13-L1 Problem 1

We return to the plane-strain compression problem where the goal is to predict von Mises stress at a node given a set of its features.

Now, you will look at ensemble methods in sklearn to determine how well they perform in this context.

```
import numpy as np
In [1]:
        import matplotlib.pyplot as plt
        from sklearn.linear_model import LinearRegression
        from sklearn.tree import DecisionTreeRegressor
        from sklearn.metrics import mean_squared_error
        from sklearn.ensemble import StackingRegressor
        def plot_shape(dataset, index, model=None, lims=None):
            x = dataset["coordinates"][index][:,0]
            y = dataset["coordinates"][index][:,1]
            if model is None:
                c = dataset["stress"][index]
            else:
                c = model.predict(dataset["features"][index])
            if lims is None:
                lims = [min(c), max(c)]
            plt.scatter(x,y,s=5,c=c,cmap="jet",vmin=lims[0],vmax=lims[1])
            plt.colorbar(orientation="horizontal", shrink=.75, pad=0,ticks=lims)
            plt.axis("off")
            plt.axis("equal")
        def plot_shape_comparison(dataset, index, model, title=""):
            plt.figure(figsize=[6,3.2], dpi=120)
            plt.subplot(1,2,1)
            plot_shape(dataset,index)
            plt.title("Ground Truth",fontsize=9,y=.96)
            plt.subplot(1,2,2)
            c = dataset["stress"][index]
            plot_shape(dataset, index, model, lims = [min(c), max(c)])
            plt.title("Prediction", fontsize=9, y=.96)
            plt.suptitle(title)
            plt.show()
        def load_dataset(path):
            dataset = np.load(path)
            coordinates = []
            features = []
            stress = []
            N = np.max(dataset[:,0].astype(int)) + 1
            split = int(N*.8)
            for i in range(N):
                idx = dataset[:,0].astype(int) == i
                data = dataset[idx,:]
                coordinates.append(data[:,1:3])
```

```
features.append(data[:,3:-1])
    stress.append(data[:,-1])

dataset_train = dict(coordinates=coordinates[:split], features=features[:split], stress=stres
    dataset_test = dict(coordinates=coordinates[split:], features=features[split:], stress=stres
    X_train, X_test = np.concatenate(features[:split], axis=0), np.concatenate(features[split:],
    y_train, y_test = np.concatenate(stress[:split], axis=0), np.concatenate(stress[split:], axis
    return dataset_train, dataset_test, X_train, X_test, y_train, y_test

def get_shape(dataset,index):
    X = dataset["features"][index]
    y = dataset["stress"][index]
    return X, y
```

Loading the data

First, complete the code below to load the data and plot the von Mises stress fields for a few shapes. You'll need to input the path of the data file, the rest is done for you.

All training node features and outputs are in X_train and y_train, respectively. Testing nodes are in X_test, y_test.

dataset_train and dataset_test contain more detailed information such as node coordinates, and they are separated by shape.

Get features and outputs for a shape by calling <code>get_shape(dataset,index)</code>. <code>N_train</code> and <code>N_test</code> are the number of training and testing shapes in each of these datasets.

```
In [2]: # YOU MAY NEED TO EDIT data_path
        data_path = "stress_nodal_features.npy"
        dataset_train, dataset_test, X_train, X_test, y_train, y_test = load_dataset(data_path)
        N_train = len(dataset_train["stress"])
        N_test = len(dataset_test["stress"])
        plt.figure(figsize=[15,3.2], dpi=150)
        for i in range(5):
            plt.subplot(1,5,i+1)
            plot_shape(dataset_train,i)
            plt.title(f"Shape {i}")
        plt.show()
            Shape 0
                                 Shape 1
                                                      Shape 2
                                                                           Shape 3
                                                                                                Shape 4
```

StackingRegressor

0.4928

0.0033

0.5793

0.0047

0.0063

A StackingRegressor consists of \$N\$ fitted regression models, which it evaluates to make \$N\$ predictions. Then, these predictions are fed into another regression model, which makes a final prediction of the target.

