

Supplementary Materials

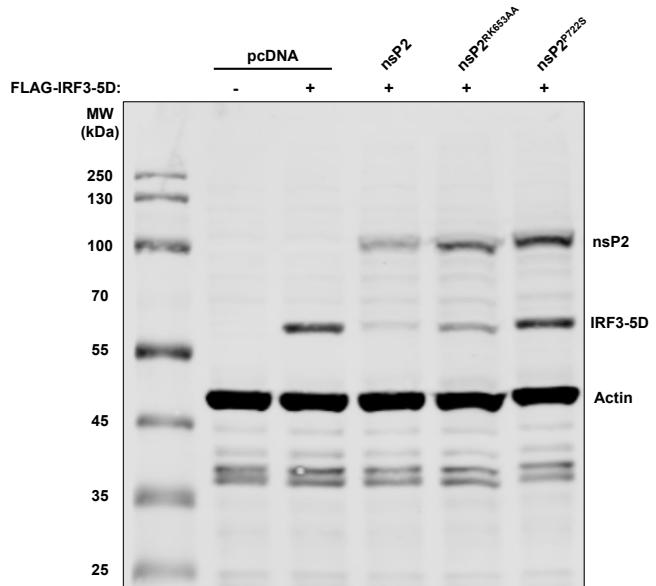


Figure S1. Full blot image from Figure 3E.

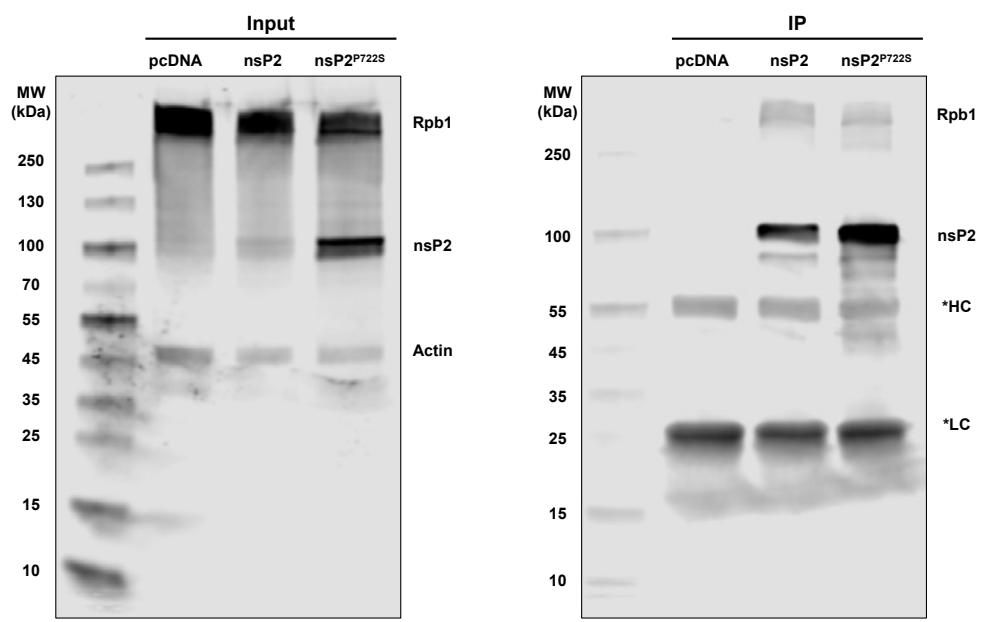


Figure S2. Full blot images from Figure 4B.

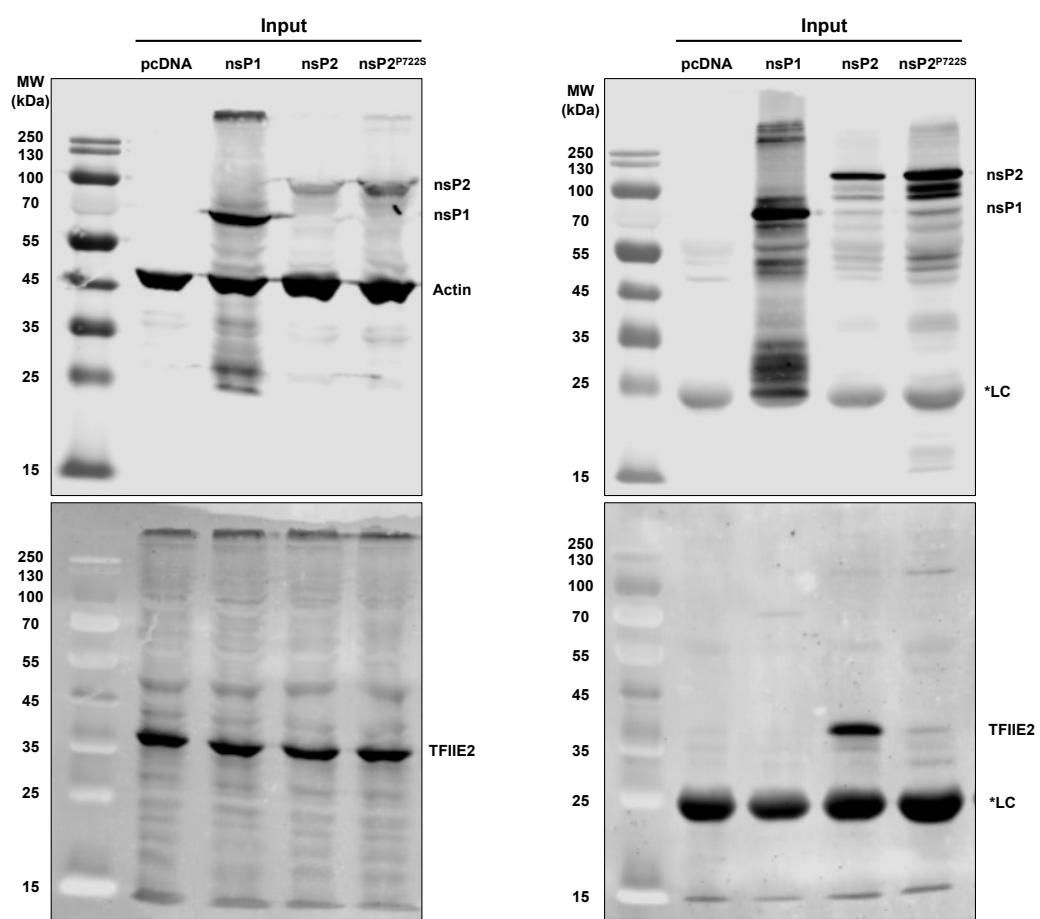


Figure S3. Full blot images from Figure 4C.

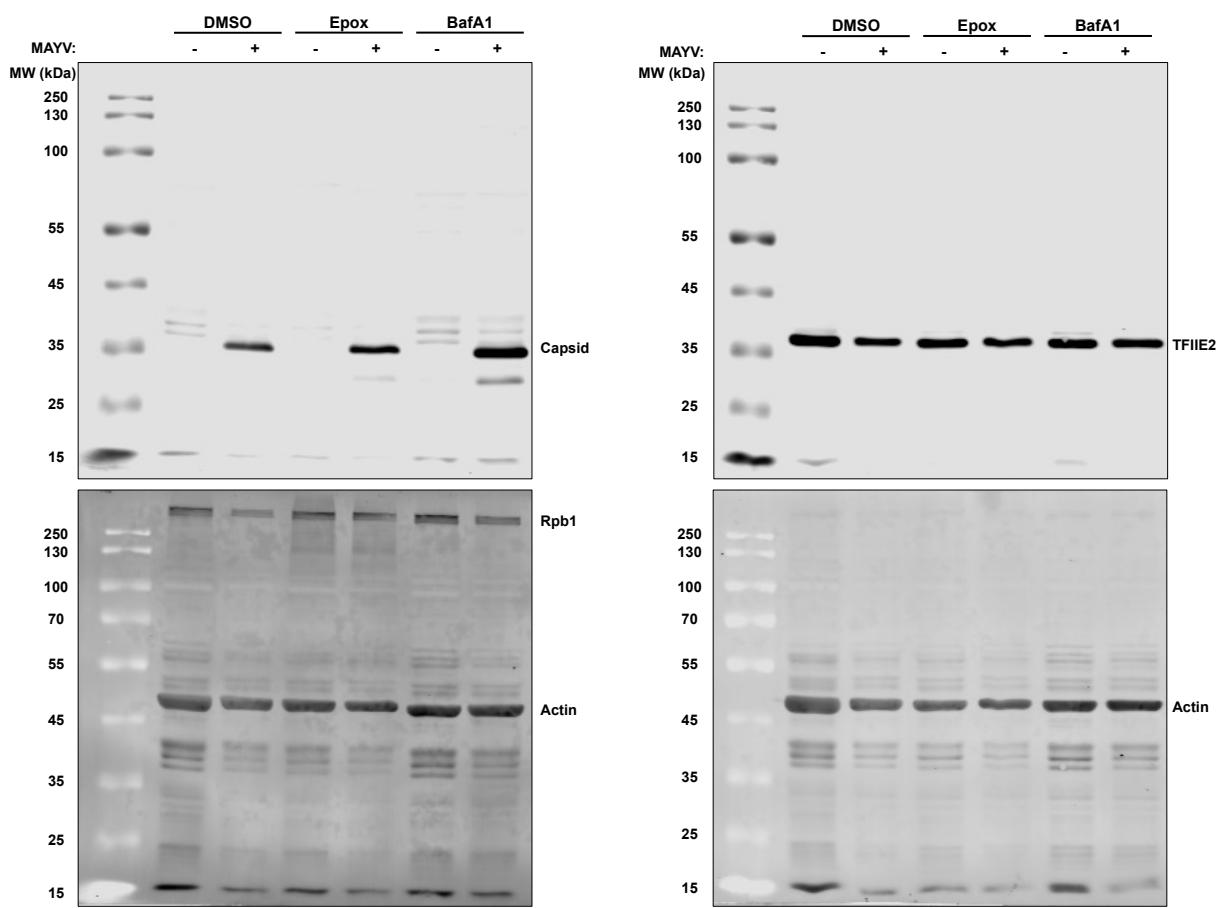


Figure S4. Full blot images from Figure 5E.

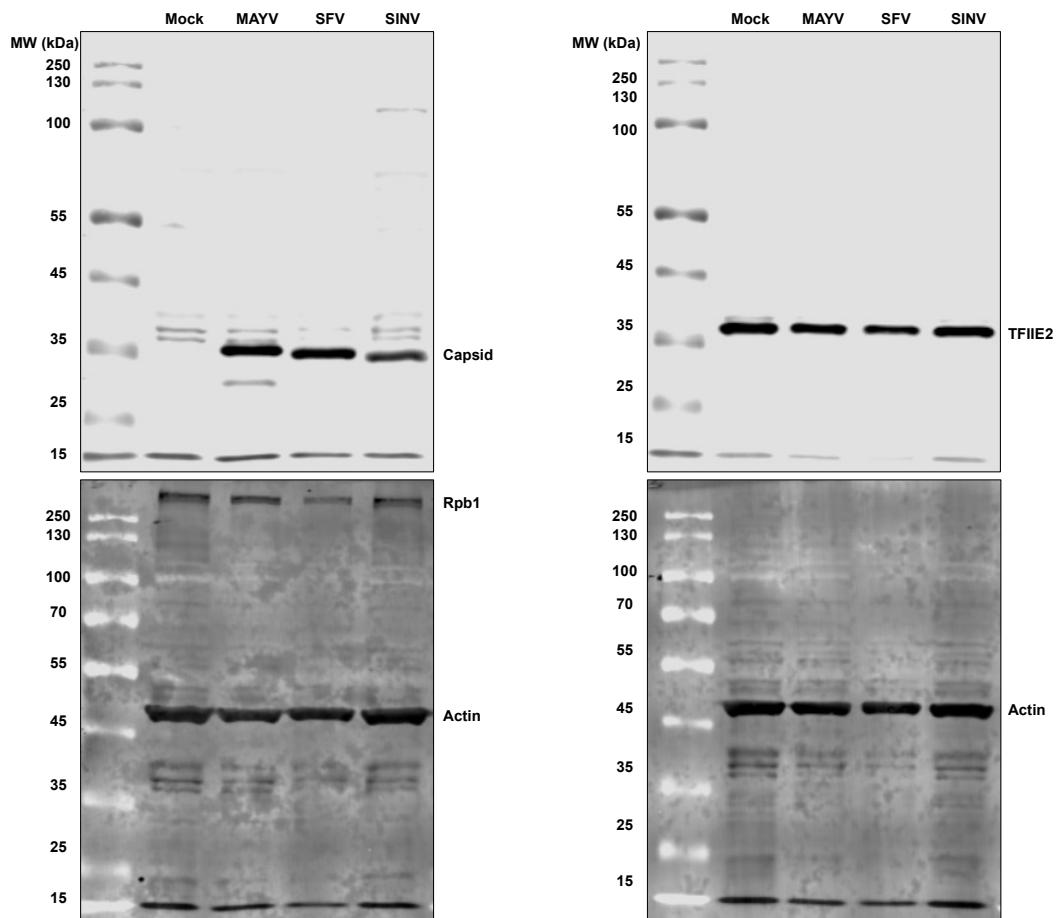


Figure S5. Full blot images of Figure 5H.

Table S1. Primers used for cloning viral proteins.

PCR Target	Primer sequences (5'→3')
nsP1	Fwd: ATATATGCTAGCATGTCGAAAGTCTTGATGATATC Rev: ATATATGGCCGGCCAAACTCCAACCTCCACATCCAC
nsP2	Fwd: ATATATGCTAGCATGCGAGCGGGAGCCGGTGTGAGA Rev: ATATATGGCCGGCCGAACCAGCTGCCGCAGCACTGT
nsP3	Fwd: ATATATGCTAGCATGGCTCCAGTGTATGCCGTTAAAAGG Rev: ATATATGGCCGGCCTCAAGATGAATTACTAATGGTTTC
nsP4	Fwd: ATATATGCTAGCATGCTAGGCCGTGCGGGGGCTATATT Rrev: ATATATGGCCGGCTTAGGACCGCCGTACAGATGTAT
E3-E2	Fwd: ATATATGCTAGCATGGCGGCCTCGACAGTGACAGCTATG Rev: ATATATGGCCGGCCTGCATGTGCTTCGGTGCAGCAACA
C	Fwd: GCTGGCTAGCATGGACTACAAAGACCATGACGGTGATTATAAGATCATGACATCGA CTACAAGGATGACGATGACAAGGGCGGGAGCGGGGGATGGACTCCTACCAACTC AAGTGT Rev: GTTTAAACTTAAGCTTGGATCCTA
6K-E1	Fwd: ATATATGCTAGCATGGCAACAAATGTCTGACACCATATG Rev: ATATATGGCCGGCCCCTCTCAAAGTTATGCAAGTAAC
nsP2 ^{Δprotease}	Fwd: CAAGCTGGCTAGCATGCGAGCGGG Rev: CCGCCGGGCCGGCCACATCCAAAGGAGCTGCTTCCTG
nsP2 ^{Δhelicase}	Fwd: AGCTGGCTAGCATGTTCCAGAATAAAGCTAAAGTGTGC Rev: CGCCGGGCCGGCCGAACCAGCTG
nsP2 ^{K197N}	Fwd: CAAGCTGGCTAGCATGCGAGCGGG Rev: TTCCAAAACGGGTACCAGGCATT Fusion fwd: GTCTGGCGTCCCCGGATCTGAAATTCAAGGTATAATCAAGAGCC TGGTT Fusion rev: AACCAAGGCTTGATTATACTGAATTCCAGATCCGGCACGCCA AGAC
nsP2 ^{C478A}	Fwd: CAAGCTGGCTAGCATGCGAGCGGG Rev: AAAACGGGTACCAGGCATTGGCCAGGCCACTTAGCTTATTCTGGAAC
nsP2 ^{RK653AA}	Fwd: AAATGCCTGGTACCCGTTGGAA Rev: CGCCGGGCCGGCCGAACCAGCTG Fusion fwd: GAGTACAACCTCATTTGCCGAGGGCAGCGGTGACGTGGATTGCTCCGCCACT Fusion rev: AGTCGGCGGAGCAATCCACGTACCGCTGCCCTGGCAAGATGAGGTTGACTC
nsP2 ^{P722S}	Fwd: AAATGCCTGGTACCCGTTGGAA Rev: CGCCGGGCCGGCCGAACCAGCTG Fusion fwd: GCACTTACCTACTGAAATCTGGGGGAAGCCTCCTTGAG Fusion rev: CTCAAAAGGAGGCTCCCCAGATTCAGTAGGTAAAGTGC

Table S2. Primers used for qRT-PCR.

Target Gene	Primer sequences (5'→3')
MAYV nsP1	Fwd: TTCCGAACCAAGTGGGATT Rev: CACTTACGTAYGGKGATGG
SeV HN	Fwd: AAAATTACATGGCTAGGAGGGAAAC Rev: GTGAATGGAATGGTTGTGACTCTTA
ACTB	Fwd: CCTGGCACCCAGCACAAT Rev: GCCGATCCACACGGAGTACT
<i>ifnb</i>	Fwd: TAGCACTGGCTGGAATGAGA Rev: TCCTTGGCCTTCAGGTAATG
<i>ifnl2</i>	Fwd: AGTTCCGGGCCTGTATCCAG Rev: GAGCCGGTACAGCCAATGGT
GAPDH	Fwd: ACAGTCAGCCGCATCTTCTT Rev: GTTAAAAGCAGCCCTGGTGA
<i>Tuba1a</i>	Fwd: GCAACAACCTCTCCTCTTCG Rev: GAATCATCTCCTCCCCAAT

Table S3. MiST analysis-based scoring of nsP2 interacting cellular proteins.

Table S4. GO enrichment analysis of nsP2 interacting cellular proteins.