

# cnn\_v10

October 15, 2022

```
[ ]: %env LD_LIBRARY_PATH=$LD_LIBRARY_PATH:$CONDA_PREFIX/lib/  
%env TF_GPU_ALLOCATOR=cuda_malloc_async
```

```
env: LD_LIBRARY_PATH=$LD_LIBRARY_PATH:$CONDA_PREFIX/lib/  
env: TF_GPU_ALLOCATOR=cuda_malloc_async
```

```
[ ]: import os  
print(os.environ["LD_LIBRARY_PATH"])
```

```
:/home/nkspartan/miniconda3/envs/tf-gpu/lib/:/home/nkspartan/miniconda3/envs/tf-gpu/lib/
```

```
[ ]: import tensorflow as tf  
import numpy as np  
import pandas as pd  
import os  
import keras  
import matplotlib.pyplot as plt  
  
from keras import Sequential, models, Input  
from keras.layers import Dense, Flatten, Conv2D, MaxPooling2D, Dropout, LeakyReLU, AveragePooling2D, GlobalAveragePooling2D, BatchNormalization  
from keras.optimizers import SGD, Adam
```

```
2022-10-15 20:05:43.700610: I tensorflow/core/platform/cpu_feature_guard.cc:193] This TensorFlow binary is optimized with oneAPI Deep Neural Network Library (oneDNN) to use the following CPU instructions in performance-critical operations: AVX2 FMA
```

To enable them in other operations, rebuild TensorFlow with the appropriate compiler flags.

```
2022-10-15 20:05:44.195466: E tensorflow/stream_executor/cuda/cuda_blas.cc:2981] Unable to register cuBLAS factory: Attempting to register factory for plugin cuBLAS when one has already been registered
```

```
2022-10-15 20:05:45.085080: W tensorflow/stream_executor/platform/default/dso_loader.cc:64] Could not load dynamic library 'libnvinfer.so.7'; dlopen: libnvinfer.so.7: cannot open shared object file: No such file or directory; LD_LIBRARY_PATH: /home/nkspartan/miniconda3/envs/tf-gpu/lib/:/home/nkspartan/miniconda3/envs/tf-
```

```

gpu/lib/
2022-10-15 20:05:45.085286: W
tensorflow/stream_executor/platform/default/dso_loader.cc:64] Could not load
dynamic library 'libnvinfer_plugin.so.7'; dLError: libnvinfer_plugin.so.7:
cannot open shared object file: No such file or directory; LD_LIBRARY_PATH:
:/home/nkspartan/miniconda3/envs/tf-gpu/lib/;/home/nkspartan/miniconda3/envs/tf-
gpu/lib/
2022-10-15 20:05:45.085292: W
tensorflow/compiler/tf2tensorrt/utils/py_utils.cc:38] TF-TRT Warning: Cannot
dlopen some TensorRT libraries. If you would like to use Nvidia GPU with
TensorRT, please make sure the missing libraries mentioned above are installed
properly.

```

```

[ ]: from tensorflow.python.client import device_lib

#print(device_lib.list_local_devices())
print('Default GPU Device: {}'.format(tf.test.gpu_device_name()))

```

Default GPU Device: /device:GPU:0

```

2022-10-15 20:05:46.641759: I tensorflow/core/platform/cpu_feature_guard.cc:193]
This TensorFlow binary is optimized with oneAPI Deep Neural Network Library
(oneDNN) to use the following CPU instructions in performance-critical
operations:  AVX2 FMA
To enable them in other operations, rebuild TensorFlow with the appropriate
compiler flags.
2022-10-15 20:05:46.672297: I
tensorflow/stream_executor/cuda/cuda_gpu_executor.cc:980] successful NUMA node
read from SysFS had negative value (-1), but there must be at least one NUMA
node, so returning NUMA node zero
2022-10-15 20:05:46.716563: I
tensorflow/stream_executor/cuda/cuda_gpu_executor.cc:980] successful NUMA node
read from SysFS had negative value (-1), but there must be at least one NUMA
node, so returning NUMA node zero
2022-10-15 20:05:46.716747: I
tensorflow/stream_executor/cuda/cuda_gpu_executor.cc:980] successful NUMA node
read from SysFS had negative value (-1), but there must be at least one NUMA
node, so returning NUMA node zero
2022-10-15 20:05:47.516134: I
tensorflow/stream_executor/cuda/cuda_gpu_executor.cc:980] successful NUMA node
read from SysFS had negative value (-1), but there must be at least one NUMA
node, so returning NUMA node zero
2022-10-15 20:05:47.516764: I
tensorflow/stream_executor/cuda/cuda_gpu_executor.cc:980] successful NUMA node
read from SysFS had negative value (-1), but there must be at least one NUMA
node, so returning NUMA node zero
2022-10-15 20:05:47.516918: I
tensorflow/stream_executor/cuda/cuda_gpu_executor.cc:980] successful NUMA node

