## Soil Layering by Cone Penetration Test Data

June 4, 2025

## 1 Soil Layering by Cone Penetration Test Data

This notebook aims to explore various machine learning approaches to automatically determine soil stratification based on CPT data. Author: Zhiyan Jiang (linkedIn.com/zhiyanjiang)

```
[31]: import warnings warnings.simplefilter("ignore") # default #%matplotlib inline
```

```
[32]: # import external libraries
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
import time
from scipy.signal import find_peaks
from scipy.signal import peak_prominences
import pickle
import io

# import internal libraries
from constants import PSF2TSF, PA2TSF, PSI2TSF
from importData import *
from plotCPT import *
from plot import *
```

#### 1.1 Step 1: configurations

```
[33]: cptFileName = "..\\Data\\NSF\\4_5.csv"
  gwtFileName = "..\\Data\\GWT data.csv"
  SBTnImgFileName = '.\\Images\\SBTn_background.jpg'

# Include multiple cptFileNames to the list, if needed
  cptFileNames = [cptFileName] #, "..\\Data\\NSF\\981.csv"]

# Clustering parameters
  randomState = None # 0 or # None, i.e. time.time()
  numberClusters = 4
```

```
# CPT-specific parameters
# area ratio
an = 0.88

# Assume uniform unit weight
soilUnitWeight = 120 # unit shall be pcf
waterUnitWeight = 62.4 # unit shall be pcf

# Applying filtering, unit of feet
# Purpose: to remove consecutive peaks
windowLength = 3
# Repeating filtering. Default = 1
filterTimes = 1

# Remove peaks
removePeaksFlag = False
```

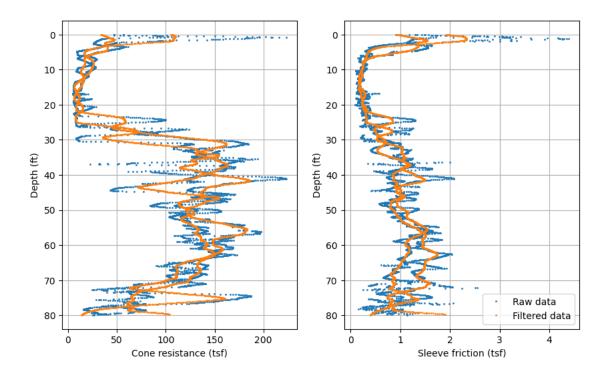
## 1.2 Step 2: Import data, averaging, and removing spikes.

```
[34]: # Import data and apply filtering
plotImportFlag = True
rawData = importCPTs(cptFileNames, windowLength, filterTimes,removePeaksFlag, plotImportFlag)

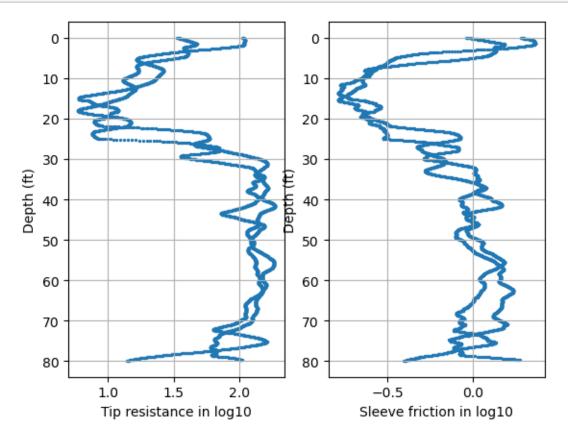
# Import GWT data
gwtData = importGWT(gwtFileName)

# Backup rawData
rawDataCopy = rawData.copy()
```

Row number is: 4866, and Column number is: 4



## 1.3 Step 3: Pre-processing



## 1.4 3.1 Sort

## [37]: # PLACE HOLDER

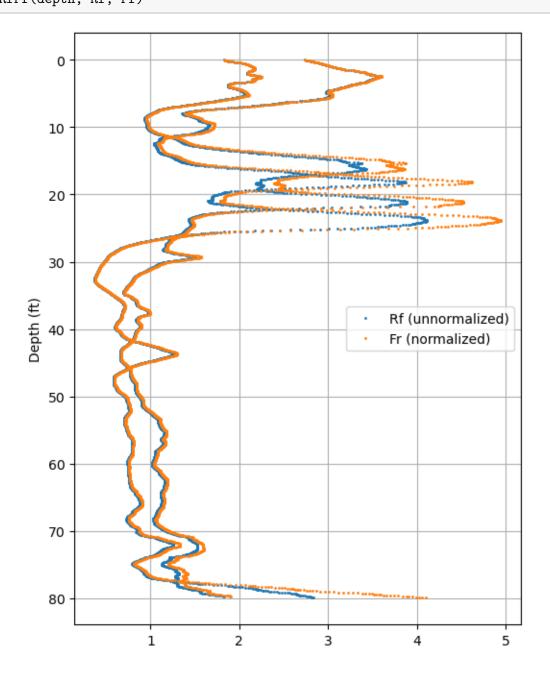
## 1.5 3.2 Calculate derived CPT parameters

