

# Biocep-R

Open Science in the cloud, towards a  
universal platform for mathematical and  
statistical computing

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*Croire possible le souhaitable est aussi dangereux que de croire souhaitable le possible. Utopies sentimentales et automatismes de la technique.*

Nicolás Gómez Dávila

*Il n'y a que le solitaire qui soit capable de penser plus que des vérités tactiques.*

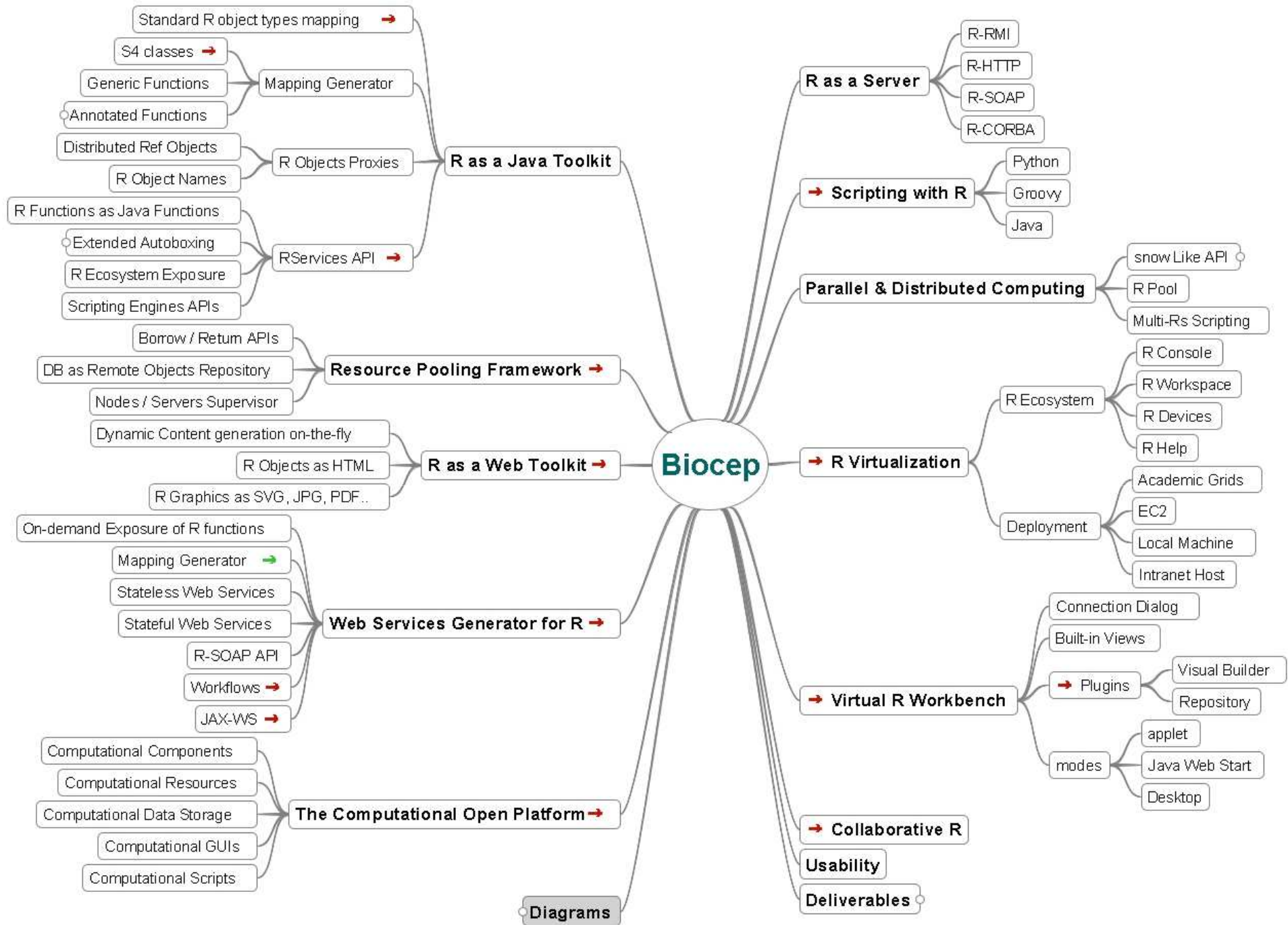
Nicolás Gómez Dávila

## **Extract from the GridSolve Description Document**

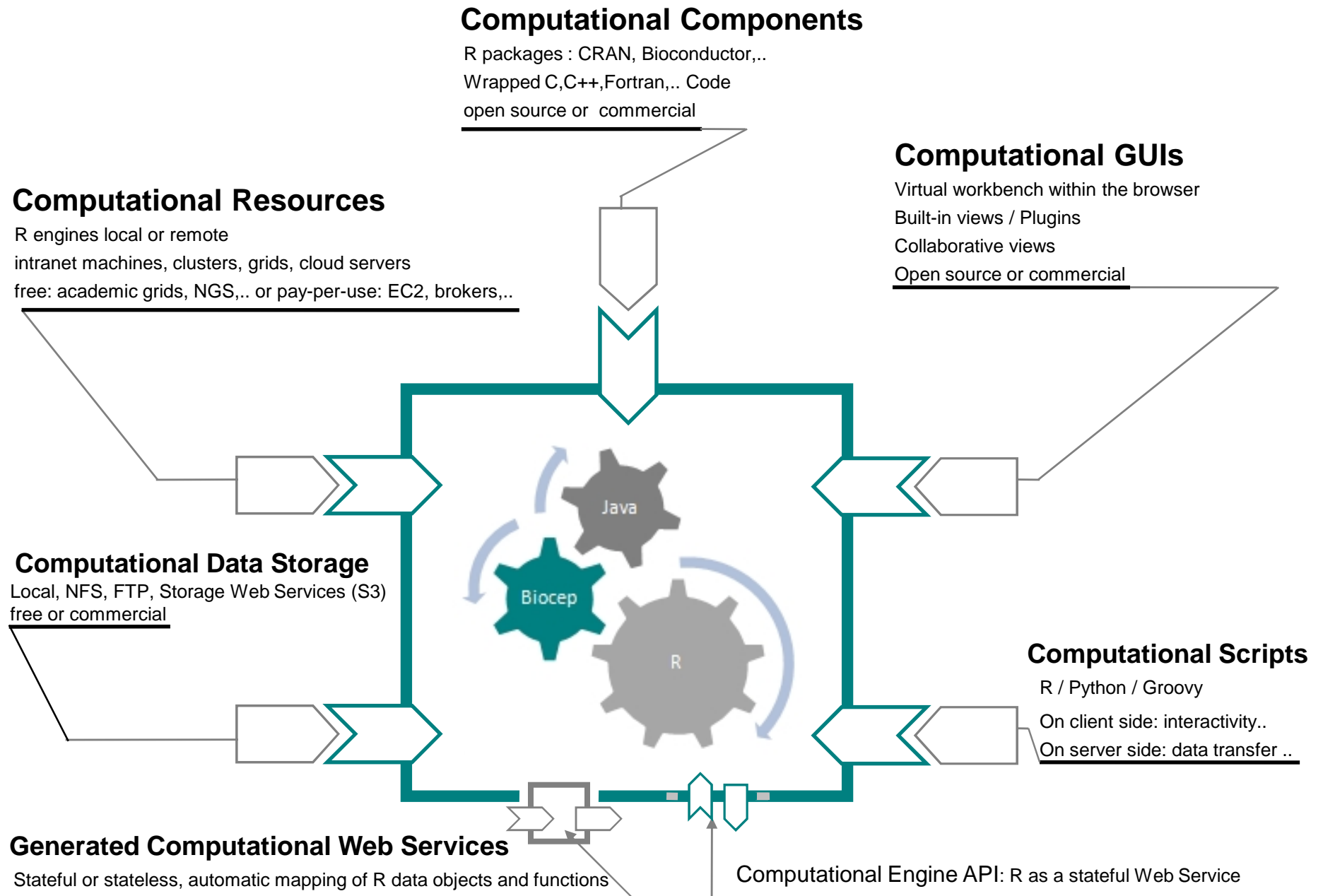
The emergence of Grid computing as the prototype of a next generation cyberinfrastructure for science has excited high expectations for its potential as an accelerator of discovery, but it has also raised questions about whether and how the broad population of research professionals, who must be the foundation of such productivity, can be motivated to adopt this new and more complex way of working.

The rise of the new era of scientific modeling and simulation has, after all, been precipitous, and many science and engineering professionals have only recently become comfortable with the relatively simple world of the uniprocessor workstations and desktop scientific computing tools. In that world, software packages such as Matlab and Mathematica represent general-purpose scientific computing environments (SCEs) that enable users — totaling more than a million worldwide — to solve a wide variety of problems through flexible user interfaces that can model in a natural way the mathematical aspects of many different problem domains.

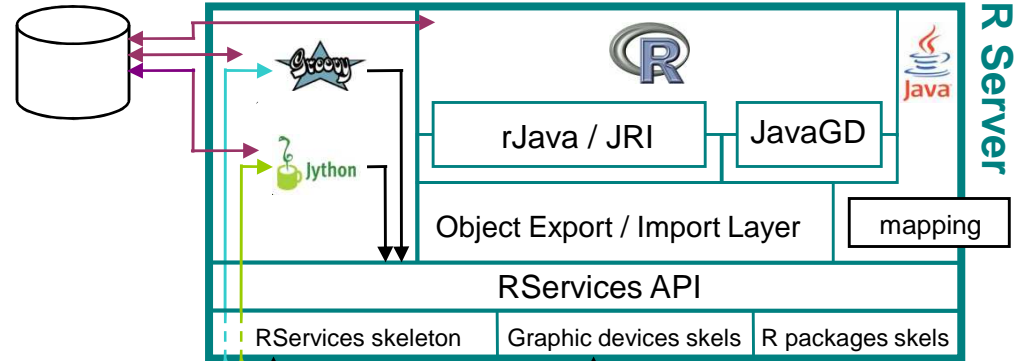
Moreover, the ongoing, exponential increase in the computing resources supplied by the typical workstation makes these SCEs more and more powerful, and thereby tends to reduce the need for the kind of resource sharing that represents a major strength of Grid computing [1]. Certainly there are various forces now urging collaboration across disciplines and distances, and the burgeoning Grid community, which aims to facilitate such collaboration, has made significant progress in mitigating the well-known complexities of building, operating, and using distributed computing environments. But it is unrealistic to expect the transition of research professionals to the Grid to be anything but halting and slow if it means abandoning the SCEs that they rightfully view as a major source of their productivity. We therefore believe that Grid computing's prospects for success will tend to rise and fall according to its ability to interface smoothly with the general purpose SCEs that are likely to continue to dominate the toolbox of its targeted user base.



# Biocep Computational Open Platform Ecosystem



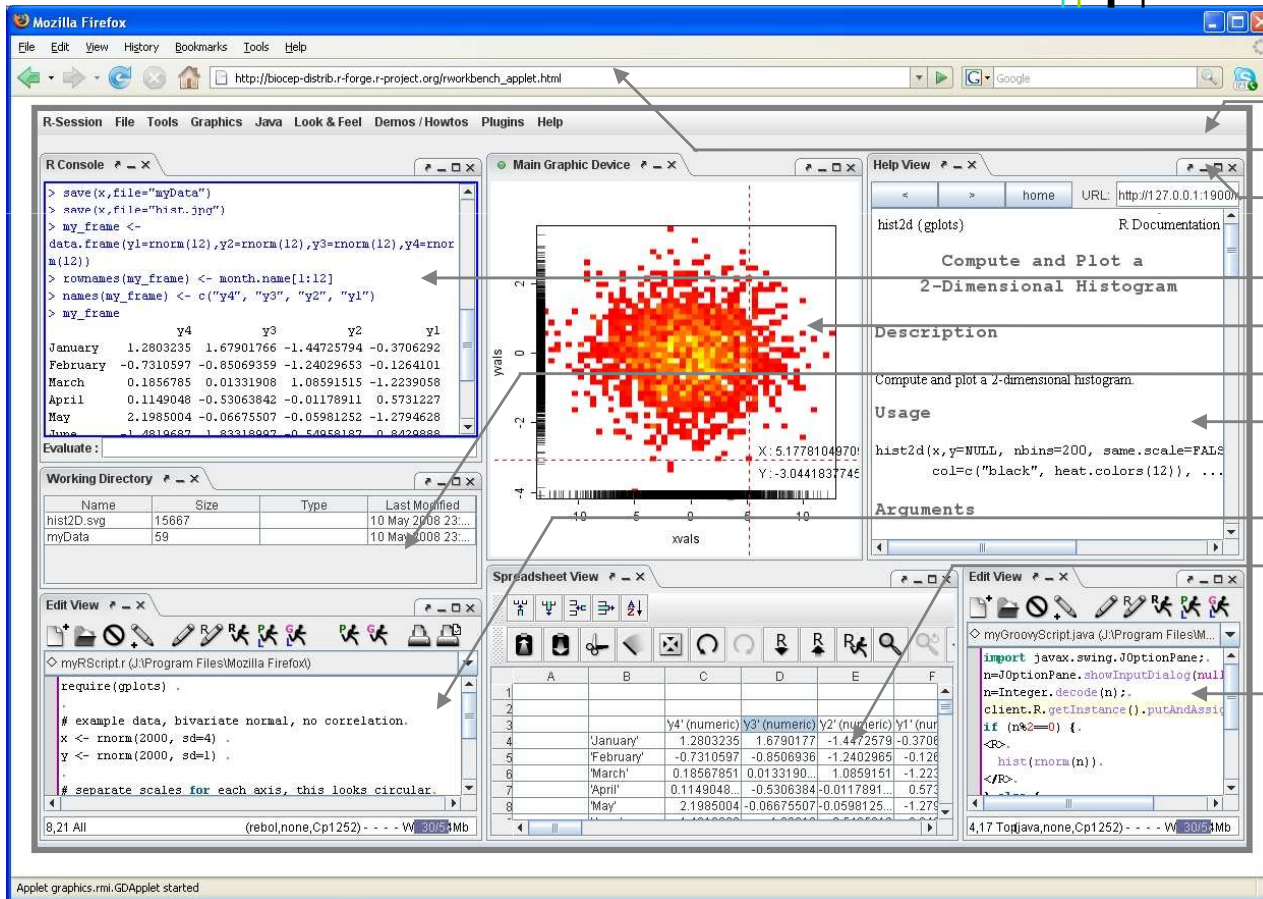
# R Virtualization



Server Side - Personal Machine, Academic Grids, Clusters, Clouds

Client Side - Internet

## Virtual R Workbench



Internet Browser

Java Applet

Virtual R Workbench URL

Docking Framework

R Console

R Graphic Device+Interactors

R Workspace

R Help Browser

R Script Editor

R Spreadsheet

Groovy / Jython Script Editor





# Integrating R - State of the art

- **SJava and rJava/JRI**
  - Basic mapping via JNI of the R C API
- **TypeInfo**
  - Plug meta descriptions to R functions
- **RWebservices**
  - Generated Java Beans for basic R Types / S4 Classes
  - Axis Web Services based on SJava and ActiveMQ
- **JavaGD**
  - R devices connection to Java (JGR)
- **Rserve**
  - TCP/IP interface to R



# What was missing ?

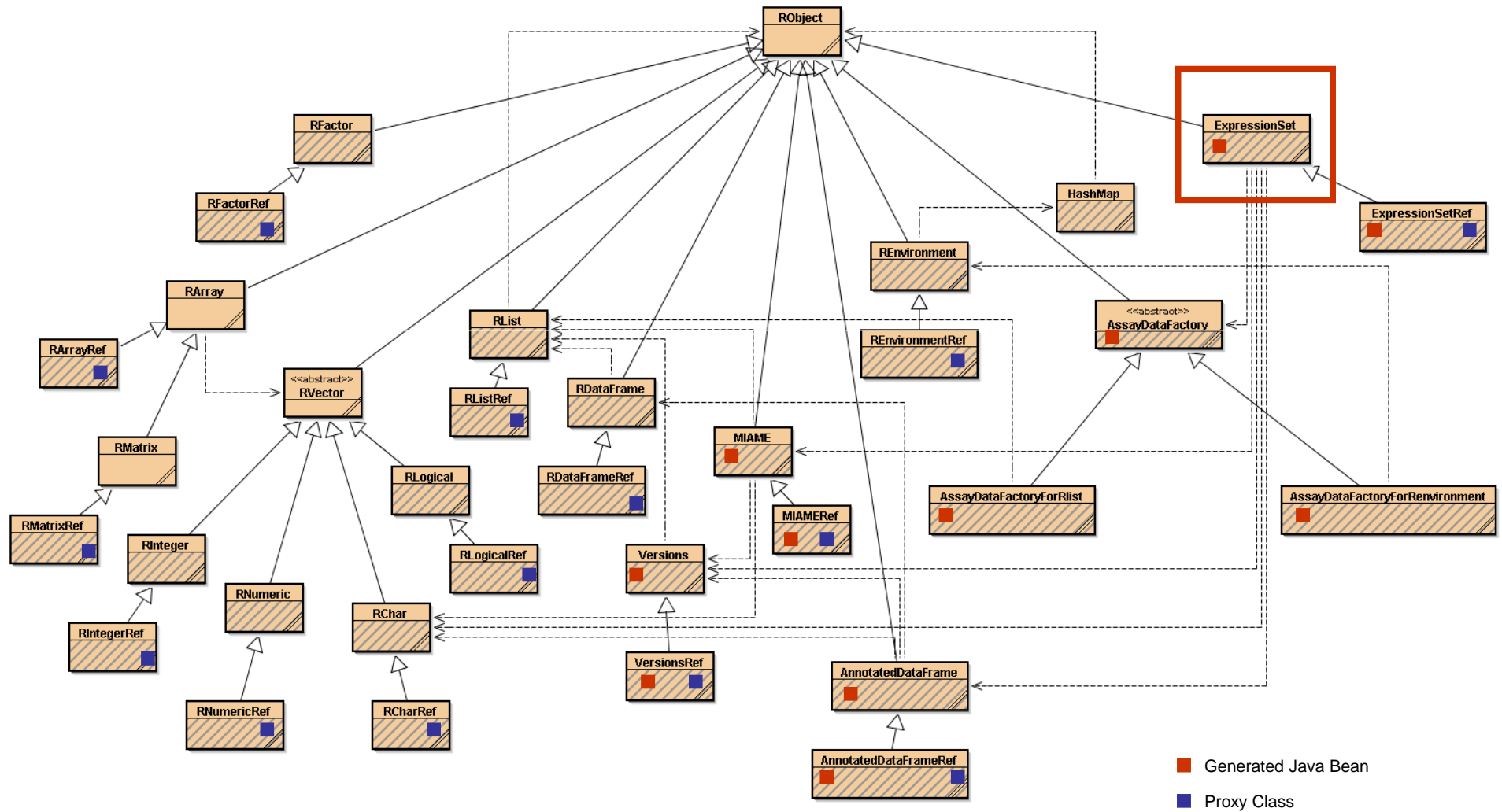
- High Level Java API for Accessing R
- Stateful, Resuable, Remotable R Components
- Scalable, Distributed, R Based Infrastructure
- Safe multiple clients framework for components usage as a pool of indistinguishable Remote Resources
- User friendly Interface for the remote resources creation, tracking and debugging

# What was missing ?

- Generated light-weight Java proxies for R Types / S4 Classes
- On-demand mapping and deployment of R packages as RMI Components or as JAX-WS Web Services
- Remotable R Graphics / Swing Components for R
- Remote R components files exchange API
- Semi-thick client (applet) for web based tools using R



# Generated beans for ExpressionSet



# RServices API - I

```
public interface RServices extends ManagedServant {
```

```
-----  
public String consoleSubmit(String expression) throws ...  
public String evaluate(String expression) throws ...  
-----
```

```
public RObject getObject(String expression) throws ...  
public Object getObjectConverted(String expression) throws ...  
public RObject getReference(String expression) throws ...  
public RObject getObjectName(String expression) throws ...  
-----
```

```
public void putAndAssign(Object obj, String name) throws ...  
public RObject putAndGetReference(Object obj) throws RemoteException;  
-----
```

```
public RObject call(String methodName, Object... args) throws ...  
public RObject callAndConvert(String methodName, Object... args) throws ...  
public RObject callAndGetReference(String methodName, Object... args) throws ...  
public RObject callAndGetObjectName(String methodName, Object... args) throws ...  
public void callAndAssign(String varName, String methodName, Object... args) throws ...  
-----
```

```
public RObject realizeObjectName(RObject objectName) throws ...  
public Object realizeObjectNameConverted(RObject objectName) throws ...  
public RObject referenceToObject(RObject refObj) throws ...  
-----
```

```
public boolean isReference(RObject obj) throws ...  
public void assignReference(String name, RObject refObj) throws ...  
-----
```

```
}
```

# RServices API - II

```
public interface RServices extends ManagedServant {
```

```
public String[] listPackages() throws ...  
public RPackage getPackage(String packageName) throws ...
```

```
public GDDevice newDevice(int w, int h) throws ...  
public GDDevice[] listDevices() throws ...
```

```
public interface GDDevice extends Remote {  
    public Vector<GDObject> popAllGraphicObjects() throws ...  
    public void fireSizeChangedEvent(int w, int h) throws ...  
    public void dispose() throws ...  
    ...  
}
```

```
public String[] getWorkingDirectoryFileNames() throws ...  
public FileDescription getWorkingDirectoryFileDescription(String fileName) throws...  
public void createWorkingDirectoryFile(String fileName) throws ...  
public void removeWorkingDirectoryFile(String fileName) throws ...  
public byte[] readWorkingDirectoryFileBlock(String name, long off, int size) throws...  
public void appendBlockToWorkingDirectoryFile(String name, byte[] block) throws...
```

```
public String getRHelpFileUri(String topic, String pack) throws ...  
public byte[] getRHelpFile(String uri) throws ...
```

```
public Vector<RAction> popRActions() throws ...
```

```
}
```

# RServices API - III

```
public interface RServices extends ManagedServant {
```

```
-----  
public void    startHttpServer(int port) throws ...  
public void    stopHttpServer() throws ...  
-----
```

```
public String  pythonExec(String pythonCommand) throws ...  
public RObject pythonEval(String pythonCommand) throws ...  
public void    pythonSet(String name, Object Value) throws ...  
-----
```

```
public String  groovyExec(String groovyCommand) throws ...  
public Object  groovyEval(String expression) throws ...  
public void    groovySet(String name, Object Value) throws ...  
-----
```

```
public void    setCallback(RCallback callback) throws ...  
-----
```

```
public String  getStatus() throws ...  
public void    stop() throws ...  
public void    freeReference(RObject refObj) throws ...  
public void    freeAllReferences() throws ...  
public String  print(String expression) throws ...  
public String  sourceFromResource(String resource) throws ...  
public String  sourceFromBuffer(StringBuffer buffer) throws ...  
public RNI     getRNI() throws ...  
-----
```

```
...  
}
```

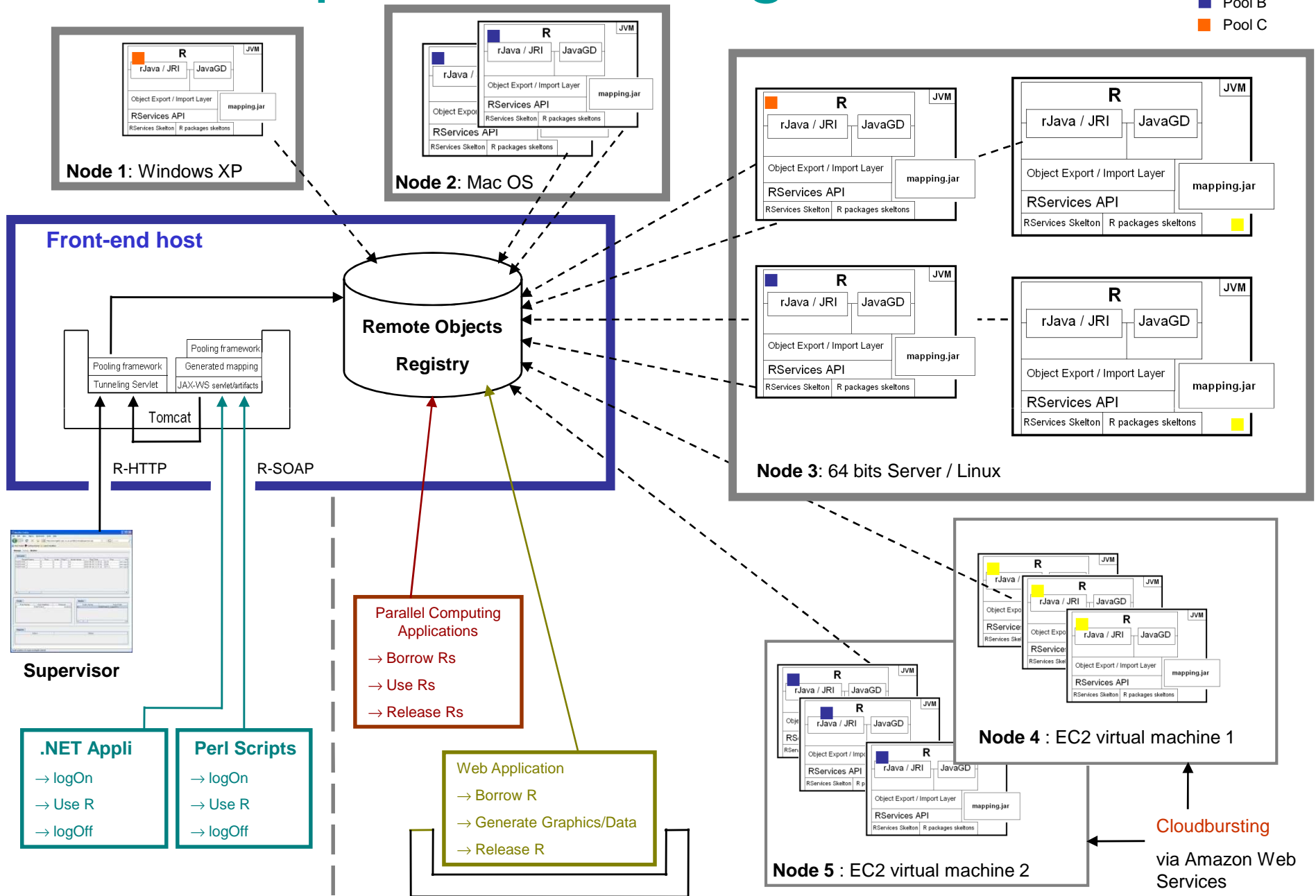


# Remote Resources Pooling Framework

- Generic Standalone framework
- Pooling of any RMI components and if combined with JNI of any library / open architecture
- New Remote Object Registry based on Derby| Oracle| MySQL
- Three implementations available
  - rmiregistry / mono-node / single client process
  - rmiregistry / multinodes / single client process
  - database ROR / multinodes / multiple client processes
- User friendly interface for the remote resources creation, tracking and debugging, nodes and pools management

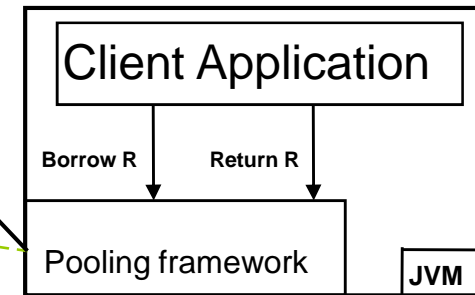
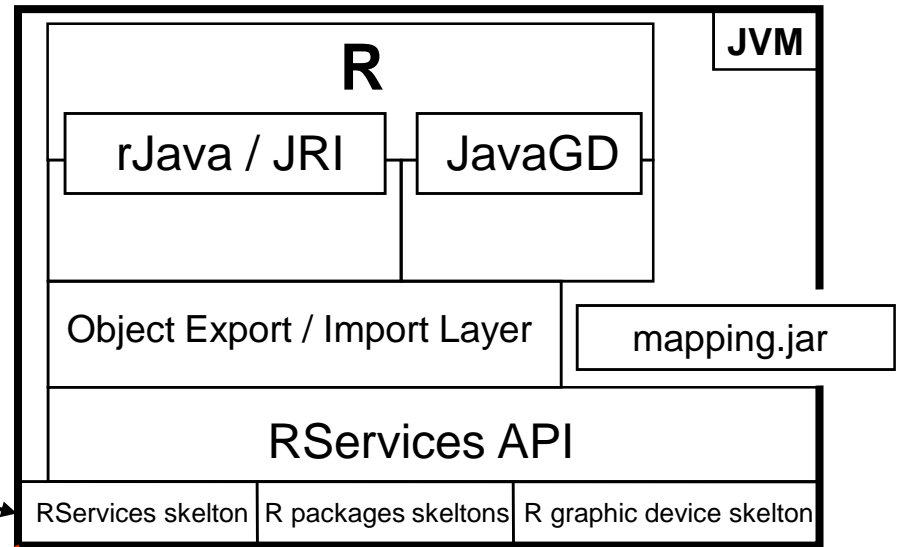
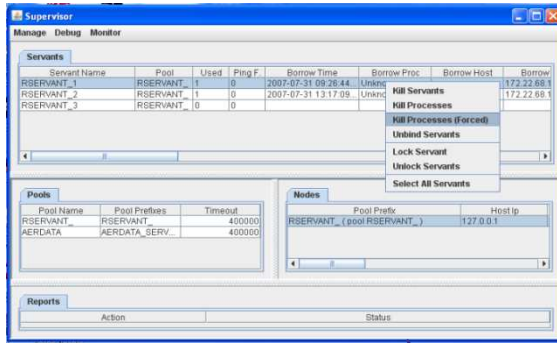
# Computational Engines Pools

■ Pool A  
■ Pool B  
■ Pool C

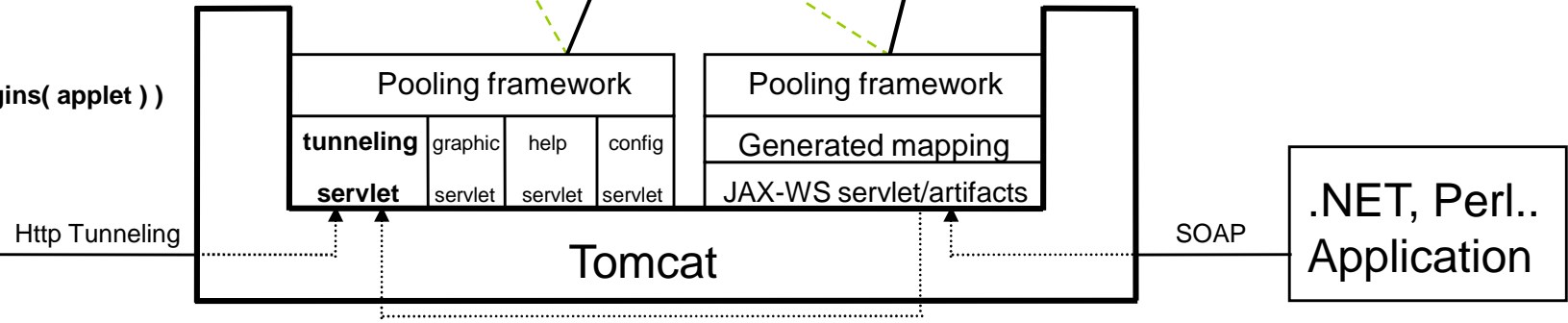
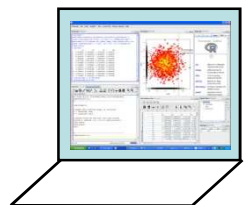


# R Pools

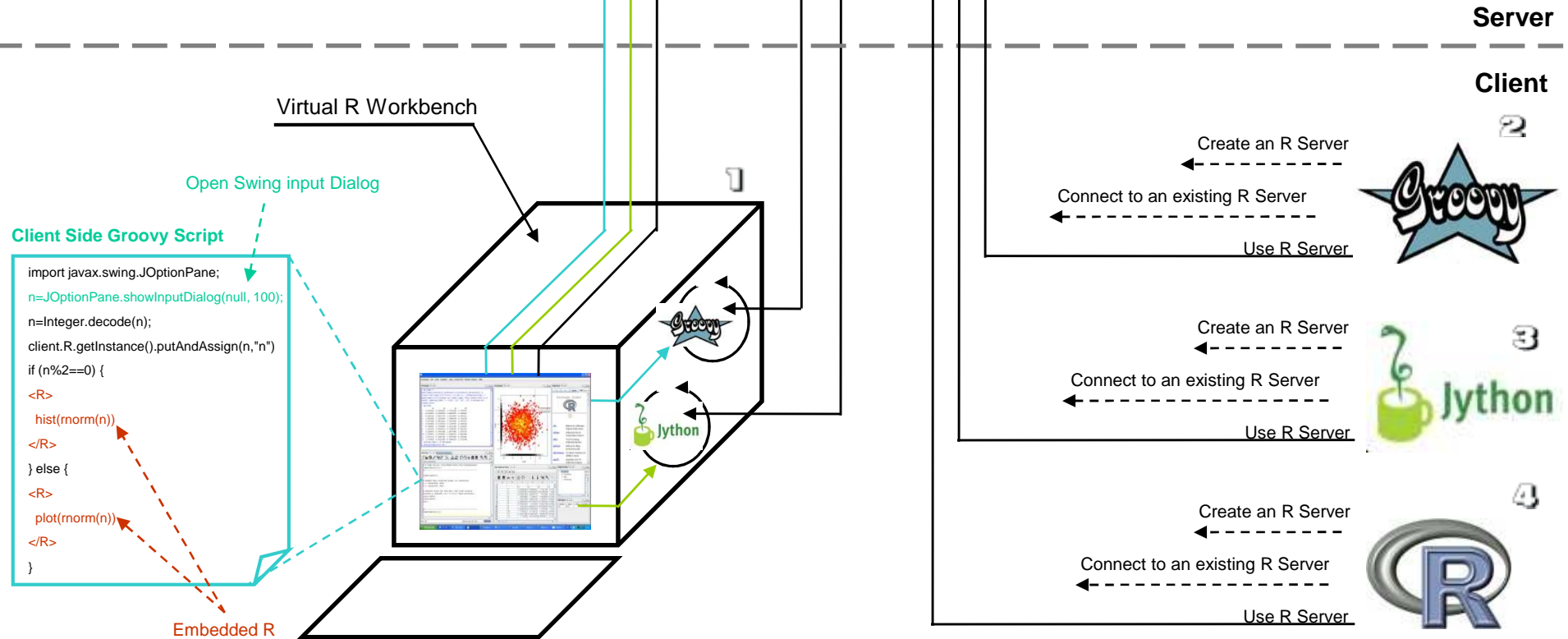
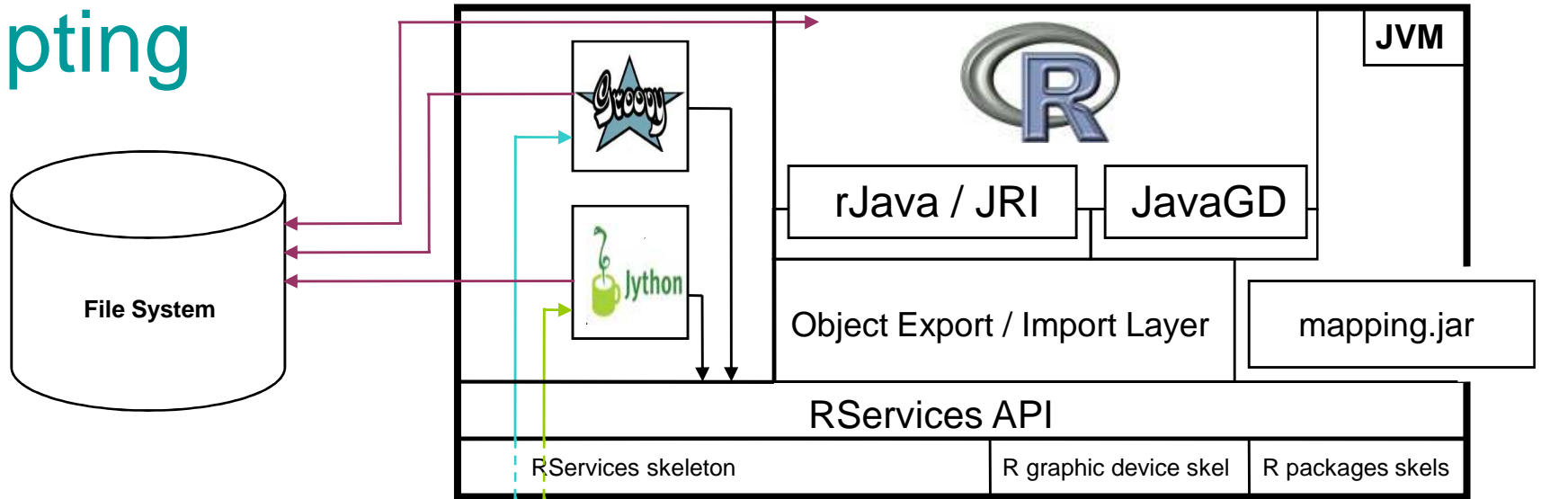
## Supervisor



## Browser( java plugins( applet ) )



# Scripting



## Client Side Groovy Script

```
import javax.swing.JOptionPane;
n=JOptionPane.showInputDialog(null, 100);
n=Integer.decode(n);
client.R.getInstance().putAndAssign(n, "n")
if (n%2==0) {
  <R>
  hist(rnorm(n))
  </R>
} else {
  <R>
  plot(rnorm(n))
  </R>
}
```

Embedded R

Open Swing input Dialog

Virtual R Workbench

Server

Client

2



3



4



# Parallel Computing

```
final double[][] m=..;
Future<Double>[] result=new Future[m.length];
ExecutorService exec = Executors.newFixedThreadPool(50);
for (int i=0; i<result.length; ++i) {
final double[] v=m[i];
result[i]= exec.submit(
    new Callable<Double>() {
public Double call() throws Exception {
    RServices r=null;
    try {
        r=(RServices)ServantProviderFactory.getFactory().getServantProvider().borrowServantProxy();
        Rnumeric mean=(RNumeric)r.call("mean", new RNumeric(v));
        return mean.getValue()[0];
    } finally { ServantProviderFactory.getFactory().getServantProvider().returnServantProxy(r); }
    }
    });
}
while(true) {
int count=0; for (int i=0; i<result.length; ++i) if (result[i].isDone()) ++count; if (count==result.length) break;
Thread.sleep(100);
}
for (int i=0; i<result.length; ++i) System.out.println(result[i].get());
```

# Snow with Biocep

*From the R Console :*

- **makeCluster**(n,...) **stopCluster**(cl)  
→ Starting and Stopping clusters
- **clusterEvalQ**(cl, expr)  
→ The expression is evaluated on the slave nodes.
- **clusterApply**(cl, seq, fun, ...)  
→ Calls the function with the first element of the list on the first node, with the second element of the list on the second node, and so on.
- **clusterExport**(cl, list)  
→ Assigns the global values on the master of the variables named in 'list' to variables of the same names in the global environments of each node.

...

# Web Services Generation

## Script / globals.r

```
square ← function(x) {return(x^2) }
typeInfo(square) ← SimultaneousTypeSpecification(
TypedSignature(x = "numeric"), returnType = "numeric")
```

## Script / rjmap.xml

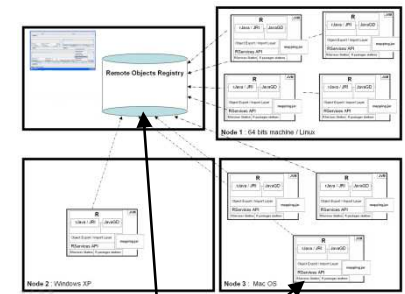
```
<rj>
<publish>
<functions> <function name="square" forWeb="true"/> </functions>
</publish>
<scripts> <initScript name="globals.r" embed="true"/> </scripts>
</rj>
```

WS generator

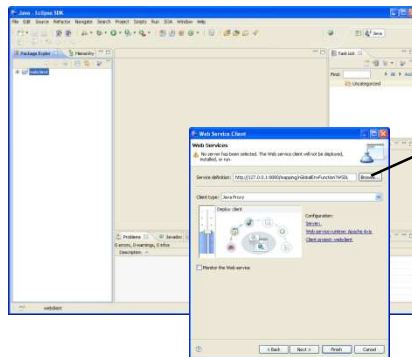
## rws.war

- + mapping.jar
- + pooling framework
- + R Java Bridge
- + JAX-WS
- Servlets
- Generated artifacts

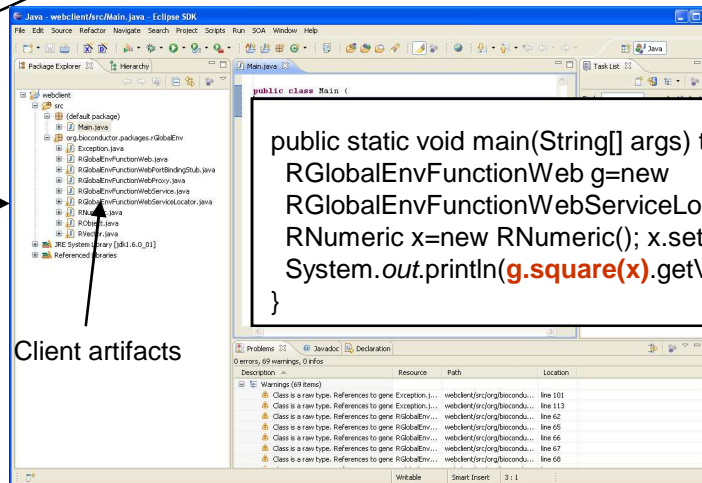
Deploy



<http://127.0.0.1:8080/rws/rGlobalEnvFunction?WSDL>



Eclipse Web Service Client Generator

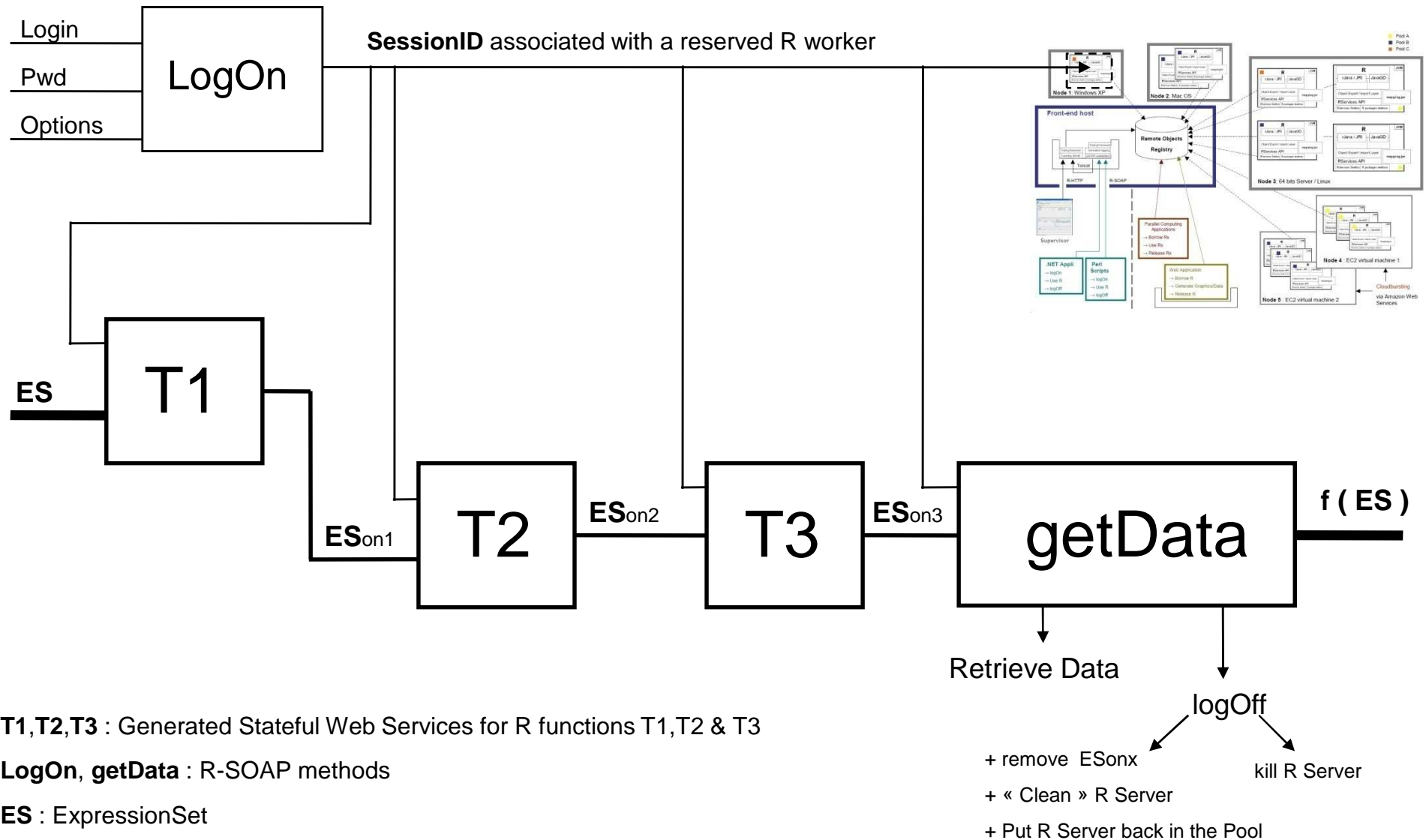


Client artifacts

```
public static void main(String[] args) throws Exception {
RGlobalEnvFunctionWeb g=new
RGlobalEnvFunctionWebServiceLocator().getRGlobalEnvFunctionWebPort();
RNumeric x=new RNumeric(); x.setValue(new Double[] {6.0});
System.out.println(g.square(x).getValue()[0]);
}
```



# Workflows with Stateful Web Services



**T1,T2,T3** : Generated Stateful Web Services for R functions T1,T2 & T3

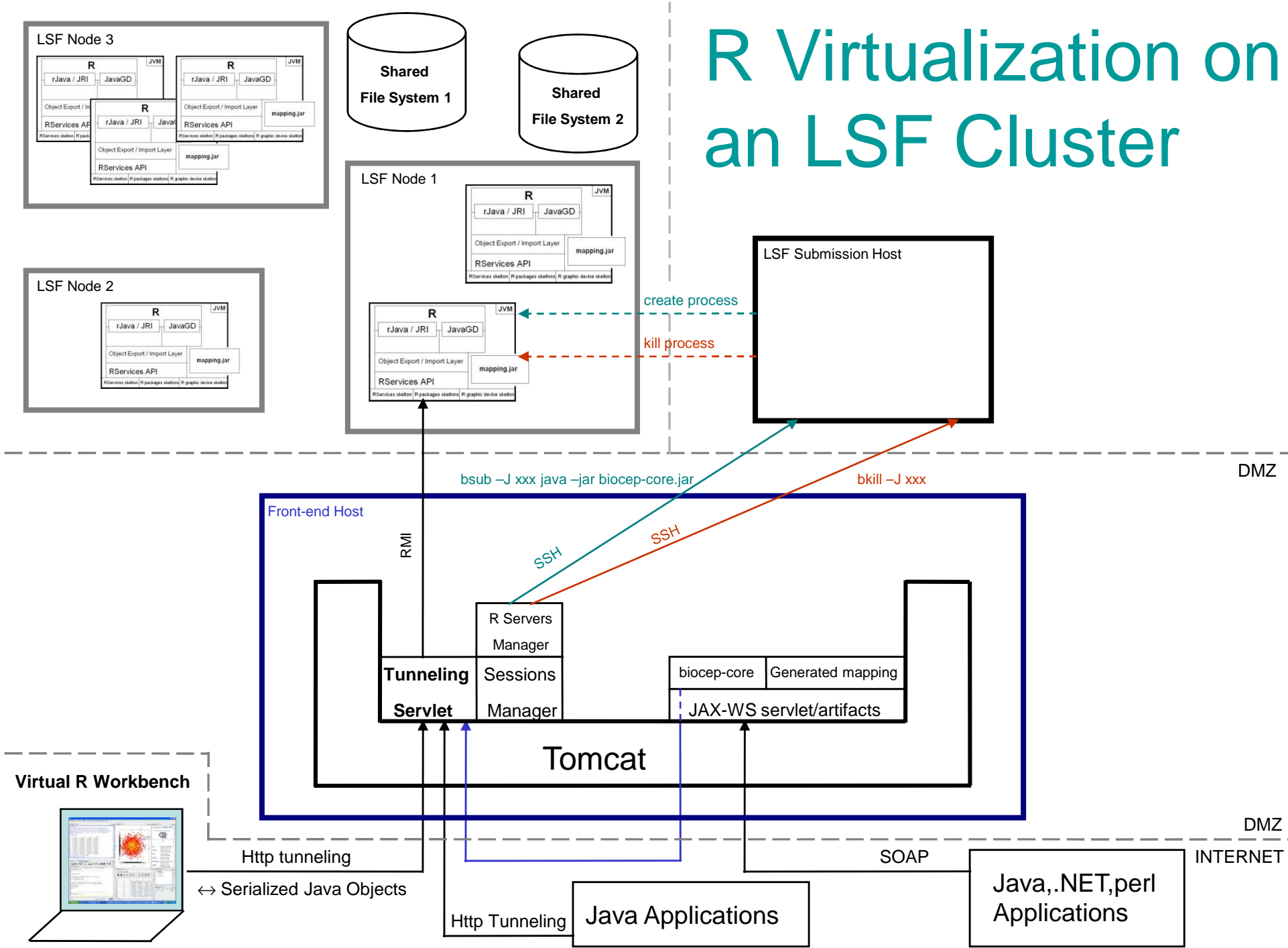
**LogOn, getData** : R-SOAP methods

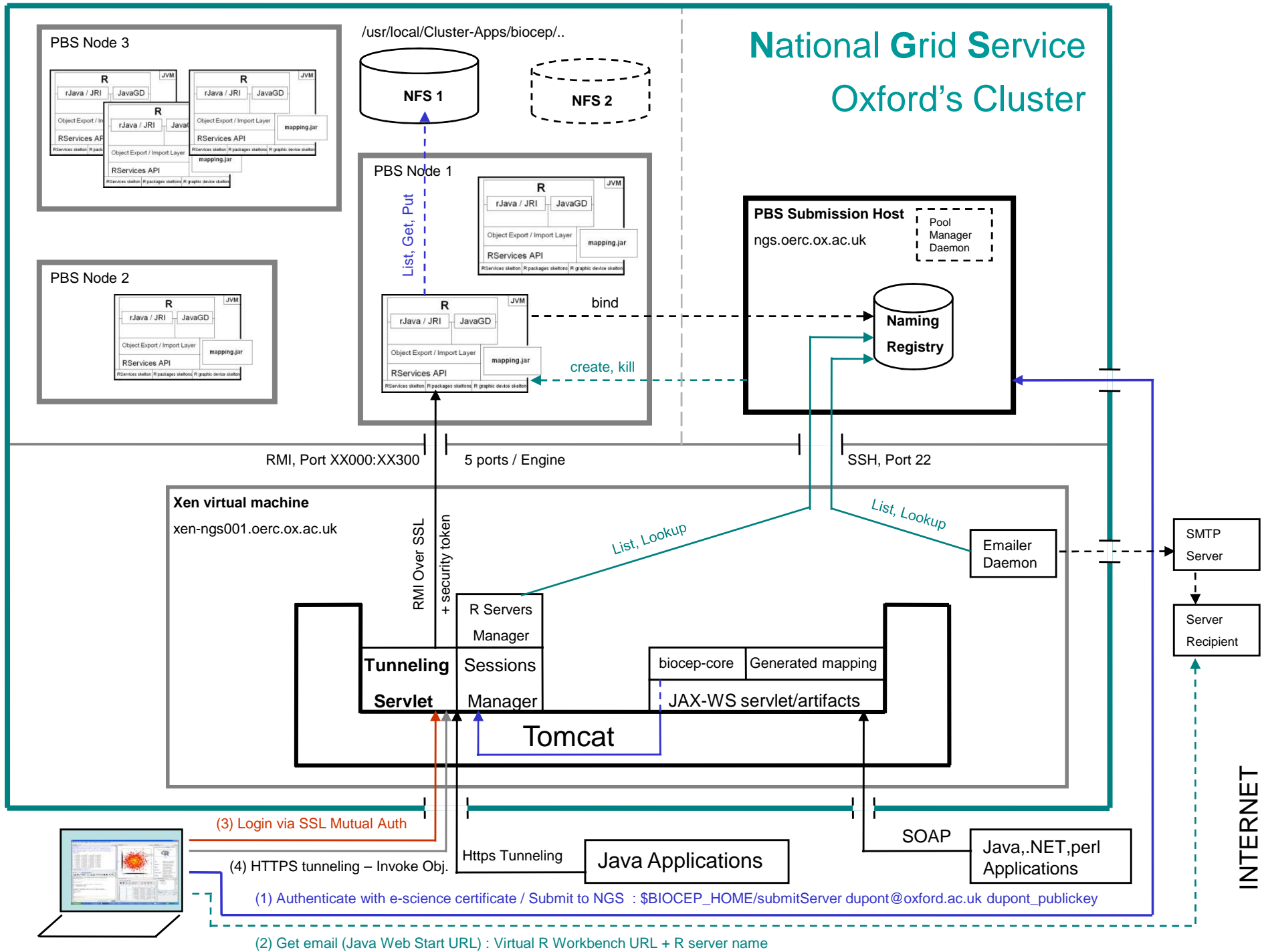
**ES** : ExpressionSet

**ESon1, ESon2, ESon3** : ExpressionSet Object Names

$f = T3 \circ T2 \circ T1$

# R Virtualization on an LSF Cluster





(2) Get email (Java Web Start URL) : Virtual R Workbench URL + R server name

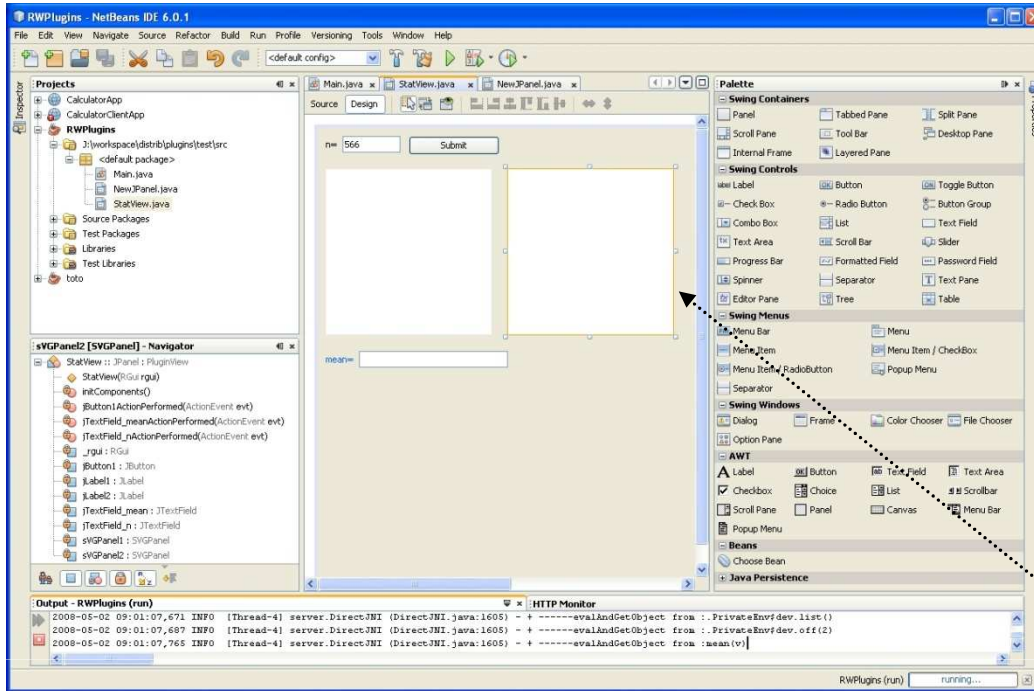
(1) Authenticate with e-science certificate / Submit to NGS : \$BIOCEP\_HOME/submitServer dupont@oxford.ac.uk dupont\_publickey

(4) HTTPS tunneling - Invoke Obj.

(3) Login via SSL Mutual Auth

# Netbeans 6 – Visual GUI builder

# GUI Plugins



Compile

**myPlugin.jar**

- + myView1
- + myView2
- + descriptor.xml

Import Plugin

Virtual R Workbench

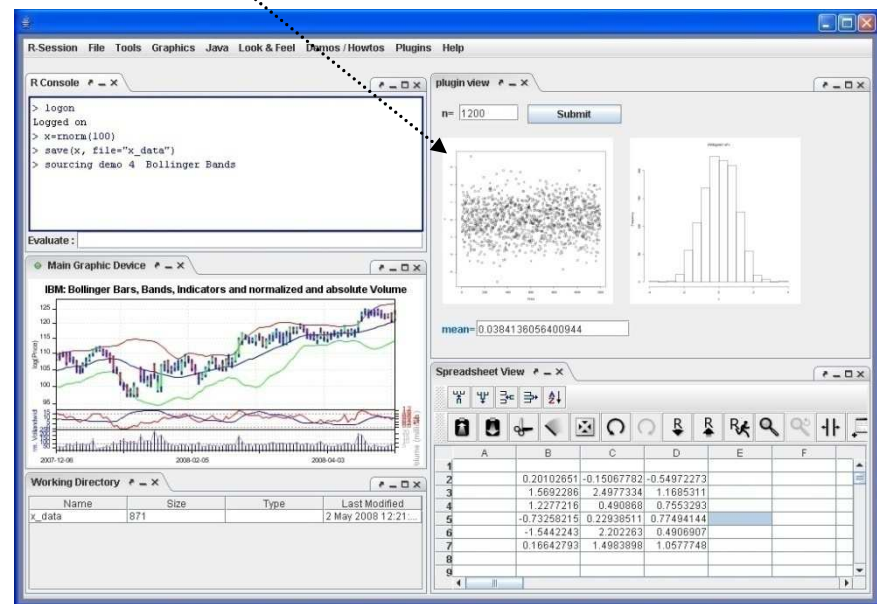
Upload plugin

**Plugins Repository**

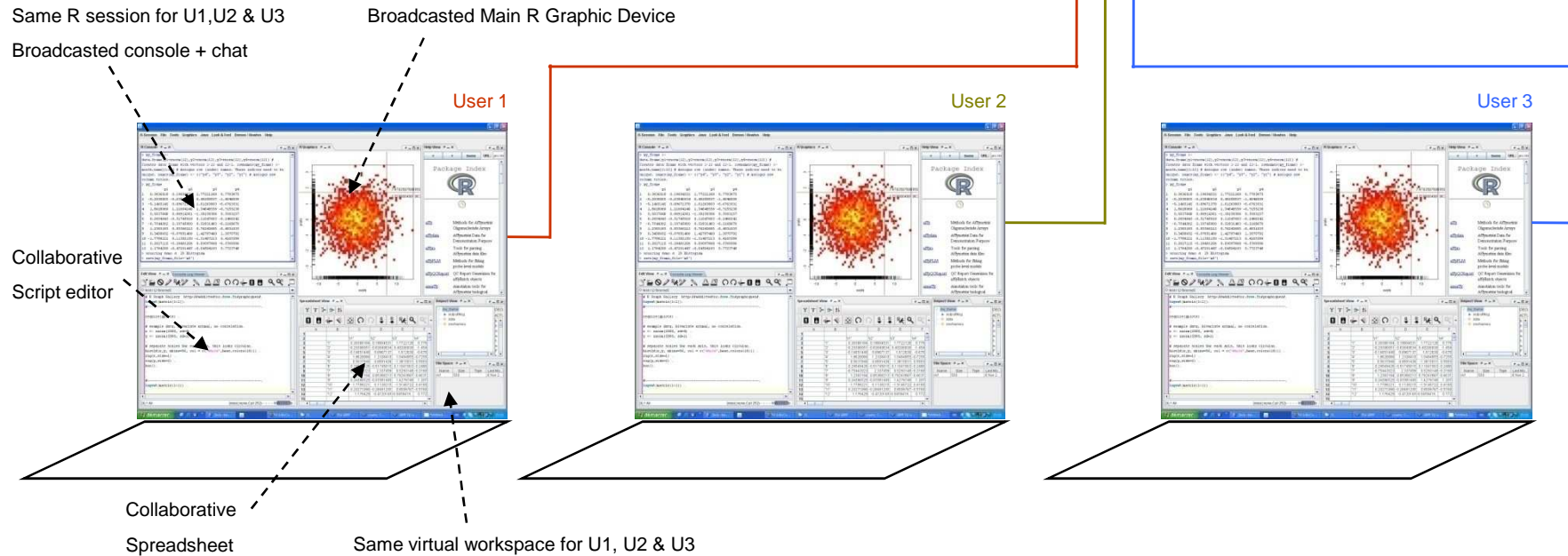
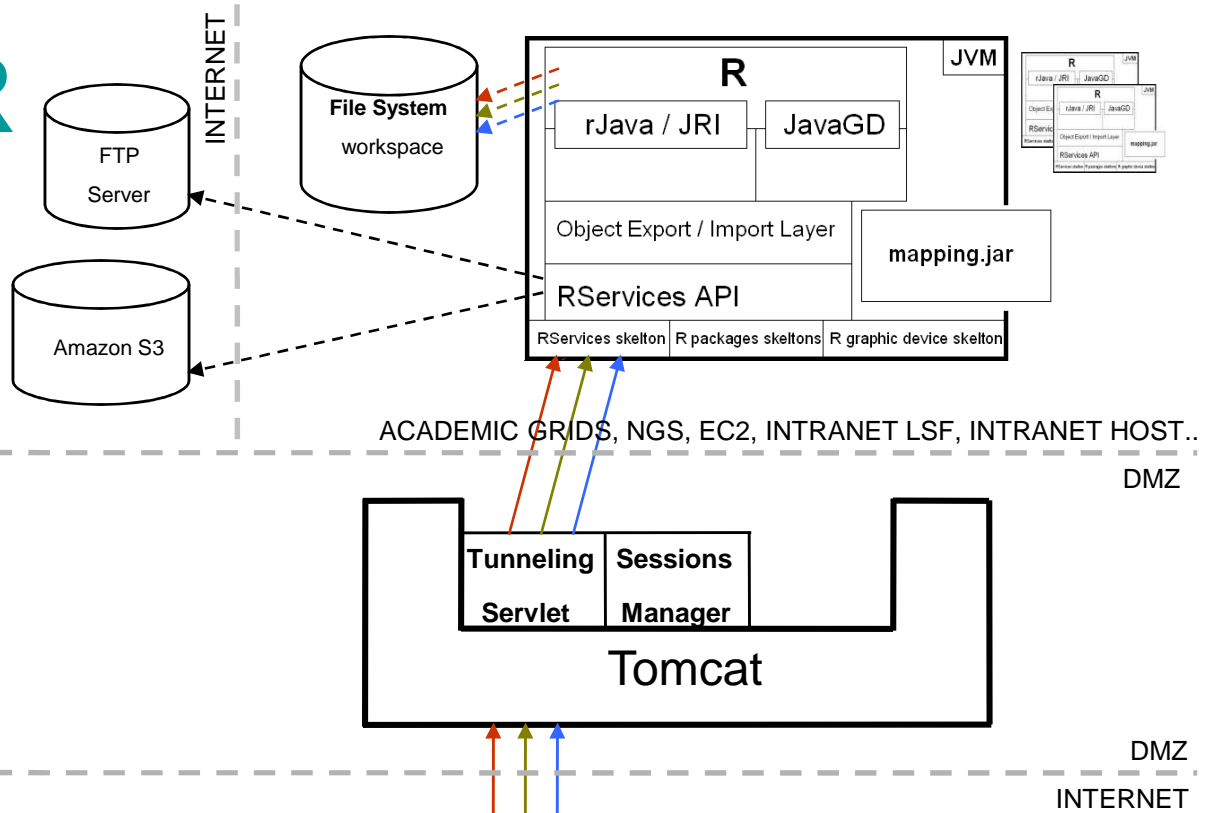
- \* myPlugin \* myDashboard
- \* Klimt \* iPlots
- \* Mondrian \* E. Profiler

Browse Repository

Download Plugin



# Collaborative R



# Ease Of Use - I

## •• Reasonable Pre-requirements

- java 5 and R $\geq$ 2.5 accessibles from the command line : to run R servers, generate mappings & Web Services, run the miniature virtualisation and the R-SOAP Web Apps..

## •• All-in-one Highly Productive Workbench

- Docking framework, spreadsheets, syntax highlighting enabled editors, objects viewer,
  - help browser, storage views, zooming system on R graphics, settings persistence..

## • Easy Computational Resource Acquisition

- Provide nothing to run R servers on local machine
- Provide HOST / PORT / LOGIN / PWD to run R Servers on remote hosts (SSH)
  - Provide URL & (LOGIN/PWD or X.509 Certificate) to Connect to Grid Rs or Cluster Rs

## • Easy Scripting

- Simple API for running/connecting to R servers
- Embeddable R code (`<R> </R>`) within scripts
- Automatic conversion from/to R Objects for common data types(standard,arrays,collections)

# Ease Of Use -II

- **Easy Plugins Integration**

Import local file / Browse Plugins repository and choose a plugin

- « **Push button** » **Web Services Generation/Web Services Deployment**

Add TypeInfo to your function / add your function name to an XML / run biocep-tools

Deploy: *java -port=80 -cp biocep-core.jar HttpServer rvirtual.war MyWebServices.war*

- **Self-contained jar & war files distribution :**

biocep.jar biocep-core.jar biocep-tools.jar rvirtual.war rws.war

- **Configurationless Parallel Computing from R console :**

makeCluster(n,..), stopCluster(cl), clusterEvalQ(cl, expr), clusterApply(cl, seq, fun, ..) ...



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