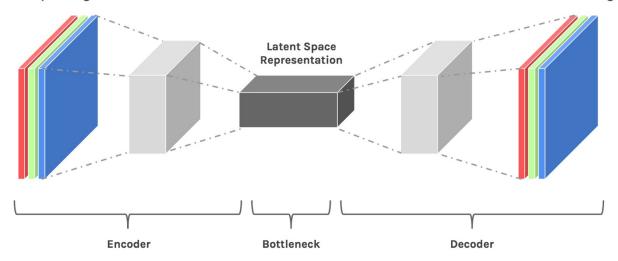
```
from google.colab import files
files.upload()
! chmod 600 kaggle.json
! kaggle datasets download -d theaayushbajaj/cbir-dataset --unzip
کتابخانه اصلی مورد نیاز برای اجرای پروژه #
!pip install barbar torchsummary
!pip install barbar
from sklearn.model selection import train test split
from torch.utils.data.dataset import Dataset
from torch.utils.data import DataLoader
from torchvision import transforms
from torchsummary import summary
from pathlib import Path
from barbar import Bar
from tqdm import tqdm
from PIL import Image
from torch import nn
%matplotlib inline
import pandas as pd # پردازش دادهه های درون فایل # CSV I/O (e.g.
pd.read csv)
برای استفاده از جبر خطی # import numpy as np
import matplotlib.pyplot as plt
import torchvision
import pickle
import time
import copy
import scipy
import torch
import cv2
import os
import gc
RANDOMSTATE = 0
cpu استفاده میکنیم اگر نبود از cuda برای بهینه سازی سرعت اموزش مدل از #
device = torch.device('cuda:0' if torch.cuda.is available() else
'cpu')
print(device)
cpu
# آماده سازی DataFrame
datasetPath = Path('/content/dataset/')
```

```
df = pd.DataFrame()
df['image'] = [f for f in os.listdir(datasetPath) if
os.path.isfile(os.path.join(datasetPath, f))]
df['image'] = '/content/dataset/' + df['image'].astype(str)
df.head()
class CBIRDataset(Dataset):
    def init (self, dataFrame):
        self.dataFrame = dataFrame
        self.transformations = transforms.Compose([
            transforms.ToTensor(),
            transforms.Normalize((0.5, 0.5, 0.5), (0.5, 0.5, 0.5))
        ])
    def getitem (self, key):
        if isinstance(key, slice):
            raise NotImplementedError('slicing is not supported')
        row = self.dataFrame.iloc[kev]
        image = self.transformations(Image.open(row['image']))
        return image
    def len (self):
        return len(self.dataFrame.index)
تابع میانی برای پردازش داده ها از کلاس بازیابی داده ها #
def prepare data(DF):
    trainDF, validateDF = train_test_split(DF, test size=0.15,
random state=RANDOMSTATE)
    train set = CBIRDataset(trainDF)
    validate set = CBIRDataset(validateDF)
    return train set, validate set
```

