

Figure S1. Comparison of the growth of the wild-type *Xanthomonas oryzae* pv. *oryzicola* (Xoc) strain Rs105 and the *rpfF* mutant in nutrient-rich broth (NB). The point highlighted with a red line represented the time-point for extraction of extracellular proteins from the supernatants of wild-type strain and *rpfF* mutant. At that time, the cell density (OD_{600nm}) of wild-type strain and *rpfF* mutant was 2.5 and 2.9, respectively. Rs105, wild-type strain of Xoc; $\Delta rpfF$, the diffusible signal factor (DSF)-deficient mutant of Xoc (11). Three replicates for each treatment were used, and the experiment was repeated three times.

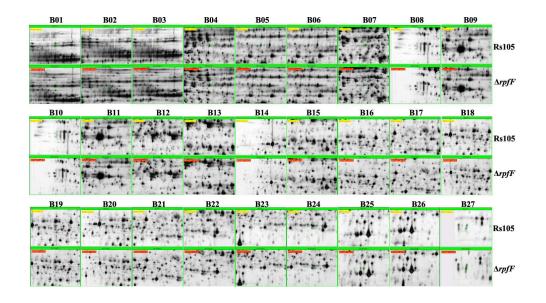


Figure S2. Identification of 27 extracellular protein spots negatively regulated by diffusible signal factor (DSF) in Xanthomonas oryzae pv. oryzicola (Xoc). These 27 protein spots

exhibited remarkably increased expression level (> 1.5 fold) in rp/F mutant compared to wild-type strain. Protein identifications are provided in Table 2. Rs105, the wild-type strain of Xoc;

 $\Delta rp/F$, the diffusible signal factor (DSF)-deficient mutant of Xoc (11). Three replicates for each treatment were used, and the experiment was repeated three times.

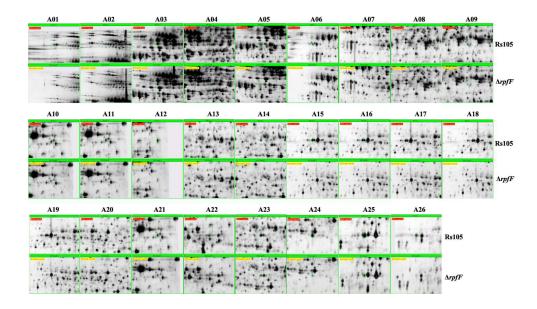


Figure S3. Identification of 26 extracellular protein spots positively regulated by diffusible signal factor (DSF) in Xanthomonas oryzae pv. oryzicola (Xoc). These 26 protein spots

exhibited significantly decreased expression level (> 1.5 fold) in rp/F mutant compared to wild-type strain. Protein identifications are provided in Table 2. Rs105, the wild-type strain of Xoc;

 $\Delta rpfF$, the diffusible signal factor (DSF)-deficient mutant of Xoc (11). Three replicates for each treatment were used, and the experiment was repeated three times.



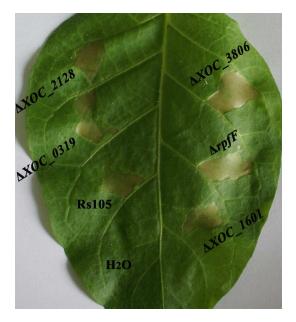


Figure S4. All the four selected genes of *Xanthomonas oryzae* pv. *oryzicola* (Xoc) were not required for triggering hypersensitive response (HR) in non-host plant (tobacco). The phenotype of HR, a programmed cell death, was observed around the inoculation sites in tobacco leaves after 24 h of infiltration with the wild-type and mutants. Rs105, the wild-type strain of Xoc; ΔXOC_0319 , the deletion mutant of XOC_0319 , which encodes a Ax21 (activator of XA21-mediated immunity)-like protein; ΔXOC_3806 , the deletion mutant of XOC_3806 , which encodes a serine protease; ΔXOC_1601 , the deletion mutant of XOC_1601 , which encodes a cysteine protease; ΔXOC_2128 , the deletion mutant of XOC_2128 , which encodes a polygalacturonase. $\Delta rpfF$, the diffusible signal factor (DSF)-deficient mutant of Xoc (11); H₂O, a negative control. Three replicates for each treatment were used, and the experiment was repeated three times.

