



FauxConda: ways to use Conda without Conda

Wednesday, March 24, 2021

We will be starting at 12:05 PM

Disclaimer

The title was just to get you here, we will NOT use Conda at all !!!

Except in the singularity image

Here we assume that:

- You can connect to Compute Canada clusters
- You are comfortable with python and python environments
- You understand how to load and search for modules in Compute Canada systems.



Why should you avoid Conda?

Conda is a useful solution but to ensure compatibility and optimal performance, should not be used. Also (list is not comprehensive):

- Anaconda might suboptimally install software we already have
- Use inappropriate binaries for our architecture.
- Makes incorrect assumptions about the location Libraries.
- Is enormous filling your \$HOME
- Installation is s slower than using Python wheels.
- It modifies the .bashrc file, which can cause conflicts.





What options do you have?

In Compute Canada systems you have two main options:

- 1. Generate a python virtual environment:
 - a. Check python dependencies
 - b. Check program dependencies
 - c. Load and install dependencies
- 2. Install your package with conda within a singularity image
 - a. Create a singularity image
 - i. If available from dockerhub, you are done!
 - b. Install conda
 - c. Install your program





Python virtual environment

- Does not manage packages
- Provides a user-writable space to install python packages (mainly through pip)
- Sets a temporary user space (when activated) with the paths exported.
- DOES NOT CLASH WITH THE SYSTEM!!!!
- It uses optimal compilation of programs





Step 0: Download the source of the program you want

For this webinar we will be using a bioinformatics pipeline called Anvi'o. You can follow their instructions. In brief clone git:

```
[jshleap@cedar1 scratch]$ git clone --recursive https://github.com/merenlab/anvio.git Cloning into 'anvio'...
remote: Enumerating objects: 246, done.
remote: Counting objects: 100% (246/246), done.
remote: Compressing objects: 100% (144/144), done.
remote: Total 77549 (delta 144), reused 179 (delta 102), pack-reused 77303
Receiving objects: 100% (77549/77549), 394.58 MiB | 2.44 MiB/s, done.
Resolving deltas: 100% (59056/59056), done.
Updating files: 100% (1175/1175), done.
```





Step 1: Identify python dependencies

Good packages will have a requirements.txt file, others, you will have to scrape conda's meta.yaml.

```
[jshleap@gra-login1 scratch]$ cd anvio/
[jshleap@gra-login1 anvio]$ ls
anvio AUTHORS.txt bin CHANGELOG.md conda-recipe Dockerfile LICENSE.txt
MANIFEST.in NEWS.md README.md requirements.txt sandbox setup.py
[jshleap@gra-login1 anvio]$ ls conda-recipe/anvio*
conda-recipe/anvio:
meta.yaml
conda-recipe/anvio-minimal:
meta.yaml
```





Step 1.1: Create a virtual environment and install python dependencies

```
[jshleap@gra-login1 anvio]$ module load StdEnv/2020 gcc/9.3.0 python/3.8 r/4.0.2
Lmod is automatically replacing "intel/2020.1.217" with "gcc/9.3.0".

Lmod is automatically replacing "intel/2020.1.217" with "gcc/9.3.0".

[jshleap@gra-login1 anvio]$ virtualenv --no-download anvio_env
created virtual environment CPython3.8.2.final.0-64 in 2184ms
    creator CPython3Posix(dest=/scratch/jshleap/anvio/anvio_env, clear=False, global=False)
    seeder FromAppData(download=False, pip=latest, setuptools=latest, wheel=latest, via=copy, ap
p_data_dir=/home/jshleap/.local/share/virtualenv/seed-app-data/v1.0.1)
    activators BashActivator,CShellActivator,FishActivator,PowerShellActivator,PythonActivator,X
onshActivator
[jshleap@gra-login1 anvio]$ source ./anvio_env/bin/activate
(anvio_env) [jshleap@gra-login1 anvio]$ pip install --no-index -r requirements.txt
```

(anvio_env) [jshleap@gra-login1 anvio]\$ pip install --no-index numpy scikit-learn pandas --upg rade





