## **Supporting Information**

Liu et al. 10.1073/pnas.1215543110

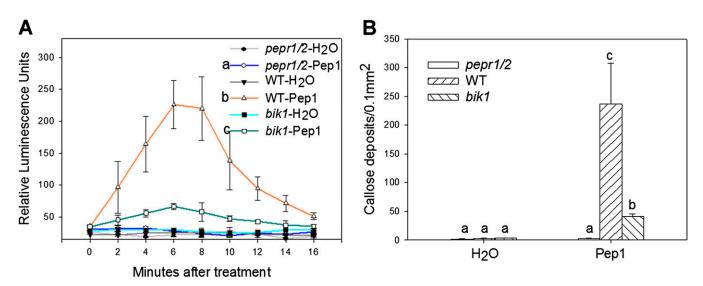


Fig. S1. Botrytis-induced kinase 1 (bik1) is compromised in Pep1-induced defenses. (A)  $H_2O_2$  production in leaf strips of bik1, pep1 receptor kinase 1/pep1 receptor kinase 2 (pep1/2), and WT treated with 1 μM Pep1. Different letters indicate significant difference (mean ± SD;  $n \ge 4$ ; P < 0.01). (B) Callose deposition in bik1, pep1/pep2 (pep1/2), and WT leaves treated with 1.5 μM Pep1. Different letters indicate significant difference (mean + SD; P < 0.01).

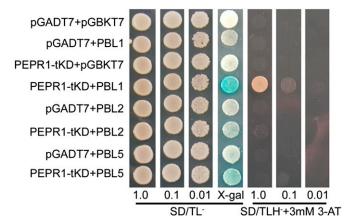


Fig. S2. Interaction of PBS1-like proteins (PBLs) with PEPR1. The truncated PEPR1 KD (PEPR1-tKD) prey plasmid was cotransfected into yeast with different PBL bait plasmids and tested for *lacZ* and *His* reporter activity as described Fig. 1A.

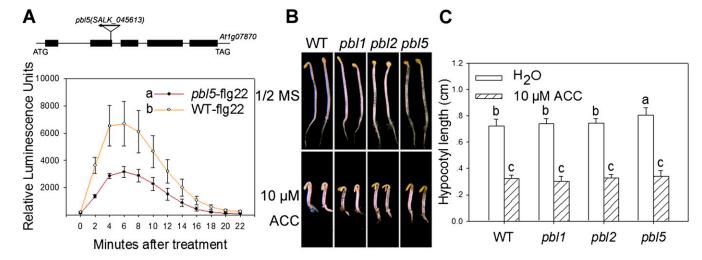


Fig. S3. (A) The pbl5 mutant is compromised in flg22-induced  $H_2O_2$  accumulation. (Upper) Schematic representation of T-DNA insertion in the PBL5 gene. Black boxes denote annotated exons from TAIR. The triangle and arrowhead indicate position and direction of T-DNA insertion, respectively. (Lower) pbl5 plants are compromised in flg22-induced oxidative burst. Different letters denote significant difference (mean  $\pm$  SD;  $n \ge 4$ ; P < 0.01). (B and C) pbl1, pbl2, and pbl5 mutants display normal triple response. (B) Photograph of 5-d-old WT, pbl1, pbl2, and pbl5 etiolated seedlings germinated on 1/2 MS medium with or without 10  $\mu$ M 1-aminocyclopropane-1-carboxylate (ACC). (C) Hypocotyl length of the seedlings (mean + SD;  $n \ge 12$ ; P < 0.01).

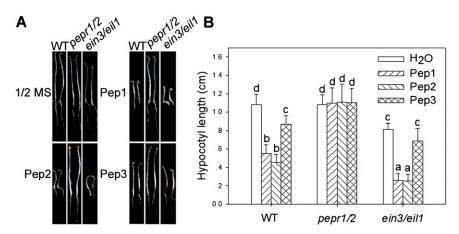


Fig. S4. Pep1-3 inhibits hypocotyl and root growth. (A) Photograph of 5-d-old etiolated seedlings grown in the presence of 10  $\mu$ M Pep1, Pep2, and Pep3. (B) Hypocotyl length. Different letters denote significant difference (mean + SD;  $n \ge 17$ ; P < 0.01).