```

```

read from SysFS had negative value (-1), but there must be at least one NUMA
node, so returning NUMA node zero
2022-10-15 20:05:47.517053: I
tensorflow/core/common_runtime/gpu/gpu_device.cc:1616] Created device
/device:GPU:0 with 4173 MB memory: -> device: 0, name: NVIDIA GeForce RTX 2060,
pci bus id: 0000:08:00.0, compute capability: 7.5

```

## 0.1 Read the csv dataset to get the values for stage and discharge of the images

```

[ ]: df = pd.read_csv("../dataset/2012_2019_PlatteRiverWeir_features_merged_all.
    ↪ csv")
df.head()

```

```

[ ]:      Unnamed: 0      SensorTime      CaptureTime  \
0          0  2012-06-09 13:15:00  2012-06-09T13:09:07
1          1  2012-06-09 13:15:00  2012-06-09T13:10:29
2          2  2012-06-09 13:45:00  2012-06-09T13:44:01
3          3  2012-06-09 14:45:00  2012-06-09T14:44:30
4          4  2012-06-09 15:45:00  2012-06-09T15:44:59

```

```

      Filename Agency  SiteNumber TimeZone  Stage  \
0  StateLineWeir_20120609_Farrell_001.jpg  USGS    6674500    MDT    2.99
1  StateLineWeir_20120609_Farrell_002.jpg  USGS    6674500    MDT    2.99
2  StateLineWeir_20120609_Farrell_003.jpg  USGS    6674500    MDT    2.96
3  StateLineWeir_20120609_Farrell_004.jpg  USGS    6674500    MDT    2.94
4  StateLineWeir_20120609_Farrell_005.jpg  USGS    6674500    MDT    2.94

```

```

      Discharge      CalcTimestamp  ...  WeirPt2X  WeirPt2Y  WwRawLineMin  \
0      916.0  2020-03-11T16:58:28  ...      -1      -1           0.0
1      916.0  2020-03-11T16:58:33  ...      -1      -1           0.0
2      873.0  2020-03-11T16:58:40  ...      -1      -1           0.0
3      846.0  2020-03-11T16:58:47  ...      -1      -1           0.0
4      846.0  2020-03-11T16:58:55  ...      -1      -1           0.0

```

```

      WwRawLineMax  WwRawLineMean  WwRawLineSigma  WwCurveLineMin  \
0           0.0           0.0           0.0           0.0
1           0.0           0.0           0.0           0.0
2           0.0           0.0           0.0           0.0
3           0.0           0.0           0.0           0.0
4           0.0           0.0           0.0           0.0

```

```

      WwCurveLineMax  WwCurveLineMean  WwCurveLineSigma
0           0.0           0.0           0.0
1           0.0           0.0           0.0
2           0.0           0.0           0.0
3           0.0           0.0           0.0
4           0.0           0.0           0.0

```

[5 rows x 60 columns]

```
[ ]: df = df[["Filename", "Stage", "Discharge"]]
```

### 0.1.1 Scale the data

```
[ ]: from sklearn.preprocessing import StandardScaler
      from joblib import load

      #scaler = StandardScaler()
      scaler = load('std_scaler.joblib')
```

```
[ ]: df[["Stage", "Discharge"]] = scaler.fit_transform(df[["Stage", "Discharge"]])
      df
```

```
[ ]:
      Filename      Stage  Discharge
0    StateLineWeir_20120609_Farrell_001.jpg  0.138117 -0.046094
1    StateLineWeir_20120609_Farrell_002.jpg  0.138117 -0.046094
2    StateLineWeir_20120609_Farrell_003.jpg  0.100875 -0.082160
3    StateLineWeir_20120609_Farrell_004.jpg  0.076046 -0.104807
4    StateLineWeir_20120609_Farrell_005.jpg  0.076046 -0.104807
...
42054 StateLineWeir_20191011_Farrell_409.jpg -0.420526 -0.450369
42055 StateLineWeir_20191011_Farrell_410.jpg -0.420526 -0.450369
42056 StateLineWeir_20191011_Farrell_411.jpg -0.420526 -0.450369
42057 StateLineWeir_20191011_Farrell_412.jpg -0.420526 -0.450369
42058 StateLineWeir_20191011_Farrell_413.jpg -0.420526 -0.450369
```

[42059 rows x 3 columns]

```
[ ]: from joblib import dump
      #dump(scaler, 'std_scaler.joblib')
```

