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Automatic report
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Study ID: Rebecca_Steward_Fragaria_ananassa_Sweden_2014

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Credit: <https://snd.gu.se/en/catalogue/study/by-topic>

Publication: 10.1016/j.baae.2016.09.006

Number of sites: 30

Year of sampling: 2014

Crop: Fragaria x ananassa

Variety: There are missing varieties (given 0 out of 30).

Location: Full information.

Country: All reported countries are OK.

Field size: Full information.

Management: There are missing results for management (given 0 out of 30).

Sampling period: There are missing sampling periods.

Richness: There is not enough taxonomic resolution to estimate richness.

species groups considered: NA

Abundance: Full information.

Visitation rate units: NA

Visitation rate: There are missing values (given 0 out of 30).

Yield units: number/treatment/site

Yield: Full information.

Alternative yield units: pollination index of strawberry

Alternative yield: There are missing values (given 27 out of 30).

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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.
• If possible, please assign a management category [organic, IPM, conventional] to each field.

- 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 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.
- 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.
- There are sites without their alternative yield values. 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.