# Evaluation of correlation methods for construction of a co-occurrence network of rice crop health survey data

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### Introduction

Surveys<sup>1</sup> conducted in rice fields of Asia have generated large data sets to characterize pests, pest injuries, and production situation.

Network analysis is widely applied in biological science and can be used to improve the efficiency of pest management. The construction of a network model is based on correlation coefficients. Methods are often selected based on the type of data to be analyzed.

### Problem

CHECK

**±** 200 −

S 100-

**5** 200 -

250 -200 -150 -100 -

tail.

Histogram of WH

Histogram of RB

Histogram of NBS

1000 2000

Which correlation measure is suitable for collected data on pest and pest injuries?

### Choosing the correlation measure

**CHECK** 

whether the data are normally distributed or not.

**COMPARE** the similarity of correlation coefficients from different correlation measures.

**IDENTIFY** 

which suitable correlation measure can capture biological relationships the most between variables.

Pairwise correlation

between pests and pest

using a suitable measure

injuries are computed

### Conclusion

Spearman's rank correlation measure is the most suitable because:

- collected data are not normally distributed (Fig. 1),
- it yields the difference results of parametric correlation measures (Fig. 2), and
- it detects naturally occurring, nonlinear relationships (Fig. 3).

### Evaluation

Correlation matrix

construction<sup>4</sup>

A. Input (pests and pest injuries data)

Visual assessments and statistical tests

the normality of data distribution (Fig. 1).

