ML/DL

Code Review ML & DL

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Overview

Feature Extraction Machine Learning

Feature Extraction

```
32
     def prebpicture(bgr):
33
       bgr = cv2.resize(bgr, (256, 256))
34
      lab = cv2.cvtColor(bgr, cv2.COLOR_BGR2LAB)
35
      lab planes = cv2.split(lab)
36
37
      clahe = cv2.createCLAHE(clipLimit=2.0,tileGridSize=(8,8))
      lab_planes[0] = clahe.apply(lab_planes[0])
38
      lab = cv2.merge(lab_planes)
       bgr = cv2.cvtColor(lab, cv2.COLOR_LAB2BGR)
40
      #save image to temp file
41
       #cv2.imwrite('/content/drive/MyDrive/Dataset/temp/clahe-output.jpg',bgr)
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      return bgr
```

Picture preparation and Applying CLAHE

We need to convert pictures into size 256*256 and then apply CLAHE to every single picture and you can save it in temp file if you need to or just recieve bgr

```
#PRE-PROCESSING AND SMOOTHING
Abo, Ago, Aro = cv2.split(image) #splitting into 3 colors
Ar = Aro - Aro.mean()
                               #Preprocessing Red
Ar = Ar - Ar.mean() - Aro.std() #Preprocessing Red
Ar = Ar - Ar.mean() - Aro.std() #Preprocessing Red
Mr = Ar.mean()
                                        #Mean of preprocessed red
SDr = Ar.std()
                                        #SD of preprocessed red
Thr = 49.5 - 12 - Ar.std()
                                        #OD Threshold
Ag = Ago - Ago.mean()
                               #Preprocessing Green
Ag = Ag - Ag.mean() - Ago.std() #Preprocessing Green
Mg = Ag.mean()
                                        #Mean of preprocessed green
SDg = Ag.std()
                                        #SD of preprocessed green
Thg = Ag.mean() + 2*Ag.std() + 49.5 + 12 #OC Threshold
filter = signal.gaussian(99, std=6) #Gaussian Window
filter=filter/sum(filter)
hist,bins = np.histogram(Ag.ravel(),256,[0,256]) #Histogram of preprocessed green channel
histr,binsr = np.histogram(Ar.ravel(),256,[0,256]) #Histogram of preprocessed red channel
smooth_hist_g=np.convolve(filter,hist) #Histogram Smoothing Green
smooth_hist_r=np.convolve(filter,histr) #Histogram Smoothing Red
```

def segment_cd(image,plotHis,plotPic):

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Feature Extraction

Cup and disc segmentation

We can segment the cup and disc using <code>cv2.split()</code> to split the picture into 3 colors (Abo - Blue , Ago-Green , Aro - Red) and we use Red for calculating treshold for disc and Green for calcualting treshold for cup then we creating Filter and histogram to plot graph if you want to see the result in histogram format

https://github.com/Atdhasiri/machineLearning/blob/main/ML/final_feature_extraction.py#L49-L78

```
if plotHis:
80
         plt.subplot(2, 2, 1)
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         plt.plot(hist)
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         plt.title("Preprocessed Green Channel")
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         plt.subplot(2, 2, 2)
85
         plt.plot(smooth_hist_g)
86
         plt.title("Smoothed Histogram Green Channel")
87
         plt.subplot(2, 2, 3)
89
         plt.plot(histr)
90
         plt.title("Preprocessed Red Channel")
92
         plt.subplot(2, 2, 4)
93
         plt.plot(smooth_hist_r)
94
         plt.title("Smoothed Histogram Red Channel")
         plt.show()
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```

Feature Extraction

Plot histogram

this *plotHis* was set in *segment_cd()* 's parameter and if it's true it will plot you some graphs