0.3841

0.0016

0.8866

0.002

List of Regressors

First, initialize 4 regression models:

- 1. Linear Regression
- 2. Decision Tree regression, max depth 4
- 3. Decision Tree regression, max depth 8
- 4. Decision Tree regression, max depth 12

Then, store these in a list called models.

```
In [3]: # YOUR CODE GOES HERE
        # Define models, and put in a list called 'models'
        models = []
        model1 = LinearRegression()
        model2 = DecisionTreeRegressor(max_depth=4)
        model3 = DecisionTreeRegressor(max_depth=8)
        model4 = DecisionTreeRegressor(max_depth=12)
        models.append(model1)
        models.append(model2)
        models.append(model3)
        models.append(model4)
        named_models = [(f"Model {i+1}", model) for i, model in enumerate(models)]
        print(*named_models, sep="\n")
       ('Model 1', LinearRegression())
       ('Model 2', DecisionTreeRegressor(max_depth=4))
       ('Model 3', DecisionTreeRegressor(max_depth=8))
       ('Model 4', DecisionTreeRegressor(max_depth=12))
```

Final Regressor

Now make one more regressor, which will take as input the other four predictions, and combine them to make an improved prediction.

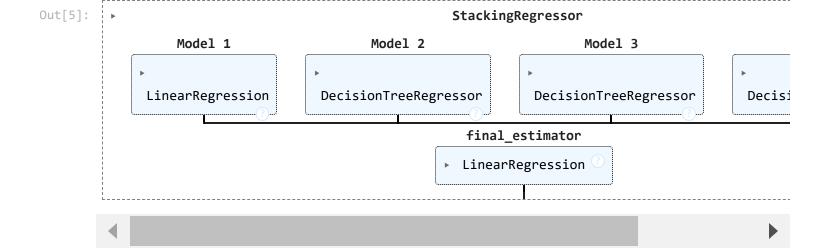
This can be another linear regression model. Call it final_model.

```
In [4]: # YOUR CODE GOES HERE
final_model = LinearRegression()
```

Creating and training the StackingRegressor

Finally, we can combine all of our models into a StackingRegressor model. We fit this just as we would fit any sklearn model. Because of the size of the dataset, this may take a few minutes.

```
In [5]: srm = StackingRegressor(named_models, final_model, verbose=True)
    srm.fit(X_train, y_train)
```



Performance of the model

Now we can investigate the performance of the model on test data, compared to constituent models. First, let's look at the performance of each individual model of our model srm. These can be accessed via srm.estimators.

```
In [6]: for i, estimator in enumerate(srm.estimators_):
            print(f"\n{named_models[i][0]}:")
            train_err = mean_squared_error(y_train, estimator.predict(X_train))
            test_err = mean_squared_error(y_test, estimator.predict(X_test))
            print(f"Training MSE: {train_err:.2e}")
            print(f" Testing MSE: {test_err:.2e}")
      Model 1:
       Training MSE: 8.11e-03
        Testing MSE: 9.78e-03
      Model 2:
      Training MSE: 1.26e-02
       Testing MSE: 1.51e-02
      Model 3:
       Training MSE: 7.56e-03
       Testing MSE: 1.03e-02
      Model 4:
       Training MSE: 3.75e-03
        Testing MSE: 8.40e-03
```

Stacking Regressor MSE on Test Data

Now compute the MSE of srm on training and testing data.

