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Automatic report  
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Study ID: Breno\_M\_Freitas\_Malpighia\_emarginata\_Brazil\_2011  
Contact: freitas@ufc.br  
Credit: Breno M. Freitas  
Publication: 10.1073/pnas.1517092112  
Number of sites: 8  
Year of sampling: 2011  
Crop: Malpighia emarginata  
Variety: There are missing varieties (given 0 out of 8).  
Location: UTM zone is needed. There are missing locations (given 0 out of 8).  
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: only bees  
Abundance: Full information.  
Visitation rate units: NA  
Visitation rate: There are missing values (given 0 out of 8).  
Yield units: % fruit set per flowers  
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 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].
- 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.
- Please, check that the brief description of your methodology (in your insect\_sampling file) is correct.
- Have *Apis Mellifera* records been excluded?