Histogram of SS

0 5 10 15 20 25

value

Histogram of RBB

0 20 40 60 80

Histogram of BPH

0 200400600800

Fig. 1. Histograms of data on pests and pest injuries. The

distribution of most variables are described as power law or long

**3** 200 -

were performed (Shapiro-Wilk test<sup>2</sup>) to check

Histogram of WM

0 300 600 900

value

Histogram of GLH

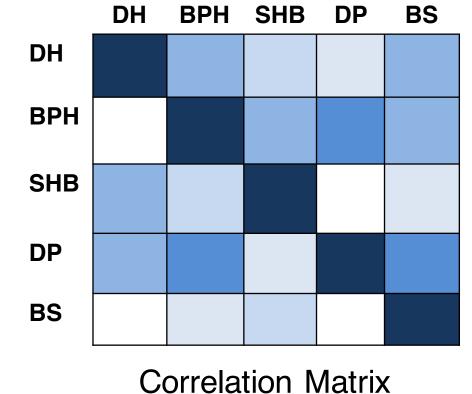
0 25 50 75 100

50 100 150

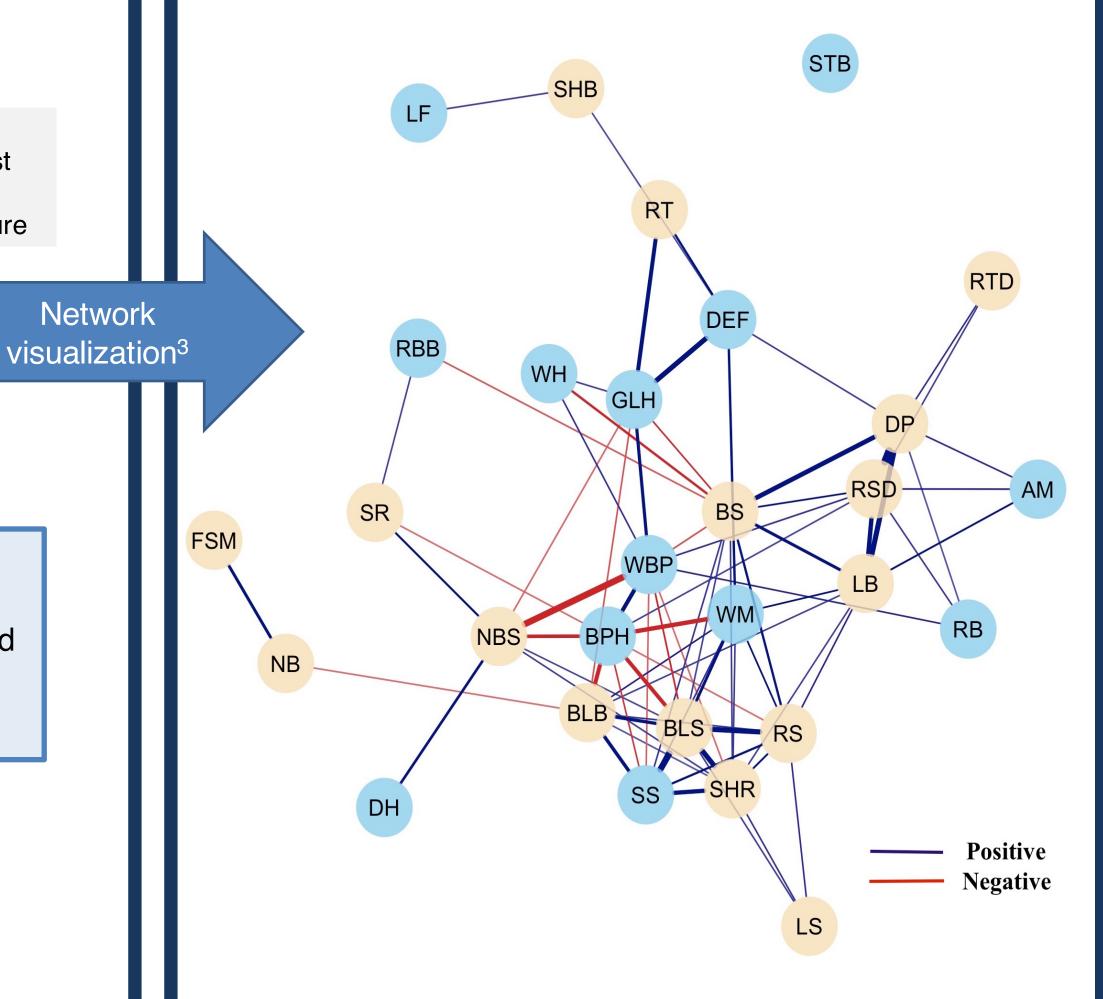
Histogram of WPH

€----- 29 types of injuries -----DH BPH SHB DP BS F1

B. Correlation Matrix

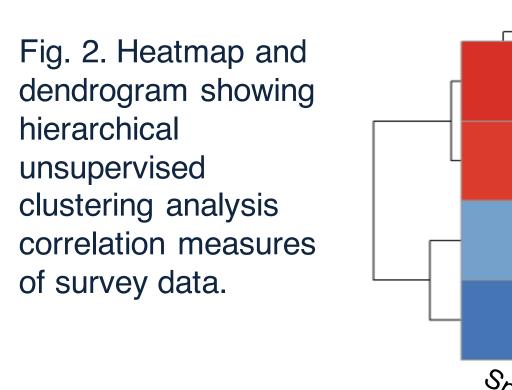


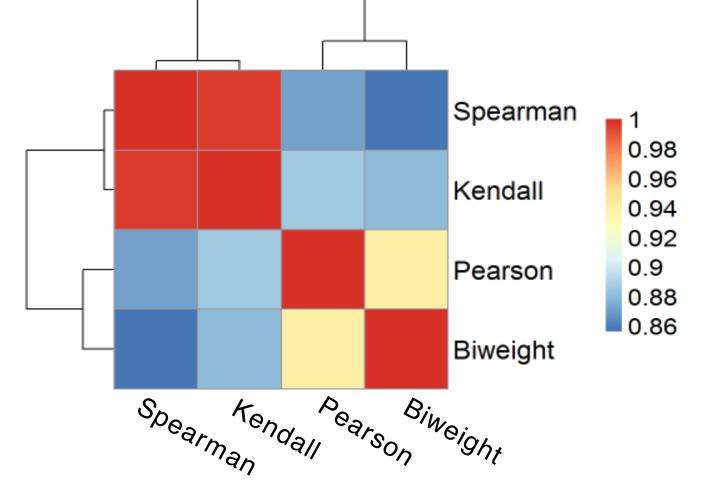
C. Network output



### COMPARE

Clustering analysis measures using Euclidian distance shows two groups of correlation methods (Fig. 2). The first is parametric correlation measures (Pearson and Biweight) and the second group is nonparametric correlation measure (Spearman and Kendall).



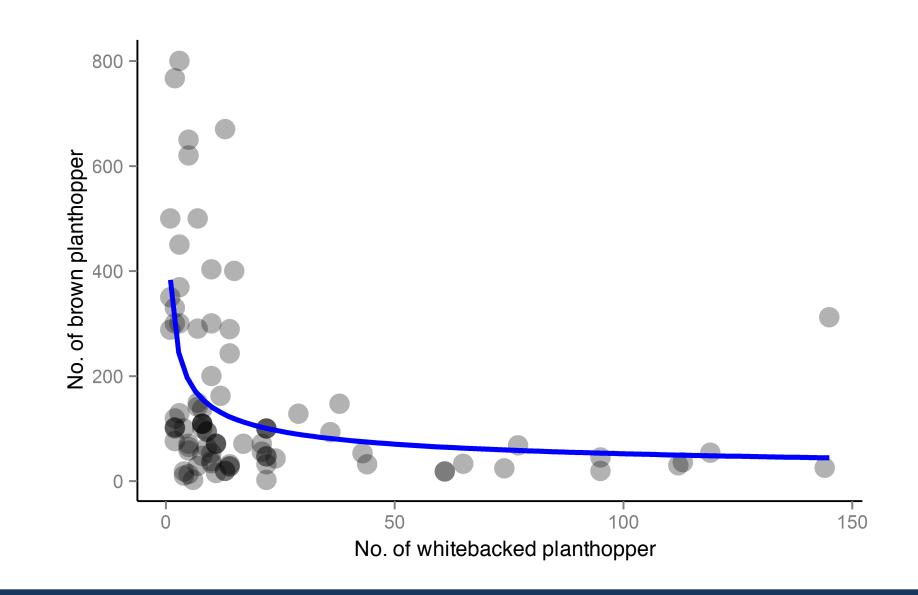


## The non-linear relationship between BPH and WBPH<sup>5</sup>

was detected by Spearman correlation, but not by Pearson, Biweight, and Kendall correlations. Other correlation measures, which are able to capture nonlinear correlation can be considered.

#### Fig. 3. Scatter plot illustrating the non-linear relationship between the brown planthopper and whitebacked planthopper.

**IDENTIFY** 



Co-occurrence network of pests - pest injuries based on Spearman's rank correlation

Blue and yellow nodes represent pest injuries and diseases, respectively. The layout of the network graph is based on the Fruchterman-Reingold algorithm, which places nodes with stronger or more connections closer to each other.

Represents animal pest group

AM = Army wormBPH = Brown planthopper DEF = Defoliators DH = Deadheart GLH = Green leafhopper LF = Leaffolder

WMA = Whorl maggot WBP = Whitebacked planthopper WH = Whitehead

SS = Silver shoot

STB = Stink bug

RT = RatRB = Rice bug RBB = Rice black bug

#### Represents disease group

BLB = Bacterial leaf blight BLS = Bacterial leaf streak BS = Brown spot DP = Dirty panicle FSM = False smut LB = Leaf blast

LS = Leaf scald

NB = Neck blast

NBS = Narrow brown spot RSD = Ragged stunt disease RTD = Rice tungro

disease SHB = Sheath blight SHR = Sheath rot SR = Stem rot

### Acknowledgments

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### References

<sup>1</sup>Savary S, Castilla NP. 2009. A survey portfolio to characterize yield-reducing factors in rice. An updated version of IRRI Discussion Paper No. 18. 32 p.

<sup>2</sup>R Core Team. 2015. R: A language and environment for statistical computing. R Foundation for Statistical Computing, Vienna, Austria. URL http://www.r-project.org/.

<sup>3</sup>Epskamp S, Cramer AO, Waldorp LJ, Schmittmann VD, Borsboom D. 2012. qgraph: Network visualizations of relationships in psychometric data. Journal of Statistical Software 48:1-18.

<sup>4</sup>Langfelder P, Horvath S. 2008. WGCNA: an R package for weighted correlation network analysis. BMC Bioinformatics 9:559. <sup>5</sup>Win SS, Muhamad R, Ahmad ZAM, Adam NA. 2011. Population fluctuations of brown planthopper, *Nilaparvata* lugens Stål. and whitebacked planthopper, Sogatella furcifera Horvath on rice. Journal of Entomology 8:183-190.



