Gene ID	Gene Structure	Description	CLE motif
AT1G69970.1		AtCLE26	RKVPRGPDPIHN
AT1G69970.2		AtCLE26	RKVPRGPDPIHN
AT1G70895.1		AtCLE17	RVVHTGPNPLHN
AT1G70895.2		AtCLE17	RVVHTGPNPLHN
AT2G27250.1		AtCLV3	RTVPSGPDPLHH
AT2G27250.2		AtCLV3	RTVPSGPDPLHH
AT2G27250.3		AtCLV3	RTVPSGPDPLHH
AT5G59305.1		AtCLE46	HKHPSGPNPTGN
AT5G59305.2		n.a.	n.a.
GRMZM2G165836_T01		ZmCLE40	REVPTGPDPIHH
GRMZM2G165836_T02		n.a.	n.a.
GRMZM2G474685_T01		ZmCLE26	RRVPNGPDPIHN
GRMZM2G474685_T02		ZmCLE26	RRVPNGPDPIHN
GRMZM5G875999_T01		n.a.	n.a.
GRMZM5G875999_T02		n.a.	RKVPNGPDPIHN

0   500   1000   1500   2000   2500   3000   3500 (bp)

■ CDS   ■ 5'/3'−UTR   — Intron   ■ CLE motif

Figure S10 Gene structure of *CLE* candidates containing alternative splicing in *A. thaliana* and *Z. may* From the left to the right: gene ID, gene structure, gene description and motif sequences of CLE candidates. containing alternative splicing in *A. thaliana* and *Z. may.* Gene structure was depicted using ggplot2-3.2.0. Orange box, CDs; navy blue box, UTR; red box, CLE motif; black line, intron; n.a., information not available.