AutoEncoder ساختار سطح بالای یک

Input image Reconstructed image



```
class ConvAutoencoder(nn.Module):
    def __init__(self):
        super(ConvAutoencoder, self). init ()
        self.encoder = nn.Sequential(# in- (N,3,512,512)
            nn.Conv2d(in channels=3,
                       out channels=16,
                       kernel size=(3,3),
                       stride=3,
                       padding=1), # (32,16,171,171)
            nn.ReLU(True),
            nn.MaxPool2d(\frac{2}{2}, stride=\frac{2}{2}), # (N, 16, 85, 85)
            nn.Conv2d(in channels=16,
                       out channels=8,
                       kernel size=(3,3),
                       stride=2,
                       padding=1), # (N, 8, 43, 43)
            nn.ReLU(True),
            nn.MaxPool2d(2, stride=1) # (N, 8, 42, 42)
        self.decoder = nn.Sequential(
            nn.ConvTranspose2d(in channels = 8,
                                 out_channels=16,
                                 kernel size=(3,3),
                                 stride=2), # (N, 16, 85, 85)
            nn.ReLU(True),
            nn.ConvTranspose2d(in channels=16,
                                 out channels=8,
                                 kernel size=(5,5),
                                 stride=3,
```

```
padding=1), # (N, 8, 255, 255)
            nn.ReLU(True),
            nn.ConvTranspose2d(in channels=8,
                                 out channels=3,
                                 kernel size=(6,6),
                                 stride=2,
                                 padding=1), # (N,3,512,512)
            nn.Tanh()
        )
    def forward(self, x):
        x = self.encoder(x)
        x = self.decoder(x)
        return x
class ConvAutoencoder v2(nn.Module):
    def init (self):
        super(ConvAutoencoder v2, self). init ()
        self.encoder = nn.Sequential(# i\overline{n}- (N, \overline{3}, 512, 512)
            nn.Conv2d(in_channels=3,
                       out_channels=64,
                       kernel size=(3,3),
                       stride=1,
                       padding=1),
            nn.ReLU(True),
            nn.Conv2d(in channels=64,
                       out channels=64,
                       kernel size=(3,3),
                       stride=1,
                       padding=1),
            nn.ReLU(True),
            nn.MaxPool2d(2, stride=2),
            nn.Conv2d(in channels=64,
                       out channels=128,
                       kernel size=(3,3),
                       stride=2,
                       padding=1),
            nn.ReLU(True),
            nn.Conv2d(in channels=128,
                       out channels=128,
                       kernel size=(3,3),
                       stride=1,
                       padding=0),
            nn.ReLU(True),
            nn.MaxPool2d(2, stride=2),
            nn.Conv2d(in channels=128,
```

```
out channels=256,
              kernel size=(3,3),
              stride=2,
              padding=1),
    nn.ReLU(True),
    nn.Conv2d(in_channels=256,
              out channels=256,
              kernel size=(3,3),
              stride=1,
              padding=1),
    nn.ReLU(True),
    nn.Conv2d(in_channels=256,
              out channels=256,
              kernel size=(3,3),
              stride=1,
              padding=1),
    nn.ReLU(True),
    nn.MaxPool2d(2, stride=2)
self.decoder = nn.Sequential(
    nn.ConvTranspose2d(in_channels = 256,
                        out channels=256,
                        kernel size=(3,3),
                        stride=1,
                       padding=1),
    nn.ConvTranspose2d(in channels=256,
                        out channels=256,
                        kernel size=(3,3),
                        stride=1,
                        padding=1),
    nn.ReLU(True),
    nn.ConvTranspose2d(in channels=256,
                        out channels=128,
                        kernel size=(3,3),
                        stride=2,
                        padding=0),
    nn.ConvTranspose2d(in channels=128,
                        out channels=64,
                        kernel size=(3,3),
                        stride=2,
                        padding=1),
    nn.ReLU(True),
    nn.ConvTranspose2d(in channels=64,
                        out channels=32,
                        kernel size=(3,3),
                        stride=2,
```

```
padding=1),
            nn.ConvTranspose2d(in channels=32,
                                out channels=32,
                                kernel size=(3,3),
                                stride=2,
                                padding=1),
            nn.ReLU(True),
            nn.ConvTranspose2d(in channels=32,
                                out channels=3,
                                kernel_size=(4,4),
                                stride=2,
                                padding=2),
            nn.Tanh()
        )
    def forward(self, x):
        x = self.encoder(x)
        x = self.decoder(x)
        return x
summary(ConvAutoencoder v2().to(device),(3,512,512))
```