Step 2: Identify software dependencies and load the modules

We carry a long list of software, scrape the names on the conda's meta.yaml, and use module spider to check if we have it:

```
(anvio_env) [jshleap@gra-login1 anvio]$ module spider prodigal
 prodigal: prodigal/2.6.3
   Description:
     Prodigal (Prokaryotic Dynamic Programming Genefinding Algorithm) is
     a microbial (bacterial and archaeal) gene finding program developed
     at Oak Ridge National Laboratory and the University of Tennessee.
   Properties:
     Bioinformatic libraries/apps / Logiciels de bioinformatique
   You will need to load all module(s) on any one of the lines below before t
he "prodigal/2.6.3" module is available to load.
     StdEnv/2020
     nixpkgs/16.09 gcc/5.4.0
     nixpkgs/16.09 gcc/7.3.0
```



Step 2.1: Add module load to activate (Optional)

If you'd like that all available requirements are loaded on activation:

```
(anvio_env) [jshleap@gra-login1 anvio]$ echo 'module load StdEnv/2020 gcc/9.3.0 scipy-stack/20
20b \' >> ${VIRTUAL ENV}/bin/activate
(anvio env) [jshleap@gra-login1 anvio]$ echo '
                                                prodigal/2.6.3 mcl/14.137 hmmer/3.2.1 diamond/
2.0.4 \' >> ${VIRTUAL ENV}/bin/activate
(anvio env) [jshleap@gra-login1 anvio]$ echo '
                                               blast+/2.11.0 spades/3.15.1 megahit/1.2.9 \' >
> ${VIRTUAL ENV}/bin/activate
(anvio env) [jshleap@gra-login1 anvio]$ echo ' bowtie2/2.4.1 bwa/0.7.17 samtools/1.11 centrif
uge/1.0.4-beta \' >> ${VIRTUAL ENV}/bin/activate
(anvio env) [jshleap@gra-login1 anvio]$ echo ' trimal/1.4 fastani/1.32 trnascan-se/2.0.7 fa
sttree/2.1.11 r/4.0.2' >> ${VIRTUAL ENV}/bin/activate
(anvio env) [jshleap@gra-login1 anvio]$ tail ${VIRTUAL ENV}/bin/activate
# be called to get it to forget past commands. Without forgetting
 past commands the $PATH changes we made may not be respected
if [ -n "${BASH-}" ] || [ -n "${ZSH VERSION-}" ] ; then
   hash -r 2>/dev/null
module load StdEnv/2020 gcc/9.3.0 scipy-stack/2020b \
 prodigal/2.6.3 mcl/14.137 hmmer/3.2.1 diamond/2.0.4 \
 blast+/2.11.0 spades/3.15.1 megahit/1.2.9 \
 bowtie2/2.4.1 bwa/0.7.17 samtools/1.11 centrifuge/1.0.4-beta \
 trimal/1.4 fastani/1.32 trnascan-se/2.0.7 fasttree/2.1.11 r/4.0.2
```

Step 2.2: Install the missing R components

Use local install:

```
(anvio_env) [jshleap@gra-login1 anvio]$ mkdir -p ${VIRTUAL_ENV}/R/lib
(anvio_env) [jshleap@gra-login1 anvio]$ export R_LIBS=${VIRTUAL_ENV}/R/lib
(anvio_env) [jshleap@gra-login1 anvio]$ echo "export R_LIBS=${VIRTUAL_ENV}/R/lib" >> ${VIRTUAL_ENV}/Bin/activate
(anvio_env) [jshleap@gra-login1 anvio]$ R -e "install.packages(c('stringi', 'magrittr', 'optpa rse', 'rlang', 'Rcpp', 'tidyverse', 'BiocManager'), depencies=TRUE, lib='${VIRTUAL_ENV}/R/lib', repos='http://cran.r-project.org', quiet=TRUE)"
```

```
(anvio_env) [jshleap@gra-login1 anvio]$ R -e "BiocManager::install('qvalue', lib='${VIRTUAL_EN
V}/R/lib')"
```

