

pre_processing

September 20, 2021

1 Classification of arrhythmia heartbeat

- Paper: <https://www.nature.com/articles/s41597-020-0386-x>
- Chinese Database: <https://physionetchallenges.org/2021/>

1.1 Importing packages

```
[2]: import numpy as np
import seaborn as sns
import pandas as pd
import matplotlib.pyplot as plt
import glob
import math
import os
import sys
import loess
from scipy.io import loadmat
from scipy import signal
from statsmodels.nonparametric.smoothers_lowess import lowess
```

1.2 Loading records and files

```
[3]: files = os.listdir(os.path.abspath('Datasets/ECGData'))

[4]: df_attrib = pd.read_excel(os.path.abspath('Datasets/AttributesDictionary.xlsx'))
df_condnames = pd.read_excel(os.path.abspath('Datasets/ConditionNames.xlsx'))
df_diagno = pd.read_excel(os.path.abspath('Datasets/Diagnostics.xlsx'))
df_rhyt = pd.read_excel(os.path.abspath('Datasets/RhythmNames.xlsx'))
```

1.3 Describing files

```
[5]: df_attrib
```

```
[5]:
```

	Attributes	Type	ValueRange	\
0	FileName	String	NaN	
1	Rhythm	String	NaN	
2	Beat	String	NaN	
3	PatientAge	Numeric	0-999	

4	Gender	String	MALE/FEMAL
5	VentricularRate	Numeric	0-999
6	AtrialRate	Numeric	0-999
7	QRSDuration	Numeric -	0-999
8	QTInterval	Numeric	0-999
9	QTCorrected	Numeric	0-999
10	RAxis	Numeric	-179~180
11	TAxis	Numeric	-179~181
12	QRSCount	Numeric	0-254
13	QOnset	Numeric	16 Bit Unsigned
14	QOffset	Numeric	17 Bit Unsigned
15	TOffset	Numeric	18 Bit Unsigned

	Description
0	ECG data file name(unique ID)
1	Rhythm Label
2	Other conditions Label
3	Age
4	Gender
5	Ventricular rate in BPM
6	Atrial rate in BPM
7	QRS duration in msec
8	QT interval in msec
9	Corrected QT interval in msec
10	R axis
11	T axis
12	QRS count
13	Q onset(In samples)
14	Q offset(In samples)
15	T offset(In samples)

```
[6]: df_condnames
```

[6]:	Acronym	Name	Full Name
0	1AVB	1 degree atrioventricular block	
1	2AVB	2 degree atrioventricular block	
2	2AVB1	2 degree atrioventricular block(Type one)	
3	2AVB2	2 degree atrioventricular block(Type two)	
4	3AVB	3 degree atrioventricular block	
5	ABI	atrial bigeminy	
6	ALS	Axis left shift	
7	APB	atrial premature beats	
8	AQW	abnormal Q wave	
9	ARS	Axis right shift	
10	AVB	atrioventricular block	
11	CCR	counterclockwise rotation	
12	CR	clockwise rotation	

13	ERV	Early repolarization of the ventricles
14	FQRS	fQRS Wave
15	IDC	Interior differences conduction
16	IVB	Intraventricular block
17	JEB	junctional escape beat
18	JPS	J point shift
19	JPT	junctional premature beat
20	LBBB	left bundle branch block
21	LBBBB	left back bundle branch block
22	LFBBB	left front bundle branch block
23	LRRI	Long RR interval
24	LVH	left ventricle hypertrophy
25	LVHV	left ventricle high voltage
26	LVQRSAL	lower voltage QRS in all lead
27	LVQRSCL	lower voltage QRS in chest lead
28	LVQRSLL	lower voltage QRS in limb lead
29	MI	myocardial infarction
30	MIBW	myocardial infarction in back wall
31	MIFW	Myocardial infarction in the front wall
32	MILW	Myocardial infarction in the lower wall
33	MISW	Myocardial infarction in the side wall
34	PRIE	PR interval extension
35	PWC	P wave Change
36	QTIE	QT interval extension
37	RAH	right atrial hypertrophy
38	RAHV	right atrial high voltage
39	RBBB	right bundle branch block
40	RVH	right ventricle hypertrophy
41	STDD	ST drop down
42	STE	ST extension
43	STTC	ST-T Change
44	STTU	ST tilt up
45	TWC	T wave Change
46	TWO	T wave opposite
47	UW	U wave
48	VB	ventricular bigeminy
49	VEB	ventricular escape beat
50	VFW	ventricular fusion wave
51	VPB	ventricular premature beat
52	VPE	ventricular preexcitation
53	VET	ventricular escape trigeminy
54	WAVN	Wandering in the atrioventricular node
55	WPW	WPW

