

Tassel and R Integration

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Calling R from Tassel

First, install rJava package as described below.

Run the command

```
> system.file('jri',package="rJava")
```

For Java development (i.e. Tassel), use the Java API provided (<http://www.rforge.net/JRI/index.html>).

Once rJava has been installed, a folder named "library/rJava/jri" can be found in the R home directory. To find the R home directory, type `R.home()` on the command line in R. In the jri folder is a file named "run", which is a script that can be used to set values for environment variables and call java, so that `java.library.path` and `classpath` are set so that the jar files and native library can be found by REngine. The simplest way to use R is to always use this script to start TASSEL.

A more flexible method is to make sure that the jri library file (`libjri.jnilib` on a Mac, a similarly named `.dll` file on Windows) is in a folder on the `java.library.path`. On a Mac, this can be accomplished by putting a copy or a link in `/Library/Java/Extensions`. On Windows, the folders listed in the `PATH` environment variable are the `java.library.path`. Alternatively, if `<jri>` = the full path to the jri folder then adding `-Djava.library.path = <jri>` to the java command line will add that folder to the library path.

In addition, the `R_HOME` environment variable must be set to the R home directory prior to running the java command line. From a terminal command line, the command "R RHOME" returns the R home directory. Probably the most practical way to enable the use of R without much user intervention is to add commands to the start scripts to test for the existence of R and the jri library, then set `R_HOME` and use `-D.library.path` to add the jri folder to the library path. From within TASSEL, code should test for value of the `R_HOME` environment variable before attempting to call R.

The examples subdirectory of jri contains `rtest.java`, which has examples showing how to pass commands to R and how to move data to and from R.

Calling Tassel from R

```
> install.packages("rJava")
```

```
Installing package(s) into '/Library/Frameworks/R.framework/Versions/2.15/Resources/library'  
(as 'lib' is unspecified)
```

```
trying URL 'http://ftp.ussg.iu.edu/CRAN/bin/macosx/leopard/contrib/2.15/rJava_0.9-3.tgz'
```

```
Content type 'application/x-gzip' length 791689 bytes (773 Kb)
```

```
opened URL
```

```
=====
```

```
downloaded 773 Kb
```

The downloaded binary packages are in

```
/var/folders/Jy/JybbW3n5GcORrFKVfrqYWk+++TI/-Tmp-//RtmpWosQU6/downloaded_packages
```

```
> library(rJava)
```

```
> .jinit()
```

```
> .jaddClassPath("/Users/terry/terry/tassel4.0_standalone/sTASSEL.jar")
```

```
> .jaddClassPath("/Users/terry/terry/tassel4.0_standalone/lib/")
```

```
> .jclassPath()
```

```
[1] "/Library/Frameworks/R.framework/Versions/2.15/Resources/library/rJava/java"
```

```
[2] "/Users/terry/terry/tassel4.0_standalone/sTASSEL.jar"
```

```
[3] "/Users/terry/terry/tassel4.0_standalone/lib"
```

```
> obj=.jnew("net.maizegenetics.pipeline.TerryTests")
```

```
> result=.jcall(obj, "[[D", "runTest")
```

```
> mat=sapply(result,.jevalArray)
```

```
> mat
```

```
  [,1] [,2]
[1,]  0  1
[2,]  1  2
[3,]  2  3
```

```
/*
```

```
 * TerryTests
```

```
*/
```

```
package net.maizegenetics.pipeline;
```

```
/**
```

```
 *
```

```
 * @author terry
```

```
*/
```

```
public class TerryTests {
```

```
    public static double[][] runTest() {
```

```
        double[][] result = new double[2][3];
```

```
        for (int x = 0; x < 2; x++) {
```

```
            for (int y = 0; y < 3; y++) {
```

```
                result[x][y] = x + y;
```

```
            }
```

```
        }
```

```
        return result;
```

```
    }
```

```
}
```

Other Resources

Darren Wilkinson's Blog

<http://darrenjw.wordpress.com/2011/01/01/calling-java-code-from-r/>

rJava Manual

<http://cran.r-project.org/web/packages/rJava/rJava.pdf>

rJava Home Page

<http://www.rforge.net/rJava/>