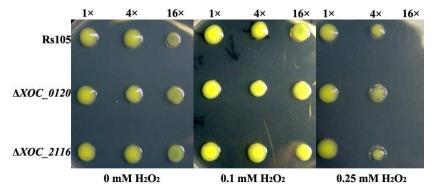


Figure S5. Mutation of XOC_0120 or XOC_2116 did not impair the resistance ability to hydrogen peroxide

(H₂O₂) in *Xanthomonas oryzae* pv. *oryzicola* (Xoc). Xoc strains were cultured to mid-logarithmic phase (OD_{600nm} = 1.0) in NB medium (1 x), and then four-fold (4 x) and sixteen-fold dilutions (16 x) were made, and grown on NB agar plates with 0, 0.1, or 0.25 mMH₂O₂. Rs105, the wild-type strain of Xoc; ΔXOC_0120 , the deletion mutant of *XOC_0120*, which encodes a glutathione peroxidase; ΔXOC_2116 , the deletion mutant of *XOC_0120*, which encodes a glutathione peroxidase for each treatment were used, and the experiment was repeated three times.

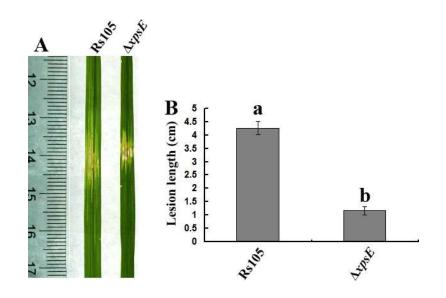


Figure S6. Mutation of *xpsE* impaired the virulence of *Xanthomonas oryzae* pv. *oryzicola* (Xoc) on host rice. (A) Representative result of water-soaking lesion lengths on the rice seedling leaves (cv. Shanyou63, 2-week old) by infiltration with wild-type strain and selected four mutants. (B) Calculated data of water-soaking lesion lengths on the leaves of rice seedling leaves. Rs105, wild-type strain of Xoc; $\Delta xpsE$, the deletion mutant of *xpsE* (*XOC_3805*). The product of *xpsE* is a component of the *xps* cluster responsible for T2SS constitution. Three replicates were used for each treatment, and the experiment was repeated three times. Vertical bars represent standard errors. Different letters above data bars indicate a significant difference between the wild-type strain and tested mutants (*P*<0.05; *t*-test).

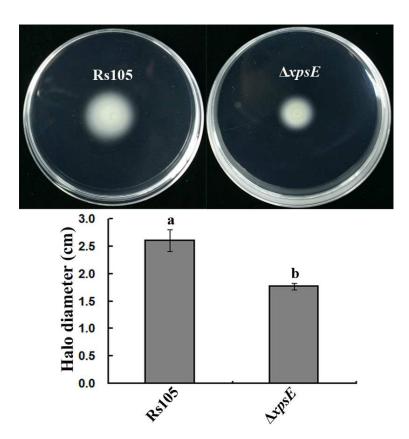


Figure S7. Mutation of *xpsE* impaired cell motility in *Xanthomonas oryzae* pv. *Oryzicola* (Xoc). (A) Representative photograph of motility halos formed by wild-type strain and the four selected mutants on XOM3 semi-solid motility medium. (B) Quantitative determination of motility halos of the wild-type strains and the four selected mutants. Rs105, the wild-type strain of Xoc; $\Delta xpsE$, the deletion mutant of *xpsE* (*XOC_3805*). The product of *xpsE* is a component of the *xps* cluster responsible for T2SS constitution. Three replicates were used for each treatment, and the experiment was repeated three times. Vertical bars represent standard errors. Different letters above data bars indicate a significant difference between the wild-type strain and the tested mutants (P<0.05; *t*-test).

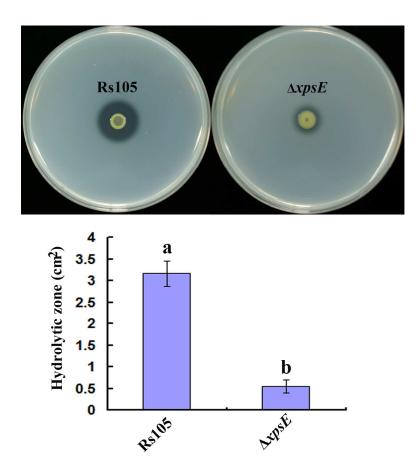


Figure S8. Mutation of *xpsE* impaired extracellular protease activity in *Xanthomonas oryzae* pv. *oryzicola* (Xoc). (A) Representative photograph of hydrolytic zone formed by the wild-type strain and the *xpsE* mutant on selective medium. (B) Quantitative determination of hydrolytic zone of the wild-type strains and *xpsE* mutant. Rs105, the wild-type strain of Xoc; $\Delta xpsE$, the deletion mutant of *xpsE* (*XOC_3805*). The product of *xpsE* is a component of the *xps* cluster responsible for T2SS constitution. Three replicates were used for each treatment, and the experiment was repeated three times. Vertical bars represent standard errors. Different letters above data bars indicate a significant difference between the wild-type strain and tested mutants (*P*<0.05; *t*-test).

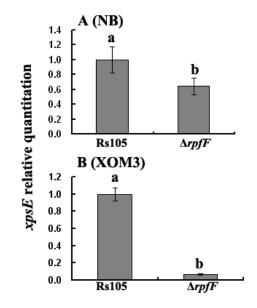


Figure S9. *xpsE* transcription was positively regulated by diffusible signal factor (DSF) in *Xanthomonas* oryzae pv. oryzicola (Xoc). (A) Comparison of transcriptional expression of *xpsE* between the *rpfF*-deletion mutant and the wild-type strain Rs105 in nutrient-rich broth (NB). Total RNA was extracted from the Rs105 and $\Delta rpfF$ at the mid-stage of growth (OD_{600nm} = 1.6). (B) Comparison of transcriptional expression of *xpsE* between the *rpfF*-deletion mutant and the wild-type Rs105 in plant-cell mimicking broth (XOM3). The Xoc strains were pre-incubated in NB medium overnight, re-suspended at OD₆₀₀ = 2.0 in XOM3 medium, and washed twice. 2 ml of the bacterial suspension was then inoculated into 50 ml of XOM3 plant-cell mimicking broth (pH 6.5) at 28 °C for 16 h.⁶⁹ Total RNA was then extracted from the Rs105 and $\Delta rpfF$. Three replicates for each treatment were used, and the experiment was repeated three times. Vertical bars represent standard errors. Different letters above bars indicate a significant difference between the wild-type strain and *rpfF*-deletion mutant (*P*<0.05; *t*-test).

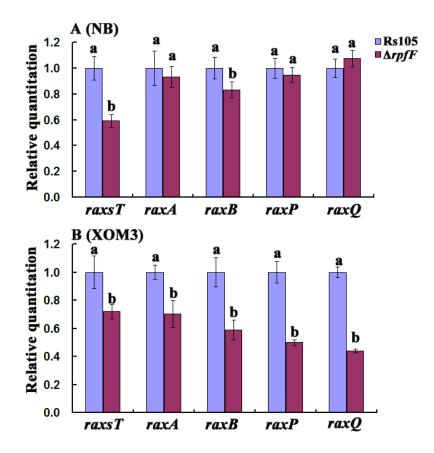


Figure S10. Transcriptional expression of the five *rax* genes of *Xanthomonas oryzae* pv. *oryzicola* (Xoc) in the *rpfF*-deletion mutant and wild-type strain. (A) Comparison of transcriptional expression of the five *rax* genes (required for Δx^{21}) between the *rpfF*-deletion mutant and the wild-type Rs105 in nutrient-rich broth (NB). Total RNA was extracted from the Rs105 and $\Delta rpfF$ at the mid-stage of growth (OD_{600nm} = 1.6). (B) Comparison of transcriptional expression of the five *rax* genes between the *rpfF*-deletion mutant and the wild-type Rs105 in plant-cell mimicking broth (XOM3). The Xoc strains were pre-incubated in NB medium overnight, re-suspended at OD_{600nm} = 2.0 in XOM3 medium, and washed twice. 2 ml of the bacterial suspension was then inoculated into 50 ml of XOM3 plant-cell mimicking broth (pH 6.5) at 28 °C for 16 h.⁶⁹ Total RNA was then extracted from the Rs105 and $\Delta rpfF$. Ax21, activator of XA21-mediated immunity. Three replicates for each treatment were used, and

the experiment was repeated three times. Vertical bars represent standard errors. Different letters above bars

indicate a significant difference between the wild-type strain and the *rpfF*-deletion mutant (P<0.05; t-test).

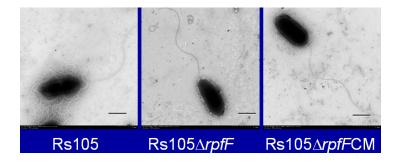


Figure S11. Effect of the *rpfF* mutation on flagella and cell shapes in *Xanthomonas oryzae* pv. *oryzicola* (Xoc).

Cells were negatively stained with 2% potassium phospotungstate. Flagella and cells were detected under a Hitachi transmission electron microscope at 80 kilovolts. Rs105, wild-type strain of Xoc; $\Delta rpfF$, the diffusible signal factor (DSF)-deficient mutant of Xoc (11). $\Delta rpfF$ CM, the corresponding complemented strain of $\Delta rpfF$. Bar represents 1µm; three replicates for each treatment were used, and the experiment was repeated three times.