

1 **Supplementary files**

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3 **The relative importance of soil moisture in predicting bacterial wilt disease occurrence**

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7 The supplementary information contains six files. The table file includes statistical information
8 about multiple regression analysis, and figure files provide further detail on the variation of
9 physicochemical properties and bacterial communities in tomato rhizosphere and how it was
10 linked to pathogen density and plant healthy in tomato rhizosphere microbiomes.

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13 **Supplementary Table 1. ANOVA table summarizing the relative importance of abiotic**
 14 **physicochemical and biotic soil properties in predicting pathogen densities in tomato**
 15 **rhizosphere samples.**

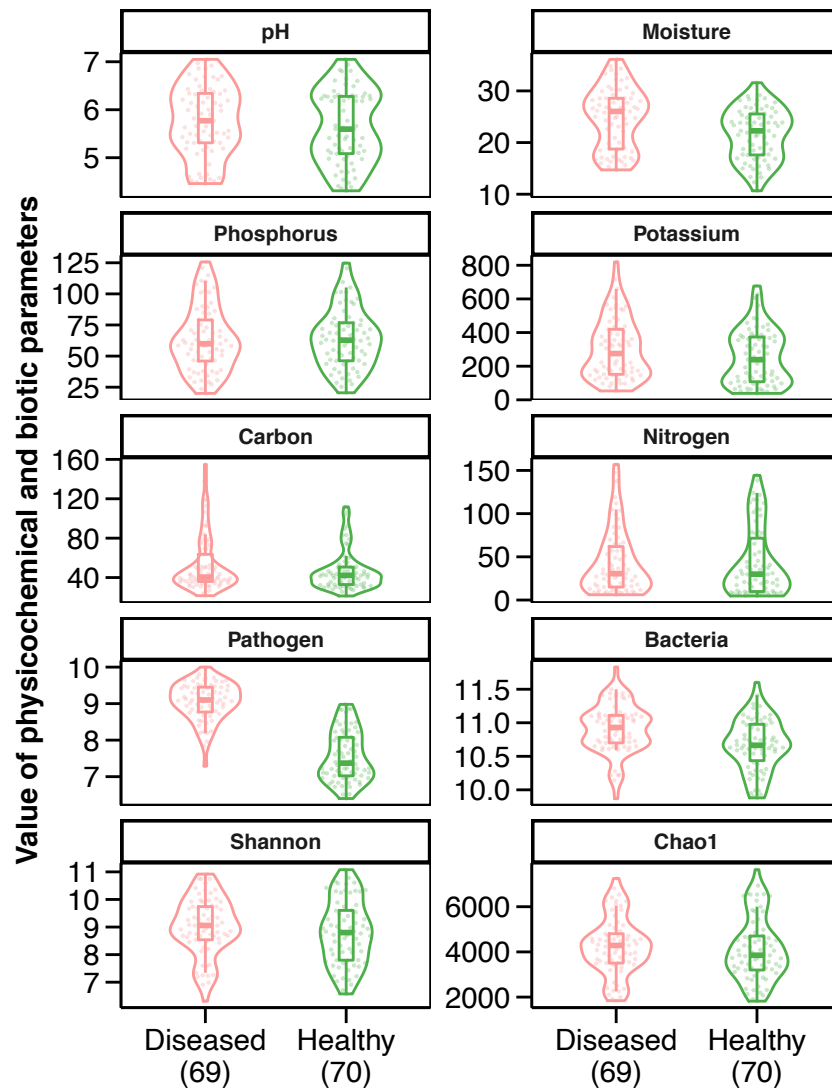
Predictor		Sum	Mean			Relative
variable	Df	Square	Square	<i>F</i> -value	<i>P</i> -value	weight
pH	1	10.29	10.29	18.68	<0.0001↓	14.59%
Moisture	1	28.45	28.45	51.66	<0.0001↑	40.36%
Phosphorus	1	0.14	0.14	0.25	0.6170↑	0.20%
Potassium	1	0.14	0.14	0.26	0.6141↑	0.20%
Carbon	1	0.07	0.07	0.12	0.7310↓	0.09%
Nitrogen	1	0.03	0.03	0.05	0.8271↓	0.04%
Bacterial						
abundance	1	16.05	16.05	29.15	<0.0001↑	22.77%
Shannon	1	1.78	1.78	3.24	0.0744↑	2.53%
Chao1	1	1.33	1.33	2.42	0.1226↓	1.89%
Bray-Curtis	1	0.11	0.11	0.21	0.6499↑	0.16%
Residuals	128	70.50	0.55			