Note how the results, particularly on test data, compare to the individual models.

```
In [7]: # YOUR CODE GOES HERE
    train_err = mean_squared_error(y_train, srm.predict(X_train))
    test_err = mean_squared_error(y_test, srm.predict(X_test))

print(f"\nStacking Regressor:")
    print(f"Training MSE: {train_err:.2e}")
    print(f"Testing MSE: {test_err:.2e}")
```

Stacking Regressor: Training MSE: 4.11e-03 Testing MSE: 6.61e-03

M13-L2 Problem 1

Once more, we will study the stress prediction problem, this time using XGBoost, a very powerful boosting method.

```
In [1]:
        import numpy as np
        import matplotlib.pyplot as plt
        import xgboost as xgb
        from xgboost import XGBRegressor
        from sklearn.metrics import mean_squared_error
        def plot_shape(dataset, index, model=None, lims=None):
            x = dataset["coordinates"][index][:,0]
            y = dataset["coordinates"][index][:,1]
            if model is None:
                c = dataset["stress"][index]
            else:
                c = model.predict(dataset["features"][index])
            if lims is None:
                lims = [min(c), max(c)]
            plt.scatter(x,y,s=5,c=c,cmap="jet",vmin=lims[0],vmax=lims[1])
            plt.colorbar(orientation="horizontal", shrink=.75, pad=0,ticks=lims)
            plt.axis("off")
            plt.axis("equal")
        def plot_shape_comparison(dataset, index, model, title=""):
            plt.figure(figsize=[6,3.2], dpi=120)
            plt.subplot(1,2,1)
            plot_shape(dataset,index)
            plt.title("Ground Truth", fontsize=9, y=.96)
            plt.subplot(1,2,2)
            c = dataset["stress"][index]
            plot_shape(dataset, index, model, lims = [min(c), max(c)])
            plt.title("Prediction", fontsize=9, y=.96)
            plt.suptitle(title)
            plt.show()
        def load_dataset(path):
            dataset = np.load(path)
            coordinates = []
            features = []
            stress = []
            N = np.max(dataset[:,0].astype(int)) + 1
            split = int(N*.8)
            for i in range(N):
                idx = dataset[:,0].astype(int) == i
                data = dataset[idx,:]
                coordinates.append(data[:,1:3])
                features.append(data[:,3:-1])
                stress.append(data[:,-1])
            dataset_train = dict(coordinates=coordinates[:split], features=features[:split], stress=stre
            dataset_test = dict(coordinates=coordinates[split:], features=features[split:], stress=stres
```

```
X_train, X_test = np.concatenate(features[:split], axis=0), np.concatenate(features[split:],
   y_train, y_test = np.concatenate(stress[:split], axis=0), np.concatenate(stress[split:], axi
    return dataset_train, dataset_test, X_train, X_test, y_train, y_test
def get_shape(dataset,index):
   X = dataset["features"][index]
   y = dataset["stress"][index]
   return X, y
def eval_model(model, verbose=False):
   pred train = model.predict(X train)
   pred_test = model.predict(X_test)
   mse_train = mean_squared_error(y_train, pred_train)
   mse_test = mean_squared_error(y_test, pred_test)
   if verbose:
        print(f"Train MSE = {mse_train:.2e}")
        print(f"Test MSE = {mse_test:.2e}")
    return mse_train, mse_test
```

Loading the data

First, complete the code below to load the data and plot the von Mises stress fields for a few shapes. You'll need to input the path of the data file, the rest is done for you.

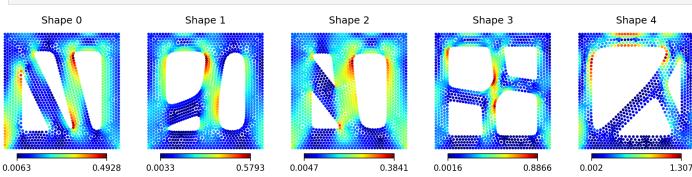
All training node features and outputs are in X_{train} and y_{train} , respectively. Testing nodes are in X_{test} , y_{test} .

dataset_train and dataset_test contain more detailed information such as node coordinates, and they are separated by shape.

Get features and outputs for a shape by calling <code>get_shape(dataset,index)</code>. <code>N_train</code> and <code>N_test</code> are the number of training and testing shapes in each of these datasets.

```
In [3]: # YOU MAY NEED TO EDIT data_path
data_path = "stress_nodal_features.npy"
dataset_train, dataset_test, X_train, X_test, y_train, y_test = load_dataset(data_path)
N_train = len(dataset_train["stress"])
N_test = len(dataset_test["stress"])

plt.figure(figsize=[15,3.2], dpi=150)
for i in range(5):
    plt.subplot(1,5,i+1)
    plot_shape(dataset_train,i)
    plt.title(f"Shape {i}")
plt.show()
```



XGBoost models, like XGBRegressor here, can be used much like sklearn models.

First, define an instance of XGBRegressor with the desired parameters; then, fit the model with model.fit . You can evaluate a fitted model with model.predict .

The provided function mse_train, mse_test = eval_model(model) to get MSE values on the train and test datasets.

```
In [5]: eta = 0.8
depth = 9

params = dict(
    eta = eta,
    max_depth = depth,
)

model = XGBRegressor(objective ='reg:squarederror', seed = 123, n_estimators = 10, **params)
model.fit(X_train, y_train)

mse_train, mse_test = eval_model(model)
print(" eta | depth | Train MSE | Test MSE")
print(f" {eta:.1f} | {depth:>2d} | {mse_train:.2e} | {mse_test:.2e}")

eta | depth | Train MSE | Test MSE
0.8 | 9 | 2.19e-03 | 6.10e-03
```

Parametric study

Now let's examine the effects of varying the parameters eta and max_depth , keeping n_estimators as 10. For every combination of eta in [0.1, 0.3, 0.5, 0.7] and max_depth in [5, 10, 15, 20], train an XGB regressor and report the train and test MSE values.

Which combination has the best performance on testing data?

```
In [6]: # YOUR CODE GOES HERE
    eta = [0.1, 0.3, 0.5, 0.7]
    max_depth = [5, 10, 15, 20]

for e in eta:
    for d in max_depth:
        params = dict(eta = e, max_depth = d)
            model = XGBRegressor(objective ='reg:squarederror', seed = 123, n_estimators = 10, **par.model.fit(X_train, y_train)

        mse_train, mse_test = eval_model(model)
        print(" eta | depth | Train MSE | Test MSE")
        print(f" {e:.1f} | {d:>2d} | {mse_train:.2e} | {mse_test:.2e}")
```

```
eta | depth | Train MSE | Test MSE
0.1 | 5 | 1.05e-02 | 1.28e-02
eta | depth | Train MSE | Test MSE
0.1 | 10 | 5.93e-03 | 8.93e-03
eta | depth | Train MSE | Test MSE
0.1 | 15 | 3.86e-03 | 7.93e-03
eta | depth | Train MSE | Test MSE
0.1 | 20 | 3.35e-03 | 7.98e-03
eta | depth | Train MSE | Test MSE
0.3 | 5 | 5.43e-03 | 7.05e-03
eta | depth | Train MSE | Test MSE
0.3 | 10 | 1.57e-03 | 4.58e-03
eta | depth | Train MSE | Test MSE
0.3 | 15 | 2.79e-04 | 4.59e-03
eta | depth | Train MSE | Test MSE
0.3 | 20 | 8.02e-05 | 4.73e-03
eta | depth | Train MSE | Test MSE
0.5 | 5 | 4.60e-03 | 6.49e-03
eta | depth | Train MSE | Test MSE
0.5 | 10 | 1.36e-03 | 4.80e-03
eta | depth | Train MSE | Test MSE
0.5 | 15 | 1.47e-04 | 5.01e-03
eta | depth | Train MSE | Test MSE
0.5 | 20 | 8.30e-06 | 5.20e-03
eta | depth | Train MSE | Test MSE
0.7 | 5 | 4.83e-03 | 7.03e-03
eta | depth | Train MSE | Test MSE
0.7 | 10 | 1.44e-03 | 5.54e-03
eta | depth | Train MSE | Test MSE
0.7 | 15 | 1.60e-04 | 5.82e-03
eta | depth | Train MSE | Test MSE
0.7 | 20 | 9.05e-06 | 6.02e-03
```

The best model is eta = 0.3 and depth = 10 for the testing data.