

NIH: Data Science with Docker

Containers Workflow Interest Group (NIH/NCI)

June-2024

A word from Docker's CEO





Scott Johnston CEO



How to participate in today's session

For online viewers:

• Please place questions into chat. We will either address as we go or discuss at the end, time pending

BUT wait... we only have a limited time today, right?

• This session is the first we intend to hold as well as more collaborative workshops. You will receive a survey which you can select topics you would like to review next session.



Your Docker account team





Dylan Tegarden Account Executive



Jessica Squizzato Customer Success

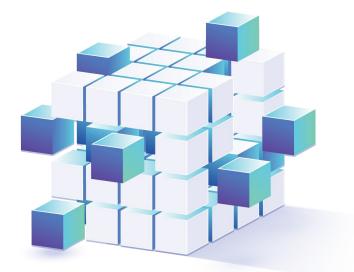


Todd Densmore Solution Architect



Agenda

- 1. Who is Docker?
- 2. What is a Container?
- 3. Modern Data Science
- 4. Data Science in the Cloud
- 5. Data Science Workflows
- 6. Kubeflow Pipelines
- 7. Closing Remarks







Who is Docker?

Docker

Docker created software "containers"

- dotCloud Inc. was founded in 2011
 - \circ $\,$ a platform as a service (PaaS) for developers
 - run their applications in the cloud
- renamed to **Docker Inc.** in 2013
- Docker containers debuted to the public in Santa Clara at PyCon in 2013





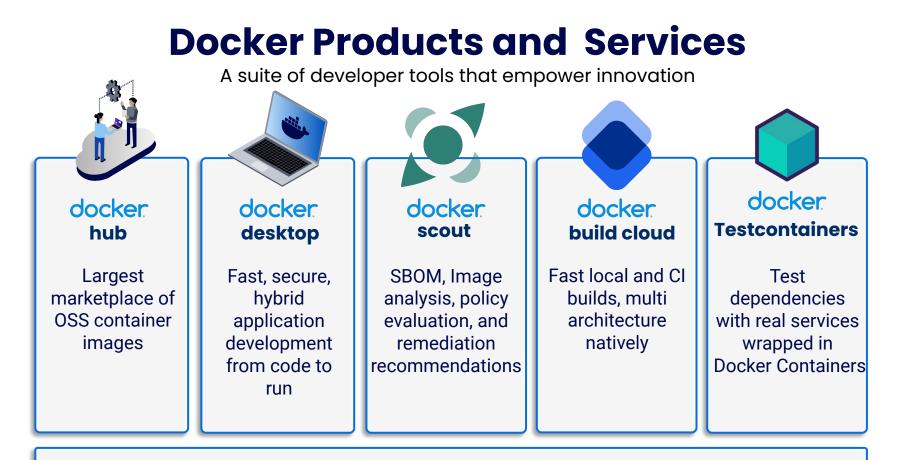
The container specifications

- Docker formed the Open Container Initiative in June 2015
- The purpose is to govern standard image formats and runtimes
- Currently run by the Linux Foundation
 - **image-spec** build images (<u>BuildKit</u>, podman, buildah)
 - **runtime-spec** run containers (containerd, CRI-O, Windows Containers)
 - **distribution-spec** share images (<u>Docker Hub</u>, Artifactory, ECR, GCR, ACR)



At Docker We Believe It Is Critical To Empower Developers Data Scientists









What is a Container?

containers

A collection of Linux technologies to package and run **lightweight** application processes **securely**. Software "containers" are similar to shipping containers

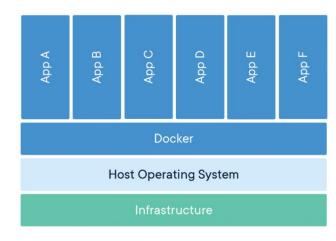


Containers run reliably across systems (MacOS, Windows, Linux) and platforms (GCP, AWS, Azure)

Container technology

Features of the Linux kernel to allow multiple processes to run in isolation

- Union filesystem single view of multiple file layers
- Chroot change the root directory for a process family (jails)
- Namespaces provides isolation for kernel resources
 - User, PID, network, mount, IPC, UTS, etc.
- **Cgroups** resource limits for (namespaced) processes
 - CPU, memory, disk I/O, network, etc.