Model Summary

AIC: 324.09; $F_{10,128} = 10.6$, $R^2 = 0.45$, $P < 0.0001$

16 Note: The significant effects ($P < 0.05$) are shown in red colour and the ‘up’ and ‘down’ arrows
 17 denote for positive and negative effects, respectively, based on multiple regression model.

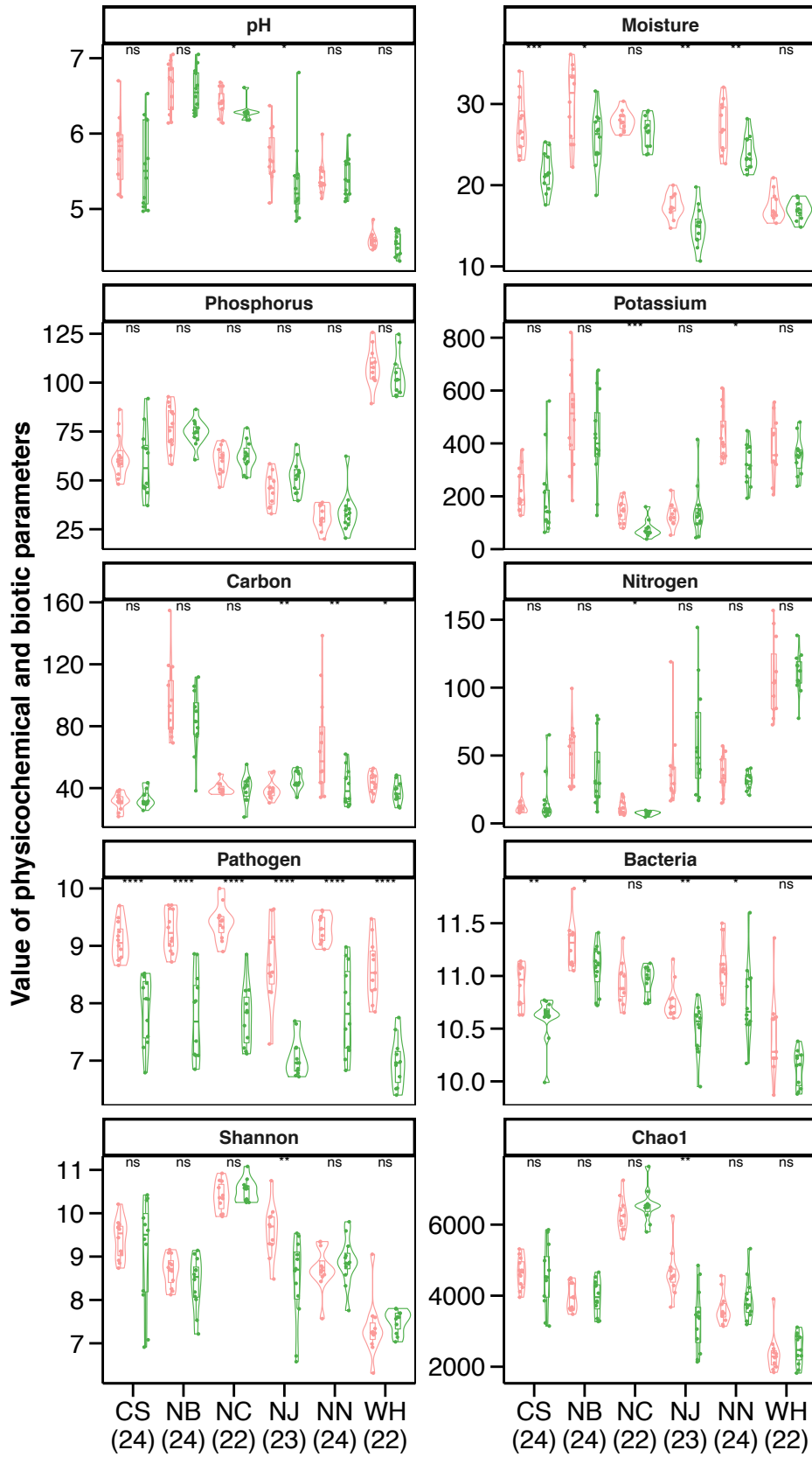
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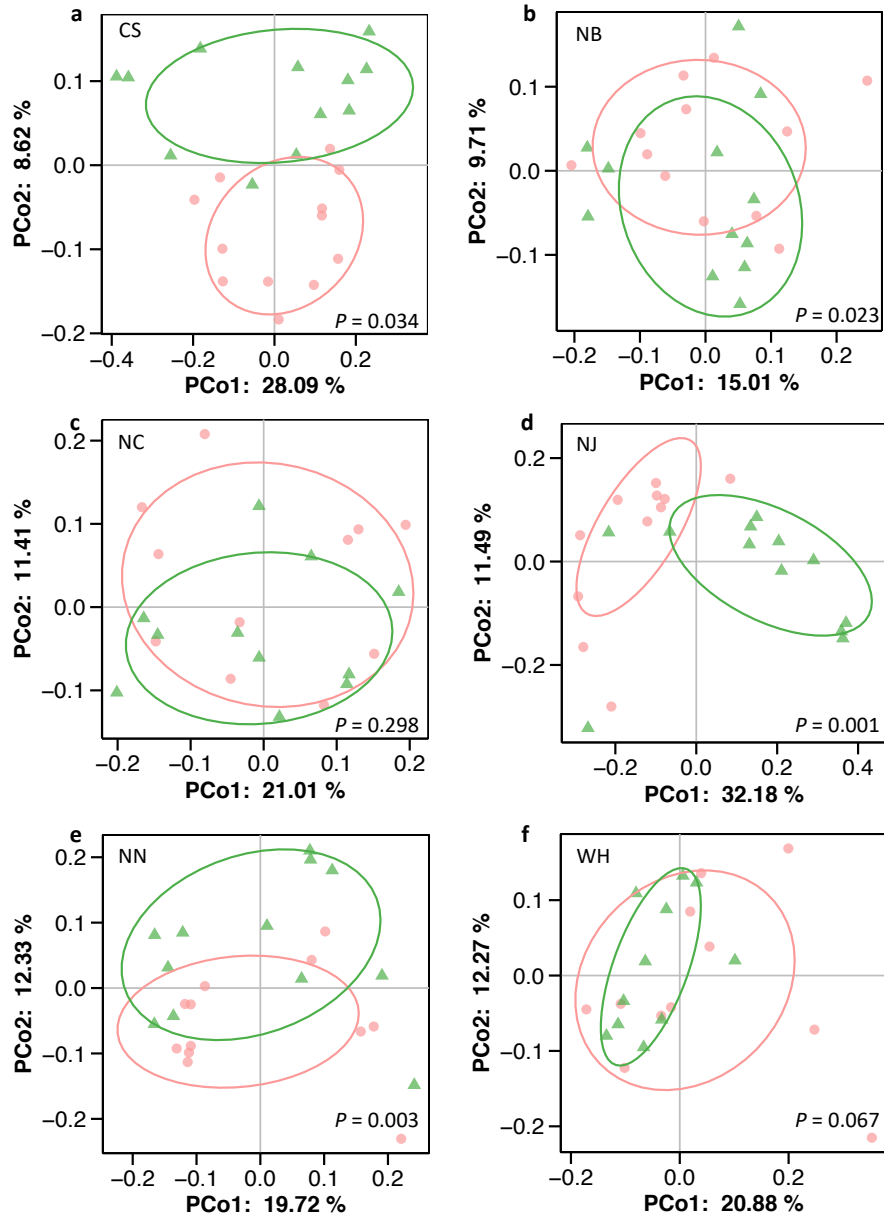
20 **Supplementary Figure 1. Differences in abiotic physicochemical and biotic soil**
 21 **properties between diseased and healthy plants.**

22 All data is pooled over sampling locations and the numbers in the parentheses denote the
 23 sample size in each group. The 'ns' denotes for non-significant difference ($P > 0.05$) and stars
 24 (*, **, *** and ****) show significant differences at levels $P < 0.05$, $P < 0.01$, $P < 0.001$ and
 25 $P < 0.0001$, respectively. Pathogen and total bacterial abundances are abbreviated as 'Pathogen'
 26 and 'Bacteria', respectively. Each violin plot shows the distribution of 69 and 70 rhizosphere
 27 soil samples in diseased and healthy plants, respectively.