```
r,c = Ag.shape
Dd = np.zeros(shape=(r,c))
Dc = np.zeros(shape=(r,c))
for i in range(1,r):
  for j in range(1,c):
    if Ar[i,j]>Thr:
      Dd[i,j]=255
    else:
      Dd[i,j]=0
for i in range(1,r):
  for j in range(1,c):
    if Ag[i,j]>Thg:
      Dc[i,j]=1
    else:
      Dc[i,j]=0
#DISPLAYING SEGMENTED OPTIC DISK AND CUP
if plotPic:
  plt.imshow(Dd, cmap = 'gray', interpolation = 'bicubic')
  plt.axis("off")
  plt.title("Optic Disk")
  plt.show()
  plt.imshow(Dc, cmap = 'gray', interpolation = 'bicubic')
  plt.axis("off")
  plt.title("Optic Cup")
  plt.show()
```

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Feature Extraction

Applying Treshold and plot picture

First, it will create zero array(0 = black color) with the same size as the picture and check every pixel of the original picture and compare to it's threshold if that pixel has more value that it's threshold it will plot the zero array in the same location as white color (255 = white color) and if it lower it will plot zero. Do this with cup aswell.

and plotPic was set in $segment_cd()$'s parameter aswell so if it's true it will plot you the picture of segmented cup and disc

```
r1 = cv2.morphologyEx(R1, cv2.MORPH_OPEN, cv2.getStructuringElement(cv2.MORPH_ELLIPSE,(7,7)), iterations = 1)
143
          R2 = cv2.morphologyEx(r1, cv2.MORPH_CLOSE, cv2.getStructuringElement(cv2.MORPH_ELLIPSE,(1,21)), iterations = 1)
144
          r2 = cv2.morphologyEx(R2, cv2.MORPH OPEN, cv2.getStructuringElement(cv2.MORPH ELLIPSE,(21,1)), iterations = 1)
145
          R3 = cv2.morphologyEx(r2, cv2.MORPH_CLOSE, cv2.getStructuringElement(cv2.MORPH_ELLIPSE,(33,33)), iterations = 1)
146
          r3 = cv2.morphologyEx(R3, cv2.MORPH_OPEN, cv2.getStructuringElement(cv2.MORPH_ELLIPSE,(43,43)), iterations = 1)
147
148
          img = clahe.apply(r3)
149
150
151
152
          ret, thresh = cv2.threshold(cup,127,255,0)
          contours, hierarchy = cv2.findContours(thresh, cv2.RETR EXTERNAL, cv2.CHAIN APPROX SIMPLE) #Get
153
          cup diameter = 0
154
          largest area = 0
155
          el cup = contours[0]
156
         if len(contours) != 0:
157
              for i in range(len(contours)):
158
                  if len(contours[i]) >= 5:
159
                      area = cv2.contourArea(contours[i]) #Getting the contour with the largest area
160
                      if (area>largest_area):
161
                          largest area=area
162
                          index = i
163
                          el cup = cv2.fitEllipse(contours[i])
164
165
          cv2.ellipse(img,el cup,(140,60,150),3) #fitting ellipse with the largest area
166
          x,y,w,h = cv2.boundingRect(contours[index]) #fitting a rectangle on the ellipse to get the length of major axis
167
          cup_diameter = max(w,h) #major axis is the diameter https://github.com/Atdhasiri/machineLearning/blob/main/ML/final_feature_extraction.py#L138-L168
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```

R1 = cv2.morphologyEx(cup, cv2.MORPH CLOSE, cv2.getStructuringElement(cv2.MORPH ELLIPSE,(2,2)), iterations = 1)

clahe = cv2.createCLAHE(clipLimit=2.0, tileGridSize=(10,10))

#morphological closing and opening operations

def cdr(cup,disc,plot):

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Feature Extraction

Calculate cup-to-disc Ratio

Recieve to picture form *segment_cd()* and create clahe to apply with *morphologyEX()* to draw outline of the cup and disc and create contours out of cup's threshold and create elipse to cover the area of cup and then use cv2.boundingReact() to get the final size of ellipse and then get max width and height to get *cup_diameter*

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```
Feature
Extraction
```

```
#morphological closing and opening operations
R1 = cv2.morphologyEx(disc, cv2.MORPH CLOSE, cv2.getStructuringElement(cv2.MORPH ELLIPSE,(2,2)), iterations = 1)
r1 = cv2.morphologyEx(R1, cv2.MORPH_OPEN, cv2.getStructuringElement(cv2.MORPH_ELLIPSE,(7,7)), iterations = 1)
R2 = cv2.morphologyEx(r1, cv2.MORPH CLOSE, cv2.getStructuringElement(cv2.MORPH ELLIPSE,(1,21)), iterations = 1)
r2 = cv2.morphologyEx(R2, cv2.MORPH OPEN, cv2.getStructuringElement(cv2.MORPH ELLIPSE,(21,1)), iterations = 1)
R3 = cv2.morphologyEx(r2, cv2.MORPH_CLOSE, cv2.getStructuringElement(cv2.MORPH_ELLIPSE,(33,33)), iterations = 1)
r3 = cv2.morphologyEx(R3, cv2.MORPH_OPEN, cv2.getStructuringElement(cv2.MORPH_ELLIPSE,(43,43)), iterations = 1)
img2 = clahe.apply(r3)
ret, thresh = cv2.threshold(disc,127,255,0)
contours, hierarchy = cv2.findContours(thresh, cv2.RETR EXTERNAL, cv2.CHAIN APPROX SIMPLE) #Getting all possible contours in the segmented image
disk diameter = 0
largest_area = 0
el disc = el cup
if len(contours) != 0:
      for i in range(len(contours)):
        if len(contours[i]) >= 5:
            area = cv2.contourArea(contours[i]) #Getting the contour with the largest area
            if (area>largest area):
                largest area=area
                index = i
                el disc = cv2.fitEllipse(contours[i])
cv2.ellipse(img2,el_disc,(140,60,150),3) #fitting ellipse with the largest area
x,y,w,h = cv2.boundingRect(contours[index]) #fitting a rectangle on the ellipse to get the length of major axis
disk diameter = max(w,h) #major axis is the diameter
```

Calculate disk_diameter

Do the same thing as *cup_diameter*

```
def adjust gamma(image, gamma=1.0):
220
         table = np.array([(i / 255.0) ** gamma) * 255
            for i in np.arange(0, 256)]).astype("uint8")
223
224
         return cv2.LUT(image, table)
225
      def extract ma(image):
226
227
          r,g,b=cv2.split(image)
          comp=255-g
228
          clahe = cv2.createCLAHE(clipLimit=5.0, tileGridSize=(8,8))
229
          histe=clahe.apply(comp)
230
          adjustImage = adjust_gamma(histe,gamma=3)
231
          comp = 255-adjustImage
232
               adjust gamma(comp,gamma=4)
233
234
          J = 255-J
         J = adjust gamma(J,gamma=4)
235
236
         K=np.ones((11,11),np.float32)
237
          L = cv2.filter2D(J,-1,K)
238
239
          ret3,thresh2 = cv2.threshold(L,125,255,cv2.THRESH BINARY|cv2.THRESH OTSU)
240
          kernel2=np.ones((9,9),np.uint8)
241
          tophat = cv2.morphologyEx(thresh2, cv2.MORPH_TOPHAT, kernel2)
242
          kernel3=np.ones((7,7),np.uint8)
243
          opening = cv2.morphologyEx(tophat, cv2.MORPH OPEN, kernel3)
244
          return opening
245
```

Feature Extraction

Calculate MA

MA is one of the feature that can detect diabetic by checking swelling in the wall of a blood vessel First, it will adjust gramma of the picture and calculate the microaneurysm and return the numpy of the result

```
def fast_glcm(img, vmin=0, vmax=255, nbit=8, kernel_size=5):
mi, ma = vmin, vmax
ks = kernel_size
```