[7]: df_rhyt

```
[7]:      Acronym Name                               Full Name
0         SB                               Sinus Bradycardia
1         SR                               Sinus Rhythm
2        AFIB                          Atrial Fibrillation
3         ST                               Sinus Tachycardia
4         AF                               Atrial Flutter
5         SI                               Sinus Irregularity
6        SVT                          Supraventricular Tachycardia
7         AT                               Atrial Tachycardia
8        AVNRT  Atrioventricular Node Reentrant Tachycardia
9         AVRT      Atrioventricular Reentrant Tachycardia
10        SAAWR      Sinus Atrium to Atrial Wandering Rhythm
```

```
[8]: df_diagno
```

```
[8]:      FileName Rhythm      Beat PatientAge Gender \
0  MUSE_20180113_171327_27000  AFIB  RBBB TWC      85  MALE
1  MUSE_20180112_073319_29000   SB      TWC      59  FEMALE
2  MUSE_20180111_165520_97000   SA     NONE      20  FEMALE
3  MUSE_20180113_121940_44000   SB     NONE      66  MALE
4  MUSE_20180112_122850_57000   AF  STDD STTC      73  FEMALE
...
10641 MUSE_20181222_204306_99000  SVT     NONE      80  FEMALE
10642 MUSE_20181222_204309_22000  SVT     NONE      81  FEMALE
10643 MUSE_20181222_204310_31000  SVT     NONE      39  MALE
10644 MUSE_20181222_204312_58000  SVT     NONE      76  MALE
10645 MUSE_20181222_204314_78000  SVT     NONE      75  MALE

      VentricularRate  AtrialRate  QRSDuration  QTInterval  QTCorrected \
0                117         234         114         356         496
1                52          52          92         432         401
2                67          67          82         382         403
3                53          53          96         456         427
4               162         162         114         252         413
...
10641            196          73         168         284         513
10642            162          81         162         294         482
10643            152          92         152         340         540
10644            175         178         128         310         529
10645            117         104         140         312         435

      RAxis  TAxis  QRSCount  QOnset  QOffset  TOffset
0         81   -27         19     208     265     386
1         76    42          8     215     261     431
2         88    20         11     224     265     415
3         34     3          9     219     267     447
4         68   -40         26     228     285     354
```

...
10641	258	244		32	177	261	319
10642	110	-75		27	173	254	320
10643	250	38		25	208	284	378
10644	98	-83		29	205	269	360
10645	263	144		19	208	278	364

[10646 rows x 16 columns]

1.4 Checking NaN values

```
[9]: df_diagno.isnull().sum()
```

```
[9]: FileName          0
     Rhythm            0
     Beat              0
     PatientAge        0
     Gender             0
     VentricularRate    0
     AtrialRate         0
     QRSDuration        0
     QTInterval         0
     QTCorrected        0
     RAxis              0
     TAxis              0
     QRSCount           0
     QOnset             0
     QOffset            0
     TOffset            0
     dtype: int64
```

```
[10]: df_diagno['Rhythm'].unique()
```

```
[10]: array(['AFIB', 'SB', 'SA', 'AF', 'SR', 'ST', 'SVT', 'AT', 'AVNRT',
            'SAAWR', 'AVRT'], dtype=object)
```

```
[11]: df_diagno.groupby(['Rhythm']).count()['FileName']
```

```
[11]: Rhythm
     AF      445
     AFIB    1780
     AT      121
     AVNRT     16
     AVRT       8
     SA      399
     SAAWR       7
     SB     3889
```

```

SR      1826
ST      1568
SVT     587
Name: FileName, dtype: int64

