

ClearML guide

Tags

clearml

mlops

To Do

- Clean and update notes
- Better logging
- Data versioning
- Deployment and monitoring
- Pipelines
- CI/CD

To learn

Setup

```
pip install clearml
```

Set up connection with the server.

```
clearml-init
```

Go to [app_clear_ml](#) and head over to [settings/workspace](#).

Click on *create new credentials*. Copy these and paste in the terminal.

ClearML Agent

Used to execute tasks on remote servers. Can also schedule tasks etc.

Reference link → https://clear.ml/docs/latest/docs/clearml_agent/

```
pip install clearml-agent  
clearml-agent init
```

- Agent can run in *pip*, *conda*, *poetry* and *docker mode*.

To run in [docker](#) mode

```
clearml-agent daemon --docker
```

After this, the `agent` will start listening for tasks on the default `queue`.

Initialize the agent before putting the task in queue.

- We can put multiple tasks in the queue and agent will run them one after the other.
- Can also pass desired image to the agent

```
clearml-agent daemon nvidia/cuda:11.7.1-cudnn8-runtime-ubuntu22.04
```

Notes

- Import and create task.

```
import tensorflow as tf
from clearml import Task

# Task.add_requirements("./requirements.txt")

task = Task.init(project_name="mnist_example", task_name="local_training")

parameters = {"lr": 0.001}

mnist = tf.keras.datasets.mnist
(x_train, y_train), (x_test, y_test) = mnist.load_data()

model = tf.keras.models.Sequential(
    [
        tf.keras.layers.Flatten(input_shape=(28, 28)),
        tf.keras.layers.Dense(128, activation="relu"),
        tf.keras.layers.Dropout(0.2),
        tf.keras.layers.Dense(10),
    ]
)

loss_fn = tf.keras.losses.SparseCategoricalCrossentropy(from_logits=True)

task.connect(parameters)

model.compile(
    optimizer=tf.keras.optimizers.Adam(learning_rate=parameters["lr"]),
    loss=loss_fn,
    metrics=["accuracy"],
)

model.fit(x_train, y_train, epochs=5)
model.evaluate(x_test, y_test, verbose=2)
```

- Command-line arguments are captured automatically.
- Logs, source codes, hyper-parameters, plots and images are all captured automatically.
- For `dict` and python `objects` we can use `Task.connect()`.

```
parameters = {"change_me": "value", "dropout": 0.5}

task.connect(parameters)
```

- We can clone the task, change hyper-parameters and execute the task remotely.
- Experiment and data tracking
 - *Dataset version control*
 - Compare multiple experiment runs, keep track of experiment configurations etc.
- **Artifacts** like model weights, processed features, files are also stored/captured
 - Model artifacts can exist as standalone entities
 - We don't need to find the original task and then try to get the attached artifact
 - We can pull the model by its `id` or `tag`
- **Scalars**
 - Loss, accuracy etc. are stored and plotted
- Compare multiple experiment using UI
- Can also add *requirements*

```
Task.add_requirements("./requirements.txt")
```

- Reporting scalars

```
Logger.current_logger().report_scalar(
    "train", "accuracy", value=train_acc, iteration=epoch
)
```

- We can create project and run multiple experiments inside that project

Experiment Tracking

Examples

https://github.com/pytholic/ClearML/blob/main/experimentation/pytorch/pytorch_test.py

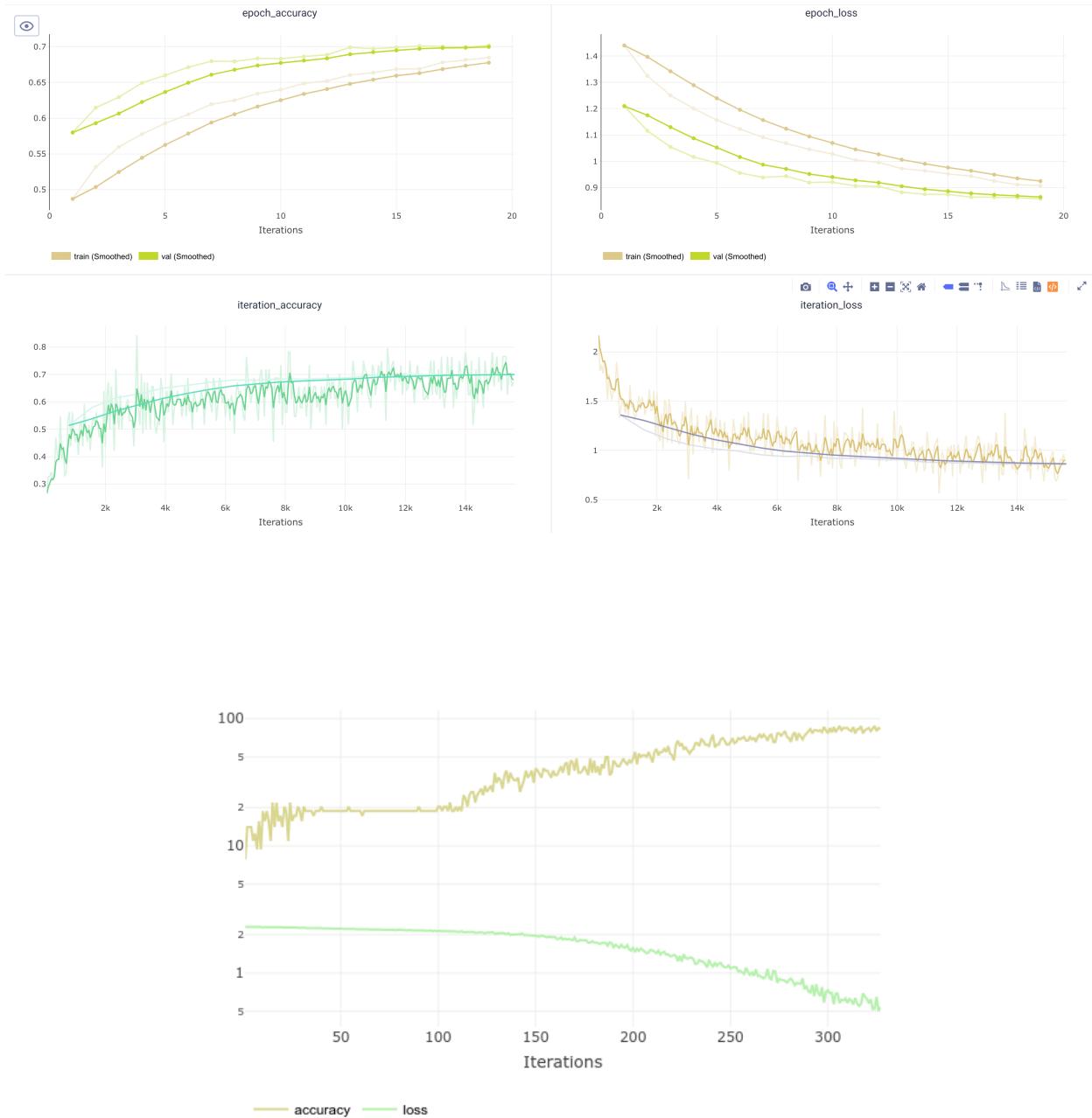
https://github.com/pytholic/ClearML/blob/main/experimentation/pytorch_lightning/cifar10_tensorboard_logger.py

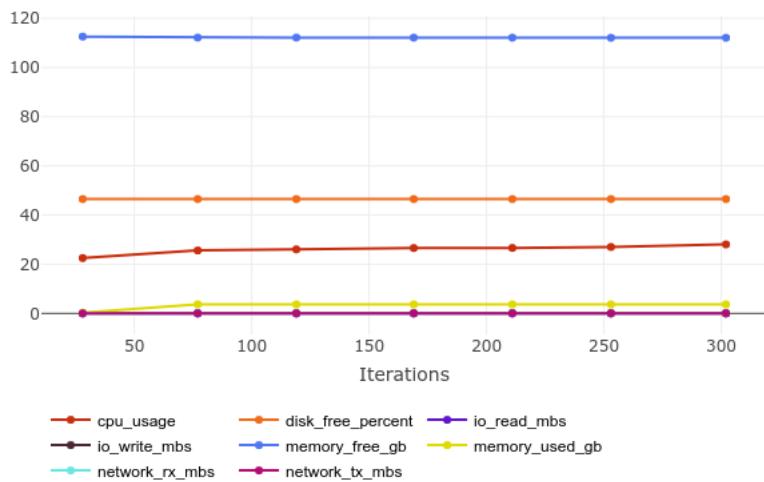
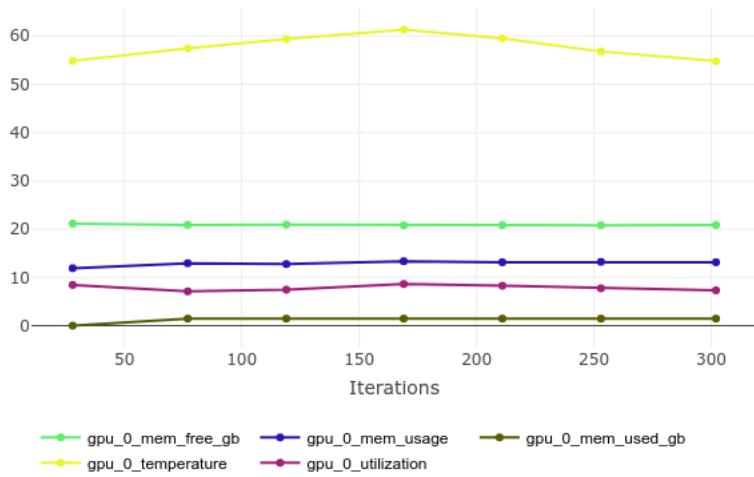
Testing on CIFAR-10 dataset. We can log loss and accuracy.

In addition, we can also get machine and gpu usage.

There are various kinds of loggers available especially in pytorch-lightning. I am using tensorboard below for better customization.

Ref → <https://pytorch-lightning.readthedocs.io/en/stable/extensions/logging.html>





Links

Tutorial

https://www.youtube.com/watch?v=kyOfwVg05EM&list=PLMdIIcuMqSTnoC45ME5_JnsJX0zWqDdlO&index=4

Hyperparameter Optimization

One way can manually writing the configurations

```
# Define the hyperparameters for each task
hyperparameters = [
    {'learning_rate': 0.01, 'batch_size': 32, 'epochs': 5},
    {'learning_rate': 0.001, 'batch_size': 64, 'epochs': 10},
    {'learning_rate': 0.0001, 'batch_size': 128, 'epochs': 20}
]

# Train the CNN with each set of hyperparameters
for hyperparameters in Hyperparameters:
    learning_rate = hyperparameters['learning_rate']
    batch_size = hyperparameters['batch_size']
    epochs = hyperparameters['epochs']
```

There are also optimizers available in clearml.

Example link

https://github.com/pytholic/ClearML/blob/main/experimentation/hyperparameter/hyperparameter_optuna.py

Ran bunch of experiments with different configurations.

```
hyper_parameters=[  
    UniformParameterRange("Args/lr", min_value=0.01, max_value=0.3, step_size=0.05),  
    DiscreteParameterRange("Args/net", values=["Net1", "Net2"]),  
    DiscreteParameterRange("Args/epochs", values=[10, 30, 50]),  
]
```

We get a nice comparison in the end with descriptive titles for each experiment. All the models artifacts are saved as well for each run.



To put a time limit or task limit on the hyper-parameter optimization process.

Can use following parameter

```
optimizer.set_time_limit(in_minutes=2.0)
```

We can also set time for each job. This will limit the execution time of a single experiment, in minutes.

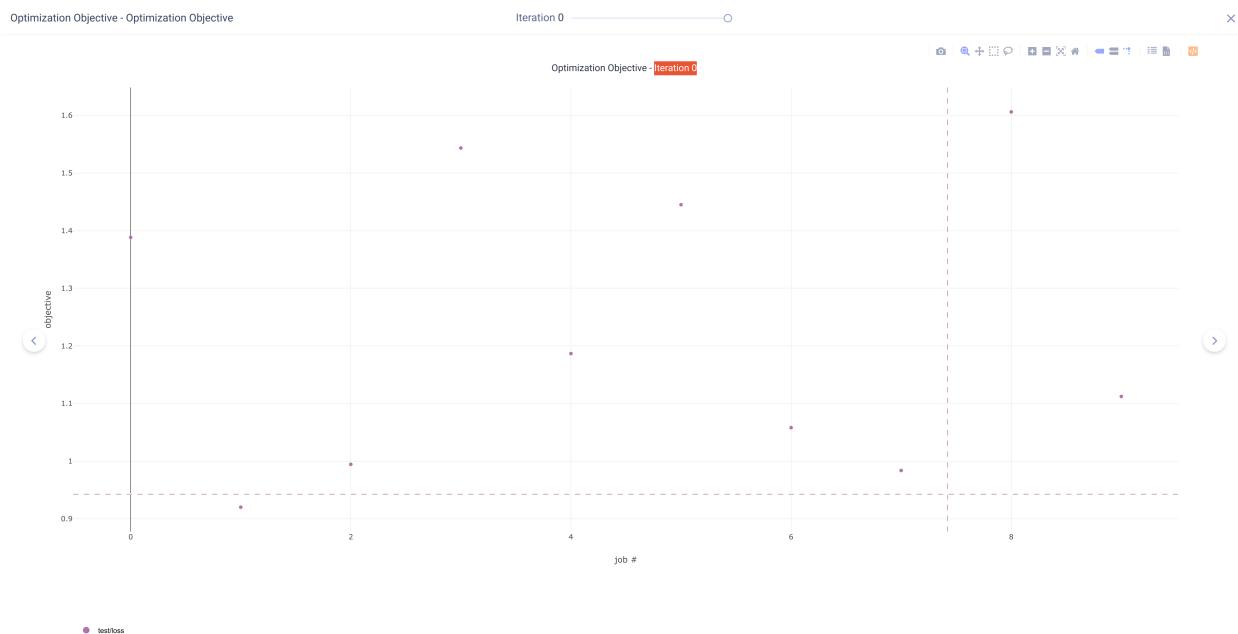
```
time_limit_per_job=10
```



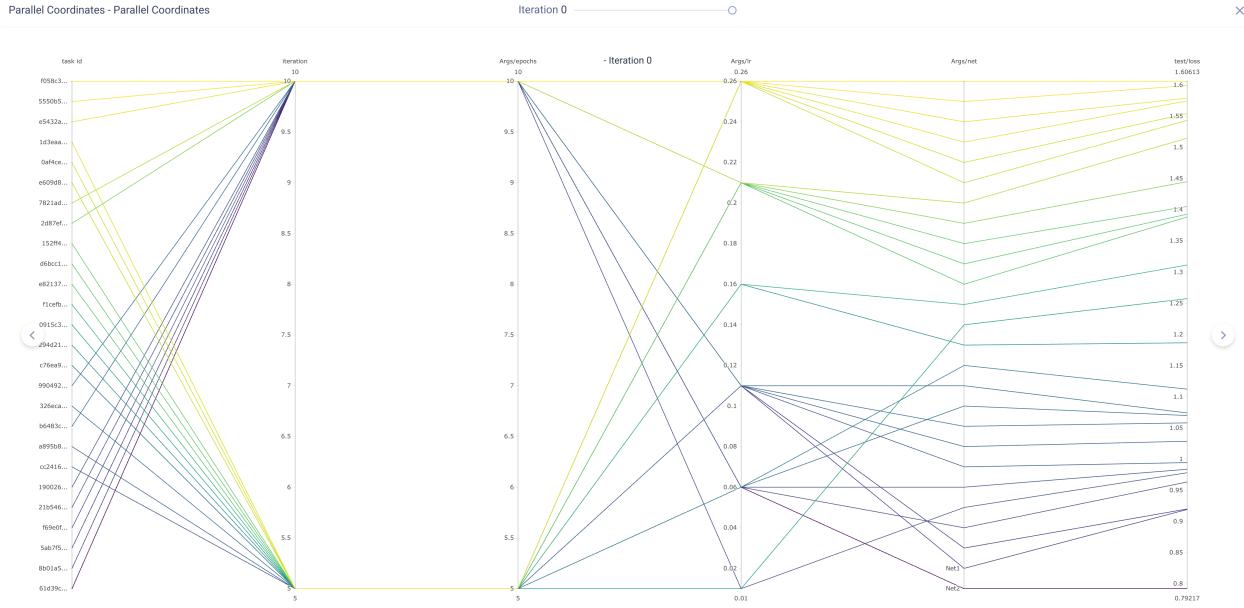
Experiments that are in progress when the optimizer stop will continue to run. However, artifacts might not be stored since the main process stopped.

Plots

Can see the final objective for all the jobs



Different task configurations and final metrics



Objective leaderboards

summary - objective: test/loss

Iteration 0

objective: test/loss - iteration 0

task_id	objective	iteration	Args/epochs	Args/lr	Args/net	status
61039c3fe8d2424aa2ed4de9be47ba	0.7921672959327698	10	10	0.060000000000000005	Net2	completed
8b01a53635fc8d49fa53ab15735e70c	0.9192112194061279	10	10	0.11	Net2	completed
5eb7f59e1b424028a3184cab2598e3	0.9200062901496887	10	10	0.11	Net2	completed
f6f0edfacc0ef4c218ac03101d280d8d	0.963354161453247	10	10	0.060000000000000005	Net1	completed
21b546ea90b54644805445509c9757	0.9781629543304443	10	10	0.01	Net2	completed
190026875942129a62bf4aecba333a	0.983741926574707	10	10	0.060000000000000005	Net1	stopped
cc2414ec4e314459e9e2226abfbcb9d	0.99442575035095	5	5	0.11	Net2	completed
a895089332324f69528ca0aa49f900	1.0286710178375245	5	5	0.11	Net2	completed
b6483ce70ff0483c8c00e244bac78493	1.0531494047954646	10	10	0.11	Net1	completed
326eead7f049487ba7a56d40c18564ab	1.0699488346099852	5	5	0.060000000000000005	Net1	completed
990497033ec4f28842524216c153b	1.074080147377793	10	10	0.11	Net1	completed
c76ea946394545sead983053a56011	1.112258788726806	5	5	0.060000000000000005	Net1	completed
2942d19a0d4cadb2aef9ce2090222926	1.1865906656188965	5	5	0.160000000000000003	Net2	completed
0915c3576bb4b3195261042025338	1.257453558700561	5	5	0.01	Net2	completed
f1c0ff5aa224af88da596336d62cc	1.3114248375892639	5	5	0.160000000000000003	Net2	completed
e821376343f6459e93332f68e3fa990b	1.3884045497971822	5	5	0.210000000000000002	Net2	completed
d6bcc13e21942588d2eeb3cc6df1711	1.3919453090896606	5	5	0.210000000000000002	Net2	completed
1527fae59565400c08f024222369e9a6	1.405269543548583	5	5	0.210000000000000002	Net1	completed
2d87ef5e23d245e094cf592708765c9	1.4450830711135863	10	10	0.210000000000000002	Net2	completed
7821e6abddc6ff4aa0266cf02537	1.5148873495101929	10	10	0.210000000000000002	Net2	completed
e60988819c9b400484e43580515e129	1.5434323788467406	5	5	0.26	Net2	completed
0a44c7a721aa7b08230aae43e176	1.5542615737391504	5	5	0.26	Net2	completed
1d3eae977224a45f8b007fd1404108765	1.574623917099	5	5	0.26	Net1	completed
e5432a019b41fc943b75aff79d702c	1.5790243570327758	10	10	0.26	Net1	completed
555050501fa4122bdddff08e9155	1.5990495563507081	10	10	0.26	Net2	completed
f058c3ab8b184691923d44e18efc30e7	1.6061320095062255	10	10	0.26	Net2	stopped

Top experiments

We can also return to `top_k` experiments which gave best performance.

```
# Return a list of top performing experiments
top_exp = optimizer.get_top_experiments(top_k=3)
print([t.id for t in top_exp]) # print just the id
```

```
# Return a dict of top performing experiments with more details
top_exp = optimizer.get_top_experiments_details(top_k=3)

[{'task_id': '3d5c98024baa47bc95e065715ca6724b', 'hyper_parameters':
 {'Args/net': 'Net2', 'Args/epochs': '10', 'Args/lr': '0.11'},
 'metrics': {'test/loss': {'metric': 'test', 'variant': 'loss',
 'value': 0.9133659139633179, 'min_value': 0.9133659139633179,
 'max_value_iteration': 10, 'max_value': 1.9955348001480102, 'max_value_iteration': 1}},
 {'task_id': '62328586c8034ae6969151e205c0a90b',
 'hyper_parameters': {'Args/net': 'Net2', 'Args/epochs': '5', 'Args/lr': '0.11'},
 'metrics': {'test/loss': {'metric': 'test', 'variant': 'loss',
 'value': 1.0147933155059814, 'min_value': 1.0147933155059814,
 'max_value_iteration': 5, 'max_value': 1.697532457923889,
 'max_value_iteration': 1}},
 {'task_id': 'db87c34586c945ffb33cbcef21a66a82',
 'hyper_parameters': {'Args/net': 'Net1', 'Args/epochs': '10', 'Args/lr': '0.11'},
 'metrics': {'test/loss': {'metric': 'test', 'variant': 'loss',
 'value': 1.1441912442207336, 'min_value': 1.1441912442207336,
 'max_value_iteration': 5, 'max_value': 1.500229824066162,
 'max_value_iteration': 1}}]
```

I had to place it between `wait` and `stop` for it to work.

```
# wait until optimization completed or timed-out
optimizer.wait()

# top_exp = optimizer.get_top_experiments(top_k=3)
# print([t.id for t in top_exp])

top_exp = optimizer.get_top_experiments_details(top_k=3)

# make sure we stop all jobs
optimizer.stop()
```

Job callbacks

We can also add `job_complete_callback` to get console message whenever a job is completed along with the experiment details.

```
Job completed! 3d5c98024baa47bc95e065715ca6724b 0.9133659139633179 10 {'status': 'in_progress', 'Args/lr': 0.11, 'Args/net': 'Net2', 'Args/total_max_jobs': 10}
Top performance experiment id: 3d5c98024baa47bc95e065715ca6724b
```

Optuna Optimizer

Need to have [optuna](#) installed.

```
pip install optuna
```

Seems like there are some required *parameters*.

```
TypeError: OptimizerOptuna.__init__() missing 2 required positional arguments: 'max_iteration_per_job' and 'total_max_jobs'
```

Parameters for `OptimizerOptuna` are slightly different from normal `HyperparameterOptimizer`.

We just set the class and parameters and run like the last example.

```
optimizer_class=OptimizerOptuna
```



One useful option `total_max_jobs` is also available with `OptimizerOptuna`. It was missing in normal `RandomSearch` optimizer.

Links

Documentation

<https://clear.ml/docs/latest/docs/fundamentals/hpo>

Tutorial

https://www.youtube.com/watch?v=aY1ImY4OnKk&list=PLMdICuMqSTnoC45ME5_JnsJX0zWqDdIO&index=6

ClearML Data

Use ClearML Data to create, manage, and version your datasets.

Accessibility → Data can be accessed from any machine

Versioning → Linking which data set version was used in which task

This helps in reproducibility.

Creating dataset with CLI

Steps

Initialize the dataset

First we need to `create` an initial dataset version.

We can do it with CLI.

```
clearml-data create --project experimentation/clearml-data --name animal-data
```

Add files

This will initialize the dataset, but it will be empty. We can then use `add` command to add files.

```
clearml-data add --files ./images
```

```
clearml-data - Dataset Management & Versioning CLI
Adding files/folder/links to dataset id 399a206cdda04949b262f11a553552a5
Generating SHA2 hash for 90 files
100%|██████████| 90/90 [00:00<00:00, 100.00B/s]
Hash generation completed
90 files added
```

This will recursively add files to the dataset.

Close

Now we need to tell the server that we are done here.

```
clearml-data - Dataset Management & Versioning CLI
Finalizing dataset id 399a206cdda04949b262f11a553552a5
Pending uploads, starting dataset upload to https://files.clear.ml
Uploading dataset changes (90 files compressed to 3.96 MiB) to https://files.clear.ml
File compression and upload completed: total size 3.96 MiB, 1 chunk(s) stored (average size 3.96 MiB)
Dataset closed and finalized
```

It will *upload* the files and change the dataset status to *finalized*. This will lock this version of the dataset.

Creating dataset with Python SDK

```
import os
from pathlib import Path

from clearml import Dataset

DATA_ROOT = "./clearml-data/sample_data/images"

# # Create dataset
ds = Dataset.create(dataset_name="animal-dataset", dataset_project="clearml-data")

# # Add files
ds.add_files(path=DATA_ROOT)

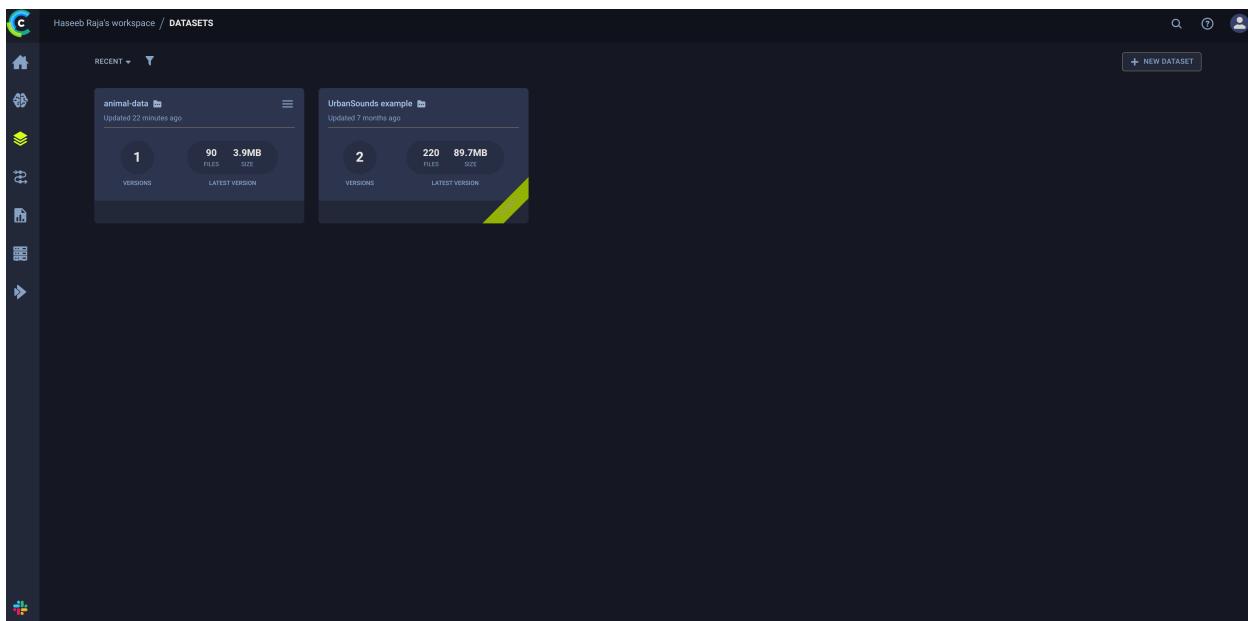
# Count the amount of samples per class
root_folder = Path(DATA_ROOT)
counts = []
folders = sorted(os.listdir(root_folder))
for folder in folders:
    counts.append([len(os.listdir(root_folder / folder))])

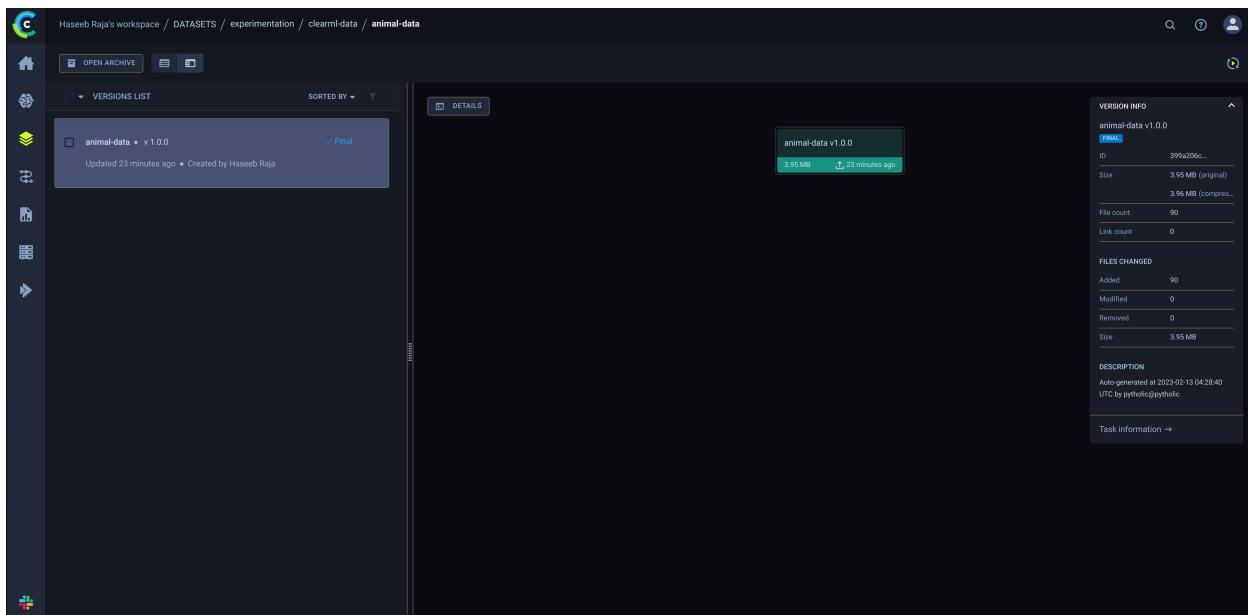
# Log some dataset statistics
ds.get_logger().report_histogram(
    title="Dataset Statistics",
    series="Train Test Split",
    labels=folders,
    values=counts,
)

# Finalize and upload
# ds.upload()
# ds.finalize()
ds.finalize(auto_upload=True)
```

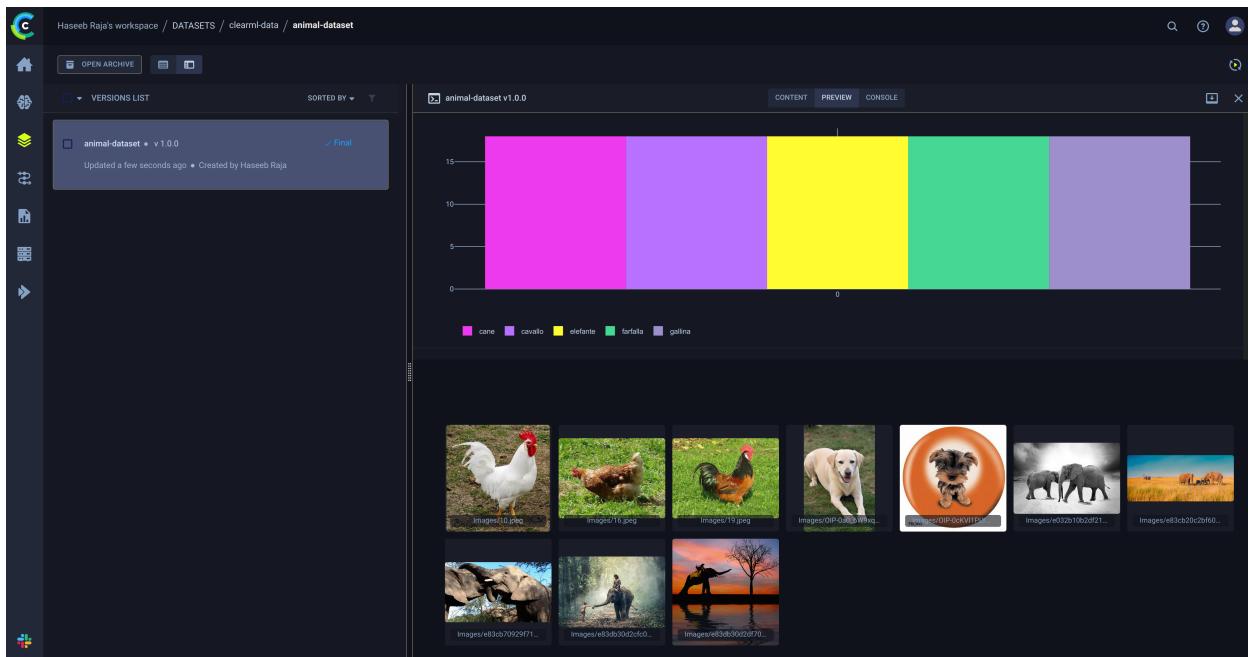
```
ClearML results page: https://app.clear.ml/projects/23aaa92afe0d4c919b74c9770620ad77/experiments/224508a1e13b4d96a1a45209a3f0c049/output/lo  
ClearML dataset page: https://app.clear.ml/datasets/simple/23aaa92afe0d4c919b74c9770620ad77/experiments/224508a1e13b4d96a1a45209a3f0c049  
Generating SHA2 hash for 90 files  
100%|██████████| 3.96M/3.96M [00:00<00:00, 1.00MB/s]  
Hash generation completed  
Pending uploads, starting dataset upload to https://files.clear.ml  
Uploading dataset changes (90 files compressed to 3.96 MiB) to https://files.clear.ml  
File compression and upload completed: total size 3.96 MiB, 1 chunk(s) stored (average size 3.96 MiB)
```

Dataset UI





We can also see *samples* and *histogram*.



Downloading dataset

You can download this version from any machine.

CLI

```
clearml-data get --id 224508a1e13b4d96a1a45209a3f0c049  
# Get in desired folder  
clearml-data get --name "animal-data" --copy /clearml-data/tmp/my_dataset_2
```

```
clearml-data - Dataset Management & Versioning CLI  
Download dataset id 224508a1e13b4d96a1a45209a3f0c049  
Dataset local copy available for files at: /home/pytholic/.clearml/cache/storage_manager/datasets/ds_224508a1e13b4d96a1a45209a3f0c049
```

You can find it from the UI or by searching.

```
clearml-data search --name "animal-data" # can also use tags/projects to search
```

```
clearml-data - Dataset Management & Versioning CLI  
Search datasets  
project | name | tags | created | id  
-----  
clearml-data | animal-dataset | | 2023-02-13 05:15:22 | 224508a1e13b4d96a1a45209a3f0c049
```

The downloaded data is cached locally. If you try to get dataset again or any other data set based on this one, it will check which files are already present and then download only the remaining ones.

SDK

The python interface has one major difference.

Getting the dataset does not mean it will be downloaded. You only capture the meta data.



This means we don't have to download the entire dataset to make changes or add files to it.

```
from clearml import Dataset  
  
ds = Dataset.get(dataset_name="animal-data")  
  
# The dataset is not download but it has all the meta data  
print(ds.list_files()[:5])
```

If you do wish to download the local copy of the dataset, it has to be done explicitly.

```
# Get read-only local copy of the data, now we will download  
local_path_read_only = ds.get_local_copy()  
print(local_path_read_only)
```

If you want to modify and make new versions

```
# Get a mutable copy of dataset to the desired folder
local_path Mutable = ds.getMutableLocalCopy(
    target_folder="/clearml-data/tmp/my_dataset"
)
```

Creating new version

We have downloaded the dataset original version.

Now we removed one folder (one category) and added a new category to create a new version of the data.

```
from clearml import Dataset

ds = Dataset.create(
    dataset_name="animal-dataset-with-mucca",
    dataset_project="clearml-data",
    parent_datasets=["224508a1e13b4d96a1a45209a3f0c049"],
)
```

We pass the id of previous original dataset version as the `parent_dataset`.

Next we modify the `metadata` object.

```
# Now we can make changes to the dataset without downloading it
ds.remove_files("./gallina")
ds.add_files("./tmp/modified/mucca")
```

It will sync everything when it is finalized.

If we already have a local copy that shows how we want the dataset to look like, then in that case it is easier to use the `sync` method.

```
# If we have a local copy that represent show we want dataset to look like
# in that case it is easier to just use the sync method
ds.sync_folder(
    local_path="./tmp/modified/"
)
```

This will compare metadata with the content of the supplied `local_path`.

Now when we call `finalize` and `upload`, it will only upload the files that are changed.

```
ds.finalize(auto_upload=True)
```

The CLI interface does not have the luxury of having metadata. It can only work with the sync command. However, it can bunch this whole process into a single command.

```
clearml-data sync --project clearml-data --id animal-dataset-with-mucca --parents 224508a1e13b4d96a1a45209a3f0c049 --folder /tmp/modified/
```

Now we can check the UI again.

Links

Documentation

https://clear.ml/docs/latest/docs/clearml_data/

Tutorial

https://www.youtube.com/watch?v=S2pz9jn26ul&list=PLMdIIcuMqSTnoC45ME5_JnsJX0zWqDdlO&index=9

Organizing

Before anything else, organizing code in the following format:

```
.
├── config
│   ├── args.json
│   └── config.py
└── logs
    ├── main.py
    └── model.py
    └── utils.py
```

We have a `main.py` file and a model definition file `model.py`. A folder for `logs` and some `config` files.

Loading args

```
args_path = config.CONFIG_DIR / "args.json"
args = Namespace(**utils.load_dict(filepath=args_path))
```

This way we have everything separated and it is easy to make changes.

Remember, since we are using external file instead of CLI arguments, we have to to `task.connect(args)` to record hyper-parameters.

Logging

Logging is the process of tracking and recording key events that occur in our applications for the purpose of inspection, debugging, etc.

```
import logging
import sys

# Create super basic logger
logging.basicConfig(stream=sys.stdout, level=logging.DEBUG)

# Logging levels (from lowest to highest priority)
logging.debug("Used for debugging your code.")
logging.info("Informative messages from your code.")
logging.warning("Everything works but there is something to be aware of.")
logging.error("There's been a mistake with the process.")
logging.critical("There is something terribly wrong and process may terminate.")
```

```
DEBUG:root:Used for debugging your code.
INFO:root:Informative messages from your code.
WARNING:root:Everything works but there is something to be aware of.
ERROR:root:There's been a mistake with the process.
CRITICAL:root:There is something terribly wrong and process may terminate.
```

Use use [RichHandler](#) for our `console` handler to get pretty formatting for the log messages.

```
pip install rich
```

```
import logging
import logging.config
from rich.logging import RichHandler
```

For my case, I just need `logger.info()` for now.

```
from config.config import logger
...
logger.info("Initializaing clearML task...")
task = Task.init(
    project_name="experimentation/logging",
    task_name=f"logging-example-{datetime.now()}"
)
```

How it looks like

```

~> ~ /pr/p/c/practice/pipelines > main !3
> python3 main.py
[02/22/23 15:13:33] INFO      Created a temporary directory at instantiator.py:21
                           /tmp/tmpgb1hm80c
[02/22/23 15:13:33] INFO      Writing instantiator.py:76
                           /tmp/tmpgb1hm80c/_remote_module_no
                           n_scriptable.py
ClearML Task: created new task id=1daf2eca76054348a405a063cb5b00bc
ClearML results page: https://app.clear.ml/projects/782628c316a547089a0391c07f186f
2b/experiments/1daf2eca76054348a405a063cb5b00bc/output/log
ClearML pipeline page: https://app.clear.ml/pipelines/782628c316a547089a0391c07f18
6f2b/experiments/1daf2eca76054348a405a063cb5b00bc
[02/22/23 15:13:50] INFO      No repository found, storing script scriptinfo.py:909
                           code instead
[02/22/23 15:13:53] INFO      No repository found, storing script scriptinfo.py:909
                           code instead
[02/22/23 15:13:56] INFO      No repository found, storing script scriptinfo.py:909
                           code instead
[02/22/23 15:14:08] INFO      Reading arguments... main.py:106
                           INFO      Preparing data... main.py:119
Launching step [prepare_data]
[02/22/23 15:15:28] INFO      Creating subsets... main.py:122
Launching step [create_subset]
[02/22/23 15:16:29] INFO      Creating model... main.py:130
                           INFO      Starting training... main.py:134
                           INFO      GPU available: True (cuda), used: rank_zero.py:45
                           True
                           INFO      TPU available: False, using: 0 TPU rank_zero.py:45
                           cores
                           INFO      IPU available: False, using: 0 IPUs rank_zero.py:45
                           INFO      HPU available: False, using: 0 HPUs rank_zero.py:45
Launching step [train]

```

Links

Logging for ML Systems

Logging for ML Systems - Made With ML

Keep records of the important events in our application.

<https://madewithml.com/courses/mlops/logging/>

ClearML Pipelines

Pipelines are useful to automate and orchestrate multiple tasks. Each task is a separate `step`.

Mainly useful when each step creates an output that is an input to another set.

- Pipelines are highly scalable.
- Really useful for ETL pipelines.

In clearML we can create pipelines from `Tasks` or `Code`. In former we can create separate task for each step. In latter we can create pipelines from codebase using decorators.

I want to create a very simple pipeline

```
Load data -> process data -> feed into model
```

I am doing three things.

- Load and transform data
- Create subset (just to add an extra step)
- Feed into model

Issue

Facing an issue

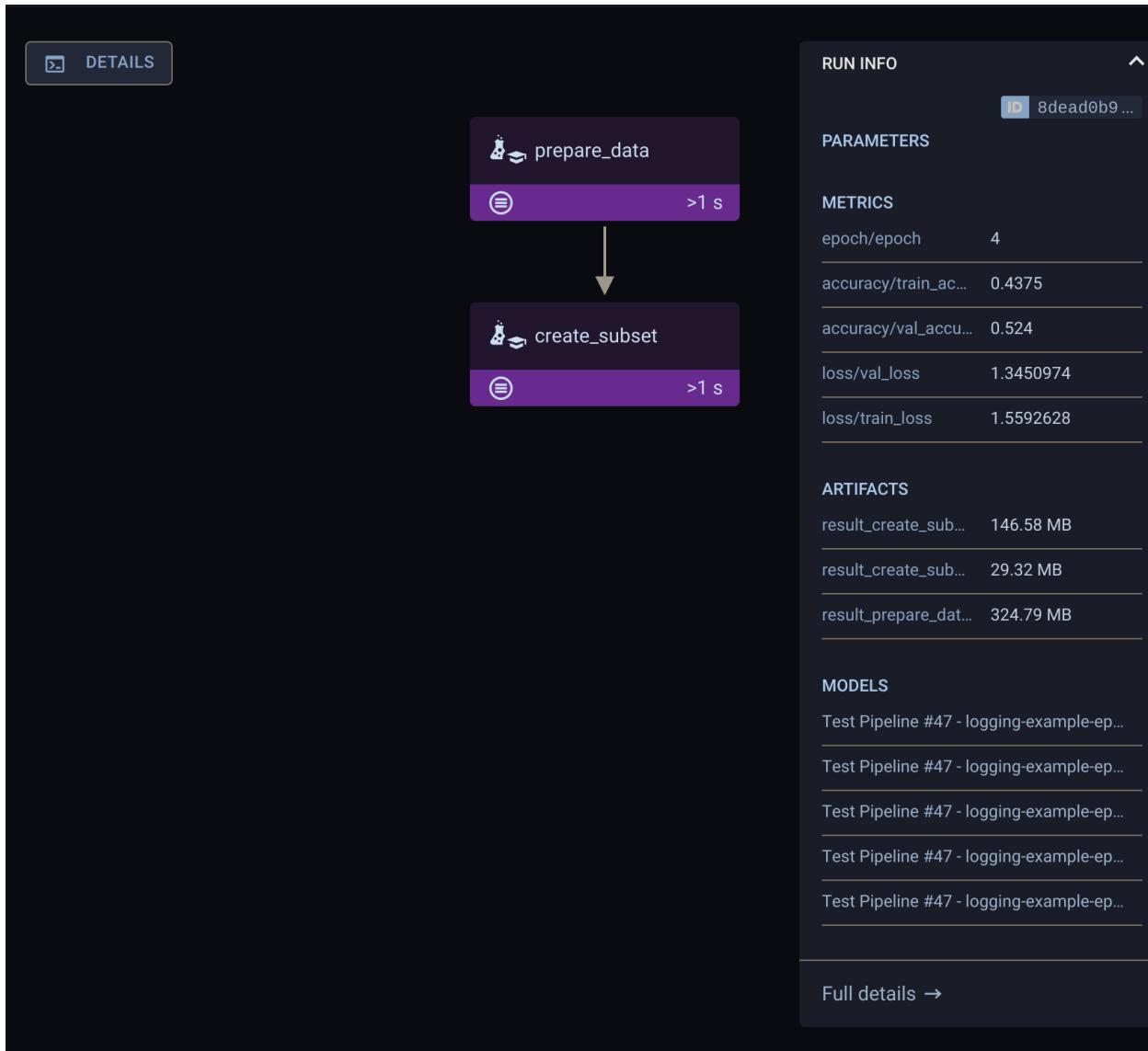
```
raise TypeError(f"`model` must be a `LightningModule`, got `{type(model).__qualname__}`")
TypeError: `model` must be a `LightningModule`, got `PosixPath`
```

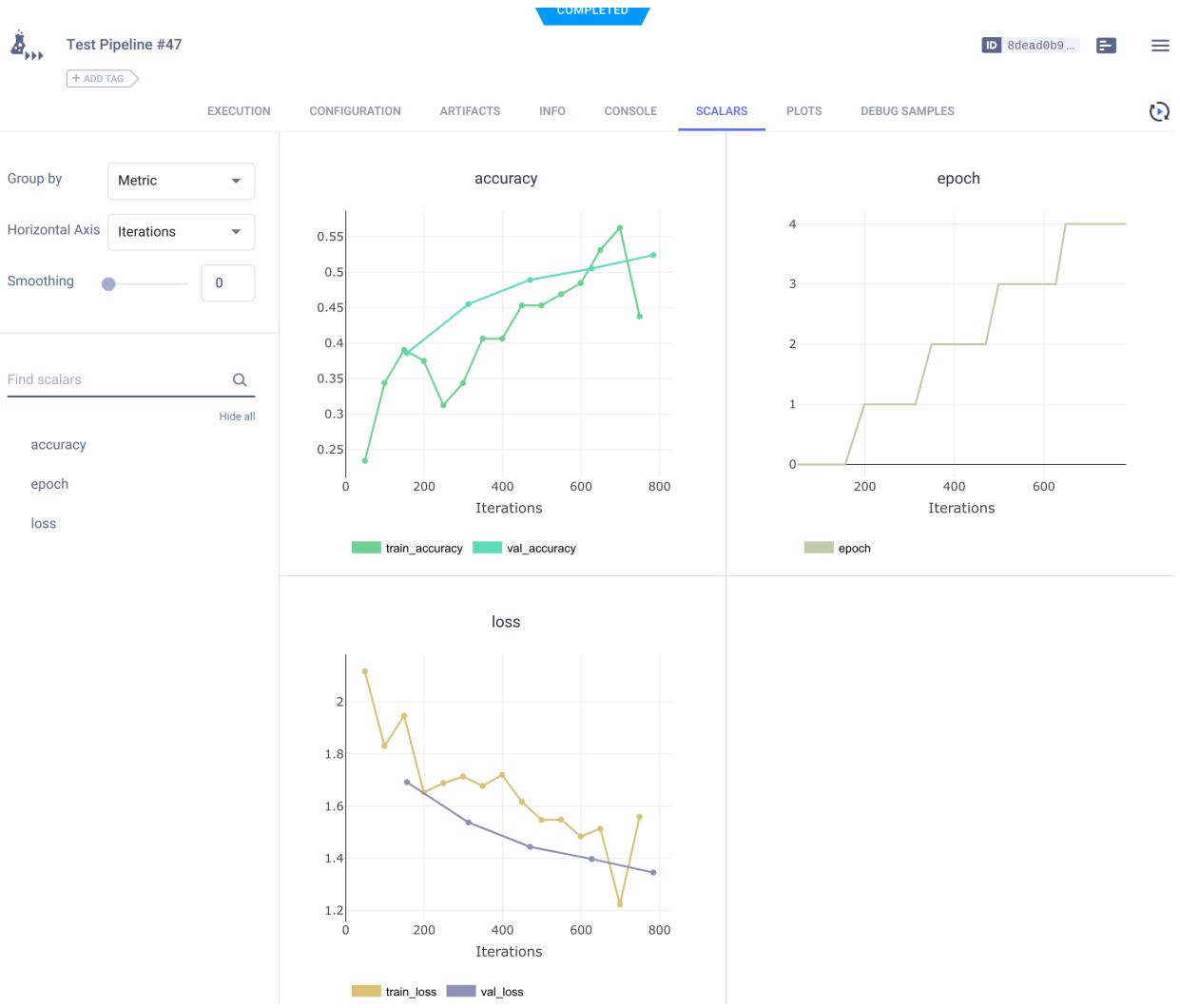
Tried printing the `classifier`

```
/home/pytholic/.clearml/cache/storage_manager/global/caa12decdd94851b2449688c2e69c588.result_train_classifier.pkl
```

It got converted to a path because of the wrapper.

Instead of creating separate train function, I ended up training it directly in the `main()` function.





You also get all the artifacts for each step.

OUTPUT MODELS

- logging-example-epoch=00-val_loss=0.00
- logging-example-epoch=01-val_loss=0.00
- logging-example-epoch=02-val_loss=0.00
- logging-example-epoch=03-val_loss=0.00
- logging-example-epoch=04-val_loss=0.00

OTHER

- result_create_subset_trainset
- result_create_subset_validset
- result_prepare_data_data_dir**

FILE PATH	https://files.clear.ml/Pipeline%20Examples/pipelines/Test%20Pipeline/Test%20Pipeline%20%252347.8dead0b9e90942a4b3987c4949e9e8d8/artifacts/result_prepare_data_data_dir/data.zip 
FILE SIZE	324.79 MB
HASH	e83115d13f98c20be31b637f73e2c5368f0fcdf1820dc63422aba930dfc1695

PREVIEW

```
Archive content /home/pytholic/projects/personal/clearML/practice/data/*:
cifar-10-batches-py/batches.meta - 158 bytes
cifar-10-batches-py/data_batch_1 - 31.04 MB
cifar-10-batches-py/data_batch_2 - 31.04 MB
cifar-10-batches-py/data_batch_3 - 31.04 MB
cifar-10-batches-py/data_batch_4 - 31.04 MB
cifar-10-batches-py/data_batch_5 - 31.04 MB
cifar-10-batches-py/readme.html - 88 bytes
cifar-10-batches-py/test_batch - 31.04 MB
cifar-10-python.tar.gz - 170.5 MB
```

Tidbits

- If you are running concurrent jobs, your console might not look pretty
 - However, you can use the Experiment UI in Clear ML app to view `console` of each experiment separately

Links

Examples

[clearml/examples at master · allegroai/clearml](https://github.com/allegroai/clearml)

You can't perform that action at this time. You signed in with another tab or window. You signed out in another tab or window. Reload to refresh your session. Reload to refresh your session.

 <https://github.com/allegroai/clearml/tree/master/examples>



Guides

[Examples | ClearML](https://clear.ml/docs/latest/docs/guides/)

To help learn and use ClearML, we provide example scripts that demonstrate how to use ClearML's various features.

 <https://clear.ml/docs/latest/docs/guides/>

Documentation

What is ClearML? | ClearML

ClearML is an open source platform that automates and simplifies developing and managing machine learning solutions

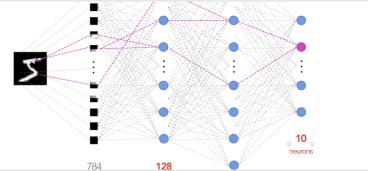
<https://clear.ml/docs/latest/docs>

Pytorch vs. Pytorch-Lightning

From PyTorch to PyTorch Lightning—A gentle introduction

This post walks through a side-by-side comparison of MNIST implemented using both PyTorch and PyTorch Lightning.

<https://towardsdatascience.com/from-pytorch-to-pytorch-lightning-a-gentle-introduction-b371b7caaf09>



ClearML SDK

Task | ClearML

```
class Task()  
    pass
```

<https://clear.ml/docs/latest/docs/references/sdk/task>

Queries

What is the difference in `task.log` and `Logger`?

The `task.log` method is a method in the `Task` class in the ClearML library, and it is used to log scalar metric values during the training of a machine learning model. The `Logger` class, on the other hand, is a more general purpose logging tool that can be used to log metric values, image, audio and other type of data in an experiment, not limited to a single task.

In summary, you can use `task.log` to log scalar metrics during training of a model, while `Logger` can be used to log a wider range of data, not just scalar metrics. Additionally, `Logger` is typically used in the context of an experiment, while `task.log` is typically used in the context of a single task.