

Q&A CanReg Webinar 6 – Customization and Management

11.12.2012

Q: The [analytical] scripts are written in R?

A: Yes.

Q: Can we develop our own modules?

A: Yes. How to extend the analytical capabilities of CanReg5 using R scripts is outlined in Appendix E of the CanReg5 [handbook](#).

Q: When I merge two patients into one and I want to keep the two tumours but only one name how do I do to delete one of the two names? Is there an easier and faster way for merging a high number of duplicates found without having to refer the ID number in each merge process?

A: You can merge two patients into one, consolidate the “best” information into one record, and delete the other. (More on this in webinar 4 - on data entry.) As of now, there is no faster way to merge patients, except exporting the data and cleaning it outside CanReg5 (using any tool that can work on tabular data) before importing it again.

Q: Can we export the data to Excel sheets?

A: Yes – please see webinar 5 for more information on that.

Q: Will the central database be online?

A: Each CanReg5 system is by default independent. You enter your cases and have full control over where they are stored. You can choose to host a CanReg5 server online and let client machines connect over standard TCP/IP, but you need to provide the security layers yourself. (VPN/SSH)

Useful links:

- The handbook: <http://www.iacr.com.fr/CanReg5/CanReg5-instructions.pdf>
- GICR Resources page: http://gicr.iarc.fr/index.php?page=resources&lang=1&item_id=4
- IACR: <http://www.iacr.com.fr/>