## 0.2 Create the dataset pipeline

```
[ ]: #IMG_SIZE = 224
      IMG_SIZE = 512
      BATCH_SIZE = 32
```

```
[ ]: from glob import glob

      def make_dataset(path, batch_size, df, seed=None):
          np.random.seed(seed)

          image_augmentation = Sequential([
```

```

    tf.keras.layers.RandomBrightness([-0.2,0.4], seed=seed),
    tf.keras.layers.RandomContrast(0.4, seed=seed),
    tf.keras.layers.RandomZoom(0.3, seed=seed),
    tf.keras.layers.RandomFlip('horizontal', seed=seed),
    tf.keras.layers.RandomTranslation(height_factor=0.1, width_factor=0.1,
↪seed=seed),
    tf.keras.layers.RandomRotation(0.1, seed=seed)
])

def random_image_augmentation(image, probability=0.5):
    if np.random.random() < probability:
        return image_augmentation(image)
    return image

def parse_image(filename):
    image = tf.io.read_file(filename)
    image = tf.image.decode_jpeg(image, channels=3)
    #image = tf.image.resize(image, [IMG_SIZE, IMG_SIZE])

    # image augmentation
    image = random_image_augmentation(image, 0.7)

    image = tf.cast(image, tf.float32)
    image /= 255
    return image

def configure_for_performance(ds):
    ds = ds.shuffle(buffer_size=100)
    ds = ds.batch(batch_size)
    ds = ds.repeat()
    ds = ds.prefetch(buffer_size=tf.data.experimental.AUTOTUNE)
    return ds

filenames = glob(path + '/*')

# make train, val and test splits of the dataset (70%, 10%, 20% split)
split1 = int(0.7 * len(filenames))
split2 = int(0.8 * len(filenames))

np.random.shuffle(filenames)
train_files = filenames[:split1] # up to split 1 (ex 70%)
val_files = filenames[split1:split2] # from ex. 70% to 80%
test_files = filenames[split2:] # from ex. 80% until the end

# create stage values
stage_train_values = [df[df.FileName == file.split('/')[0]].Stage.values for
↪file in train_files]

```

```

stage_val_values = [df[df.Filename == file.split('/')[0]].Stage.values for
→file in val_files]
stage_test_values = [df[df.Filename == file.split('/')[0]].Stage.values for
→file in test_files]

# create discharge values
discharge_train_values = [df[df.Filename == file.split(
    '/')[-1]].Discharge.values for file in train_files]
discharge_val_values = [df[df.Filename == file.split(
    '/')[-1]].Discharge.values for file in val_files]
discharge_test_values = [df[df.Filename == file.split(
    '/')[-1]].Discharge.values for file in test_files]

# join stage and discharge values
stage_discharge_train_values = [[np.squeeze(s), np.squeeze(d)] for s, d in
→zip(stage_train_values, discharge_train_values)]
stage_discharge_val_values = [[np.squeeze(s), np.squeeze(d)] for s, d in
→zip(stage_val_values, discharge_val_values)]
stage_discharge_test_values = [[np.squeeze(s), np.squeeze(
    d)] for s, d in zip(stage_test_values, discharge_test_values)]

# create images dataset (train, val, test)
filenames_train_ds = tf.data.Dataset.from_tensor_slices(train_files)
filenames_val_ds = tf.data.Dataset.from_tensor_slices(val_files)
filenames_test_ds = tf.data.Dataset.from_tensor_slices(test_files)

images_train_ds = filenames_train_ds.map(parse_image, num_parallel_calls=8)
images_val_ds = filenames_val_ds.map(parse_image, num_parallel_calls=8)
images_test_ds = filenames_test_ds.map(parse_image, num_parallel_calls=8)

# create stage and discharge dataset (train, val, test)
stage_discharge_train_ds = tf.data.Dataset.
→from_tensor_slices(stage_discharge_train_values)
stage_discharge_val_ds = tf.data.Dataset.
→from_tensor_slices(stage_discharge_val_values)
stage_discharge_test_ds = tf.data.Dataset.from_tensor_slices(
    stage_discharge_test_values)

# create tensorflow dataset of images and values (train, val, test)
train_ds = tf.data.Dataset.zip((images_train_ds, stage_discharge_train_ds))
train_ds = configure_for_performance(train_ds)
val_ds = tf.data.Dataset.zip((images_val_ds, stage_discharge_val_ds))
val_ds = configure_for_performance(val_ds)
test_ds = tf.data.Dataset.zip((images_test_ds, stage_discharge_test_ds))
test_ds = configure_for_performance(test_ds)

```

```

    return train_ds, len(train_files), val_ds, len(val_files), test_ds,
    ↪len(test_files)

```

```

[ ]: path = "../../dataset/images"

train_ds, train_size, val_ds, val_size, test_ds, test_size = make_dataset(path,
    ↪BATCH_SIZE, df, 10)

```

```

[ ]: input_shape = 0
    output_shape = 0

    for image, stage_discharge in train_ds.take(1):
        print(image.numpy().shape)
        print(stage_discharge.numpy().shape)

        input_shape = image.numpy().shape[1:]
        output_shape = stage_discharge.numpy().shape[1:]

```

```

(32, 512, 512, 3)
(32, 2)

```

```

[ ]: print(input_shape)
    print(output_shape)

```

```

(512, 512, 3)
(2,)

