git production:

https://svn.bcgsc.ca/stash/projects/PF/repos/pipeline-components/browse

ssh:

ssh beast.cluster.bccrc.ca

ssh rocks3.cluster.bccrc.ca

- /share/lustre/

- /share/scratch/

- /genesis/extscratch/shahlab/ 70T

- /shahlab/ 350T

pipeline factory:

/share/scratch/pipeline\_temp/factory/

pipeline runner

/share/scratch/pipeline\_temp/factory/pipeline-tools/pipeline\_runner/pipeline\_runner.py

help wiki:

https://svn.bcgsc.ca/stash

https://www.bcgsc.ca/jira/

http://wiki.mo.bccrc.ca/

/share/scratch/pipeline\_temp/apps/anaconda/bin/python pipeline-tools/pipeline\_runner/pipeline\_runner.py -w example/ -c pipelines/convert\_bam\_to\_fastq/convert\_bam\_to\_fastq.yaml --setup ./pipelines/convert\_bam\_to\_fastq/convert\_bam\_to\_fastq\_setup\_rocks.txt --sample example/sample.txt -r test --drmaa\_library\_path /opt/gridengine/lib/lx26-amd64/libdrmaa.so --num\_jobs 40 --num\_pipelines 10 -q ' -q all.q -l mem\_free={mem},mem\_token={mem},h\_vmem={mem}' -b drmaa --results\_dir example/ --python\_installation /share/scratch/pipeline\_temp/apps/anaconda/bin/python

setups:

\_\_OPTIONS\_\_ factory\_dir /share/scratch/pipeline\_temp/factory/pipeline-factory/pipeline\_factory/

\_\_OPTIONS\_\_ components\_dir /share/scratch/jrosner\_temp/factory/components/

\_\_OPTIONS\_\_ drmaa\_library\_path /opt/gridengine/lib/lx26-amd64/libdrmaa.so

\_\_GENERAL\_\_ python /share/scratch/pipeline\_temp/apps/anaconda/bin/python

\_\_GENERAL\_\_ samtools /share/scratch/pipeline\_temp/apps/samtools-0.1.18/samtools

/share/lustre/kshumansky/Alignments/Huntsman/FF/wtss/diffExpr/deseq/ff\_deseq\_epsr\_filtered\_naturesamples\_deseq\_dispersionEstimation\_counts\_retry.R