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Automatic report
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Study ID: Natacha_Chacoff_Citrus_paradisi_Argentina_2000
Contact: nchacoff@gmail.com
Credit: Natacha Chacoff (Instituto de Ecologia Regional. CONICET UNT)
Publication: 10.1111/j.1365-2664.2005.01116.x,10.1098/rspb.2007.1547
Number of sites: 4
Year of sampling: 2000
Crop: Citrus paradisi
Variety: Full information.
Location: Full information.
Country: All reported countries are OK.
Field size: There are missing field sizes (given 0 out of 4).
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: There are missing values (given 0 out of 4).
Visitation rate units: (average number of) visits per 100 flowers and hour
Visitation rate: Full information.
Yield units: fruit set (%)
Yield: Full information.
Alternative yield units: NA
Alternative yield: NA

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