Step 3: Install the missing software

(anvio_env) [jshleap@gra-login1 anvio]\$ cd \${VIRTUAL_ENV}

Sometimes we do not have the required software, so you can install it in your virtual environment. In our case IQTree:

```
(anvio env) [jshleap@gra-login1 anvio env]$ git clone https://github.com/Cibiv/IO-TREE.git
Cloning into 'IQ-TREE'...
remote: Enumerating objects: 861, done.
remote: Counting objects: 100% (861/861), done.
remote: Compressing objects: 100% (474/474), done.
remote: Total 30979 (delta 626), reused 565 (delta 387), pack-reused 30118
Receiving objects: 100% (30979/30979), 65.23 MiB | 23.96 MiB/s, done.
Resolving deltas: 100% (23141/23141), done.
Updating files: 100% (638/638), done.
(anvio_env) [jshleap@gra-login1 anvio_env]$ cd IQ-TREE && mkdir build && cd build
(anvio env) [jshleap@gra-login1 build]$ module load eigen/3.3.7
(anvio_env) [jshleap@gra-login1 build]$ cmake -DIQTREE_FLAGS=omp -DEIGEN3_INCLUDE_DIR=${EBROOT
EIGEN} -DCMAKE INSTALL PREFIX=${VIRTUAL ENV} ..
(anvio_env) [jshleap@gra-login1 build]$ make -j 4
(anvio env) [jshleap@gra-login1 build]$ chmod +x iqtree
(anvio_env) [jshleap@gra-login1 build]$ cp iqtree $VIRTUAL_ENV/bin
(anvio_env) [jshleap@gra-login1 build]$ cd ../..
```



Step 4: Install Anvi'o

Now that we have all the requisites, installing avio is simple:

```
(anvio_env) [jshleap@gra-login1 anvio]$ pip install .
Ignoring pip: markers 'python_version < "3"' don't match your environment
Looking in links: /cvmfs/soft.computecanada.ca/custom/python/wheelhouse/gentoo/avx2, /cvmfs/soft.computecanada.ca/cust
ft.computecanada.ca/custom/python/wheelhouse/gentoo/generic, /cvmfs/soft.computecanada.ca/cust
om/python/wheelhouse/generic
Processing /scratch/jshleap/anvio</pre>
```

We can test it, but first deactivate and reactivate the environment:

```
(anvio_env) [jshleap@gra-login1 anvio]$ deactivate
[jshleap@gra-login1 anvio]$ source anvio_env/bin/activate

Lmod is automatically replacing "intel/2020.1.217" with "gcc/9.3.0".

Lmod is automatically replacing "intel/2020.1.217" with "gcc/9.3.0".

(anvio env) [jshleap@gra-login1 anvio]$ anvi-self-test
```



Step 1: Build a singularity image

Some programs can be found in <u>dockerhub</u> like <u>Anvi'o</u>. In those cases your life just about to become a lot easier:

```
[jshleap@gra-login1 jshleap]$ module load singularity
[jshleap@gra-login1 jshleap]$ singularity build anvio.sif docker://meren/anvio
INFO: Starting build...
```

If not, don't worry, we can still build a singularity image:

```
[jshleap@gra-login1 jshleap]$ singularity build --sandbox anvio-dir docker://centos INFO: Starting build...
```

Singularity Image with conda

- Self contained space
- All variables and binary paths are set for you
- Easier to set up

 It is suboptimal to run in our systems (compilation)



Step 2: If not in docker hub, install conda

[jshleap@gra-login1 scratch]\$ singularity build --sandbox anvio_from_source-dir docker://centos INFO: Starting build...



Step 2.1: Activate conda

```
Singularity> mkdir -p /localhome
Singularity> echo 'export HOME=/localhome' >> /environment
Singularity> source /environment
Singularity> source /opt/conda/bin/activate
(base) Singularity> conda init
no change /opt/conda/condabin/conda
no change /opt/conda/bin/conda
no change /opt/conda/bin/conda-env
          /opt/conda/bin/activate
no change
             /opt/conda/bin/deactivate
no change
             /opt/conda/etc/profile.d/conda.sh
no change
no change
             /opt/conda/etc/fish/conf.d/conda.fish
             /opt/conda/shell/condabin/Conda.psm1
no change
             /opt/conda/shell/condabin/conda-hook.ps1
no change
             /opt/conda/lib/python3.8/site-packages/xontrib/conda.xsh
no change
no change
             /opt/conda/etc/profile.d/conda.csh
modified
             /localhome/.bashrc
==> For changes to take effect, close and re-open your current shell. <==
(base) Singularity> echo ". /localhome/.bashrc" >> /environment
```



Step 3: Download, install, and activate your recipe

```
(base) Singularity> curl https://merenlab.org/files/anvio-conda-environments/anvio-environment
-7-LINUX.yaml --output anvio-environment-7.yaml
% Total % Received % Xferd Average Speed Time Time Current
Dload Upload Total Spent Left Speed
100 9650 100 9650 0 0 13049 0 --:--:-- 13040
(base) Singularity> conda env remove --name anvio-7
(base) Singularity> conda env create -f anvio-environment-7.yaml
Collecting package metadata (repodata.json): \
```

```
[jshleap@gra-login1 jshleap]$ singularity shell -e -w anvio-dir
Singularity> echo "conda init" >> /environment
Singularity> echo "conda activate anvio-7" >> /environment
Singularity> . /environment
no change /opt/conda/condabin/conda
no change
             /opt/conda/bin/conda
no change
              /opt/conda/bin/conda-env
no change
              /opt/conda/bin/activate
              /opt/conda/bin/deactivate
no change
              /opt/conda/etc/profile.d/conda.sh
no change
              /opt/conda/etc/fish/conf.d/conda.fish
no change
              /opt/conda/shell/condabin/Conda.psm1
no change
              /opt/conda/shell/condabin/conda-hook.ps1
no change
              /opt/conda/lib/python3.8/site-packages/xontrib/conda.xsh
no change
              /opt/conda/etc/profile.d/conda.csh
no change
              /localhome/.bashrc
no change
No action taken.
(anvio-7) Singularity> conda install -y -c bioconda hmmer=3.2.1
```

Step 3.1: Keep following instructions

Most programs will have a set of steps to follow... just follow those. For Anvi'o:

```
Singularity> cd /opt/
Singularity> wget https://github.com/merenlab/anvio/releases/download/v7/anvio-7.tar.gz

Singularity> pip install anvio-7.tar.gz

Processing ./anvio-7.tar.gz

Singularity> pip install mistune==0.8.4

Requirement already satisfied: mistune==0.8.4 in ./conda/envs/anvio-7/lib/python3.6/site-packa ges (0.8.4)
```



Step 4: Make it an image (optional) and test

```
[jshleap@gra-login1 jshleap]$ chmod -R 755 anvio-dir/
[jshleap@gra-login1 jshleap]$ singularity build anvio.sif anvio-dir
INFO: Starting build...
INFO: Creating SIF file...
INFO: Build complete: anvio.sif
[jshleap@gra-login1 jshleap]$ singularity run -e anvio.sif anvi-self-test
```

[jshleap@gra-login1 jshleap]\$ singularity run -e -w anvio-dir anvi-self-test



Q&A

Help can also be reached at support@computecanada.ca or help@sharcnet.ca.

Commands can be found at:

- 1. Python environment https://staff.sharcnet.ca/jshleap/GI_webinars/FauxConda/Commands_environment.sh
- 2. Singularity https://staff.sharcnet.ca/jshleap/Gl_webinars/FauxConda/Commands_singularity.sh

This presentation (minus the show and tell) can be found: https://staff.sharcnet.ca/jshleap/Gl_webinars/FauxConda/FauxConda.pdf

Recording of this presentation can be found at (but give it some time): https://youtube.sharcnet.ca