## Training Function

```
def load ckpt(checkpoint fpath, model, optimizer):
    # load check point
    checkpoint = torch.load(checkpoint fpath)
    # initialize state dict from checkpoint to model
    model.load state dict(checkpoint['model state dict'])
    # initialize optimizer from checkpoint to optimizer
    optimizer.load state dict(checkpoint['optimizer state dict'])
    # initialize valid loss min from checkpoint to valid loss min
    #valid loss min = checkpoint['valid loss min']
    # return model, optimizer, epoch value, min validation loss
    return model, optimizer, checkpoint['epoch']
def save checkpoint(state, filename):
    """Save checkpoint if a new best is achieved"""
    print ("=> Saving a new best")
    torch.save(state, filename) # save checkpoint
def train model(model,
```

```
criterion,
            optimizer,
            #scheduler,
            num epochs):
since = time.time()
best_model_wts = copy.deepcopy(model.state_dict())
best loss = np.inf
for epoch in range(num epochs):
    print('Epoch {}/{}'.format(epoch, num_epochs))
    print('-' * 10)
    # Each epoch has a training and validation phase
    for phase in ['train', 'val']:
        if phase == 'train':
            model.train() # Set model to training mode
        else:
            model.eval() # Set model to evaluate mode
        running loss = 0.0
        # Iterate over data.
        for idx,inputs in enumerate(Bar(dataloaders[phase])):
            inputs = inputs.to(device)
            # zero the parameter gradients
            optimizer.zero grad()
            # forward
            # track history if only in train
            with torch.set grad enabled(phase == 'train'):
                outputs = \overline{model(inputs)}
                loss = criterion(outputs, inputs)
                # backward + optimize only if in training phase
                if phase == 'train':
                    loss.backward()
                    optimizer.step()
            # statistics
            running loss += loss.item() * inputs.size(0)
        #if phase == 'train':
            scheduler.step()
        epoch_loss = running_loss / dataset_sizes[phase]
        print('{} Loss: {:.4f}'.format(
            phase, epoch loss))
```

```
# deep copy the model
            if phase == 'val' and epoch loss < best loss:
                best loss = epoch loss
                best model wts = copy.deepcopy(model.state dict())
                save checkpoint(state={
                                     'epoch': epoch,
                                    'state dict': model.state dict(),
                                    'best_loss': best loss,
'optimizer state dict':optimizer.state dict()
                                },filename='ckpt epoch {}.pt'.format(e
poch))
        print()
    time elapsed = time.time() - since
    print('Training complete in {:.0f}m {:.0f}s'.format(
        time elapsed // 60, time elapsed % 60))
    print('Best val Loss: {:4f}'.format(best_loss))
    # load best model weights
    model.load_state dict(best model wts)
    return model, optimizer, epoch_loss
EPOCHS = 150
NUM BATCHES = 32
RETRAIN = False
train set, validate set = prepare data(DF=df)
dataloaders = {'train': DataLoader(train set, batch size=NUM BATCHES,
shuffle=True, num workers=1) ,
                'val':DataLoader(validate set, batch size=NUM BATCHES,
num workers=1)
                }
dataset sizes = {'train': len(train set), 'val':len(validate set)}
model = ConvAutoencoder v2().to(device)
criterion = nn.MSELoss()
# Observe that all parameters are being optimized
optimizer = torch.optim.Adam(model.parameters(), lr=3e-4)
# Decay LR by a factor of 0.1 every 7 epochs
#exp lr scheduler = lr scheduler.StepLR(optimizer, step size=7,
qamma=0.1
# If re-training is required:
# Load the old model
if RETRAIN == True:
```

```
# load the saved checkpoint
    model, optimizer, start epoch =
load ckpt('/content/conv autoencoder.pt', model, optimizer)
    print('Checkpoint Loaded')
from google.colab import drive
drive.mount('/gdrive')
model, optimizer, loss = train model(model=model,
                    criterion=criterion,
                    optimizer=optimizer,
                    #scheduler=exp lr scheduler,
                    num epochs=EPOCHS)
model.save('modelh5', save format="h5")
# Save the Trained Model
torch.save({
            'epoch': EPOCHS,
            'model state dict': model.state dict(),
            'optimizer state dict': optimizer.state dict(),
            'loss': loss.
            }, 'conv autoencoderv2 200ep.pt')
```

