

Figure 1S

1	MFCLLQ	LARC	DRFAVLIAGS	I	NDFYNYRHQA	II	DIFNMYQQLV	III	KRGFDDQHIT
51	MMAYDDIALS	IV	SENPFGRGKVF	V	HTLKHVNIYP	GSSKINYAHN	VI	SVTADQFYTV	VII
101	LTTLKSTTSD	VIII	NVYIYYDNHG	IX	GPGILGVDPG	VPGGYIEAEP	X	LAKAFDTMEA	
151	KGLYGKLF	FFG	IEACYSGSVA	AVFRAKNMCT	ITAANDESS	YAAVYDSTVG			
201	VYLSNEFSNY	F	MAYLDSNPQ	NTIGNLYTKV	KAQTTGSHVC	YYGDVNMKNL			
251	KLSDFLGTPN	EVVAPKADAK	IDIIPHYLAT	KSTLYQLAQS	TDAKIAGRAK				
301	VALHEVIAAA	EKLDLTLT	TSI	AEILEPETKN	VLRACGKIT	PEYFEVLHYF			
351	TEKYGVVKGD	DMIKLRVLVN	LALKHKVADI	KAIDAIC					

### Supplementary figure legend

Figure 1S. Matched peptides identified in the three protein spots recognized by the anti-TvLEGU-1r antibody in the deduced amino acid sequence of the TvLEGU-1 protein. Boxes in gray show the matched peptides obtained from tryptic digestion and mass spectrometry (Table 1). Consecutive Roman numbers (I-X) were assigned to the identified peptides.