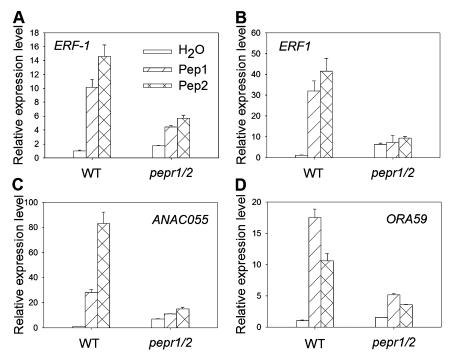


Fig. S5. Pep1 and Pep2 induce defense-related genes expression. (A–D) Real-time RT-PCR analysis of Pep1- and Pep2-induced ERF-1, ERF1, ANAC055, and ORA59 genes expression. The 8-d-old WT and pepr1/pepr2 seedlings were sprayed with  $H_2O$  or 10  $\mu$ M Pep1 or Pep2 containing 0.01% silwet L-77, and RNA was isolated 10 h later.

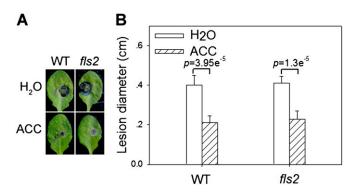


Fig. S6. FLS2 is not required for ACC-induced resistance to *Botrytis cinerea*. (A) Symptoms 2 d after *B. cinerea* inoculation. The experiments were performed as described in Fig. 5E. (B) Lesion size of indicated plants (mean + SD;  $n \ge 7$ ). P value is given above the bars.

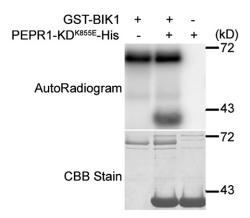


Fig. 57. BIK1 phosphorylates PEPR1 KD in vitro. His-tagged PEPR1 KD<sup>K855E</sup> was incubated with GST-tagged BIK1 in kinase buffer containing <sup>32</sup>P-γ-ATP, and protein phosphorylation was detected by autoradiography. CBB-stain shows protein loading. The experiment was performed as described in Fig. 6A.

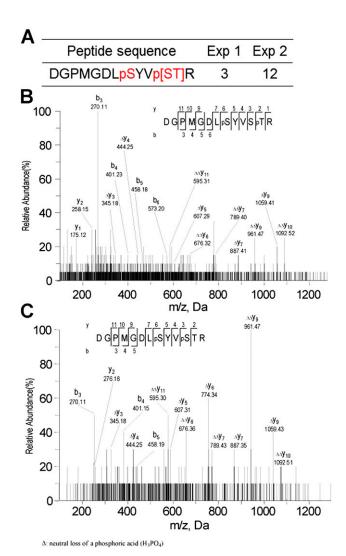


Fig. S8. Identification of BIK1 amino acids phosphorylated by PEPR1 KD. The GST-BIK1<sup>K105E</sup> protein, coexpressed with PEPR1-KD-His in *Escherichia coli*, was isolated and subjected to LC-MS/MS analysis. (*A*) Number of phosphorylated peptides corresponding to the activation loop identified in two independent experiments. Parenthesis indicates that the phosphorylation on serine 236 or threonine 237 cannot be easily distinguished. (*B* and *C*) Representative collision-induced dissociation (CID) spectra of the phosphopeptide DGPMGDLpSYVp[ST]R. The major b- and y-type ions are indicated in the mass graphs. Δ, neutral loss of a phosphoric acid.

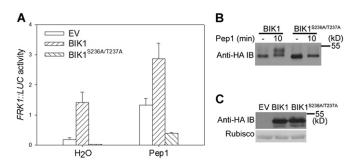


Fig. S9. Serine 236 and threonine 237 are required for Pep1-induced BIK1 phosphorylation and signaling. (*A*) Ala substitution of Ser-236 and Thr-237 in BIK1 blocks Pep1-induced *FRK1* reporter expression in protoplasts. Empty vector (EV), WT BIK1, and BIK1<sup>S236A/T237A</sup> plasmids were transfected into WT protoplasts along with *35S::R-LUC* and *FRK1::LUC*, treated with H<sub>2</sub>O or 1  $\mu$ M Pep1 for 3 h, and the *FRK1::LUC* activity was determined. (*B*) The BIK1<sup>S236A/T237A</sup> mutant is not phosphorylated upon Pep1 treatment. (*C*) BIK1 and BIK1<sup>S236A/T237A</sup> protein accumulation in *A*.