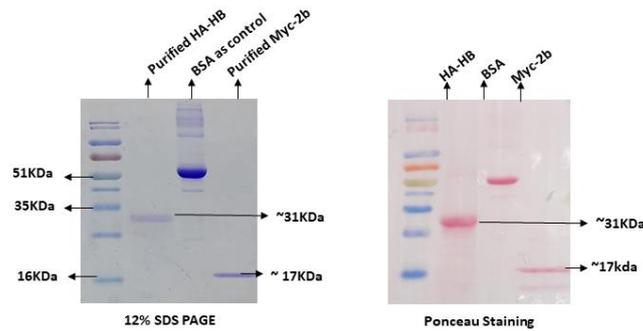
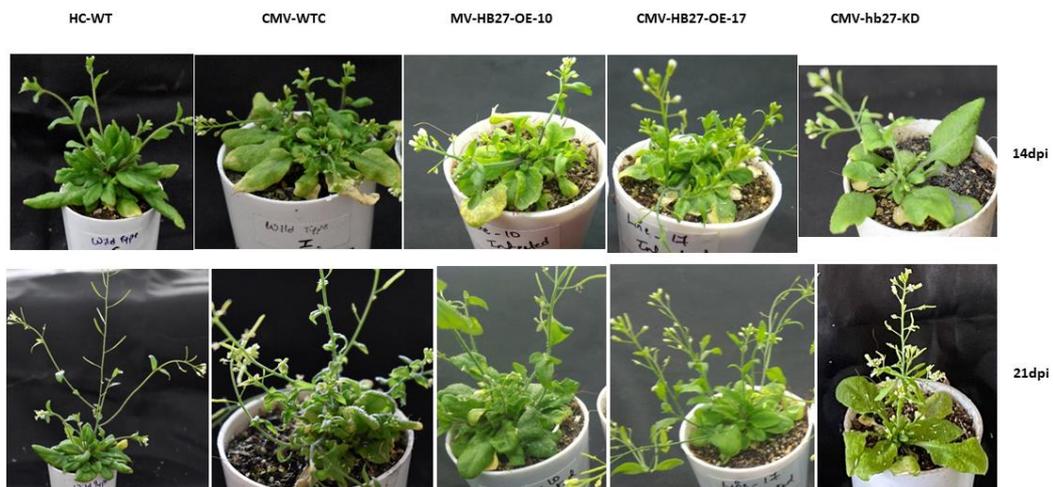


### Supplementary Figure S1:



Far western blotting: Purified HB27-HA-pET28a and CMV2b-MYC-pET28a proteins were run along with BSA run on SDS-PAGE gel and stained with coomassie brilliant blue (a). the ponceau staining was done to ensure the transfer of proteins and consider as loading control (b).

### Supplementary Figure S2:



At 14 and 21 dpi, the symptoms of CMV on HB27-OE, hb27-KD, and wild type plants were compared to healthy wild type plants. When compared to HB27-KD and wild type plants, the severity of CMV infection in HB27-OE is lower.

**Table S1:** List of primers used in this study. Underlined sequences represent the restriction sites. Italicized sequences showed the HA and MYC tag sequences.

|                  |  |
|------------------|--|
|                  | <b>Y2H construct</b>   |
| HB27             | FP 5' <u>AGATCT</u> ATGGATGAGATAAAACCAA 3'<br>RP 5' <u>CTCGAGTCATTCATCAATTATAATTT</u> 3'                                   |
| CMV 2b           | FP 5' <u>CATATGATGGATGTGTTGA</u> 3'<br>RP 5' <u>CCCGGGTCAAACGACCCTTC</u> 3'  |
|                  | <b>BiFC construct</b>  |
| HB27             | FP 5' <u>CTCGAGATGGATGAGATAAAACCAAAC</u> 3'<br>RP 5' <u>ACTAGTTTCATCAATTATAATTTTATCC</u> 3'                                |
| 2b               | FP 5' <u>GGATCCATGGATGTGTTGACAGTAGTG</u> 3'<br>RP 5' <u>CTCGAGAAACGACCCTTCGGCCATTCG</u> 3'                                 |
|                  | <b>Far western construct</b>   |
| HB27             | FP 5' GCAG <u>CCATATGTACCCATACGATGTTCCAGATTACGCTGATGAGATAAAACCAAAG</u> 3'<br>RP 5' <u>TGGTGCTCGAGTCATTCATCAATTATAAT</u> 3' |
| CMV 2b           | FP 5' <u>GGATCCATGGAACAAAACTCATCTCAGAAGAGGATCTGGATGTGTTGACAGTAG</u> 3'<br>RP 5' <u>TGGTGCTCGAGTTATCAAACGACCCTTCG</u>       |
|                  | <b>Overexpression construct</b>  |
| HB27             | FP 5' AAT <u>AGATCTTATGGATGAGATAAAACCAAAG</u> 3'<br>RP 5' <u>TAACTAGTTTCATCAATTATAATTTTATCC</u> 3'                         |
| GFP              | FP 5' GGAGTTGTCCCAATTCTTGT 3'<br>RP 5' GTCTCTCTTTTCGTTGGGATC 3'  |
| Hygromycin       | FP 5' CATAACAAGCCAACCACGGCCTCC 3'<br>RP 5' GCGTGGATATGTCCTGCGGGTA 3'   |
|                  | <b>Mutant genotyping</b>   |
| HB27 full length | FP 5' ATGGATGAGATAAAACCAAAGAAAGAAG 3'<br>RP 5' TTCATCAATTATAATTTTATCCTTGTG 3'  |
| LBb1.3           | 5' ATTTTGCCGATTCGGAAC 3'   |
|                  | <b>Real time Primers</b>   |
| 18S              | FP 5' TCTGCCCGTTGCTCTGATG 3'<br>RP 5' TCACCCGTCACCACCATAG 3'   |
| HB27             | FP 5' TCCTCGACTCAATCTTCAA 3'<br>RP 5' TTCTCCGGGATTAACCTCTC 3'  |
| CMV              | FP 5' ACCCTGAAACCGCCTGAAAT 3'<br>RP 5' TCCGAAGTAAACCCACACG 3'  |