

SUPPLEMENTAL TABLE 1
Synthetic oligonucleotides used in this study

Restriction sites are underlined. Start codons are in green and stop codons in red. Sequences complementary to the pET28b vector are in blue and the FLAG sequence is in italics.

Co-Expression & BiFC Constructs (5'-3')		
P1	GATCGATCCATATGAGTTGTCCTGCGCCGGAC	
P2	TGATCGTTCGACCTACATAGAGCTCGAGGAAGA	
P3	GATCGATCCATATGAGCTGTCCGGCGACTGAT	
P4	TGATCCTCGAGCTACAGAGAGCTCAAGGTAGGACG	
P5	GATCGATCCATATGGGCTGTCCTTTCCTGAC	
P6	TGATCCTCGAGCTAACAGAGTCCATAGCAGG	
P7	TATACCATGGACTACAAAGACGATGACGACAAGAGCAGCGGCCCTGGTGC	
P8	TGATCCTCGAGGATAACAATCCCCCTCTAG	
P9	ATAAGAATGCGGCCGCTGATCGGATCCCTACATAGAGCTCGAGGAAGA	
P10	TGATCGGATCCGATAACAATCCCCCTCTAG	
P11	ATAAGAATGCGGCCGCTACAGAGAGCTCAAGGTAGGACG	
P12	GATCGATCCATATGGTGAGCAAGGGCGAGG	
P13	TGATCCTCGAGTGATCTCTAGACTAGGCCATGATATAGACGTTGTGG	
P14	GATCGATCCATATGGACAAGCAGAAGAACGGCATC	
P15	TGATCCTCGAGTGATCTCTAGACTACTTGACAGCTCGTCCATGCC	
P16	TATACTGCAGATGGAAACCAATTTTTGATTCC	
P17	TATACTGCAGCTCAGAGAGCTCAAGGTAGGACGC	
P18	TATACCATGGGAAACTACCTTTCGATTTC	
P19	TATACCATGGTCATAGAGCTCGAGGAAGAATGTGG	
P20	TATACCATGGACGACGGTAGAGTTAGGTAGGAATTA	
Cloning of full length cDNAs		
	<i>Forward (5'-3')</i>	<i>Reverse (5'-3')</i>
GGH1	CTAGTCCAGTACTCAGATATAAGTTACG	GTGGTCCAGAACGGATAAGCTTA
GGH2	GCATGTTCAATACTCACAGACC	TCAGAGACTTGTGCTCCAATTC
GGH3	ATGTCTAACTACTTCCTCATTTC	CGATTGCTTGAAGCATGAATG
3' UTR Hybridization Probes		
	<i>Forward (5'-3')</i>	<i>Reverse (5'-3')</i>
GGH1	TTAGTGGATGTTGATCACGGGTT	CACTTGGAAAGCATAAAGGTTTCAG
GGH2	TTAGTGGATGTTGATCACGGGTT	CACTTGGAAAGCATAAAGGTTTCAG
GGH3	CAGAGGAAATGCATATGACCTAAG	GTTCAATTACAACCTCGAAATATCTC