Corona Lung Dataset-CNN In [24]: import numpy as np import cv2 import os import warnings warnings.filterwarnings("ignore") from concurrent.futures import ThreadPoolExecutor In [25]: images=[os.path.join(os.getcwd(),"train",image) for image in os.listdir("./train/") if image.endswith("jpg") or image.endswith(".png") or image.endswith(".jpeg")] print(images[10]) covid=[] normal=[] for image in images: if image.split("\\")[-1].startswith("covid"): covid.append(image) else: normal.append(image) print(f"covid:- {covid[10]}") print(f"covid:-{normal[10]}") C:\Users\User\train\covid 108.jpg covid:- C:\Users\User\train\covid 108.jpg covid:-C:\Users\User\train\normal 108.jpeg In [13]: len(covid) 151 Out[13]: In [26]: %time print("100") Wall time: 0 ns 100 In [27]: # covid import time start=time.time() def covid_save(img, label): path=os.path.join("Data_", "train", label, img.split("\\")[-1]) img=cv2.imread(img) cv2.imwrite(f"{path}", img) for cov in covid: covid_save(cov, "cov") print(time.time()-start) 10.249018669128418 In [28]: # normal start=time.time() def normal_save(img): path=os.path.join("Data_", "train", "normal", img.split("\\")[-1]) img=cv2.imread(img) cv2.imwrite(f"{path}", img) with ThreadPoolExecutor(max_workers=100) as executor: executor.map(normal_save,normal) print(time.time()-start) 7.142252445220947 In [29]: import keras from keras.layers import Dense, Activation, Conv2D, MaxPool2D, Dropout from keras.models import Sequential from keras.optimizers import RMSprop, SGD, Adam from keras.preprocessing import image from keras.layers import BatchNormalization, Flatten from keras.layers import ZeroPadding2D #DataGenerator train_gen=image.ImageDataGenerator(rescale=1./255, featurewise_center=True, samplewise_center=True, featurewise_std_normalization=True, samplewise_std_normalization=True, zca_whitening=False, zca_epsilon=1e-06, rotation_range=40, width_shift_range=0.0, height_shift_range=0.0, brightness_range=(0.2, 0.2), shear_range=0.2, zoom_range=0.2, channel_shift_range=0.0, fill_mode='nearest', cval=0.0, horizontal_flip=True, vertical_flip=True, data_format=None, validation_split=0.0, dtype=None) val_gen=image.ImageDataGenerator(rescale=1./255, featurewise_center=True, samplewise_center=True, featurewise_std_normalization=True, samplewise_std_normalization=True, zca_whitening=False, zca_epsilon=1e-06, rotation_range=40, width_shift_range=0.0, height_shift_range=0.0, brightness_range=(0.2, 0.2), shear_range=0.2, zoom_range=0.2, channel_shift_range=0.0, fill_mode='nearest', cval=0.0, horizontal_flip=True, vertical_flip=True, data_format=None, validation_split=0.0, dtype=None) data_train=train_gen.flow_from_directory(r"C:\Users\User\Data\train\\", target_size=(32,32), classes=["covid", "dog"], class_mode="categorical", batch_size=64, seed=1) #Data val data_val=val_gen.flow_from_directory(r"C:\Users\User\Data\val\\", target_size=(32,32), classes=["covid", "dog"], class_mode="categorical", batch_size=64, seed=1) #Model model=Sequential() model.add(ZeroPadding2D(padding=(2,2), input_shape=(32,32,3))) model.add(Conv2D(5, kernel_size=(3,3), strides=(1,1), padding="SAME", activation="relu")) model.add(MaxPool2D(pool_size=(2,2), strides=(1,1),padding="SAME")) model.add(BatchNormalization(axis=-1)) model.add(Dropout(0.3)) #conv2 