Breast cancer classification with Keras and Deep Learning **Import Packages** In $[1]\colon$ \mid # set the matplotlib backend so figures can be saved in the background import matplotlib matplotlib.use("Agg") # import the necessary packages from tensorflow.keras.preprocessing.image import ImageDataGenerator from tensorflow.keras.callbacks import LearningRateScheduler from tensorflow.keras.models import Sequential from tensorflow.keras.layers import BatchNormalization from tensorflow.keras.layers import SeparableConv2D from tensorflow.keras.layers import MaxPooling2D from tensorflow.keras.layers import Activation from tensorflow.keras.layers import Flatten from tensorflow.keras.layers import Dropout from tensorflow.keras.layers import Dense from tensorflow.keras.optimizers import Adagrad from tensorflow.keras import backend as K from tensorflow.keras.utils import to_categorical from imutils import paths import matplotlib.pyplot as plt import numpy as np import argparse import random import shutil import sklearn import sklearn.metrics import os Our Config class In [2]: | class Config: # initialize the path to the *original* input directory of images ORIG_INPUT_DATASET = "data" # initialize the base path to the *new* directory that will contain # our images after computing the training and testing split BASE_PATH = "data" # derive the training, validation, and testing directories TRAIN_PATH = os.path.sep.join([BASE_PATH, "training"]) VAL_PATH = os.path.sep.join([BASE_PATH, "validation"]) TEST_PATH = os.path.sep.join([BASE_PATH, "testing"]) # define the amount of data that will be used training TRAIN_SPLIT = 0.8 # the amount of validation data will be a percentage of the # *training* data VAL_SPLIT = 0.1 # initialize our config class config = Config() Building the breast cancer image dataset # and shuffle them imagePaths = list(paths.list_images(config.ORIG_INPUT_DATASET)) random.seed(42) random.shuffle(imagePaths) # compute the training and testing split i = int(len(imagePaths) * config.TRAIN_SPLIT) trainPaths = imagePaths[:i] testPaths = imagePaths[i:] # we'll be using part of the training data for validation i = int(len(trainPaths) * config.VAL_SPLIT) valPaths = trainPaths[:i] trainPaths = trainPaths[i:] # define the datasets that we'll be building datasets = [("training", trainPaths, config.TRAIN_PATH), ("validation", valPaths, config.VAL_PATH), ("testing", testPaths, config.TEST_PATH) # for (dType, imagePaths, baseOutput) in datasets: # # show which data split we are creating # print("[INFO] building '{}' split".format(dType)) # # if the output base output directory does not exist, create it

```
In [3]: # grab the paths to all input images in the original input directory
In [4]: # Loop over the datasets
               if not os.path.exists(baseOutput):
                  print("[INFO] 'creating {}' directory".format(baseOutput))
                  os.makedirs(baseOutput)
         # # loop over the input image paths
              for inputPath in imagePaths:
                 # extract the filename of the input image and extract the
                  # class label ("0" for "negative" and "1" for "positive")
                  filename = inputPath.split(os.path.sep)[-1]
                  label = filename[-5:-4]
                  # build the path to the label directory
                  labelPath = os.path.sep.join([baseOutput, label])
                  # if the label output directory does not exist, create it
                   if not os.path.exists(labelPath):
                      print("[INFO] 'creating {}' directory".format(labelPath))
                      os.makedirs(labelPath)
                  # construct the path to the destination image and then copy
                  # the image itself
                  p = os.path.sep.join([labelPath, filename])
                  shutil.copy2(inputPath, p)
```