```
Feature
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h,w = img.shape 252 254 # digitize 255 bins = np.linspace(mi, ma+1, nbit+1) gl1 = np.digitize(img, bins) - 1 256 gl2 = np.append(gl1[:,1:], gl1[:,-1:], axis=1) 258 259 # make glcm glcm = np.zeros((nbit, nbit, h, w), dtype=np.uint8) 260 for i in range(nbit): 261 for j in range(nbit): 262 mask = ((gl1==i) & (gl2==j))263 264 glcm[i,j, mask] = 1265 kernel = np.ones((ks, ks), dtype=np.uint8) 266 for i in range(nbit): 267 for j in range(nbit): 268 glcm[i,j] = cv2.filter2D(glcm[i,j], -1, kernel)269 270

glcm = glcm.astype(np.float32)

return glcm

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Calculate GLCM

This function will create GLCM but first you need to change your picture into greyscale before using fast_glcm() and it will return numpy array as a result

```
def fast_glcm_mean(img, vmin=0, vmax=255, nbit=8, ks=5):
276
          calc glcm mean
278
          h,w = img.shape
279
          glcm = fast glcm(img, vmin, vmax, nbit, ks)
280
          mean = np.zeros((h,w), dtype=np.float32)
281
          for i in range(nbit):
282
              for j in range(nbit):
283
                  mean += glcm[i,j] * i / (nbit)**2
284
285
286
          return mean
287
288
      def fast glcm std(img, vmin=0, vmax=255, nbit=8, ks=5):
289
          1 1 1
290
          calc glcm std
          h,w = img.shape
293
          glcm = fast_glcm(img, vmin, vmax, nbit, ks)
294
          mean = np.zeros((h,w), dtype=np.float32)
          for i in range(nbit):
296
              for j in range(nbit):
297
                  mean += glcm[i,j] * i / (nbit)**2
298
299
          std2 = np.zeros((h,w), dtype=np.float32)
300
          for i in range(nbit):
301
              for j in range(nbit):
302
                  std2 += (glcm[i,j] * i - mean)**2
303
304
          std = np.sqrt(std2)
305
          return std
```

Feature Extraction

Calculate GLCM using mean and std methods

the Mean method is using mean to calculate GLCM and the std method is using standard deviation to calculate GLCM

```
def fast glcm contrast(img, vmin=0, vmax=255, nbit=8, ks=5):
   calc glcm contrast
   h,w = img.shape
   glcm = fast_glcm(img, vmin, vmax, nbit, ks)
   cont = np.zeros((h,w), dtype=np.float32)
   for i in range(nbit):
       for j in range(nbit):
            cont += glcm[i,j] * (i-j)**2
    return cont
def fast_glcm_dissimilarity(img, vmin=0, vmax=255, nbit=8, ks=5):
   calc glcm dissimilarity
   h,w = img.shape
   glcm = fast_glcm(img, vmin, vmax, nbit, ks)
   diss = np.zeros((h,w), dtype=np.float32)
   for i in range(nbit):
       for j in range(nbit):
           diss += glcm[i,j] * np.abs(i-j)
```

Feature Extraction

Calculate GLCM using contrast and dissimilarity methods

Contrast is measure of variation in intensity.

Dissimilarity is the measure of relationship between pairs with similar and different intensities, Higher is the value of dissi, more is the dissimilarity between the pairs of pixels

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```
def fast glcm homogeneity(img, vmin=0, vmax=255, nbit=8, ks=5):
    calc glcm homogeneity
    h,w = img.shape
    glcm = fast_glcm(img, vmin, vmax, nbit, ks)
    homo = np.zeros((h,w), dtype=np.float32)
   for i in range(nbit):
        for j in range(nbit):
            homo += glcm[i,j] / (1.+(i-j)**2)
    return homo
def fast glcm ASM(img, vmin=0, vmax=255, nbit=8, ks=5):
    1.1.1
    calc glcm asm, energy
   h,w = img.shape
    glcm = fast glcm(img, vmin, vmax, nbit, ks)
    asm = np.zeros((h,w), dtype=np.float32)
   for i in range(nbit):
        for j in range(nbit):
            asm += glcm[i,j]**2
    ene = np.sqrt(asm)
    return asm, ene
```

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Feature Extraction

Calculate GLCM using homogeneity and ASM methods

Homogeneity is the metric used for analysis of homogeneity in an image.

Feature Extraction

```
def fast_glcm_max(img, vmin=0, vmax=255, nbit=8, ks=5):
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368
          calc glcm max
         glcm = fast_glcm(img, vmin, vmax, nbit, ks)
370
         max_ = np.max(glcm, axis=(0,1))
         return max
374
      def fast glcm entropy(img, vmin=0, vmax=255, nbit=8, ks=5):
376
         calc glcm entropy
377
378
          glcm = fast_glcm(img, vmin, vmax, nbit, ks)
379
         pnorm = glcm / np.sum(glcm, axis=(0,1)) + 1./ks**2
380
          ent = np.sum(-pnorm * np.log(pnorm), axis=(0,1))
382
          return ent
```

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Calculate GLCM using Max and **Entropy methods**

Max method is to find the max value of neighbor Entro is the measure of randomness or uncertainty

create machine learning model

```
def create_model(model_name):
    if(model_name == "svm"):
        model = SVC(kernel='rbf',probability=True)
        #kernel = {'linear', 'poly', 'rbf', 'sigmoid', 'precomputed'}
    elif(model_name == "knn"):
        model = KNeighborsClassifier(n_neighbors=5)
    elif(model_name == "rf"):
        model = RandomForestClassifier(n_estimators = 400, criterion = "gini")
    return model
```

- Do 5-fold cross validation.
- separate data to train set and validate set.
- scaling feature.

- Create model
- Do feature selection by important weight of spacific model
- Apply those feature to classifier
- change model to be one vs rest
- train the model

 select best model of all fold by compare sum of micro average AUC and macro average AUC

```
if roc_auc["micro"]+roc_auc["macro"] > best_sum_average_AUC:
    best_sum_average_AUC = roc_auc["micro"]+roc_auc["macro"]
    best_sum_3class_AUC = roc_auc[0] + roc_auc[1] + roc_auc[2]
    bestModel = model
    bestFold = i+1

elif roc_auc["micro"]+roc_auc["macro"] == best_sum_average_AUC and best_sum_3class_AUC < roc_auc[0] + roc_auc[1] + roc_auc[2]:
    best_sum_average_AUC = roc_auc["micro"]+roc_auc["macro"]
    best_sum_3class_AUC = roc_auc[0] + roc_auc[1] + roc_auc[2]
    bestModel = model
    bestFold = i+1</pre>
```

plot mean ROC curve and calculate SD

```
plt.show()
plt.figure()
plt.plot(mean_fpr, mean_tpr, color='b',
        label=r'Mean ROC (AUC = %0.2f $\pm$ %0.2f)' % (mean_auc, std_auc),
        lw=2)
for i in range(5):
    plt.plot(fprs[i], tprs[i],label='micro-average ROC curve of fold{} (area = {:0.2f})'.format(i+1,aucs[i]), alpha=.5)
plt.plot([0, 1], [0, 1], 'k--', lw=lw)
plt.xlim([0.0, 1.0])
plt.ylim([0.0, 1.05])
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Mean ROC of all fold')
plt.legend(loc="lower right")
plt.show()
```

mean_tpr = np.mean(interp_tpr_all_fold, axis=0)

std_tpr = np.std(interp_tpr_all_fold, axis=0)

tprs_upper = np.minimum(mean_tpr + std_tpr, 1)

tprs_lower = np.maximum(mean_tpr - std_tpr, 0)

mean_auc = auc(mean_fpr, mean_tpr)

 $mean_tpr[-1] = 1.0$

std_auc = np.std(aucs)

https://github.com/Atdhasiri/machineLearning/blob/main/ML/Machine%20learning.py#L287-L312



Deep Learning

Overview

Convert Image
Deep Learning

Convert Image

```
def save_to_numpy(imgs,label,path,filename):
    np.save(os.path.join(path, filename + '.npy'), imgs)
    np.save(os.path.join(path, filename + '_label' + '.npy'), label)
    print('Numpy files have been saved')
```

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Save picture in numpy

This function will create .npy file in the path that you define

https://github.com/Atdhasiri/machineLearning/blob/main/DL/convert%20image%20to%20numpy.py#L23-L26

classes = ['Glaucoma', 'Normal', 'Other']

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```
num_folders = ['1','2','3','4','5']
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33
     temp_x = []
34
    temp_y = []
35
     x=[]
     y=[]
```

Convert Image

Declaration

Declare the Classes and Fold that you have and create temp array for x (data) and temp array for y (label) and the real x and y. I created temp because I will create nested array which mean len(x) will answer 5 (Fold) and in each it will contain pictures

```
for k in range(len(num folders)):
  for i in range(len(classes)):
    images lst = os.listdir('/content/drive/MyDrive/Dataset/Train/' + classes[i] + '/Fold-' + num folders[k])
    #print(images lst)
    count = 0
    for j in images lst:
      count += 1
     if count == 2:
       break
     print('\n' + j + ' ----> ', count, '/', len(images lst))
     print('class : ' + classes[i])
     print('fold : ' + num_folders[k])
     img = cv2.imread('/content/drive/MyDrive/Dataset/Train/' + classes[i] + '/Fold-' + num folders[k]+'/'+j)
     img = cv2.resize(img, (256, 256))
     temp x.append(np.array(img))
     temp y.append(i)
 x.append(temp x)
 temp x = []
 y.append(temp y)
 temp y = []
#save to .npy
#output path = '/content/drive/MyDrive/Dataset/np data/'
#save to numpy(x,y,output path,'Train')
print('-----')
print('done')
```

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Convert Image

Loop for every pictures in Train

It will loop every Fold that you have and then every Classes that you have and then it will resize the image to 256*256 and append to temp_x first and temp_y will appen the number of class

After the finished a Fold the temp_x and temp_y will append to real x and y to add a nested array and do this to another Fold

```
\#temp_y = []
77 y=[]
78
     for i in range(len(classes)):
       images lst = os.listdir('/content/drive/MyDrive/Dataset/Test/' + classes[i])
       #print(images lst)
       count = 0
83
       for j in images_lst:
84
         count += 1
         print('\n' + j + ' ----> ', count, '/', len(images lst))
         print('class : ' + classes[i])
         #print('fold : ' + num_folders[k])
88
         img = cv2.imread('/content/drive/MyDrive/Dataset/Test/' + classes[i] +'/'+ j)
89
         img = cv2.resize(img, (256, 256))
90
         x.append(np.array(img))
91
         if classes[i] == 'Glaucoma':
92
93
           y.append('0')
           print('Y appended: 0')
94
         elif classes[i] == 'Normal':
           y.append('1')
           print('Y appended : 1')
97
98
         else:
           y.append('2')
99
           print('Y appended : 2')
103
    #save to .npy
     #output_path = '/content/drive/MyDrive/Dataset/np_data/'
105
     #save to numpy(x,y,output path, 'test folder not the training')
106
107
108
     print('done')
```

classes = ['Glaucoma', 'Normal', 'Other']

#num folders = ['1','2','3','4','5']

74 $\#temp_x = []$

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Convert lmage

Loop for every pictures in Test Folder

For dataset Test we don't have Fold so we just loop every pictures and add it directly to the x(data) array and y(label) array


```
Deep learning
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load data

```
y = np.load("/content/drive/MyDrive/Colab Notebooks/Dataset/np_data/TRAIN_label.npy",allow_pickle=True)
X_test = np.load("/content/drive/MyDrive/Colab Notebooks/Dataset/np_data/test.npy",allow_pickle=True)
y_test = np.load("/content/drive/MyDrive/Colab Notebooks/Dataset/np_data/test_label.npy",allow_pickle=True)

X = np.array([np.array(fold) for fold in X])
y = np.array([np.array(fold) for fold in y])
X_test = np.array(X_test)
y_test = np.array(y_test)
```

create model

```
def build model(base model name):
    print('Creating Model : {}......'.format(base model name.upper()))
    if base_model_name == 'DenseNet121' :
       init_model = DenseNet121(input_shape= (height, width ,3), include_top=False, weights='imagenet', pooling='avg')
       x = init_model.output
       out_layer = Dense(num_output, activation="softmax")(x)
       model = Model(init_model.input, out_layer)
    elif base_model_name == 'ResNet101V2' :
       init model = ResNet101V2(input shape= (height, width ,3), include top=False, weights='imagenet', pooling='avg')
       x = init_model.output
       out_layer = Dense(num_output, activation="softmax")(x)
       model = Model(init model.input,out layer)
   else :
        sys.exit("Model Not Support")
   return→model
```

- create early stop function to early stop training if validation loss not decrease for 10 epochs
- create reduce learning rate function to reduce learning rate if validation loss not decrease for 7 epochs

- Create and compile model
- apply optimizer(adam) and loss algorithm

save weight at the best epoch(focus on val_loss)

```
checkpoint_path = "/content/drive/MyDrive/Colab Notebooks/Dataset/weight/{}-weights_fold-{}.hdf5".format(model_name,i+1)
mcp_save = ModelCheckpoint(checkpoint_path, save_best_only=True, monitor='val_loss', mode='min',save_weights_only=True)
```

```
history = model.fit(X_train,y_train,
  steps_per_epoch = (len(X_train) / batch_size ),
  epochs = 50,
  validation_data=(X_val, y_val),
  callbacks=[earlyStopping, mcp_save, reduce_lr_loss]
)
```

train model

 select best model of all fold by compare sum of micro average AUC and macro average AUC

```
if roc_auc["micro"]+roc_auc["macro"] > best_sum_average_AUC:
    best_sum_average_AUC = roc_auc["micro"]+roc_auc["macro"]
    best_sum_3class_AUC = roc_auc[0] + roc_auc[1] + roc_auc[2]
    bestModel = model
    bestFold = i+1

elif roc_auc["micro"]+roc_auc["macro"] == best_sum_average_AUC and best_sum_3class_AUC < roc_auc[0] + roc_auc[1] + roc_auc[2]:
    best_sum_average_AUC = roc_auc["micro"]+roc_auc["macro"]
    best_sum_3class_AUC = roc_auc[0] + roc_auc[1] + roc_auc[2]
    bestModel = model
    bestFold = i+1</pre>
```

Evaluation

```
def calculateFprTprAuc(binary_y_test, y_score):
    # Compute ROC curve and ROC area for each class
    fpr = dict()
    tpr = dict()
    roc_auc = dict()
    for i in range(n_classes):
        fpr[i], tpr[i], _ = roc_curve(binary_y_test[:, i], y_score[:, i])
        roc_auc[i] = auc(fpr[i], tpr[i])

# Compute micro-average ROC curve and ROC area
    fpr["micro"], tpr["micro"], _ = roc_curve(binary_y_test.ravel(), y_score.ravel())
    roc_auc["micro"] = auc(fpr["micro"], tpr["micro"])
    return fpr , tpr , roc_auc
```

Calculate FPR TPR and AUC score of each classes and micro average

```
def plotROC(fpr,tpr,roc_auc,title):
    #aggregate all false positive rates
    all_fpr = np.unique(np.concatenate([fpr[i] for i in range(n_classes)]))
    #interpolate all ROC curves
    mean_tpr = np.zeros_like(all_fpr)
    for i in range(n_classes):
        mean tpr += interp(all fpr, fpr[i], tpr[i])
    #average tpr and compute AUC
    mean_tpr /= n_classes
    fpr["macro"] = all fpr
    tpr["macro"] = mean_tpr
    roc_auc["macro"] = auc(fpr["macro"], tpr["macro"])
    # Plot all ROC curves
    plt.figure()
    plt.plot(fpr["micro"], tpr["micro"],
             label='micro-average ROC curve (area = {0:0.2f})'
                   ''.format(roc auc["micro"]),
             color='deeppink', linestyle=':', linewidth=4)
    plt.plot(fpr["macro"], tpr["macro"],
             label='macro-average ROC curve (area = {0:0.2f})'
                   ''.format(roc_auc["macro"]),
             color='navy', linestyle=':', linewidth=4)
    colors = cycle(['aqua', 'darkorange', 'cornflowerblue'])
    for i, color in zip(range(n_classes), colors):
        plt.plot(fpr[i], tpr[i], color=color, lw=lw,
                 label='ROC curve of {0} (area = {1:0.2f})'
                 ''.format(classes_list[i], roc_auc[i]))
    plt.plot([0, 1], [0, 1], 'k--', lw=lw)
```

plt.xlim([0.0, 1.0])

Evaluation

calculate macro average and Plot ROC curve of each classes(One vs Rest) and plot micro average and macro average of them

```
title=None,
                      cmap=plt.cm.Blues):
np.set printoptions(precision=2)
if not title:
    title = 'Normalized confusion matrix'
# Compute confusion matrix
cm = confusion_matrix(y_true, y_pred)
# Only use the labels that appear in the data
classes = classes[unique labels(y true, y pred)]
cm_normalize = cm.astype('float') / cm.sum(axis=1)[:, np.newaxis]
print("Normalized confusion matrix")
#print(cm)
fig, ax = plt.subplots()
im = ax.imshow(cm_normalize, interpolation='nearest', cmap=cmap)
ax.figure.colorbar(im, ax=ax)
# We want to show all ticks...
ax.set(xticks=np.arange(cm_normalize.shape[1]),
       yticks=np.arange(cm normalize.shape[0]),
       # ... and label them with the respective list entries
       xticklabels=classes, yticklabels=classes,
       title=title,
       ylabel='True label',
       xlabel='Predicted label')
```

def plot_confusion_matrix(y_true, y_pred, classes,

Evaluation

Plot 3 classes confusion metrix with normalization(percentage)

2

1. https://github.com/Atdhasiri/machineLearning/blob/main/DL/Deep%20learning.py#L118-L150

2. https://github.com/Atdhasiri/machineLearning/blob/main/DL/Deep%20learning.py#L153-L167

Evaluation

```
# Sensitivity, hit rate, recall, or true positive rate
TPR = TP/(TP+FN)
# Specificity or true negative rate
TNR = TN/(TN+FP)
# Precision or positive predictive value
PPV = TP/(TP+FP)
                                                           Calculate and print sensitivity, specificity
# Negative predictive value
NPV = TN/(TN+FN)
                                                           and etc. by the value from confusion
# Fall out or false positive rate
FPR = FP/(FP+TN)
                                                            matrix
# False negative rate
FNR = FN/(TP+FN)
# False discovery rate
FDR = FP/(TP+FP)
# F1-score
F1 = 2*((PPV*TPR)/(PPV+TPR))
# Overall accuracy
ACC = (TP+TN)/(TP+FP+FN+TN)
print("="*60)
print("{:<26} | {:>8} {:>8} {:>8}".format("value", "Glaucoma", "Normal", "Other"))
print("="*60)
print("{:<26} | {:>8.2f} {:>8.2f} {:>8.2f}".format("Sensitivity(TPR)", TPR[0], TPR[1], TPR[2]))
print("{:<26} | {:>8.2f} {:>8.2f} {:>8.2f}".format("Specificity(TNR)", TNR[0], TNR[1], TNR[2]))
print("{:<26} | {:>8.2f} {:>8.2f} {:>8.2f}".format("Precision(PPV)", PPV[0], PPV[1], PPV[2]))
print("{:<26} | {:>8.2f} {:>8.2f} {:>8.2f}".format("Negative predictive value", NPV[0], NPV[1], NPV[2]))
print("{:<26} | {:>8.2f} {:>8.2f} {:>8.2f}".format("False positive rate", FPR[0], FPR[1], FPR[2]))
print("{:<26} | {:>8.2f} {:>8.2f} {:>8.2f}".format("False negative rate", FNR[0], FNR[1], FNR[2]))
print("{:<26} | {:>8.2f} {:>8.2f} {:>8.2f}".format("False discovery rate", FDR[0], FDR[1], FDR[2]))
print("{:<26} | {:>8.2f} {:>8.2f} {:>8.2f}".format("F1-score", F1[0], F1[1], F1[2]))
print("{:<26} | {:>8.2f} {:>8.2f} {:>8.2f}".format("Overall accuracy", ACC[0], ACC[1], ACC[2]))
```

https://github.com/Atdhasiri/machineLearning/blob/main/DL/Deep%20learning.py#L181-L210

ML&DL

Thanks for attending!