

## Additions and enhancements to BRAHMS for v 7.2 and 7.3.

The following additions to BRAHMS are now available.

WEB TOOLBARS

TAG HIGHLIGHTING

TEXT REPORTER

PRIVATE FIELDS

FORMATTING TOOLS FOR REVISIONS

EXPORT OPTIONS

CONSERVATION ASSESSMENTS

UPDATE FROM RDE

CRIA DATA SUBMISSIONS

HOST PLANTS

BARCODE SEARCHING ON TOOLBAR

LIVING COLLECTIONS

EXTRACT FILE CONTROL

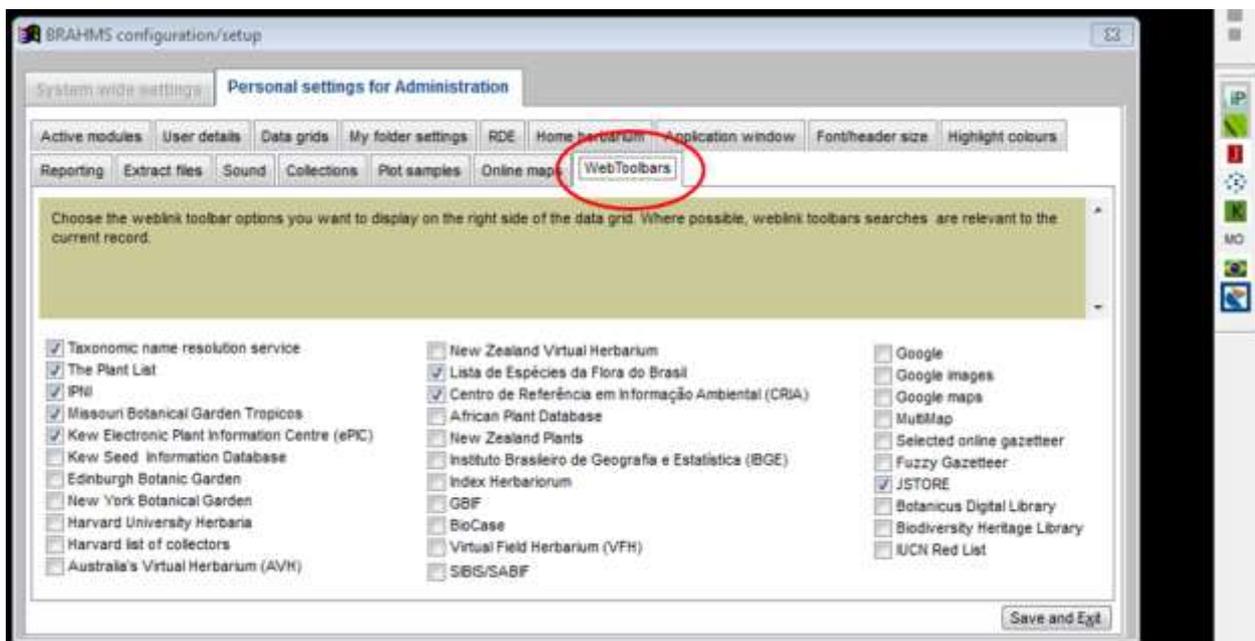
LATITUDE GRADIENTS

MATCHING RECORDS IN SECOND DATABASE

SOFTWARE SPEED ON NETWORKS

### Web Toolbars

A greater range of web link toolbars is now available and you can select the ones you wish to be active in the right side toolbar. Further options are being added.



## Tag highlighting

You can now display different tag highlight colours using characters or numbers as defined on **Utilities > My setup > Highlight colours**.

Click to select colours	Default/suggested colours
<input type="button" value="Tag highlight * or 1"/>	<input type="button" value="Restore default"/>
<input type="button" value="tag highlight \$ or 2"/>	<input type="button" value="Restore default"/>
<input type="button" value="tag highlight ! or 3"/>	<input type="button" value="Restore default"/>
<input type="button" value="Tag highlight = or 4"/>	<input type="button" value="Restore default"/>
<input type="button" value="Tag highlight + or 5"/>	<input type="button" value="Restore default"/>
<input type="button" value="Column highlight"/>	<input type="button" value="Restore default"/>
<input type="button" value="Zoom window colour"/>	<input type="button" value="Restore default"/>

tag	del	mergeto	snumber	taxstat	syncal	synnote	synof	synlot	family	genus
			0	5342	acc		memo	0	Pinaceae	Pinus
			0	5355	acc		memo	0	Cupressaceae	Juniperus
1			0	5381	acc		memo	0	Pinaceae	Larix
2			0	5424	acc		memo	0	Pinaceae	Pinus
3			0	5508	acc		memo	0	Taxaceae	Torreya
4			0	5514	acc		memo	0	Taxaceae	Torreya
5			0	5531	acc		memo	0	Cupressaceae	Cupressus
			0	5532	acc		memo	0	Cupressaceae	Juniperus
*			0	5541	acc		memo	0	Cupressaceae	Cupressus
\$			0	5555	acc		memo	0	Cupressaceae	Juniperus
!			0	5558	acc		memo	0	Pinaceae	Abies
=			0	5560	acc		memo	0	Taxaceae	Amentotaxa
+			0	5565	acc		memo	0	Pinaceae	Pinus

## Text reporter

A heavily revised text reporter, now separated from the visual reporter with its own toolbar.

Documentation on text reporting is provided in

<http://herbaria.plants.ox.ac.uk/bol/Content/Groups/brahms/Resources/Textreporting.pdf>.

The screenshot shows a window titled "Editing report: Sample taxa revision (c:\brahms\data-conifers\myreports/sample\_taxa\_revision.dbf)". It features a menu bar with "Quick summaries", "Text reports", "Edit template", and "Help". Below the menu is a table with columns: Position, Lines before, Text before, Print code, Spaces, Field name, Print code, Caps, Decimal places, Text after, Print code, Spaces, Lines after, Suppress identical, and Exc. The table lists fields like "tout.family", "tout.fullname", "tout.synnum", "tout.descr", "tout.distr", "linked ecology", and "tout.speclat".

Below the table is a configuration panel with several sections:

- Buttons:** "Add new line", "Delete line".
- File-field name:** "tout", "speclat", "Print setting": "None", "Add to current".
- Options:** "Print data in CAPS", "Exclude item if empty/null", "Suppress identical (only print when value changes)".
- Only print when:** A text input field.
- Lines before/after:** Input fields for "Lines before" and "Lines after".
- Text before/after:** "Material examined:-", "Print setting": "None", "+ spaces": "1".
- Remove 'Text after' after the final item:** A checkbox.
- Sort data for report using command:** A large text input area.
- Expression grouping:** "Group expression in position 25 with one in position", "Text at group start", "Text at group end".
- Max grid:** A button.
- Bottom bar:** "Generate report", "Tagged only", "File/document (default)", "Exit".

*Sample text report template being edited.*

## Private fields

Private fields can now be set to visible but read-only rather than just visible. Private fields can only be edited by the user who owns these fields. To enable the use of private fields, select **Admin > Project configuration > System wide settings > PrivateFields**.



## Update from RDE

RDE botanical record files can now be submitted to update existing records rather than import as new. Additional specimens can be added from RDE to botanical records already stored in BRAHMS.

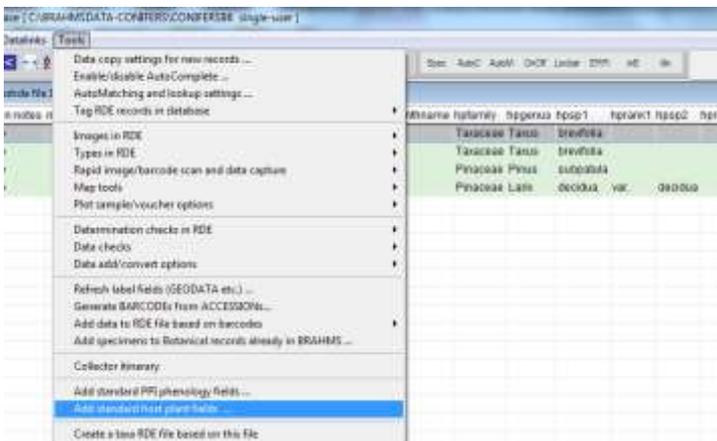
## CRIA data submissions

Improved export option for uploading to CRIA <http://www.cria.org.br/> (Brazil projects).



## Host plants

Host plant species names can be added to RDE using a new set of fields HPfamily, HPgenus, etc. An RDE tools option is provided to add the host plant fields. Useful for bryophytes, lichens, fungi, parasitic plants and the like.



