

Table S2. Oligonucleotide primers coding for the predicted cleavage sites of SARS-CoV-2 Mpro.
After annealing of the complementary oligonucleotide primers, the 5' and 3' ends of the resulting short double-stranded DNAs represented the sticky ends complementary to those formed by PacI and NheI restriction endonucleases, respectively. FWD: forward; REV: reverse. # The sequence of the sequencing primer was published previously: Gazda LD, Joóné Matúz K, Nagy T, Mótyán JA, Tőzsér J. Biochemical characterization of Ty1 retrotransposon protease. *PLoS One*. 2020; 15(1): e0227062.

Protein	Cleavage site	Primer	Oligonucleotide primer sequence
Q14697_GANAB	AVVLQ*TKGSP	FWD	5' -TAAAGCGGTGGTGCTGCAGACCAAAGGCTCTCCAG-3'
		REV	5' - CTAGCTGGAGAGCCTTGGTCTGCAGCACCCAGCGTTAAT-3'
P16949_STMN1	RASGQ*AFELI	FWD	5' -TAAACGTGCGTCTGCCAGCGTTGAAGTATCG-3'
		REV	5' - CTAGCGATCAGTCACGCTGGCCAGACGACGTTAAT-3'
Q9H9B4_SFVN1	EAELQ*AKIQE	FWD	5' -TAAAGAACGGAAACTGCAGCGAAAATCCAGGAAG-3'
		REV	5' - CTAGCTTCTGGATTTGCCCTGCAGTCCGTTTAAT-3'
Q15717_ELAV1	GLRLQ*SKTIK	FWD	5' -TAAAGCCTGCGTCTGCAGTCTAAACCATCAAAG-3'
		REV	5' - CTAGCTTGATGGTTTAGACTGCAGACGCAGGGCTTAAT-3'
P28066_PSA5	ESVTQ*AVSNL	FWD	5' -TAAAGAACATCTGTGACCCAGGCGGTCTAACCTGG-3'
		REV	5' - CTAGCCAGGTTAGACACCGCCTGGGTACAGATTCTTAAT-3'
P01024_CO3	LVSLQ*SGYLF	FWD	5' -TAAACTGGTCTCTGCAGTCTGGCTATCTGTTG-3'
		REV	5' - CTAGCAAAACAGATAGCCAGACTGCAGAGACACCAGTTAAT-3'
P54578_UPB14	SPTLQ*RNALY	FWD	5' -TAAATCTCCAACCCTGCAGCGTAACCGCCTGTATG-3'
		REV	5' - CTAGCATAACAGCGCTTACGCTGCAGGGTTGGAGATTAAAT-3'
P01023_A2MG	RAVDQ*SVLLM	FWD	5' -TAAACGTGCGGTGGATCAGTCTGTGCTGATGG-3'
		REV	5' - CTAGCCATCAGCAGCACAGACTGATCCACCGCACGTTAAT-3'
Q9UBT2_SAE2	GSRLQ*ADDFL	FWD	5' -TAAAGGCTCTCGTCTGCAGGCGATGATTTCTGG-3'
		REV	5' - CTAGCCAGAAAATCATCCGCTGCAGACGAGAGCCTTAAT-3'
O14964_HGS	YAQLQ*AMPA	FWD	5' -TAAATATGCGCAGCTGCAGGGCGATGCCAGCGGG-3'
		REV	5' - CTAGCCGCCGCTGGCATCGCTGCAGCTGCGCATATTTAAT-3'
P22314_UBA1	GSDLQ*EKLGK	FWD	5' -TAAAGGCTCTGATCTGCAGGAAAAACTGGCAAAG-3'
		REV	5' - CTAGCTTGCCCAGTTTCTGCAGATCAGAGCCTTAAT-3'
	ATFLQ*SVQVP	FWD	5' -TAAAGCGACCTTCTGCAGTCTGTGAGGTGCCAG-3'
		REV	5' - CTAGCTGGCACCTGCACAGACTGCAGAAAGGTCGCTTAAT-3'
sequencing primer #		FWD	5' -GATGAAGCCCTGAAAGACGCGCAG-3'