EX. NO: 7

INSTALLATION GUIDE FOR R AND R STUDIO

Step 1 - Install R

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1. Download the R installer from https://cran.r---project.org/

What are R and CRAN?

The Comprehensive R Archive Network

Figure 1. Screenshot of http://cran.csiro.au/

2. Run the installer. Default settings are fine. If you do not have admin rights on your laptop, then ask you local IT support. In that case, it is important that you also ask them to give you full permissions to the R directories. Without this, you will not be able to install additional packages later

Step 2 – Install RStudio

R Studio Products Pricing About Us Blog HD H Do you need support or a commercial license? Check out our commercial offerings RStudio Desktop 0.99.467 — Release Notes RStudio requires R 2.11.1 (or higher). If you don't already have R, you can download it here. **Installers for Supported Platforms** RStudio 0.99.467 - Windows Vista/7/8 73.9 MB 2015-07-15 5c0bf6987adcfb6dd441326ecc67f6e8 RStudio 0.99.467 - Mac OS X 10.6+ (64-bit) 56.2 MB 2015-07-15 RStudio 0.99.467 - Ubuntu 12.04+/Debian 8+ (32-bit) Oca919255495cc87112df12a1cff7e29 RStudio 0.99.457 - Ubuntu 12.04+/Debian 8+ (64-bit) 83.9 MB 2015-07-15 dd64fc165de55a0be229f2362cd776da RStudio 0.99.467 - Fedora 19+/RedHat 7+/openSUSE 13.1+ (32-bit) 76.8 MB 2015-07-15 le152bafa8b6c5355a2ec0f6822abdff RStudio 0.99.467 - Fedora 19+/RedHat 7+/openSUSE 13.1+ (64-bit) 77.6 MB 2015-07-15 a82a27b113184e1790ec5bd3c36e2137 Zip/Tarballs Date 105.5 MB 2015-07-15 b88bbbc042aaa04b83a1786322981c5f RStudio 0.99.467 - Windows Vista/7/8 2015-07-15 171de4b2e08edd33988ab1b75d571ae6 RStudio 0.99.467 - Ubuntu 12.04+/Deblan 8+ (32-bit) 78.1 MB RStudio 0.99.467 - Ubuntu 12.04+/Deblan 8+ (64-bit) 84.8 MB 2015-07-15 85ca4bc15c2e5e535594bcf586604324 RStudio 0 99.467 - Fedora 19+/RedHat 7+/openSUSE 13.1+ (32-bit) 77.4 MB 2015-07-15 6abc51b45cdd1fa0c14a991b5336a8ad RStudio 0.99.467 - Fedora 19+/RedHat 7+/openSUSE 13.1+ (64-bit) 78.4 MB 2015-07-15 9d693e5e5ed1b2003e1b18691f31574b

1. Download RStudio: https://www.rstudio.com/products/rstudio/download/

Figure 2. Download RStudio on https://www.rstudio.com/products/rstudio/download/

A tarball containing source code for RStudio v0.99.467 can be downloaded from here

- 2. Once the installation of R has completed successfully (and not before), run the RStudio installer.
- 3. If you do not have administrative rights on your laptop, step 2 may fail. Ask your IT Support or download a pre---built zip archive of RStudio which doesn't need installing. The link for this is towards the bottom of the download page, highlighted in Image 2.
 - a. Download the appropriate archive for your system (Windows/Linux only the Mac version can be installed into your personal "Applications" folder without admin rights).
 - Double clicking on the zip archive should automatically unpack it on most Windows machines.

Step 3 – Check that R and RStudio are working

Source Code

- 1. Open RStudio. It should open a window that looks similar to image 3 below.
- 2. In the left hand window, by the '>'sign, type '4+5' (without the quotes) and hit enter. An output line reading '[1] 9' should appear. This means that R and RStudio are working.
- 3. If this is not successful, contact us or your local IT support for further advice

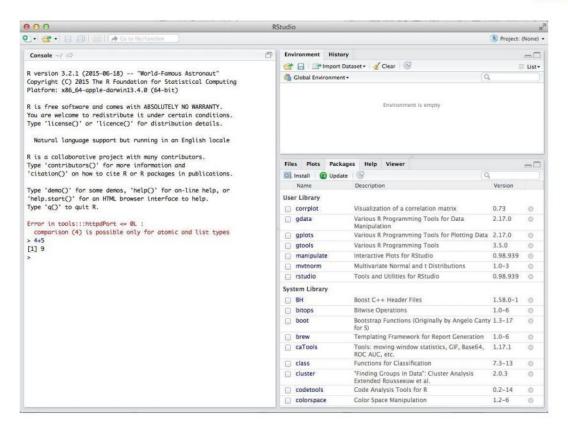


Figure 3. Running R with RStudio

Step 4 – Install R packages required for the workshop

- Click on the tab 'Packages' then 'Install' as shown in Image 4. Or Tools ---> Install packages.
- Install the following packages: mixOmics version 6.1.0, mvtnorm, RColorBrewer, corrplot, igraph (see Image 4). For apple mac users, if you are unable to install the mixOmics imported library rgl, you will need to install the XQuartz software first https://www.xquartz.org/
- 3. Check that the packages are installed by typing 'library(mixOmics)' (without the quotes) in the prompt and press enter (see Image 5).
- 4. Then type 'sessionInfo()' and check that mixOmics version 6.1.0 has been installed (image 6).



Figure 4. Click on Install to install R packages.



Figure 5. Specify the list of packages to be installed

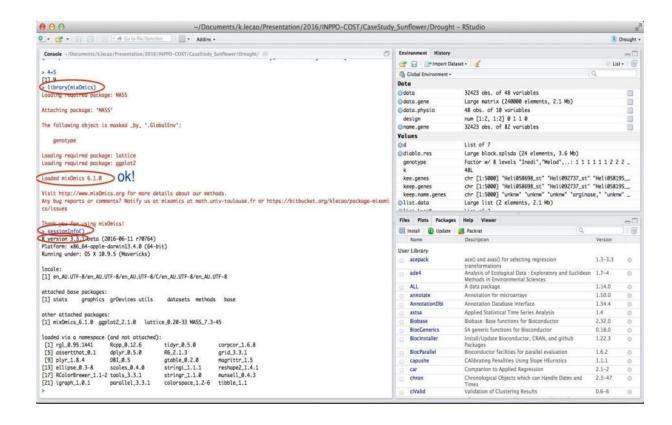


Figure 6. Check that the package mixOmics is installed and has the version 6.1.0.