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Automatic report  
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Study ID: Yael\_Mandelik\_Citrullus\_lanatus\_Israel\_2009

Contact: gidpisa79@yahoo.com

Credit: Gideon Pisanty, Yael Mandelik

Publication: 10.1890/14-0910.1

Number of sites: 17

Year of sampling: 2009

Crop: Citrullus lanatus

Variety: There are missing varieties (given 0 out of 17).

Location: Full information.

Country: Reported countries may contain errors.

Field size: There are missing field sizes (given 0 out of 17).

Management: There are missing results for management (given 0 out of 17).

Sampling period: Full information.

Richness: Full information.

species groups considered: only non-managed bees

Abundance: Full information.

Visitation rate units: NA

Visitation rate: There are missing values (given 0 out of 17).

Yield units: NA

Yield: NA

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 area of each field [in hectares].

- If possible, please assign a management category [organic, IPM, conventional] to each field.
- If possible, check the latitude and longitude that were included in your `field_level_data` (csv) file. According to our automatic tests, some locations do not belong to the reported country/countries.
- 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.
- If possible, please, provide yield information.
- There are fields with two (25x25 m<sup>2</sup>) plots. Since we don’t know their corresponding Ids, we set to NA the sampling parameters, namely: total sampling area and total sampling time