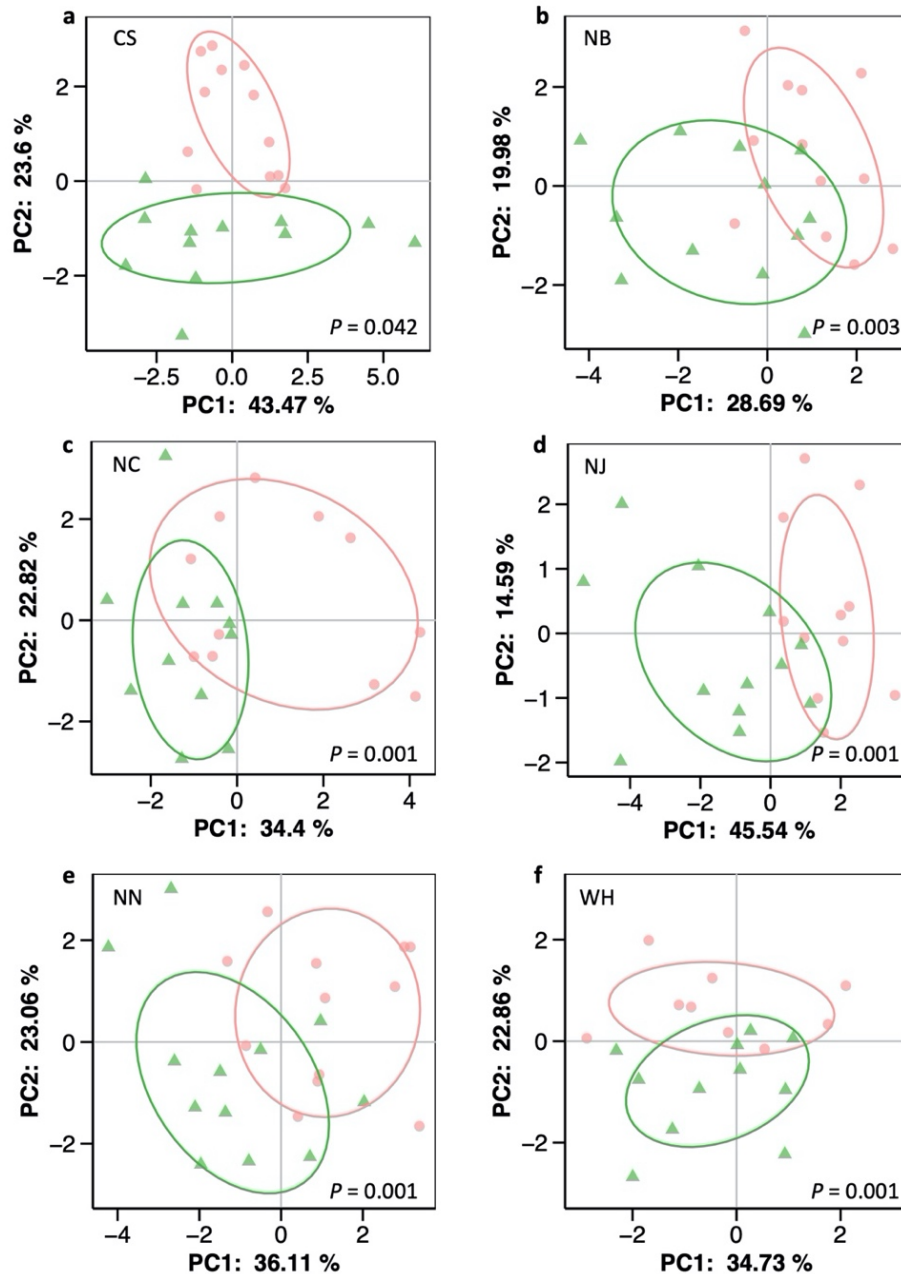
29 **Supplementary Figure 2. Differences in abiotic physicochemical and biotic soil**
30 **properties of diseased and healthy plants in different sampling locations (provinces).**

31 Numbers in the parentheses show sample size in each group. The 'ns' denotes for non-
32 significant difference ($P > 0.05$) and stars (*, **, *** and ****) show significant differences
33 at levels $P < 0.05$, $P < 0.01$, $P < 0.001$ and $P < 0.0001$, respectively. Pathogen and total bacterial
34 abundances are abbreviated as 'Pathogen' and 'Bacteria', respectively. Each violin plot shows
35 the distribution of rhizosphere soils in each province. Sampling locations are abbreviated as
36 follows: CS = Changsha, NB = Ningbo, NC = Nanchang, NJ = Nanjing, NN = Nanning and
37 WH = Wuhan.



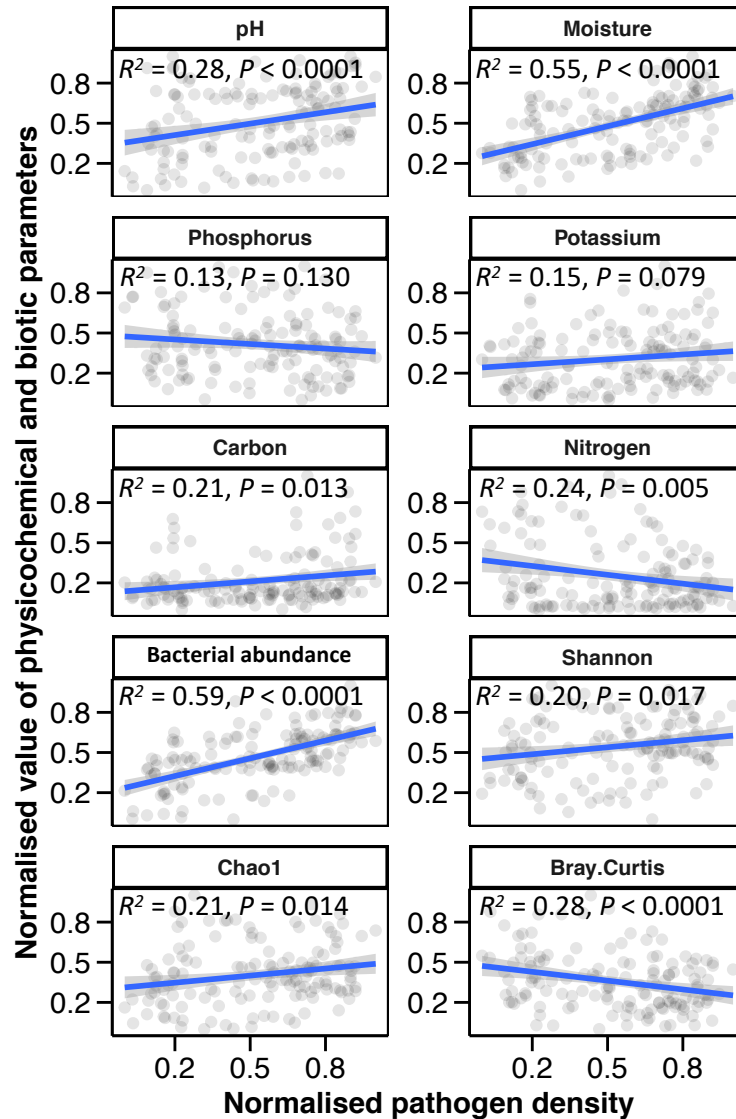
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 39 **Supplementary Figure 3. Differences in bacterial community composition (PCoA)**
 40 **between healthy (green) and diseased (red) rhizosphere soil samples in different**
 41 **sampling locations (a-f).**

42 PERMANOVA was used to identify microbial composition difference between the rhizosphere
 43 soil of diseased and healthy plants in each location based on Bray–Curtis distance matrices. *P*-
 44 values are indicated in each panel. Green triangles and red circles denote healthy and diseased
 45 plants, respectively. Sampling locations are abbreviated as follows: CS = Changsha, NB =
 46 Ningbo, NC = Nanchang, NJ = Nanjing, NN = Nanning and WH = Wuhan.



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 48 **Supplementary Figure 4. Differences in physicochemical and biotic parameters (PCA)**
 49 **between healthy (green) and diseased (red) rhizosphere soil samples in different**
 50 **sampling locations (a-f).**

51 PERMANOVA was used to identify the environmental difference between the rhizosphere
 52 soil of diseased and healthy plants in each location based on Euclidean distance matrices. *P*-
 53 values are indicated in each panel. Green triangles and red circles denote healthy and diseased
 54 plants, respectively. Sampling locations are abbreviated as follows: CS = Changsha, NB =
 55 Ningbo, NC = Nanchang, NJ = Nanjing, NN = Nanning and WH = Wuhan.

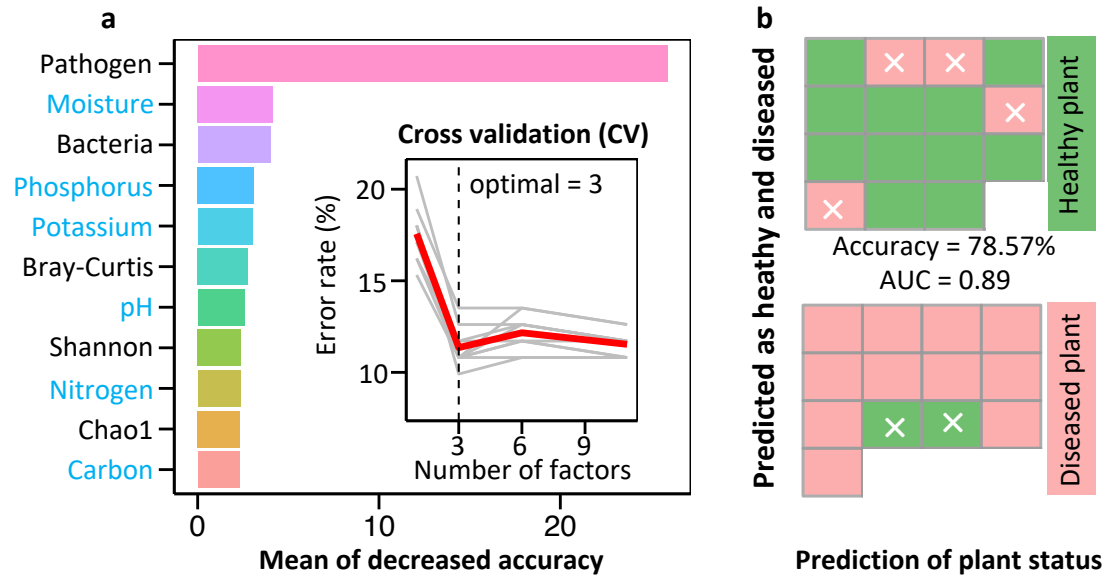


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57 **Supplementary Figure 5. Linear correlations between the normalised pathogen**
 58 **densities and abiotic physicochemical and biotic soil properties (averaged over healthy**
 59 **and diseased plant samples).**

60 Blue lines indices the linear fitting of curves, while R^2 indicates the Spearman correlation
 61 coefficient of the linear regression and P -values the significance of each correlation.

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64 **Supplementary Figure 6. Comparing the relative importance of all soil parameters in**
 65 **predicting plant health status.**

66 (a) Relative importance rank of overall abiotic physicochemical (blue) and biotic (black) soil
 67 properties and ten-fold cross-validation of random forest model (inset in a) based on the
 68 training set (80% of randomly selected rhizosphere samples). Pathogen and total bacterial
 69 abundances are abbreviated as ‘Pathogen’ and ‘Bacteria’, respectively. (b) Validation of
 70 random-forest model with a test set (20% of remaining samples) predicting plant disease
 71 outcomes based on soil properties: green and red filled cells denote for correct predictions,
 72 while filled cells with white crosses denote for false predictions. The overall model gained an
 73 average accuracy (78.6%) in classifying plant status, with 84.6% accuracy for diseased and
 74 73.3% accuracy for health plants (AUC = 0.89).

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