

Institute of Engineering & Technology



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Mini Project Report

On

Leaf Detection

Submitted by

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Declaration

I hereby declare that the work which is being presented in the Mini Project Titled: **“Leaf Detection”**, in partial fulfillment of the requirements for Mini-Project LAB, is an authentic record of our own work carried under the supervision of **Dr. Manoj Vashney, Technical Trainer, GLA University, Mathura.**

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CERTIFICATE

This is to certify that the project entitled “**Leaf Dtection**” carried out in Mini Project – II Lab is a bonafide work done by **Sazal Goyal(171500307), Mudit Singhania(17150052), Sidharth(171500195)** and is submitted in partial fulfillment of the requirements for the award of the degree Bachelor of Technology (Computer Science & Engineering)

Signature of Supervisor:

Name of Supervisor: Dr. Manoj Vashney

Date: 16/04/2020

ABSTRACT

The main objective is to develop for helping the farmers. Automated systems for plant recognition can be used to classify plants into appropriate taxonomies. Such information can be useful for botanists, industrialists, food engineers and physicians. In this work, a recognition system capable of identifying plants by using the images of their leaves has been developed. A mobile application was also developed to allow a user to take pictures of leaves and upload them to a server. The server runs pre-processing and feature extraction techniques on the image before a pattern matcher compares the information from this image with the ones in the database in order to get potential matches.

The different features that are extracted are the length and width of the leaf, the area of the leaf, the perimeter of the leaf, the hull area, the hull perimeter, a distance map along the vertical and horizontal axes, a colour histogram and a centroid-based radial distance map. A k-Nearest Neighbour classifier was implemented and tested on 640 leaves belonging to 32 different species of plants. An accuracy of 83.5% was obtained. The system was further enhanced by using information obtained from a colour histogram which increased the recognition accuracy to 87.3%. Furthermore, our system is simple to use, fast and highly scalabl

ACKNOWLEDGEMENT

It gives us a great sense of pleasure to present the report of the B. Tech Mini Project undertaken during B. Tech. Third Year. This project in itself is an acknowledgement to the inspiration, drive and technical assistance contributed to it by many individuals. This project would never have seen the light of the day without the help and guidance that we have received.

Our heartiest thanks to **Dr. (Prof). Anand Singh Jalal**, Head of Dept., Department of CEA for providing us with an encouraging platform to develop this project, which thus helped us in shaping our abilities towards a constructive goal.

We owe special debt of gratitude to **Dr. Manoj Vashney** , Technical Trainer, Department of CEA, for his constant support and guidance throughout the course of our work. His sincerity, thoroughness and perseverance have been a constant source of inspiration for us. He has showered us with all his extensively experienced ideas and insightful comments at virtually all stages of the project & has also taught us about the latest industry-oriented technologies.

We also do not like to miss the opportunity to acknowledge the contribution of all faculty members of the department for their kind guidance and cooperation during the development of our project. Last but not the least, we acknowledge our friends for their contribution in the completion of the project

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Project Information:

Title Of Project	Leaf Detection												
Role & Responsibility													
Technical Details	<p>Hardware Requirements:</p> <table> <tr> <td>Main Processor</td><td>Core I3</td></tr> <tr> <td>Hard-disk Capacity</td><td>1 G.B</td></tr> <tr> <td>RAM</td><td>2 GB</td></tr> <tr> <td>Clock Speed</td><td>2.8 Hz</td></tr> </table> <p>Keyboard 104 Key</p> <p>Software Requirements:</p> <table> <tr> <td>Operating System</td><td>Windows 10</td></tr> <tr> <td>Language</td><td>Python,Machine Learning Algorithms techniques</td></tr> </table>	Main Processor	Core I3	Hard-disk Capacity	1 G.B	RAM	2 GB	Clock Speed	2.8 Hz	Operating System	Windows 10	Language	Python,Machine Learning Algorithms techniques
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Project Implementation Details	Fully Implemented												

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CHAPTER 1

INTRODUCTION

Introduction: When leaving a town and entering the suburbs, we may encounter many kinds of trees. We may be able to identify those trees that often grow on urban streets, however most of the trees and plants found in city suburbs will be unknown to the majority of us. There are approximately 100,000 species of trees on earth, which account for about 25% of all plants. Many of the trees are in tropical regions, and because only limited botanical research has been carried out in these areas, it is believed that there are many undiscovered species [1]. It is clear that identifying large numbers of such trees is a complex process

Motivation: An example of the complexity of tree identification can be seen with plums and apricots. These are very similar in leaf shape, the shape of the tree, and even in the shape of the young fruit. The flower shape is also very similar, and the tree type can only be identified by determining whether the calyx is attached, or inverted relative to the petal. Additionally, some trees are not easily distinguishable except at particular times; for example, when they bloom or bear fruit. To identify trees like these, considerable information is required, including leaf shape, shape of the leaves that are directly attached to branches, branch shape, shape of the whole tree, tree size, flower shape, flowering time, and fruit.

Objective: When using branches of biology such as cell biology, molecular biology, phytochemistry, or morphologic anatomy, it may be possible to distinguish plants without time constraints. However, it is unrealistic for the general public to identify the names of trees or plants using these methods when, for example, they are walking in a woodland.

1.1 Scope Of Project : It will give increase the cost of barber and decrease the work of Farmers to identify the leaf .

CHAPTER 2

Technology Used

2.1 Dataset

In this project we will apply K-Means on a small dataset of 1600 binary leaf images with different shapes and try to get a feel for the distribution of leaf images using different visualizations that clarify different aspects about how one can interpret K-Means results.

Examples:



2.2 Machine Learning

Machine learning is a growing technology which enables computers to learn automatically from past data. Machine learning uses various algorithms for building mathematical models and making predictions using historical data or information. Currently, it is being used for various tasks such as image recognition, speech recognition, email filtering, Facebook auto-tagging, recommender system, and many more.

This machine learning tutorial gives you an introduction to machine learning along with the wide range of machine learning techniques such as Supervised, Unsupervised, and Reinforcement learning. You will learn about regression and classification models, clustering methods, hidden Markov models, and various sequential models.

Machine learning is a buzzword for today's technology, and it is growing very rapidly day by day. We are using machine learning in our daily life even without knowing it such as Google Maps, Google assistant, Alexa, etc. Below are some most trending real-world applications of Machine Learning.

2.3 Python

Python is a widely used high-level programming language for general-purpose programming. Apart from being open source programming language, python is a great object-oriented, interpreted, and interactive programming language. Python combines remarkable power with very clear syntax. It has modules, classes, exceptions, very high level dynamic data types, and dynamic typing.

Python 3.0 was released in 2008. Although this version is supposed to be backward incompatibles, later on many of its important features have been backported to be compatible with version 2.7. This tutorial gives enough understanding on Python 3 version programming language. Please refer to this link for our Python 2 tutorial.

Python is Interactive – You can actually sit at a Python prompt and interact with the interpreter directly to write your programs.

Python is Object-Oriented – Python supports Object-Oriented style or technique of programming that encapsulates code within objects.