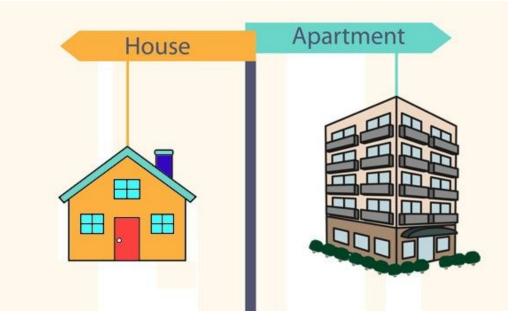
Houses vs Apartments

House has its own

- Water
- Heat
- Electricity

Apartment shares

- Water
- Heat
- Electricity





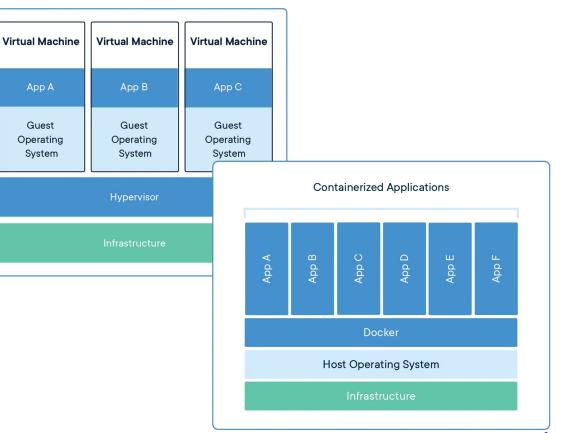
VMs vs Containers

VM virtualizes its own

- CPU
- Memory
- Disk

Container shares

- CPU
- Memory
- Disk



Benefits of Software Containers

Efficient

Containers require less system resources (they don't include operating system things)

• Portable

Can be run on multiple different operating systems / hardware platforms / clouds

Consistent

applications in containers will run the same (they include dependencies)

• Scalable

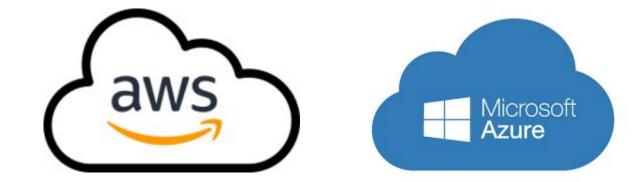
applications can be rapidly deployed, patched, or scaled

• Application development

Containers support accelerate development, test, and production cycles

Result: Containers Power every Cloud





Google Cloud







Modern Data Science

Golden Age of Data Science

Everything is connected to the Internet

Massive amounts of data being generated (<u>zettabytes</u>)

- Mobile devices
- Cars
- Medical data

Improvements in cloud computing

• Massive compute power available on-demand

Advances in Artificial intelligence





Some challenges



Problem: Managing Data

Data sets are becoming VERY large (GB-TB)

Providing data sets quickly (on-demand)

- Imaging Data Commons
- Genomic Data Commons

Storing (archiving) data sets

Multiple formats

Data Integrity (Accurate, complete, and valid)



Genomic Data

The original human genome project (Celera) is based on 5 anonymous donors, however this is being expanded.

The current reference human genome is 3.1 billion base pairs

GRCh38.p14 (July 2023) is <u>3.1 Gb (2 Gb compressed)</u>

But... the volume of genomics data is doubling every 7 months

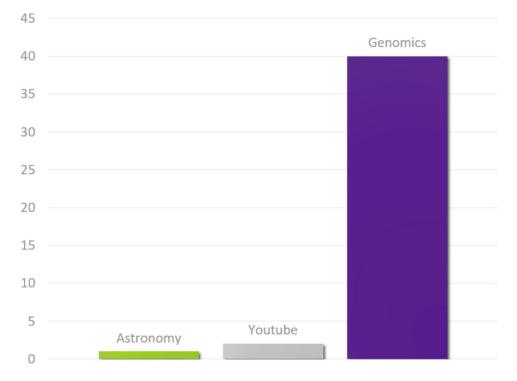
By 2025, the predicted total genomics data capacity is 40 exabytes

(40,000 petabytes)



Genomic Data vs. YouTube

Projected Annual Storage 2025 (Exabytes/Year)







Data Science in the Cloud

Benefits: Cloud Platforms for Data Science (1)

Compute scalability

- Enables massive bursts of computation
- close to the data easily attach to terabytes of data

Accessibility

- Share data easily (14 million medical images with annotations)
- Share data globally

On-demand specialized Tools

• Machine learning tools



Benefits: Cloud Platforms for Data Science (2)

Cost savings

• Pay by usage (hour / query / terabyte)

Reliability

- Availability guaranteed uptime (99.999% == 5 mins 15 seconds)
- Fault tolerance / redundancy (data and services)

Security

• data security and isolation (regions are very important)



Example: Machine Learning

Specialized Hardware: GPUs

Example: a Google a3-megagpu-8g machine has:

- 8 NVIDIA A100 GPU (208 virtual CPUs)
- 1,872 GB RAM
- 6.4 TB storage

Specialized Frameworks:

- Google Cloud AI Platform
- Tensorflow



NIH Cloud Platforms: STRIDES

STRIDES - Science and Technology Research Infrastructure for Discovery, Experimentation and Sustainability

https://cloud.nih.gov/resources/cloudlab/

- Amazon Web Services
- Google Cloud
- Microsoft Azure





Data Science Workflows

What is Kubernetes (K8S)?

The standard way to define and manage containerized applications



- Open Source
- Works across all cloud providers (or on-prem)
- Originally created by Google (Borg)
- Greek word for "helmsman" or "(ship) pilot"
- Handles networking, secrets, scaling, data storage, failover, deployments, etc



What is Kubeflow?

A simple way to define and run ML workflows

Runs on Kubernetes (based on containers)

Standardizes ML Ops

Provides best-in-class tools for data scientists

- Notebooks (JupyterLab, RStudio, and Visual Studio Code)
- Data pipelines (ML workflows)
- Model <u>training</u> using <u>PyTorch</u>, <u>TensorFlow</u>, <u>XGBoost</u>
 - or BYOM (<u>HuggingFace</u>, <u>DeepSpeed</u>, or <u>Megatron-LM</u>)
- Model serving



Kubeflow





Kubeflow Pipelines

What is a Workflow?

A formal series of (software) steps

Each steps performs a specific task:

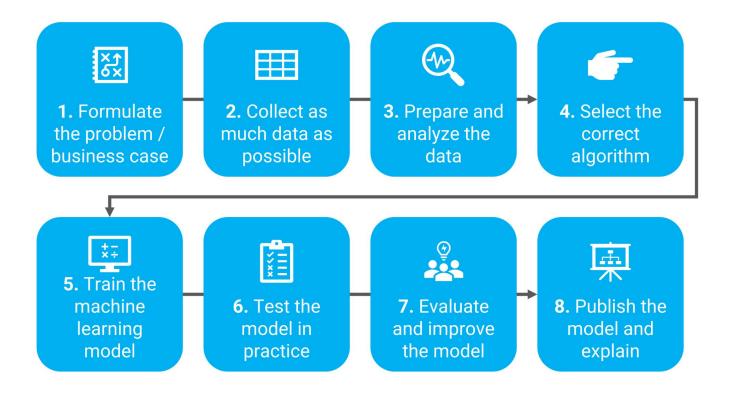
- Input massive amounts of data
- Process (transform) and analyze the data
- Outputs (including metadata and data provenance)

Guidelines

FAIR principles - findability, accessibility, interoperability, and reuse (of assets) for humans and computers



