

=====
Automatic report
=====

Study ID: Simon_Potts_Vicia_faba_UK_2005

Contact: s.g.potts@reading.ac.uk

Credit: Simon Potts (University of Reading)

Publication: 10.1016/j.agee.2009.05.001

Number of sites: 10

Year of sampling: 2005

Crop: Vicia faba

Variety: There are missing varieties (given 0 out of 10).

Location: Full information.

Country: All reported countries are OK.

Field size: There are missing field sizes (given 0 out of 10).

Management: Full information.

Sampling period: There are missing sampling periods.

Richness: Full information.

species groups considered: only bees

Abundance: Full information.

Visitation rate units: NA

Visitation rate: There are missing values (given 0 out of 10).

Yield units: mean weight per plant

Yield: Full information.

Alternative yield units: NA

Alternative yield: NA

=====
Queries
=====

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.

- =====
 - Please check that credit information is correct and add the corresponding affiliations and acknowledge-ments/funding information in your 'Data_ownership' (excel) file.
 - If your study is already published, please check that its DOI is correct. CORRECT
 - If possible, please provide the names of missing crop varieties. ADDED
 - If possible, please provide the area of each field [in hectares]. NOT AVAILABLE
=====

- 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). ADDED
- 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. DATA NOT COLLECTED
- Please, check that the brief description of your methodology (in your insect_sampling file) is correct. CORRECT
- We found differences between this dataset (in Dainese et al., 2019) and the one from Garratt (in Kleijn et al. 2015). Please, confirm that the referred datasets are different. THESE WERE DIFFERENT STUDIES