COSC522: Final Project - Catifier

Cameron Adkins, Purnachandra Anirudh Gajjala, Gabriel Abeyie

```
In [1]: # Numpy.
        import numpy as np
        from numpy.lib.stride tricks import sliding window view
        # Need plots.
        import matplotlib.pyplot as plt
        # Pandas.
        import pandas as pd
        # Machine learning toolkit.
        from sklearn.model_selection import KFold, cross_val_score
        from sklearn.ensemble import RandomForestClassifier
        from sklearn.svm import SVC
        from sklearn.tree import DecisionTreeClassifier
        from sklearn.model selection import train test split
        from sklearn.preprocessing import *
        from sklearn.preprocessing import Normalizer
        from sklearn.metrics import confusion matrix
        from sklearn.metrics import classification report, accuracy score
        # Scipy for fft's and the like.
        import scipy as sc
        import scipy.io.wavfile as wavfile
        from scipy import signal
        from scipy.fftpack import fft, fftfreq
        from scipy import stats
        # Seaborn for plots.
        import seaborn as sns
        # IPython for basic visual output types.
        import IPython
        # Imbalanced learn for rebalancing.
        from imblearn.over sampling import SMOTE
        # Standard Python libs.
        import os
        import glob
        import csv
        import xml.etree.ElementTree as et
        from dataclasses import dataclass
        # Pillow.
        import PIL as pil
        from PIL import ImageDraw
        # Tensorflow
        #import tensorflow as tf
        #from tensorflow.keras.preprocessing.image import ImageDataGenerator
        #from tensorflow.keras.datasets import fashion mnist
        #from tensorflow.keras.layers import Dense
        #from tensorflow.keras.optimizers import Adam
        #from tensorflow.keras.layers import Conv2D, Flatten, Dense, AveragePooling2D, GlobalAveragePooling2D,Dropout
        #from tensorflow.keras.applications.resnet import ResNet50
        #from tensorflow.keras.models import Sequential
        #from tensorflow.keras.preprocessing.image import ImageDataGenerator
        # For images
        from skimage.color import rgb2gray
        import cv2
        from scipy import ndimage
```

```
In [2]: # Utility functions.
        def image_load(filename):
            loader = pil.Image.open(filename);
            ret = loader.copy();
            loader.close();
            return ret;
        def image_resize(image, new_width):
            new height = int(new width * (image.height / image.width));
            return image.resize((new_width, new_height), pil.Image.Resampling.LANCZOS);
        def trimap_to_mask(trimap, include_border = True):
            trimap_data = trimap.getdata();
            mask_data = np.zeros((trimap.height, trimap.width), dtype=np.uint8);
            for x in range(0, trimap.width):
                for y in range(0, trimap.height):
                    idx = x + (y * trimap.width);
                    tri = trimap_data[idx];
                    if (tri == 1 or (include_border == True and tri == 3)):
                        mask_data[y, x] = 255;
                    else:
                        mask data[y, x] = 0;
            mask = pil.Image.fromarray(mask data);
            return mask;
```

```
In [3]: # Class definitions.
        @dataclass
        class Point:
            x: int;
            y: int;
        @dataclass
        class BoundingBox:
            ll: Point;
            lr: Point;
            ul: Point;
            ur: Point;
        class CatBreedSample:
            def __init__(self, label, image_file, mask_file = None, bb_file = None):
                self.label = label;
                self.image file = image file;
                self.mask_file = mask_file;
                self.bb file = bb file;
                # Load the image
                self.image = image_load(self.image_file);
                # Composite if a mask is available.
                if (self.mask file):
                    self.mask = trimap to mask(image load(self.mask file), False);
                    background = pil.Image.new("RGB", self.mask.size, 0);
                    self.masked image = pil.Image.composite(self.image, background, self.mask);
