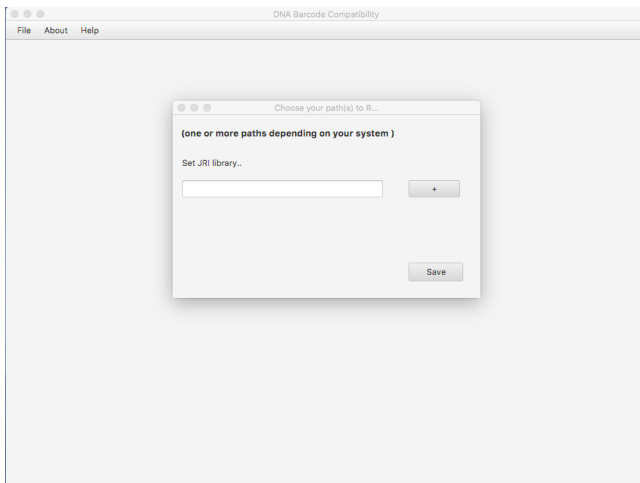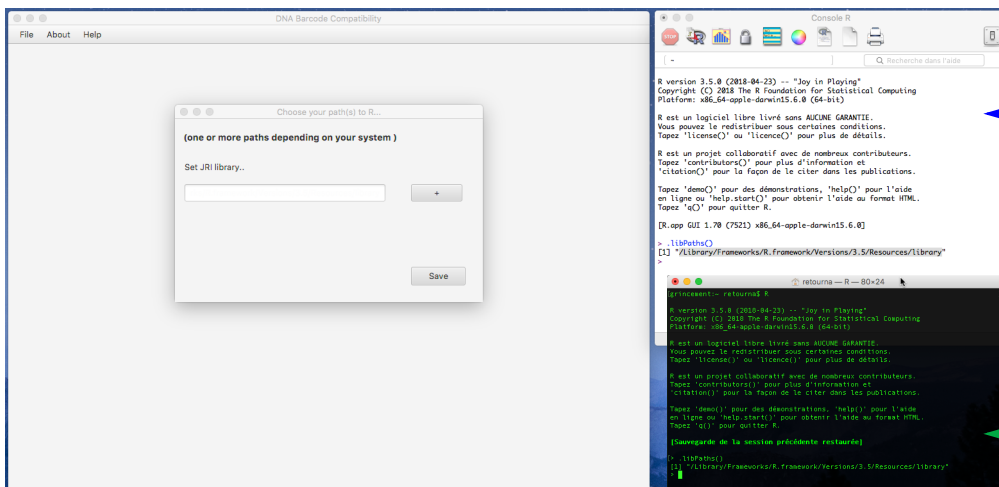# Configure the DNABarcodeCompatibility interface to enable the communication with the R environment

**1**

When you first start the DNABarcodeCompatibility user graphical interface, a window pops up to ask you for the R library paths.
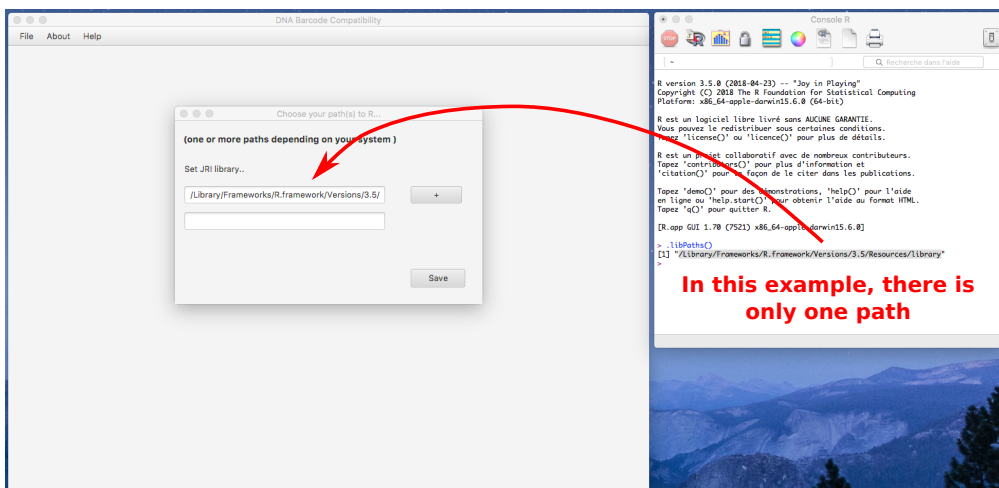
**2**

On Windows and MacOSx, clicking the R application opens an R console in which to type in **.libPaths()** to list all possible library paths.

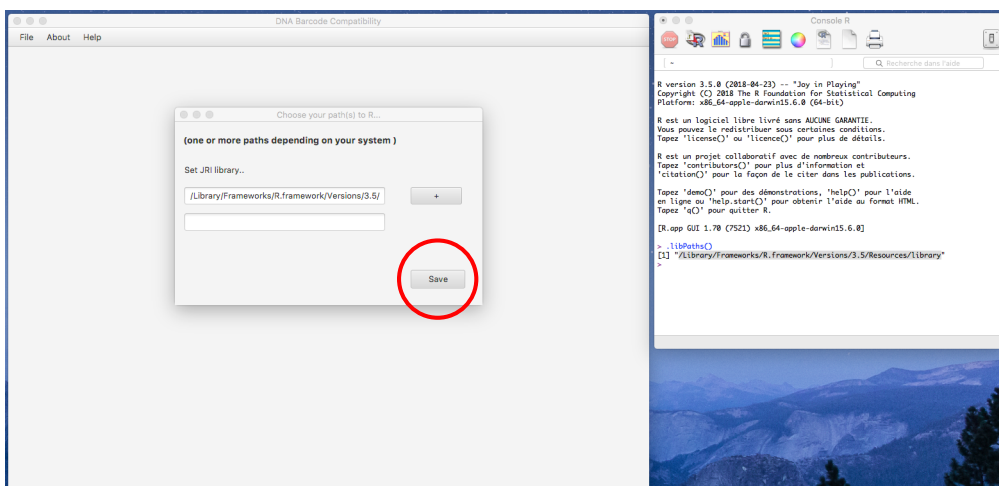Alternatively, on Linux and MacOSx, run R within a terminal and type in **.libPaths()**

**3**

Copy and paste one-by-one all paths **without quotes** into each empty field of the pop up window

**In this example, there is only one path**

In this pop up window, you can add en empty field by clicling on the "+" icon.

**4**

Click the save button and then close the pop up explicitly