Homework 3

```
###################
# supercomputer.py #
#!/usr/bin/env python3
from collections import deque
import pickle
import os
import sys
import json
import subprocess
import itertools
import matplotlib.pyplot as plt
import numpy as np
from tensorflow.keras.callbacks import EarlyStopping
from models import dnn
from symbiotic_metrics import FractionOfVarianceAccountedFor
def main():
  """Spits out training jobs for each configuration"""
  # Create error file
  with open("error/err.txt", "w") as f:
    pass
  # Create option dictionary
  options = {
    "rotation": list(range(20)),
    "n_train_folds": [1, 2, 3, 5, 10, 18],
    "dropout": [0, .3, .6, .8]
    #"I2": [0, .001, .01, .1]
  }
  option combinations = create combinations(options)
  option combinations = create index log(options, option combinations)
```

```
# Start a job for each hyperparameter
  for option combo in option combinations:
    start training_job(**option_combo)
def create combinations(option dictionaries):
  Used to create a list of dictionaries containing all possible combinations
    of input dictionary arguments
  Found on tutorial website: https://riptutorial.com/python/example/10160/all-combinations-
of-dictionary-values
  keys = option_dictionaries.keys()
  values = (option dictionaries[key] for key in keys)
  combinations = [dict(zip(keys, combination)) for combination in itertools.product(*values)]
  for i in range(len(combinations)):
    combinations[i]["experiment num"] = i
  return combinations
def create index log(options, option combinations):
  """Write index to file that describes experiment hyperparameters"""
  fbase = "results/"
  if not os.path.exists(fbase):
    os.mkdir(fbase)
  batch num = 0
  while ( os.path.exists("{}batch_{{}}/".format(fbase, batch_num)) ):
    batch num += 1
  fbase = "{}batch {}/".format(fbase, batch num)
  os.mkdir(fbase)
  with open('{}index.txt'.format(fbase), 'w') as f:
    f.write("Number of experiments: {}\n".format(len(option combinations)))
    json.dump(options, f)
    f.write("\n")
    for i in range(len(option combinations)):
      option_combinations[i]["batch_num"] = batch_num
      json.dump(option combinations[i], f)
      f.write("\n")
```

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return option combinations
def start_training_job(**kwargs):
  Starts a job for the fed arguments. This takes the form of a subprocess,
  whether on a normal computer or supercomputer
  print("Starting job:\n\t{}".format(kwargs))
  # Decide which script to run
  if "-s" in sys.argv:
    script_to_run = ["sbatch", "supercomputer_job.sh", "-s"]
    script to run = ["./standard job.sh"]
  # Build script with hyperparameters
  full command = [
    *script to run,
    "-job"
  ]
  for key, value in kwargs.items():
    full_command.append("--{}={}".format(key, value))
  # Run chosen script with correct arguments
  process = subprocess.Popen(full command)
  # Wait if not parallel
  if "-p" not in sys.argv:
    process.wait()
def parse args():
  # Parse the hyperparameter arguments
  kwargs = {}
  for arg in sys.argv:
    if "--" in arg:
      arg = arg.replace("--", "")
      key, value = arg.split("=")
      kwargs[key] = value
  return kwargs
```

```
def train(**kwargs):
  print("PARAMETERS: {}".format(kwargs))
  # Unpack relevant kwargs
  rotation = int(kwargs["rotation"])
  n train folds = int(kwargs["n train folds"])
  experiment num = int(kwargs["experiment num"])
  batch num = int(kwargs["batch num"])
  if "dropout" in kwargs.keys():
    dropout = float(kwargs["dropout"])
  else:
    dropout = 0
  if "I2" in kwargs.keys():
    12 = float(kwargs["I2"])
  else:
    12 = 0
  # Rotate indices based on current rotation
  rotation indices = get rotation indices(n folds=20, rotation=rotation)
  # Get the training, validation, and test fold indices
  fold inds = get set indices(rotation indices=rotation indices, n train folds=n train folds)
  " Load data
  Key MI, Length 20, Shape (1193, 960)
  Key theta, Length 20, Shape (1193, 2)
  Key dtheta, Length 20, Shape (1193, 2)
  Key ddtheta, Length 20, Shape (1193, 2)
  Key torque, Length 20, Shape (1193, 2)
  Key time, Length 20, Shape (1193, 1)
  if "-s" in sys.argv:
    data path = "/home/fagg/ml datasets/bmi/bmi dataset.pkl"
  else:
    data path = "../homework 2/bmi dataset.pkl"
  with open(data path, "rb") as fp:
    hw2 dataset = pickle.load(fp)
  # Splits the data into its respective train, validation, and test sets / ins and outs
  processed data = process dataset(hw2 dataset, fold inds)
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# Build model
  model = dnn(
    input_size=(processed_data["train"]["ins"].shape[1],),
    hidden sizes=[300, 150, 100, 50, 10],
    output_size=processed_data["train"]["outs"].shape[1],
    hidden act="elu",
    output act="linear",
    dropout=dropout,
    12=12)
  # Compile model with fvaf metric
  fvaf = FractionOfVarianceAccountedFor(processed data["test"]["outs"].shape[1])
  model.compile(optimizer="adam", loss="mse", metrics=[fvaf], verbose=2)
  model.summary()
  # Callbacks
  es callback = EarlyStopping(
               monitor="val loss",
               patience=20,
               restore_best_weights=True,
               min delta=.0001)
  # Train model
  history = model.fit(
      x=processed data["train"]["ins"],
      y=processed data["train"]["outs"],
      validation_data = (processed_data["val"]["ins"], processed_data["val"]["outs"]),
      epochs=10000,
      batch_size=32,
      callbacks=[es callback]
      )
  # Log results
  log(model, processed data, kwargs)
  # Plot the torque and save figure
  if experiment_num == 0:
    plot shoulder orientation(model, processed data, kwargs)
def plot shoulder orientation(model, data, kwarg dict):
  """Plots the torque graph"""
  # Create results directory
```

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save path = "results/"
  if not os.path.exists(save_path):
    os.mkdir(save path)
  save path += "batch {}/".format(kwarg dict["batch num"])
  if not os.path.exists(save_path):
    os.mkdir(save path)
  # Create specific experiment directory
  save_path += "experiment_{}/".format(kwarg_dict["experiment_num"])
  if not os.path.exists(save_path):
    os.mkdir(save_path)
  true orientation = data["test"]["outs"][:, 0]
  predicted_orientation = model.predict(data["test"]["ins"])[:, 0]
  # Create and configure plot
  fig = plt.figure()
  ax = fig.add subplot(1, 1, 1)
  ax.plot(data["test"]["time"], true orientation, label="True Orientation")
  ax.plot(data["test"]["time"], predicted_orientation, label="Predicted Orientation")
  ax.legend()
  plt.ylabel("Orientation")
  plt.xlabel("Time")
  # Save plot
  fig.savefig(save path + f"orientation plot.png", dpi=fig.dpi)
def log(model, data, kwarg dict):
  """Log results to file"""
  print("Logging results")
  # Generate results
  results = {}
  results['predict train'] = model.predict(data["train"]["ins"])
  results['eval train'] = model.evaluate(data["train"]["ins"], data["train"]["outs"])
  results['predict_val'] = model.predict(data["val"]["ins"])
  results['eval val'] = model.evaluate(data["val"]["ins"], data["val"]["outs"])
  results['predict test'] = model.predict(data["test"]["ins"])
  results['eval test'] = model.evaluate(data["test"]["ins"], data["test"]["outs"])
  for key, value in kwarg dict.items():
    results[key] = value
```

```
# Create results directory
  fbase = "results/"
  if not os.path.exists(fbase):
    os.mkdir(fbase)
  fbase += "batch {}/".format(kwarg dict["batch num"])
  if not os.path.exists(fbase):
    os.mkdir(fbase)
  fbase += "experiment {}/".format(kwarg dict["experiment num"])
  if not os.path.exists(fbase):
    os.mkdir(fbase)
  # Save results
  with open("{}results dict.pkl".format(fbase), "wb") as fp:
    pickle.dump(results, fp)
    fp.close()
  # Create model directory
  if not os.path.exists("{}/model/".format(fbase)):
    os.mkdir("{}/model/".format(fbase))
  # Save model
  model.save("{}/model/".format(fbase))
def process dataset(dataset, fold inds):
  Process the dataset into the train, validation, and test folds;
  Also split into ins & out sets
  processed_data = {}
  for key in fold inds.keys():
    processed data[key] = split dataset(dataset, fold inds[key])
  return processed data
def split dataset(dataset, inds):
  # Placeholder for data splits
  processed data = {
    "ins": None,
    "outs": [],
```

```
"time": None
  }
  for key in dataset.keys():
    # Get folds for this key
    folds = [dataset[key][ind] for ind in inds]
    # Join the folds
    joined = np.concatenate((folds), axis=0)
    # See if the key is for the ins or outs of the dataset
    if key == "MI":
       processed data["ins"] = joined
    elif key == "time":
       processed_data["time"] = joined
    elif key == "theta":
       processed data["outs"] = np.expand dims(joined[:, 0], axis=1)
  return processed_data
def get_set_indices(rotation_indices, n_train_folds):
  """Get the fold indices for each set"""
  inds = \{\}
  inds["train"] = [rotation indices[i] for i in range(n train folds)]
  inds["val"] = [rotation indices[len(rotation indices)-2]]
  inds["test"] = [rotation indices[len(rotation indices)-1]]
  return inds
def get rotation indices(n folds, rotation=0):
  """Rotate folds to get the right indices"""
  fold list = list(range(n folds))
  fold list = deque(fold list)
  fold list.rotate(rotation)
  fold list = list(fold list)
  return fold list
if name == " main ":
  # If this is a subprocess, run the training program
  if "-job" in sys.argv:
```

```
try:
    train(**kwargs)

# If any exception occurs, write to error folder to differentiate between all the job outputs
except Exception as e:
    fbase = "error/"
    if not os.path.exists(fbase):
        os.mkdir(fbase)

with open("{}err.txt".format(fbase), "a") as f:
        err_str = "Error: {}\n".format(e)
        f.write(err_str)

else:
    main()
```

```
########
# local.py #
########
#!/usr/bin/env python3
import sys
import json
import pickle
import os
import subprocess
import matplotlib.pyplot as plt
import numpy as np
def main():
  # Use -s argument to scp results from supercomputer before continuing
  if "-s" in sys.argv:
    script to run = [
      "scp",
      "-r",
      "jwspaeth@schooner.oscer.ou.edu:/home/jwspaeth/workspaces/advanced-
ml/homework_3/results",
      "./"]
    process = subprocess.Popen([*script to run])
    process.wait()
  # Load dropout experiments and results
  dropout_options, dropout_experiments = load_index_log(batch_num=0)
  dropout results = [load result from experiment(experiment) for experiment in
dropout experiments]
  # Load I2 experiments and results
  12 options, I2 experiments = load_index_log(batch_num=1)
  12_results = [load_result_from_experiment(experiment) for experiment in l2_experiments]
  # Compute fvaf curves for each key value
  # (training folds x hyperparameter value), so average across rotations
  dropout avg fvafs = compute avg fvaf curves(dropout results,
                         dropout options["n train folds"],
                         dropout_options["dropout"],
                         "dropout",
                         dropout options["rotation"])
```

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12 avg fvafs = compute avg fvaf curves(12 results,
                        12 options["n train folds"],
                        12 options["I2"],
                        "l2",
                        12 options["rotation"])
  # Plot fvaf curves for each key value
  plot fvaf curves(dropout options["n train folds"],
           dropout options["dropout"],
           dropout avg fvafs["val"],
           "Validation Dropout")
  plot_fvaf_curves(I2_options["n_train_folds"],
          12 options["I2"],
          12 avg fvafs["val"],
           "Validation L2")
  # Get argmax for the best hyperparameter values
  dropout argmax fvafs = np.argmax(dropout avg fvafs["val"], axis=1)
  12 argmax_fvafs = np.argmax(I2_avg_fvafs["val"], axis=1)
  # Build arrays containing test values for best validation models
  test dropout = []
  test |2 = []
  for i in range(dropout_argmax_fvafs.shape[0]):
    test dropout.append(dropout avg fvafs["test"][i, dropout argmax fvafs[i]])
    test |2.append(|2 avg fvafs["test"][i, |2 argmax fvafs[i]])
  test dropout = np.expand dims(np.concatenate(test dropout), axis=1)
  test |2 = np.expand dims(np.concatenate(test |2), axis=1)
  test = np.concatenate((test_dropout, test_l2), axis=1)
  # Plot the test set fvaf for the argmaxes
  plot fvaf curves(dropout options["n train folds"],
           ["Dropout", "L2"],
           test,
           "Test Curves")
def load result from experiment(experiment):
  """Load result of given experiment"""
  file str =
"results/batch {}/experiment {}/results dict.pkl".format(experiment["batch num"],
experiment["experiment num"])
  with open(file str, "rb") as fp:
    return pickle.load(fp)
```

```
def load index log(batch num):
  """Load the index log of the batch"""
  experiments = []
  with open("results/batch_{}/index.txt".format(batch_num), "r") as f:
    contents = f.read().split("\n")
    options = json.loads(contents[1])
    for i, experiment str in enumerate(contents[2:len(contents)-1]):
      experiments.append(json.loads(experiment str))
    return options, experiments
def plot_fvaf_curves(n_train_folds_list, key_list, curves, plot_name):
  """Plot fvaf based on the given curves"""
  # Create results directory
  save path = "results/"
  if not os.path.exists(save_path):
    os.mkdir(save_path)
  # Create plots directory
  save_path += "fvaf_plots/"
  if not os.path.exists(save path):
    os.mkdir(save path)
  # Create and configure plot
  fig = plt.figure()
  ax = fig.add_subplot(1, 1, 1)
  for i in range(curves.shape[1]):
    ax.plot(n train folds list, curves[:, i], label=str(key list[i]))
  plt.legend()
  plt.ylabel("Average FVAF")
  plt.xlabel("Number of Training Folds")
  plt.title(plot name)
  # Save
  fig.savefig("{}{} fvaf plot.png".format(save path, plot name), dpi=fig.dpi)
def get_matching_result(results, n_train_fold, key_val, key_name, rotation_val):
  """Find the result that matches the given values"""
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```
for i, result in enumerate(results):
    if (int(result["n train folds"]) == n train fold) and (float(result[key name]) == key val) and
(int(result["rotation"]) == rotation val):
       return result
def compute avg fvaf curves(results, n train fold list, key list, key name, rotation list):
  """Gets the average fvaf across rotations for both the validation and test sets"""
  # Create 3d array of results
  fvaf curves = {
    "val": np.zeros(shape=(len(n train fold list), len(key list), len(rotation list)),
dtype=object),
    "test": np.zeros(shape=(len(n train fold list), len(key list), len(rotation list)),
dtype=object)
  }
  for i, n train fold in enumerate(n train fold list):
    for j, key val in enumerate(key list):
       for k, rotation val in enumerate(rotation list):
         # Get result with the matching parameters
         result = get matching result(results, n train fold, key val, key name, rotation val)
         # Store the validation or test fvaf
         fvaf curves["val"][i, j, k] = result["eval val"][1]
         fvaf curves["test"][i, j, k] = result["eval test"][1]
  # Average across rotations
  fvaf curves["val"] = np.average(fvaf curves["val"], axis=2)
  fvaf_curves["test"] = np.average(fvaf_curves["test"], axis=2)
  return fvaf curves
if __name__ == "__main__":
  main()
```

```
##########
# models.py #
##########
from tensorflow.keras.models import Model
from tensorflow.keras.layers import Input, Dense, Dropout
from tensorflow.keras import regularizers
def pipe model(inputs, layers):
  """Pipes an input through a model to obtain the output hook"""
  for i in range(len(layers)):
    if i == 0:
      carry out = layers[i](inputs)
      carry out = layers[i](carry out)
  return carry out
def dnn(input size, hidden sizes, output size, hidden act="sigmoid", output act="tanh",
dropout=0, I2=0):
  """Construct a simple deep neural network"""
  layers = []
  # Dropout layer if applicable
  if dropout > 0:
    layers.append(Dropout(rate=dropout))
  # Add hidden layers with respective dropout and I2 values
  layers.append(
    hidden stack(hidden sizes, hidden act, dropout=dropout, I2=I2)
    )
  # 12 regularization if applicable
  if 12 > 0:
    layers.append(
      Dense(
         output size,
         activation=output act,
        kernel regularizer=regularizers.l2(l2)
  else:
```

```
layers.append(
      Dense(
        output size,
        activation=output_act
      )
  # Pipe model by feeding through input placeholder
  inputs = Input(shape=input size)
  outputs = pipe_model(inputs, layers)
  return Model(inputs=inputs, outputs=outputs)
def hidden stack(hidden sizes, hidden act="sigmoid", dropout=0, l2=0):
  """Represents a stack of neural layers"""
  layers = []
  for size in hidden sizes:
    # Apply I2 if applicable
    if I2 > 0:
      layers.append(Dense(
           size,
           activation=hidden_act,
           kernel regularizer=regularizers.l2(l2)
           )
    else:
      layers.append(Dense(
           size,
           activation=hidden act,
           )
        )
    # Apply dropout if applicable
    if dropout > 0:
      layers.append(Dropout(rate=dropout))
  def hidden stack layer(inputs):
    """Layer hook for stack"""
    for i in range(len(layers)):
      if i == 0:
         carry out = layers[i](inputs)
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else:
        carry_out = layers[i](carry_out)

return carry_out

return hidden_stack_layer
```

supercomputer_job.sh #

#!/bin/bash

#SBATCH --partition=normal #SBATCH --ntasks=1 #SBATCH --mem=2000 #SBATCH --output=job-output/subprocess-%j-stdout.txt #SBATCH --error=job-output/subprocess--%j-stderr.txt #SBATCH --time=7:00:00

#SBATCH --job-name=subprocess-%j

#SBATCH --mail-user=john.w.spaeth-1@ou.edu

#SBATCH --mail-type=ALL

#SBATCH --chdir=/home/jwspaeth/workspaces/advanced-ml/homework 3/

#SBATCH --wait

python3 supercomputer.py \$@

#!/bin/bash

python3 supercomputer.py \$@