                else:
                    self.mask = None;
                    self.masked image = None;
                # Get a bounding box.
                if (self.bb file):
                    tree = et.parse(self.bb file);
                     root = tree.getroot();
                    xmin = int(root.findall("./object/bndbox/xmin")[0].text);
                    xmax = int(root.findall("./object/bndbox/xmax")[0].text);
                    ymin = int(root.findall("./object/bndbox/ymin")[0].text);
                    ymax = int(root.findall("./object/bndbox/ymax")[0].text);
                    self.bb = BoundingBox(0, 0, 0, 0);
                    self.bb.ll = Point(xmin, ymin);
                    self.bb.lr = Point(xmax, ymin);
                    self.bb.ul = Point(xmin, ymax);
                    self.bb.ur = Point(xmax, ymax);
                    if (self.masked image):
                         self.bounded image = self.masked image.crop((xmin, ymin, xmax, ymax));
                    else:
                         self.bounded image = self.image.crop((xmin, ymin, xmax, ymax));
                else:
                    self.bb = None;
                    self.bounded image = None;
                self.image = image resize(self.image, 256);
                self.masked_image = image_resize(self.masked_image, 256);
                self.bounded_image = image_resize(self.bounded_image, 256);
            def display(self):
                print("Image:", self.image file);
                print("Label:", self.label);
                display(self.image);
```

```
if (self.mask):
        display(self.masked image);
   if (self.bb):
        display(self.bounded image);
def features(self):
    fv = [];
   # Determine eye positions.
   # Determine ear positions.
   # We just estimate by starting at corners of the region segmentation and working
   # our way towards the center. The first non-black pixel is our x/y.
    region segmentation = self.segmentation regionbased();
   #Left
   left ear = (0, 0);
    #print(len(region_segmentation), len(region_segmentation[0]))
    #print(seg data[0], seg data[1], seg data[2], seg data[3]);
    for y in range(0, int(len(region segmentation) / 2)):
        for x in range(0, int(len(region segmentation[0]) / 2)):
            \#print(region\_segmentation[y, x], end = "")
            if (region_segmentation[x, y] != 0):
                #print(x, y, "=", region segmentation[y, x]);
                left ear = (y, x);
                break:
        if (left ear != (0, 0)):
            break;
   #print(left ear);
    #Right
    right ear = (0, 0);
    for y in range(0, int(len(region segmentation) / 2)):
        for x in reversed(range(int(\overline{len}(region\ segmentation[0])\ /\ 2), int(\overline{len}(region\ segmentation[0])))):
            #print(region_segmentation[y, x], end = " ")
            if (region segmentation[y, x] != 0):
                \#region\ segmentation[y, x] = 0.5;
                right ear = (x, y);
                break;
        if (right ear !=(0, 0)):
            break;
    #print(right ear);
    #plt.imshow(region segmentation, cmap='gray');
    #draw = ImageDraw.Draw(self.bounded image);
   #ellipse = ((left ear[0] - 2, left ear[1] - 2, left ear[0] + 2, left ear[1] + 2));
   #draw.ellipse(ellipse, fill = "red", outline = "red");
   \#ellipse = ((right\ ear[0] - 2,\ right\ ear[1] - 2,\ right\ ear[0] + 2,\ right\ ear[1] + 2));
   #draw.ellipse(ellipse, fill = "red", outline = "red");
    fv.append(right ear[0] - left ear[0]);
    fv.append(right ear[1] - left ear[1]);
   # Bin the colors that make up this image.
    #region segmentation = self.segmentation colorclustering();
    #unique, counts = np.unique(region segmentation, return counts = True);
    #unique colors = dict(zip(unique, counts));
    #counts = counts[1:];
    # . counts = np.unique(counts, return counts = True):
```

```
cv2image = np.array(self.masked image);
    channels = cv2.split(cv2image);
    for i in range(0, 3):
        hist = cv2.calcHist([channels[i]], [0], None, [5], [0, 255]);
        for val in hist:
           fv.append(int(val));
        #print(hist);
    #plt.figure();
    #plt.plot(hist);
    #plt.xlim([0, 256])
    #plt.ylim([0, 1200])