The equation for normalized soil behavior type can be found in: Robertson, P.K. Guide to Cone Penetration Testing, 6th Ed., 2015

```
[38]: # Calculate friciton ratio

Rf = calculateRf(data.iloc[:,1], data.iloc[:,2])

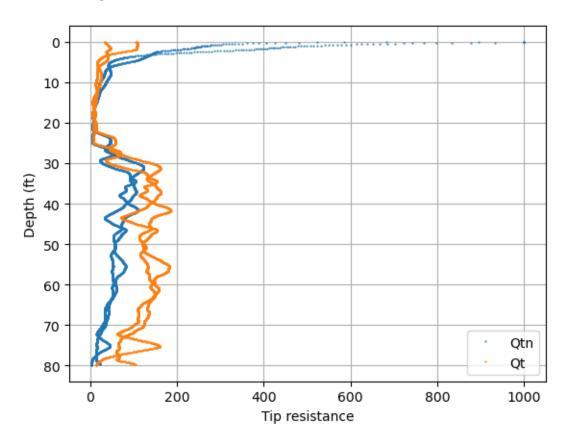
# calculate normalized friction ratio
```



```
[39]: # Calculate stresses
sigma_vo = calculateSigma_vo(data.iloc[:,0].to_frame(), soilUnitWeight)
```

Qtn converged in 4 times

[39]: Text(0.5, 0, 'Tip resistance')



```
[40]: # Calculate and plot NORMALIZED Soil behavior type Index

Ic = calculateIc(Qtn, Fr)
```

## 1.6 Step 4: Clustering

#### 1.7 4.1 Decision Tree

```
[41]: from performDecisionTreeClustering import *
```

#### 1.7.1 4.1.1 Decision Tree on log of [tip resistance, sleeve friction]

The score by Decision tree regression: 0.7923484234428955

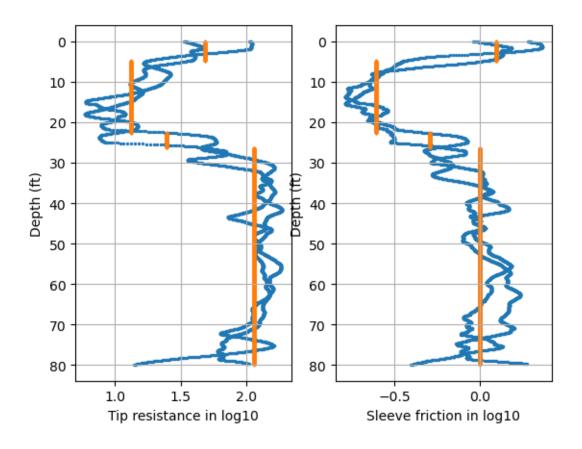
The layer interface depths resulting from Decision tree method is below:

Decision Tree Criteria

0 4.839239

0 4.839239 1 22.621390 2 26.295932

Below is the results by Decision tree regression



#### 1.7.2 4.1.2 Decision Tree on Ic

```
[43]: # Perform decision tree on Ic
      dataIc = pd.concat([data.iloc[:,0], Ic], axis = 1)
      performDecisionTreeFlag = "regression"
      decisionTreeInput = [dataIc, numberClusters, randomState]
      decisionTreeObj, decisionTreeResult =_
       performDecisionTree(performDecisionTreeFlag, decisionTreeInput)
      # Extract decisionTreeCriteria
      decisionTreeCriteria = getDecisionTreeCriteria(decisionTreeObj)
      print(f"The layer interface depths resulting from Decision tree method is below:
       ")
      print(decisionTreeCriteria)
      print()
      # plot decision tree results
      IcAxes = plotIc(data.iloc[:,0].to_frame(), Ic)
      plotDecisionTreeResult(decisionTreeResult, dataIc, IcAxes)
      # Plot Ic histogram
```

```
IcHistogramAxes = plotIcHistogram(Ic)
IcCounts, _ = np.histogram(Ic, bins = 30)

# Plot Icthreshold
plotIcThreadholds(np.max(IcCounts), IcHistogramAxes)

#IcAxes[0].figure
```

The score by Decision tree regression: 0.6668726596924246

The layer interface depths resulting from  $\operatorname{Decision}$  tree  $\operatorname{method}$  is  $\operatorname{below}$ :

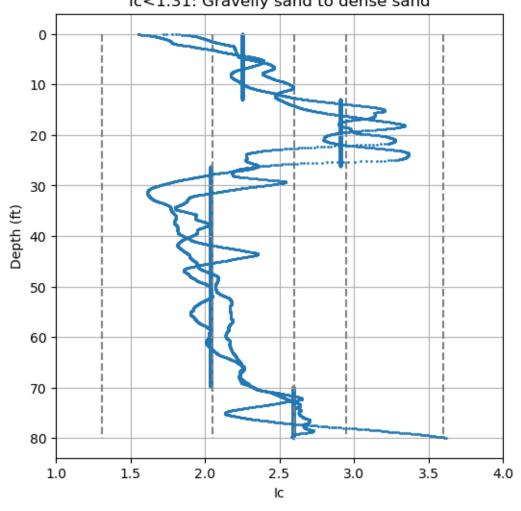
Decision Tree Criteria
0 12.942913
1 26.197507
2 69.832680

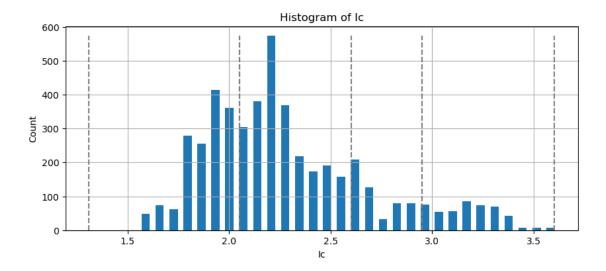
Below is the results by Decision tree regression

[43]: [<Axes: title={'center': 'Histogram of Ic'}, xlabel='Ic', ylabel='Count'>]

lc>3.6: Organic Soils - Clay lc=[2.95,3.6]: Clays - silty clay to clay lc=[2.6,2.95]: silt

Ic=[2.05,2.6]: Sand mixtures - silty sand to sandy silt Ic=[1.31,2.05]: Sands - clean sand to silty sand Ic<1.31: Gravelly sand to dense sand





#### 1.7.3 4.1.3 Decision tree on SBTn type

Infer soil behavior type based on SBTn type, then apply decision tree

```
[44]: # Obtain sbtn dataSBTn1D = calculateSBTn1D(dataIc)
```

4866

```
[45]: # Perform decisiontree regression on SBTn
      performDecisionTreeFlag = "classification" # when using SBTn, must use_
       \hookrightarrow "classification"
      decisionTreeInput = [dataSBTn1D, numberClusters, randomState]
      decisionTreeObj, decisionTreeResult =__
       performDecisionTree(performDecisionTreeFlag, decisionTreeInput)
      # Extract decisionTreeCriteria
      decisionTreeCriteria = getDecisionTreeCriteria(decisionTreeObj)
      print(f"The layer interface depths resulting from Decision tree method is below:
       " )
      print(decisionTreeCriteria)
      print()
      # plot decision tree results
      IcAxes = plotIc(data.iloc[:,0].to_frame(), Ic)
      # Obtain SBTn corresponded middle Ic value
      decisionTreeResultSBTn1DIc = calculateSBTn1DIc(decisionTreeResult)
      plotDecisionTreeResult(decisionTreeResultSBTn1DIc, dataIc, IcAxes)
```

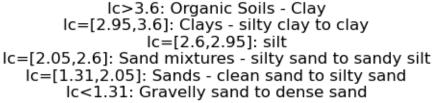
The score by Decision tree regression: 0.6623510069872586

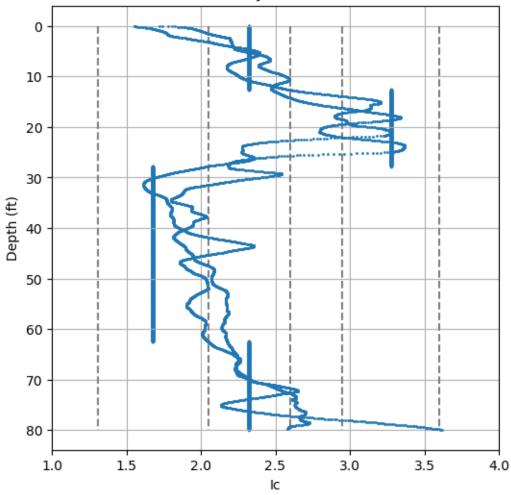
The layer interface depths resulting from  $\operatorname{Decision}$  tree  $\operatorname{method}$  is  $\operatorname{below}$ :

	Decision	Tree Criteria
0		12.647637
1		27.772309
2		62.483595

Below is the results by Decision tree regression

[45]: [<Axes: title={'center': 'Ic>3.6: Organic Soils - Clay\nIc=[2.95,3.6]: Clays - silty clay to clay\nIc=[2.6,2.95]: silt\nIc=[2.05,2.6]: Sand mixtures - silty sand to sandy silt\nIc=[1.31,2.05]: Sands - clean sand to silty sand\nIc<1.31: Gravelly sand to dense sand'}, xlabel='Ic', ylabel='Depth (ft)'>]





#### 1.8 4.2 Random Forest

```
[46]:  # Perform Random Forest
from performRandomForestClustering import *
```

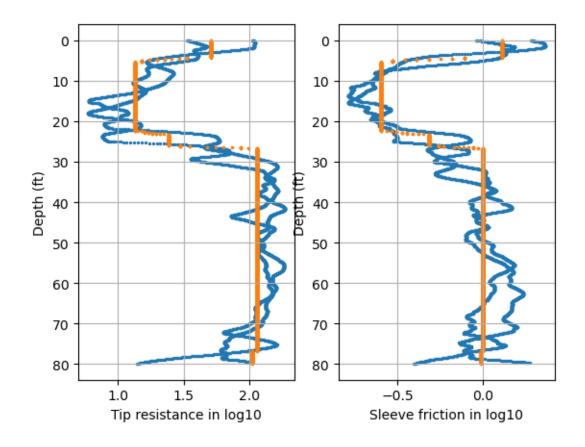
## 1.8.1 4.2.1 Perform random forest regression on log [tip resistance, sleeve friction]

```
[47]: # Perform Random Forest on log of [Tip resistance, sleeve friction]

performRandomForestFlag = "regression"
numberTrees = 10
maxLeafNodes = numberClusters
randomForestInput = [dataLog, numberTrees, maxLeafNodes, randomState]

randomForestObj, randomForestResult = ____
performRandomForest(performRandomForestFlag, randomForestInput)

# plot random forest result
dataLogAxes = plotAggregate(dataLog, labels = dataLog.columns, markerSize = 2,____
axes = None)
plotRandomForestResult(randomForestResult, dataLog, dataLogAxes)
```



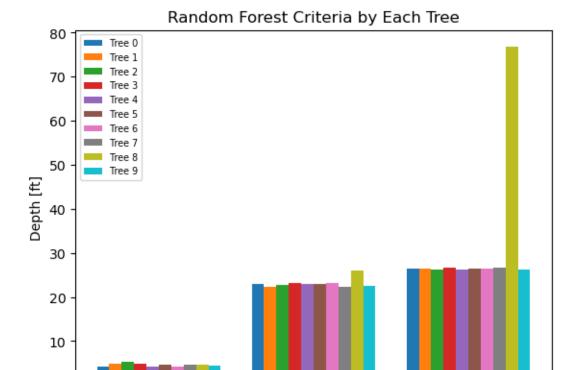
The reduced Random Forest criteria by median is:

0 4.552166 1 22.941273 2 26.410761 dtype: float64 The reduced Random Forest criteria by majority is: Random Forest Criteria 1D  $\,$ 

Labels	
1	4.594816
0	24.680550
2	76.755249

0

0.0



Senstivity analysis on number of layers, i.e., number of leaf nodes

0.5

```
[49]: # test multiple max_leaf_nodes
from testMaxLeafNodes import *

testDecisionTreeFlag = "regression"
testRandomForestFlag = "regression"
testObjFlags = [testDecisionTreeFlag, testRandomForestFlag]
leafNodesRange = [2, 20]

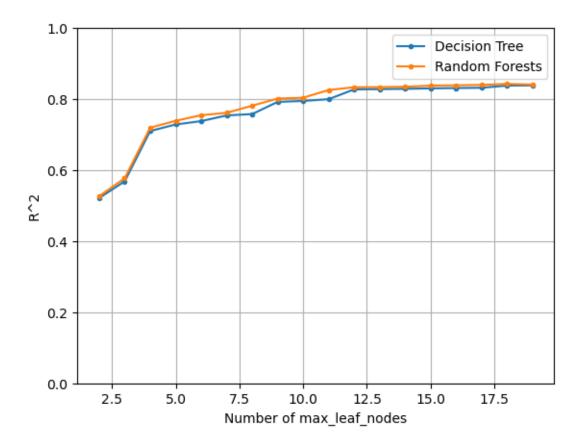
testMaxLeafNodes(leafNodesRange, testObjFlags, data)
```

1.0

1.5

2.0

2.5

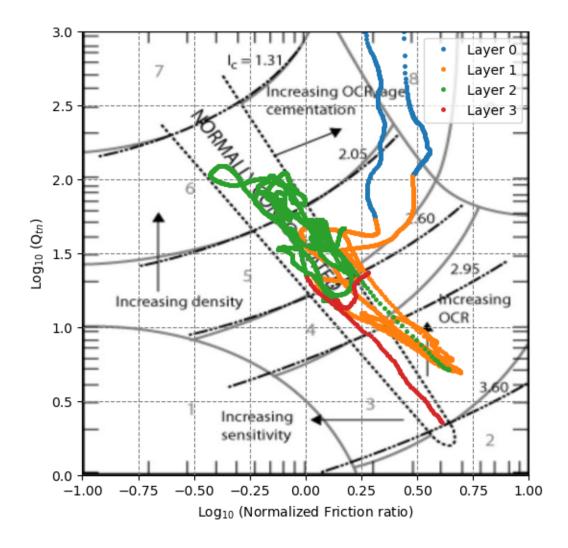


Notes: If only a limit number of CPT soundings is used, need to avoid overfitting. Plot random forest results on Peter Robertson Soil Behavior Type Chart

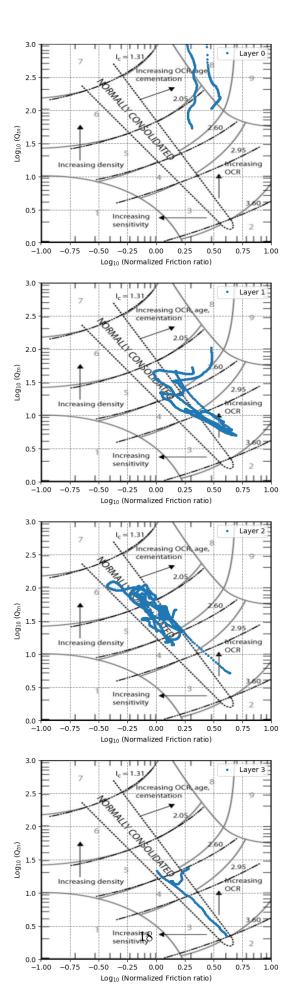
ylabel='Log\$\_{10}\$ (Q\$\_{tn}\$)'>,

ylabel='Log\$\_{10}\$ (Q\$\_{tn}\$)'>,

<Axes: xlabel='Log\$\_{10}\$ (Normalized Friction ratio)',</pre>



<Figure size 640x480 with 0 Axes>



#### 1.8.2 4.2.2 Perform random forest regression on Ic

```
[51]: # Perform Random Forest on Ic
      performRandomForestFlag = "regression"
      numberTrees = 10
      maxLeafNodes = numberClusters
      randomForestInput = [dataIc, numberTrees, maxLeafNodes, randomState]
      randomForestObj, randomForestResult =
       performRandomForest(performRandomForestFlag, randomForestInput)
      # plot random forest result
      IcAxes = plotIc(data.iloc[:,0].to_frame(), Ic)
      plotRandomForestResult(randomForestResult, dataIc, IcAxes)
      # Extract layer interface depths from non-leaf nodes
      randomForestCriteria = getRandomForestCriteria(randomForestObj)
      # plot results by each tree using bar chart
      plt.figure()
      plotRandomForestCriteria(randomForestCriteria)
      # reduce randomForestCriteria as median
      randomForestCriteriaReduced = randomForestCriteriaMedian(randomForestCriteria)
      print()
      print(f"The reduced Random Forest criteria by median is:
       →\n{randomForestCriteriaReduced}")
      # reduce randomForestCriteria as majority
      print()
      randomForestCriteriaReduced = randomForestCriteriaMajority(randomForestCriteria)
      print(f"The reduced Random Forest criteria by majority is:
       →\n{randomForestCriteriaReduced}")
     Use Random Forest regressor.
     The score by Random Forest: 0.6727148693902694
     Below is the results by Random forest regression
     The reduced Random Forest criteria by median is:
          13.106956
     1
          26.148294
          69.816273
     dtype: float64
```