#### Inference

1. Indexing

```
transformations = transforms.Compose([
            transforms.ToTensor(),
            transforms.Normalize((0.5, 0.5, 0.5), (0.5, 0.5, 0.5))
        ])
# Load Model in Evaluation phase
model = ConvAutoencoder v2().to(device)
model.load state dict(torch.load('/content/conv autoencoderv2 200ep.pt
', map location=device)['model state dict'], strict=False)
model.eval()
def get latent features(images, transformations):
    latent features = np.zeros((4738, 256, 16, 16))
    \#latent\ features = np.zeros((4738,8,42,42))
    for i,image in enumerate(tqdm(images)):
        tensor = transformations(Image.open(image)).to(device)
        latent features[i] =
model.encoder(tensor.unsqueeze(0)).cpu().detach().numpy()
```

```
del tensor
   gc.collect()
   return latent_features

images = df.image.values
latent_features = get_latent_features(images, transformations)

indexes = list(range(0, 4738))
feature_dict = dict(zip(indexes, latent_features))
index_dict = {'indexes':indexes,'features':latent_features}

# write the data dictionary to disk
with open('features.pkl', "wb") as f:
   f.write(pickle.dumps(index_dict))
```

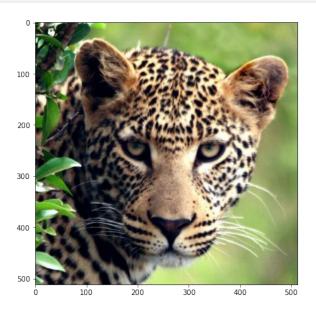
- 1. Image Retrieval¶ This will be approached with two ways as discussed in the start:
- Euclidean Search:
  - Identifying the Latent Features
  - Calculating the Euclidean Distance between them
  - Returning the closest N indexes (of images)
- Locality Sensitive Hashing
  - Create hashes of the feature vector from Encoder
  - Store it in a Hashing Table
  - Identify closest images based on hamming distance
- 2.1 Euclidean Search Method¶