    #region segmentation
    #print(unique);
    return fv;
def segmentation regionbased(self):
    gray = rgb2gray(np.array(self.bounded_image));
    #plt.imshow(gray, cmap = 'gray')
    gray_r = gray.reshape(gray.shape[0]*gray.shape[1])
    for i in range(gray_r.shape[0]):
        if gray r[i] > gray r.mean():
           gray r[i] = 1
        else:
           gray_r[i] = 0
    gray = gray r.reshape(gray.shape[0],gray.shape[1])
    #plt.figure();
    #plt.imshow(gray, cmap='gray')
    return gray;
    # The darker region (black) represents the background and the brighter (white) region is the foreground. We can define multiple thresholds as well to detect multiple ob
    # gray r = gray.reshape(gray.shape[0]*gray.shape[1])
    #for i in range(gray_r.shape[0]):
        #if gray_r[i] > gray_r.mean():
           \#gray r[i] = 3
        \#elif\ gray_r[i] > 0.5:
           \#gray_r[i] = 2
        #elif gray r[i] > 0.25:
           \#gray_r[i] = 1
        #else:
           \#gray \ r[i] = 0
    #gray = gray_r.reshape(gray.shape[0],gray.shape[1])
    #plt.imshow(gray, cmap='gray')
def segmentation_edgebased(self):
    gray = rgb2gray(np.array(self.bounded image));
    #plt.figure();
    #plt.imshow(gray, cmap='gray');
    # defining the sobel filters
    # [
    # [1 2 1]
    # [ 0 0 0]
    # [-1 -2 -1]
```

```
# is a kernel for detecting horizontal edges
   # [
   # [-1 0 1]
   # [-2 0 2]
   # [-1 0 1]
   # 1
   # is a kernel for detecting vertical edges
    sobel horizontal = np.array([np.array([1, 2, 1]), np.array([0, 0, 0]), np.array([-1, -2, -1])])
    print('Kernel for detecting horizontal edges:\n', sobel horizontal)
    sobel\_vertical = np.array([np.array([-1, \ 0, \ 1]), \ np.array([-2, \ 0, \ 2]), \ np.array([-1, \ 0, \ 1])])
    print('Kernel for detecting vertical edges:\n', sobel vertical)
    out h = ndimage.convolve(gray, sobel horizontal, mode='reflect')
    out v = ndimage.convolve(gray, sobel vertical, mode='reflect')
    # here mode determines how the input array is extended when the filter overlaps a border.
    plt.figure();
    plt.imshow(out h, cmap='gray')
    plt.imshow(out v, cmap='gray')
    # Here, we are able to identify the horizontal as well as the vertical edges. There is one more type of filter that can detect both horizontal and vertical edges at the
   # [
   # [1 1 1]
    # [1 -8 1]
    # [1 1 1]
    # ]
    kernel\ laplace = np.array([np.array([1, 1, 1]), np.array([1, -8, 1]), np.array([1, 1, 1])])
    print("Laplacian kernel:\n", kernel laplace)
    out l = ndimage.convolve(gray, kernel laplace, mode='reflect')
    plt.figure();
    plt.imshow(out l, cmap='gray')
def segmentation colorclustering(self):
    # According to wikipedia the R, G, and B components of an object's color in a digital image are all correlated with the amount of light hitting the object,
    # and therefore with each other, image descriptions in terms of those components make object discrimination difficult.
   # Descriptions in terms of hue/lightness/chroma or hue/lightness/saturation are often more relevant. So, we need to convert our image from RGB Colours Space to HSV to we
    cv2img = np.array(self.masked image);
    vectorized = np.float32(cv2img.reshape((-1,3)))
   vectorized.shape
    criteria = (cv2.TERM CRITERIA EPS + cv2.TERM CRITERIA MAX ITER, 20, 1.0)
   K = 8
    attempts=10
    ret, label, center = cv2.kmeans(vectorized, K, None, criteria, attempts, cv2.KMEANS PP CENTERS)
    center = np.uint8(center)
    res = center[label.flatten()]
    result image = res.reshape((cv2img.shape))
    # result image is the output result. Lets see how the image looks after k-means clustering
    #figure size = 15
    #plt.figure(figsize=(figure_size, figure_size))
    #plt.subplot(2,3,1),plt.imshow(cv2img)
    #plt.title('Original Image'), plt.xticks([]), plt.yticks([])
    #plt.subplot(2,3,2),plt.imshow(result image)
    #plt.title('Segmented Image when K = %i' % K), plt.xticks([]), plt.yticks([])
    #plt.show()
    return result image;
```

```
def haarcascade classifier(self):
                cv2img = np.array(self.image);
                cv2gray = cv2.cvtColor(cv2img, cv2.COLOR RGB2GRAY);
                cascade = cv2.CascadeClassifier("pretrained/haarcascade frontalcatface.xml");
