
anesthPlot

Release beta

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WELCOME TO ANESTHPLOTS DOCUMENTATION!

anesthPlot is a python package developped to extract, manipulate and plots anesthesia data recorded from the Monitor Software to be used mostly in a teaching environment.

Warning: This project is:

- a work in progres
- the processes are mainly focused on horses anesthesia
- in our environment the data recorded came from an as3 or as5 anesthesia machine monitoring ekg, invasive pressure, etCO2, halogenate, spirometry.

1.1 Features

- **load** recordings from a trend or a wave recordings

– **from command line:**

```
python anesthPlot/anesplot/__main__.py
```

- * build a **standard debriefing** (trends) **plot series** (script usage)
 - global histograms (cardiovascular and anesthesia summary)
 - cardiovascular trends time based plots
 - respiratory trends time based plots
 - anesthesia trends time based plots
- * build a **plot for wave** recording
 - one or two waves on the same plot (script usage)

- can also be used as a **python package**

– **usage :**

```
import anesplot.record_main as rec
trendname = 'a_full_path_to_csv_file'
trends = rec.MonitorTrend(trendname)
wavename = rec.trendname_to_wavename(trendname)
waves = rec.MonitorWave(trends)
```

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```
trends.show_graphs() # -> set of plots for debriefing purposes  
waves.plot_waves() # -> one or two traces  
waves.define_a_roi() # -> to register the plotting scales  
waves.animate_fig() #-> to build an animation using these parameters
```

- additional functions are available to extract instantaneous heart rate
 - * see anesplot/treatrec/ekg_to_hr.py

MAIN SCRIPT

2.1 anesplot.record_main module

main script/module to load and display an anesthesia record

can be runned as a script:: python record_main.py

or imported as a package:: import anesplot.record_main as rec

`anesplot.record_main.choosefile_gui(dir_path=None)`

Select a file via a dialog and return the (full) filename.

Parameters `dir_path` (*str*) – location to place the gui (generally paths[data]) else home

Returns `fname[0]` – filename

Return type *str*

`anesplot.record_main.trendname_to_wavename(name)`

just compute the supposed name

`anesplot.record_main.select_type(question=None, items=None, num=0)`

select the recording type:

Returns `kind` – kind of recording in [monitorTrend, monitorWave, taphTrend, telvet]

Return type *str*

`anesplot.record_main.select_wave(waves, num=1)`

select the recording type:

Returns `kind` – kind of recording in [monitorTrend, monitorWave, taphTrend, telvet]

Return type *str*

`anesplot.record_main.build_param_dico(file=None, asource=None, pathdico={'cwd':`

`'/Users/cdesbois/pg/chrisPg/anesthPlot', 'data':`

`'/Users/cdesbois/enva/clinique/recordings/anesthRecords/onPanelPcRecorded',`

`'recordMain': '/Users/cdesbois/pg/chrisPg/anesthPlot/anesplot',`

`'root': '/Users/cdesbois', 'sBg': '/Users/cdesbois/ownCloud', 'sFig':`

`'/Users/cdesbois/ownCloud', 'save': '/Users/cdesbois/ownCloud',`

`'utils': '/Users/cdesbois/pg'})`

initialise a dict save parameters -> TODO see min vs sec

Parameters

- **file** (*str*) – the recording filename

- **source** (*str*) – the origin of the recording

Returns

dico –

a dictionary describing the situation [item, xmin, xmax, ymin, ymax, path, unit, save, memo, file, source]

Return type dict

`anesplot.record_main.plot_trenddata(df, header, param_dico)`

clinical main plots of a trend recordings

parameters df : pdDataframe

recorded data (MonitorTrend.data)

header [dict] recording parameters (MonitorTrend.header)

param_dico [dict] plotting parameters (MonitorTrend.param)

Returns **afig_dico**

Return type dict of name:fig

class `anesplot.record_main.Waves(filename=None)`

Bases: object

the base object to store the records.

class `anesplot.record_main.SlowWave(filename=None)`

Bases: [anesplot.record_main.Waves](#)

class for slowWaves = trends

file [str] short name

filename [str] long name

clean_trend : **external**

clean the data

show_graphs : **external**

plot clinical main plots

clean_trend()

clean the data, remove irrelevant, input = self.data, output = pandas dataframe nb doesnt change the obj.data in place

show_graphs()

basic clinical plots

class `anesplot.record_main.MonitorTrend(filename=None, load=True)`

Bases: [anesplot.record_main.SlowWave](#)

monitor trends recordings:

input = filename : path to file load = boolean to load data (default is True)

file [str] short name

filename [str] long name

header [dict] record parameters

source [str] recording apparatus (default = monitor)

fs [float] sampling rate

param [dict] display parameters

clean_trend [external] clean the data

show_graphs [external] plot clinical main plots

class `anesplot.record_main.TaphTrend(filename=None)`

Bases: `anesplot.record_main.SlowWave`

taphonius trends recordings

input FILLME

attributes FILLME

load_header()

load the header -> pandas.dataframe

extract_taph_events()

extract Taph events

Parameters **data** (*pandas dataframe*) – record df form taphonius recording)

Returns events dataframe

Return type eventdf pandas dataframe

class `anesplot.record_main.FastWave(filename=None)`

Bases: `anesplot.record_main.Waves`

class for Fastwaves = continuous recordings.

plot_wave(*traces_list=None, dt=None*)

simple choose and plot for a wave input = none -> GUI, or list of waves to plot (max=2)

define_a_roi(*erase=False*)

define a Region Of Interest (roi).

input : erase (boolean) default=False takes the figure attribute return a dictionary containing:

dt : xscale datetime location pt: xscale point location sec: xscale seconde location ylims: ylimits
traces: waves used to draw the figure fig : the related figure

animate_fig(*speed=1, save=False, savedir=~*)

build a video the previous builded figure

use .fig attribute (builded through .plot_wave()) and .roi attribute (builded thourhg .define_a_roi())

Parameters

- **speed** (*int, optional*) – speed of the video, defaults to 1
- **save** (*boolean, optional*) – save or just display, defaults to False
- **savedir** (*str, optional*) – directory to save the animation, defaults to ~

Returns video file

Return type mp4

class `anesplot.record_main.TelevetWave`(*filename=None*)

Bases: `anesplot.record_main.FastWave`

class to organise teleVet recordings transformed to csv files.

class `anesplot.record_main.MonitorWave`(*filename=None, load=True*)

Bases: `anesplot.record_main.FastWave`

class to organise monitorWave recordings. input : filename = path to file load = boolean to load data (default is True)

attibutes FILLME

methods FILLME

`anesplot.record_main.main()`

main script called from command line call : `python anesthPlot/anesplot/__main__.py` args : optional filename (fullname)

return: set of plots for either monitorTrend, monitorWave oe televet recording

MODULES

3.1 anesplot package

3.1.1 Subpackages

anesplot.config package

Submodules

anesplot.config.build_recordRc module

build a recordRc.yaml configuration file to adapt to a specific computer location at the root of anesplot

- input <-> data : to load the records
- output <-> save : to save the plots

`anesplot.config.build_recordRc.filedialog(kind="", directory="/Users/cdesbois/pg/chrisPg/anesthPlot/anesplot/config", for_open=True, fmt="", is_folder=False)`
general dialog function.

`anesplot.config.build_recordRc.read_config()`
locate & load the yaml file.

`anesplot.config.build_recordRc.write_configfile(path)`
record the yaml file.

