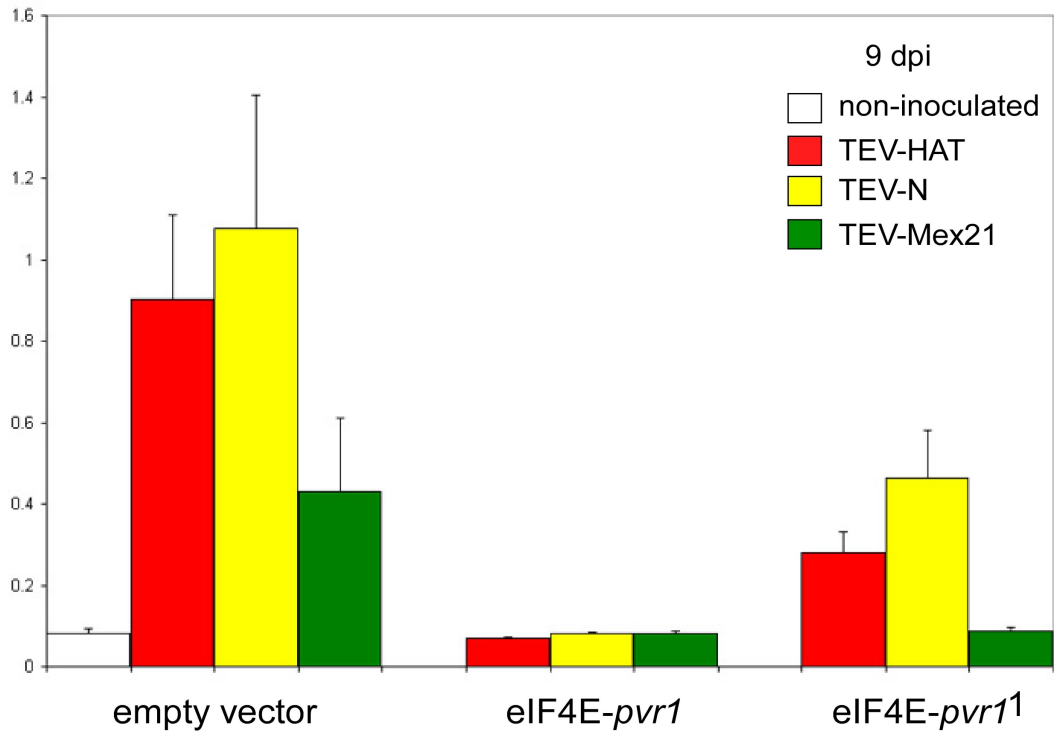


### Supplemental Figure 1

Confocal microscope images from bimolecular fluorescence complementation assay (BiFC) using *Agrobacterium*-mediated transient expression assay in *N. benthamiana*.

(A) eIF4E proteins fused with YN; YN:EE:eIF4E-*Pvr1*<sup>+</sup>, YN:EE:eIF4E-*pvr1* YN:EE:eIF4E-*pvr1*<sup>1</sup> YN:EE:eIF4E-*pvr1*<sup>2</sup>, YN:EE:eIF4E-G107R, YN:EE:eIF4E-*pvr1*<sup>1</sup>+*pvr1*<sup>2</sup>, were transiently expressed in *N. benthamiana* leaf tissue with and without VPg protein fused with YC; YN:HA:VPg (TEV-HAT). Yellow fluorescent signal generated by the protein-protein interaction were detected in mesophyll cells 60 hours post-infiltration. Chloroplast autofluorescence is shown in red. Scale bar = 10 micron.

(B) Immunoblot image from coimmunoprecipitation assays. Total protein extracts were pulled down with anti-HA agarose beads and immunoblotted with an antibody for *Capsicum*-eIF4E.



Supplemental Figure 2

Accumulation of TEV coat protein determined by ELISA for transgenic tomatoes containing empty vector, eIF4E-*pvr1*, or eIF4E-*pvr1*<sup>1</sup>.

ELISA was performed for T2 plants containing empty vector, eIF4E-*pvr1*, or eIF4E-*pvr1*<sup>1</sup>, 9 days after inoculating with TEV-HAT, TEV-N and TEV-Mex21. Accumulation of TEV coat protein determined by ELISA of tissue sampled from upper un-inoculated leaves at 9 dpi.

**Supplemental Table 1.** Primer sequences used for site-directed mutagenesis.

Primer name	Sequence (5' → 3')
T51Af	caa aga aat agc <b>agc</b> aaa gca tcc at
T51Ar	at gga tgc ttt gct <b>gct</b> att tct ttg
P66Tf	ctg gtt tga taa <b>ta</b> c agt ggc gaa atc
P66Tr	gat ttc gcc act <b>gta</b> tta tca aac cag
G107Rf	gca agt tag ttg <b>tga</b> gag cag act tac attg
G107Rr	caa tgt aag tct gct <b>ctc</b> aca act aac ttgc
V67Ef	ctg gtt tga taa tcc <b>aga</b> ggc gaa atc gaaac
V67Er	gtt tgc att tgc <b>cct</b> ctg gat tat caa accag
L79Rf	ggg tag ctc <b>gcg</b> tgc caa cgt cta cac
L79Rr	gtg tag acg ttg cga <b>cgc</b> gag cta ccc
D109Nf	gtt agt tgt ggg agc <b>aaa</b> ctt aca ttg
D109Nr	caa tgt aac <b>ttt</b> gct ccc aca act aac
G107R_D109Nf	gtt agt tgt <b>gag</b> agc <b>aaa</b> ctt aca ttg
G107R_D109Nr	caa tgt aac <b>ttt</b> gct <b>ctc</b> aca act aac
P66T_L79Rf	ggt ttg ata <b>ata</b> cag <b>agg</b> cga aat cg
P66T_L79Rr	cga ttt cgc <b>ctc</b> tgt <b>ttt</b> atc aaa cc

**Supplemental Table 2.** Primers sequences used for various plasmid construction.

Primer name	Primer sequence (5' → 3')	Cloning vector
VPgMex-xr	ccctcgagctattcaaacatcaactcct	pEG202/pJG4-5
VPgNW-xr	ccctcgagctattcaaacgtcaactcct	pEG202/pJG4-5
eIF4E-sf	ccgagctcatggcaacagctgaaatgg	pSY735/pSY736
eIF4E-br	tccggatccctatacgggtgaacg	pSY735/pSY736
VPgHAT-sf	gagctcatggggaagaagaatcagaa	pSY735/pSY736
VPgMex-sf	gagctcatggggaagaagaatcag aa	pSY735/pSY736
VPgMex-bsr	ggatccactattcaaacatcaactcct	pSY735/pSY736
VPgNNW-bsr	ggatccactattcaaacgtcaactcct	pSY735/pSY736
VPgHAT-bsr	ggatccactattcaaacgtcaactcct	pSY735/pSY736
VPgHAT-BamHIeraserF	cattaggtttgtggaccattgacaggtcac	pSY735/pSY736
VPgHAT-BamHIeraserR	gtgacctgtcaatgggtccacaaacctaatg	pSY735/pSY736
VPgHAT-HindIIIeraserF	cagaagcacaagctgaagatgagagaggcg	pCAMBIA
VPgHAT-HindIIIeraserR	cgcctctctcatcttcagcttgctctctg	pCAMBIA
eIF4E_SmaI F	tccccgggatggcaacagctgaaatgg	pBI121
eIF4E_SacIR	tccgagctctatacgggtgaacg	pBI121