                bounding rects = cascade.detectMultiScale(cv2gray, scaleFactor = 1.3, minNeighbors = 1, minSize = (25, 25));
                print(bounding rects);
                for (x, y, w, h) in bounding rects:
                    cv2.rectangle(cv2gray, (x, y), (x + w, y + h), (0, 0, 255), 2);
                plt.figure();
                plt.imshow(cv2gray, cmap='gray');
            def contours(self):
                #region segmentation = self.segmentation regionbased();
                region segmentation = np.array(self.bounded image);
                region_segmentation = cv2.cvtColor(region_segmentation, cv2.COLOR RGB2GRAY)
                ret, threshold = cv2.threshold(region segmentation, 16, 255, cv2.THRESH BINARY);
                image, contours, hierarchy = cv2.findContours(threshold, cv2.RETR TREE, cv2.CHAIN APPROX SIMPLE);
                cv2.drawContours(threshold, contours, -1, (0, 255, 0), 3);
                approx = cv2.approxPolyDP(contours[1], 0.01 * cv2.arcLength(contours[1], True), True)
                print(approx);
                plt.figure();
                plt.imshow(threshold);
                #i = 0;
                #for contour in contours:
                # # here we are ignoring first counter because
                    # findcontour function detects whole image as shape
                    if i == 0:
                        i = 1
                         continue
                    # cv2.approxPloyDP() function to approximate the shape
                     #approx = cv2.approxPolyDP(contour, 0.01 * cv2.arcLength(contour, True), True)
                     print(contour)
                     # using drawContours() function
                     cv2.drawContours(region segmentation, [contour], 0, (0, 0, 255), 5)
                #plt.figure();
                #plt.imshow(region segmentation);
In [4]: #test sample = CatBreedSample(
```

```
# "Abyssinian",
# "/home/cva/catifier/training_data/Abyssinian_115.jpg",
# "/home/cva/catifier/training_data/Abyssinian_115_mask.png",
# "/home/cva/catifier/training_data/Abyssinian_115_mask.png",
# "/home/cva/catifier/training_data/Abyssinian_115_bb.xml"
#)

#test_sample.segmentation_regionbased();
#test_sample.segmentation_edgebased();
#test_sample.segmentation_colorclustering();
#test_sample.haarcascade_classifier();
#test_sample.contours();

#fv = test_sample.features();
#test_sample.display();
```

```
In [5]: # Load everything.
         def load samples(samples dir):
             class dirs = glob.glob(samples dir + "/*")
             samples = [];
             for class dir in class dirs:
                 label = os.path.basename(class dir);
                 class_dir_glob = glob.glob(class_dir + "/*.jpg");
                 print("Reading files for label '" + label + "'");
                 for sample_image in class_dir_glob:
                     basename = os.path.splitext(sample image)[0];
                     mask_file = basename + "_mask.png";
bb_file = basename + "_bb.xml";
                     if (not os.path.exists(mask_file)):
                         mask file = None;
                     if (not os.path.exists(bb_file)):
                         continue;
                         #bb file = None;
                     print(
                         len(samples),
                         ":",
                         os.path.basename(sample image),
                         "\t(mask:",
                         (mask file != None),
                         "| bb:",
                         (bb_file != None),
                         ")"
                     );
                     sample = CatBreedSample(label, sample image, mask file, bb file);
                     samples.append(sample);
             return samples;
        PWD = os.getcwd();
        TRAINING_DATA = PWD + "/training_data";
         samples = load samples(TRAINING DATA);
        (... trimmed for brevity ...)
        Reading files for label 'Abyssinian'
        Reading files for label 'Sphynx'
        Reading files for label 'Egyptian Mau'
        Reading files for label 'Persian'
        Reading files for label 'Siamese'
        Reading files for label 'Birman'
        Reading files for label 'Bombay'
        Reading files for label 'Russian_Blue'
```

