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Automatic report  
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Study ID: Smitha\_Krishnan\_Coffea\_canephora\_India\_2014

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Credit: Smitha Krishnan, ETH Zurich and Ms. Maike Nesper (The PI of the project MOCA is Prof. Jaboury Ghazoul)

Publication: NA

Number of sites: 25

Year of sampling: 2014

Crop: Coffea canephora

Variety: There are missing varieties (given 0 out of 25).

Location: Full information.

Country: All reported countries are OK.

Field size: There are missing field sizes (given 0 out of 25).

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: (average number of) visits per inflorescence and hour

Visitation rate: Full information.

Yield units: kg/ha (dried and hulled beans)

Yield: Full information.

Alternative yield units: kg/ha (dried berries)

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).
- 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.
- When processing your raw data, we aggregated the measurements at different distances within each site. Is it correct?