A workflow for implementing HTCondor Pipelines

First Mondays: November 6, 2017

Problem: Building binaries for CHTC is problematic.

Workflow: Two ideas to alleviate the pain:

- 1. Build on a CentOS 7 platform that's more friendly to users.
- 2. Use Git to manage the environment.

High-level view:

- Step 1: Get the binary running on pink.botany.wisc.edu. Capture the environment in a (local) git repository.
- Step 2: Propogate the git repository to a "remote" server.
- Step 3: Fire up an interactive HTCondor job, as usual.
- Step 4: Clone the repo on the execute node and test. Push changes to the remote repo.
- Step 5: Deploy as a noninteractive job, as usual.

A workflow for implementing HTCondor Pipelines

First Mondays: November 6, 2017

Problem: Building binaries for CHTC is problematic.

Workflow: Two ideas to alleviate the pain:

- 1. Build on a CentOS 7 platform that's more friendly to users.
- 2. Use Git to manage the environment.

Some Details:

Step 1: Get it running on pink.

Ask Steve or Henry to set up a login.

1. Connect to pink:

```
ssh pink.botany.wisc.edu
```

- 2. Make directory and cd into it.
- 3. Initialize a git repo:

```
git init
```

4. Follow the directions for building bwa:

https://github.com/lh3/bwa

Hint: google "bwa github"

- 5. Test the binary.
 - a. Run the command:

./bwa

b. Construct test data.

head -10 reference.fasta >
test/indir/reference.test.fasta
head -1000 reads.fastq > test/indir/reads.test.fastq

Commit to the repo early and often!

```
git status
git add
git commit -m "description"
```

Step 2: Propagate to remote repository. (Actually, this is also on pink – but it could be on another server, e.g. github.com.)

1. Create a home for your remotes.

```
mkdir ~/gitRepos
cd ~/gitRepos
git --bare init runbwa.git
```

*Hint: google "git setup bare repo"*https://gist.github.com/joahking/780877

2. Connect your local git repo to this new remote:

```
cd - ### change to the previous dir!
git remote add origin ~/gitRepos/runbwa
git push origin master
```

Step 3: Fire up an interactive job, as usual.

```
condor_submit -i queue/submit/interactive.submit
```

Hints: Use this as a starting template.

http://chtc.cs.wisc.edu/inter-submit.shtml

```
Use screen session to launch a job that persists.
screen -S "name"
```

Step 4 (method 1): Go to town in the interactive job.

```
./bwa
```

- ./bwa index indir/reference.fasta
- ./bwa mem indir/reference.fasta indir/reads*.fastq > pe.sam

Step 4 (method 2): Go to town in the interactive job.

git clone pink.botany.wisc.edu:gitRepos/runbwa

- ./bin/bwa
- ./bin/bwa index test/indir/reference.fasta

etc.

Continue to develop your pipeline in the interactive job and then push to the remote.

```
git push origin master
```

You can clone Steve's repo:

```
git clone
pink.botany.wisc.edu:/home/sgoldstein/gitRepos/runbwa
```