

```
pwd
```

Out[1]:

```
'/home/wsuser/work'
```

In [2]:

```
import warnings
warnings.filterwarnings("ignore")
```

In [3]:

```
!pip install imutils

Collecting imutils
  Downloading imutils-0.5.4.tar.gz (17 kB)
Building wheels for collected packages: imutils
  Building wheel for imutils (setup.py) ... done
  Created wheel for imutils: filename=imutils-0.5.4-py3-none-any.whl size=2
5860 sha256=9b27083f9cc4fd096c0ed96e02d664a4fd9cbef45c07309a802fd276271febc
4
  Stored in directory: /tmp/wsuser/.cache/pip/wheels/4b/a5/2d/4a070a801d3a3
d93f033d3ee9728f470f514826e89952df3ea
Successfully built imutils
Installing collected packages: imutils
Successfully installed imutils-0.5.4
```

## Image Pre-processing

### Importing the necessary libraries

In [4]:

```
import cv2
import numpy as np
from sklearn.preprocessing import LabelEncoder
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import confusion_matrix
from skimage import feature
from imutils import paths
import os
import pickle
```

### Functions to load and quantify the images

In [5]:

```
import os, types
import pandas as pd
from botocore.client import Config
import ibm_boto3

def __iter__(self): return 0

# @hidden_cell
# The following code accesses a file in your IBM Cloud Object Storage. It
# includes your credentials.
# You might want to remove those credentials before you share the notebook.
cos_client = ibm_boto3.client(service_name='s3',
                              ibm_api_key_id='rlyEPAR18kPTMGZRBByQGAWk_2w81tdr_D7ky9zEjT6ja',
                              ibm_auth_endpoint="https://iam.cloud.ibm.com/oidc/token",
```

```

        config=Config(signature_version='oauth'),
        endpoint_url='https://s3.private.us.cloud-object-
storage.appdomain.cloud')

bucket = 'parkinson39sdiseasedetection-donotdelete-pr-9rhldkrukkwg0y'
object_key = 'dataset.zip'

streaming_body_1 = cos_client.get_object(Bucket=bucket,
Key=object_key)['Body']

# Your data file was loaded into a botocore.response.StreamingBody object.
# Please read the documentation of ibm_boto3 and pandas to learn more about
the possibilities to load the data.
# ibm_boto3 documentation: https://ibm.github.io/ibm-cos-sdk-python/
# pandas documentation: http://pandas.pydata.org/

```

In [6]:

```

from io import BytesIO
import zipfile
unzip = zipfile.ZipFile(BytesIO(streaming_body_1.read()), 'r')
file_paths = unzip.namelist()
for path in file_paths:
    unzip.extract(path)

```

In [7]:

```
pwd
```

Out[7]:

```
'/home/wsuser/work'
```

In [8]:

```

def quantify_image(image):
    features = feature.hog(image,
                           orientations=9,
                           pixels_per_cell=(5,5),
                           cells_per_block=(2,2),
                           transform_sqrt=True,
                           block_norm="L1")

    return features

```

In [9]:

```

def load_split(path):
    path_images = list(paths.list_images(path))
    data=[]
    labels=[]

    for path_image in path_images:
        label = path_image.split(os.path.sep)[-2]
        image = cv2.imread(path_image)
        image = cv2.cvtColor(image, cv2.COLOR_BGR2GRAY)
        image = cv2.resize(image, (200,200))
        image = cv2.threshold(image, 0, 225, cv2.THRESH_BINARY_INV |
cv2.THRESH_OTSU) [1]

        features = quantify_image(image)
        data.append(features)
        labels.append(label)

    return (np.array(data), np.array(labels))

```

## Using spiral images

### Defining the path for training data and testing data

```
path_training_data = r"/home/wsuser/work/dataset/spiral/training"
path_testing_data = r"/home/wsuser/work/dataset/spiral/testing"
```

In [10]:

### Loading the training and testing data

```
(x_train, y_train) = load_split(path_training_data)
(x_test, y_test) = load_split(path_testing_data)
```

In [11]:

### Label Encoding

```
label_encoder = LabelEncoder()
y_train = label_encoder.fit_transform(y_train)
y_test = label_encoder.transform(y_test)
print(x_train.shape, y_train.shape)
# 0:healthy,1:Parkinson
(72, 54756) (72,)
```

In [12]:

## Building the model

### Training the model

```
model = RandomForestClassifier(n_estimators=100)
model.fit(x_train, y_train)
```

In [13]:

```
RandomForestClassifier()
```

Out[13]:

### Testing the model

```
testingPaths = list(paths.list_images(path_testing_data))
idxs = np.arange(0, len(testingPaths))
idxs = np.random.choice(idxs, size=(25,), replace=False)
images = []
```

In [14]:

```
for i in idxs:
    # loading the testing image, clone it, and resize it
    image = cv2.imread(testingPaths[i])
    output = image.copy()
    output = cv2.resize(output, (128, 128))

    # pre-processing the image
    image = cv2.cvtColor(image, cv2.COLOR_BGR2GRAY)
    image = cv2.resize(image, (200, 200))
```

In [15]:

```

    image = cv2.threshold(image, 0, 255, cv2.THRESH_BINARY_INV |
cv2.THRESH_OTSU) [1]

    # quantify the image and make predictions based on the extracted
    # features using the last trained Random Forest
    features = quantify_image(image)
    preds = model.predict([features])

    label = label_encoder.inverse_transform(preds)[0]

    # draw the colored class label on the output image and add it to the
    set of output images
    if label == "healthy":
        color = (0, 255, 0)
    else:
        color = (0, 0, 255)
    cv2.putText(output, label, (3, 20), cv2.FONT_HERSHEY_SIMPLEX, 0.5,
color, 2)
    images.append(output)

```

## Evaluating the model

```

In [16]:
predictions = model.predict(x_test)    # predictions on the testing data

cm = confusion_matrix(y_test, predictions).flatten ()    # computing the
confusion matrix
print(cm)
(tn, fp, fn, tp) = cm

accuracy = (tp + tn) / float(cm.sum())    # computing the accuracy
print(accuracy)

[12  3  5 10]
0.7333333333333333

```

## Saving the model

```

In [17]:
pickle.dump(model, open('parkinson.pkl', 'wb'))

In [18]:
!tar -zcvf parkinsonsmodel.tgz parkinson.pkl
parkinson.pkl

In [19]:

ls
dataset/ parkinson.pkl parkinsonsmodel.tgz

In [20]:
!pip install watson-machine-learning-client --upgrade
from ibm_watson_machine_learning import APIClient
wml_credentials = {
    "url" : "https://us-south.ml.cloud.ibm.com",

```

```

        "apikey" :
        "K36t7CdPrC_R4VWU_knCWisv6ePwSofTJbYqDvwtHL54"
    }
client = APIClient(wml_credentials)

```

In [22]:

```

def guid_from_space_name(client, space_name):
    space = client.spaces.get_details()
    print(space)
    return (next(item for item in space['resources'] if
item['entity']['name']==space_name) ['metadata']['id'])

```

In [23]:

```

space_uid = guid_from_space_name(client, 'ParkinsonsDiseaseDetection')
print('Space UID = ' + space_uid)
client.set.default_space(space_uid)

```

Out[24]:

```
'SUCCESS'
```

In [25]:

```
client.software_specifications.list()
```

NAME	ASSET_ID	TYPE
default_py3.6	0062b8c9-8b7d-44a0-a9b9-46c416adcbd9	base
kernel-spark3.2-scala2.12	020d69ce-7ac1-5e68-ac1a-31189867356a	base
pytorch-onnx_1.3-py3.7-edt	069ea134-3346-5748-b513-49120e15d288	base
scikit-learn_0.20-py3.6	09c5a1d0-9c1e-4473-a344-eb7b665ff687	base
spark-mllib_3.0-scala_2.12	09f4cff0-90a7-5899-b9ed-1ef348aebdee	base
pytorch-onnx_rt22.1-py3.9	0b848dd4-e681-5599-be41-b5f6fccc6471	base
ai-function_0.1-py3.6	0cdb0f1e-5376-4f4d-92dd-da3b69aa9bda	base
shiny-r3.6	0e6e79df-875e-4f24-8ae9-62dcc2148306	base
tensorflow_2.4-py3.7-horovod	1092590a-307d-563d-9b62-4eb7d64b3f22	base
pytorch_1.1-py3.6	10ac12d6-6b30-4ccd-8392-3e922c096a92	base
tensorflow_1.15-py3.6-ddl	111e41b3-de2d-5422-a4d6-bf776828c4b7	base
runtime-22.1-py3.9	12b83a17-24d8-5082-900f-0ab31fbfd3cb	base
scikit-learn_0.22-py3.6	154010fa-5b3b-4ac1-82af-4d5ee5abbc85	base
default_r3.6	1b70aec3-ab34-4b87-8aa0-a4a3c8296a36	base
pytorch-onnx_1.3-py3.6	1bc6029a-cc97-56da-b8e0-39c3880dbbe7	base
kernel-spark3.3-r3.6	1c9e5454-f216-59dd-a20e-474a5cdf5988	base
pytorch-onnx_rt22.1-py3.9-edt	1d362186-7ad5-5b59-8b6c-9d0880bde37f	base
tensorflow_2.1-py3.6	1eb25b84-d6ed-5dde-b6a5-3fbdf1665666	base
spark-mllib_3.2	20047f72-0a98-58c7-9ff5-a77b012eb8f5	base
tensorflow_2.4-py3.8-horovod	217c16f6-178f-56bf-824a-b19f20564c49	base
runtime-22.1-py3.9-cuda	26215f05-08c3-5a41-a1b0-da66306ce658	base
do_py3.8	295addb5-9ef9-547e-9bf4-92ae3563e720	base
autoai-ts_3.8-py3.8	2aa0c932-798f-5ae9-abd6-15e0c2402fb5	base
tensorflow_1.15-py3.6	2b73a275-7cbf-420b-a912-eae7f436e0bc	base
kernel-spark3.3-py3.9	2b7961e2-e3b1-5a8c-a491-482c8368839a	base
pytorch_1.2-py3.6	2c8ef57d-2687-4b7d-acce-01f94976dac1	base
spark-mllib_2.3	2e51f700-bca0-4b0d-88dc-5c6791338875	base
pytorch-onnx_1.1-py3.6-edt	32983cea-3f32-4400-8965-dde874a8d67e	base
spark-mllib_3.0-py37	36507ebe-8770-55ba-ab2a-eafe787600e9	base
spark-mllib_2.4	390d21f8-e58b-4fac-9c55-d7ceda621326	base
xgboost_0.82-py3.6	39e31acd-5f30-41dc-ae44-60233c80306e	base
pytorch-onnx_1.2-py3.6-edt	40589d0e-7019-4e28-8daa-fb03b6f4fe12	base
default_r36py38	41c247d3-45f8-5a71-b065-8580229facf0	base
autoai-ts_rt22.1-py3.9	4269d26e-07ba-5d40-8f66-2d495b0c71f7	base

autoai-obm_3.0	42b92e18-d9ab-567f-988a-4240baled5f7	base
pmml-3.0_4.3	493bcb95-16f1-5bc5-bee8-81b8af80e9c7	base
spark-mllib_2.4-r_3.6	49403dff-92e9-4c87-a3d7-a42d0021c095	base
xgboost_0.90-py3.6	4ff8d6c2-1343-4c18-85e1-689c965304d3	base
pytorch-onnx_1.1-py3.6	50f95b2a-bc16-43bb-bc94-b0bed208c60b	base
autoai-ts_3.9-py3.8	52c57136-80fa-572e-8728-a5e7cbb42cde	base
spark-mllib_2.4-scala_2.11	55a70f99-7320-4be5-9fb9-9edb5a443af5	base
spark-mllib_3.0	5c1b0ca2-4977-5c2e-9439-ffd44ea8ffe9	base
autoai-obm_2.0	5c2e37fa-80b8-5e77-840f-d912469614ee	base
spss-modeler_18.1	5c3cad7e-507f-4b2a-a9a3-ab53a21dee8b	base
cuda-py3.8	5d3232bf-c86b-5df4-a2cd-7bb870a1cd4e	base
autoai-kb_3.1-py3.7	632d4b22-10aa-5180-88f0-f52dfb6444d7	base
pytorch-onnx_1.7-py3.8	634d3cdc-b562-5bf9-a2d4-ea90a478456b	base
spark-mllib_2.3-r_3.6	6586b9e3-ccd6-4f92-900f-0f8cb2bd6f0c	base
tensorflow_2.4-py3.7	65e171d7-72d1-55d9-8ebb-f813d620c9bb	base
spss-modeler_18.2	687eddc9-028a-4117-b9dd-e57b36f1efa5	base

-----

Note: Only first 50 records were displayed. To display more use 'limit' parameter.