The reduced Random Forest criteria by majority is:

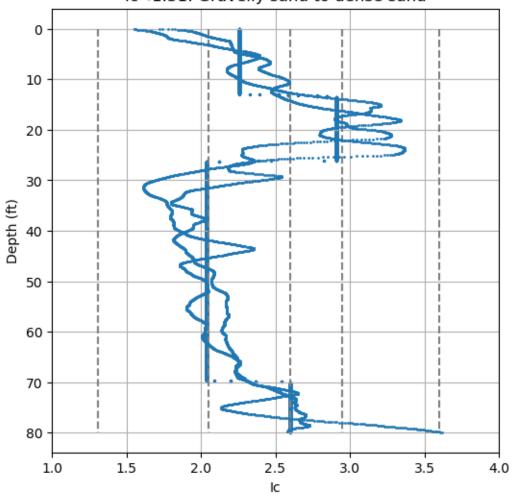
Random Forest Criteria 1D

T	·а	h	e	٦	s

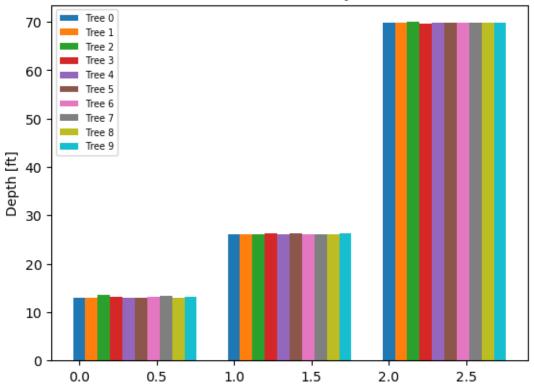
0	13.134842
2	26.163058
1	69.826115

lc>3.6: Organic Soils - Clay lc=[2.95,3.6]: Clays - silty clay to clay lc=[2.6,2.95]: silt

Ic=[2.05,2.6]: Sand mixtures - silty sand to sandy silt Ic=[1.31,2.05]: Sands - clean sand to silty sand Ic<1.31: Gravelly sand to dense sand





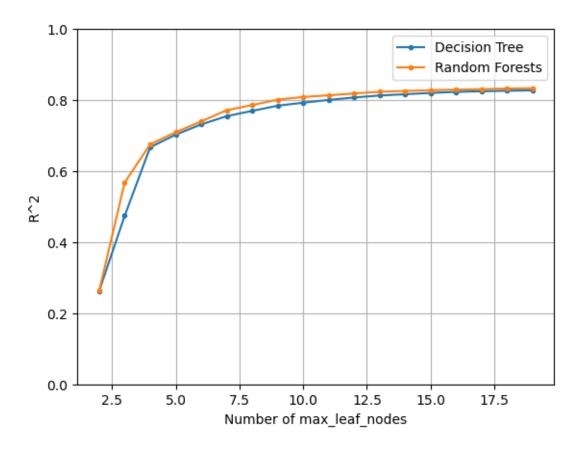


Senstivity analysis on number of layers, i.e., number of leaf nodes

```
[52]: # test multiple max_leaf_nodes
from testMaxLeafNodes import *

testDecisionTreeFlag = "regression"
testRandomForestFlag = "regression"
testObjFlags = [testDecisionTreeFlag, testRandomForestFlag]
leafNodesRange = [2, 20]

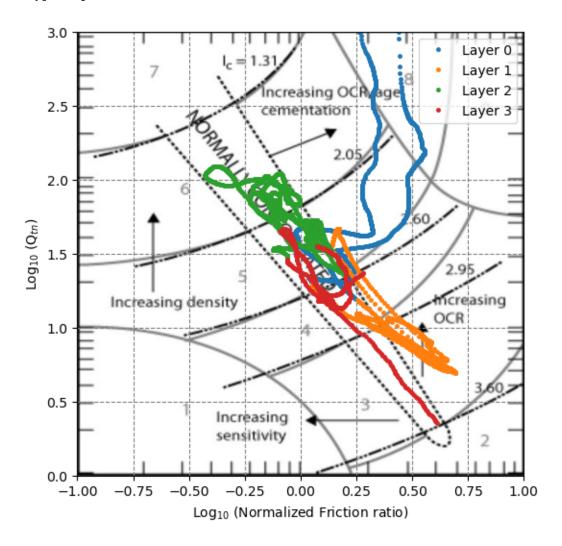
testMaxLeafNodes(leafNodesRange, testObjFlags, dataIc)
```



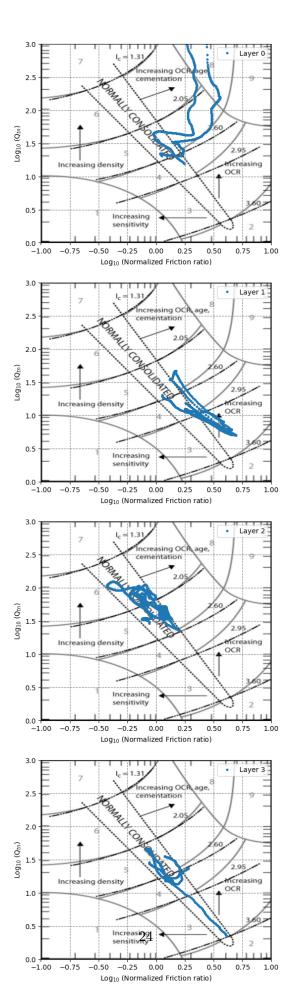
Plot random forest results on Peter Robertson Soil Behavior Type Chart

```
[53]: from applyCriteria import *
      # Get strata index of each data point
      randomForestStrataIndex = getStrataIndex(randomForestCriteriaReduced, dataIc)
      plotSBTnAllinOne(Fr, Qtn, numberClusters, randomForestStrataIndex,
       →SBTnImgFileName)
      plt.figure()
      plotSBTnAllinAll(Fr, Qtn, numberClusters, randomForestStrataIndex, u
        ⇒SBTnImgFileName)
[53]: array([<Axes: xlabel='Log$_{10}$ (Normalized Friction ratio)',
      ylabel='Log$_{10}$ (Q$_{tn}$)'>,
             <Axes: xlabel='Log$_{10}$ (Normalized Friction ratio)',</pre>
      ylabel='Log$_{10}$ (Q$_{tn}$)'>,
             <Axes: xlabel='Log$_{10}$ (Normalized Friction ratio)',</pre>
      ylabel='Log$_{10}$ (Q$_{tn}$)'>,
             <Axes: xlabel='Log$_{10}$ (Normalized Friction ratio)',</pre>
      ylabel='Log$_{10}$ (Q$_{tn}$)'>],
```

dtype=object)



<Figure size 640x480 with 0 Axes>



#### 1.8.3 4.2.3 Perform random forest regression on Ic-correlated SBTn

```
[54]: # Perform random forest regressino on SBTn
      performRandomForestFlag = "classification" # when using SBTn, must use_
       ⇔"classification"
      numberTrees = 10
      maxLeafNodes = numberClusters
      randomForestInput = [dataSBTn1D, numberTrees, maxLeafNodes, randomState]
      randomForestObj, randomForestResult = __
       performRandomForest(performRandomForestFlag, randomForestInput)
      # plot random forest result
      IcAxes = plotIc(data.iloc[:,0].to_frame(), Ic)
      randomForestResultSBTn1DIc = calculateSBTn1DIc(randomForestResult)
      plotRandomForestResult(randomForestResultSBTn1DIc, dataSBTn1D, IcAxes)
      # Extract layer interface depths from non-leaf nodes
      randomForestCriteria = getRandomForestCriteria(randomForestObj)
      # plot results by each tree using bar chart
      plt.figure()
      plotRandomForestCriteria(randomForestCriteria)
      # reduce randomForestCriteria as median
      randomForestCriteriaReduced = randomForestCriteriaMedian(randomForestCriteria)
      print()
      print(f"The reduced Random Forest criteria by median is:
       →\n{randomForestCriteriaReduced}")
      # reduce randomForestCriteria as majority
      randomForestCriteriaReduced = randomForestCriteriaMajority(randomForestCriteria)
      print(f"The reduced Random Forest criteria by majority is:

¬\n{randomForestCriteriaReduced}")
     Use Random Forest classifier.
     The score by Random Forest: 0.6703658035347307
     Below is the results by Random forest regression
     The reduced Random Forest criteria by median is:
          12.795276
          27.829724
     1
          62.450787
     dtype: float64
```

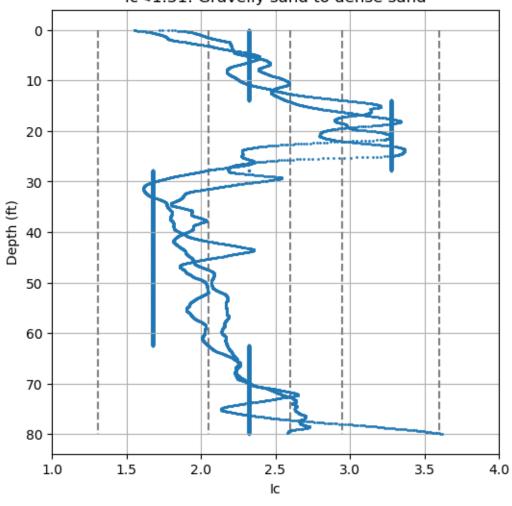
The reduced Random Forest criteria by majority is: Random Forest Criteria 1D  $\,$ 

#### Labels

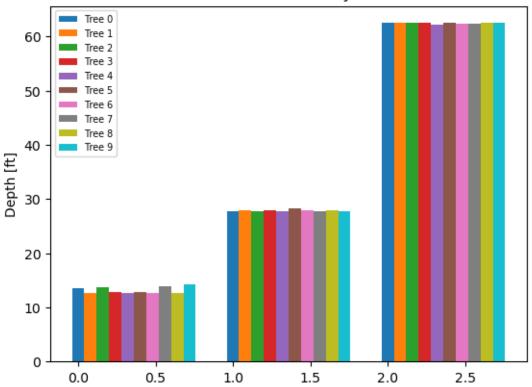
0	13.169291
2	27.870735
1	62.406495

lc>3.6: Organic Soils - Clay lc=[2.95,3.6]: Clays - silty clay to clay lc=[2.6,2.95]: silt

lc=[2.05,2.6]: Sand mixtures - silty sand to sandy silt lc=[1.31,2.05]: Sands - clean sand to silty sand lc<1.31: Gravelly sand to dense sand</pre>

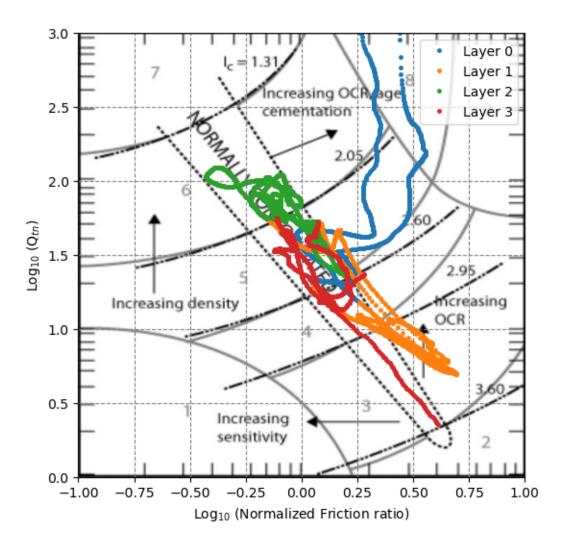


## Random Forest Criteria by Each Tree

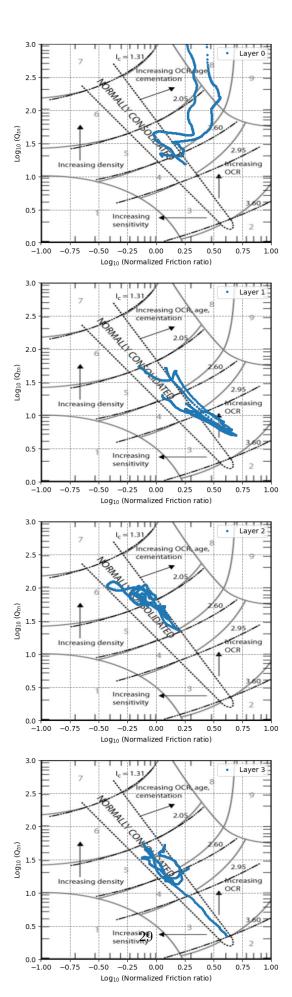


```
[55]: # Get strata index of each data point
randomForestStrataIndex = getStrataIndex(randomForestCriteriaReduced, dataIc)

plotSBTnAllinOne(Fr, Qtn, numberClusters, randomForestStrataIndex,
SBTnImgFileName)
plt.figure()
plotSBTnAllinAll(Fr, Qtn, numberClusters, randomForestStrataIndex,
SBTnImgFileName)
```



<Figure size 640x480 with 0 Axes>



## 1.9 4.3 Agglomerative Clustering

```
[56]: # PLACE HOLDER
```

## 1.10 4.4 Clustering based on 2D SBTn chart

```
[]: import shapely
from shapely.geometry import Point, Polygon
print("External package Shapely is loaded. Version:")
print(shapely.__version__)
```

External package Shapely has been loaded. Version: 2.0.7

```
[58]: # Load digitized shape file
SBTnShapeFile = "..\\Digitize SBTn chart\\SBTn zone shapes.csv"

SBTnShapeData = importSBTnChart(SBTnShapeFile)

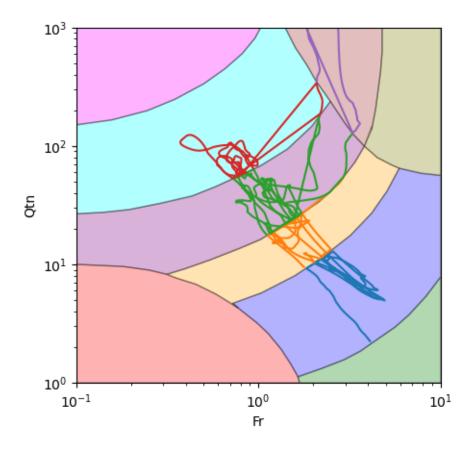
SBTnShapeCoords = digitizeSBTnChart(SBTnShapeData)
```

```
[59]: # Determine SBTn 2D zone index for each point
SBTn2D = calculateSBTn2D(Fr, Qtn, SBTnShapeCoords)

# Verify SBTn2D
verifySBTn2D(Fr, Qtn, SBTn2D, SBTnShapeCoords)

dataSBTn2D = pd.concat([depth, SBTn2D], axis = 1)
```

To verify SBTn2D, Points in each zone shall have the same color.



```
[60]: # Perform random forest on 2D SBTn
      performRandomForestFlag = "classification" # when using 2D SBTn, must use_
       ⇔"classification"
      numberTrees = 10
      maxLeafNodes = numberClusters
      randomForestInput = [dataSBTn2D, numberTrees, maxLeafNodes, randomState]
      randomForestObj, randomForestResult =
       →performRandomForest(performRandomForestFlag, randomForestInput)
      # plot random forest result
      IcAxes = plotIc(data.iloc[:,0].to_frame(), Ic)
      randomForestResultSBTn1DIc = calculateSBTn1DIc(randomForestResult)
      plotRandomForestResult(randomForestResultSBTn1DIc, dataSBTn2D, IcAxes)
      # Extract layer interface depths from non-leaf nodes
      randomForestCriteria = getRandomForestCriteria(randomForestObj)
      # plot results by each tree using bar chart
      plt.figure()
```

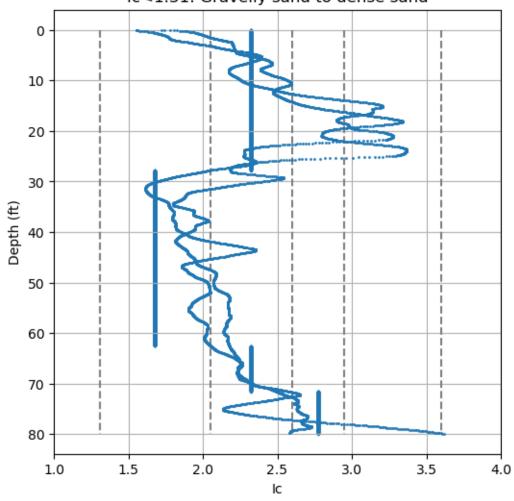
```
plotRandomForestCriteria(randomForestCriteria)
      # reduce randomForestCriteria as median
      randomForestCriteriaReduced = randomForestCriteriaMedian(randomForestCriteria)
      print()
      print(f"The reduced Random Forest criteria by median is:
       →\n{randomForestCriteriaReduced}")
      # reduce randomForestCriteria as majority
      print()
      randomForestCriteriaReduced = randomForestCriteriaMajority(randomForestCriteria)
      print(f"The reduced Random Forest criteria by majority is:⊔
       →\n{randomForestCriteriaReduced}")
      # Get strata index of each data point
      randomForestStrataIndex = getStrataIndex(randomForestCriteriaReduced,__
       →dataSBTn2D)
      plotSBTnAllinOne(Fr, Qtn, numberClusters, randomForestStrataIndex, u
       →SBTnImgFileName)
      plt.figure()
      plotSBTnAllinAll(Fr, Qtn, numberClusters, randomForestStrataIndex,

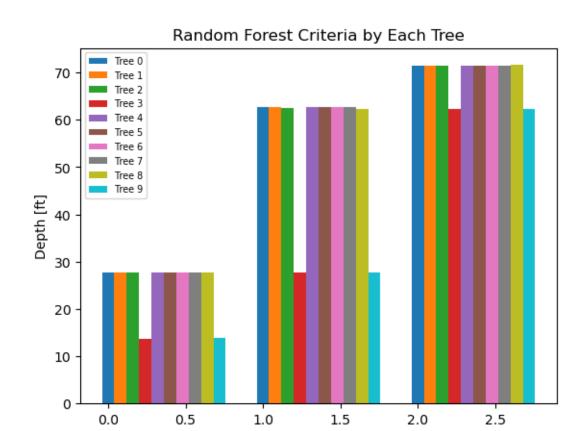
→SBTnImgFileName)

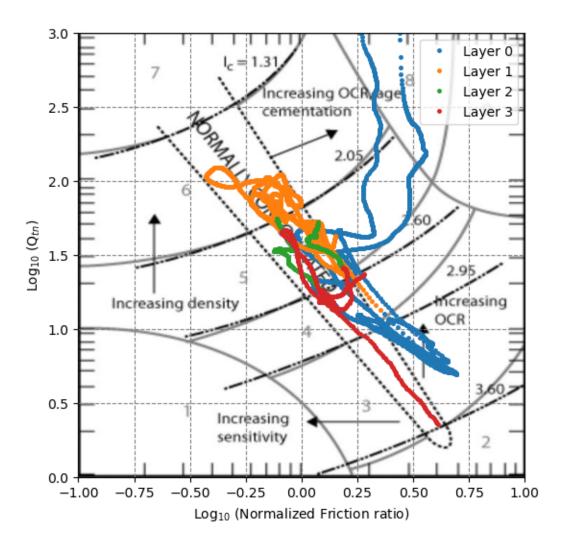
     Use Random Forest classifier.
     The score by Random Forest: 0.6196054254007398
     Below is the results by Random forest regression
     The reduced Random Forest criteria by median is:
          27.772309
          62.582022
          71.432087
     dtype: float64
     The reduced Random Forest criteria by majority is:
             Random Forest Criteria 1D
     Labels
     0
                              25.426509
                              62.491798
     1
                             71.460793
[60]: array([<Axes: xlabel='Log$_{10}$ (Normalized Friction ratio)',
     ylabel='Log$_{10}$ (Q$_{tn}$)'>,
             <Axes: xlabel='Log$_{10}$ (Normalized Friction ratio)',</pre>
      ylabel='Log$_{10}$ (Q$_{tn}$)'>,
             <Axes: xlabel='Log$_{10}$ (Normalized Friction ratio)',</pre>
      ylabel='Log$_{10}$ (Q$_{tn}$)'>,
```

<Axes: xlabel='Log\$\_{10}\$ (Normalized Friction ratio)',</pre>  $ylabel='Log_{10}\ (Q_{tn}\)'>],$ dtype=object)

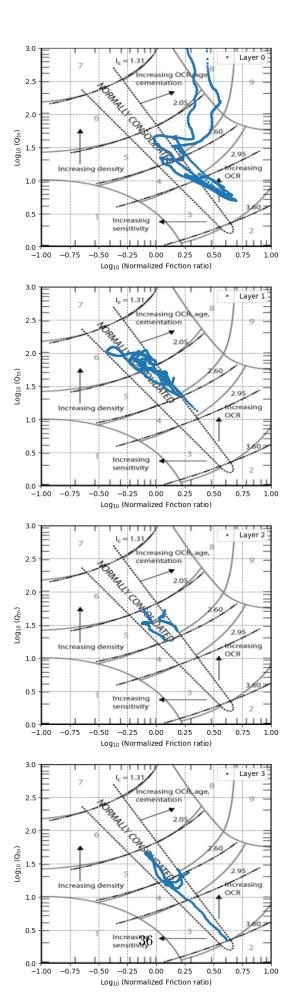
lc>3.6: Organic Soils - Clay
Ic=[2.95,3.6]: Clays - silty clay to clay
Ic=[2.6,2.95]: silt
Ic=[2.05,2.6]: Sand mixtures - silty sand to sandy silt
Ic=[1.31,2.05]: Sands - clean sand to silty sand
Ic<1.31: Gravelly sand to dense sand







<Figure size 640x480 with 0 Axes>



# 2 END