CancerNet: Our breast cancer prediction CNN

```
In [5]: | class CancerNet:
              @staticmethod
              def build(width, height, depth, classes):
                 # initialize the model along with the input shape to be
                 # "channels last" and the channels dimension itself
                 model = Sequential()
                 inputShape = (height, width, depth)
                 chanDim = -1
                 # if we are using "channels first", update the input shape
                 # and channels dimension
                 if K.image_data_format() == "channels_first":
                     inputShape = (depth, height, width)
                     chanDim = 1
                 # CONV => RELU => POOL
                 model.add(SeparableConv2D(32, (3, 3), padding="same",
                     input_shape=inputShape))
                 model.add(Activation("relu"))
                 model.add(BatchNormalization(axis=chanDim))
                  model.add(MaxPooling2D(pool_size=(2, 2)))
                 model.add(Dropout(0.25))
                 # (CONV => RELU => POOL) * 2
                 model.add(SeparableConv2D(64, (3, 3), padding="same"))
                 model.add(Activation("relu"))
                 model.add(BatchNormalization(axis=chanDim))
                 model.add(SeparableConv2D(64, (3, 3), padding="same"))
                 model.add(Activation("relu"))
                 model.add(BatchNormalization(axis=chanDim))
                 model.add(MaxPooling2D(pool_size=(2, 2)))
                 model.add(Dropout(0.25))
                 # (CONV => RELU => POOL) * 3
                 model.add(SeparableConv2D(128, (3, 3), padding="same"))
                 model.add(Activation("relu"))
                 model.add(BatchNormalization(axis=chanDim))
                 model.add(SeparableConv2D(128, (3, 3), padding="same"))
                 model.add(Activation("relu"))
                 model.add(BatchNormalization(axis=chanDim))
                 model.add(SeparableConv2D(128, (3, 3), padding="same"))
                 model.add(Activation("relu"))
                 model.add(BatchNormalization(axis=chanDim))
                 model.add(MaxPooling2D(pool_size=(2, 2)))
                 model.add(Dropout(0.25))
                 # first (and only) set of FC => RELU layers
                 model.add(Flatten())
                 model.add(Dense(256))
                 model.add(Activation("relu"))
                 model.add(BatchNormalization())
                 model.add(Dropout(0.5))
                 # softmax classifier
                 model.add(Dense(classes))
                 model.add(Activation("softmax"))
                 # return the constructed network architecture
                 return model
```

Our training script

Found 255556 images belonging to 2 classes. Found 42596 images belonging to 2 classes.

```
In [6]: # # construct the argument parser and parse the arguments
         # ap = argparse.ArgumentParser()
         # ap.add_argument("-p", "--plot", type=str, default="plot.png",
         # help="path to output loss/accuracy plot")
         # args = vars(ap.parse_args())
         # since we are using Jupyter Notebooks we can replace our argument
         # parsing code with *hard coded* arguments and values
         args = {
             "plot": "plot.png",
In [7]: # initialize our number of epochs, initial learning rate, and batch
         # size
         NUM_EPOCHS = 40
         INIT_LR = 1e-2
         BS = 32
         # determine the total number of image paths in training, validation,
         # and testing directories
         trainPaths = list(paths.list_images(config.TRAIN_PATH))
         totalTrain = len(trainPaths)
         totalVal = len(list(paths.list_images(config.VAL_PATH)))
         totalTest = len(list(paths.list_images(config.TEST_PATH)))
         # calculate the total number of training images in each class and
         # initialize a dictionary to store the class weights
         trainLabels = [int(p.split(os.path.sep)[-2]) for p in trainPaths]
         trainLabels = to_categorical(trainLabels)
         classTotals = trainLabels.sum(axis=0)
         classWeight = dict()
         # loop over all classes and calculate the class weight
         for i in range(0, len(classTotals)):
            classWeight[i] = classTotals.max() / classTotals[i]
```

In [8]: # initialize the training data augmentation object trainAug = ImageDataGenerator(rescale=1 / 255.0, rotation_range=20, zoom_range=0.05, width_shift_range=0.1, height_shift_range=0.1, shear_range=0.05, horizontal_flip=True, vertical_flip=True, fill_mode="nearest") # initialize the validation (and testing) data augmentation object valAug = ImageDataGenerator(rescale=1 / 255.0)

In [9]: # initialize the training generator trainGen = trainAug.flow_from_directory(config.TRAIN_PATH, class_mode="categorical", target_size=(48, 48), color_mode="rgb", shuffle=True, batch_size=BS) # initialize the validation generator valGen = valAug.flow_from_directory(config.VAL_PATH, class_mode="categorical", target_size=(48, 48), color_mode="rgb", shuffle=False, batch_size=BS) # initialize the testing generator testGen = valAug.flow_from_directory(config.TEST_PATH, class_mode="categorical", target_size=(48, 48), color_mode="rgb", shuffle=False, batch_size=BS)

```
In [10]: | ### initialize our CancerNet model and compile it
      model = CancerNet.build(width=48, height=48, depth=3,
        classes=2)
      opt = Adagrad(lr=INIT_LR, decay=INIT_LR / NUM_EPOCHS)
      model.compile(loss="binary_crossentropy", optimizer=opt,
        metrics=["accuracy"])
      # fit the model
      H = model.fit(
        x=trainGen,
        steps_per_epoch=totalTrain // BS,
        validation_data=valGen,
        validation steps=totalVal // BS,
        class_weight=classWeight,
        epochs=NUM_EPOCHS)
      # save model
      model.save("Breast_cancer_model")
     c:\users\leobr\appdata\local\programs\python\python39\lib\site-packages\tensorflow\python\keras\optimizer_v2\optimizer_v2.py:374: UserWarning: The `lr` argument is deprecated, use `learning_rate` instead.
      WARNING:tensorflow:From c:\users\leobr\appdata\local\programs\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\python\pyt
     Instructions for updating:
     The `validate_indices` argument has no effect. Indices are always validated on CPU and never validated on GPU.
     Epoch 1/40
     Epoch 2/40
     Epoch 3/40
      Epoch 4/40
     Epoch 5/40
      Epoch 6/40
      Epoch 7/40
      Epoch 8/40
     Epoch 9/40
     Epoch 10/40
     Epoch 11/40
      Epoch 12/40
     Epoch 13/40
      Epoch 14/40
      Epoch 15/40
      Epoch 16/40
     Epoch 17/40
     Epoch 18/40
     Epoch 19/40
      Epoch 20/40
     Epoch 21/40
      Epoch 22/40
      Epoch 23/40
      Epoch 24/40
      Epoch 25/40
      Epoch 26/40
      Epoch 27/40
      Epoch 28/40
     Epoch 29/40
     Epoch 30/40
     Epoch 31/40
      Epoch 32/40
     Epoch 33/40
     Epoch 34/40
     Epoch 35/40
      Epoch 36/40
     Epoch 37/40
     Epoch 38/40
      Epoch 39/40
      Epoch 40/40
     INFO:tensorflow:Assets written to: Breast_cancer_model\assets
In [11]: \mid # reset the testing generator and then use our trained model to
      # make predictions on the data
      print("[INFO] evaluating network...")
      testGen.reset()
      predIdxs = model.predict(x=testGen, steps=(totalTest // BS) + 1)
      # for each image in the testing set we need to find the index of the
      # label with corresponding largest predicted probability
      predIdxs = np.argmax(predIdxs, axis=1)
      # show a nicely formatted classification report
      print(sklearn.metrics.classification_report(testGen.classes, predIdxs,
        target_names=testGen.class_indices.keys()))
     [INFO] evaluating network...
             precision recall f1-score support
                0.92
                      0.85
                           0.89
                0.69
                      0.82
                           0.75
                                28448
                           0.85 99743
       accuracy
                0.81 0.84
       macro avg
                          0.82 99743
                0.86 0.85 0.85 99743
      weighted avg
In [12]: # compute the confusion matrix and and use it to derive the raw
      # accuracy, sensitivity, and specificity
      cm = sklearn.metrics.confusion_matrix(testGen.classes, predIdxs)
      total = sum(sum(cm))
      acc = (cm[0, 0] + cm[1, 1]) / total
      sensitivity = cm[0, 0] / (cm[0, 0] + cm[0, 1])
      specificity = cm[1, 1] / (cm[1, 0] + cm[1, 1])
      # show the confusion matrix, accuracy, sensitivity, and specificity
      print(cm)
      print("acc: {:.4f}".format(acc))
      print("sensitivity: {:.4f}".format(sensitivity))
      print("specificity: {:.4f}".format(specificity))
     [[60915 10380]
      [ 5019 23429]]
      acc: 0.8456
     sensitivity: 0.8544
In [13]: # plot the training loss and accuracy
      N = NUM_EPOCHS
      plt.style.use("ggplot")
      plt.figure()
      plt.plot(np.arange(0, N), H.history["loss"], label="train_loss")
      plt.plot(np.arange(0, N), H.history["val_loss"], label="val_loss")
      plt.plot(np.arange(0, N), H.history["accuracy"], label="train_acc")
      plt.plot(np.arange(0, N), H.history["val_accuracy"], label="val_acc")
      plt.title("Training Loss and Accuracy on Dataset")
      plt.xlabel("Epoch #")
```

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plt.ylabel("Loss/Accuracy")
plt.legend(loc="lower left")
plt.savefig(args["plot"])

Found 99743 images belonging to 2 classes.

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