## ccuds4s7s

## February 14, 2024

#### $0.1 \ 2a >$

```
[9]: import matplotlib.pyplot as plt
import os
import cv2
from pathlib import Path
from skimage.color import rgb2gray
from skimage import io, exposure
import xml.etree.ElementTree as ET
from PIL import Image

myBreeds = [
    'n02099712-Labrador_retriever',
    'n02110185-Siberian_husky',
    'n02113799-standard_poodle',
    'n02113186-Cardigan'
]
```

```
[10]: class AllRequiredPaths:
    myBreeds = [
        'n02099712-Labrador_retriever',
        'n02110185-Siberian_husky',
        'n02113799-standard_poodle',
        'n02113186-Cardigan'
    ]

    root_dir = ""
    imgs_folder = ""
    annotation_folder = ""
    imgs_folder_sub_folders = [f"images/{name}" for name in myBreeds]

    def __init__(self):
        self.root_dir = os.getcwd()
        self.imgs_folder = os.path.join(self.root_dir, "images")
        self.annotation_folder = os.path.join(self.root_dir, "Annotation")
```

```
class CropAndResizeImages:
   def __init__(self):
        self.all_paths = AllRequiredPaths()
   def _get_bbox(self, annot):
       xml = annot
       tree = ET.parse(xml)
       root = tree.getroot()
        objects = root.findall('object')
       bbox = []
       for o in objects:
            bndbox = o.find("bndbox")
            xmin = int(bndbox.find("xmin").text)
            ymin = int(bndbox.find("ymin").text)
            xmax = int(bndbox.find("xmax").text)
            ymax = int(bndbox.find('ymax').text)
            bbox.append((xmin, ymin, xmax, ymax))
        return bbox
   def _get_images_from_classes(self, annot):
        image_name = f"{os.path.basename(annot)}.jpg"
       breed_name = os.path.basename(os.path.dirname(annot))
       return os.path.join(self.all_paths.imgs_folder, breed_name, image_name)
   def crop save images(self, source path, boxes):
        original_image = Image.open(source_path)
        for index, bounding box in enumerate(boxes):
            cropped_img = original_image.crop(bounding_box).resize((128, 128),_
 →Image.LANCZOS)
            original_filename = os.path.basename(source_path)
            new file name = f"{os.path.
 ⇒splitext(original_filename)[0]}_section_{index}{os.path.
 ⇔splitext(original_filename)[1]}"
            save_path = os.path.join("CroppedImagesFolder", new_file_name)
            Path(save_path).parent.mkdir(parents=True, exist_ok=True)
            cropped_img.convert('RGB').save(save_path)
   def process(self, myBreeds):
        annotation_files = []
        for myBreed in myBreeds:
            for file_name in os.listdir(os.path.join(self.all_paths.
 →annotation_folder, myBreed)):
                if not os.path.isdir(os.path.join(self.all_paths.
 →annotation_folder, myBreed, file_name)):
```

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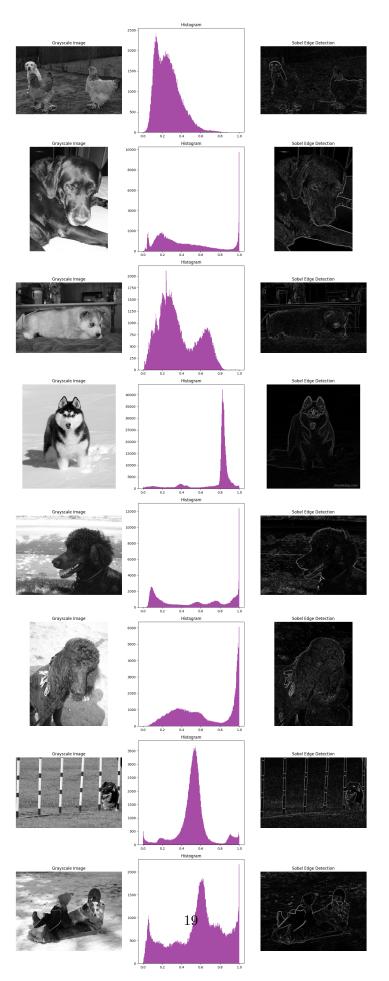
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     0.2 	ext{ 2b} >
 [4]: import numpy as np
      from skimage import io, color, filters, exposure
      import matplotlib.pyplot as plt
[12]: class ImageProcess:
          def __init__(self):
              self.all_paths = AllRequiredPaths()
          def _fetch_two_images_from_each_class(self, class_dir):
              img_files = []
              for file in os.listdir(class_dir):
                  if file.endswith(".jpg"):
                      img_files.append(file)
              images = []
              for img in img_files[3:5]:
                  images.append(os.path.join(class_dir, img))
              return images
          def _get_and_process_img(self):
```