`anesplot.config.build_recordRc.main()`
main function for script execution.

anesplot.config.load_recordRc module

load an already generated recordRc.yaml configuration file

- input <-> data : to load the records
- output <-> save : to save the plots

`anesplot.config.load_recordRc.build_paths()`

read the yaml configuration file.

`anesplot.config.load_recordRc.adapt_with_syspath(path_dico)`

add the folder location to the system path.

Module contents

anesplot.loadrec package

Submodules

anesplot.loadrec.explore module

Created on Thu Mar 12 16:52:13 2020

@author: cdesbois

`anesplot.loadrec.explore.gui_choosefile(paths=None)`

select a file via a dialog and return the file name.

anesplot.loadrec.loadmonitor_trendrecord module

Created on Wed Jul 24 13:43:26 2019 @author: cdesbois

load a monitor trend recording:

- choose a file
- load the header to a dictionary
- load the data into a pandas dataframe

`anesplot.loadrec.loadmonitor_trendrecord.choosefile_gui(dir_path=None)`

select a file using a dialog.

Parameters `dir_path` (*str*) – optional location of the data (paths[data])

Returns filename (full path)

Return type str

`anesplot.loadrec.loadmonitor_trendrecord.loadmonitor_trendheader(filename)`

load the file header.

Parameters `filename` (*str*) – full name of the file

Returns header

Return type dict

`anesplot.loadrec.loadmonitor_trendrecord.loadmonitor_trenddata(filename, headerdico)`

load the monitor trend data

Parameters

- `filename` (*str*) – fullname

- **headerdico** (*dict*) – fileheader

Returns df = trends data

Return type pandas.DataFrame

anesplot.loadrec.loadmonitor_waverecord module

Created on Wed Jul 24 14:56:58 2019 @author: cdesbois

load a monitor wave recording:

- choose a file
- load the header to a pandas dataframe
- load the date into a pandas dataframe

anesplot.loadrec.loadmonitor_waverecord.**choosefile_gui**(*dir_path=None*)
select a file using a dialog.

Parameters **dir_path** (*str*) – optional location of the data (paths[data])

Returns filename (full path)

Return type str

anesplot.loadrec.loadmonitor_waverecord.**loadmonitor_waveheader**(*filename=None*)
load the wave file header.

Parameters **filename** (*str*) – full name of the file

Returns header

Return type pandas.DataFrame

anesplot.loadrec.loadmonitor_waverecord.**loadmonitor_wavedata**(*filename=None*)
load the monitor wave csvDataFile.

Parameters **filename** (*str*) – full name of the file

Returns df = trends data

Return type pandas.DataFrame

anesplot.loadrec.loadtaph_trendrecord module

Created on Wed Jul 24 15:30:07 2019 @author: cdesbois

load a taphonius data recording:

- choose a file
- load the patient datafile to a dictionary
- load the physiological date into a pandas dataframe

anesplot.loadrec.loadtaph_trendrecord.**choosefile_gui**(*dir_path=None*)
select a file using a dialog.

Parameters `dir_path` (*str*) – optional location of the data (paths[data])

Returns filename (full path)

Return type str

`anesplot.loadrec.loadtaph_trendrecord.loadtaph_trenddata(filename)`

load the taphoniusData trends data.

Parameters `filename` (*str*) – fullname

Returns df = trends data

Return type pandas.DataFrame

`anesplot.loadrec.loadtaph_trendrecord.loadtaph_patientfile(headername)`

load the taphonius patient.csv file

Parameters `headername` (*str*) – fullname

Returns descr = patient_data

Return type dict

anesplot.loadrec.loadtelevet module

Created on Wed Jul 31 16:22:06 2019 @author: cdesbois

load televet exported (csv) data: to be developped

`anesplot.loadrec.loadtelevet.choosefile_gui(dir_path=None)`

select a file using a dialog.

Parameters `dir_path` (*str*) – optional location of the data (paths[data])

Returns filename (full path)

Return type str

