Review of Docker commands

Images

docker images #list images

docker run ubuntu <cmd_in_container>

docker save -o img.tar ubuntu

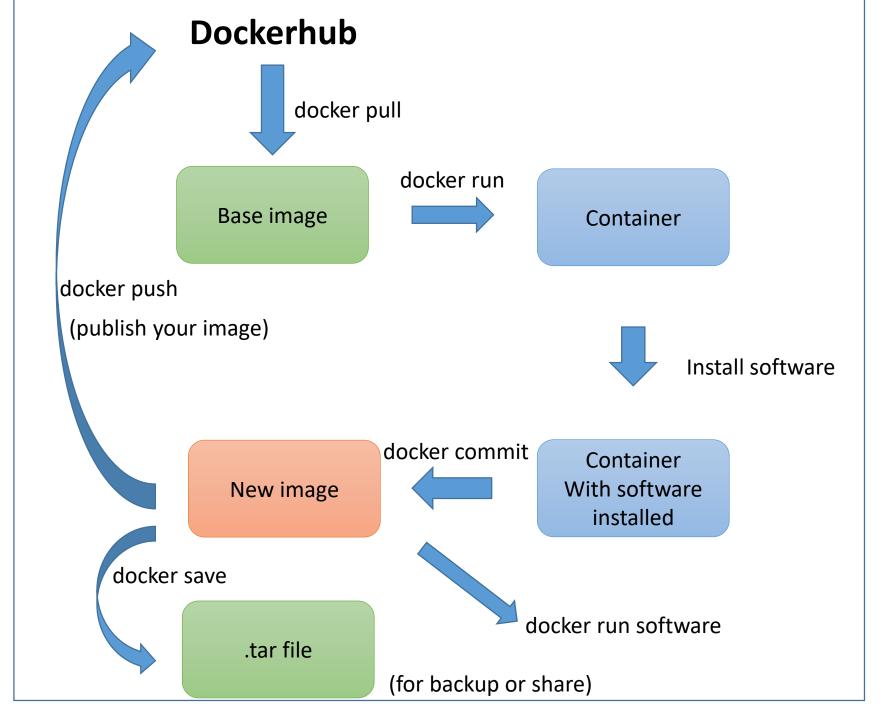
Containers

docker ps -a #list containers

docker exec 39sadf0a93 <cmd_in_container>

docker commit 39sadf0a93 img_name

Typical workflow



Review of Docker commands

docker run --rm -v /workdir/qisun/data:/data -w /data ezlabgva/busco:v5.4.3_cv1 busco -h

Command on host side

Image name

Command in container

export BUSCO="docker run --rm -v /workdir/qisun/data:/data -w /data ezlabgva/busco:v5.4.3_cv1"

\$BUSCO busco -h

Commonly used command options

```
docker run -dit ubuntu:20.04
                                       # -dit: start a docker container, running in background
docker run -it --rm myimg myscript.py
                                            # --rm: remove container after job finished; -it: in interactive mode
docker run --rm -v /workdir/$USER/data:/data myimg myscript.py
                                                                             #-v: mount a volume
docker run --rm -v /workdir/$USER/data:/data -w /data myimg myscript.py
                                                                                       #-w: current dir
docker exec -it asd9f7dasfa bash
                                         # execute a command (e.g. "bash") in a running container, interactively
```

Conda in Docker

Why?How?

Run BUSCO in Conda

source ~/miniconda3/bin/activate

conda create -n busco busco

conda activate busco

busco

~/miniconda3/env/busco:

• 3.1 GB, 41442 files

Run BUSCO in Docker

docker run --rm ezlabgva/busco:v5.4.3_cv1 busco

Image size:

• 2.28 GB, single file

Run BUSCO in Singularity

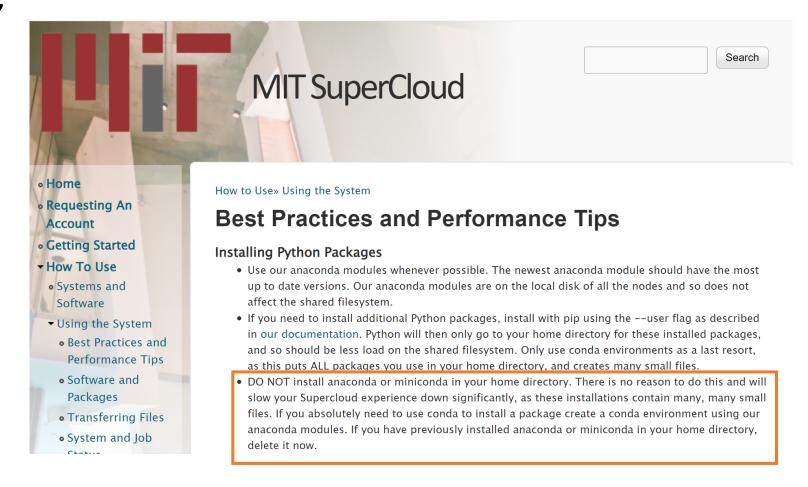
singularity build busco.sif docker://ezlabgva/busco:v5.4.3_cv1
./busco.sif busco

Image size:

• 0.73 GB, single file

Why is Conda bad for the cloud?

- To many small files in Conda is a huge burden for the cloud storage;
- Not fully encapsulated;



Solution: Put Conda in Docker/Singularity

Two base images for building Conda in Docker:

continuumio/miniconda3 P

Published by Anaconda

mambaorg/micromamba

Published by Mamba

Conda in Docker/Singularity

Protocol: https://biohpc.cornell.edu/doc/CondaInContainer.html

Using Dockerfile

create a Docker file

```
FROM continuumio/miniconda3
RUN conda install -y -c bioconda -c conda-forge \
    python=3.9.1 \
    samtools \
    bwa
RUN conda clean --all --yes
```

#build image

docker1 build -t samimage /workdir/qisun/

Interactively

```
docker run --dit continuumio/miniconda3
docker exec -it 348q0840qaf bash
conda install -y -c bioconda -c conda-forge \
   python=3.9.1
  samtools \
   hwa
conda clean --all yes
exit
conda commit 348q0840qaf samimage
```

#after the image is created, run command

docker run --rm samimage samtools

- Conda directory in Docker: /opt/conda;
- Install in Conda "base". No need to create environment;
- The "base" environment is activated automatically;

Advantage of Conda in Docker

1.One single file, instead of thousands or even millions;

2. Fully encapsulated, portable and reproducible;

Strongly recommended !!!



Singularity (new name Apptainer)



Another container system;

Why do we need it?

Docker vs Singularity

Problem of Docker:

- Run as root;
- Not supported in most HPC systems;
- Docker1 on BioHPC is a restricted Docker implementation;

Advantage of Singularity:

- Run as same user ID as the host;
- Supported in almost all HPC systems;

Disadvantage of Singularity:

 A newcomer, no community supported large database like
 Dockerhub

Build singularity images

Singularity image file: *.sif

Convert from Docker image

Docker image in Dockerhub

singularity build myimg.sif docker://rocker/r-ver

Is -I myimg.sif

An executable file

-rwxr-x--- 1 qisun qisun 319188992 Oct 26 20:28 myimg.sif

Build singularity images

Singularity image file: *.sif

From Docker image

A local Docker image tar file

singularity build myimg.sif docker-archive://myimg.tar

Build a Singularity image without Docker

From a Singularity .def file

Create a text file myapp.def

BootStrap: library
From: ubuntu:focal
%environment
%files
%post
apt -y update
apt install -y build-essential

Build command:

singularity build --fakeroot myapp.sif myapp.def

Interactively with a Singularity sandbox

#create sandbox
singularity build --fakeroot --sandbox myUbuntu myUbuntu.def

#start a shell in sandbox
singularity shell --fakeroot --writable myUbuntu

install all needed modules in the sandbox

convert the sandbox into a sif file
singularity build --fakeroot myUbuntu.sif myUbuntu

* I found it easier to build a Docker image, and then convert the Docker image into a Singularity image.

Run software in Singularity

singularity run img.sif cmd

* -C: Contain all. By default, environment and home directory are carried into the container;

singularity run -C img.sif cmd

* --bind: mount host directory into container. By default, home directory is mounted;

singularity run --bind /workdir:/data img.sif cmd

* --pwd: set default directory

singularity run --pwd /data img.sif cmd

Shortcuts in "singularity" commands

Full command: safest

```
singularity run -C \
--bind /workdir/$USER:/data \
--pwd /data \
img.sif cmd
```

Shortest command

```
./img.sif
```

- Home directory and environment are carried from host;
- No data file bound;
- Default command and current directory setting are built into the image;

Lots of shortcuts in "singularity" commands

Full command: safest

```
singularity run -C \
--bind $PWD \
--pwd $PWD \
img.sif cmd
```

- \$PWD: current directory
- If no ":", source and destination directories are the same

Shortest command

./img.sif

- Home directory and environment are carried from host;
- No data file bound;
- Default command and current directory setting are built into the image;

By default, Singularity is not fully contained.

These items are carried into the container:

- 1. Home directory (R and Python modules, setting files in home directory);
- 2. Environment variables (e.g. PYTHONPATH, PATH)
- 3. Current directory (rule changed in latest Singularity)

If you run into problems, do "singularity run -C" to contain everything, and then explicitly bind the directories and set default PWD

```
singularity run -C \
--bind /workdir/$USER:/data \
--pwd /data \
img.sif cmd
```

"singularity shell": an interactive interface

singularity shell ubuntu.sif

Apptainer> pwd /home/qisun

Apptainer> apt update
Permission denied

- Within container, only the mounted directories are writable;
- You cannot install extra software.
- To install software, the image needs to be converted into "Singularity sandbox" first.
- I found it easier to do all software installation in Docker first, and then to convert the Docker image to a Singularity image.

My protocol of using Docker and Singularity

- 1. Identify the base Docker image;
- 2. Build a Docker image interactively. Make sure to document every step;
- 3. (optional) Create a Docker file based on documented steps;
- 4. Convert Docker image into a Singularity image;
- 5. Save a copy of the Docker image .tar file;

6. Run software with Singularity image;