	1				2	6
MLYIPGLIAI	IIFYLLILLV	GFWAARKSRS	NPAGTTETED	VMLAGRNIGL	IVGIFTMTAT	-
WVGGGYINGT	AENTFRPGQG	LVWCQAPIGY	AASLVLGGFF	FAKRMRSLAF	VTMLDPFQNK	126
YGERMCGLLF	IPALLGETFW	SAAILSALGA	TLGVIVELDH	TTSVIISACI	ALLYTLFGGL	186
YSVAYTDVVQ	LFCIFVGLWI	6 SLPFALLNKA	TVPISQTWDR	WKGSVEAVDA	/ GIYVDNLLML	240
TFGGIPWQVY	FQRVLSCRTA	KQAQILSFVA 9	8 AIGCLIMALP	SVLIGAVGAS	TDWNMTGYGR	300
AENFTIPDGE	AKLVLPLVLR	YLCPVWVSFV	GLGAVSAAVM	SSSDSSVLSA	ASMFARNVVK 11	360
PVFWQSATER	QIIWVMRASI	FGVGALSCLL	AICIQSIYGL	WYLCSDLIYV	ILFPQLICVL	426
YVRFSNTYGS 13	LIGYILGLFV	RLTSGEALLG	LPALFKYPYF	YDDPQDFYQR	FPCKTFAMLC	486
TLVVTIVVSY	FTDLYFRKDS	KRLKYDFFHC	YKQRASASGS	IHIGNRFNRP	SIRTAKLPIQ	546
VPEVDSENAD	VDENSNKNSN	TLDNNKPLA*				600

Supplementary Fig. S2. Reverse transcription PCR for CsChT segments and overlapping PCR for cDNA preparation. (A) The size of the cDNA segments was expected to be 599, 508, and 684 bp, each overlapping with the adjacent segment for 20 bp. (B) The size of the full-length of cDNA was expected to be 1,710 bp.