGCA	Eco RI
5941-98R3	GAAGCCATGGTGTTTTGGAAAGGAGAAAAAAAAAACTAAT	Nco I
GFPF1	AGGTCGACTCCAGAGGATCCATGGTGAGCAA	Bam HI
GFPR1	GGAATCTAGAGCTCTTACTTGTACAGCTCGTCCAT	Xba I

Supplemental Table 2. Primers used to generate the 5' and 3' deletions of the *DD11* promoter.

5' endpoint	3' endpoint	Forward primer (5' to 3')	Forward restriction enzyme	Reverse primer (5' to 3')	Reverse restriction enzyme
398	(+21)	GATTACGCCTGCAGTTTTCCTTAACGGCGAACGA	Pst I	GCTCACCATGGATCCTATGAGAATTGCTTTCTCCATTT	Bam HI
352	(+21)	GATTACGCCTGCAGGGAAGAAGAAACAGAGTGGC	Pst I	GCTCACCATGGATCCTATGAGAATTGCTTTCTCCATTT	Bam HI
290	(+21)	GATTACGCCTGCAGAATGGGCTTCAATGGTTTTG	Pst I	GCTCACCATGGATCCTATGAGAATTGCTTTCTCCATTT	Bam HI
234	(+21)	GATTACGCCTGCAGCACAAAAGCCCCTTCTAAGTC	Pst I	GCTCACCATGGATCCTATGAGAATTGCTTTCTCCATTT	Bam HI
174	(+21)	GATTACGCCTGCAGGAAATGTGACTTAATTATTAATC	Pst I	GCTCACCATGGATCCTATGAGAATTGCTTTCTCCATTT	Bam HI
137	(+21)	GATTACGCCTGCAGTGTAAACATGTCAAAA	Pst I	GCTCACCATGGATCCTATGAGAATTGCTTTCTCCATTT	Bam HI
87	(+21)	GATTACGCCTGCAGAAAGCTAGTCGTTACTC	Pst I	GCTCACCATGGATCCTATGAGAATTGCTTTCTCCATTT	Bam HI
352	ATG (-1)	GATTACGCAAGCTTGGAAGAAGAAACAGAGTGGC	Hind III	TTACGCCCTCGAGTTTCTTTTTCTTGTAATGAAGAAG	Xho I
352	- 59	GATTACGCAAGCTTGGAAGAAGAAACAGAGTGGC	Hind III	TTACGCCCTCGAGTTAATTTGTAACGAGTAACG	Xho I
352	- 98	GATTACGCAAGCTTGGAAGAAGAAACAGAGTGGC	Hind III	TTACGCCCTCGAGGTCATTAAAGGATTGTTTGA	Xho I
352	- 139	GATTACGCAAGCTTGGAAGAAGAAACAGAGTGGC	Hind III	TTACGCCCTCGAGATGTTACTTACTTGATTAAT	Xho I
352	- 179	GATTACGCAAGCTTGGAAGAAGAAACAGAGTGGC	Hind III	TTACGCCCTCGAGTATACACATTTTTTCATGTT	Xho I