

Supplementary information, Table S1 Primers used in this work

Name	Sequence 5' → 3'	Construct
CBL2GFPfor	<u>tttctagagg</u> atcctgaatgtcgcagtcggtga	CBL2.GFP CBL2.HA
CBL2revGFP	tt <u>cccgggg</u> tatcttcaacctgagaatgg	
CBL2dntermF	<u>ttggatcc</u> aaaatgggggtgtttgatcttgattgtat aaa	CBL2Dn.GFP
GFPCBL2nterm	<u>ttggatcctg</u> aaaaatgtcgcagtcggtgacggt atcaagcatttatgcacctctgtttaagtaaagga gaagaactttcacactttcac	CBL2n16.GFP
GFPCBL2nterm22aa	atgtcgcagtcggtgacggtatcaagcattatgc acctctgtttagggtgtttgatctgatagtaaagg agaagaactttcac	CBL2n22.GFP
CBL2forC4S	<u>ttggatcctg</u> aaaaatgtcgcagtcggtgacg	CBL2C4S.GFP CBL2C4S.HA CBL2nC4S.GFP
CBL2forC12S	<u>ttggatcctg</u> aaaaatgtcgcagtcggtgacggt atcaagcatttaagcacctctg	CBL2C12S.GFP CBL2C12S.HA
CBL2forC18S	<u>ttggatcctg</u> aaaaatgtcgcagtcggtgacggt atcaagcatttatgcacctctgtttaggagtttga tc	CBL2C18S.GFP CBL2C18S.HA
GFPCBL2nterm22 CxxxS	<u>ttggatcctg</u> aaaaatgtcgcagtcggtgacggt atcaagcatttaagcacctctgtttaggagtttg atctgatagtaaaggagaagaactttcac	CBL2nC4,12,18S.GFP
CBL2Q3Efor	<u>ttggatcctg</u> aaaaatgtcggagtcggtgacg	CBL2Q3E.GFP
OFPSpforBamHI	<u>tttggatcc</u> aaaatgaagactaatcttttctttct catctttcactctcctatcattatcctcggccgagtt cgccgcctcctccgaggacg	OFP.TM23
OFPTM23rev	aaaagagctcttaagatcttttctccaacaaat atgcaataagaacaataagaacaagtccagca agagcacctccaactgcaataggaataagagcc tcagcttctctcctcaatagtagaggcgcgggtg gagtgg	
CBL2Prom For Hind	<u>tttaagctt</u> accggtactggattgttcggtat	CBL2 promoter
CBL2Prom Rev Spe	tt <u>tactagt</u> gatatacttaagcagcaaatcaaaac tct	
GFPforXmal	ttt <u>cccggg</u> gatgagtaaaggagaagaactttca ct	GFP (wild type)
GFPrevSacl	aaaagagctcttattgtatagttcatccatgcca	
TPK1_BamF	<u>ttggatcc</u> atgtcagtgatgcagctc	TPK1.GFP
TPK1_XhoR	tt <u>ctcgagc</u> ctttgaatctgagacgtgg	
Actin2 RT for	gtaagagacatcaaggagaagctctc	Actin transcript analysis
Actin2 RT rev	ggagatccacatctgctggaatg	

Restriction sites are underlined;