```

1.5 Grouping rhythms

```

[ ]: df_1 = df_diagno[df_diagno['Rhythm'].isin(['AF','AFIB'])]
df_1['Rhythm_grouped'] = ['AFIB' for _ in range(len(df_1))]

df_2 = df_diagno[df_diagno['Rhythm'].isin(['SVT','AT','_
↳ 'SA','SAAWR','ST','AVNRT','AVRT'])]
df_2['Rhythm_grouped'] = ['GSVT' for _ in range(len(df_2))]

df_3 = df_diagno[df_diagno['Rhythm'].isin(['SB'])]
df_3['Rhythm_grouped'] = ['SB' for _ in range(len(df_3))]

df_4 = df_diagno[df_diagno['Rhythm'].isin(['SR','SI'])]
df_4['Rhythm_grouped'] = ['SR' for _ in range(len(df_4))]

df_total = [df_1,df_2,df_3,df_4]
df = pd.concat(df_total)

```

```

[13]: df.groupby(['Rhythm_grouped']).count()['FileName']

```

```

[13]: Rhythm_grouped
AFIB      2225
GSVT      2706
SB         3889
SR         1826
Name: FileName, dtype: int64

```

```

[48]: df

```

```

[48]:
      FileName Rhythm  Beat  PatientAge  Gender \
0  MUSE_20180113_171327_27000  AFIB  RBBB TWC      85  MALE
4  MUSE_20180112_122850_57000   AF  STDD STTC      73  FEMALE
6  MUSE_20180114_075026_69000  AFIB      TWC      80  FEMALE
15 MUSE_20180113_133901_16000  AFIB  STTC      67  FEMALE
22 MUSE_20180116_123940_90000  AFIB      TWC      81  MALE
...
9878 MUSE_20180209_123628_18000  SR      NONE      66  MALE
9879 MUSE_20180209_122145_99000  SR      NONE      63  FEMALE
9887 MUSE_20180209_132636_13000  SR      NONE      46  MALE
9888 MUSE_20180210_123100_20000  SR      TWC      57  FEMALE
9891 MUSE_20180209_132228_29000  SR      NONE      71  FEMALE

```

	VentricularRate	AtrialRate	QRSDuration	QTInterval	QTCorrected	\
0	117	234	114	356	496	
4	162	162	114	252	413	
6	98	86	74	360	459	
15	72	65	90	416	455	
22	79	150	92	404	463	
...	
9878	73	73	96	416	458	
9879	72	72	72	402	440	
9887	78	78	100	348	396	
9888	68	68	96	400	425	
9891	90	90	68	360	440	

	RAxis	TAxis	QRSCount	QOnset	QOffset	TOffset	Rhythm_grouped
0	81	-27	19	208	265	386	AFIB
4	68	-40	26	228	285	354	AFIB
6	69	83	17	215	252	395	AFIB
15	-1	-15	12	228	273	436	AFIB
22	-15	-51	13	218	264	420	AFIB
...	
9878	43	42	12	209	257	417	SR
9879	13	45	12	214	250	415	SR
9887	73	73	13	218	268	392	SR
9888	18	60	11	210	258	410	SR
9891	70	48	15	226	260	406	SR

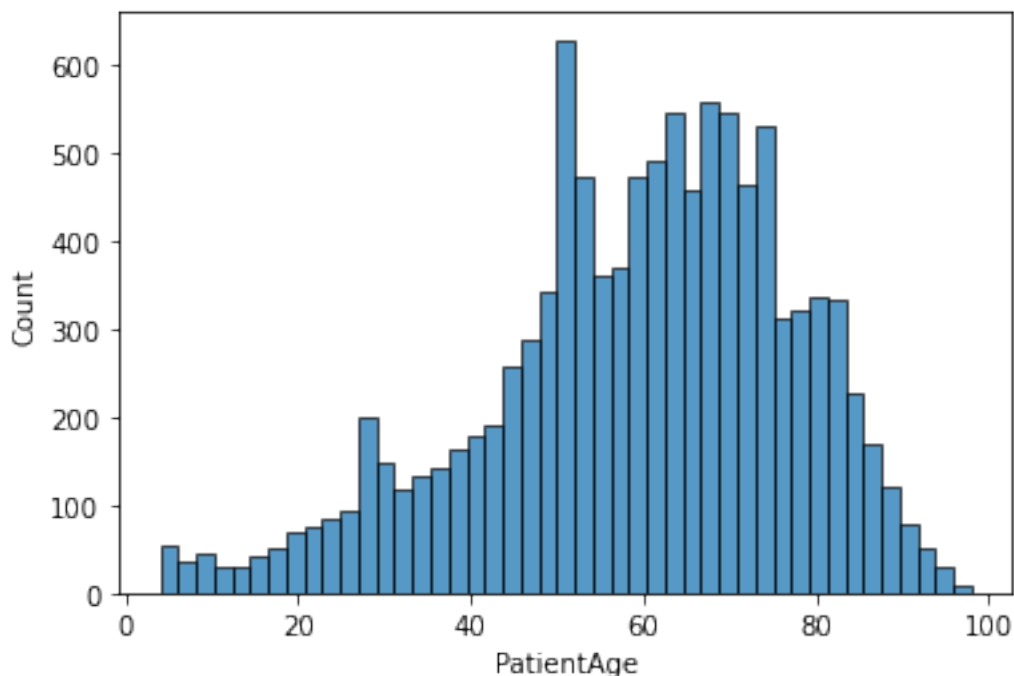
[10646 rows x 17 columns]

```
[14]: df_diagno['Gender'].unique()
```

```
[14]: array(['MALE', 'FEMALE'], dtype=object)
```

```
[15]: sns.histplot(df_diagno['PatientAge'])
```

```
[15]: <AxesSubplot:xlabel='PatientAge', ylabel='Count'>
```



1.6 Loading a patient's ECG

```
[38]: patient_denoised = pd.read_csv('Datasets/ECGDataDenoised/'+ files[0])
      patient_denoised.columns =
      ↪ ['I', 'II', 'III', 'aVR', 'aVL', 'aVF', 'V1', 'V2', 'V3', 'V4', 'V5', 'V6' ]
      patient_denoised
```

```
[38]:
```

	I	II	III	aVR	aVL	aVF	V1	V2	\
0	-150.75	-336.810	-114.980	251.410	-23.8830	-217.52	441.74	646.47	
1	-136.69	-315.560	-108.630	233.450	-19.1260	-204.36	436.06	656.31	
2	-123.74	-296.230	-103.090	217.230	-14.8150	-192.29	432.27	666.14	
3	-112.57	-279.750	-98.611	203.530	-11.2130	-181.87	431.08	676.31	
4	-103.74	-266.700	-95.229	192.840	-8.5498	-173.50	432.76	687.31	
...	
4994	-190.32	52.396	170.760	74.543	-170.8900	107.63	310.73	398.59	
4995	-178.98	65.438	171.870	62.325	-165.6500	114.41	302.32	398.80	
4996	-171.04	74.181	172.080	53.876	-161.4900	118.93	297.25	401.02	
4997	-165.46	79.570	171.300	48.203	-157.8400	121.64	293.97	403.86	
4998	-160.69	83.461	169.810	43.624	-154.1300	123.48	290.95	406.44	
	V3	V4	V5	V6					
0	642.56	35.389	-269.510	-532.21					
1	665.95	76.572	-225.830	-495.39					
2	688.05	113.510	-186.000	-461.84					


```

3      708.47  144.300 -151.680 -432.56
4      726.98  167.550 -124.030 -408.07
...
4994  538.11  230.970   87.684 -249.88
4995  541.18  245.270  110.940 -226.42
4996  544.16  254.480  129.340 -207.20
4997  546.18  259.510  144.140 -190.64
4998  547.24  262.570  157.620 -174.67

```

[4999 rows x 12 columns]

```
[50]: patient_normal = pd.read_csv('Datasets/ECGData/'+ files[0])
      patient_normal
```

```
[50]:
```

