

### **C.1.1 Educational goals of the modules**

### **C.1.2 Module subject material**

We propose to develop two collections of modules, **Improving the Reproducibility of Experimental Data Recording** and **Improving the Reproducibility of Experimental Data Pre-Processing**.

The **Improving the Reproducibility of Experimental Data Recording** collection will include the following modules:

1. The principals of “tidy” data
2. Creating spreadsheet templates for experimental data collection
3. Example spreadsheet template: A template for collecting CPU data for a tuberculosis study [make more specific]
4. Choosing a spreadshet program for reproducible data collection: Excel, Google Sheets, and RStudio
5. Organizing data recording and meta-data recording through RStudio “Projects”
6. Creating “Project Templates” for consistency across projects
7. Example of creating a “Project Template”: A project template for a lab group studying tuberculosis [make more specific]
8. Harnessing version control (git and GitLab) to improve transparency in data recording
9. Using git from RStudio
10. Using GitLab for version controlled collaborations

The **Improving the Reproducibility of Experimental Data Pre-Processing** collection will include the following modules:

1. An introduction to R code scripts
2. The benefits of scripts for data pre-processing
3. Getting started with RMarkdown
4. The relationship between R code scripts and RMarkdown documents
5. Creating reproducible data pre-processing protocols using Rmarkdown
6. Example of a reproducible data pre-processing protocol: Automated gating for flow cytometry data
7. Example of a reproducible data pre-processing protocol: Measuring metabolite feature intensities for metabolomics LC/MS data
8. Complex data types in R and their use in Bioconductor packages
9. Converting from complex data types to “tidy” formats for data analysis and visualization with R’s “tidy” data tools