`anesplot.loadrec.loadtelevet.loadtelevet(fname=None, all_traces=False)`

load the televetCsvExportedFile.

Parameters

- **file** (*str*) – name of the file
- **all_traces** (*bool*) – load all the derivations

Returns df = recorded traces

Return type pandas.DataFrame

Module contents

anesplot.plot package

Submodules

anesplot.plot.trend_plot module

Created on Tue Apr 19 09:08:56 2016 @author: cdesbois

collection of functions to plot the trend data

`anesplot.plot.trend_plot.color_axis(ax, spine='bottom', color='r')`
change the color of the label & tick & spine.

Parameters

- **ax** (*matplotlib.pyplot.axis*) – the axis
- **spine** (*str*) – optional location in [bottom, left, top, right]
- **colors** (*str*) – optional color

`anesplot.plot.trend_plot.append_loc_to_fig(ax, dt_list, label='g')`
append vertical lines to indicate a location eg: arterial blood gas

Parameters

- **ax** (*matplotlib.pyplot.axis*) – the axis
- **dt_list** (*[datetime]*) – list of datetime values
- **label** (*str*) – a key to add to the label (default is g)

Returns **res** a dictionary containing the locations

Return type dict

`anesplot.plot.trend_plot.save_graph(path, ext='png', close=True, verbose=True)`
Save a figure from pyplot. :param path: The path (and filename, without the extension) to save the figure to.

Parameters

- **ext** (*string (default='png')*) – The file extension. This must be supported by the active matplotlib backend (see matplotlib.backends module). Most backends support png, pdf, ps, eps, and svg.
- **close** (*boolean (default=True)*) – Whether to close the figure after saving. If you want to save the figure multiple times (e.g., to multiple formats), you should NOT close it in between saves or you will have to re-plot it.
- **verbose** (*boolean (default=True)*) – Whether to print information about when and where the image has been saved.

`anesplot.plot.trend_plot.plot_header(descr, param=None)`
plot the header of the file.

Parameters

- **descr** (*dict*) – header of the recording
- **param** (*dict*) – dictionary of parameters

Returns **fig** plot of the header

Return type `pyplot.figure`

`anesplot.plot.trend_plot.hist_cardio(data, param=None)`
mean arterial pressure histogramme using matplotlib.

Parameters

- **data** (*pandas.DataFrame*) – the recorded trends data (keys used : `ip1m` and `hr`),
- **param** (*dict*) – parameters (`save=boolean`, `path`: path to directory)

Returns **fig** `matplotlib.pyplot.figure`

`anesplot.plot.trend_plot.plot_one_over_time(x, y, colour)`
plot y over x using colour

`anesplot.plot.trend_plot.hist_co2_iso(data, param=None)`
CO2 and iso histogramme (NB CO2 should have been converted from % to mmHg)

Parameters

- **data** (*pandas.DataFrame*) – the trends recorded data
- **param** (*dict*) – dictionary of parameters

Returns **fig** `pyplot.figure`

`anesplot.plot.trend_plot.cardiovasc(data, param=None)`
cardiovascular plot

Parameters

- **data** (*pandas.DataFrame*) – the recorded trends data keys used : [`ip1s`, `ip1m`, `ip1d`, `hr`]
- **param** (*dict*) – dict(`save`: boolean, `path[save]`, `xmin`, `xmax`, `unit`, `dtime` = boolean for time display in HH:MM format)

Returns **fig**= `pyplot.figure`

`anesplot.plot.trend_plot.cardiovasc_p1p2(data, param=None)`
cardiovascular plot with central venous pressure (p2)

Parameters

- **data** (*pandas.DataFrame*) – the trends recorded data keys used : [`ip1s`, `ip1m`, `ip1d`, `hr`, `ip2s`, `ip2m`, `ip2d`]
- **param** (*dict*) – dict(`save`: boolean, `path[save]`, `xmin`, `xmax`, `unit`, `dtime` = boolean for time display in HH:MM format)

Returns **fig**= `pyplot.figure`

`anesplot.plot.trend_plot.co2iso(data, param=None)`
anesth plot (CO2/iso)

Parameters

- **data** (*pandas.DataFrame*) – the recorded data keys used : [`ip1s`, `ip1m`, `ip1d`, `hr`]
