

M. Minutolo, M. Cinque, G. Altamura, F. Di Serio, D. Alioto, B. Navarro (2021). Identification, full-length genome sequencing, and field survey of citrus vein enation virus in Italy. *Phytopathologia Mediterranea* 60(2): 293-301. doi: 10.36253/phyto-12180

ORF2 aa position		83	104	113
CVEV_isolate_14Q	Italy	EEKL	PKGKESSTPC	SERTSAHENG
CVEV_isolate_VE-1	Spain	EEKL	PKGKESSTSC	SERTSAHENG
CVEV_isolate_SM	China	EEKL	PKGKESSTPC	SERTSAHENG
CVEV_isolate_VE823	USA	EEKF	PKGKESSTSC	SERTSAHENG
CVEV_isolate_VE704	USA	EEKF	PKGKESSTSC	SERTSAHENG
CVEV_isolate_VE705	USA	EEKF	PKGKESSTSC	SERTSAHENG
CVEV_isolate_VE706	USA	EEKF	PKGKESSTSC	SEG TSAHENG
CVEV_isolate_VE708	USA	EEKF	PKGKESSTSC	SERTSAHENG
CVEV_isolate_VE707	USA	EEKF	PKGKESSTSC	SERTSAHENG
CVEV_isolate_VE703	USA	EEKF	PKGKESSTSC	SERTSAHENG
CVEV_isolate_VE702	USA	EEKF	PKGKESSTSC	SEG TSAHENG
CVEV_isolate_VE701	USA	EEKF	PKGKESSTSC	SEG TSAHENG
CVEV_isolate_VE709	Australia	EEKF	PKGKESSTSC	SERTSAHENG
CVEV_isolate_STM-1	Japan	EEKF	PKGKESSTSC	SERTSIHENG
CVEV_isolate_STM-2	Japan	EEKF	PKGKESSTSC	SERTSIHENG
CVEV_isolate_IBK	Japan	EEKF	PKGKESSTSC	SERTSIREN
CVEV_isolate_NGS	Japan	EEKF	PKGKESSTSC	SERTSIREN
CVEV_isolate_YM1	Japan	EEKF	PKGKESSTSC	SERTSIREN
CVEV_isolate_PTC	Korea	EEKF	PKGKESSTSC	SERTSTRENG
CVEV_isolate_PCJ	Korea	EEKF	PKGKESSTSC	SERTSIHENG
CVEV_isolate_JJ	Korea	EEKF	PKGKESSTSC	SEKTSIREN

**Supplementary Figure 1.** Multiple alignment of the amino acid (aa) region encoded by ORF2 between positions 80-115. Positions 83, 104 and 113, described by Nakazono-Nagaoka *et al.* (2017) as differential between isolates are marked, with green colour indicating aa considered as conserved in Spanish isolates and grey indicating aa conserved in Japanese isolates. Geographical origins of the isolates are indicated in the middle column.