
Automatic report

Study ID: Juliana_Hipólito_Coffea_arabica_Brazil_2013

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Number of sites: 31 Year of sampling: 2013 Crop: Coffea arabica

Variety: There are missing varieties (given 0 out of 31).

Location: UTM zone is needed. There are missing locations (given 0 out of 31).

Country: There are no locations to check.

Field size: Full information.

Management: Full information.

Sampling period: There are missing sampling periods.

Richness: Full information.

species groups considered: No taxa restrictions have been identified.

Abundance: Full information.

Visitation rate units: visits per 100 flowers and hour

Visitation rate: Full information.

Yield units: bags/hectare Yield: Full information.

Alternative yield units: fruit set

Alternative yield: There are missing values (given 30 out of 31).

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 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).
- If possible, please provide the latitude and longitude of the missing locations [in decimal degrees].
- Please check that our information about the species groups considered is correct.
- There are sites without their alternative yield values. 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.