- **param** (*dictionary*) – dict(`save`: boolean, `path[save]`, `xmin`, `xmax`, `unit`, `dtime` = boolean for time display in HH:MM format)

:returns fig= pyplot.figure

anesplot.plot.trend_plot.**func**(ax, x, y1, y2, color='tab:blue', x0=38)

anesplot.plot.trend_plot.**co2o2**(data, param)

respiratory plot (CO2 and Iso)

Parameters

- **data** (*pandas.DataFrame*) – recorded trends data keys used :[ip1s, ip1m, ip1d, hr]
- **param** (*dict*) – dict(save: boolean, path[save], xmin, xmax, unit, dtime = boolean for time display in HH:MM format)

Returns fig= pyplot.figure

anesplot.plot.trend_plot.**ventil**(data, param)

plot ventilation parameters (.tvInsp, .pPeak, .pPlat, .peep, .minVexp, .co2RR, .co2exp)

Parameters

- **data** (*pandas.DataFrame*) – recorded data, keys used :[ip1s, ip1m, ip1d, hr]
- **param** (*dict*) – dict(save: boolean, path[save], xmin, xmax, unit, dtime = boolean for time display in HH:MM format)

Returns fig= pyplot.figure

anesplot.plot.trend_plot.**recrut**(data, param)

display a recrut manoeuver (.pPeak, .pPlat, .peep, .tvInsp)

Parameters

- **data** (*pandas.DataFrame*) – recorded data keys used :[ip1s, ip1m, ip1d, hr]
- **param** (*dict*) – dict(save: boolean, path[save], xmin, xmax, unit, dtime = boolean for time display in HH:MM format)

:returns fig= pyplot.figure

anesplot.plot.trend_plot.**ventil_cardio**(data, param)

build ventilation and cardiovascular plot

Parameters

- **data** (*pandas.DataFrame*) – teh recorded trends data keys used :[ip1s, ip1m, ip1d, hr]
- **param** (*dict*) – dict(save: boolean, path[save], xmin, xmax, unit, dtime = boolean for time display in HH:MM format)

Returns fig= pyplot.figure

anesplot.plot.trend_plot.**save_distri**(data, path)

save as **O_.** the 4 distributions graphs for cardiovasc annd respi

anesplot.plot.trend_plot.**fig_memo**(path, fig_name)

append latex citation commands in a txt file inside the fig folder create the file iif it doesnt exist

anesplot.plot.wave_plot module

Created on Tue Apr 19 09:08:56 2016

@author: cdesbois

`anesplot.plot.wave_plot.color_axis(ax, spine='bottom', color='r')`
change the color of the label & tick & spine.

Parameters

- **ax** (*matplotlib.pyplot.axis*) – the axis
- **spine** (*str*) – optional location in [bottom, left, top, right]
- **colors** (*str*) – optional color

`anesplot.plot.wave_plot.plot_wave(data, keys, param)`
plot the waves recorded (from as5)

Parameters

- **data** (*pandas.DataFrame*) – the recorded trends data
- **keys** (*list*) – one or two in [wekg,ECG,wco2,wawp,wflow,wap]
- **{mini}** (*dict*) – limits in point value (index), maxi: limits in point value (index)}

Returns fig plt.figure the plot

Returns lines plt.line2D the line to animate

(Nb plot data/index, but the xscale is indicated as sec)

`anesplot.plot.wave_plot.get_a_roi(waves)`
use the drawn figure to extract the relevant data in order to build an animation

Parameters waves (*MonitorWave object*) – a wave recording

Returns a dictionary containing ylims, xlims(point, dtime and sec),

traces used to build the plot, the fig object :rtype: dictionary

`anesplot.plot.wave_plot.create_video(waves, speed=1, save=False, savedir='~')`
create a video from a figure

Module contents

Created on Tue Apr 19 09:08:56 2016

functions to plot the trend data

@author: cdesbois

anesplot.treatrec package

Submodules

anesplot.treatrec.clean_data module

Created on Wed Jul 31 16:05:29 2019

@author: cdesbois

`anesplot.treatrec.clean_data.clean_trenddata(df)`
remove artifacts in the recorded trends

anesplot.treatrec.ekg_to_hr module

Created on Wed Feb 12 16:52:00 2020 @author: cdesbois

function used to treat an EKG signal and extract the heart rate typically (copy, paste and execute line by line)

0. after

```
:: import pandas as pd
    import anesplot.record_main as rec from anesplot.treatrec import ekg_to_hr as tohr
```

1. load the data in a pandas dataframe:

(through classes `rec.MonitorTrend` & `rec.MonitorWave`)

```
trendname = '' # fullname
or
trendname = rec.choosefile_gui()
```

```
wavename = rec.trendname_to_wavename(trendname)
-
# load the data
trends = rec.MonitorTrend(trendname)
waves = rec.MonitorWave(wavename)
-
# format the name
name = trends.header['Patient Name'].title().replace(' ', '')
name = name[0].lower() + name[1:]
```

2. treat the ekg wave:

- get parameters
- build a dataframe to work with (waves)
- low pass filtering
- build the beat locations (beat based dataframe):

```
params = waves.param
ekg_df = pd.DataFrame(waves.data.wekg)
ekg_df['wekg_lowpass'] = rec.wf.fix_baseline_wander(ekg_df.wekg,
                                                    waves.param['fs'])
beat_df = tohr.detect_beats(ekg_df.wekg_lowpass, mult=1)
```

3. perform the manual adjustments required:

- based on a graphical display of beat locations, an rr values
- build a container for the manual corrections:

```
figure = tohr.plot_beats(ekg_df.wekg_lowpass, beat_df)
to_change_df = pd.DataFrame(columns=beat_df.columns.insert(0, 'action'))
```

- remove or add peaks : zoom on the figure to observe only one peak, then:

```
to_change_df = tohr.remove_beat(beat_df, ekg_df, to_change_df, figure)
or
to_change_df = tohr.append_beat(beat_df, ekg_df, to_change_df, figure,
                                yscale=1)
```

- combine to update the beat_df with the manual changes:

```
beat_df = tohr.update_beat_df(beat_df, to_change_df,
                              path_to_file="", from_file=False)
```

- save the peaks locations:

```
tohr.save_beats(beat_df, to_change_df, savename='', savepath=None)
(# or reload
beat_df = pd.read_hdf('beatDf.hdf', key='beatDf') )
```

4. go from points values to continuous time:

```
beat_df = tohr.compute_rr(beat_df)
ahr_df = tohr.interpolate_rr(beat_df)
tohr.plot_rr(ahr_df, params)
```

5. append intantaneous heart rate to the initial data:

```
ekg_df = tohr.append_rr_and_ihr_to_wave(ekg_df, ahr_df)
waves.data = tohr.append_rr_and_ihr_to_wave(waves.data, ahr_df)
trends.data = tohr.append_ihr_to_trend(trends.data, waves.data, ekg_df)
```

6. save:

```
tohr.save_trends_data(trends.data, savename=name, savepath='data')
tohr.save_waves_data(waves.data, savename=name, savepath='data')
```

`anesplot.treatrec.ekg_to_hr.detect_beats(ser, fs=300, species='horse', mult=1)`
detect the peak locations

Parameters

- **ser** (*pandas.series*) – the data
- **fs** (*integer*) – sampling frequency
- **species** (*string*) – in [horse]
- **mult** (*float*) – correction / 1 for qRs amplitude

Returns df=pandas.DataFrame

`anesplot.treatrec.ekg_to_hr.plot_beats(ecg, beats)`
plot ecg waveform + beat location

`anesplot.treatrec.ekg_to_hr.append_beat(beatdf, ekgdf, tochange_df, fig, lim=None, yscale=1)`
locate the beat in the figure, append to a dataframe[toAppend]

Parameters

- **beatdf** (*pandas.DataFrame*) – contains the point based location (pLocs)
- **ekgdf** (*pandas dataframe*) – contains the wave recording ((wekg_lowpass)
- **tochange_df** (*pandas.DataFrame*) – to store the beats toAppend or toRemove
- **fig** (*pyplot.Figure*) – figure to find time limits
- **lim** (*integer*) – ptBasedLim optional to give it manually
- **yscale** (*float*) – amplitude mutliplication factor for detection (default=1)

Returns tochange_df: incremented changedf (pt location)

Return type pandasDataFrame

methods :

locate the beat in the figure, append to a dataframe[toAppend] 0.: if not present : build a dataframe:

```
>>> to_change_df = pd.DataFrame(columns=['toAppend', 'toRemove'])
```

1.: locate the extra beat in the figure (cf plot_beats()) and zoom to observe only a negative peak

2.: call the function:

```
>>> to_change_df = remove_beat(beatdf, ekgdf, tochange_df, fig)
-> the beat parameters will be added the dataframe
```

.in the end of the manual check, update the beat_df

- first : save beat_df and to_change_df
- **second** [run:]

```
>>> beat_df = update_beat_df()
```

`anesplot.treatrec.ekg_to_hr.remove_beat(beatdf, ekgdf, tochange_df, fig, lim=None)`
locate the beat in the figure, append to a dataframe[toRemove]

0.: if not present build a dataframe:

```
>>> to_change_df = pd.DataFrame(columns=['toAppend', 'toRemove'])
```

1.: locate the extra beat in the figure (cf plot_beats()) and zoom to observe only a negative peak

2.: call the function:::

```
>>> to_change_df = remove_beat(beatdf, ekgdf, tochange_df, fig)
-> the beat parameters will be added the dataframe
```

.(in the end of the manual check, update the beat_df

- first : save beat_df and to_change_df
- **second** [run]

```
>>> beat_df = update_beat_df()
```

`anesplot.treatrec.ekg_to_hr.save_beats(beatdf, tochangedf, savename="", dirpath=None)`
save the beats locations as csv and hdf5 file

Parameters

- **beatde** (*pd.dataframes*) –
- **tochangedf** (*pandas.dataframe*) –
- **savename** (*filename*) –
- **dirpath** (*path to save in*) –
- **output** –
- **-----** –
- **file** (*hdf*) –
- **key**='beatDf' –

`anesplot.treatrec.ekg_to_hr.update_beat_df(beatdf, tochangedf, path_to_file="", from_file=False)`
implement in the beat location the manual corrections fromFile = True force the disk loading of the dataframes

`anesplot.treatrec.ekg_to_hr.compute_rr(beatdf, fs=None)`
compute rr intervals (from pt to time)

Parameters

- **beatdf** (*pd.DataFrame*) – with pLoc

- **fs** (*integer*) – sampling frequency

Returns with: $rr = rr \text{ duration}$ $rrDiff = rrVariation$ $rrSqDiff = rrVariation^2$

Return type `pd.DataFrame`

`anesplot.treatrec.ekg_to_hr.interpolate_rr(beatdf, kind=None)`
interpolate the beat_df (pt -> time values)

Parameters

- **beatDf** (*pd.DataFrame*) –
- **kind** (*str*) – linear or cubic(default)

Returns $espts = \text{evenly spaced points}$ $rrInterpol = \text{interpolated } rr$

Return type `pd.DataFrame` with evenly spaced data

`anesplot.treatrec.ekg_to_hr.plot_rr(ahr_df, param, HR=False)`
plot RR vs pt values + $rrSqDiff$

Parameters

- **pdDataFrame** (*hr_df* =) –
- **params** – dict containing fs as key

`anesplot.treatrec.ekg_to_hr.append_rr_and_ihr_to_wave(wave, ahrdf)`
append rr and ihr to the waves based on pt value (ie index)

`anesplot.treatrec.ekg_to_hr.plot_agreement(trenddf)`
plot ip1HR & ihr to check agreement

`anesplot.treatrec.ekg_to_hr.append_ihr_to_trend(trenddf, wavedf, ekcdf)`
append ihr (instantaneous heart rate) to the trends

`anesplot.treatrec.ekg_to_hr.save_trends_data(trenddf, savename="", dirpath='data')`

save the trends data to a csv and hd5 file, including an ihr column

`trenddf` : `pd.dataframes` `savename` : `str` `dirpath` : `str`

path to save in (default= current working directory)

hdf file, key=trends_data

`anesplot.treatrec.ekg_to_hr.save_waves_data(wavedf, savename="", dirpath='data')`
save the trends data to a hd5 file, including an ihr column

Parameters

- **trenddf** (*pd.dataframes*) –
- **savename** (*str*) – `dirpath` : path to save in (default=data)
- **output** –
- ----- –
- **hdf_file** –
- **key='waves_data'** –

anesplot.treatrec.extract_hypotension module

Spyder Editor

This is a temporary script file.

`anesplot.treatrec.extract_hypotension.extract_hypotension(atrend, pamin=70)`
 return a dataframe with the beginning and ending phses of hypotension

Parameters

- **atrend** (*MonitorTrend object*) –
- **pamin** (*float= threshold de define hypotension on mean arterial pressure*) –
- **70** (*((default is)*) –

Returns **durdf** – transitions (up and down, in seconds from beginning) and duration in the hypotension state (in seconds)

Return type pandas DataFrame containing

`anesplot.treatrec.extract_hypotension.plot_hypotension(atrend, durdf, durmin=15, pamin=70)`
 plot the hupotentions phases

Parameters

- **atrend** (*TYPE*) – DESCRIPTION.
- **durdf** (*TYPE*) – DESCRIPTION.
- **durmin** (*TYPE, optional*) – DESCRIPTION. The default is 15.

Returns **fig** – DESCRIPTION.

Return type TYPE

`anesplot.treatrec.extract_hypotension.scatter_length_meanhypo(atrend, durdf)`
 draw a scatter plot (hypotensive arterial value vs duration of hypotension) :param trends: :type trends: MonitorTrend :param durdf: :type durdf: pandas dataframe containing the value and duration

Returns **fig**

Return type matplotlib.pyplot figure

`anesplot.treatrec.extract_hypotension.plot_all_dir_hypo(dirname=None, scatter=False)`
 walk throught the folder and plot the values

anesplot.treatrec.hr_to_hrv module

`anesplot.treatrec.hr_to_hrv.build_hrv_limits(spec='horse')`
 return a dico containing HRV limits (VLF, LF, HF) input : spec in [horse, man]

anesplot.treatrec.wave_func module

Created on Fri Dec 8 12:46:41 2017

@author: cdesbois

`anesplot.treatrec.wave_func.fix_baseline_wander(data, fs=500)`

BaselineWanderRemovalMedian.m from ecg-kit. Given a list of amplitude values (data) and sample rate (sr), it applies two median filters to data to compute the baseline. The returned result is the original data minus this computed baseline.

`anesplot.treatrec.wave_func.rol_mean(ser, win_length=1, fs=500)`

returns a rolling mean of a RR serie

Parameters

- **pd.Series** (*ser*) –
- **win_length** (*integer*) – window lenght for averaging (in sec),
- **fs** (*int*) – sampling frequency

`anesplot.treatrec.wave_func.return_points(df, fig)`

return a tuple containing the point values of ROI

Parameters

- **df** (*anesthesia record dataframe*) –
- **fig** (*pyplot.figure*) –

Returns ROI

Return type dict

`anesplot.treatrec.wave_func.restrict_time_area(df1, mini=None, maxi=None)`

return a new dataframe with reindexation

Parameters

- **df1** (*pandas.DataFrame*) –
- **mini** (*integer*) – miniPointValue
- **maxi** (*integer*) – maxiPointValue

Returns

Return type pandas.DataFrame

Module contents

3.1.2 Submodules

3.1.3 Module contents

anesthPlot is a package to plot/use clinical anesthesia records for teaching

three way to use it:

1. **run directly anesplot from a terminal** -> PYTHONPATH=<pathToAnesthPlot> python -m anesplot -> generate a quick plotting of most interestings parts

2. **from an ipython terminal** -> `import anesthPlot.anesplot.recordmain as rec -> trends = rec.MonitorTrend() -> waves = rec.MonitorWave() ->` and use the objects trends and waves
3. import the module in a python environment (see below)

(the presets are actually designed

- for use with equine anesthesia
- to load data from a Monitor generated datex AS3/5 monitoring machine)

typical use when importing the module to build a clinical case

```
import os import sys
```

```
import numpy as np import pandas as pd
```

```
import anesplot.record_main as rec sys.path.append(os.path.expanduser("~/pg/utills)) from utills import saveGraph im-
port bloodGases.bgmain_manual as bgman
```

```
paths = rec.paths paths[save] = os.path.expanduser("~/toPlay/temp/) os.chdir(paths[save])
```

```
## globals def save_plot(name):
```

```
    filename = os.path.join(paths[save], fig, name) saveGraph(filename, ext=png, close=False, verbose=True)
```

def explore_hdf(filename):

```
    try: hdf = pd.HDFStore(filename) keys= [key.replace(/, ) for key in hdf.keys()] print( found h5_file { }
        that contains { }.format(filename, keys))
```

```
        hdf.close()
```

```
    except: print({ } is not an h5 file.format(filename))
```

```
saveName = os.path.join(paths[save], data, aname.h5)
```

```
explore_hdf(saveName)
```

```
## load and work trendName = rec.choosefile_gui(paths[data]) WaveName = rec.choosefile_gui(paths[data])
```

```
# build objects with headers trends = rec.MonitorTrend(trendName, load=True) waves = rec.MonitorWave(waveName,
load=True)
```

```
# or append data (pretreated ones) #trends.data = pd.read_hdf(saveName, trend_df) #waves.data =
pd.read_hdf(saveName, wave_df)
```

```
#remove filenames del waveName, trendName
```

```
# now you are ready to work with loaded trends and waves
```


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