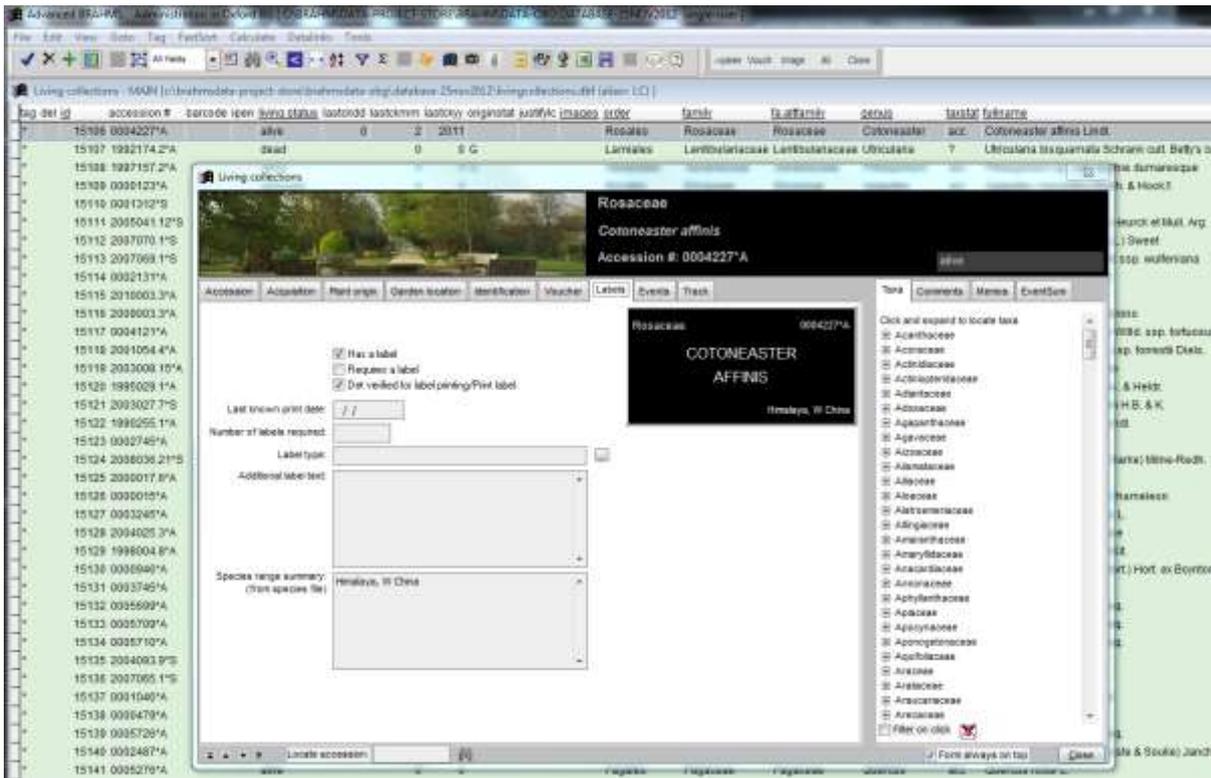
## Barcode searching on toolbar



A new fast access barcode search in the main botanical record and specimen files.

## Living collections

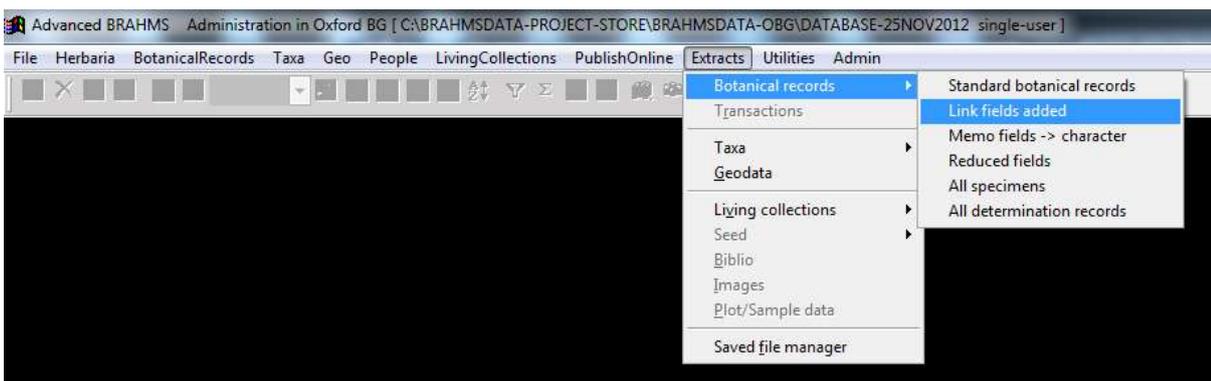
The Living Collection module has been upgraded with more options available in the events file.



*Living collections file with form*

## Extract file control

Extract files now provide separate menu options to open extracts with link fields added, memo fields converted to character, and other options as relevant to the data category.



## Latitude gradients

Function to calculate latitude gradient bands in a selected band width added to the Botanical Records extract file (Tools > Data summaries and indexes > Latitude band scan).



Example map for all conifers with taxa and genera totals calculated in 10 degree bands

## Matching records in second database

A new set of functions to locate matching Botanical Records in separate BRAHMS databases and optionally update specimens in one database from another. See

[http://herbaria.plants.ox.ac.uk/bol/content/documentation/MatchingBotanicalRecords\\_Jan2013.pdf](http://herbaria.plants.ox.ac.uk/bol/content/documentation/MatchingBotanicalRecords_Jan2013.pdf)

Advanced BRAHMS Administration in INPA [ C:\INPA\DATABASE single-user ]

File Edit View Goto Tag FastSort Calculate Datalinks Tools

Collection extract file [ c:\tempfiles-brahms-1\inpa\extracts\collextract.dbf (alias= COUT) ]

tag	dbid	brahms	collector	number	day	month	year	family	species
INPA		85910	Carreira, L.M.M.	287	27	1	1983	Leguminosae-mim	Senegalia multipinnata
*	HAMAB	85910	Carreira, L.M.M.	287	27	1	1983	Leguminosae-mim	Acacia multipinnata
INPA		17465	Carreira, L.M.M.	304	27	1	1983	Rutaceae	Spathelia excelsa
*	HAMAB	17465	Carreira, L.M.M.	304	27	1	1983	Rutaceae	Sohnreyia excelsa
INPA		87298	Carreira, L.M.M.	319	28	1	1983	Leguminosae-mim	Inga pilosula
*	HAMAB	87298	Carreira, L.M.M.	319	28	1	1983	Leguminosae-mim	Inga nitida
INPA		21041	Carreira, L.M.M.	328	28	1	1983	Piperaceae	Piper divaricatum
*	HAMAB	21041	Carreira, L.M.M.	328	28	1	1983	Piperaceae	Piper divaricatum

The above screen shows an example sort. The INPA database has been match with HAMAB. The HAMAB BRAHMS numbers have been edited to be the same as the matched INPA records to allow updating. NB the original HAMAB BRAHMS number is stored in the MYSTRING memo as this can then be used to draw further data including specimens from that database. You can see, even if this small example, differences are being picked up with determinations.

An example list of matched records form two databases.

## Software speed on networks

Inconveniently slower speeds are sometimes encountered on certain types of local network when running BRAHMS in multiple user mode. The slow speeds can be noted when using data grids, usually in the main species and people files but potentially elsewhere. Slower speeds are not noted in single user mode or when using RDE, even on the same network.

A typical networked installation of BRAHMS will have the software and database folder installed on a shared drive – e.g. N:\BRAHMS\BRAHMS7 and N:\BRAHMS\BRAHMSDATA-MYDATA\DATABASE. Several workstations will have access to these folders. The tempfiles-brahms folders are normally stored on the local workstation – e.g. C:\TEMPFILES-BRAHMS-A.

The symptoms of slow speed in multiple user mode are a) slow opening of files to display data grids b) a general stickiness of the cursor moving within the grid c) slow tag and filtering functions d) slow Fastsort operations.

If you encounter such problems, you can greatly speed up BRAHMS performance in general by selecting some or all of the following options:

### Field views:

Using field views to reduce the number of columns displayed in data grids speeds up performance. Files such as the species file include many fields that you will not need day to day. Create a field view (using the  toolbar) that removes these. You can probably reduce the number of fields by at least 50% without the loss of routine editing tasks. Note that you can set your preference on Utilities > My setup/profile > Data grids > Restore last used field view.

### Disable dynamic formatting:

Dynamic formatting is when a row colour or font is edited in relation to something in that row, for example a different colour when a record is tagged. This formatting can slow up speeds as you can experiment by disabling one or more of these options by choosing:

- Utilities > My setup/profile > Data grids > Highlight tagged records : de-select
- Utilities > My setup/profile > Data grids > Disable strikeouts font : select
- Utilities > My setup/profile > Data grids > Do not highlight accepted names: select

### Private fields

The use of private fields (fields that can only be seen and/or edited by selected users) can cause a slowing of grid opening. See Admin > Project configuration > System wide settings > PrivateFields.