2.4 K-Means Clustering

k-means is one of the simplest unsupervised learning algorithms that solve the well known clustering problem. The procedure follows a simple and easy way to classify a given data set through a certain number of clusters (assume k clusters) fixed apriori. The main idea is to define k centers, one for each cluster. These centers should be placed in a cunning way because of different location causes different result. So, the better choice is to place them as much as possible far away from each other. The next step is to take each point belonging to a given data set and associate it to the nearest center. When no point is pending, the first step is completed and an early group age is done. At this point we need to re-calculate k new centroids as barycenter of the clusters resulting from the previous step. After we have these k new centroids, a new binding has to be done between the same data set points and the nearest new center. A loop has been generated. As a result of this loop we may notice that the k centers change their location step by step until no more changes are done or in other words centers do not move any more.

Leaf Detection

Visualizing K-Means with Leaf Dataset

This script is about perhaps the simplest and most popular **unsupervised learning algorithm** out there: the K-Means clustering algorithm.

In this script we will apply K-Means on a small dataset of 1600 binary leaf images with different shapes and try to get a feel for the distribution of leaf images using different visualizations that clarify different aspects about how one can interpret K-Means results.

We will then continue to see if the K-Means features (distances from cluster centers) are informative in terms of classifying leafs and determine what is the optimal K (number of clusters) for the sake of leaf type classification.

CHAPTER 3

Softwares

3.1 Jupyter

Project Jupyter is a nonprofit organization created to "develop open-source software, open-standards, and services for interactive computing across dozens of programming languages". Spun-off from IPython in 2014 by Fernando Pérez, Project Jupyter supports execution environments in several dozen languages. Project Jupyter's name is a reference to the three core programming languages supported by Jupyter, which are Julia, Python and R, and also a homage to Galileo's notebooks recording the discovery of the moons of Jupiter. Project Jupyter has developed and supported the interactive computing products Jupyter Notebook, JupyterHub, and JupyterLab, the next-generation version of Jupyter Notebook.

3.2 Pycharm

PyCharm is an integrated development environment used in computer programming, specifically for the Python language. It is developed by the Czech company JetBrains.

3.3 Import Library

```
import pandas as pd
```

```
import numpy as np
```

```
import matplotlib
```

```
import matplotlib.image as mpimg
```

```
import matplotlib.pyplot as plt
```

```
import matplotlib.gridspec as gridspec
```

```
from sklearn import model_selection
```

```
from sklearn import decomposition
```

```
from sklearn import linear_model
```

```
from sklearn import cluster
```

```
Leaf Detection
from sklearn import ensemble

from sklearn import neighbors

from sklearn.preprocessing import LabelEncoder

from sklearn.neighbors import KernelDensity

from sklearn.manifold import TSNE

from sklearn.metrics import accuracy_score

from skimage.transform import rescale

from scipy import ndimage as ndi

matplotlib.style.use('fivethirtyeight')
```

CHAPTER 4

Project Implementation

4.1 Import Libraries

```
import pandas as pd
import numpy as np
import matplotlib
import matplotlib.image as mpimg
import matplotlib.pyplot as plt
import matplotlib.gridspec as gridspec
from sklearn import model_selection
from sklearn import decomposition
from sklearn import linear_model
from sklearn import cluster
from sklearn import ensemble
from sklearn import neighbors
from sklearn.preprocessing import LabelEncoder
from sklearn.neighbors import KernelDensity
from sklearn.manifold import TSNE
from sklearn.metrics import accuracy_score

from skimage.transform import rescale
from scipy import ndimage as ndi

matplotlib.style.use('fivethirtyeight')
```

4.2 Store Dataset

```
trainData = pd.read_csv('train.csv')

classEncoder = preprocessing.LabelEncoder()

trainLabels = classEncoder.fit_transform(trainData.loc[:, 'species'])

trainIDs = np.array(trainData.loc[:, 'id'])

# show some random images

plt.figure(figsize=(14, 12))

dataDir = '../demo/'

plt.suptitle('Original Images (with variable image sizes)', fontsize=22)

for k in range(28):

    randTrainInd = np.random.randint(80)

    randomID = trainIDs[randTrainInd]

    path = dataDir + 'images/' + str(randomID) + '.jpg'

    imageFilename = path

    plt.subplot(4, 7, k+1)

    plt.imshow(mpimg.imread(imageFilename), cmap='gray')

    plt.title(classEncoder.classes_[trainLabels[randTrainInd]], fontsize=10)

    plt.axis('off')
```


4.3 Store Data into 2-D Array And Train Data

```

numImages = 80

dataDir='../demo/'

shapesMatrix = np.zeros((2,numImages))

listOfImages = []

for k in range(numImages):

    imageFilename = dataDir + 'images/' + str(k+1) + '.jpg'

    currImage = mpimg.imread(imageFilename)

    shapesMatrix[:,k] = np.shape(currImage)

    listOfImages.append(currImage)

# calculate the shape of an image that will contain all original images within it

maxShapeSize = shapesMatrix.max(axis=1)

for k in range(len(maxShapeSize)):

    if maxShapeSize[k] % 2 == 0:

        maxShapeSize[k] += 311

    else:

        maxShapeSize[k] += 310

# place all original images at the center of the large reference frame

fullImageMatrix3D = np.zeros(np.hstack((maxShapeSize,
np.shape(shapesMatrix[1]))).astype(int),dtype=np.dtype('u1'))

destXc = (maxShapeSize[1]+1)/2; destYc = (maxShapeSize[0]+1)/2

for k, currImage in enumerate(listOfImages):

    Yc, Xc = ndi.center_of_mass(currImage)

    Xd = destXc - Xc; Yd = destYc - Yc

    rowIndLims = (int(round(Yd)),int(round(Yd)+np.shape(currImage)[0]))

```

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```
colIndLims = (int(round(Xd)),int(round(Xd)+np.shape(currImage)[1]))

fullImageMatrix3D[rowIndLims[0]:rowIndLims[1],colIndLims[0]:colIndLims[1],k] = currImage

'''

# make sure nothing was ruined in the process
plt.figure(figsize=(14,7))
plt.suptitle('Processed Images (fixed size)', fontsize=22)
for k in range(28):

    randInd = np.random.randint(np.shape(fullImageMatrix3D)[2])

    plt.subplot(4,7,k+1); plt.imshow(fullImageMatrix3D[:, :, randInd], cmap='gray'); plt.axis('off')

'''

# re crop according to rows and columns that don't have zeros in them in any image
xValid = fullImageMatrix3D.mean(axis=2).sum(axis=0) > 0
yValid = fullImageMatrix3D.mean(axis=2).sum(axis=1) > 0
xLims = (np.nonzero(xValid)[0][0],np.nonzero(xValid)[0][-1])
yLims = (np.nonzero(yValid)[0][0],np.nonzero(yValid)[0][-1])
fullImageMatrix3D = fullImageMatrix3D[yLims[0]:yLims[1],xLims[0]:xLims[1],:]

# make sure nothing was ruined in the process
plt.figure(figsize=(14,7))
plt.suptitle('Final Processed Images (with fixed image size)', fontsize=22)
for k in range(28):

    randInd = np.random.randint(np.shape(fullImageMatrix3D)[2])

    plt.subplot(4,7,k+1); plt.imshow(fullImageMatrix3D[:, :, randInd], cmap='gray'); plt.axis('off')

    if randInd < len(trainLabels):

        plt.title(classEncoder.classes_[trainLabels[randInd]], fontsize=10)

    else:
```

Leaf Detection

```
plt.title('test data sample', fontsize=10)

# scale down all images to be in normal size
rescaleFactor = 0.15

scaledDownImage = rescale(fullImageMatrix3D[:, :, 0], rescaleFactor)
scaledDownImages = np.zeros(np.hstack((np.shape(scaledDownImage),
                                         np.shape(fullImageMatrix3D)[2])), dtype=np.dtype('f4'))

for imInd in range(np.shape(fullImageMatrix3D)[2]):
    scaledDownImages[:, :, imInd] = rescale(fullImageMatrix3D[:, :, imInd], rescaleFactor)

del fullImageMatrix3D
```

4.3 K-Means Clustering

class KmeansModel:

```
def __init__(self, X, numClusters=10, objectPixels=None):
    """
    inputs:
        X                - numSamples x numDimensions matrix
        numClusters      - number of clusters to use
        objectPixels (optional) - an binnary mask image used for presentation
                               will be used as Im[objectPixels] = dataSample
                               must satisfy objectPixels.ravel().sum() = X.shape[1]
    """
    numDataSamples = X.shape[0]
    self.numClusters = numClusters
    if objectPixels is None:
        self.objectPixels = np.ones((1, X.shape[1]), dtype=np.bool)
    else:
        self.objectPixels = objectPixels
    assert(self.objectPixels.ravel().sum() == X.shape[1])
```

```

KmeansModel = cluster.KMeans(n_clusters=numClusters, n_init=5)
self.dataRepresentation = KmeansModel.fit_transform(X)
self.KmeansModel = KmeansModel

# calculate cluster frequency
clusterInds = KmeansModel.labels_
clusterFrequency = []
for clusterInd in range(numClusters):
    clusterFrequency.append((clusterInds == clusterInd).sum()/float(numDataSamples))
self.clusterFrequency = np.array(clusterFrequency)
self.sortedTemplatesByFrequency = np.flipud(np.argsort(clusterFrequency))

def RepresentUsingModel(self, X, representationMethod='distFromAllClusters'):

    if representationMethod == 'distFromAllClusters':
        return self.KmeansModel.transform(X)
    if representationMethod == 'clusterIndex':
        return self.KmeansModel.predict(X)
    if representationMethod == 'oneHotClusterIndex':
        clusterAssignment = self.KmeansModel.predict(X)
        X_transformed = np.zeros((X.shape[0],self.numClusters))
        for sample in range(X.shape[0]):
            X_transformed[sample,clusterAssignment[sample]] = 1
        return X_transformed

def ReconstructUsingModel(self, X_transformed, representationMethod='distFromAllClusters'):

```

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```
if representationMethod == 'clusterIndex':
    clusterAssignment = X_transformed

if representationMethod == 'oneHotClusterIndex':
    clusterAssignment = np.argmax(X_transformed,axis=1)

if representationMethod == 'distFromAllClusters':
    clusterAssignment = np.argmin(X_transformed,axis=1)


X_reconstructed = np.zeros((X_transformed.shape[0],self.KmeansModel.cluster_centers_.shape[1]))
for sample in range(X_transformed.shape[0]):
    X_reconstructed[sample,:] = self.KmeansModel.cluster_centers_[clusterAssignment[sample],:]


return X_reconstructed


def InterpretUsingModel(self, X, representationMethod='clusterIndex'):
    return self.ReconstructUsingModel(\
        self.RepresentUsingModel(X,representationMethod),representationMethod)


# shows the cluster centers

def ShowTemplates(self, numTemplatesToShow=16):
    numTemplatesToShow = min(numTemplatesToShow, self.numClusters)

    numFigRows = np.ceil(np.sqrt(numTemplatesToShow));
    numFigCols = np.ceil(np.sqrt(numTemplatesToShow));
    numTemplatesPerFigure = int(numFigRows*numFigCols)
    numFigures = int(np.ceil(float(numTemplatesToShow)/numTemplatesPerFigure))

    for figureInd in range(numFigures):
        plt.figure()
```

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```
for plotInd in range(numTemplatesPerFigure):

    templateInd = self.sortedTemplatesByFrequency[numTemplatesPerFigure*figureInd + plotInd]

    if templateInd >= self.numClusters:

        break

    templateImage = np.zeros(np.shape(self.objectPixels))

    templateImage[self.objectPixels] = \

        self.KmeansModel.cluster_centers_[templateInd,:].ravel()


    plt.subplot(numFigRows,numFigCols,plotInd+1)

    if np.shape(self.objectPixels)[0] == 1:

        plt.plot(templateImage)

    else:

        plt.imshow(templateImage,cmap='hot'); plt.axis('off')

    plt.title(str(100*self.clusterFrequency[templateInd]):4 + '% frequency');

plt.tight_layout()
```

shows several random model reconstructions

```
def ShowReconstructions(self, X, numReconstructions=6):

    assert(np.shape(X)[1] == self.objectPixels.ravel().sum())

    numSamples = np.shape(X)[0]

    numReconstructions = min(numReconstructions, numSamples)


    originalImage    = np.zeros(np.shape(self.objectPixels))

    reconstructedImage = np.zeros(np.shape(self.objectPixels))


    numReconstructionsPerFigure = min(6, numReconstructions)

    numFigures = int(np.ceil(float(numReconstructions)/numReconstructionsPerFigure))
```

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```
for figureInd in range(numFigures):  
    plt.figure()  
    for plotCol in range(numReconstructionsPerFigure):  
        dataSampleInd = np.random.randint(numSamples)  
        originalImage[self.objectPixels] = X[dataSampleInd,:].ravel()  
        reconstructedImage[self.objectPixels] = \  
            self.InterpretUsingModel(np.reshape(X[dataSampleInd,:],[1,-1])).ravel()  
        diffImage = abs(originalImage - reconstructedImage)  
  
        # original image  
        plt.subplot(3,numReconstructionsPerFigure,0*numReconstructionsPerFigure+plotCol+1)  
        if np.shape(self.objectPixels)[0] == 1:  
            plt.plot(originalImage); plt.title('original signal')  
        else:  
            plt.imshow(originalImage, cmap='gray');  
            plt.title('original image'); plt.axis('off')  
  
        # reconstred image  
        plt.subplot(3,numReconstructionsPerFigure,1*numReconstructionsPerFigure+plotCol+1)  
        if np.shape(self.objectPixels)[0] == 1:  
            plt.plot(reconstructedImage); plt.title('reconstructed signal')  
        else:  
            plt.imshow(reconstructedImage, cmap='gray');  
            plt.title('reconstructed image'); plt.axis('off')  
  
        # diff image  
        plt.subplot(3,numReconstructionsPerFigure,2*numReconstructionsPerFigure+plotCol+1)  
        if np.shape(self.objectPixels)[0] == 1:
```

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```
plt.plot(diffImage); plt.title('abs difference signal')

else:

    plt.imshow(diffImage, cmap='gray');

    plt.title('abs difference image'); plt.axis('off')

plt.tight_layout()
```

shows distribution along the distance from a particular cluster and several examples for that distance

```
def ShowSingleTemplateDistances(self, X, listOfTemplates=[0,1]):
```

```
    showAsTraces = (np.shape(self.objectPixels)[0] == 1)
```

```
    assert(all([(x in range(self.numClusters)) for x in listOfTemplates]))
```

```
    X_rep = self.RepresentUsingModel(X, representationMethod='distFromAllClusters')
```

```
    percentilesToShow = [1,5,10,30,60,99]
```

```
    numReadDataSamplePerPercentile = 4
```

```
    representationPercentiles = []
```

```
    for percentile in percentilesToShow:
```

```
        representationPercentiles.append(np.percentile(self.dataRepresentation, percentile, axis=0))
```

```
    medianRepVec = np.percentile(self.dataRepresentation, 50, axis=0)
```

```
    for templateInd in listOfTemplates:
```

```
        plt.figure(); gs = gridspec.GridSpec(numReadDataSamplePerPercentile+2,
```

```
                                                len(percentilesToShow))
```

```
        # calculate the Gaussian smoothed distribution of values along the eigenvector direction
```

```
        sigmaOfKDE = (representationPercentiles[-1][templateInd] -
```


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```
representationPercentiles[1][templateInd])/100.0

pdfStart = representationPercentiles[1][templateInd] - 15*sigmaOfKDE
pdfStop = representationPercentiles[-1][templateInd] + 15*sigmaOfKDE
xAxis = np.linspace(pdfStart,pdfStop,200)
PDF_Model = KernelDensity(kernel='gaussian', \
    bandwidth=sigmaOfKDE).fit(self.dataRepresentation[:,templateInd].reshape(-1,1))
logPDF = PDF_Model.score_samples(xAxis.reshape(-1,1))
percentileValuesToShow = \
    [representationPercentiles[x][templateInd] for x in range(len(representationPercentiles))]
percentilesToShowLogPDF = \
    PDF_Model.score_samples(np.array(percentileValuesToShow).reshape(-1,1))

# show distribution of distance from current template and red dots at the list of precentiles to show
plt.subplot(gs[0,:])
plt.fill(xAxis, np.exp(logPDF), fc='b', alpha=0.9);
plt.scatter(percentileValuesToShow, np.exp(percentilesToShowLogPDF), c='r',s=40);
plt.title(str(100*self.clusterFrequency[templateInd]):4 + '% assignment frequency');

for plotCol, currPrecentile in enumerate(percentilesToShow):
    currPrecentileRepVec = medianRepVec.copy()
    currPrecentileRepVec[templateInd] = representationPercentiles[plotCol][templateInd]

    currPrecentileImage = np.zeros(np.shape(self.objectPixels))
    currPrecentileRepVec = currPrecentileRepVec[:,np.newaxis].T
    currPrecentileImage[self.objectPixels] = \
        self.ReconstructUsingModel(currPrecentileRepVec).ravel()

# show the median image with current precentile as activation of the curr image
```

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```
plt.subplot(gs[1,plotCol]);
if showAsTraces:
    plt.plot(currPrecentileImage);
    plt.title('precentile: ' + str(percentilesToShow[plotCol]) + '%')
else:
    plt.imshow(currPrecentileImage, cmap='hot');
    plt.title('precentile: ' + str(percentilesToShow[plotCol]) + '%'); plt.axis('off')

# find the most suitable candidates in X for current precentile
distFromPercentile = abs(X_rep[:,templateInd] -
                        representationPercentiles[plotCol][templateInd])
X_inds = np.argpartition(distFromPercentile, \
                        numReadDataSamplePerPercentile)[:numReadDataSamplePerPercentile]
for k, X_ind in enumerate(X_inds):
    currNearestPrecentileImage = np.zeros(np.shape(self.objectPixels))
    currNearestPrecentileImage[self.objectPixels] = X[X_ind,:].ravel()

plt.subplot(gs[2+k,plotCol]);
if showAsTraces:
    plt.plot(currNearestPrecentileImage);
    plt.title('NN with closest percentile');
else:
    plt.imshow(currNearestPrecentileImage, cmap='gray');
    plt.title('NN with closest percentile'); plt.axis('off')
plt.tight_layout()
```

```
def ShowDataScatterPlotsWithTSNE(self, X=None, y=None, tSNE_perplexity=30.0, colorMap='Paired'):
```

Leaf Detection

show the distance from 2 most frequent clusters and the tSNE of the entire "distance form template" space

if X is None:

 X_rep = self.dataRepresentation

else:

 X_rep = self.RepresentUsingModel(X)

if y is None:

 y = np.ones(X_rep.shape[0])

tSNE_KmeansModel = TSNE(n_components=2, perplexity=tSNE_perplexity, random_state=0)

X_rep_tSNE = tSNE_KmeansModel.fit_transform(X_rep)

take the two most frequent patterns

mostFrequent = self.sortedTemplatesByFrequency[:2]

plt.figure()

plt.subplot(1,2,1);

plt.scatter(X_rep[:,mostFrequent[0]], \

 X_rep[:,mostFrequent[1]],c=y,cmap=colorMap,s=10,alpha=0.9)

plt.title('"distance form template" representation');

plt.xlabel('distance from template 1'); plt.ylabel('distance from template 2')

plt.subplot(1,2,2);

plt.scatter(X_rep_tSNE[:,0],X_rep_tSNE[:,1],c=y,cmap=colorMap,s=15,alpha=0.9)

plt.title('t-SNE of Kmeans representation'); plt.xlabel('t-SNE axis1'); plt.ylabel('t-SNE axis2')

def ShowTemplatesInPCASpace(self, X, y=None, tSNE_perplexity=30.0, colorMap='Paired'):

Leaf Detection

```
# show the templates in the 2PC space and the tSNE of the entire PCA space

# build PCA model and project the data onto the PCA space
PCAModel = decomposition.PCA(n_components=60, whiten=False)
X_rep = PCAModel.fit_transform(X)

# project the Kmeans templates onto the PCA space
templates_rep = PCAModel.transform(templateModel.KmeansModel.cluster_centers_)

if y is None:
    y = self.RepresentUsingModel(X, representationMethod='clusterIndex')

tSNE_PCAModel = TSNE(n_components=2, perplexity=tSNE_perplexity, random_state=0)
X_rep_tSNE = tSNE_PCAModel.fit_transform(np.vstack((X_rep, templates_rep)))

plt.figure()
plt.subplot(1,2,1); plt.scatter(X_rep[:,0],X_rep[:,1],c=y,cmap=colorMap,s=15,alpha=0.9)
plt.scatter(templates_rep[:,0],templates_rep[:,1],c='k',cmap=colorMap,s=50)
plt.title('PCA representation'); plt.xlabel('PC1 coeff'); plt.ylabel('PC2 coeff')

nC = templates_rep.shape[0]
plt.subplot(1,2,2);
plt.scatter(X_rep_tSNE[:,-nC,0],\
            X_rep_tSNE[:,-nC,1],c=y,cmap=colorMap,s=15,alpha=0.9)
plt.scatter(X_rep_tSNE[-nC:,0],\
            X_rep_tSNE[-nC:,1],c='k',cmap=colorMap,s=50)
plt.title('t-SNE of PCA representation'); plt.xlabel('t-SNE axis1'); plt.ylabel('t-SNE axis2')
```

4 Efficiency

```
matplotlib.rcParams['font.size'] = 12
```

```
matplotlib.rcParams['figure.figsize'] = (12,9)
```

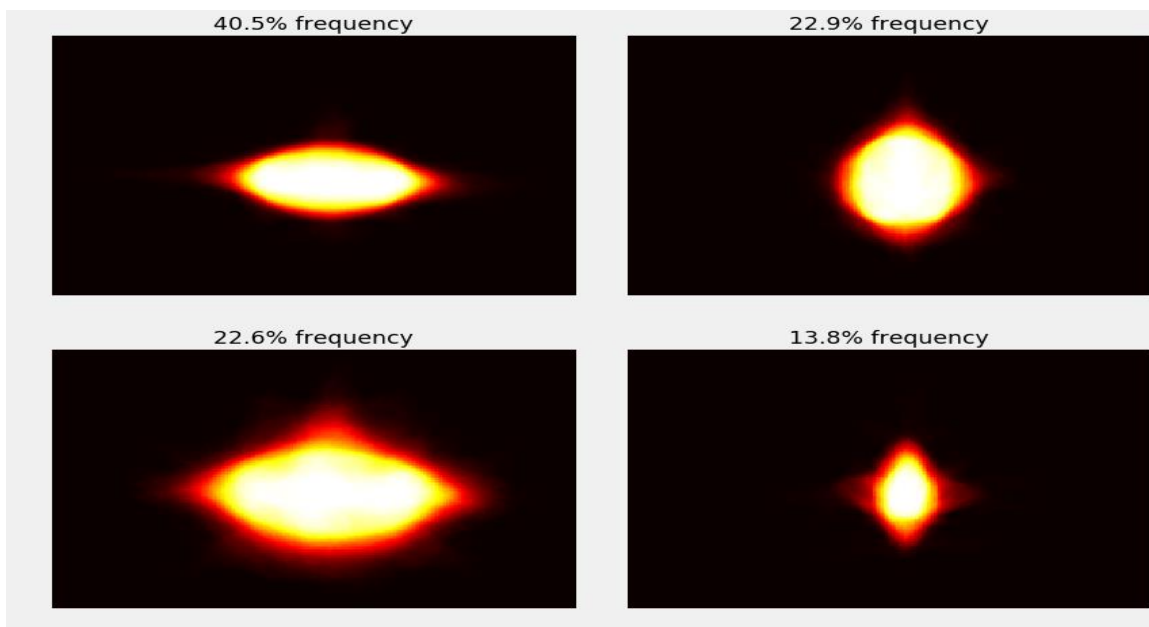
```
objectPixels = np.ones((np.shape(scaledDownImages)[0],np.shape(scaledDownImages)[1])) == 1
```

```
sampleDim = np.shape(scaledDownImages)[0]*np.shape(scaledDownImages)[1]
```

```
X = scaledDownImages.reshape(sampleDim,-1).T
```

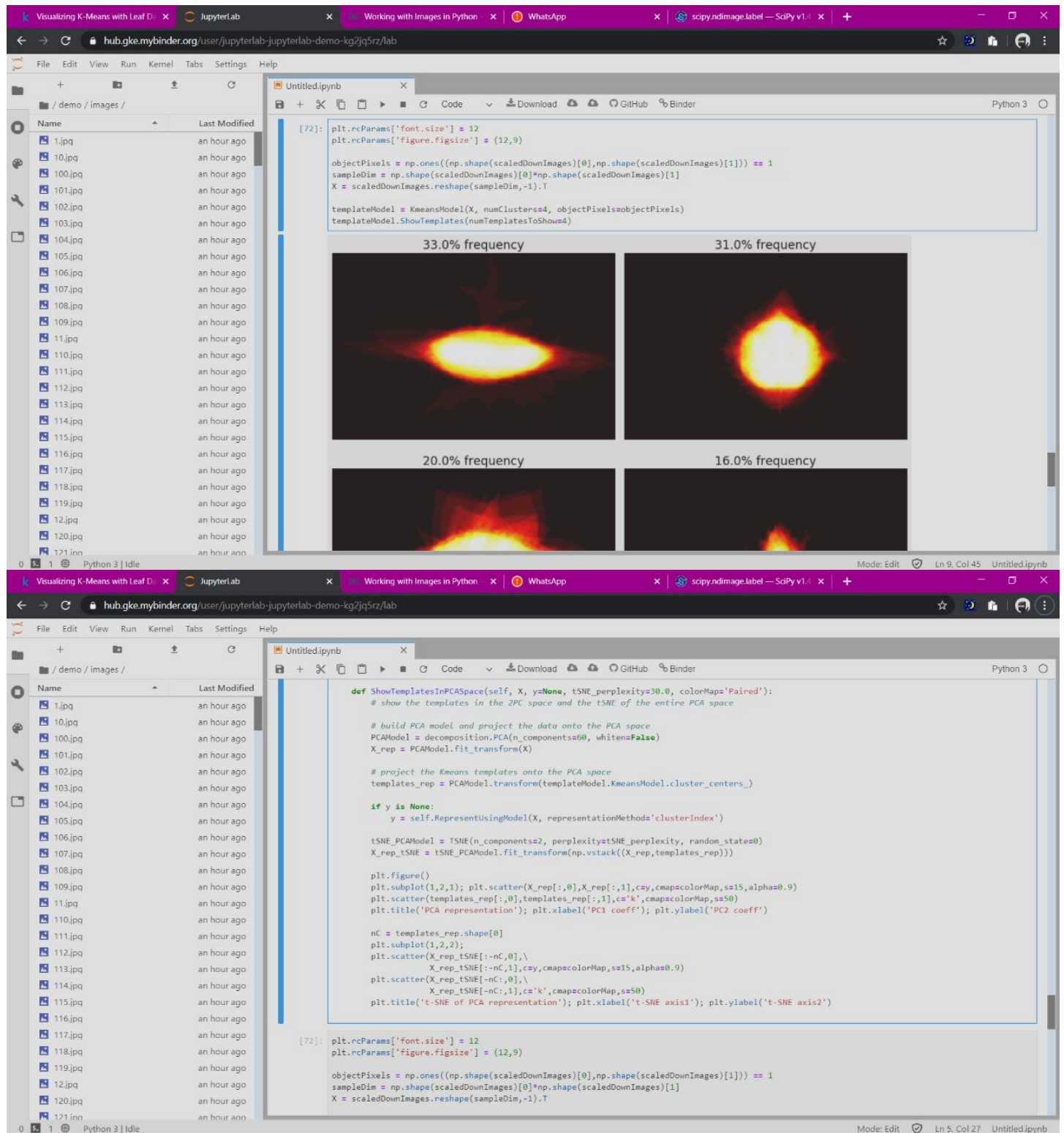
```
templateModel = KmeansModel(X, numClusters=4, objectPixels=objectPixels)
```

```
templateModel.ShowTemplates(numTemplatesToShow=4)
```



Leaf Detection

Screen Shot And Layouts



Leaf Detection

The image displays two screenshots of a JupyterLab environment, showing code for leaf detection. The interface includes a file browser on the left, a code editor in the center, and a terminal at the bottom.

Top Screenshot: ShowDataScatterPlotsWithTSNE

```
def ShowDataScatterPlotsWithTSNE(self, X=None, y=None, tSNE_perplexity=30.0, colorMap='Paired'):
    # show the distance from 2 most frequent clusters and the tSNE of the entire "distance from template" space

    if X is None:
        X_rep = self.dataRepresentation
    else:
        X_rep = self.RepresentUsingModel(X)

    if y is None:
        y = np.ones(X_rep.shape[0])

    tSNE_KmeansModel = TSNE(n_components=2, perplexity=tSNE_perplexity, random_state=0)
    X_rep_tSNE = tSNE_KmeansModel.fit_transform(X_rep)

    # take the two most frequent patterns
    mostFrequent = self.sortedTemplatesByFrequency[:2]

    plt.figure()
    plt.subplot(1,2,1);
    plt.scatter(X_rep[:,mostFrequent[0]], \
                X_rep[:,mostFrequent[1]],c=y,cmap=colorMap,s=10,alpha=0.9)
    plt.title("distance from template" representation");
    plt.xlabel('distance from template 1'); plt.ylabel('distance from template 2')
    plt.subplot(1,2,2);
    plt.scatter(X_rep_tSNE[:,0],X_rep_tSNE[:,1],c=y,cmap=colorMap,s=15,alpha=0.9)
    plt.title('t-SNE of Kmeans representation'); plt.xlabel('t-SNE axis1'); plt.ylabel('t-SNE axis2')

def ShowTemplatesInPCASpace(self, X, y=None, tSNE_perplexity=30.0, colorMap='Paired'):
    # show the templates in the 2PC space and the tSNE of the entire PCA space

    # build PCA model and project the data onto the PCA space
    PCAModel = decomposition.PCA(n_components=60, whiten=False)
    X_rep = PCAModel.fit_transform(X)

    # project the Kmeans templates onto the PCA space
    templates_rep = PCAModel.transform(templateModel.KmeansModel.cluster_centers_)
```

Bottom Screenshot: ShowTemplatesInPCASpace

```
for plotCol, currPercentile in enumerate(percentilesToShow):
    currPercentileRepVec = medianRepVec.copy()
    currPercentileRepVec[templateInd] = representationPercentiles[plotCol][templateInd]

    currPercentileImage = np.zeros(np.shape(self.objectPixels))
    currPercentileRepVec = currPercentileRepVec[:,np.newaxis].T
    currPercentileImage[self.objectPixels] = \
        self.ReconstructUsingModel(currPercentileRepVec).ravel()

    # show the median image with current percentile as activation of the curr image
    plt.subplot(gs[1,plotCol]);
    if showAsTraces:
        plt.plot(currPercentileImage);
        plt.title('percentile: ' + str(percentilesToShow[plotCol]) + '%')
    else:
        plt.imshow(currPercentileImage, cmap='hot');
        plt.title('percentile: ' + str(percentilesToShow[plotCol]) + '%'); plt.axis('off')

    # find the most suitable candidates in X for current percentile
    distFromPercentile = abs(X_rep[:,templateInd] -
                            representationPercentiles[plotCol][templateInd])
    X_inds = np.argsort(distFromPercentile)[:numReadDataSamplePerPercentile]

    for k, X_ind in enumerate(X_inds):
        currNearestPercentileImage = np.zeros(np.shape(self.objectPixels))
        currNearestPercentileImage[self.objectPixels] = X[X_ind,:].ravel()

        plt.subplot(gs[2+k,plotCol]);
        if showAsTraces:
            plt.plot(currNearestPercentileImage);
            plt.title('NN with closest percentile');
        else:
            plt.imshow(currNearestPercentileImage, cmap='gray');
            plt.title('NN with closest percentile'); plt.axis('off')
    plt.tight_layout()
```


Leaf Detection

The image displays two screenshots of a JupyterLab interface, showing code for leaf detection. The interface includes a file browser on the left, a code editor in the center, and a terminal at the bottom.

Top Screenshot: The code editor shows the `ShowSingleTemplateDistances` function. The function takes `X` and `listOfTemplates` as input. It calculates the Gaussian smoothed distribution of values along the eigenvector direction. The code includes comments and assertions to ensure the input is valid. The function uses `plt.figure()` to create a plot and `plt.scatter()` to show the distribution of distance from current template and red dots at the list of percentiles to show.

```
def ShowSingleTemplateDistances(self, X, listOfTemplates=[0,1]):  
    showAsTraces = (np.shape(self.objectPixels)[0] == 1)  
    assert(all([x in range(self.numClusters) for x in listOfTemplates]))  
  
    X_rep = self.RepresentUsingModel(X, representationMethod='distFromAllClusters')  
  
    percentilesToShow = [1,5,10,30,60,99]  
    numReadDataSamplePerPercentile = 4  
    representationPercentiles = []  
    for percentile in percentilesToShow:  
        representationPercentiles.append(np.percentile(self.dataRepresentation, percentile, axis=0))  
    medianRepVec = np.percentile(self.dataRepresentation, 50, axis=0)  
  
    for templateInd in listOfTemplates:  
        plt.figure(); gs = gridspec.GridSpec(numReadDataSamplePerPercentile+2,  
                                            len(percentilesToShow))  
  
        # calculate the Gaussian smoothed distribution of values along the eigenvector direction  
        sigmaOfKDE = (representationPercentiles[-1][templateInd] -  
                      representationPercentiles[1][templateInd])/100.0  
        pdfStart = representationPercentiles[1][templateInd] - 15*sigmaOfKDE  
        pdfStop = representationPercentiles[-1][templateInd] + 15*sigmaOfKDE  
        xAxis = np.linspace(pdfStart,pdfStop,200)  
        PDF_Model = KernelDensity(kernel='gaussian', \n  
                                bandwidth=sigmaOfKDE).fit(self.dataRepresentation[:,templateInd].reshape(-1,1))  
        logPDF = PDF_Model.score_samples(xAxis.reshape(-1,1))  
        percentileValuesToShow = \n  
        [representationPercentiles[x][templateInd] for x in range(len(representationPercentiles))]  
        percentilesToShowLogPDF = \n  
        PDF_Model.score_samples(np.array(percentileValuesToShow).reshape(-1,1))  
  
        # show distribution of distance from current template and red dots at the list of percentiles to show  
        plt.subplot(gs[0,:])  
        plt.fill(xAxis, np.exp(logPDF), fc='b', alpha=0.9);  
        plt.scatter(percentileValuesToShow, np.exp(percentilesToShowLogPDF), c='r', s=40);  
        plt.title(str(100*self.clusterFrequency[templateInd]:4] + '% assignment frequency'));
```

Bottom Screenshot: The code editor shows the `ShowSingleTemplateDistances` function and the `ShowSingleTemplateDistances` function. The function takes `X` and `listOfTemplates` as input. It calculates the Gaussian smoothed distribution of values along the eigenvector direction. The code includes comments and assertions to ensure the input is valid. The function uses `plt.figure()` to create a plot and `plt.scatter()` to show the distribution of distance from current template and red dots at the list of percentiles to show.

```
# original image  
plt.subplot(3,numReconstructionsPerFigure,0*numReconstructionsPerFigure+plotCol+1)  
if np.shape(self.objectPixels)[0] == 1:  
    plt.plot(originalImage); plt.title('original signal')  
else:  
    plt.imshow(originalImage, cmap='gray');  
    plt.title('original image'); plt.axis('off')  
  
# reconstrued image  
plt.subplot(3,numReconstructionsPerFigure,1*numReconstructionsPerFigure+plotCol+1)  
if np.shape(self.objectPixels)[0] == 1:  
    plt.plot(reconstructedImage); plt.title('reconstructed signal')  
else:  
    plt.imshow(reconstructedImage, cmap='gray');  
    plt.title('reconstructed image'); plt.axis('off')  
  
# diff image  
plt.subplot(3,numReconstructionsPerFigure,2*numReconstructionsPerFigure+plotCol+1)  
if np.shape(self.objectPixels)[0] == 1:  
    plt.plot(diffImage); plt.title('abs difference signal')  
else:  
    plt.imshow(diffImage, cmap='gray');  
    plt.title('abs difference image'); plt.axis('off')  
plt.tight_layout()  
  
# shows distribution along the distance from a particular cluster and several examples for that distance  
def ShowSingleTemplateDistances(self, X, listOfTemplates=[0,1]):  
    showAsTraces = (np.shape(self.objectPixels)[0] == 1)  
    assert(all([x in range(self.numClusters) for x in listOfTemplates]))  
  
    X_rep = self.RepresentUsingModel(X, representationMethod='distFromAllClusters')  
  
    percentilesToShow = [1,5,10,30,60,99]  
    numReadDataSamplePerPercentile = 4  
    representationPercentiles = []  
    for percentile in percentilesToShow:
```


Leaf Detection

The image displays two screenshots of a JupyterLab environment, showing code for leaf detection. The interface includes a file browser on the left, a code editor in the center, and a terminal at the bottom.

Top Screenshot: The code editor shows the following code:

```
# shows the cluster centers
def ShowTemplates(self, numTemplatesToShow=16):
    numTemplatesToShow = min(numTemplatesToShow, self.numClusters)

    numFigRows = np.ceil(np.sqrt(numTemplatesToShow));
    numFigCols = np.ceil(np.sqrt(numTemplatesToShow));
    numTemplatesPerFigure = int(numFigRows*numFigCols);
    numFigures = int(np.ceil(float(numTemplatesToShow)/numTemplatesPerFigure))

    for figureInd in range(numFigures):
        plt.figure()
        for plotInd in range(numTemplatesPerFigure):
            templateInd = self.sortedTemplatesByFrequency[numTemplatesPerFigure*figureInd + plotInd]
            if templateInd >= self.numClusters:
                break
            templateImage = np.zeros(np.shape(self.objectPixels))
            templateImage[self.objectPixels] = \
                self.KmeansModel.cluster_centers_[templateInd,:].ravel()

            plt.subplot(numFigRows,numFigCols,plotInd+1)
            if np.shape(self.objectPixels)[0] == 1:
                plt.plot(templateImage)
            else:
                plt.imshow(templateImage,cmap='hot'); plt.axis('off')
            plt.title(str(100*self.clusterFrequency[templateInd][:4] + '% frequency'));
        plt.tight_layout()

# shows several random model reconstructions
def ShowReconstructions(self, X, numReconstructions=6):
    assert(np.shape(X)[1] == self.objectPixels.ravel().sum())
    numSamples = np.shape(X)[0]
    numReconstructions = min(numReconstructions, numSamples)

    originalImage = np.zeros(np.shape(self.objectPixels))
    reconstructedImage = np.zeros(np.shape(self.objectPixels))

    numReconstructionsPerFigure = min(6, numReconstructions)
    numFigures = int(np.ceil(float(numReconstructions)/numReconstructionsPerFigure))
```

Bottom Screenshot: The code editor shows the following code:

```
def RepresentUsingModel(self, X, representationMethod='distFromAllClusters'):

    if representationMethod == 'distFromAllClusters':
        return self.KmeansModel.transform(X)
    if representationMethod == 'clusterIndex':
        return self.KmeansModel.predict(X)
    if representationMethod == 'oneHotClusterIndex':
        clusterAssignment = self.KmeansModel.predict(X)
        X_transformed = np.zeros((X.shape[0],self.numClusters))
        for sample in range(X.shape[0]):
            X_transformed[sample,clusterAssignment[sample]] = 1
        return X_transformed

def ReconstructUsingModel(self, X_transformed, representationMethod='distFromAllClusters'):

    if representationMethod == 'clusterIndex':
        clusterAssignment = X_transformed
    if representationMethod == 'oneHotClusterIndex':
        clusterAssignment = np.argmax(X_transformed,axis=1)
    if representationMethod == 'distFromAllClusters':
        clusterAssignment = np.argmax(X_transformed,axis=1)

    X_reconstructed = np.zeros((X_transformed.shape[0],self.KmeansModel.cluster_centers_.shape[1]))
    for sample in range(X_transformed.shape[0]):
        X_reconstructed[sample,:] = self.KmeansModel.cluster_centers_[clusterAssignment[sample],:]

    return X_reconstructed

def InterpretUsingModel(self, X, representationMethod='clusterIndex'):
    return self.ReconstructUsingModel(\
        self.RepresentUsingModel(X,representationMethod),representationMethod)

# shows the cluster centers
def ShowTemplates(self, numTemplatesToShow=16):
    numTemplatesToShow = min(numTemplatesToShow, self.numClusters)

    numFigRows = np.ceil(np.sqrt(numTemplatesToShow));
    numFigCols = np.ceil(np.sqrt(numTemplatesToShow));
```

Leaf Detection

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demo / images /

Name	Last Modified
1.jpg	an hour ago
10.jpg	an hour ago
100.jpg	an hour ago
101.jpg	an hour ago
102.jpg	an hour ago
103.jpg	an hour ago
104.jpg	an hour ago
105.jpg	an hour ago
106.jpg	an hour ago
107.jpg	an hour ago
108.jpg	an hour ago
109.jpg	an hour ago
11.jpg	an hour ago
110.jpg	an hour ago
111.jpg	an hour ago
112.jpg	an hour ago
113.jpg	an hour ago
114.jpg	an hour ago
115.jpg	an hour ago
116.jpg	an hour ago
117.jpg	an hour ago
118.jpg	an hour ago
119.jpg	an hour ago
12.jpg	an hour ago
120.jpg	an hour ago
121.jpg	an hour ago

Untitled.ipynb

```
[70]: class KmeansModel:
def __init__(self, X, numClusters=10, objectPixels=None):
'''
inputs:
X - numSamples x numDimensions matrix
numClusters - number of clusters to use
objectPixels (optional) - an binary mask image used for presentation
will be used as Im[objectPixels] = dataSample
must satisfy objectPixels.ravel().sum() = X.shape[1]
'''
numDataSamples = X.shape[0]
self.numClusters = numClusters
if objectPixels is None:
self.objectPixels = np.ones((1,X.shape[1]),dtype=np.bool)
else:
self.objectPixels = objectPixels
assert(self.objectPixels.ravel().sum() == X.shape[1])
KmeansModel = cluster.KMeans(n_clusters=numClusters, n_init=5)
self.dataRepresentation = KmeansModel.fit_transform(X)
self.KmeansModel = KmeansModel
# calculate cluster frequency
clusterInds = KmeansModel.labels_
clusterFrequency = []
for clusterInd in range(numClusters):
clusterFrequency.append((clusterInds == clusterInd).sum()/float(numDataSamples))
self.clusterFrequency = np.array(clusterFrequency)
self.sortedTemplatesByFrequency = np.flipud(np.argsort(clusterFrequency))
def RepresentUsingModel(self, X, representationMethod='distFromAllClusters'):
if representationMethod == 'distFromAllClusters':
return self.KmeansModel.transform(X)
if representationMethod == 'clusterIndex':
return self.KmeansModel.predict(X)
if representationMethod == 'oneHotClusterIndex':
```

Mode: Edit | Ln 5, Col 27 | Untitled.ipynb

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demo / images /

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106.jpg	an hour ago
107.jpg	an hour ago
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11.jpg	an hour ago
110.jpg	an hour ago
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114.jpg	an hour ago
115.jpg	an hour ago
116.jpg	an hour ago
117.jpg	an hour ago
118.jpg	an hour ago
119.jpg	an hour ago
12.jpg	an hour ago
120.jpg	an hour ago
121.jpg	an hour ago

Untitled.ipynb

Final Processed Images (with fixed image size)

Acer Rufinerve	Quercus Crassifolia	Cornus Chinensis	Betula Austrosinensis	Quercus Agrifolia	Liriodendron Tulipifera	Quercus Palustris
Zelkova Serrata	Celtis Koraiensis	Quercus Pyrenaica	Liriodendron Tulipifera	Liquidambar Styraciflua	Alnus Maximowiczii	Quercus Hartwissiana
Acer Saccharinum	Acer Platanoids	Quercus Canariensis	Acer Circinatum	Alnus Sieboldiana	Cotinus Coggygria	Quercus Pyrenaica
Quercus Canariensis	Betula Austrosinensis	Quercus Pontica	Quercus Texana	Cornus Chinensis	Quercus Hartwissiana	Acer Circinatum

```
[70]: class KmeansModel:
def __init__(self, X, numClusters=10, objectPixels=None):
'''
inputs:
X - numSamples x numDimensions matrix
numClusters - number of clusters to use
objectPixels (optional) - an binary mask image used for presentation
will be used as Im[objectPixels] = dataSample
must satisfy objectPixels.ravel().sum() = X.shape[1]
'''
```

Mode: Command | Ln 1, Col 16 | Untitled.ipynb

Leaf Detection

The image displays two screenshots of a JupyterLab environment, showing a file browser on the left and a code editor on the right.

Top Screenshot: The file browser shows a directory named `demo / images /` containing files from `1.jpg` to `121.jpg`. The code editor shows the following Python code:

```
# make sure nothing was ruined in the process
plt.figure(figsize=(14,7))
plt.suptitle('Processed Images (fixed size)', fontsize=22)
for k in range(28):
    randInd = np.random.randint(np.shape(fullImageMatrix3D)[2])
    plt.subplot(4,7,k+1); plt.imshow(fullImageMatrix3D[:, :, randInd], cmap='gray'); plt.axis('off')
    ...

# re crop according to rows and columns that don't have zeros in them in any image
xValid = fullImageMatrix3D.mean(axis=2).sum(axis=0) > 0
yValid = fullImageMatrix3D.mean(axis=2).sum(axis=1) > 0
xLims = (np.nonzero(xValid)[0][0], np.nonzero(xValid)[0][-1])
yLims = (np.nonzero(yValid)[0][0], np.nonzero(yValid)[0][-1])
fullImageMatrix3D = fullImageMatrix3D[yLims[0]:yLims[1], xLims[0]:xLims[1], :]

# make sure nothing was ruined in the process
plt.figure(figsize=(14,7))
plt.suptitle('Final Processed Images (with fixed image size)', fontsize=22)
for k in range(28):
    randInd = np.random.randint(np.shape(fullImageMatrix3D)[2])
    plt.subplot(4,7,k+1); plt.imshow(fullImageMatrix3D[:, :, randInd], cmap='gray'); plt.axis('off')
    if randInd < len(trainLabels):
        plt.title(classEncoder.classes_[trainLabels[randInd]], fontsize=10)
    else:
        plt.title('test data sample', fontsize=10)

# scale down all images to be in normal size
rescaleFactor = 0.15

scaledDownImage = rescale(fullImageMatrix3D[:, :, 0], rescaleFactor)
scaledDownImages = np.zeros((np.shape(scaledDownImage),
                             np.shape(fullImageMatrix3D)[2]), dtype=np.dtype('f4'))
for imInd in range(np.shape(fullImageMatrix3D)[2]):
    scaledDownImages[:, :, imInd] = rescale(fullImageMatrix3D[:, :, imInd], rescaleFactor)

del fullImageMatrix3D
```

Bottom Screenshot: The file browser shows the same directory. The code editor shows the following Python code:

```
[69]: numImages = 100
dataDir = './demo/'
shapesMatrix = np.zeros((2, numImages))
listOfImages = []
for k in range(numImages):
    imageFilename = dataDir + 'images/' + str(k+1) + '.jpg'
    currImage = mpimg.imread(imageFilename)
    shapesMatrix[:, k] = np.shape(currImage)
    listOfImages.append(currImage)

# calculate the shape of an image that will contain all original images within it
maxShapeSize = shapesMatrix.max(axis=1)
for k in range(len(maxShapeSize)):
    if maxShapeSize[k] % 2 == 0:
        maxShapeSize[k] += 311
    else:
        maxShapeSize[k] += 310

# place all original images at the center of the large reference frame
fullImageMatrix3D = np.zeros((maxShapeSize, np.shape(shapesMatrix[1])), dtype=np.dtype('u1'))
destXc = (maxShapeSize[1]+1)/2; destYc = (maxShapeSize[0]+1)/2
for k, currImage in enumerate(listOfImages):
    Yc, Xc = ndi.center_of_mass(currImage)
    Xd = destXc - Xc; Yd = destYc - Yc
    rowIndims = (int(round(Yd)), int(round(Yd)+np.shape(currImage)[0]))
    colIndims = (int(round(Xd)), int(round(Xd)+np.shape(currImage)[1]))
    fullImageMatrix3D[rowIndims[0]:rowIndims[1], colIndims[0]:colIndims[1], k] = currImage
    ...

# make sure nothing was ruined in the process
plt.figure(figsize=(14,7))
plt.suptitle('Processed Images (fixed size)', fontsize=22)
for k in range(28):
    randInd = np.random.randint(np.shape(fullImageMatrix3D)[2])
    plt.subplot(4,7,k+1); plt.imshow(fullImageMatrix3D[:, :, randInd], cmap='gray'); plt.axis('off')
    ...
```


Leaf Detection

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demo / images /

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10.jpg	an hour ago
100.jpg	an hour ago
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12.jpg	an hour ago
120.jpg	an hour ago
121.jpg	an hour ago

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```

path=dataDir+'images/'+str(randomID)+'.jpg'
imagefilename = path
plt.subplot(4,7,k+1)
plt.imshow(mpimg.imread(imagefilename), cmap='gray')
plt.title(classEncoder.classes_[trainLabels[randTrainInd]], fontsize=10)
plt.axis('off')

```

Original Images (with variable image sizes)

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119.jpg	an hour ago
12.jpg	an hour ago
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```

[62]: import pandas as pd
import numpy as np
import matplotlib
import matplotlib.image as mpimg
import matplotlib.pyplot as plt
import matplotlib.gridspec as gridspec

from sklearn import model_selection
from sklearn import decomposition
from sklearn import linear_model
from sklearn import cluster
from sklearn import ensemble
from sklearn import neighbors
from sklearn.preprocessing import LabelEncoder
from sklearn.neighbors import KernelDensity
from sklearn.manifold import TSNE
from sklearn.metrics import accuracy_score

from skimage.transform import rescale
from scipy import ndimage as ndi

matplotlib.style.use('fivethirtyeight')

[60]: trainData = pd.read_csv('train.csv')
classEncoder = preprocessing.LabelEncoder()
trainLabels = classEncoder.fit_transform(trainData.loc[:, 'species'])
trainIDs = np.array(trainData.loc[:, 'id'])
# show some random images
plt.figure(figsize=(14,12))
dataDir='../demo/'
plt.suptitle('Original Images (with variable image sizes)', fontsize=22)
for k in range(20):
    randTrainInd = np.random.randint(100)
    randomID = trainIDs[randTrainInd]
    path=dataDir+'images/'+str(randomID)+'.jpg'
    imagefilename = path

```

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CHAPTER 5

REFERENCES

- <https://www.google.com/>
- <https://www.javatpoint.com/>
- <https://stackoverflow.com/>
- <https://www.kaggle.com/>