Reading files for label 'Bengal'
Reading files for label 'Maine_Coon'
Reading files for label 'Ragdoll'

Reading files for label 'British Shorthair'

```
In [6]: #sample id = 120;
        #samples[sample id].display();
        #samples[sample id].segmentation regionbased();
        #samples[sample_id].segmentation_edgebased();
        #samples[sample id].segmentation colorclustering();
In [8]: # Try the model.
        fv = [];
        labels = [];
        for sample in samples:
            print("Extract features from:", sample.image_file);
            fv.append(sample.features());
            labels.append(sample.label);
        # Rebalance.
        oversampler = SMOTE();
        (fv, labels) = oversampler.fit resample(fv, labels);
        # Scale.
        scaler = RobustScaler():
        fv = scaler.fit transform(fv);
        # Split the data.
        x_train, x_test, y_train, y_test = train_test_split(fv, labels, test_size = 0.30, random_state = 64);
        # Train.
        dt = RandomForestClassifier();
        dt.fit(x_train, y_train);
        # Testing the model.
        cv scores = cross val score(dt, x train, y train, cv = 10);
        print('Average Cross Validation Score from Training:', cv scores.mean(), sep = '\n', end = '\n\n\n');
        y pred = dt.predict(x test);
        cm = confusion matrix(y test, y pred);
        cr = classification report(y test, y pred);
        print('Confusion Matrix:', cm, sep = '\n', end = '\n\n\n');
        print('Missing classifications (if any):', set(y test) - set(y pred));
        print('Test Statistics:', cr, sep = '\n', end = '\n\n\n');
        print('Testing Accuracy:', accuracy_score(y_test, y_pred));
```

(... omit loading messages for brevity ...)
Average Cross Validation Score from Training:
0.3547619047619047

(100	nfı	ısio	n ľ	1atı	rix:							
	[[:	10	6	1	1	1	1	3	1	0	2	1	3]
	[4	10	2	3	0	0	5	2	0	0	1	0]
	[0	1	6	1	2	4	1	2	6	1	5	1]
	[0	0	0	24	1	0	1	0	0	1	0	0]
	[0	2	1	1	15	1	5	3	3	4	0	0]
	[1	1	2	0	0	7	0	0	0	5	7	2]
	[5	7	2	1	0	0	4	4	2	1	3	1]
	[4	1	0	0	1	0	1	9	4	2	1	3]
	[3	0	4	0	1	4	0	5	6	0	5	3]
	[1	1	0	2	8	4	1	1	1	11	0	0]
	[0	5	7	1	0	3	7	1	3	0	5	3]
	1:	10	1	3	0	2	3	2	1	1	1	2	811

Missing classifications (if any): set()
Test Statistics:

	precision	recall	f1-score	support
Abyssinian	0.26	0.33	0.29	30
Bengal	0.29	0.37	0.32	27
Birman	0.21	0.20	0.21	30
Bombay	0.71	0.89	0.79	27
British_Shorthair	0.48	0.43	0.45	35
Egyptian Mau	0.26	0.28	0.27	25
Maine_Coon	0.13	0.13	0.13	30
Persian	0.31	0.35	0.33	26
Ragdoll	0.23	0.19	0.21	31
Russian_Blue	0.39	0.37	0.38	30
Siamese	0.17	0.14	0.15	35
Sphynx	0.33	0.24	0.28	34
accuracy			0.32	360
macro avg	0.31	0.33	0.32	360
weighted avg	0.31	0.32	0.31	360

Testing Accuracy: 0.3194444444444444

In []: