## **Problem 2**

Data-driven field prediction models can be used as a substitute for performing expensive calculations/simulations in design loops. For example, after being trained on finite element solutions for many parts, they can be used to predict nodal von Mises stress for a new part by taking in a mesh representation of the part geometry.

Consider the plane-strain compression problem shown in "data/plane-strain.png".

In this problem you are given node features for 100 parts. These node features have been extracted by processing each part shape using a neural network. You will perform feature selection to determine which of these features are most relevant using feature selection tools in sklearn.

You are welcome to use any of the code provided in the lecture activities.

### Summary of deliverables:

SciKit-Learn Models: Print Train and Test MSE

- LinearRegression() with all features
- DecisionTreeRegressor() with all features
- LinearRegression() with features selected by RFE()
- DecisionTreeRegressor() with features selected by RFE()

Feature Importance/Coefficient Visualizations

- Feature importance plot for Decision Tree using all features
- Feature coefficient plot for Linear Regression using all features
- Feature importance plot for DT showing which features RFE selected
- Feature coefficient plot for LR showing which features RFE selected

Stress Field Visualizations: Ground Truth vs. Prediction

- Test dataset shape index 8 for decision tree and linear regression with all features
- Test dataset shape index 16 for decision tree and linear regression with RFE features

#### Questions

• Respond to the 5 prompts at the end

#### **Imports and Utility Functions:**

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.tree import DecisionTreeRegressor
from sklearn.linear_model import LinearRegression
from sklearn.metrics import mean_squared_error
from sklearn.feature_selection import RFE

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], s
    dataset_test = dict(coordinates=coordinates[split:], features=features[split:], st
    X train, X test = np. concatenate (features[:split], axis=0), np. concatenate (features
    y train, y test = np. concatenate(stress[:split], axis=0), np. concatenate(stress[s]
    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 plot importances(model, selected = None, coef=False, title=""):
    plt. figure (figsize= (6, 2), dpi=150)
    y = model.coef_ if coef else model.feature_importances_
    N = 1 + 1 en(y)
    x = np. arange(1, N)
    plt. bar (x, y)
    if selected is not None:
        plt. bar(x[selected], y[selected], color="red", label="Selected Features")
        plt. legend()
    plt. xlabel ("Feature")
```

```
plt.ylabel("Coefficient" if coef else "Importance")
plt.xlim(0,N)
plt.title(title)
plt.show()
```

## 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  $X_{train}$ .

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 get\_shape(dataset,index). N\_train and N\_test are the number of training and testing shapes in each of these datasets.

```
In [61]:
           # YOUR CODE GOES HERE
           # Define data path
           data_path = "data\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
```

## Fitting models with all features

0.0033

0.4928

0.0063

Create two models to fit the training data X\_train , y\_train :

- 1. A LinearRegression() model
- 2. A DecisionTreeRegressor() model with a max\_depth of 20

0.5793

0.0047

0.0016

0.8866

0.002

0.3841

Print the training and testing MSE for each.

```
In [62]:
          # YOUR CODE GOES HERE
          linear_model = LinearRegression()
          linear model. fit (X train, y train)
          linear_train = linear_model.predict(X_train)
          linear_test= linear_model.predict(X_test)
          mse_linear_train = mean_squared_error(y_train, linear_train)
          mse_linear_test = mean_squared_error(y_test, linear_test)
          print("Linear_train:", mse_linear_train)
           print("Linear_test:", mse_linear_test)
           dt model = DecisionTreeRegressor(max depth=20)
           dt model. fit (X train, y train)
          dt_model.predict(X_train)
          dt_train = dt_model.predict(X_train)
          dt_test= dt_model.predict(X_test)
          mse_dt_train = mean_squared_error(y_train, dt_train)
          mse_dt_test = mean_squared_error(y_test, dt_test)
          print("df_train:", mse_dt_train)
          print("df_test:", mse_dt_test)
          Linear train: 0.008110601
```

Linear\_test: 0.009779482 df\_train: 0.0004944875978805109 df test: 0.008283868036732832

#### Visualization

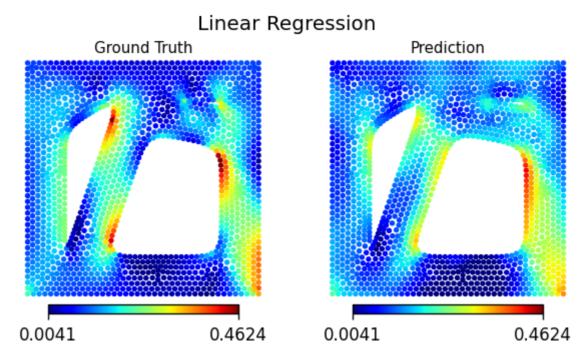
Use the plot\_shape\_comparison() function to plot the index 8 shape results in dataset\_test for each model.

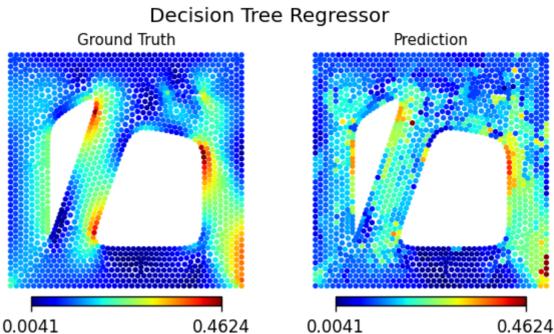
Include titles to indicate which plot is which, using the title argument.

```
In [63]: test_idx = 8

# YOUR CODE GOES HERE

plot_shape_comparison(dataset_test, test_idx, linear_model, title = "Linear Regression")
plot_shape_comparison(dataset_test, test_idx, dt_model, title = "Decision Tree Regressor")
```





# Feature importance

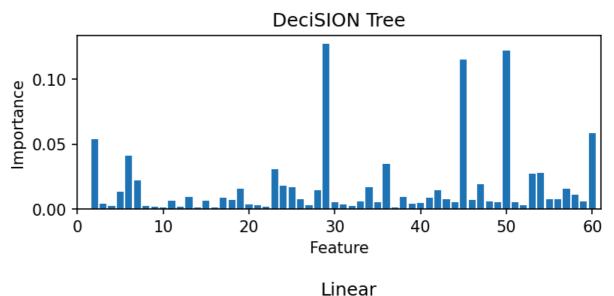
For a tree methods, "feature importance" can be computed, which can be done for an sklearn model using <code>.feature\_importances\_</code>.

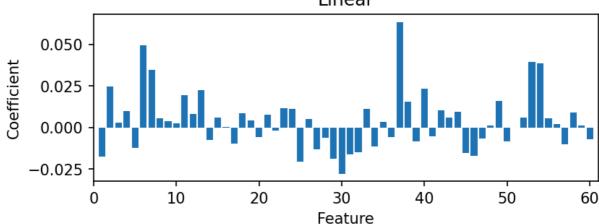
Use the provided function plot\_importances() to visualize which features are most important to the final decision tree prediction.

Then create another plot using the same function to visualize the linear regression coefficients by setting the "coef" argument to True .

```
# YOUR CODE GOES HERE

plot_importances(dt_model, selected = None, coef=False, title="DeciSION Tree")
plot_importances(linear_model, selected = None, coef=True, title="Linear")
```





# Feature Selection by Recursive Feature Elimination

Using RFE() in sklearn, you can iteratively select a subset of only the most important features.

For both linear regression and decision tree (depth 20) models:

- 1. Create a new model.
- 2. Create an instance of RFE() with n\_features\_to\_select set to 30.
- 3. Fit the RFE model as you would a normal sklearn model.
- 4. Report the train and test MSE.

Note that the decision tree RFE model may take a few minutes to train.

Visit https://scikit-learn.org/stable/modules/generated/sklearn.feature\_selection.RFE.html for more information.

```
In [65]: # YOUR CODE GOES HERE
    rfe_linear = RFE(LinearRegression(), n_features_to_select=30)
    rfe_linear.fit(X_train, y_train)
    rfe_linear_train = rfe_linear.predict(X_train)
    rfe_linear_test = rfe_linear.predict(X_test)
    rfe_mse_linear_train = mean_squared_error(y_train, rfe_linear_train)
    rfe_mse_linear_test = mean_squared_error(y_test, rfe_linear_test)
    print("Linear_train:", rfe_mse_linear_train)
    print("Linear_test:", rfe_mse_linear_test)

    rfe_dt = RFE(DecisionTreeRegressor(max_depth=20), n_features_to_select=30)
```

```
rfe_dt.fit(X_train, y_train)
rfe_dt_train = rfe_dt.predict(X_train)
rfe_dt_test = rfe_dt.predict(X_test)
rfe_mse_dt_train = mean_squared_error(y_train, rfe_dt_train)
rfe_mse_dt_test = mean_squared_error(y_test, rfe_dt_test)
print("rf_train:", rfe_mse_dt_train)
print("rf_test:", rfe_mse_dt_test)
```

Linear\_train: 0.008508718 Linear\_test: 0.010150376 rf\_train: 0.0005586730433575364 rf\_test: 0.009001621507497446

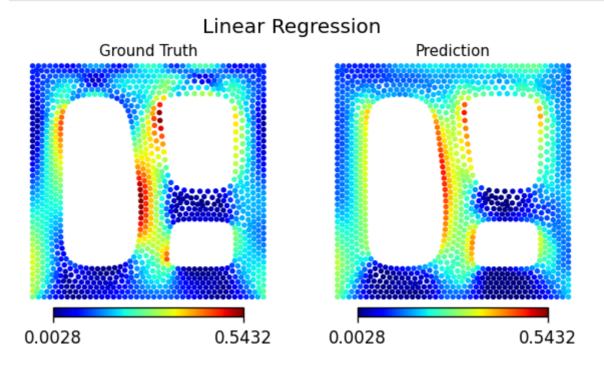
## Visualization

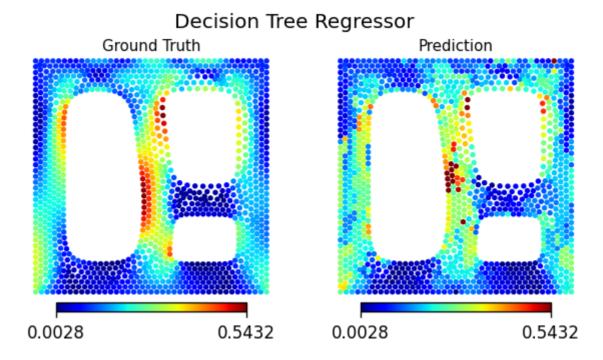
Use the plot\_shape\_comparison() function to plot the index 16 shape results in dataset\_test for each model.

As before, include titles to indicate which plot is which, using the title argument.

```
In [66]:
    test_idx = 16

plot_shape_comparison(dataset_test, test_idx, rfe_linear, title = "Linear Regression")
    plot_shape_comparison(dataset_test, test_idx, rfe_dt, title = "Decision Tree Regressor")
```





# Feature importance with RFE

Recreate the 2 feature importance/coefficent plots from earlier, but this time highlight which features were ultimately selected after performing RFE by coloring those features red. You can do this by setting the selected argument equal to an array of selected indices.

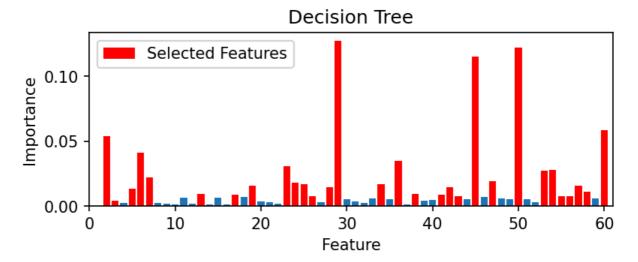
For an RFE model rfe, the selected feature indices can be obtained via rfe.get\_support(indices=True).

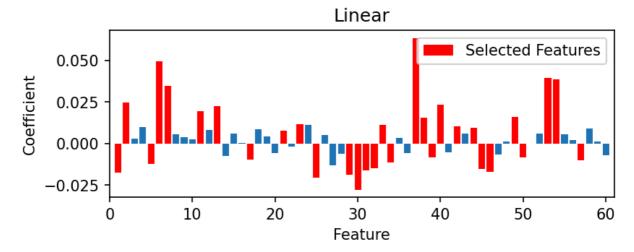
```
# YOUR CODE GOES HERE

dt_select = rfe_dt.get_support(indices = True)
linear_select = rfe_linear.get_support(indices = True)

plot_importances(dt_model, selected = dt_select, coef = False, title="Decision Tree")

plot_importances(linear_model, selected = linear_select, coef = True, title="Linear")
```





## Questions

- 1. Did the MSE increase or decrease on test data for the Linear Regression model after performing RFE?
- 1. Did the MSE increase or decrease on test data for the Decision Tree model after performing RFE?
- 1. Describe the qualitative differences between the Linear Regression and the Decision Tree predictions.
- 1. Describe how the importance of features that were selected by RFE compare to that of features that were eliminated (for the decision tree).
- 1. Describe how the coefficients that were selected by RFE compare to that of features that were eliminated (for linear regression).
- 1. The MSE decrease after performing RFE
- 2. The MSE decrease after performing RFE
- 3. MSE of Decision tree model is always smaller than that of lienar regression model
- 4. The feature that were selected have much higher coefficient than non-selected ones.
- 5. The feature that were selected have relatively higher coefficient than non-selected ones.