

Supplementary Table 1. Analysis of genetic diversity on different regions of beet soil-borne virus genome

	Size (nt)	Pi
Genomic segment		
RNA1	5,835	0.02513
RNA2	3,454	0.0279
RNA3	3,005	0.02997
Genomic region		
5'-UTR, RNA1	129	0.03565
Rep, RNA1	5,337	0.02634
3'-UTR, RNA1	394	0.02154
5'-UTR, RNA2	382	0.0135
CP	495	0.01781
CPRT, RNA2	2,796	0.03093
3'-UTR, RNA2	276	0.01649
5'-UTR, RNA3	404	0.03443
TGB1, RNA3	1,281	0.03085
TGB2, RNA3	384	0.02239
TGB3, RNA3	573	0.034
TGB, RNA3	2,057	0.03058
3'-UTR, RNA3	544	0.02449

UTR, untranslated region; CPRT, coat protein read through; TGB, triple gene block.