intro-to-ecg

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1 ECG Library

1.1 Suggested pipeline

1.1.1 Load ECG images

```
[4]: import os
     from PIL import Image
[5]: dirname = 'examples/example_images/'
[6]: images = os.listdir(dirname)
[7]: images = sorted(images, key=lambda a: int(a.split('.')[0]))
[8]:
    images
[8]: ['1.png',
      '2.png',
      '3.png',
      '4.png',
      '5.png',
      '6.png',
      '7.png',
      '8.png',
      '9.png',
      '10.png',
      '11.png',
      '12.png']
[9]: images = [Image.open(dirname + filename) for filename in images]
[7]:
     images[1]
[7]:
```

1.1.2 Convert images to signal

```
[2]: import ECG.api as api import numpy as np
```

```
[10]: signal = [api.convert_image_to_signal(np.asarray(image)[:,:,:3]) for image in____
images]
```

1.1.3 display explanations

```
[11]: def display_text_explanation(explanation:api.TextExplanation):
    print('Text explanation:', explanation.content)

def display_text_and_image_explanation(explanation:api.TextAndImageExplanation):
    print('Text explanation:', explanation.text)
    print('GradCAM visualization:')
    return explanation.image
```

1.1.4 Get signal of all 12 ECG leads

Recommended sampling rate is 500

```
[12]: sampling_rate = 500
```

```
[13]: mm_per_mv = 10
```

```
[14]: signal = np.asarray([i / mm_per_mv for i in signal])
```

```
[15]: assert len(signal.shape) == 2
assert signal.shape[0] == 12
```

1.1.5 Check whether ST-elevation is present

```
[16]: res = api.check_ST_elevation(signal, sampling_rate=sampling_rate)
print('Result:', res[0])
print('Explanation:')
display_text_explanation(res[1])
```

Result: ElevatedST.Abscent

Explanation:

Text explanation: ST elevation value in lead V3 (0.04126003416910934 mV) did not exceed the threshold 0.2, therefore ST elevation was not detected.

```
[17]: res = api.check_ST_elevation_with_NN(signal)
      print('Result:', res[0])
      print('Explanation:')
      display_text_and_image_explanation(res[1])
     Load model at ./ECG/NN_based_approach/Models/Conv1_ste_model.pt
     Result: ElevatedST.Abscent
     Explanation:
     Text explanation: Significant ST elevation probability is 0.3325
     GradCAM visualization:
「17]:
     1.1.6 Evaluate risk markers
[18]: api.evaluate_risk_markers(signal, sampling_rate=sampling_rate)
[18]: RiskMarkers(Ste60_V3=0.04126003416910934, QTc=417, RA_V4=0.9481276953640756)
     1.1.7 Perform differential diagnosis
[19]: res = api.diagnose_with_risk_markers(signal, sampling_rate=sampling_rate)
      print('Result:', res[0])
      print('Explanation:')
      display_text_explanation(res[1])
     Result: Diagnosis.BER
     Explanation:
     Text explanation: Criterion value calculated as follows: (1.196 * [STE60 V3 in
     mm]) + (0.059 * [QTc in ms]) - (0.326 * [RA V4 in mm])) = 22.00557372177566 did
     not exceed the threshold 23.4, therefore the diagnosis is Benign Early
     Repolarization
[20]: res = api.check_BER_with_NN(signal)
      print('Result:', res[0])
      print('Explanation:')
      display_text_and_image_explanation(res[1])
     Load model at ./ECG/NN_based_approach/Models/Conv_ber_model.pt
     Result: True
     Explanation:
     Text explanation: BER probability is 0.9278
     GradCAM visualization:
[20]:
```

```
[21]: res = api.check_MI_with_NN(signal)
    print('Result:', res[0])
    print('Explanation:')
    display_text_and_image_explanation(res[1])

Load model at ./ECG/NN_based_approach/Models/Conv_mi_model.pt
    Result: False
    Explanation:
    Text explanation: MI probability is 0.0017
    GradCAM visualization:
[21]:
```

1.1.8 Check if the ECG is normal or not

```
[25]: from ECG.ecghealthcheck.enums import ECGClass

[27]: signal = signal[:, :4000]
    signal.shape

[27]: (12, 4000)

[29]: res = api.check_ecg_is_normal(signal, ECGClass.ALL)
    print('Result:', res[0])
    display_text_explanation(res[1])
```

Result: True

Text explanation: The signal is ok