In [26]:

```
software_spec_uid=client.software_specifications.get_uid_by_name("default_py3.6")
software_spec_uid
```

Out[26]:

```
'0062b8c9-8b7d-44a0-a9b9-46c416adcbd9'
```

In [ ]:

```
model_details =
client.repository.store_model(model='parkinsonsmodel.tgz',meta_props={
client.repository.ModelMetaNames.NAME: "parkinson",
client.repository.ModelMetaNames.TYPE: "default_py3.6",
client.repository.ModelMetaNames.SOFTWARE_SPEC_UID: software_spec_uid})
model_id = client.repository.get_model_uid(model_details)
```

## Using wave images

### Defining the path for wave training data and testing data

In [58]:

```
path_training_data = r"dataset/wave/training"
path_testing_data = r"dataset/wave/testing"
```

### Loading the training and testing data

In [61]:

```
(x_train, y_train) = load_split(path_training_data)
(x_test, y_test) = load_split(path_testing_data)
```

### Label Encoding

In [62]:

```
label_encoder = LabelEncoder()
y_train = label_encoder.fit_transform(y_train)
y_test = label_encoder.transform(y_test)
print(x_train.shape, y_train.shape)
```

```
# 0:healthy,1:Parkinson
(72, 54756) (72,)
```

## Building the model

### Training the model

```
model = RandomForestClassifier(n_estimators=100)
model.fit(x_train, y_train)
```

In [63]:

```
RandomForestClassifier()
```

Out[63]:

### Testing the model

```
testingPaths = list(paths.list_images(path_testing_data))
idxs = np.arange(0, len(testingPaths))
idxs = np.random.choice(idxs, size=(25,), replace=False)
images = []
```

In [64]:

```
for i in idxs:
    # loading the testing image, clone it, and resize it
    image = cv2.imread(testingPaths[i])
    output = image.copy()
    output = cv2.resize(output, (128, 128))

    # pre-processing the image
    image = cv2.cvtColor(image, cv2.COLOR_BGR2GRAY)
    image = cv2.resize(image, (200, 200))
    image = cv2.threshold(image, 0, 255, cv2.THRESH_BINARY_INV |
cv2.THRESH_OTSU) [1]

    # quantify the image and make predictions based on the extracted
    # features using the last trained Random Forest
    features = quantify_image(image)
    preds = model.predict([features])

    label = label_encoder.inverse_transform(preds)[0]

    # draw the colored class label on the output image and add it to the
    set of output images
    if label == "healthy":
        color = (0, 255, 0)
    else:
        color = (0, 0, 255)
    cv2.putText(output, label, (3, 20), cv2.FONT_HERSHEY_SIMPLEX, 0.5,
color, 2)
    images.append(output)
```

In [65]:

### Evaluating the model

In [66]:

```
predictions = model.predict(x_test)    # predictions on the testing data

cm = confusion_matrix(y_test, predictions).flatten ()    # computing the
confusion matrix
print(cm)
(tn, fp, fn, tp) = cm

accuracy = (tp + tn) / float(cm.sum())    # computing the accuracy
print(accuracy)
[10  5  5 10]
0.6666666666666666
```

## Saving the model

In [67]:

```
pickle.dump(model, open('parkinson_w.pkl', 'wb'))
```

In [68]:

```
!tar -zcvf parkinsons-detection-model_s.tgz parkinson_w.pkl
parkinson_w.pkl
```

In [69]:

```
ls
dataset/      parkinsons-detection-model_s.tgz  parkinson_w.pkl
parkinson.pkl  parkinsonsmodel.tgz
```