

¹ RMAVIS v1.0: a R Shiny application for the analysis of
² vegetation survey data and assignment to GB NVC
³ communities.

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Authors of papers retain copyright and release the work under a Creative Commons Attribution 4.0 International License ([CC BY 4.0](#)). In the Great Britain (GB) the development of computational methods and programs for the assignment of vegetation survey data to NVC communities began with the development of TABLEFIT ([Hill, 1989](#); [Marrs, 2019](#)) and was followed by MATCH ([Malloch, 1998](#)). The most recent program, the Modular Analysis of Vegetation Information System (MAVIS), was developed as a windows application in 2000 and was a DEFRA-funded output of the ECOFACT project ([Bunce et al., 1999](#)), with the latest version released in 2016 ([Smart et al., 2016](#)).

Summary

RMAVIS is a R Shiny application for the assignment of vegetation sample plot data to British National Vegetation Classification (NVC) communities ([Rodwell, 1991, 1992a, 1992b, 1995, 2000](#)).

The assignment of vegetation sample plot data to established vegetation classification units using computational methods is a well established and recognised practice ([Maciejewski et al., 2020](#)). The results of this assignment process are used in various ways, including assisting in the phase 2 habitat survey (or NVC survey) ([Rodwell, 2006](#)) process; establishing an ecological baseline and identifying important ecological features such as protected habitats ([CIEEM, 2022](#)); and in ecological restoration by providing a proxy for historical reference ecosystems to target and against which to measure restoration progress ([Gann et al., 2019; Pywell et al., 2002; Sturbois et al., 2023](#)).

Statement of need

The requirement for a new program in the form of a R Shiny ([Chang et al., 2024](#)) application arises from several needs, namely to: 1) accommodate updates to the NVC; 2) provide a means to easily reproduce and attribute the results of the NVC assignment process; 3) broaden and simplify access through the provision of a web-based Graphical User Interface (GUI); and 4) facilitate the continuous development of such NVC assignment software. We developed RMAVIS considering these needs, with the view to providing a reliable system for use by the GB ecology and conservation community, analogous to the Engine for Relevés to Irish Communities Assignment (ERICA) tool ([Perrin et al., 2018; Perrin, 2019](#)) developed for Ireland.

37 Application structure

38 Inspired by the extensible structure of the species niche and distribution modelling R shiny
 39 application wallace (Kass et al., 2023), we constructed RMAVIS with a modular architecture,
 40 enabling both the easy maintenance of existing modules, and easy development of additional
 41 modules in the future.

42 RMAVIS currently contains a total of eighteen modules, with the fourteen main modules
 43 summarised in the following table.

Module	Description
Sidebar	Acts as the control module, containing the options for each module.
Data Input	Facilitates the entry of data, three methods are provided: manual, upload, and example.
Data Validation	Checks the format of the inputted data and provides means to adjust incorrectly formatted entries.
Data Structure	Checks the structure of the inputted data, displaying the availability of data by species and number of plots per year and group.
NVC	Displays the results of the NVC assignment process.
Assignment	
Habitat Correspondence	Displays the habitats from alternative habitat classifications associated with the top-fitted NVC communities.
Floristic Tables	Allows the comparison of floristic tables composed from the inputted data with the NVC's floristic tables.
Frequency	Summarises the frequency of occurrence of each species across all plots over time.
EIVs	Displays the mean Hill-Ellenberg ecological indicator values (EIVs) for each plot, group of plots, and all plots.
Diversity	
MVA National	Displays a number of diversity metrics for each plot and group of plots. Constructs an ordination space using all NVC pseudo-quadrats, with sample plots added in passively.
MVA Local, restricted	Constructs an ordination space using the top-fitted NVC communities pseudo-quadrats, with sample plots added in passively.
MVA Local, unrestricted	Constructs an ordination space using the top-fitted NVC communities pseudo-quadrats and sample plots.
Report	Provides the user with a downloadable report, containing user-selected outputs from the app session.



Figure 1: The MVA module of RMAVIS, showing the trajectory of all sample plots from the Leith Hill Wood example dataset in the ordination spaces over time.

44 Data sources and dependencies

45 RMAVIS depends on a number of individual datasets. The list of accepted species was constructed
 46 using the vascular plant (*Tracheophyta* and *Pteridophyta*), moss (*Bryophyta*), liverwort
 47 (*Marchantiophyta*), and hornwort (*Anthocerotophyta*) taxa present in the UKSI, retrieved
 48 from the National Biodiversity Network (NBN), ([NBN Trust, 2024d, 2024e, 2024b, 2024c, 2024a](#))
 49 filtered to include taxa at the species, species hybrid, species aggregate, subspecies,
 50 species *sensu lato*, and genus ranks; and the limited number of lichen (*Lecanoromycetes*)
 51 and charophyte (*Charophyta*) taxa, along with one unspecified 'algae' taxon present in the
 52 NVC floristic tables. The NVC communities present in RMAVIS are composed from Rodwell
 53 ([1991](#)), Rodwell ([1992a](#)), Rodwell ([1992b](#)), Rodwell ([1995](#)), Rodwell ([2000](#)), Wallace & Prosser
 54 ([2017](#)), Prosser et al. ([2023](#)), and Wallace et al. ([2023](#)). Data for habitat correspondences
 55 is derived from: the Joint Nature Conservation Committee (JNCC) Spreadsheet of Habitat
 56 Correspondences ([JNCC, 2008](#)), UKHab V1.1 ([Butcher et al., 2020](#)), and the National Plant
 57 Monitoring Scheme (NPMS) habitat correspondences ([Pescott et al., 2019](#)). Data for the
 58 vascular plant Hill-Ellenberg values were retrieved from the BSBI checklists, specifically the
 59 Nitrogen Score (N) ([BSBI, 2017c](#)), Moisture Score (F) ([BSBI, 2017d](#)), Reaction Score (R)
 60 ([BSBI, 2017a](#)), Salinity Score (S) ([BSBI, 2017b](#)), and Light Score (L) ([BSBI, 2017e](#)) checklists.
 61 The corresponding data for bryophytes was taken from BRYOATT ([Hill et al., 2007](#)). Four
 62 example datasets are bundled with RMAVIS: 1) Parsonage Down ([Riddig et al., 2020](#)), 2) Leith
 63 Hill Place Wood ([Smart et al., 2024; Wood et al., 2015](#)), 3) Whitwell Common ([Smart, 2000](#)),
 64 and 4) Newborough Warren ([Wallace & Jones, 2018](#)).

65 RMAVIS depends on the rhandsontable ([Owen, 2021](#)), reactable ([Lin, 2024](#)), bslib ([Sievert
 66 et al., 2024](#)), shinyWidgets ([Perrier et al., 2024](#)), shinyjs ([Attali, 2023](#)), shinybusy ([Meyer
 67 & Perrier, 2024](#)), rmarkdown ([Allaire et al., 2023](#)), bookdown ([Xie, 2024](#)), tidyverse ([Wickham
 68 et al., 2019](#)), plotly ([Sievert, 2020](#)), kableExtra ([Zhu, 2024](#)), writexl ([Ooms, 2024](#)),
 69 and vegan ([Oksanen et al., 2022](#)) R packages.

70 Conclusion

71 RMAVIS provides a web-based, easily accessible GUI for the assignment of vegetation sample plot
 72 data to GB NVC communities. RMAVIS also provides a number of other exploratory analyses,

⁷³ which compliment the NVC assignment results. We plan to maintain, optimise, and expand
⁷⁴ the functionality found in v1.0 of RMAVIS and hope that it acts as a useful tool for the GB
⁷⁵ ecology and conservation community.

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