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Automatic report
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Study ID: Nicolas_J_Vereecken_several_crops_several_countries_several_years_3

Contact: Nicolas.Vereecken@ulb.be, Timothy.Weekers@ulb.ac.be

Credit: Nicolas J. Vereecken, Timothy Weekers

Publication: NA

Number of sites: 11

Year of sampling: 2019

Crop: Malus domestica

Variety: There are missing varieties (given 8 out of 11).

Location: Full information.

Country: All reported countries are OK.

Field size: Full information.

Management: Full information.

Sampling period: Full information.

Richness: There are missing values (given 8 out of 11).

species groups considered: only bees

Abundance: There are missing values (given 8 out of 11).

Visitation rate units: NA

Visitation rate: There are missing values (given 0 out of 11).

Yield units: kg/ha

Yield: There are missing values (given 2 out of 11).

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+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.

- 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.

- 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.

- There are sites without richness estimations. Please, check that such information is correct.

- There are sites without yield. Please, check that such information is correct. If possible, please provide the missing values (or the corresponding z-scores).

- Please, check that the brief description of your methodology (in your insect_sampling file) is correct.