Example: Data Science Workflow





Example Kubeflow pipeline



Kubeflow Pipeline Code: Python

```
import kfp
from kfp import dsl
```

```
def load_data_op():
    return dsl.ContainerOp(
        name="Load Data",
        image="python:3.7",
        command=["sh", "-c"],
        arguments=["echo 'Loading data' && sleep 5"],
```

```
def preprocess_data_op():
```

```
return dsl.Container0p(
    name="Preprocess Data",
    image="python:3.7",
    command=["sh", "-c"],
    arguments=["echo 'Preprocessing data' && sleep 5"],
```

def train_model_op():

```
return dsl.Container0p(
    name="Train Model",
    image="python:3.7",
    command=["sh", "-c"],
    arguments=["echo 'Training model' && sleep 5"],
```

@dsl.pipeline(

name="My First Pipeline", description="A simple pipeline that demonstrates loading, preprocessing, and training steps."

```
def my_first_pipeline():
```

```
load_data = load_data_op()
preprocess_data = preprocess_data_op().after(load_data)
train_model = train_model_op().after(preprocess_data)
```

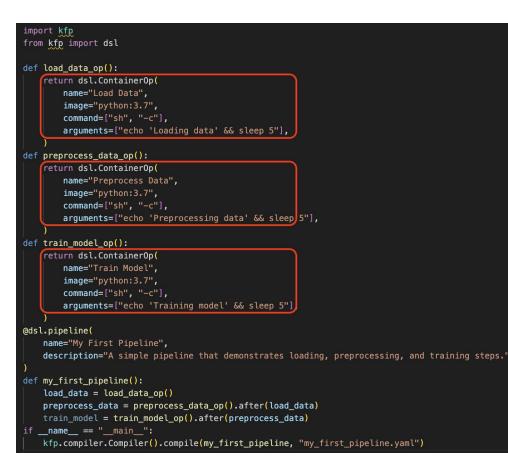
Write a pipeline in familiar Python

Each python function is a step in the pipeline

Can have as many steps as needed to load, transform and analyze data



Kubeflow Pipeline Code: Python



The KFP library converts each python function into a versioned container

The whole pipeline is transformed into protobuf YAML

Can import directly into Kubeflow or build pipelines in a Jupyter Notebook.

Docker benefits for Workflows

Portable

- Facilitates research collaboration
- Regulatory questions answered / data retention

Reproducible

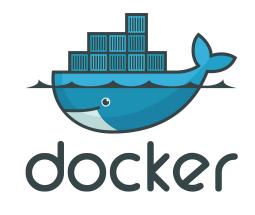
- Generate the same result from the same data, software (packages, versions), compute environment
- Workflow results can be independently verified

Scalable

- Containers consume less resources than VMs
- Quickly scaled up or down to accommodate different data sets

Secure

• Containers are run in isolation





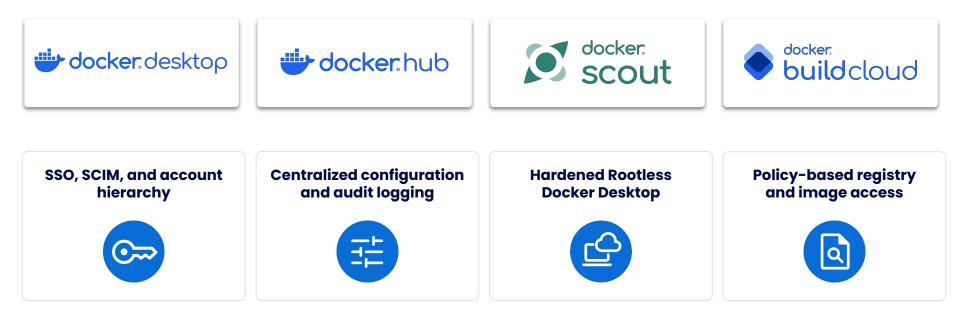


Closing Remarks



Docker Business Subscription

A suite of developer tools that empower innovation



Contact dylan.tegarden@docker.com for pricing





Thank You!