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Automatic report  
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Study ID: Bryony\_Willcox\_Mangifera\_indica\_Australia\_2016

Contact: bwillcox@myune.edu.au

Credit: Bryony Willcox (The University of New England)

Publication: NA

Number of sites: 9

Year of sampling: 2016

Crop: Mangifera indica

Variety: There are missing varieties (given 0 out of 9).

Location: Full information.

Country: All reported countries are OK.

Field size: There are missing field sizes (given 0 out of 9).

Management: Full information.

Sampling period: There are missing sampling periods.

Richness: There is not enough taxonomic resolution to estimate richness.

species groups considered: NA

Abundance: Full information.

Visitation rate units: NA

Visitation rate: There are missing values (given 0 out of 9).

Yield units: average fruit set per 100 flowers

Yield: There are missing values (given 7 out of 9).

Alternative yield units: average fruit number per metre<sup>2</sup> of tree canopy just prior to harvest

Alternative yield: Full information.

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Queries  
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Instructions: Please, edit this file and answer the following queries within the document, one by one. Then send the edited 'Summary\_report\_and\_queries' (with your comments) and your 'Data\_ownership' (excel) file to alfonso.allen.perkins+observeddataset@gmail.com before the 20th of July.

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• Please check that credit information is correct and add the corresponding affiliations and acknowledgements/funding information in your 'Data\_ownership' (excel) file.  
• If your study is already published, please check that its DOI is correct.  
• If possible, please provide the names of missing crop varieties.  
• If possible, please provide the area of each field [in hectares].

- If possible, please provide the sampling start month and sampling end month for each field, respectively, and use a numeric format (for example, 1 for January, 2 for February and so on).
- There are sites without visitation rate records. Please, check that such information is correct. See also the information about OBServ data processing in your ‘First read me - General report’ pdf file.
- According to our raw data, there is not enough taxonomic resolution to estimate richness. Please, check that such information is correct. See also the information about OBServ data processing in your ‘First read me - General report’ pdf file.
- There are sites without yield. Please, check that such information is correct. If possible, please provide the missing values (or the corresponding z-scores).
- Please, check that the brief description of your methodology (in your insect\_sampling file) is correct.