```
all_images = []
       for img_dir in self.all_paths.imgs_folder_sub_folders:
           for img in self._fetch_two_images_from_each_class(img_dir):
               all_images.append(img)
      original_imgs = [io.imread(path) for path in all_images]
      gray_scale_imgs = [color.rgb2gray(img) for img in original_imgs]
      edge_imgs = [filters.sobel(img) for img in gray_scale_imgs]
      return original_imgs, gray_scale_imgs, edge_imgs
  def show results(self):
      original_imgs, grayscale_imgs, edge_imgs = self._get_and_process_img()
      total_imgs = len(grayscale_imgs)
      fig, axes = plt.subplots(total_imgs, 3, figsize = (15, 5 * total_imgs))
      for i in range(total_imgs):
           # show grayscale images
           axGray = axes[i, 0] if total_imgs > 1 else axes[0]
           axGray.imshow(grayscale_imgs[i], cmap="gray")
           axGray.set_title("Grayscale Image")
           axGray.axis("off")
           # show histrogram graph for each images
           axHistogram = axes[i, 1] if total_imgs > 1 else axes[1]
           axHistogram.hist(grayscale_imgs[i].ravel(), bins=256, range=[0,__
→1],color='purple', alpha=0.7)
           axHistogram.set_title("Histogram")
           # show edge detection results
           axEdge = axes[i, 2] if total_imgs > 1 else axes[2]
           axEdge.imshow(edge imgs[i], cmap="gray")
           axEdge.set_title("Sobel Edge Detection")
           axEdge.axis("off")
      plt.tight_layout()
      plt.show()
```

```
[13]: image_proecess = ImageProcess()
  image_proecess.show_results()
```



### $0.3 \ 2c >$

```
[7]: import os
from skimage import io, color, filters, exposure
import numpy as np
import matplotlib.pyplot as plt
import matplotlib.patches as patches
from PIL import Image
```

```
[8]: class EdgeHistogramAnalysis:
         def __init__(self, all_images):
             self.all_paths = AllRequiredPaths()
             self.all_images = all_images
         def _angle(self, dx, dy):
             return np.mod(np.arctan2(dx, dy), np.pi)
         def show result(self):
             imgs_subdir = self.all_paths.imgs_folder
             for catg, img_name in self.all_images:
                 img_path = os.path.join(imgs_subdir, catg, img_name)
                 img = io.imread(img_path)
                 # grayscale image
                 grayscale_img = color.rgb2gray(img)
                 # edge orientation angles
                 dx = filters.sobel_h(grayscale_img)
                 dy = filters.sobel_v(grayscale_img)
                 ang_img = self._angle(dx,dy)
                 hist, hist centers = exposure histogram(ang img, nbins=36,

¬source_range="dtype")
                 plt.figure(figsize=(10,4))
                 plt.subplot(1,2,1)
                 plt.imshow(grayscale_img, cmap="gray")
                 plt.title(f"{catg}")
                 plt.axis("off")
                 plt.subplot(1,2,2)
                 plt.bar(hist_centers, hist, width=0.1, align="center", color =_

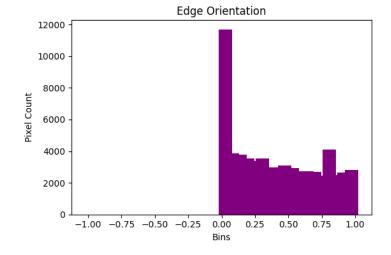
¬"purple")
```

```
plt.xlabel("Bins")
plt.ylabel("Pixel Count")
plt.title("Edge Orientation")

