```
================
Automatic report
================
```

Study ID: Romina_Rader_Mangifera_indica_Australia_2014

Contact: rrader@une.edu.au

Credit: Dr Romina Rader (The University of New England)

Publication: NA

Number of sites: 10

Year of sampling: 2014

Crop: Mangifera indica

Variety: There are missing varieties (given 0 out of 10).

Location: Full information.

Country: All reported countries are OK.

Field size: There are missing field sizes (given 0 out of 10).

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 10).

Yield units: average fruit number per metre2 of tree canopy just prior to harvest

Yield: Full information.

Alternative yield units: NA

Alternative yield: NA

```
========================================================
Queries
========================================================
```
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+observdataset@gmail.com
before the 20th of July.
```
========================================================
```

- 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.

- Please, check that the brief description of your methodology (in your insect_sampling file) is correct.