# Diabetic Retinopathy Report

### 1. Overview

Diabetic Retinopathy (DR) is an eye disease associated with long-standing diabetes. Around 40% to 45% of Americans with diabetes have some stage of the disease. A clinician has rated the presence of diabetic retinopathy in each image on a scale of 0 to 4, according to the following scale:

- 0 No DR
- 1 Mild
- 2 Moderate
- 3 Severe
- 4 Proliferative DR

Our task is to create an automated analysis system capable of assigning a score based on this scale.

### 2. Dataset

Training data consists of 35126 images which are further classified into following five categories:

Category	Scaling	Number of images
No DR	0	25810
Mild DR	1	2443
Moderate DR	2	5292
Severe DR	3	873
Proliferative DR	4	708

## 3. Softwares

• Google Cloud instance with following specifications is used:

a) Zone: us-east1-d

b) Machine type: 24 vCPU

c) Boot disk: Ubuntu 17.04 with 400GB size

• Python version: 2.7

• Libraries used are given in the code

## 4. Image Preprocessing and Data augmentation:

We used opency to perform foolowing modifications in the existing data

- Resized the images to (385\*385) pixels.
- Cropped horizontally the images by 40 pixels.
- Randomly rotated the images between 0 and 360 degrees.
- Subtracted the local average color, the avgerage gets mapped to 50% gray.
- Applied horizontal and vertical flip.



 $\begin{array}{ccc} {\rm Image:} \ 16\_{\rm left} \\ {\rm Rating:} \ 4 \end{array}$ 

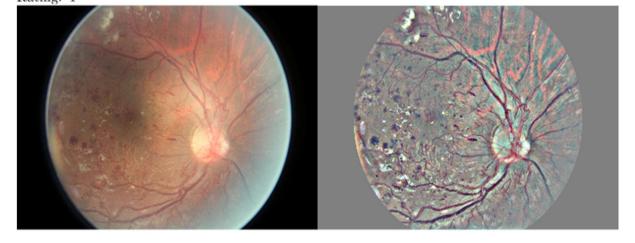


Figure 4: Two images from the training set. Original images on the left and preprocessed images on the right.

```
import cv2 , glob , numpy ,os
   TRAIN DIR='/home/vishal041196/train'
   def scaleRadius(img,scale):
       x=img[img.shape[0]/2,:,:].sum(1)
       r=(x>x.mean()/10).sum()/2
       s=scale*1.0/r
       return cv2.resize(img,(0,0),fx=s,fy=s)
   scale=300
   for img in os.listdir(TRAIN DIR):
       for f in glob.glob(img):
          try:
              print(f)
              a=cv2.imread(f)
              #scaleimg to a given radius
              a=scaleRadius(a,scale)
              #subtractlocal mean color
              a=cv2.addWeighted(a,4,
                               cv2. Gaussian Blur(a,(0,0),scale/30),-4,
              #remove o u t e r 10%
              b=numpy.zeros(a.shape)
              cv2.circle(b,(a.shape[1]/2,a.shape[0]/2),
              int(scale*0.9),(1,1,1),-1,8,0)
              a=a*b+128*(1-b)
              cv2.imwrite(str(scale)+"_"+f,a)
          except:
              print(f)
9 Code for creating training data:
   import cv2 ,os ,pandas as pd ,numpy as np ,random
   from random import shuffle
   from tqdm import tqdm
   IMG SIZE=600
   TRAIN DIR='/home/vishal041196/train1'
   training_data=[]
   for img in tqdm(os.listdir(TRAIN_DIR)):
       label=[1,0]
       num=img.split('_')[1]+'_'+img.split('_')[2].split('.')[0]
       print(num)
       data=pd.read_csv('trainLabels.csv')
       a=data.level[data.image==num]
       if (a.values[0] == 0):label=[1,0]
       else:label=[0,1]
       path = os.path.join(TRAIN_DIR,img)
       img = cv2.imread(path,cv2.IMREAD_UNCHANGED)
       img = cv2.resize(img, (IMG_SIZE,IMG_SIZE))
       training_data.append([np.array(img),np.array(label)])
       rows, cols = 600,600
       r=random.random()*360
       M = cv2.getRotationMatrix2D((cols/2,rows/2),r,1)
       dst = cv2.warpAffine(img,M,(cols,rows))
       training_data.append([np.array(dst),np.array(label)])
   shuffle(training_data)
   np.save('train data preprocess2.npy', training data)
Convolution neural network model:
```

#### **Structure:-**

The core of our solution was a deep convolutional neural network. Although we started with fairly shallow models — 4 convolutional layers, we quickly discovered that adding more layers, and filters inside layers helps a lot. Our best single model consisted of 16 convolutional layers. The detailed architecture is:

Type	Units	Filter	Stride
Input			
Conv	16	3	1
Conv	16	1	1
MaxPool	-	3	2
Conv	32	3	1
Conv	32	1	1
MaxPool	-	3	2
Conv	64	3	1
Conv	64	1	1
MaxPool	-	3	2
Conv	128	3	1
Conv	128	1	1
MaxPool	-	3	2
Conv	256	3	1
Conv	256	1	1
MaxPool	-	3	2
Conv	384	3	1
Conv	384	3	1
MaxPool	-	3	2
Conv	512	3	1
Conv	512	3	1
MaxPool	-	3	2
Conv	512	3	1
Conv	512	3	1
MaxPool	-	3	2
Dropout		0.5	-
fully_connected	1024	-	-
fully_connected	1024	-	-
fully_connected	2	-	-

```
from sklearn.metrics import confusion matrix
from sklearn import metrics
import sklearn as sk
import cv2
import numpy as np
import os
from random import shuffle
from tadm import tadm
#TRAIN_DIR = '/home/vishal041196/train'
LR = 1e-3
MODEL_NAME = 'retinopathy-{}-{}.newmodel'.format(LR, '2conv-basic')
IMG_SIZE = 600
import tflearn
from tflearn.layers.conv import conv_2d, max_pool_2d
from tflearn.layers.core import input_data, dropout, fully_connected
from tflearn.layers.estimator import regression---
convnet = input_data(shape=[None, IMG_SIZE, IMG_SIZE, 3], name='input')
convnet = conv_2d(convnet, 16, 3, activation='relu')
convnet = conv_2d(convnet, 16, 1, activation='relu')
convnet = max_pool_2d(convnet, 3,strides=2)
convnet = conv_2d(convnet, 32, 3, activation='relu')
convnet = conv_2d(convnet, 32, 1, activation='relu')|
convnet = max_pool_2d(convnet, 3, strides=2)
convnet = conv_2d(convnet, 64, 3, activation='relu')
convnet = conv_2d(convnet, 64, 1, activation='relu')
convnet = max_pool_2d(convnet, 3,strides=2)
convnet = conv_2d(convnet, 128, 3, activation='relu')
convnet = conv_2d(convnet, 128, 1, activation='relu')
convnet = max_pool_2d(convnet, 3, strides=2)
convnet = conv_2d(convnet, 256, 3, activation='relu')
convnet = conv_2d(convnet, 256, 1, activation='relu')
convnet = max_pool_2d(convnet, 3,strides=2)
convnet = conv_2d(convnet, 384, 3, activation='relu')
convnet = conv_2d(convnet, 384, 1, activation='relu')
convnet = max_pool_2d(convnet, 3, strides=2)
convnet = conv_2d(convnet, 512, 3, activation='relu')
convnet = conv_2d(convnet, 512, 3, activation='relu')
convnet = max pool 2d(convnet, 3,strides=2)
convnet = conv 2d(convnet, 512, 3, activation='relu')
convnet = conv 2d(convnet, 512, 3, activation='relu')
convnet = max pool 2d(convnet, 3,strides=2)
convnet = dropout(convnet, 0.5)
convnet = fully connected(convnet, 1024, activation='softmax')
convnet = fully_connected(convnet, 1024, activation='softmax')
convnet = fully connected(convnet, 2, activation='softmax')
convnet = regression(convnet, optimizer='adam', learning_rate=LR, loss='categorical_crossentropy', name='targets')
model = tflearn.DNN(convnet, tensorboard dir='log')
train data=np.load('train data preprocess2.npy')
train = train data[:-7000]
test = train data[-7000:]
X = np.array([i[0] for i in train]).reshape(-1,IMG SIZE,IMG SIZE,3)
Y = [i[1]  for i in train]
test_x = np.array([i[0] for i in test]).reshape(-1,IMG_SIZE,IMG_SIZE,3)
test y = [i[1] \text{ for } i \text{ in } test]
if os.path.exists('{}.meta'.format(MODEL NAME)):
    model.load(MODEL NAME)
    print('model loaded!')
model.fit({'input': X}, {'targets': Y}, n epoch=1, validation set=({'input': test x}, {'targets': test y}),
    snapshot step=500, show metric=True, run id=MODEL NAME)
model.save(MODEL NAME)
```

**9** Code for checking the predictions:

```
y_true=[]
y_pred=[]
for j in test_y:
    if (np.argmax(j)):y_true.append(1)
    else:y_true.append(0)
for i in test_x:
    pred=model.predict([i])[0]
    if(np.argmax(pred)==0):y_pred.append(0)
    else:y_pred.append(1)

print("Precision", sk.metrics.precision_score(y_true, y_pred))
print("Recall", sk.metrics.recall_score(y_true, y_pred))
print("f1_score", sk.metrics.f1_score(y_true, y_pred))
print("confusion_matrix")
print(sk.metrics.confusion_matrix(y_true, y_pred))
fpr, tpr, tresholds = sk.metrics.roc_curve(y_true, y_pred)
```

### **Results and Conclusions:**

Number of images in Training set = 64890Number of images in Validation set = 7500

After 70 epochs the results are as follows:

```
Training accuracy = .9615
Validation accuracy = .7299
Precision = .753
Recall = .6909
fl_score = .7297
```

#### Confusion matrix:

2860	857
1160	2614