plt.tight_layout()
plt.show()
```

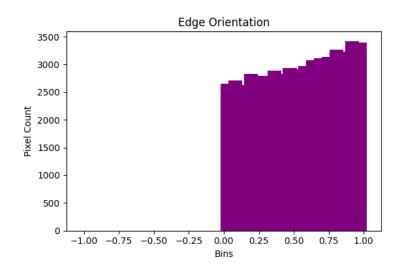
n02099712-Labrador\_retriever





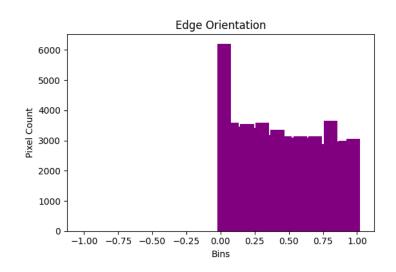
n02110185-Siberian\_husky



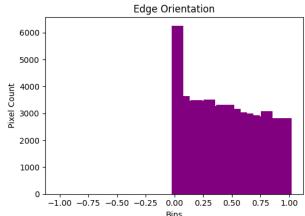


n02113799-standard\_poodle









### 0.4 2d >

```
[17]: class HistorgramComparison:
          def __init__(self, selected_classes):
              self.all_paths = AllRequiredPaths()
              self.selected_classes = selected_classes
          def _angle(self, dx, dy):
              return np.mod(np.arctan2(dx,dy), np.pi)
          def _edge_histogram(self, image):
              gray_img = color.rgb2gray(image)
              angles = self._angle(filters.sobel_h(gray_img), filters.
       →sobel_v(gray_img))
              hist, _ = np.histogram(angles, bins=36, range=(-np.pi, np.pi))
              return hist
          def _process_images(self):
              edge_hist = {}
              for catg, img_name in self.selected_classes:
                  img_path = os.path.join(self.all_paths.imgs_folder, catg, img_name)
                  img = io.imread(img_path)
                  hist = self._edge_histogram(img)
                  edge_hist[img_name] = hist
              return edge_hist
          def compare_hist(self, title):
              edge_hist = self._process_images()
              histos = list(edge_hist.values())
              img_names = list(edge_hist.keys())
              euclidean_dist = euclidean_distances(histos)
              manhattan_dist = manhattan_distances(histos)
              cosine_dist = cosine_distances(histos)
              print(title)
              for i in range(len(img_names)):
                  for j in range(i + 1, len(img_names)):
```

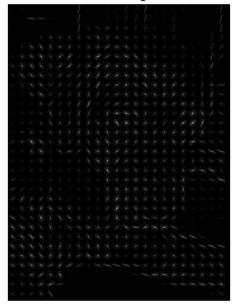
```
print(f"Euclidean: {euclidean_dist[i][j]}")
                      print(f"Manhattan: {manhattan_dist[i][j]}")
                      print(f"Cosine: {cosine_dist[i][j]}")
      histogram_compare1 = HistorgramComparison([
          ("n02110185-Siberian_husky", "n02110185_10047.jpg"),
          ("n02110185-Siberian_husky", "n02110185_10844.jpg")
      1)
      histogram_compare1.compare_hist("Comparison between two images of same class")
      print("")
      histogram_compare2 = HistorgramComparison([
          ("n02110185-Siberian_husky", "n02110185_10047.jpg"),
          ("n02113186-Cardigan", "n02113186_1030.jpg")
      ])
      histogram compare2.compare hist("Comparison between two images of different ⊔
       ⇔class")
     Comparison between two images of same class
     Euclidean: 4678.233213511272
     Manhattan: 16230.0
     Cosine: 0.005141347337101587
     Comparison between two images of different class
     Euclidean: 10356.438383923307
     Manhattan: 35606.0
     Cosine: 0.024073880505602663
     0.5 \ 2e >
[18]: import matplotlib.pyplot as plt
      from skimage import io, exposure
      from skimage.transform import resize
      from skimage.feature import hog
[19]: class HOGAnalysis:
          def __init__(self, images):
              self.all_path = AllRequiredPaths()
              self.images = images
          def _fetchAndPlot(self, img_path, orientations = 8, pixels_per_cell =_
       \hookrightarrow (16,16), cells_per_block = (1,1)):
              img = io.imread(img path)
```

```
fd, hog_img = hog(img, orientations=orientations,__
 ⇔pixels_per_cell=pixels_per_cell, cells_per_block=cells_per_block,_
 ⇔visualize=True, channel_axis=-1)
        hog_img_rescaled = exposure.rescale_intensity(hog_img,_
 \rightarrowout range=(0,225))
        plt.figure(figsize=(10,5))
        plt.subplot(1,2,1)
        plt.imshow(img, cmap=plt.cm.gary if img.ndim == 2 else None)
        plt.title("Original Image")
        plt.axis('off')
        plt.subplot(1,2,2)
        plt.imshow(hog_img_rescaled, cmap="gray")
        plt.title("HOG Image")
        plt.axis('off')
        plt.show()
    def show_result(self):
        for catg, img_name in self.images:
            img_path = f"images/{catg}/{img_name}"
            self._fetchAndPlot(img_path)
images = [
    ("n02099712-Labrador_retriever", "n02099712_129.jpg"),
    ("n02110185-Siberian_husky", "n02110185_10047.jpg"),
    ("n02113799-standard_poodle", "n02113799_1155.jpg"),
    ("n02113186-Cardigan", "n02113186_1030.jpg")
1
hogAnalysis = HOGAnalysis(images)
hogAnalysis.show_result()
```

Original Image



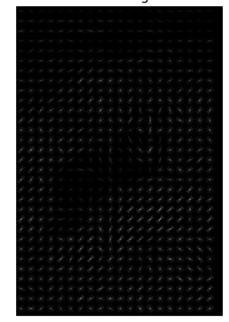
**HOG Image** 



Original Image



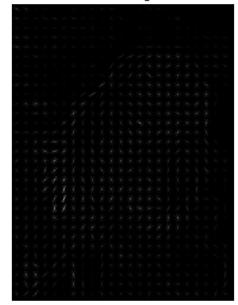
HOG Image



Original Image



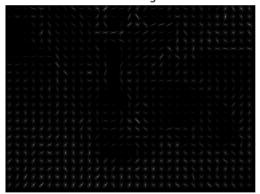
**HOG Image** 



Original Image



**HOG Image** 



# 0.6 2f>

```
[1]: import os
  import numpy as np
  import matplotlib.pyplot as plt
  from skimage import io, color, filters
  from sklearn.decomposition import PCA
  from sklearn.preprocessing import StandardScaler
```

```
[20]: class PCAAnalysis:
          def __init__(self):
              self.all_paths = AllRequiredPaths()
          def _angle(self, dx, dy):
              return np.mod(np.arctan2(dx,dy), np.pi)
          def _edge_histogram(self, image):
              gray_img = color.rgb2gray(image)
              angles = self._angle(filters.sobel_h(gray_img), filters.
       →sobel_v(gray_img))
              hist, _ = np.histogram(angles, bins=36, range=(-np.pi, np.pi))
              return hist
          def fetch_images(self, folder_name):
              allImage = []
              for img name in os.listdir(folder name):
                  if img_name.lower().endswith(".jpg"):
                      allImage.append(os.path.join(folder_name, img_name))
              return allImage
          def perform_pca_and_show_result(self, class1, class2):
              hists = []
              labels = []
              classes = class1 + class2
              for img_path in classes:
                  img = io.imread(img_path)
                  hist = self._edge_histogram(img)
                  hists.append(hist)
                  labels.append("Class 1" if img_path in class1 else "Class 2")
              histogram = np.array(hists)
              std_scalar = StandardScaler()
              hist_std = std_scalar.fit_transform(histogram)
              pca = PCA(n_components=2)
              hist_pca = pca.fit_transform(hist_std)
              pca1 = hist_pca[:len(class1)]
              pca2 = hist_pca[:len(class2)]
              plt.scatter(pca1[:, 0], pca1[:, 1], color = "red", label = "Class-1")
              plt.scatter(pca2[:, 0], pca2[:, 1], color = "blue", label = "Class-2")
              plt.xlabel("Principal Component 1")
```

```
plt.ylabel("Principal Component 2")
    plt.legend(["Class 1", "Class 2"])
    plt.title("PCA of Image")
    plt.show()

pca_analysis = PCAAnalysis()
pca1_folder = pca_analysis.fetch_images("images/n02110185-Siberian_husky")
pca2_folder = pca_analysis.fetch_images("images/n02113186-Cardigan")

pca_analysis.perform_pca_and_show_result(pca1_folder, pca2_folder)
```

# PCA of Image