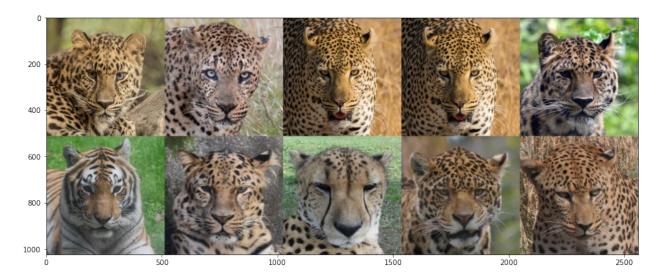
```
def euclidean(a, b):
    # compute and return the euclidean distance between two vectors
    return np.linalg.norm(a - b)
def cosine distance(a,b):
    return scipy.spatial.distance.cosine(a, b)
def perform search(queryFeatures, index, maxResults=64):
    results = []
    for i in range(0, len(index["features"])):
        # compute the euclidean distance between our query features
        # and the features for the current image in our index, then
        # update our results list with a 2-tuple consisting of the
        # computed distance and the index of the image
        d = euclidean(queryFeatures, index["features"][i])
        results.append((d, i))
    # sort the results and grab the top ones
    results = sorted(results)[:maxResults]
    # return the list of results
    return results
```

```
def build montages(image list, image shape, montage_shape):
    if len(image shape) != 2:
        raise Exception('image shape must be list or tuple of length 2
(rows, cols)')
    if len(montage shape) != 2:
        raise Exception('montage shape must be list or tuple of length
2 (rows, cols)')
    image montages = []
    # start with black canvas to draw images onto
    montage image = np.zeros(shape=(image shape[1] *
(montage shape [1]), image shape [0] * montage shape [0], [3]),
                          dtype=np.uint8)
    cursor pos = [0, 0]
    start new img = False
    for img in image_list:
        if type(img). module != np. name :
            raise Exception('input of type {} is not a valid numpy
array'.format(type(img)))
        start new img = False
        img = cv2.resize(img, image shape)
        # draw image to black canvas
        montage_image[cursor_pos[1]:cursor_pos[1] + image shape[1],
cursor pos[0]:cursor pos[0] + image shape[0]] = img
        cursor pos[0] += image shape[0] # increment cursor x position
        if cursor pos[0] >= montage shape[0] * image shape[0]:
            cursor pos[1] += image shape[1] # increment cursor y
position
            cursor pos[0] = 0
            if cursor pos[1] >= montage shape[1] * image shape[1]:
                cursor pos = [0, 0]
                image montages.append(montage image)
                # reset black canvas
                montage_image = np.zeros(shape=(image_shape[1] *
(montage shape [1]), image shape [0] * montage shape [0], 3),
                                      dtype=np.uint8)
                start new img = True
    if start new img is False:
        image montages.append(montage image) # add unfinished montage
    return image montages
# take the features for the current image, find all similar
# images in our dataset, and then initialize our list of result
# images
fig, ax = plt.subplots(nrows=2, figsize=(15,15))
queryIdx = 3166# Input Index for which images
MAX RESULTS = 10
queryFeatures = latent features[queryIdx]
```

```
results = perform search(queryFeatures, index dict,
maxResults=MAX RESULTS)
imgs = []
# loop over the results
for (d, j) in results:
    img = np.array(Image.open(images[j]))
    print(j)
    imgs.append(img)
# display the query image
ax[0].imshow(np.array(Image.open(images[queryIdx])))
# build a montage from the results and display it
montage = build montages(imgs, (512, 512), (5, 2))[0]
ax[1].imshow(montage)
testpath = Path('../input/testcbir/Test Images')
testdf = pd.DataFrame()
testdf['image'] = [f for f in os.listdir(testpath) if
os.path.isfile(os.path.join(testpath, f))]
testdf['image'] = '../input/testcbir/Test Images/' +
testdf['image'].astype(str)
testdf.head()
testimages = testdf.image.values
test latent features = get latent features(testimages,
transformations)
test latent features.shape
fig, ax = plt.subplots(nrows=2, figsize=(15, 15))
MAX RESULTS = 10
queryIdx = 12
queryFeatures = test latent features[queryIdx]
results = perform search(queryFeatures, index dict,
maxResults=MAX RESULTS)
imgs = []
# loop over the results
for (d, j) in results:
    img = np.array(Image.open(images[j]))
    print(j)
    imgs.append(img)
# display the query image
ax[0].imshow(np.array(Image.open(testimages[queryIdx])))
```

# build a montage from the results and display it
montage = build\_montages(imgs, (512, 512), (5, 2))[0]
ax[1].imshow(montage)





## 2.2 LSHashing Method

```
!pip install lshashpy3
from lshashpy3 import LSHash
# Locality Sensitive Hashing
params
k = 12 # hash size
L = 5 # number of tables
```

```
d = 14112 # Dimension of Feature vector
lsh = LSHash(hash size=k, input dim=d, num hashtables=L)
# LSH on all the images
for idx,vec in tgdm(feature dict.items()):
    lsh.index(vec.flatten(), extra_data=idx)
# Exporting as pickle
pickle.dump(lsh, open('lsh.p', "wb"))
def get similar item(idx, feature dict, lsh variable, n items=10):
    response =
lsh variable.query(feature dict[list(feature dict.keys())
[idx]].flatten(),
                     num results=n items+1, distance func='hamming')
    imgs = []
    for i in range(1, n items+1):
        imgs.append(np.array(Image.open(images[response[i][0][1]])))
    return imgs
fig, ax = plt.subplots(nrows=2, figsize=(15,15))
queryIdx = 5
ax[0].imshow(np.array(Image.open(images[queryIdx])))
montage = build montages(get similar item(queryIdx, feature dict,
lsh, 10), (512, 512), (5, 2))[0]
ax[1].imshow(montage)
```

#### **End Notes**

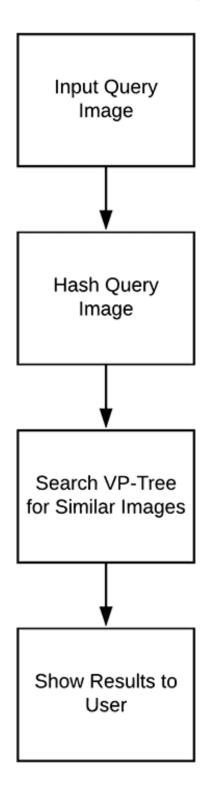
We started with the approach of AutoEncoders for Image Latent Features extraction followed by Image retrieval using Euclidean Distance which was an O(NlogN) approach (Time-Complexity) to Hashing which gave us an O(logN) approach

Another approach was to use Hashing on features obtained from SIFT, SURF, OBS and building the VP Trees ans search the images in it.

# Indexing

# Dataset of Images Hash Each Image Store Hashes **Build VP-Tree**

## Searching



## Clustering of Images

```
from sklearn.cluster import KMeans, MiniBatchKMeans
from scipy.spatial.distance import cdist
from sklearn.metrics import silhouette samples, silhouette score
import matplotlib.cm as cm
%matplotlib inline
def get_latent_features1D(images, transformations):
    latent features1d = []
    for i,image in enumerate(tgdm(images)):
        tensor = transformations(Image.open(image)).to(device)
latent features1d.append(model.encoder(tensor.unsqueeze(0)).cpu().deta
ch().numpy().flatten())
    del tensor
    qc.collect()
    return latent features1d
images = df.image.values
latent features1d = get latent features1D(images, transformations)
latent features1d = np.array(latent features1d)
distortions = []
inertias = []
mapping1 = \{\}
mapping2 = \{\}
K = range(4, 10)
for k in tqdm(K):
    #Building and fitting the model
    kmeanModel = KMeans(n clusters=k).fit(latent features1d)
    distortions.append(sum(np.min(cdist(latent features1d,
kmeanModel.cluster_centers_,
                       'euclidean'),axis=<mark>1</mark>)) /
latent features1d.shape[0])
    inertias.append(kmeanModel.inertia )
    mapping1[k] = sum(np.min(cdist(latent features1d,
kmeanModel.cluster_centers_,
                  'euclidean'),axis=1)) / latent features1d.shape[0]
    mapping2[k] = kmeanModel.inertia
plt.plot(K, distortions, 'bx-')
plt.xlabel('Values of K')
```

```
plt.ylabel('Distortion')
plt.title('The Elbow Method using Distortion')
plt.show()
X = np.array(latent features1d)
K = range(3, 10)
for n clusters in tqdm(K):
    # Create a subplot with 1 row and 2 columns
    fig, (ax1, ax2) = plt.subplots(1, 2)
    fig.set size inches(18, 7)
    # The 1st subplot is the silhouette plot
    # The silhouette coefficient can range from -1, 1 but in this
example all
    # lie within [-0.1, 1]
    ax1.set xlim([-0.1, 1])
    # The (n clusters+1)*10 is for inserting blank space between
silhouette
    # plots of individual clusters, to demarcate them clearly.
    ax1.set ylim([0, len(X) + (n clusters + 1) * 10])
    # Initialize the clusterer with n clusters value and a random
generator
    # seed of 10 for reproducibility.
    clusterer = KMeans(n clusters=n clusters,
random state=RANDOMSTATE)
    cluster labels = clusterer.fit predict(X)
    # The silhouette score gives the average value for all the
samples.
    # This gives a perspective into the density and separation of the
formed
    # clusters
    silhouette avg = silhouette score(X, cluster labels)
    print("For n clusters =", n clusters,
          "The average silhouette score is :", silhouette avg)
    # Compute the silhouette scores for each sample
    sample silhouette values = silhouette samples(X, cluster labels)
    y lower = 10
    for i in range(n clusters):
        # Aggregate the silhouette scores for samples belonging to
        # cluster i, and sort them
        ith cluster_silhouette_values = \
            sample silhouette values[cluster labels == i]
        ith cluster silhouette values.sort()
```

```
size cluster i = ith cluster silhouette values.shape[0]
        y upper = y lower + size cluster i
        color = cm.nipy spectral(float(i) / n clusters)
        ax1.fill betweenx(np.arange(y_lower, y_upper),
                          0, ith cluster silhouette values,
                          facecolor=color, edgecolor=color, alpha=0.7)
        # Label the silhouette plots with their cluster numbers at the
middle
        ax1.text(-0.05, y lower + 0.5 * size cluster i, str(i))
        # Compute the new y lower for next plot
        y lower = y upper + 10 # 10 for the 0 samples
    ax1.set title("The silhouette plot for the various clusters.")
    ax1.set_xlabel("The silhouette coefficient values")
    ax1.set_ylabel("Cluster label")
    # The vertical line for average silhouette score of all the values
    ax1.axvline(x=silhouette avg, color="red", linestyle="--")
    ax1.set yticks([]) # Clear the yaxis labels / ticks
    ax1.set xticks([-0.1, 0, 0.2, 0.4, 0.6, 0.8, 1])
    # 2nd Plot showing the actual clusters formed
    colors = cm.nipy spectral(cluster labels.astype(float) /
n clusters)
    ax2.scatter(X[:, 0], X[:, 1], marker='.', s=30, lw=0, alpha=0.7,
                c=colors, edgecolor='k')
    # Labeling the clusters
    centers = clusterer.cluster centers
    # Draw white circles at cluster centers
    ax2.scatter(centers[:, 0], centers[:, 1], marker='o',
                c="white", alpha=1, s=200, edgecolor='k')
    for i, c in enumerate(centers):
        ax2.scatter(c[0], c[1], marker='$%d$' % i, alpha=1,
                    s=50, edgecolor='k')
    ax2.set_title("The visualization of the clustered data.")
    ax2.set xlabel("Feature space for the 1st feature")
    ax2.set ylabel("Feature space for the 2nd feature")
    plt.suptitle(("Silhouette analysis for KMeans clustering on sample
data "
                  "with n clusters = %d" % n clusters),
                 fontsize=14, fontweight='bo\overline{l}d')
```

```
plt.show()
```

- The Silhouette score isn't significant for any cluster since its close to 0 for every k, that translates to less differentiability for a point to belong to a particular cluster.
- GMM can help in this case because animals share a lot of similar traits with each other in terms of appearance but we have to get the bottleneck case since an animal can only belong to one cluster, so kmeans will be the way to go but a different feature/keypoint detection might help identify right number of clusters.

## Using SIFT/SURF/ORB technique

```
def build dictionary(xfeatures2d, images, n clusters):
    #print('Computing descriptors..')
    desc list = []
    for image path in images:
        image = cv2.imread(image path)
        gray = cv2.cvtColor(image, cv2.COLOR BGR2GRAY)
        kp, dsc = xfeatures2d.detectAndCompute(gray, None)
        desc list.extend(dsc)
    desc = np.array(desc list)
    #print('Creating BoW dictionary using K-Means clustering with
k={}..'.format(n clusters))
    dictionary = MiniBatchKMeans(n clusters=n clusters,
batch_size=100, verbose=0)
    dictionary.fit(desc)
    distortion = sum(np.min(cdist(desc, dictionary.cluster centers,
                      'euclidean'),axis=1)) / desc.shape[0]
    return distortion
orb = cv2.0RB create()
images = df.image.values
K = range(4, 10)
distortions = []
for k in tqdm(K):
    distortions.append(build dictionary(orb, images, n clusters=k))
plt.plot(K, distortions, 'bx-')
plt.xlabel('Values of K')
```

```
plt.ylabel('Distortion')
plt.title('The Elbow Method using Distortion')
plt.show()
```

• The ORB technique tells us there are 6/7 major clusters that are persistent in the data