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Automatic report
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Study ID: Dara_St Stanley_Brassica_napus_Ireland_2010
Contact: darastanley@gmail.com
Credit: Dara Stanley NUI Galway, Jane Stout (Trinity College Dublin)
Publication: 10.1007/s10841-013-9599-z, 10.1007/s11258-014-0301-7
Number of sites: 3
Year of sampling: 2010
Crop: Brassica napus
Variety: There are missing varieties (given 0 out of 3).
Location: Full information.
Country: All reported countries are OK.
Field size: There are missing field sizes (given 0 out of 3).
Management: Full information.
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 3).
Yield units: seed set
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 area of each field [in hectares].

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