	I	II	III	aVR	aVL	aVF	V1	V2	V3	\
0	-214.72	-229.36	-14.64	224.48	-102.48	-122.00	614.88	814.96	912.56	
1	-200.08	-209.84	-9.76	204.96	-97.60	-112.24	605.12	819.84	927.20	
2	-190.32	-195.20	-4.88	195.20	-92.72	-102.48	600.24	829.60	956.48	
3	-165.92	-165.92	0.00	165.92	-82.96	-82.96	590.48	844.24	971.12	
4	-161.04	-156.16	4.88	161.04	-82.96	-78.08	585.60	844.24	985.76	
...	
4995	-117.12	9.76	126.88	53.68	-122.00	68.32	439.20	634.40	800.32	
4996	-102.48	14.64	117.12	43.92	-112.24	63.44	424.56	634.40	819.84	
4997	-92.72	29.28	122.00	34.16	-107.36	73.20	424.56	644.16	829.60	
4998	-92.72	34.16	126.88	29.28	-112.24	78.08	424.56	644.16	829.60	
4999	-87.84	29.28	117.12	29.28	-102.48	73.20	424.56	653.92	834.48	

	V4	V5	V6
0	126.88	-239.12	-507.52
1	165.92	-195.20	-463.60
2	204.96	-151.28	-429.44
3	239.12	-117.12	-400.16
4	273.28	-78.08	-370.88
...
4995	278.16	0.00	-273.28
4996	297.68	24.40	-248.88
4997	312.32	43.92	-229.36
4998	312.32	58.56	-204.96
4999	312.32	63.44	-200.08

[5000 rows x 12 columns]

1.7 Records that do not meet the premises below are excluded.

- lead II = lead I + lead II
- lead aVR + aVL + aVF = 0

Authors: “It is well known that the right hand electrode and left hand electrode could have their

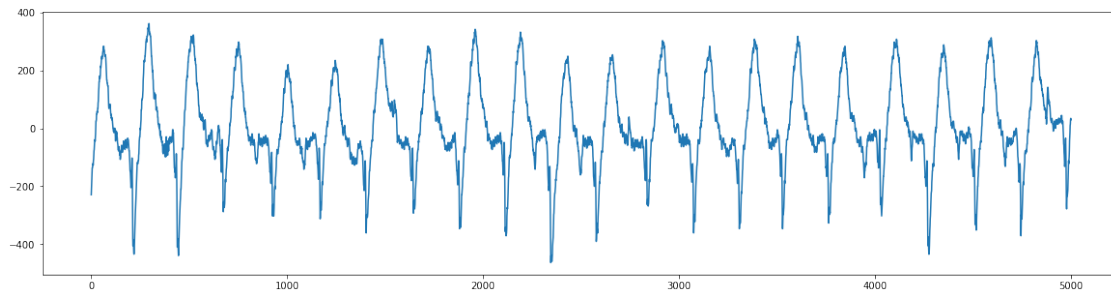
positions switched by operators without a change on corresponding ECG data. Moreover, some of the electrodes could slip off during the test resulting in ECGs displaying a straight line.”

```
[40]: #for i in range(len(patient1)):
#      if (patient1.II[i] != (patient1.I[i] + patient1.III[i])) or ((patient1.
↪aVR[i] + patient1.aVL[i] + patient1.aVF[i]) != 0):
#          patient1.drop(i, inplace=True)
#patient1.reset_index(drop=True, inplace=True)
```

```
[41]: #patient1.reset_index(drop=True, inplace=True)
#patient1
```

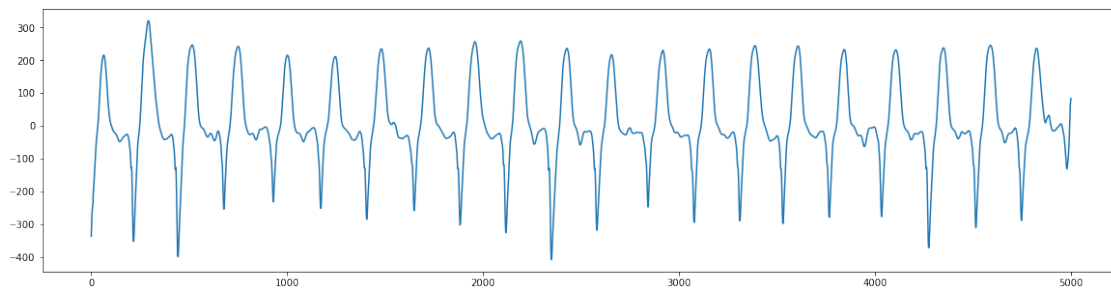
```
[51]: plt.figure(figsize=(20,5))
patient_normal['II'].plot()
```

[51]: <AxesSubplot:>



```
[43]: plt.figure(figsize=(20,5))
patient_denoised['II'].plot()
```

[43]: <AxesSubplot:>



1.8 Denoise signals

- Butterworth low pass filter

- LOcally WEighted Scatterplot Smoothing (LOWESS) curve fitting
- Non local means(NLM)

```
[44]: def NLM_1dDarbon(signal,Nvar,P,PatchHW):
    if isinstance(P,int): # scalar has been entered; expand into patch sample
    ↪index vector
        P = P-1 #Python start index from 0
        Pvec = np.array(range(-P,P+1))
    else:
        Pvec = P # use the vector that has been input
    signal = np.array(signal)
    #debug = [];
    N = len(signal)

    denoisedSig = np.empty(len(signal)) #NaN * ones(size(signal));
    denoisedSig[:] = np.nan
    # to simplify, don't bother denoising edges
    iStart = PatchHW+1
    iEnd = N - PatchHW
    denoisedSig[iStart: iEnd] = 0

    #debug.iStart = iStart;
    #debug.iEnd = iEnd;

    # initialize weight normalization
    Z = np.zeros(len(signal))
    cnt = np.zeros(len(signal))

    # convert lambda value to 'h', denominator, as in original Buades papers
    Npatch = 2 * PatchHW + 1
    h = 2 * Npatch * Nvar**2

    for idx in Pvec: # loop over all possible differences: s - t
        # do summation over p - Eq.3 in Darbon
        k = np.array(range(N))
        kplus = k + idx
        igood = np.where((kplus >=0) & (kplus < N)) # ignore OOB data; we could
    ↪also handle it
        SSD = np.zeros(len(k))
        SSD[igood] = (signal[k[igood]] - signal[kplus[igood]])**2
        Sdx = np.cumsum(SSD)

        for ii in range(iStart,iEnd): # loop over all points 's'
            distance = Sdx[ii + PatchHW] - Sdx[ii - PatchHW-1] #Eq 4;this is in
    ↪place of point - by - point MSE
            # but note the - 1; we want to icnlude the point ii - iPatchHW
```

```

w = math.exp(-distance/h) # Eq 2 in Darbon
t = ii + idx # in the papers, this is not made explicit

if t>0 and t<N:
    denoisedSig[ii] = denoisedSig[ii] + w * signal[t]
    Z[ii] = Z[ii] + w
    #cnt[ii] = cnt[ii] + 1
    #print('ii',ii)
    #print('t',t)
    #print('w',w)
    #print('denoisedSig[ii]', denoisedSig[ii])
    #print('Z[ii]',Z[ii])
# loop over shifts

# now apply normalization
denoisedSig = denoisedSig/(Z + sys.float_info.epsilon)
denoisedSig[0: PatchHW+1] =signal[0: PatchHW+1]
denoisedSig[- PatchHW: ] =signal[- PatchHW: ]
#debug.Z = Z;

return denoisedSig    #,debug

```

```

[52]: Fs=500
fp=50
fs=60
rp=1
rs=2.5
wp=fp/(Fs/2)
ws=fs/(Fs/2)

[n,wn]= signal.buttord(wp,ws,rp,rs)
[bz,az] = signal.butter(n,wn)
LPassDataFile= signal.filtfilt(bz,az,patient_normal['II'])

t = range(0,len(LPPassDataFile))
yy2 = lowess(t,LPassDataFile,frac=0.1)
#yy2 = signal.savgol_filter(LPpassDataFile,2,1)
BWRemoveDataFile = (LPassDataFile-yy2[:,1])
#BWRemoveDataFile = (LPassDataFile-yy2)
Dl1=BWRemoveDataFile

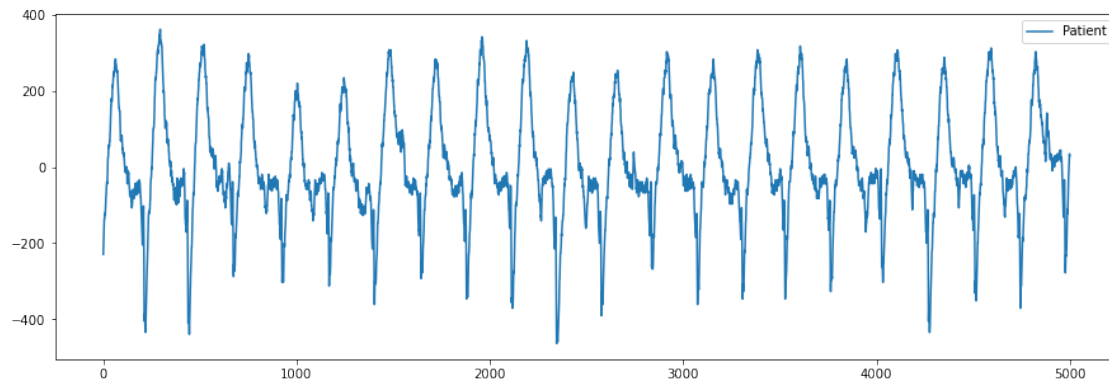
for k in range(2,len(Dl1)-1):
    Dl1[k]=(2*Dl1[k]-Dl1[k-1]-Dl1[k+1])/math.sqrt(6)

NoisSTD = 1.4826*np.median(np.abs(Dl1-np.median(Dl1)))
Denoising_Patient_1= NLM_1dDarbon(BWRemoveDataFile,(1.5)*(NoisSTD),5000,10)

```

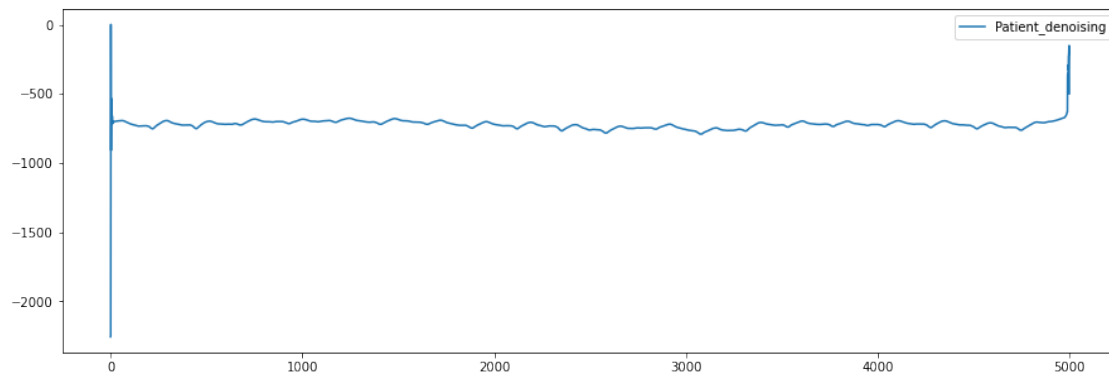
```
[54]: plt.figure(figsize=(15,5))
      #plt.xlim(1000,1500)
      patient_normal['II'].plot()
      plt.legend(['Patient'])
```

[54]: <matplotlib.legend.Legend at 0x24ea473d3d0>



```
[55]: plt.figure(figsize=(15,5))
      #plt.xlim(1000,1500)
      df_denoised = pd.DataFrame(Denoising_Patient_1,columns=['II'])
      df_denoised['II'].plot()
      plt.legend(['Patient_denoising'])
```

[55]: <matplotlib.legend.Legend at 0x24ea47a90d0>



[]: