Proposed design to the Bioc Build System

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I. INTRODUCTION

The BBS (BioC Build System) consists of 2 subsystems: the *Build System* (running on several *build nodes*) and the *Publishing System* (running on the *publishing node*).

The src pkgs (source packages) and bin pkgs (binary packages) produced by the build nodes are called build products. All other files and directories produced by the build nodes are called build results.

Names of build products and build results are represented in bold.

During the builds, all build products and build results for a given BioC version (branch) are placed in a central location referred to as the BBS_LATEST_BASEURL. For the 1.8 builds, BBS_LATEST_BASEURL is something like http://gopher5/.../1.8/latest (directory).

Depending on its role in the Build System, a build node might need write access here (using mv, rsync of ftp). All nodes (build nodes + publishing node) must be able to retrieve files from there via HTTP.

II. MAIN TASKS

1) The Build System does the following:

The "src build" task

On each build node, build the BioC src pkgs from the pkg dirs:

- COMMAND: Run 'R CMD build <pkg dir>' on each pkg dir.
- BUILD PRODUCTS (1 file per pkg dir): <pkg>_x.y.z.tar.gz (the src pkg).

- BUILD RESULTS (3 files per pkg dir): <pkg>.src.out,
 <pkg>.src.err and <pkg>.src.report.
- Put the above products and results under BBS LATEST BASEURL/<build node>/src/

The "src check" task

On each *build node*, **check** the BioC *src pkgs* produced during the "src build" task:

- COMMAND: Run 'R CMD check < src pkg>' on each src pkg.
- BUILD PRODUCTS: None.
- Put the above results under BBS LATEST BASEURL/

 build node>/

The "bin build" task

On some chosen build nodes only, build the BioC bin pkgs:

- COMMAND: Run 'R CMD INSTALL -build src pkq>' on each src pkq.
- BUILD PRODUCTS (1 file per src dir): <pkg>_x.y.z<platform
 suffix> (the bin pkg).
- BUILD RESULTS (3 files per src pkg): <pkg>.bin.out,
 <pkg>.bin.err and <pkg>.bin.report.
- Put the above products and results under BBS_LATEST_BASEURL/<build node>/bin/

NB: Bin pkgs have a platform specific suffix: ".zip" on Windows, ".tgz" on OS X, " R x86 64-unknown-linux-gnu.tar.gz" on x86 64 Linux, etc...

2) The Publishing System does the following:

The "publishing" task

Publish the following stuff to the Bioconductor website:

- Publish all *build results* produced by the *Build System* to the "nightly build" section of the website.
- Publish src pkgs (from one chosen build node only) + bin pkgs (from one Windows build node only) to the repos section of the website. Replace a previously published pkg by the new one only if the new one was build with no errors and has a higher version string.
- Publish all the vignettes extracted from the *src pkgs* (from one chosen *build node* only) somewhere on the website so that people can download them without having to download the corresponding *src pkgs*.

III. NODE ROLES

A build node can have several roles, depending on what we want it to do:

- It can be "the prelim repos builder" (see STEP 1 below). Only one build node needs to perform this step.
- It can be "a src builder" (see STEPS 2,3 below) i.e. a machine that builds src pkgs. In theory 2 src pkgs build from the same pkg dir on 2 different build nodes should be (almost) identical, except for timestamps and (on very rare cases) for vignettes (due to obscure platform related issues).

- It can be "a src checker" (see STEPS 2,3,4 below) i.e. a machine that checks src pkqs.
- It can be "a bin builder" (see STEPS 2,3,5 below) i.e. a machine that builds the bin pkgs from the src pkgs.

The *publishing node* is the node that performs the "publishing" steps (see STEPS 6,7,8 below).

IV. LOCAL LAYOUT

On each build node, a given BioC branch (e.g. 1.8) is build under its own directory called the BBS_WORK_TOPDIR. For efficiency, the BBS WORK TOPDIR should be **local** to the build node (not NFS mounted).

On "the prelim repos builder", the BBS_WORK_TOPDIR layout looks like this:

```
BBS_WORK_TOPDIR/
   madman-wc/
   madman/
On "a src checker" or "a bin builder", it looks like this:
   BBS_WORK_TOPDIR/
   madman/
No BBS WORK TOPDIR is needed on the publishing node.
```

V. 8 STEPS

It's convenient to define the 8 following steps (elementary tasks) that we can put together in order to achieve the 4 main tasks described at the beginning:

STEPS 1 and 2 are preliminary steps. STEP 1 is a global preliminary step: it needs to be performed on one build node only ("the prelim repos builder") before any other build node can start to do anything. STEP 2 is a local preliminary step: it must be run on each build node before anything else (after STEP 1 has successfully terminated on the "prelim repos builder").

STEPS 3,4,5 correspond to the "src build", "src check" and "bin build" tasks.

STEPS 6,7,8 are the "publishing" steps (performed on one the *publishing* node only).

STEP 1: Make "madman.tgz" and "latest prelim repos"

```
Do a "rolling backup" of BBS_LATEST_BASEURL

cd BBS_WORK_TOPDIR

rm -rf madman madman.tgz

svn up madman-wc

svn export madman-wc madman

tar zcf madman.tgz madman

mv madman.tgz BBS_LATEST_BASEURL/

mkdir BBS_LATEST_BASEURL/src

mkdir BBS_LATEST_BASEURL/src/contrib

cd madman/Rpacks

Get list of pkgs from manifest file

for each pkg in list do:

R CMD build --no-vignettes <pkg>

mv <pkg>_x.y.z.tar.gz BBS_LATEST_BASEURL/src/contrib/

Make the "latest prelim repos" control file
```

STEP 2: Install all pkgs from "latest prelim repos"

-- no details for now --

STEP 3: Build the src pkgs

STEP 4: Check the src pkgs

STEP 5: Build the bin pkgs

STEP 6: Publish the "src build", "src check" and "bin build" results to the Bioconductor website.

```
-- no details for now --
```

STEP 7: Publish the src pkgs and bin pkgs to a public repos (update only).

```
-- no details for now --
```

STEP 8: Publish the vignettes to the Bioconductor website.

```
-- no details for now --
```

VI. SCHEDULING THE NODES

Here is the list of nodes we use and the roles they have:

	platform	prelim repos builder	src builder	src checker	bin builder	publis	her
gopher5	Linux x86_64	1	2 3	4		6 7	8
wellington	Linux i686		2* 3	4			
churchill	sparc		2* 3	4			
lemming	Windows 2003 Server		2* 3	4	5		
walpople	Windows XP		2 3	4	5		

We could also setup a "bin builder" node only (no checks) like this:

binbuilderonly: 2* 3 5

On each node a step can only be started if the preceding step terminated successfully. A node might also have to wait that some step terminates on **another** node before it can start to perform a step e.g. 2* here means that STEP 2 can only be started if STEP 1 terminated successfully on gopher5.

It would be nice to implement a cross machine locking/logging mechanism. Not the first priority though...