model.add(ZeroPadding2D(padding=(2,2))) model.add(Conv2D(5, kernel_size=(3,3), strides=(1,1), padding="SAME", activation="relu")) model.add(MaxPool2D(pool_size=(2,2), strides=(1,1),padding="SAME")) model.add(BatchNormalization(axis=-1)) model.add(Dropout(0.3)) #Conv 3 model.add(ZeroPadding2D(padding=(2,2))) model.add(Conv2D(3, kernel_size=(3,3), strides=(1,1), padding="SAME", activation="relu")) model.add(MaxPool2D(pool_size=(2,2), strides=(1,1),padding="SAME")) model.add(BatchNormalization(axis=-1)) model.add(Dropout(0.3)) #conv 4 model.add(ZeroPadding2D(padding=(2,2))) model.add(Conv2D(3, kernel_size=(2,2), strides=(1,1), padding="SAME", activation="relu")) model.add(MaxPool2D(pool_size=(2,2), strides=(1,1),padding="SAME")) model.add(BatchNormalization(axis=-1)) model.add(Dropout(0.3)) #conv 5 model.add(ZeroPadding2D(padding=(2,2))) model.add(Conv2D(3, kernel_size=(2,2), strides=(1,1), padding="SAME", activation="relu")) model.add(MaxPool2D(pool_size=(2,2), strides=(1,1),padding="SAME")) model.add(BatchNormalization(axis=-1)) model.add(Dropout(0.3)) **#FLATTEN** model.add(Flatten()) #Dense 1 model.add(Dense(128, activation='relu')) model.add(BatchNormalization(axis=-1)) model.add(Dropout(0.2)) #Dense 2 model.add(Dense(64, activation='relu')) model.add(BatchNormalization(axis=-1)) model.add(Dropout(0.3)) #Dense 3 model.add(Dense(32, activation='relu')) model.add(BatchNormalization(axis=-1)) model.add(Dropout(0.5)) #Densed model.add(Dense(2, activation="sigmoid")) #model summary model.summary() #optimizers ad=Adam(1r=0.01)#compile model model.compile(optimizer=Ad, loss="categorical_crossentropy", metrics=["accuracy"]) #fitting Model history=model.fit_generator(data_train, steps_per_epcoh=100, epcohs=3, validation_data=data_val, validation_steps=5) Found 151 images belonging to 2 classes. Found 0 images belonging to 2 classes. Model: "sequential_5" Layer (type) Output Shape Param # zero_padding2d_21 (ZeroPadd (None, 36, 36, 3) ing2D) conv2d_20 (Conv2D) (None, 36, 36, 5) 140 max_pooling2d_20 (MaxPoolin (None, 36, 36, 5) 0 batch_normalization_25 (Bat (None, 36, 36, 5) 20 chNormalization) dropout_23 (Dropout) (None, 36, 36, 5) 0 zero_padding2d_22 (ZeroPadd (None, 40, 40, 5) 0 ing2D) conv2d_21 (Conv2D) (None, 40, 40, 5) 230 max_pooling2d_21 (MaxPoolin (None, 40, 40, 5) g2D) batch_normalization_26 (Bat (None, 40, 40, 5) 20 chNormalization) dropout_24 (Dropout) 0 (None, 40, 40, 5) zero_padding2d_23 (ZeroPadd (None, 44, 44, 5) 0 ing2D) conv2d_22 (Conv2D) (None, 44, 44, 3) 138 max_pooling2d_22 (MaxPoolin (None, 44, 44, 3) 0 g2D) batch_normalization_27 (Bat (None, 44, 44, 3) 12 chNormalization) dropout_25 (Dropout) (None, 44, 44, 3) 0 zero_padding2d_24 (ZeroPadd (None, 48, 48, 3) ing2D) conv2d_23 (Conv2D) (None, 48, 48, 3) 39 max_pooling2d_23 (MaxPoolin (None, 48, 48, 3) 0 batch_normalization_28 (Bat (None, 48, 48, 3) 12 chNormalization) dropout_26 (Dropout) (None, 48, 48, 3) zero_padding2d_25 (ZeroPadd (None, 52, 52, 3) 0 ing2D) conv2d_24 (Conv2D) (None, 52, 52, 3) 39 max_pooling2d_24 (MaxPoolin (None, 52, 52, 3) g2D) batch_normalization_29 (Bat (None, 52, 52, 3) 12 chNormalization) dropout_27 (Dropout) (None, 52, 52, 3) 0 flatten_4 (Flatten) (None, 8112) dense_7 (Dense) (None, 128) 1038464 batch_normalization_30 (Bat (None, 128) 512 chNormalization) dropout_28 (Dropout) (None, 128) 0 (None, 64) dense_8 (Dense) 8256 batch_normalization_31 (Bat (None, 64) 256 chNormalization) dropout_29 (Dropout) (None, 64) dense_9 (Dense) (None, 32) 2080 batch_normalization_32 (Bat (None, 32) 128 chNormalization) dropout_30 (Dropout) (None, 32) dense_10 (Dense) 66 (None, 2) ______ Total params: 1,050,424 Trainable params: 1,049,938 Non-trainable params: 486 Traceback (most recent call last) ~\AppData\Local\Temp/ipykernel_8116/950378390.py in <module> 129 130 #compile model --> 131 model.compile(optimizer=Ad, loss="categorical_crossentropy", 132 metrics=["accuracy"]) 133 NameError: name 'Ad' is not defined In [1]: import matplotlib.pyplot as plt history=model.fit() print(history.history.keys()) #summarize history for accuracy plt.plot(history.history['accuracy']) plt.plot(history.history['val_accuracy']) plt.title('model_accuracy') plt.ylabel('accuracy') plt.xlabel('epoch') plt.legend(['train','test'],loc='upper left') plt.show() **#Summarize history for loss** plt.plot(history.history['loss']) plt.plot(history.history['val_loss']) plt.title('model_loss') plt.ylabel('loss') plt.legend(["train", "test"], loc="upper left") plt.show() Traceback (most recent call last) ~\AppData\Local\Temp/ipykernel_8888/1681388505.py in <module> 1 import matplotlib.pyplot as plt ----> 2 history=model.fit() 3 print(history.history.keys()) 5 #summarize history for accuracy NameError: name 'model' is not defined In [31]: test_images=[os.path.join(os.getcwd(),"test",image) for image in os.listdir("./test/") $\textbf{if} \ \texttt{image.endswith(".jpg")} \ \textbf{or} \ \texttt{image.endswith(".png")}$ or image.endswith(".jpeg")] In [1]: import os.path if os.path.isfile("covid_normal_model.h5") is False: model.save("covid_normal_model.h5") In []: from tensor In [32]: import matplotlib.pyplot as plt classes=["covid", "normal"] for i in range(0,10): img=cv2.imread(test_images[i]) img=cv2.resize(img, (32, 32)) img=np.expand_dims(img, axis=0) print(model.predict(img), np.argmax(model.predict(img))) print(classes[np.argmax(model.predict(img))]) plt.imshow(cv2.imread(test_images[i])) plt.show() 1/1 [=======] - 0s 334ms/step [[0.6305605 0.94976306]] 1 1/1 [=======] - 0s 35ms/step normal 200 400 600 800 1000 1200 1400 0 250 500 750 1000 1250 1500 1750 1/1 [========] - 0s 33ms/step [[0.21272689 0.85885704]] 1 normal 200 400 600 800 600 800 [[0.8097797 0.9284845]] 1 normal 50 100 150 200 150 1/1 [=======] - 0s 34ms/step [[0.9941465 0.980127]] 0 1/1 [=======] - 0s 34ms/step covid 50 100 150 200 100 150 200 250 1/1 [=======] - 0s 41ms/step [[0.94431317 0.90047544]] 0 1/1 [=======] - Os 33ms/step covid 50 100 150 200 200 1/1 [=======] - 0s 35ms/step 1/1 [=======] - Os 33ms/step [[0.22269739 0.9258754]] 1 1/1 [=======] - 0s 33ms/step normal 50 100 150 200 250 100 150 200 250 1/1 [=======] - 0s 35ms/step 1/1 [======] - 0s 33ms/step [[0.31283256 0.8605097]] 1 1/1 [=======] - Os 32ms/step normal 50 100 150 200 250 150 200 250 300 1/1 [=======] - 0s 35ms/step 1/1 [=======] - 0s 35ms/step [[0.26781756 0.9589037]] 1 1/1 [=======] - 0s 35ms/step normal 50 100 150 200 250 100 150 200 250 1/1 [=======] - 0s 34ms/step 1/1 [=======] - 0s 34ms/step [[0.25042903 0.8952273]] 1 1/1 [=======] - 0s 36ms/step normal 50 100 150 200 250 300 350 400 300 200 1/1 [=======] - 0s 33ms/step [[0.942094 0.9835716]] 1 1/1 [=======] - 0s 35ms/step normal 0 -50 100 150 200 250 300 350 400 200 300 In []: