





Project Goal

使用MIT-BIT的DATA SET, 透過深度學習做出 判斷心律不整的模型。

訓練與測試

Reference annotations	Signals	Header	
100.atr	100.dat	100.hea	
101.atr	<u>101.dat</u>	101.hea	
102.atr	102.dat	102.hea	
103.atr	103.dat	103.hea	
104.atr	<u>104.dat</u>	104.hea	
105.atr	105.dat	105.hea	
106.atr	106.dat	106.hea	
107.atr	107.dat	107.hea	
108.atr	108.dat	108.hea	
109.atr	109.dat	109.hea	
111.atr	111.dat	111.hea	

https://www.physionet.org/physiobank/database/mitdb/ 總共48筆資料,取前38筆為訓練資料 後10筆為測試資料 每人於6/26號期末展示結果

MIT-BIH Dataset

美國麻省理工學院提供, 研究心律失常的數據庫。 目前國際上公認可作為標準 的心電數據庫之一。

MIT-BIH ARRHYTHMIA DATABASE

This database is described in

Moody GB, Mark RG. The impact of the MIT-BIH Arrhythmia Database. *IEEE Eng in Med and Biol* 20(3):45-50 (May-June 2001). (PMID: 11446209)

Please cite this publication when referencing this material, and also include the standard citation for PhysioNet:

Goldberger AL, Amaral LAN, Glass L, Hausdorff JM, Ivanov PCh, Mark RG, Mietus JE, Moody GB, Peng C-K, Stanley HE. PhysioBank, PhysioToolkit, and PhysioNet: Components of a New Research Resource for Complex Physiologic Signals. *Circulation* 101(23):e215-e220 [Circulation Electronic Pages; http://circ.ahajournals.org/content/101/23/e215.full

[☑]; 2000 (June 13).



MIT-BIH Dataset

麻煩到令人想放棄的資料集

MIT-BIH 的數據格式:

- MIT-BIH 為了節省文件長度和存儲空間,使用了自定義的格式。一個心電記錄由三個部分組成:
- (1)頭文件[.hea],存儲方式ASCII碼字符。
- (2)數據文件[.dat],按三進制存儲,每三個字節存儲兩個數,一個數12bit。
- (3) 註釋文件[.atr],按二進制存儲。

```
000000000h: 23
             33
                F3
                       33
                          F3 E3 33
                                   F3
                                      E3 33
                    33
                                   E8 33 FO
                                   33 F2 E0
                                33 F6 E1 33
00000030h: DC 33 F5 DB 33 F6 DE
           33 F5 DA 33 F1 DC 33 F0 E1 33 EF
                                            E.5
              DA 33
           CE 33 EA CC
                          E8
                                33
                                   E3
                       33
                             C9
```

看到數據集後的反應

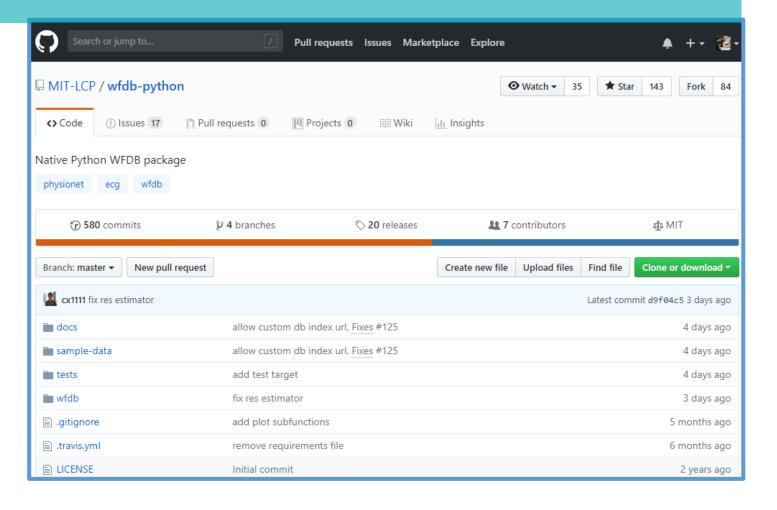


上帝為你關了一扇門

那是因為他要你吹冷氣

冷氣來啦! WFDB

waveform-database (WFDB) package



Introduction

The native Python waveform-database (WFDB) package.

A library of tools for reading, writing, and processing WFDB signals and annotations.

Core components of this package are based on the original WFDB specifications. This package does not contain the exact same functionality as the original WFDB package. It aims to implement as many of its core features as possible, with user-friendly APIs.

Additional useful physiological signalprocessing tools are added over time.

CODE片段

```
# Load Data
 In [5]:
         record = wfdb.rdsamp('mitdb/100',)
         annotation = wfdb.rdann('mitdb/100', 'atr')
         record
Out[5]: (array([[-0.145, -0.065],
              [-0.145, -0.065],
              [-0.145, -0.065],
              [-0.675, -0.365],
              [-0.765, -0.335],
              [-1.28, 0. ]]),
         {'base date': None,
          'base time': None,
          'comments': ['69 M 1085 1629 x1', 'Aldomet, Inderal'],
          'fs': 360.
          'n sig': 2,
          'sig len': 650000,
          'sig name': ['MLII', 'V5'],
          'units': ['mV', 'mV']})
```

Introduction to Data Structure of WFDB

• record: 記錄.dat檔案中的數據,每 筆.dat檔案有兩個探測點(各蒐集65萬筆 資料)

- annotation: 紀錄.atr檔案中的資料,該 筆資料包含了兩個部分:
 - 1.採樣點位置

6 6

2.該位置的症狀判斷



註釋代碼

註釋代碼		說明	備註
0		No TQRS	
1	N	Normal beat	正常搏動
2	L	Left bundle branch block beat	左束支傳導阻滯
3	R	Right bundle branch block beat	右束支傳導阻滯
4	a	Aberrated atrial premature beat	異常 <u>房性早搏</u>
5	V	Premature	室性早搏
		ventricular contraction	
6	F	Fusion of ventricular and normal beat	心室融合心跳
7	J	Nodal (junctional) premature beat	<u>交界性早搏</u>
8	A	Atrial premature beat	<u>房性早搏</u>
9	S	Premature or ectopic supraventricular beat	室上性 <u>早搏</u> 或異常
10	Е	Ventricular escape beat	室性 <u>逸搏</u>

Introduction to Data Structure of WFDB

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• record: 記錄.dat檔案中的數據,每 筆.dat檔案有兩個探測點(各蒐集65萬筆 資料)

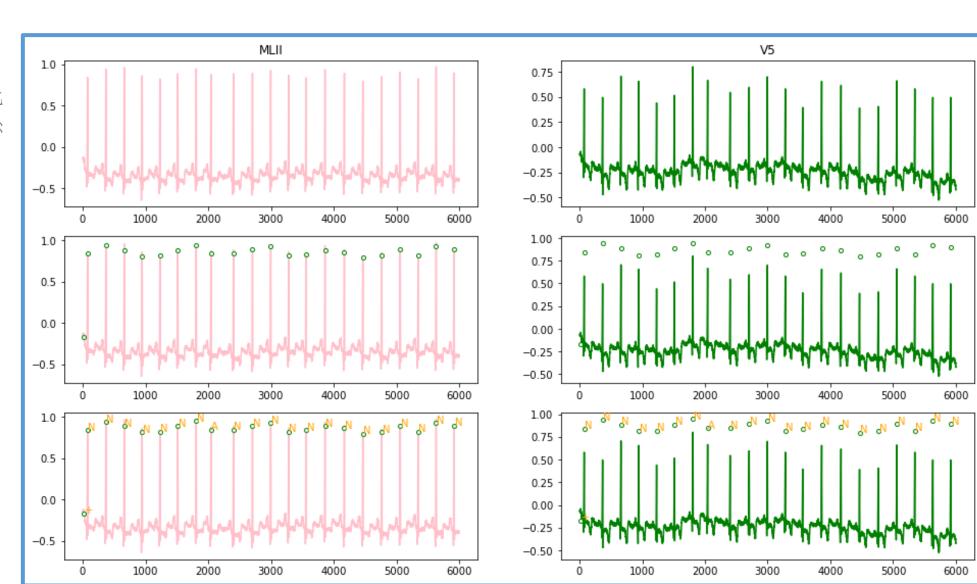
- annotation: 紀錄.atr檔案中的資料,該 筆資料包含了兩個部分:
 - 1.採樣點位置
 - 2.該位置的症狀判斷



Data

Visualization

透過資料視覺化 觀察數據與標記的關係



Label

statistical analysis

透過資料視覺化觀察數據與標記的關係

We need: N. L. R. a. V

```
In [28]:
         for file name in train file name:
            # Load file
            record = wfdb.rdsamp('mitdb/' + file name,)
            annotation = wfdb.rdann('mitdb/' + file name, 'atr')
            print('file name:', file name, '
                                                     describes: ',describe annotation(annotation))
         file name: 100
                                   describes: {'N': 2239, '+': 1, 'V': 1, 'A': 33}
         file name: 101
                                   describes: {'Q': 2, 'A': 3, '+': 1, '~': 4, '|': 4, 'N': 1860}
         file name: 102
                                   describes: {'/': 2028, 'N': 99, '+': 5, 'V': 4, 'f': 56}
         file name: 103
                                   describes: {'N': 2082, '+': 1, 'A': 2, '~': 6}
                                   describes: {'f': 666, '+': 45, 'Q': 18, '~': 37, '/': 1380, 'V': 2, 'N': 163}
         file name: 104
         file name: 105
                                   describes: {'+': 1, '~': 88, '|': 30, 'Q': 5, 'V': 41, 'N': 2526}
         file name: 106
                                   describes: {'N': 1507, '+': 41, 'V': 520, '~': 30}
         file name: 107
                                   describes: {'/': 2078, 'V': 59, '+': 1, '~': 2}
         file name: 108
                                   describes: {'F': 2, 'x': 11, '+': 1, 'A': 4, '~': 41, '|': 8, 'V': 17, 'N': 1739, 'j': 1}
         file name: 109
                                   describes: {'F': 2, 'V': 38, '+': 1, 'L': 2492, '~': 2}
         file name: 111
                                   describes: {'V': 1, '+': 1, 'L': 2123, '~': 8}
         file name: 112
                                   describes: {'N': 2537, '+': 1, 'A': 2, '~': 10}
         file name: 113
                                   describes: {'N': 1789, '+': 1, 'a': 6}
         file name: 114
                                   describes: {'F': 4, '+': 3, 'J': 2, 'A': 10, '~': 7, '|': 1, 'V': 43, 'N': 1820}
         file name: 115
                                   describes: {'|': 6, 'N': 1953, '+': 1, '~': 2}
         file name: 116
                                   describes: {'N': 2302, '+': 1, 'V': 109, 'A': 1, '~': 8}
         file name: 117
                                   describes: {'N': 1534, '+': 1, 'A': 1, '~': 3}
                                   describes: {'x': 10, '+': 1, 'R': 2166, 'A': 96, '~': 12, 'V': 16}
         file name: 118
         file name: 119
                                   describes: {'N': 1543, '+': 103, 'V': 444, '~': 4}
          file name: 121
                                    docoriboo: (INII: 1061 Ltl: 1 IV/: 1 IAI: 1 Ltl: 10)
```



