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Automatic report

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Study ID: Alexandra\_Maria\_Klein\_Prunus\_dulcis\_USA\_2008

 $Contact: \ alexandra.klein@nature.uni-freiburg.de$ 

Credit: Claire Brittain, Claire Kremen and Alexandra-Maria Klein

Publication: 10.1111/gcb.12043

Number of sites: 23 Year of sampling: 2008 Crop: Prunus dulcis

Variety: There are missing varieties (given 0 out of 23).

Location: There are missing locations (given 0 out of 23).

Country: There are no locations to check.

Field size: There are missing field sizes (given 0 out of 23).

Management: Full information.

Sampling period: Full information.

Richness: Full information.

species groups considered: No taxa restrictions have been identified.

Abundance: There are missing values (given 0 out of 23).

Visitation rate units: visits per flower and second

Visitation rate: Full information.

Yield units: fruitset (%)
Yield: Full information.
Alternative yield units: NA

Alternative yield: NA

## 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+observdataset@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 latitude and longitude of the missing locations [in decimal degrees].
- 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.
- Please check that our information about the species groups considered is correct.
- Please, check that the brief description of your methodology (in your insect\_sampling file) is correct.