```

### 0.3 Check images

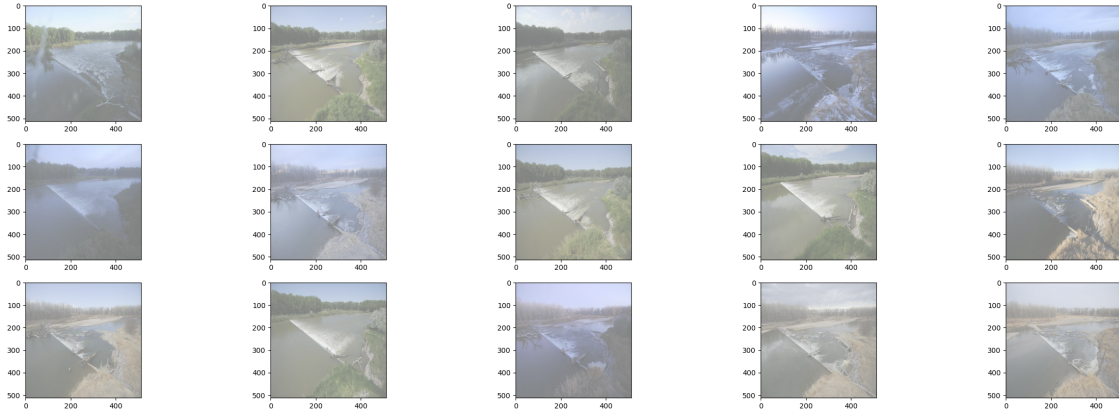
```

[ ]: fig, ax = plt.subplots(nrows=3, ncols=5, figsize=(30, 10))

    for image, stage_discharge in test_ds.take(1):
        images = image[:15]
        for img, ax in zip(images, ax.flatten()):
            img = img.numpy()
            img = img / 2 + 0.5      # unnormalize
            ax.imshow(img)

plt.show()

```



## 0.4 Create model

```
[ ]: def create_model(input_shape, output_shape, transfer_learning=False):
    model = Sequential()

    if (transfer_learning == True):
        base_model = tf.keras.applications.MobileNetV2(include_top=False,
                                                         weights='imagenet',
                                                         input_shape=input_shape)

        base_model.trainable = False
        base_model._name = 'base_model_MobileNet'

        model.add(base_model)
        model.add(Dropout(0.6))
        model.add(GlobalAveragePooling2D())

        model.add(Dense(512, activation='tanh'))
        model.add(Dense(256, activation='tanh'))
        model.add(Dense(256, activation='tanh'))
        model.add(Dense(64, activation='tanh'))
    else:
        model.add(Input(shape=input_shape))

        model.add(Conv2D(64, kernel_size=(4, 4), strides=(2, 2),
        ↪padding='same', activation=LeakyReLU()))
        model.add(MaxPooling2D(pool_size=(4, 4)))

        model.add(Conv2D(64, kernel_size=(4, 4), strides=(2, 2),
        ↪activation=LeakyReLU(), padding='same'))
        model.add(MaxPooling2D(pool_size=(2, 2)))
```



```

        model.add(Conv2D(32, kernel_size=(3, 3), activation=LeakyReLU(0.2),
        ↳padding='same'))
        #model.add(AveragePooling2D(pool_size=(2, 2)))

        model.add(Conv2D(32, kernel_size=(3, 3), activation='relu'))

        model.add(Conv2D(32, kernel_size=(2, 2), activation='relu'))
        model.add(AveragePooling2D(pool_size=(2, 2)))

        model.add(Flatten())
        model.add(Dense(128, activation='tanh'))
        model.add(Dropout(0.3))
        model.add(Dense(64, activation='tanh'))
        model.add(Dense(32, activation='tanh'))
        model.add(Dense(32, activation='tanh'))

        model.add(Dense(output_shape, activation='linear')) # linear regression
        ↳output layer

    return model

```

```
[ ]: model = create_model(input_shape, output_shape[0], False)
```

```
[ ]: model.summary()
```

Model: "sequential\_6"

Layer (type)	Output Shape	Param #
conv2d_5 (Conv2D)	(None, 256, 256, 64)	3136
max_pooling2d_2 (MaxPooling 2D)	(None, 64, 64, 64)	0
conv2d_6 (Conv2D)	(None, 32, 32, 64)	65600
max_pooling2d_3 (MaxPooling 2D)	(None, 16, 16, 64)	0
conv2d_7 (Conv2D)	(None, 16, 16, 32)	18464
conv2d_8 (Conv2D)	(None, 14, 14, 32)	9248
conv2d_9 (Conv2D)	(None, 13, 13, 32)	4128
average_pooling2d_1 (Averag	(None, 6, 6, 32)	0

ePooling2D)

flatten_1 (Flatten)	(None, 1152)	0
dense_10 (Dense)	(None, 128)	147584
dropout_2 (Dropout)	(None, 128)	0
dense_11 (Dense)	(None, 64)	8256
dense_12 (Dense)	(None, 32)	2080
dense_13 (Dense)	(None, 32)	1056
dense_14 (Dense)	(None, 2)	66

```
=====
Total params: 259,618
Trainable params: 259,618
Non-trainable params: 0
-----
```

```
[ ]: def compile_model(loss_func, optimizer, metrics=["accuracy"]):
    model.compile(loss=loss_func, optimizer=optimizer, metrics=metrics)

[ ]: sgd = SGD(learning_rate=0.01, decay=1e-4, momentum=0.9, nesterov=True)
adam = Adam(learning_rate=1e-3, decay=1e-3 / 200)

compile_model('mse', adam, [
    'mse', tf.keras.metrics.RootMeanSquaredError(name='rmse'), 'mae',
    'mape'])

[ ]: def fit_model(training_values, validation_values=None, epochs=10, steps=32,
    val_steps=32, callbacks=[]):
    return model.fit(training_values, validation_data=validation_values,
    epochs=epochs, steps_per_epoch=steps, validation_steps=val_steps,
    callbacks=callbacks)

[ ]: import datetime

date_actual = datetime.datetime.now().strftime("%Y%m%d-%H%M%S")
log_dir = "logs/fit/" + date_actual
tensorboard_callback = tf.keras.callbacks.TensorBoard(log_dir=log_dir,
    histogram_freq=1)
```

```
checkpoint_callback = tf.keras.callbacks.  
    ↳ModelCheckpoint(filepath=f"model_weights/{date_actual}_cnn_best_weights.  
    ↳hdf5",  
                    monitor='val_mse',  
                    verbose=1,  
                    save_best_only=True)
```

```
[ ]: # batch_size = 0 because we already have batch size in tf dataset  
model_h = fit_model(train_ds, val_ds, epochs=20, steps=np.ceil(train_size /  
    ↳BATCH_SIZE), val_steps=np.ceil(val_size / BATCH_SIZE),  
    ↳callbacks=[tensorboard_callback, checkpoint_callback])
```

## 0.5 Evaluate model

```
[ ]: print(date_actual)
```

20221015-204855

```
[ ]: best_model = models.load_model(f'model_weights/{date_actual}_cnn_best_weights.  
    ↳hdf5')  
#best_model = models.load_model(f'best_models_weights/cnn_best_weights_v9.hdf5')
```

```
[ ]: def evaluate_model(model, test_values, steps):  
    score = model.evaluate(test_values, steps=steps)  
    return score
```

```
[ ]: test_loss, test_mse, test_rmse, test_mae, test_mape =  
    ↳evaluate_model(best_model, test_ds, steps=np.ceil(test_size / BATCH_SIZE))
```

263/263 [=====] - 17s 62ms/step - loss: 0.0030 - mse:  
0.0030 - rmse: 0.0547 - mae: 0.0386 - mape: 15.8871

```
[ ]: #predictions = best_model.predict(test_ds, steps=np.ceil(test_size /  
    ↳BATCH_SIZE))
```

```
[ ]: for image, stage_discharge in test_ds.take(1):  
    predictions = best_model.predict(x=image)  
  
    stage_discharge_test_values = stage_discharge[:2].numpy()  
    predictions_values = predictions[:2]  
  
    diff = predictions_values.flatten() - stage_discharge_test_values.  
    ↳flatten()  
    percentDiff = (diff / stage_discharge_test_values.flatten()) * 100  
    absPercentDiff = np.abs(percentDiff)  
    # compute the mean and standard deviation of the absolute percentage  
    # difference
```

```

mean = np.mean(absPercentDiff)
std = np.std(absPercentDiff)
# finally, show some statistics on our model
print(mean)
print(std)

stage_discharge_test_values = stage_discharge[:10]
predictions_values = predictions[:10]

for i in range(len(stage_discharge_test_values.numpy())):
    print(f"pred stage: {scaler.
→inverse_transform(predictions_values)[i][0]}, actual stage: {scaler.
→inverse_transform(stage_discharge_test_values)[i][0]}")
    print(f"pred discharge: {scaler.
→inverse_transform(predictions_values)[i][1]}, actual discharge: {scaler.
→inverse_transform(stage_discharge_test_values)[i][1]}")

```

```

1/1 [=====] - 0s 106ms/step
5.142239835017092
0.9115972227498049
pred stage: 1.9042773246765137, actual stage: 1.94
pred discharge: 30.319110870361328, actual discharge: 73.60000000000002
pred stage: 2.4251697063446045, actual stage: 2.45
pred discharge: 296.9564514160156, actual discharge: 336.0
pred stage: 3.692131757736206, actual stage: 3.72
pred discharge: 1960.296630859375, actual discharge: 2050.0
pred stage: 3.468950033187866, actual stage: 3.43
pred discharge: 1495.0885009765625, actual discharge: 1440.0
pred stage: 2.826493263244629, actual stage: 2.83
pred discharge: 753.9630737304688, actual discharge: 704.0
pred stage: 2.2473175525665283, actual stage: 2.25
pred discharge: 196.7310333251953, actual discharge: 214.0
pred stage: 2.5455737113952637, actual stage: 2.58
pred discharge: 385.1839904785156, actual discharge: 403.0
pred stage: 2.2392189502716064, actual stage: 2.27
pred discharge: 177.2609405517578, actual discharge: 206.0
pred stage: 2.4840328693389893, actual stage: 2.46
pred discharge: 334.6043395996094, actual discharge: 315.0
pred stage: 2.1623551845550537, actual stage: 2.17
pred discharge: 131.10499572753906, actual discharge: 169.0

```

### 0.5.1 Residual analysis

```

[ ]: y_predictions = np.empty(shape=(1, 2))
     y_real = np.empty(shape=(1, 2))

"""for image, stage_discharge in test_ds.take(100):

```

```

y_predictions = np.concatenate((y_predictions, best_model.predict(x=image)))
y_real = np.concatenate((y_real, stage_discharge.numpy()))"""

```

```

[ ]: 'for image, stage_discharge in test_ds.take(100):\n    y_predictions =
np.concatenate((y_predictions, best_model.predict(x=image)))\n    y_real =
np.concatenate((y_real, stage_discharge.numpy()))'

```

```

[ ]: residuals = y_real - y_predictions
residuals_std = residuals/residuals.std()

y_real_stage = np.array([i[0] for i in y_real])
residual_stage = np.array([i[0] for i in residuals])

y_real_discharge = np.array([i[1] for i in y_real])
residual_discharge = np.array([i[1] for i in residuals])

plt.scatter(y_real_stage, residual_stage / residual_stage.std(), label="stage_
↳residuals")
plt.scatter(y_real_discharge, residual_discharge / residual_discharge.std(),
↳label="discharge residuals")
plt.axhline(y=0.0, color='r', linestyle='-')
plt.xlabel("Fitted values")
plt.ylabel("Standarized residuals")

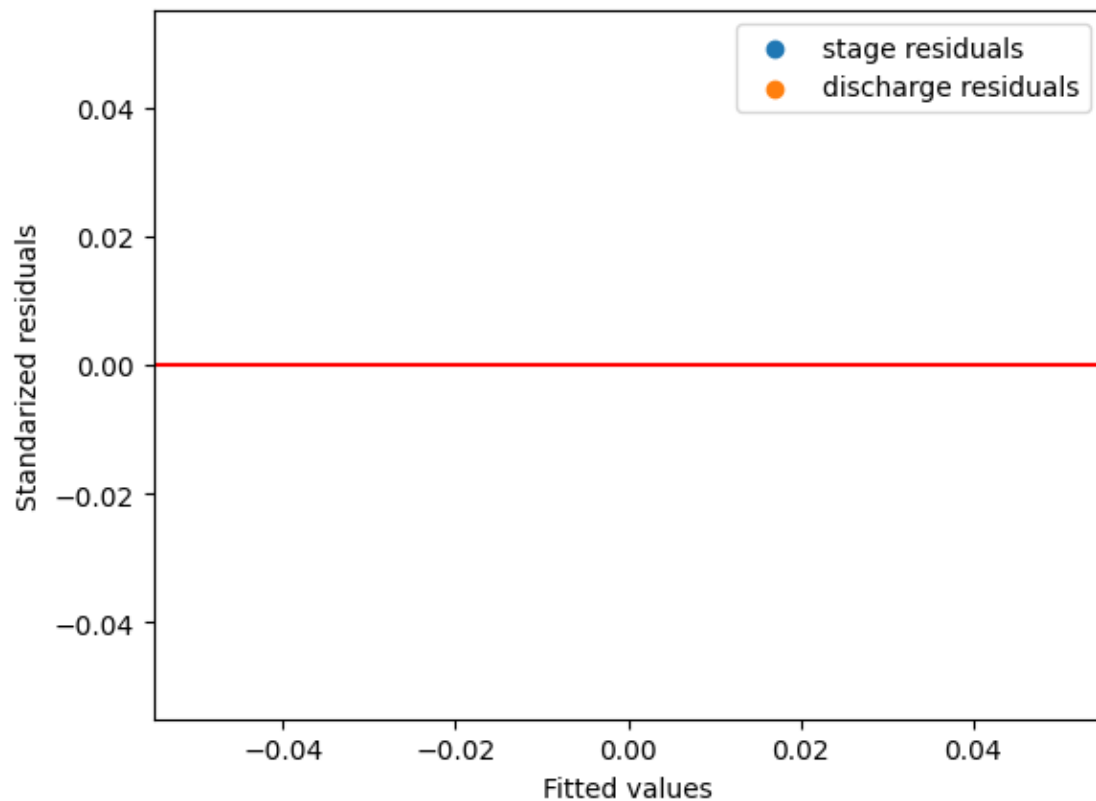
plt.legend()
plt.show()

```

```

/tmp/ipykernel_67186/3264406076.py:10: RuntimeWarning: divide by zero
encountered in divide
    plt.scatter(y_real_stage, residual_stage / residual_stage.std(), label="stage
residuals")
/tmp/ipykernel_67186/3264406076.py:11: RuntimeWarning: divide by zero
encountered in divide
    plt.scatter(y_real_discharge, residual_discharge / residual_discharge.std(),
label="discharge residuals")

```

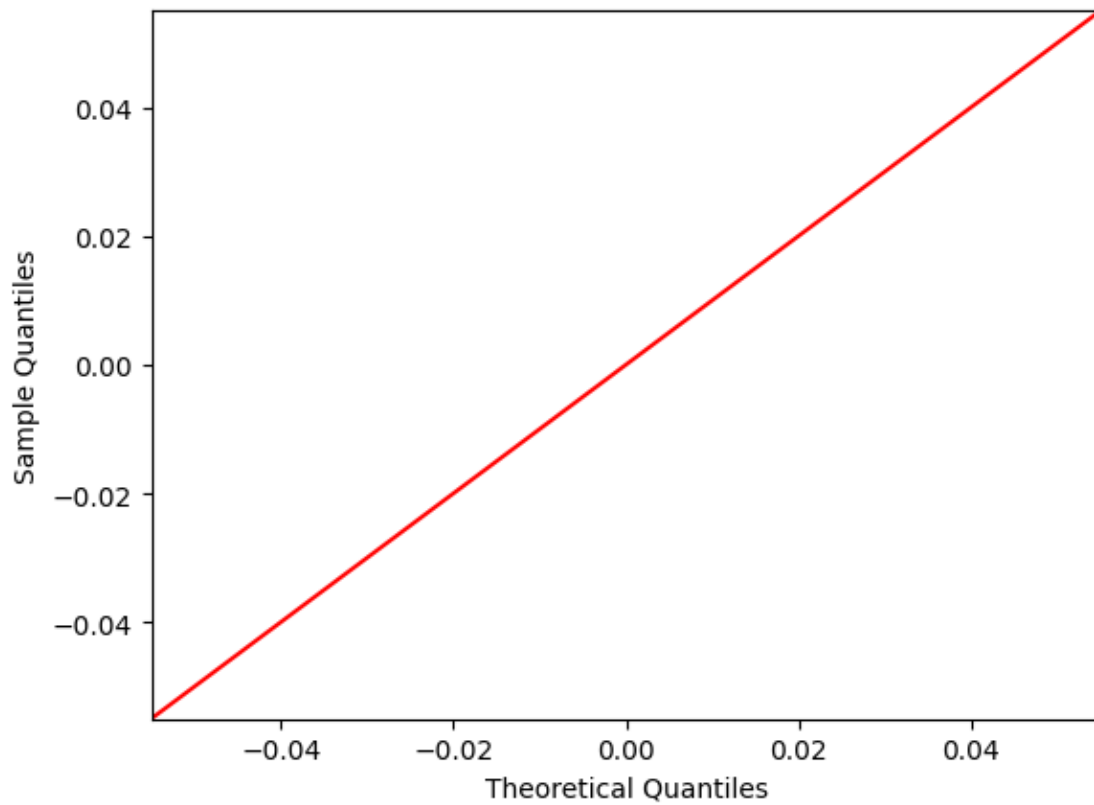


```
[ ]: import statsmodels.api as sm
from statsmodels.stats.diagnostic import normal_ad

figure = sm.qqplot(residual_stage / residual_stage.std(), line='45',
    ↪label='stage')
plt.show()
```

/tmp/ipykernel\_67186/3505247562.py:4: RuntimeWarning: divide by zero encountered in divide

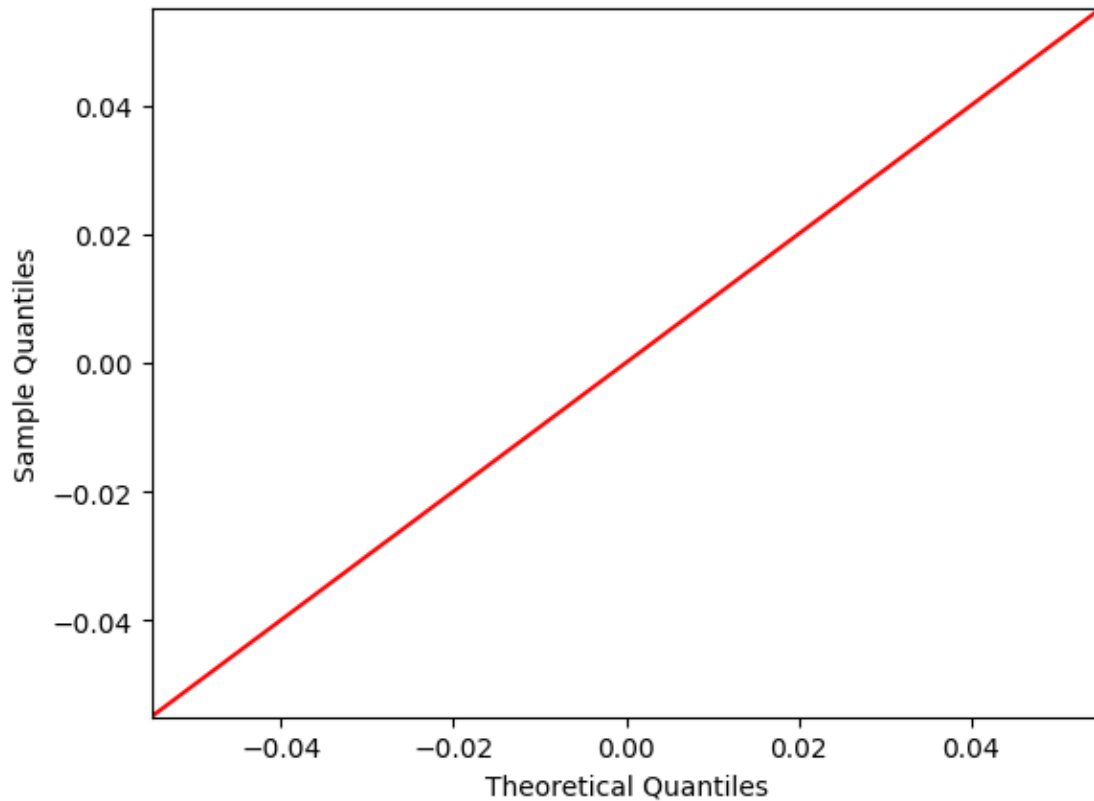
```
figure = sm.qqplot(residual_stage / residual_stage.std(), line='45',
label='stage')
```



```
[ ]: figure = sm.qqplot(residual_discharge / residual_discharge.std(), line='45',  
    ↪label='discharge')  
plt.show()
```

/tmp/ipykernel\_67186/1192211388.py:1: RuntimeWarning: divide by zero encountered  
in divide

```
    figure = sm.qqplot(residual_discharge / residual_discharge.std(), line='45',  
label='discharge')
```



```
[ ]: import seaborn as sns

#sns.histplot(residuals, kde=True, bins = 10)

[ ]: stat, pval = normal_ad(residual_stage)
print("p-value:", pval)

if pval<0.05:
    print("Hay evidencia de que los residuos no provienen de una distribución ↵
    ↵normal.")
else:
    print("No hay evidencia para rechazar la hipótesis de que los residuos ↵
    ↵vienen de una distribución normal.")
```

p-value: 0.0

Hay evidencia de que los residuos no provienen de una distribución normal.

/home/nkspartan/miniconda3/envs/tf-gpu/lib/python3.10/site-packages/numpy/core/\_methods.py:265: RuntimeWarning: Degrees of freedom <= 0 for slice

ret = \_var(a, axis=axis, dtype=dtype, out=out, ddof=ddof,  
/home/nkspartan/miniconda3/envs/tf-gpu/lib/python3.10/site-



```
packages/numpy/core/_methods.py:257: RuntimeWarning: invalid value encountered
in double_scalars
    ret = ret.dtype.type(ret / rcount)
```

```
[ ]: stat, pval = normal_ad(residual_discharge)
print("p-value:", pval)

if pval < 0.05:
    print("Hay evidencia de que los residuos no provienen de una distribución
    ↪normal.")
else:
    print("No hay evidencia para rechazar la hipótesis de que los residuos
    ↪vienen de una distribución normal.")
```

p-value: 0.0

Hay evidencia de que los residuos no provienen de una distribución normal.

## 0.6 Visualize layers

```
[ ]: layer_outputs = [layer.output for layer in best_model.layers[:12]]
# Extracts the outputs of the top 12 layers
activation_model = models.Model(inputs=best_model.input, outputs=layer_outputs)
    ↪# Creates a model that will return these outputs, given the model input
```

```
[ ]: activations = activation_model.predict(test_ds.take(1))
```

1/1 [=====] - 0s 174ms/step

```
[ ]: import matplotlib.pyplot as plt

layer_names = []
for layer in best_model.layers[:12]:
    layer_names.append(layer.name) # Names of the layers, so you can have them
    ↪as part of your plot

images_per_row = 16

for layer_name, layer_activation in zip(layer_names, activations): # Displays
    ↪the feature maps
    n_features = layer_activation.shape[-1] # Number of features in the feature
    ↪map
    size = layer_activation.shape[1] #The feature map has shape (1, size, size,
    ↪n_features).
    n_cols = n_features // images_per_row # Tiles the activation channels in
    ↪this matrix
    display_grid = np.zeros((size * n_cols, images_per_row * size))
```

```

print(layer_name)
if "flatten" in layer_name or "dense" in layer_name: break

for col in range(n_cols): # Tiles each filter into a big horizontal grid
    for row in range(images_per_row):
        channel_image = layer_activation[0,
                                         :, :,
                                         col * images_per_row + row]
        channel_image -= channel_image.mean() # Post-processes the feature
        ↪to make it visually palatable
        channel_image /= channel_image.std()
        channel_image *= 64
        channel_image += 128
        channel_image = np.clip(channel_image, 0, 255).astype('uint8')
        display_grid[col * size : (col + 1) * size, # Displays the grid
                     row * size : (row + 1) * size] = channel_image

scale = 1. / size
plt.figure(figsize=(scale * display_grid.shape[1],
                    scale * display_grid.shape[0]))
plt.title(layer_name)
plt.grid(False)
plt.imshow(display_grid, aspect='auto', cmap='viridis')

```

```

conv2d_5
max_pooling2d_2
conv2d_6
max_pooling2d_3
conv2d_7
conv2d_8
conv2d_9
average_pooling2d_1
flatten_1

```

/tmp/ipykernel\_67186/1031377702.py:24: RuntimeWarning: invalid value encountered in divide

```
channel_image /= channel_image.std()
```