Data Preprocessing

先對資料作一些初步的處理

数據載人

透過wfdb可以很方便地將所有 MIT-BIH資料集下載並讀取。 欄位

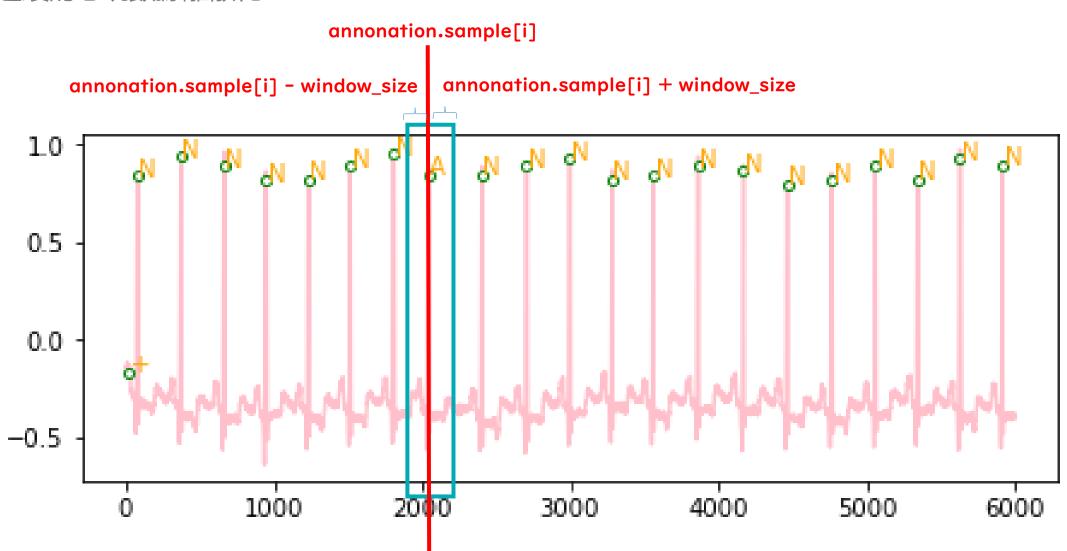
每一筆.dat資料存有兩個不同部位 的測量數據,在此只採用其中一筆 作為data。

亂數 排序 資料集的分布較不平均,如果只取 前面作為訓練資料可能會有問題, 所以一次性讀取後再打散排序。 資料 切割

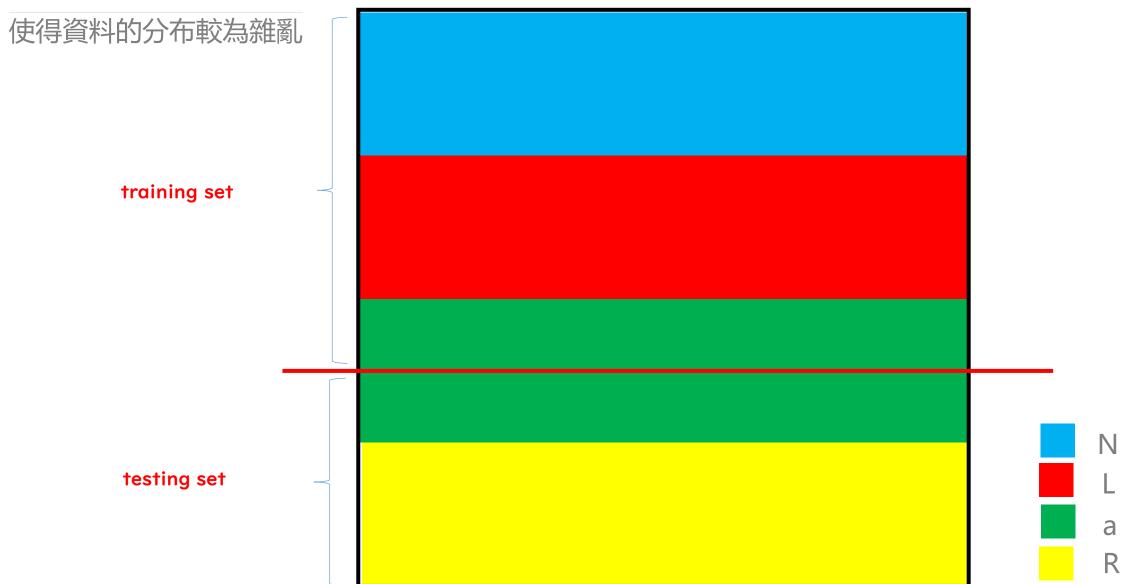
由於心跳的數據是連續性的,必須 切成一筆一筆的資料,並且配對正 確的標籤,之後才能拿來訓練。

資料切割

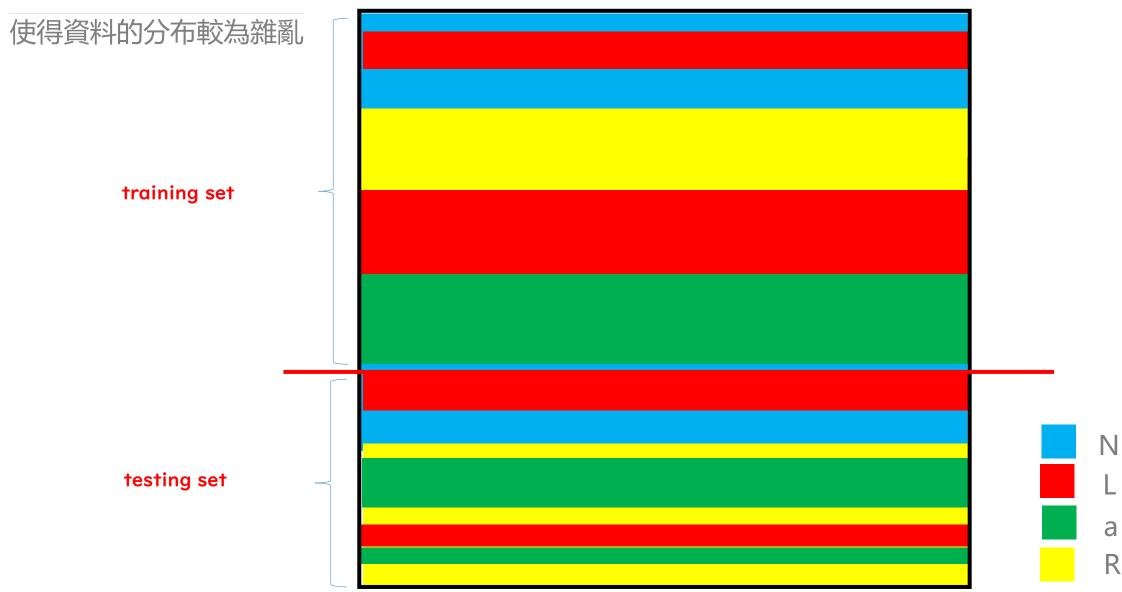
將連續的心跳數據離散化



亂數排序

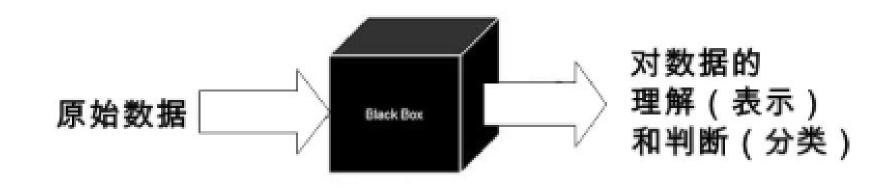


亂數排序



LSTM Model

機器學習就像是黑盒子一樣..







LR = 0.01 ITERATIONS = 100000 NUM_UNITS = 100

Model: BasicLSTMCell loss function: softmax_cross_entropy

Optimizer :AdadeltaOptimizer

BATCH_SIZE = 100 TIME_STEP = 10 INPUT_SIZE = 20

先說結論

In conclusion

_EXTRACTION_SAMPLE_SIZE =
#initi: lization
popula ion = [] #母群
pool = [] #否记池

Result of

LS TM

Training

train_years in range(1, _CroSS_VALIDATION_ALL_

透過系上提供的GPU工作站進行Data # Training,由於tf自帶Optimizer,在 learning rate上不用那麼著墨,比較注 重在訓練次數的調整上。

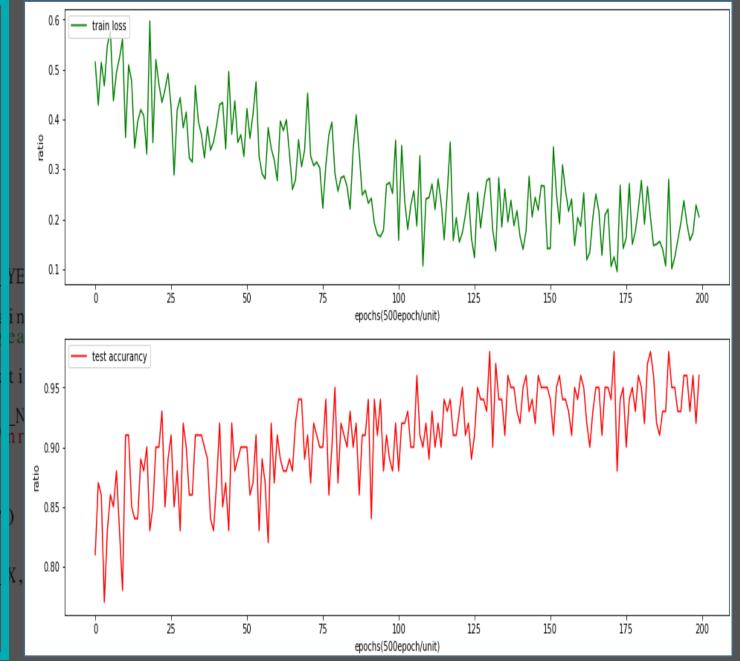
#p在Epochs數量少時正確率還不是那麼明顯,不過把Epochs大幅提高(從8000->100000次)後,有了明顯的提升

/mutation(train_X, train_y, train_years, #print('目前最好基因:'.best gene.dec value)

跑了大約 I 個小時(順便看世足),最後得 到了平均95%以上的Accurancy。

print(Training Tinish. print(training_return)

print(testing_return)



```
EXTRACTION SAMPLE SIZE = 200
    ion = [] #母群
popula
      Result of
testing
      Training
      透過系上提供的GPU工作站進行Data
      Training,由於tf自帶Optimizer,在
     Tlearning trate上不用那麼著墨,比較注 [18]
      重在訓練次數的調整上。
      在Epochs數量少時正確率還不是那麼
      明顯,不過把Epochs大幅提高(從
      8000->100000次)後,有了明顯的提
      mutation(train_X, train_y, train_years,
      跑了大約 | 個小時(順便看世足), 最後得
      到了平均95%以上的Accurancy。
     print(training return)
```

print(testing_return)

```
accuracy = sess.run(accuracy, {tf_train_x: batch_test_X, tf_train_y: batch_test_y})
        if batch test X.shape[0] < BATCH SIZE:</pre>
            test current iter = 0
        print('train loss: %.4f' %loss_, 'Itest accuracy: %.2f' % accuracy_)
        loss_list.append(loss_)
        accurancy_list.append(accuracy_)
train loss: 0.1903 | test accuracy: 0.92
train loss: 0.2655 | test accuracy: 0.97
train loss: 0.2030 | test accuracy: 0.98
train loss: 0.1474 | test accuracy: 0.96
train loss: 0.1498 | test accuracy: 0.92
train loss: 0.1555 | test accuracy: 0.91
train loss: 0.1400 | test accuracy: 0.93
train loss: 0.1066 | test accuracy: 0.93
train loss: 0.2797 | test accuracy: 0.98
train loss: 0.1009 | test accuracy: 0.95
train loss: 0.1235 | test accuracy: 0.95
train loss: 0.1580 | test accuracy: 0.93
train loss: 0.1927 | test accuracy: 0.93
train loss: 0.2371 | test accuracy: 0.96
train loss: 0.1913 | test accuracy: 0.96
train loss: 0.1576 | test accuracy: 0.93
train loss: 0.1721 | test accuracy: 0.96
train loss: 0.2284 | test accuracy: 0.92
train loss: 0.2050 | test accuracy: 0.96
```

特別感謝

宗鴻跟淙垣在技術上的諸多指導