

Table S2: Primer pairs used in this study.

Primer name Fw	Fw sequence 5' to 3'	Primer name Rv	Rv sequence 5' to 3'
pRS426-CreA-5UTR	GTAACGCCAGGGTTTTCCCAG TCACGACGTCTTTTCTTTTGC CCTTTCG	CreA Rv Luc	ATGTTCTTGGCGTCTCCATGAAC CTCTCAGCCAAGTC
Luc Fw	ATGGAGGACGCCAAGAAC	Luc Rv PyrG	TGCCTCCTCTCAGACAGAATCTAG ACGGCGATCTTGCC
3UTR CreA Fw PyrG	GCATTGTTTGAGGCGAATTCTC CGGCCAAAAAATTTCG	3UTR-CreA- pRS426	GTAACGCCAGGGTTTTCCCAGTC ACGACGCCGCAATACAGAAATAC ACTGG
5UTR ClrA Fw	GTAACGCCAGGGTTTTCCCAG TCACGACGGGTTATAGAGTTG CACCAGG	ClrA Rv Luc	ATGTTCTTGGCGTCTCCATTACT CTGCTTCCCGAGGATCTGAGA
3UTR ClrA PyrG Fw	GCATTGTTTGAGGCGAATTCTT TTGGACGTAGGAATCAACT	3UTR ClrA Rv	GTAACGCCAGGGTTTTCCCAGTC ACGACGCATATGCCGATTCGCTCT T
5UTR ClrB Fw	GTAACGCCAGGGTTTTCCCAG TCACGACGGGCTCAATCTGAT GCAA	ClrB Rv Luc	ATGTTCTTGGCGTCTCCATCGAA TAATATGGAATAGATGAATTAGTG AAAAAAGGCGGTTC
3UTR ClrB PyrG Fw	CATTGTTTGAGGCGAATTCTCA GGCTTATGTGGATACCA	3UTR ClrB Rv	GTAACGCCAGGGTTTTCCCAGTC ACGACGAGAGAAGAAGACTGTGCG GA
CreA fw Alan OH	GAAACAAGGCTCAACACCTGC AAGATGGTAGCGCGA	CreA rv Alan	CAGGTGTTGAGCCTTG
CreA fw Acid OH	TGAGCCGTTCCATTTCGCACTC ACATCGCGTCAAGCGTTC	CreA rv Acid	GAACGCTTGACGCGATGTGAGTG CGAATGGGAACGGCTCA
CreA fw Consv OH	AGGATGAGGATTCTTATGCGA AGCCATTGTCGCCG	CreA rv Consv	CGCATAAGAATCCTCATCC
CreA fw Repr OH	TCACCTATCGCTTACCACAGC ATAAGCGATATCATGTCT	CreA rv Repr	TGTGGTGAAGCGATAGG
Spacer GFP Fw	GGAACACGGGAATGAGTAA AGGAGAAGAACT	GFP Afu Rv	CTCAGACAGAATACGCCAAGCTT GCATGC
GFP PyrG Fw	GCATGCAAGCTTGGCGTATTCT GTCTGAGAGGAGGC	PyrG Rv	GAATTCGCCTCAAACAATGCTCTT CACC
PyrG CreA 3UTR Fw	ATTGTTTGAGGCGAATTCTCCG GCCAAAAAATTTCG	CreA spacer GFP Rv	GGTGACTTGGCTGAGAGGTTTCG GAACACGGGGAATGAGTAAAGG AGAAGAACT
KapC Fw	AGAATGGTTGCTCTATCAGAC	KapC Rv	CCTGAACCTAGCGAGTATCA
KapG Fw	GCAGCAGTGAGGGGA	KapG Rv	TCGCCAATAAATCACGC
KapH Fw	GCTGTCTCAATATGGTCTAGC	KapH Rv	TCATTGTCGGCTGTCATTT
KapL Fw	ATACGGTCTGACTTGCTCT	KapL Rv	GGGTATATGCGTATCGGAGA
KapL Fw	CAACTCAGCCTCTCTCCATA	KapL Rv	TACTTATCTGCTTTCTCGTGC
XlnA ChIP Fw	CAGTAAGTGGTCTAGCGGT	XlnA ChIP Rv	TCAATCAGAGTTTAGGGCAATC
CreA qRT Fw	GAGGATGAGGATTCTTATGCG	CreA qRT Rv	TATCGCTTATGCTTGGGC
XlnA qRT Fw	ATCAACTACGGCGGAAG	XlnA qRT Rv	CAGTAATAGAAGCCGACCC
EglA qRT Fw	CTCACCGATCCCCAAGATA	EglA qRT Rv	CCCAGTTTCCCGTTCTCT
XlnR qRT Fw	AATAGCCTCGCCCTAGC	XlnR qRT Rv	GAGGAATGACTCGGAAGAGA