```
In [19]: from google.colab import drive
          drive.mount('/content/drive')
         Mounted at /content/drive
In [26]:
         orig = cv.imread("/content/drive/My Drive/food/training/food/856.jpg")
         # Convert image to RGB from BGR (another way is to use "image = image[:, :, ::-1]" code)
In [27]:
          orig = cv.cvtColor(orig, cv.COLOR BGR2RGB)
          # Resize image to 224x224 size
          image = cv.resize(orig, (224, 224)).reshape(-1, 224, 224, 3)
          # We need to preprocess imageto fulfill ResNet50 requirements
          image = preprocess input(image)
          # Extracting our features
          features = model.predict(image)
          features.shape
Out[27]: (1, 7, 7, 2048)
         n features = features.shape[-1]
In [28]:
          fig = plt.figure(figsize = (17, 8))
          gs = gridspec.GridSpec(1, 2, figure = fig)
          sub gs = gridspec.GridSpecFromSubplotSpec(3, 3, subplot spec=qs[1])
          ax1 = fig.add subplot(gs[0])
          ax1.imshow(orig)
          for i in range(3):
              for j in range(3):
                  ax2 = fig.add subplot(sub gs[i, j])
                  plt.axis('off')
                  plt.imshow(features[0, :, :, np.random.randint(n features)], cmap = 'gray')
```



```
for s, d in (zip(sets, data)):
             # Defining path to categories ['non food', 'food']
              path to cat = os.path.join(path, s)
              # Loop through categories in split
             for cat in os.listdir(path to cat):
                  # Defining path to images in category
                  path to images = os.path.join(path to cat, cat)
                  # Defining labels
                  if cat == 'food':
                      label = 1
                  else:
                      label = 0
                  # Loop through images in category
                  for i in os.listdir(path to images):
                     # Path to image
                     image path = os.path.join(path to images, i)
                      # Reading and preprocessing image
                      image = cv.imread(image path)
                      image = cv.cvtColor(image, cv.COLOR BGR2RGB)
                      image = cv.resize(image, (224, 224)).reshape(-1, 224, 224, 3)
                      image = preprocess input(image)
                      # Extracting features
                     features = model.predict(image).reshape(100352)
                      # Store features and label in our lists
                      d[0].append(features)
                      d[1].append(label)
         # Shuffle data in each split
In [31]:
          random state = 666
          train X, train Y = shuffle(train X, train Y, random state = random state)
          val X, val Y = shuffle(val X, val Y, random state = random state)
          eval X, eval Y = shuffle(eval X, eval Y, random state = random state)
         # Convert data to numpy arrays
In [32]:
          train X = np.array(train X)
```

```
train Y = np.array(train_Y)
        val X = np.array(val X)
        val Y = np.array(val Y)
        eval X = np.array(eval X)
        eval Y = np.array(eval Y)
In [35]: # Creating model
        # Number of nodes were defined using rule:
        # take the square root of the previous number of nodes in the layer and then find the closest power of 2
        # (Before FC layers in ResNet50 we have 7*7*2048 = 100352 nodes (after flatten layer), so if we take square root from
        # this number we get 316.78 and closes power of 2 is 256, it's number of nodes in 1st FC layer in our network,
        # to define number of nodes in second layer we are getting swuare root from 256 = 16)
        # Number of nodes in last layer = 1, because it's binary classification problem and our labels have only two values (
        model = Sequential()
        model.add(Dense(256, input shape = (100352,), activation = 'relu', kernel initializer = 'he normal'))
        model.add(Dense(16, activation = 'relu', kernel initializer = 'he normal'))
        model.add(Dense(1, activation = 'sigmoid'))
        # Checkpoint to save best model
        checkpoint = ModelCheckpoint('best model.hdf5', monitor = 'val accuracy', verbose = 1, save best only = True)
        model.compile(optimizer = 'adam', loss = 'binary crossentropy', metrics = ['accuracy'])
        model.fit(train X, train Y, batch size = 64, epochs = 10, validation data = (val X, val Y), callbacks = [checkpoint])
        Epoch 1/10
       uracy: 0.9760
       Epoch 00001: val accuracy improved from -inf to 0.97600, saving model to best model.hdf5
       Epoch 2/10
       uracy: 0.9670
       Epoch 00002: val accuracy did not improve from 0.97600
       Epoch 3/10
       uracy: 0.9780
       Epoch 00003: val accuracy improved from 0.97600 to 0.97800, saving model to best model.hdf5
       Epoch 4/10
```

```
accuracy: 0.9790
    Epoch 00004: val accuracy improved from 0.97800 to 0.97900, saving model to best model.hdf5
    Epoch 5/10
    uracy: 0.9740
    Epoch 00005: val accuracy did not improve from 0.97900
    Epoch 6/10
    uracy: 0.9770
    Epoch 00006: val accuracy did not improve from 0.97900
    Epoch 7/10
    uracy: 0.9780
    Epoch 00007: val accuracy did not improve from 0.97900
    Epoch 8/10
    uracy: 0.9770
    Epoch 00008: val accuracy did not improve from 0.97900
    Epoch 9/10
    accuracy: 0.9730
    Epoch 00009: val accuracy did not improve from 0.97900
    Epoch 10/10
    uracy: 0.9750
    Epoch 00010: val accuracy did not improve from 0.97900
Out[35]: <keras.callbacks.History at 0x7fa3b9f10a50>
     # Plot losses and accuracies
In [36]:
     history = model.history.history
     fig = plt.figure(figsize = (8, 5))
     plt.plot(history['accuracy'], label = 'acc')
     plt.plot(history['val accuracy'], label = 'val acc')
     plt.plot(history['loss'], label = 'acc')
```

```
plt.plot(history['val loss'], label = 'acc')
         plt.grid()
         plt.legend()
         plt.show()
                                                         acc
                                                         val acc
                                                         acc
        2.0
                                                         acc
        1.5
        1.0
        0.5
        0.0
         # Loading best model and evaluate it
In [37]:
         # Here I want to compare keras.evaluate and sklearn.accuracy score methods
         model.load weights('best model.hdf5')
         model.evaluate(eval X, eval Y)
        Out[37]: [0.7975539565086365, 0.98333333492279053]
In [38]:
         # Making predictions using evaluation dataset
         preds = model.predict(eval X)
         # If our prediction more than 0.5 - we round it to 1, else to 0
         preds = [1 if i > 0.5 else 0 for i in preds]
         # Calculating accuracy score
         accuracy = accuracy score(eval Y, preds)
```

print(f'Accuracy: {round(accuracy \* 100, 4)}%')

```
# Plotting confusion matrix
l = ['non_food', 'food']
confusion = confusion_matrix(eval_Y, preds)
sns.heatmap(confusion, square = True, fmt = 'd', xticklabels = l, yticklabels = l, annot = True)
```

Accuracy: 98.3333%

Out[38]: <matplotlib.axes.\_subplots.AxesSubplot at 0x7fa3b4423850>

