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Using JDemetra+ in R: from version 2 to version 3 Presentation 3: Wrangling workspaces in R

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- 1. Introduction
- 2. Using workspaces and R packages in conjuction
- 3. Instant reading
- 4. Automating operations
- 5. Conclusion

Context

- a workspace is a JD+ specific data structure (xml files) which allows to use the GUI and cruncher
- in R we don't need this stucture to run an SA process (cf P2)
- we could think sa processing with JD+ algorithms can happen in two disntict worlds
 - GUI and cruncher using workspaces OR
 - R functions (without workpaces, TS object or numeric vectors for HF)

Still there are benefits in using workspaces and R packages in conjunction We will highlight them in the remaining of this presentation

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Using workspaces and R packages in conjuction

Two (?) classes of benefits

- instant reading: immersing results in the R world (functions, plots..)
- automating potential manual operations

De manière générale, aucune compatibilité entre v2 et v3

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Instant reading

v2 : RJDemetra::load_workspace() -> creation of RJD object (java object pointer) v3 : rjdemetra3::load_workspace() -> readable R object (list format)

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Automating operations

On me voit?

Automating operations on workspaces

Goal: to produce and to reproduce the workspace structure without the JDemetra+ graphical interface but with R and keeping the read, write and refresh properties both by the graphical interface and by the cruncher.

Cruncher and GUI react the same way to a WS: - if the structure is not good: they won't be able to read anything - if the structure is minimal: they will manage to only read and compute the SA-items - if the ws are well-structured and have all the necessairy information: they will succeed to read, compute and refresh with the chosen policy our sa-items.

Outline of the automation

Presentation in v2

operations - creation - export / write - modifications / dynamic update

Creation

```
RJDemetra::new_workspace(): creation of workspace
RJDemetra::new_multiprocessing(): creation of multiprocessing
```

```
To create a spec, there is several functions: RJDemetra::x13_spec(), RJDemetra::tramoseats_spec(), ... and also in v3 with packages rjd3tramoseats and rjd3x13
```

To create an SA-item, there is several functions depending on the chosen spec: RJDemetra::x13(), RJDemetra::tramoseats(), ... and also in v3 with packages rjd3tramoseats and rjd3x13

Finally you can use RJDemetra::add_sa_item() to add the created SA-item to a multprocessing in your workspace

Export

The function RJDemetra::save_workspace() export and create a physical workspace with a structure in folder and xml files. But it only creates what the previous functions constructed.

For example, the function RJDemetra::x13()

Dynamic update

Attention, the exported workspace

Automating operations on series

("functions that operate on one workpace") - series object - spec - meta data in v2 RJDemetra in v3 rjdworkpace and other rjd3...

Merging workspaces

("functions that operate two workpaces") context: annual rev in v2 RJDemetra in v3 rjdworkpace and other rjd3...

Preserving desirable properties ?

two features to preserve - GUI readable - Crunchable When are the preserved and when not ?

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Summary of functionnalities

- available
- missing

Package's perimeter and ressources

 V2 vs V3: not clear cut yet not many features are available in sheer v3 rjdworkspace (predating v3) functions will be (partly) integrated into rjdemetra3 package