Table S3. Primers used in this study

Name	Sequence (from 5' to 3')
RT-PCR primers	
SL1	GGTTTAATTACCCAAGTTTGAG
Cbri_SL2*	GGTTTTAACCCAGTTACTCAAG
Cbri_SL3*	GGTTTTAACCCAGTTAACCAAG
Cbri_SL4*	GGTTTTAACCCAGTTTAACCAAG
Cbri_SL10*	GGTTTTAACCCAAGTTAACCAAG
Cbri_SL13*	GGATTTATCCCAGATAACCAAG
Cbri_SL14*	GGTTTTTACCCTGATAACCAAG
*CBG02961*_R	GGTACGACCCTGGCCACGA
*CBG02962*_R	CTGCGCTCAAGGAAATCCTCGAC
*CBG08350*_R**	GGCTTTGCGCTGAGCGATGATG
*CBG08348*_R	GTGAAGAATCCAATGGTTTTTCCCAC
*CBG08349*_R	GGTCCAATGAATCGAAACCGAAATCC
*CBG16611*_R	CTTGTGGCTCCGTGTTCATTCTG
*CBG06074*_R	CCGATCGTCATCAGATGAATCGGA
Sequencing primers	
Crem-*F46B6.6*_L	CAGAAAGTTGTCAACCTGAGAAT
Crem-*F46B6.6*_R	TCACACACTCAATCACACTTTTTC
Cbri-*F37C12.2*_L	GTTCCGAATCCAAGGAAATATG
Cbri-*F37C12.2*_R	ATTTATTCTGATGTTCTGGTTTGC

*: We used mixture of these primers to amplify SL2 *trans*-spliced genes.
**: This primer is used as positive controls for SL1 *trans*-spicing and negative controls for SL2 *trans*-spicing.