Containers Walkthrough

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(SLAC Shared) Scientific Data Facility



- Greenfield deployment of hardware, software and policies for Scientific Computing at SLAC
 - Initially led by LCLS, CryoEM and ML initiatives
- Governed by newly proposed SDF Steering Committee members whom are 'Owners' in the SDF
- New Storage
 - Most 'legacy' storage will be migrated to new disks
 - Manual process, will need to work with you to determine best way to move data (archive data)
 - Baseline 'Free' storage to be determined.

(SLAC Shared) Scientific Data Facility



- New Hardware
 - All GPUs migrated from LSF to Slurm
 - 88 x AMD Rome Dual 64-core ordered (~12k cores)
 - 200GB/sec "HDR" Infiniband network
- 'Baseline' paid out of indirect costs and investments
 - Every SLAC (computer) account will have
 - Some allotment of compute CPU and GPU 'for free'
 - Some allotment of storage 'for free'
 - Balancing act of resources!

SDF Compute Usage



- Priority Access to Compute Resources
 - 'Owners' can contribute compute hardware and/or FTE to help support planning, operations and development
 - Unused (compute) cycles from 'Owners' hardware will be made available to all Users → 'Shared'
 - Owners will have pre-emptive access to their own hardware contribution
 - Minimum contribution: 1 server
- Extra 'Shared' (compute) cycles bourne from lab direct investment
 - E.g. ocio-gpuXX machines
- Hardware spec to be reviewed and governed annually by the SDF Steering
 Committee

SDF Storage



- Designed and built upon DDN Exascaler Platform
- High speed controllers (~70GB/sec each)
- Current deployment
 - 2 sets of controllers, plan to expand to more.
 - 6+PB of spinning disk storage (2PB LCLS, 2PB CryoEM, 2PB Baseline)
- Calls for more storage will be pooled each period (per quarter?)
 - Minimum pool size ~500TB, approx 25k/5 years (TBC)
 - Possible chargebacks for % fair usage of shared hardware and service
- Migration
 - Plan to be de-facto storage for SLAC
 - Will need to work with everyone to existing data into SDF
 - Limited legacy cross mounts will be provided

- Spend a long time installing applications and dependencies on each new cluster
 - Where do install? Do i have an existing script for installs?
- Sometimes certain versions of libraries not available
 - python3?! boost, tensorflow v1 code etc.
 - `module load` only helps a little often maintained by cluster admins, may not have specific versions available (cuda?), conflicts between modules etc.
 - virtualenv/conda complementary to containers
- Typical to install into group/home directories
 - End up hard coding paths what if filesystem isn't available where you need it?
- Difficult to determine versions etc. (unless CVMFS tree structure like)
 - How to version control the application? Does new library create different results?
 - Output Description

 How to use different versions of the application?

- Consistent Environment
 - Self documenting
 - Shareable! Fork, pull etc.
 - Repeatable, reproducible
- Run anywhere
 - No need to compile for each cluster
- (Much) smaller and more efficient than Virtual Machines
- Faster than rebuilding everywhere with conda, pip etc.
 - But can utilize standard scripts and methods to create container
- Don't like using yum? Use apt instead, even if Host OS is CentOS.

Container Challenges

- Numerous different technologies
 - Docker, singularity, shifter, charliecloud, podman, enroot...
- Most solutions require sudo/root to build images
- GPU support can be... interesting
- Primarily a Linux only technology
 - Most cross platform solutions are VM based (exception WSL)
- Mostly aimed at Enterprise use cases (running webservers) rather than HPC (massively parallel workloads? Multi-user systems?)

Docker



- The 'de facto' container technology
- Dockerhub allows free storage and searching of containers
 - Create account at https://dockerhub.com
 - Chances are someone's already ported what you want
- Uses a local cache to store images, so each computer needs a duplicate copy
- Needs sudo/root to build and run (some 'rootless' features in latest version)
- Running docker containers typically have full root

- Container technology aimed at HPC applications
- Uses local '.sif' file per container that can be copied and moved like any other file
- Can convert from a docker image to singularity file
- Supported at most academic and laboratory institutions
 - Major exceptions: NERSC
- Containers can be pull'd and ran without sudo/root
 - Great for multi-user environments like our clusters
- Running singularity container has same uid/gids as on host OS (inside == outside)
- Simple integration with GPUs, MPI, X etc.
- Some (persistent) issues on AFS systems.

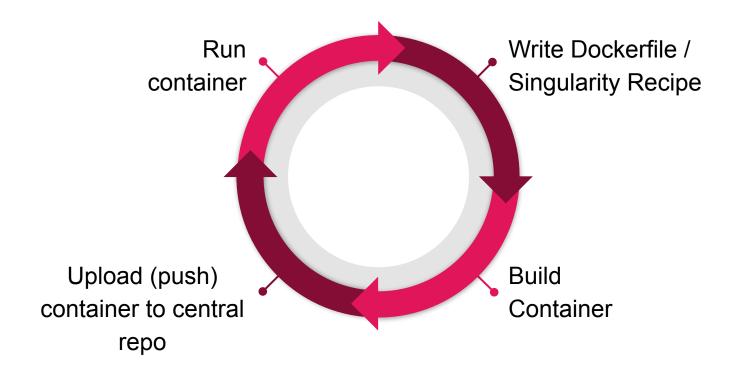
Why use Docker + Singularity?

- Use Docker for building
 - Simple language, well documented, plenty of examples
 - Support for cached 'layers' to allow for quick changes
 - Many different container technologies allow conversion from Docker images directly
 - le use the same image to run at NERSC
- Use singularity for running
 - No need for sudo/root
 - Can share applications as a single file

Demo Outline



- Create a simple python based application
- Build a Docker container from in
- Push container to Dockerhub
- Convert the Docker container to a Singularity container
- Deal with bind-mounts
- Run the singularity container
- Note about GPUs



Not covering singularity image create - not self documenting

Setup

-SLAC

	Mac	Linux
Install Docker	https://docs.docker.com/docker-for-mac/	https://docs.docker.com/engine/install/
Install Singularity	https://sylabs.io/singularity-desktop-macos/ or Vagrant (see later)	https://sylabs.io/guides/3.0/user-guide/installation.html

Vagrant Install

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- Need sudo! :(
- We can't give sudo on HPC nodes
- Only viable solution currently is for you to use your own laptop/desktop

on ubuntu

- Use a VM (not necessary on Linux)
 - Control it Vagrant

```
# on mac
brew cask install virtualbox
brew cask install vagrant
```

```
# https://www.vagrantup.com/downloads.html
```

```
sudo apt install virtualbox
dpkg install vagrant_2.2.7_x86_64.deb

# centos
sudo curl
http://download.virtualbox.org/virtualbox/rpm/rhel/virt
ualbox.repo > /etc/yum.repo.d/virtualbox.repo
yum update && yum install VirtualBox-5.1
yum install vagrant 2.2.7 linux amd64.zip
```

Start a Vagrant VM

```
# on host OS, start a new VM
mkdir centos7
vagrant init centos/7 # vagrant init sylabs/singularity-3.5-centos-7-64
vagrant up
# enter VM
vagrant ssh
ls /vagrant # directory from host OS shared here
# install docker and singularity
sudo -s
yum install -y epel-release
yum-config-manager --add-repo https://download.docker.com/linux/centos/docker-ce.repo
yum install -y singularity
yum install -y docker-ce && systemctl start docker
# exit out of VM and delete it
^d^d
vagrant snapshot save clean-install # vagrant destroy
```

Lets create a pytorch container!

- Find useful/official/collaborator's image (or build your own from scratch):
 - https://hub.docker.com/search?q=pytorch&type=image
 - Numerous versions: look at TAGs to determine differences
- Pull pytorch image from Dockerhub and run using singularity:
 - o singularity pull \
 docker://pytorch/pytorch:1.5-cuda10.1-cudnn7-runtime
 - Creates new container image at

```
pytorch_1.5-cuda10.1-cudnn7-runtime.sif
```

- A container, in singularity, is a single file
- You need singularity installed on the host that you intend to run it on
- All of your programs are inside the container
- So to run the containerized application, you need to

```
o singularity exec \
    pytorch_1.5-cuda10.1-cudnn7-runtime.sif \
    python [args...]
```

Can also bring up a shell inside the container with

```
o singularity shell \
    pytorch_1.5-cuda10.1-cudnn7-runtime.sif
```

- By default, only the working directory is 'mounted' into the container
- Need to add 'bind mounts' to singularity command
- singularity shell \

```
pytorch_1.5-cuda10.1-cudnn7-runtime.sif
Singularity> ls /gpfs
ls: cannot access '/gpfs': No such file or directory
```

• singularity shell -B /gpfs \
 pytorch_1.5-cuda10.1-cudnn7-runtime.sif
 Singularity> ls /gpfs
 automountdir slac

Run pytorch container with GPU

```
$ singularity exec
pytorch 1.5-cuda10.1-cudnn7-runtime.sif
python
Python 3.7.7 (default, Mar 23 2020,
22:36:06)
[GCC 7.3.0] :: Anaconda, Inc. on linux
Type "help", "copyright", "credits" or
"license" for more information.
>>> import torch
>>> print('Using device:',
torch.device('cuda' if
torch.cuda.is available() else 'cpu'))
Using device: cpu
```

```
$ singularity exec --nv
pytorch 1.5-cudal0 1-cudnn7-runtime.sif
python
Python 3.7.7 (default, Mar 23 2020,
22:36:06)
[GCC 7.3.0] :: Anaconda, Inc. on linux
Type "help", "copyright", "credits" or
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>>> print('Using device:',
torch.device('cuda' if
torch.cuda.is available() else 'cpu'))
Using device: cuda
```

But I need uproot!



- ... or any other library or module or binary
- You could create a singularity 'sandbox' or --writeable and add it all manually, but...
 - You would be left with an non-reproducible container image
 - You'll have to manually find ways to version control etc.
- Contribute back!
 - Build new docker image with changes
 - Push new container into dockerhub (with tags and metadata)
 - Pull image back as singularity image to run
- Why not use Singularity directly? (ie write Singularity recipe)

Create a Dockerfile with additions

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• \$ cat Dockerfile

```
FROM pytorch/pytorch:1.5-cuda10.1-cudnn7-runtime RUN pip install uproot
```

Build it with docker

>>>

- sudo docker build . -t slaclab/pytorch-test:1.0
- Push it to dockerhub (need account + docker login)
 - \$ docker push slaclab/pytorch-test:1.0
- Pull new image with singularity
 - \$ singularity pull docker://slaclab/pytorch-test:1.0
- Run
 - \$ singularity exec pytorch-test_latest.sif python
 ...
 >>> import uproot

Limitations

- Building can take a while...
 - But ultimately worth it as a means to
 - Document installation methods (Dockerfile) git commit!
 - Have multiple versions as things evolve
 - Build once, run anywhere
 - Pushing and Pulling images network limited
 - Can minimise size through intermediate images etc.
- Compiling GPU code requires GPUs
 - requires GPU node + sudo + nvidia-docker

sudo?

- Use TravisCI (or other) to auto build and push docker images
- podman still early days, little gpu support

Useful Docker images

- Base Images
 - https://hub.docker.com/r/continuumio/miniconda/
 - https://hub.docker.com/_/python
 - https://hub.docker.com/r/tensorflow/tensorflow/
 - https://hub.docker.com/r/pytorch/pytorch
 - https://hub.docker.com/r/nvidia/cuda
 - https://hub.docker.com/ /centos
 - https://hub.docker.com/ /ubuntu
- Ubuntu images generally easier, more up-to-date
- Try to avoid alpine for scientific (musl instead of glibc)

- Each RUN, COPY, RUN command creates a new 'layer'
 - Keeps a snapshot of all changes caused (eg on FS) by that command
 - More layers means a larger container
 - Building containers etc. will
 - Leave behind gcc, make, etc
 - yum/apt cache files and other temp files
 - Ensure you delete at the end of the RUN command
 - Can use multi-stage builds to COPY files from one image to another
- Copy sensitive files:
 - Use BuildKit:
 - Add to Dockerfile

```
# syntax=docker/dockerfile:experimental
RUN --mount=type=secret,id=license id
```

■ sudo DOCKER_BUILDKIT=1 docker build \
--secret id=license id,src=./license id.txt .

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Let's submit a job



- Slurm will replace LSF on SDF
- All GPUs in slurm
- Use a GPFS space (for now) for your working directory
- AFS (Home) not readily available (no tokens)

Slurm - example submission script

SLAC

```
#!/bin/bash
#SBATCH --account=shared --partition=shared
#
#SBATCH --job-name=test
#SBATCH --output=output-%j.txt --error=output-%j.txt
#
#SBATCH --nodes=1 --ntasks=1
#SBATCH --cpus-per-task=1
#SBATCH --mem-per-cpu=1q
#SBATCH --time=10:00
#SBATCH --gpus=1
# cd /gpfs/slac/cryo/fs1/u/ytl/containers/
singularity exec --nv -B /gpfs pytorch-test 1.0.sif \
     python test.py
```

Whilst we're testing...:

```
$ ssh
slacgpu.slac.stanford.edu
$ module load slurm
```

Submit with

\$ sbatch <script_path>

Slurm - common commands

sbatch	Submit batch job		
squeue	Look at the job list		
scancel <job id=""></job>	Cancel job id		
scontrol show job	Show detailed job information		
sstat	Show per time step job usage		
scontrol update job	Update job details		
srun	Connect to an allocated job		

Slurm - requesting GPUs

		Available	GPU Memory	FP32 (TFLOPS)	FP64 (TFLOPS)
gpus=1	Any available	~325			
gpus=geforce_gtx_1080_ti:1	Geforce 1080ti	~100	11GB	~11.3	~0.3
gpus=geforce_rtx_2080_ti:1	Geforce 2080ti	~200	11GB	~13.4	~0.4
gpus=v100:1	Tesla v100	~24	32GB	~15.7	~7.8

Summary



- SDF will be our new platform for Scientific Compute and Storage
 - Feedback and suggestions please!
- Containers are a great way of packaging programs
 - Shareable and repeatable (spend less time getting things working)
 - Build once, run anywhere (version control!)
 - Small additional prefix to run
- Still a little early...
 - sudo required
 - Network limited (some containers can be 10GB+)
 - GPU support (especially during building) under development
- Submit to SLURM