

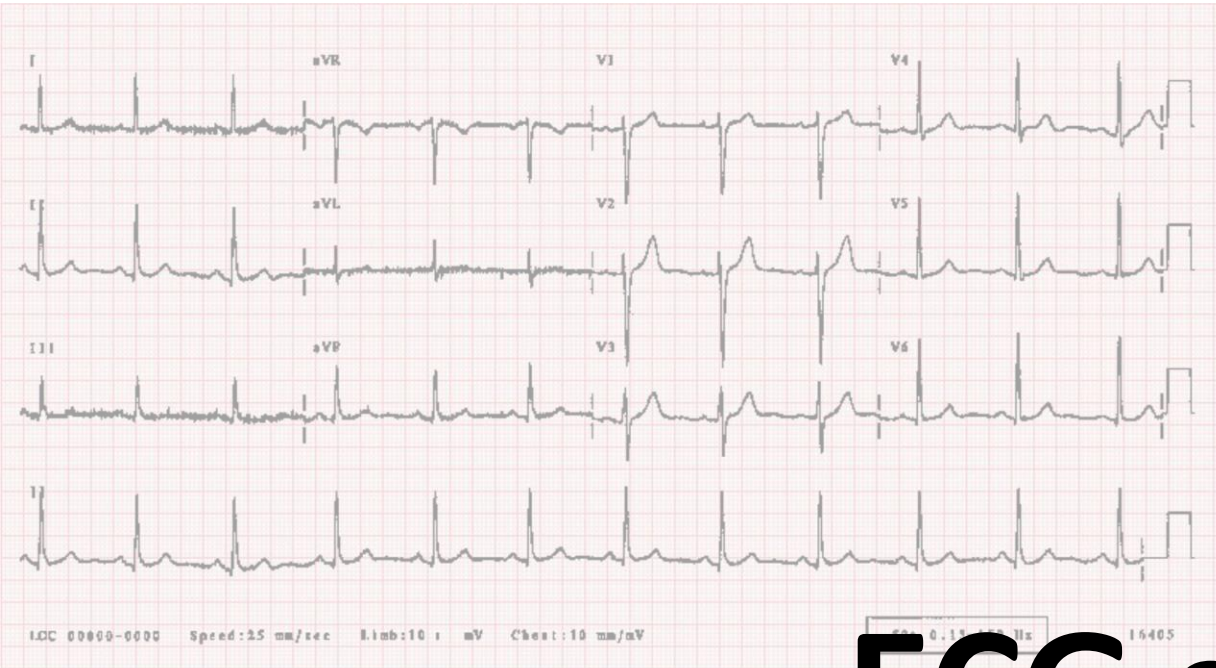
# AML Project 2: Time-series classification

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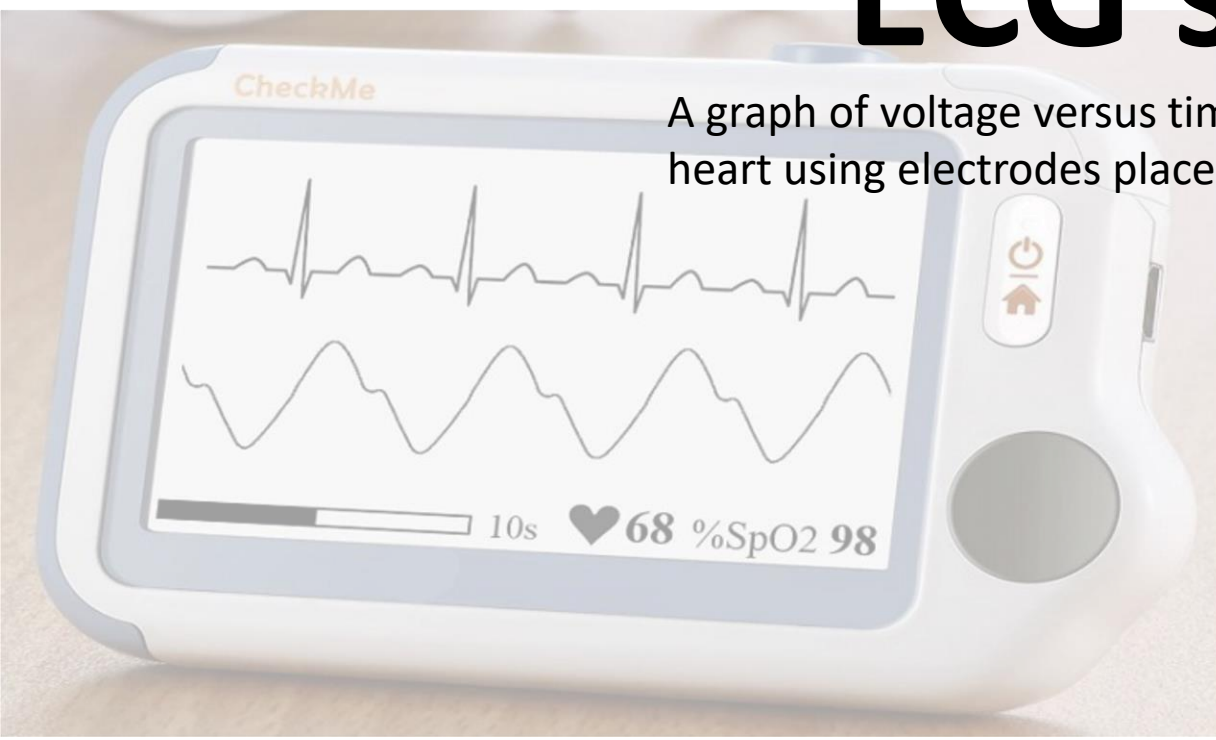
Nov. 11 – Dec. 2

Advanced Machine Learning, Autumn 2024

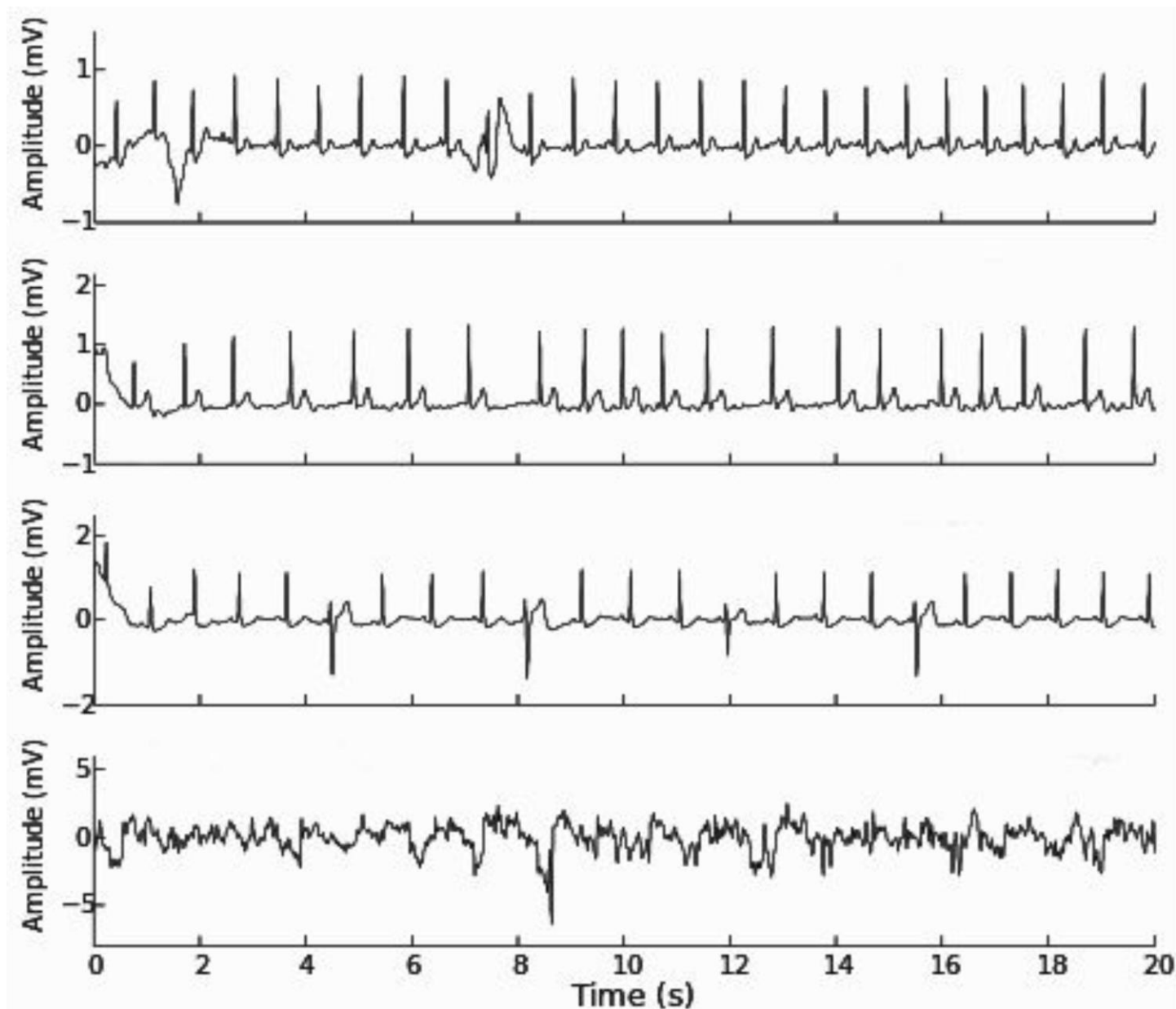


# ECG signal

A graph of voltage versus time of the electrical activity of the heart using electrodes placed on the skin.



## Input (time series)



## Output (classification)

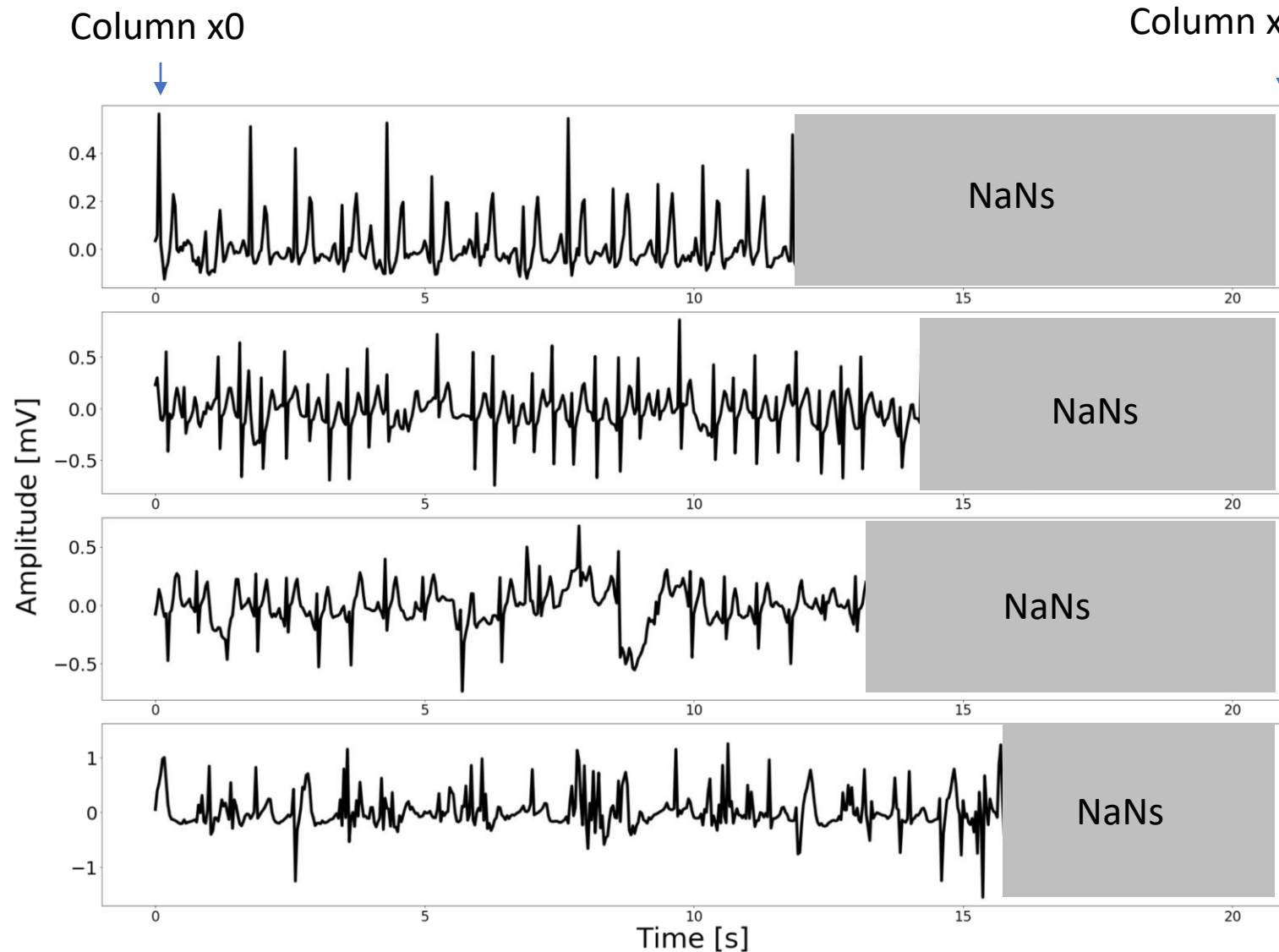
→ **Class 0**

→ **Class 1**

→ **Class 2**

→ **Class 3**

# ⚠ Variable signal length



The ECG data is a raw signal that corresponds to measurements at time stamps 0 to 17806.

A NaN shows up not because of a missing value but because of the variable length of the signal. All signals are sampled in the same frequency of 300Hz.

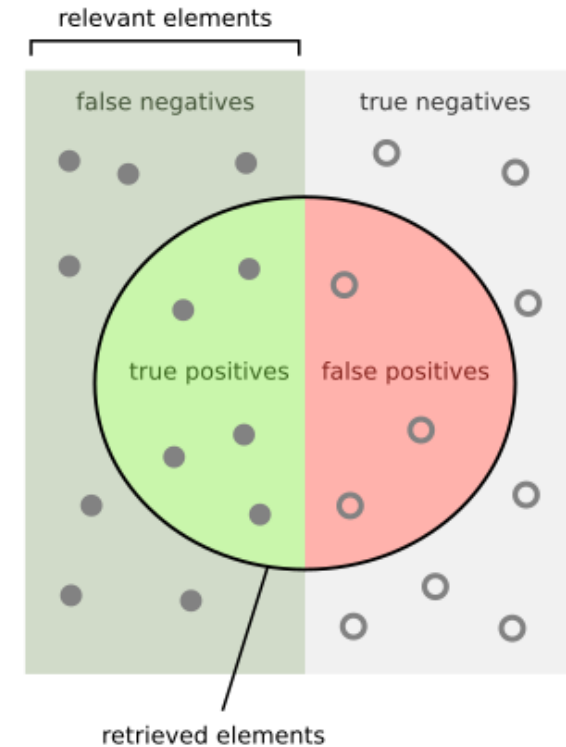
All columns **after the last measurement** have NaN values.

# Evaluation Metrics

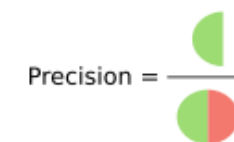
$$F1 = \frac{2 \cdot \textit{precision} \cdot \textit{recall}}{\textit{precision} + \textit{recall}}$$

In python:

```
from sklearn.metrics import f1_score  
  
F1 = f1_score(y_true, y_pred, average='micro')
```

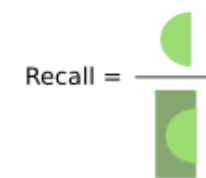


How many retrieved items are relevant?



Precision =

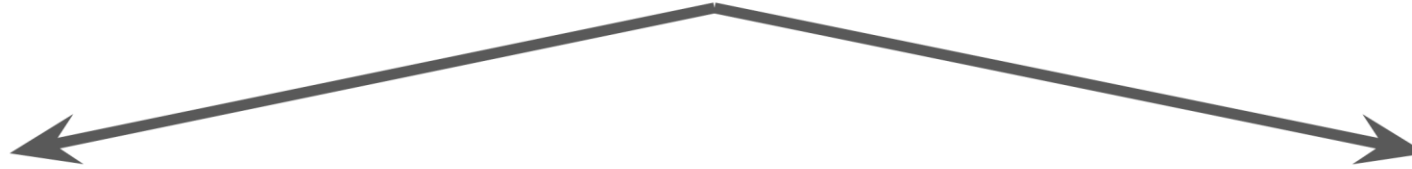
How many relevant items are retrieved?



Recall =

<https://en.wikipedia.org/wiki/F-score>

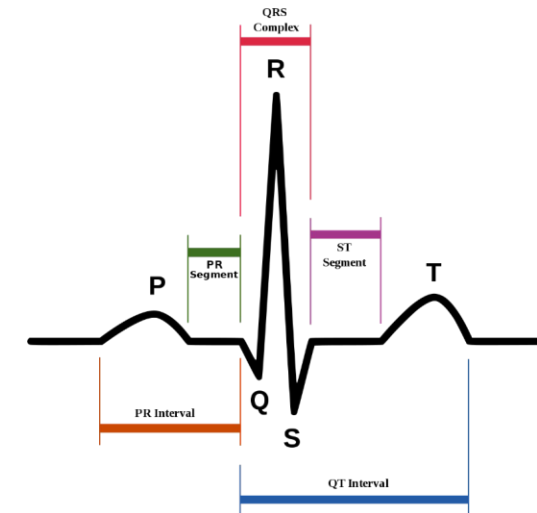
# ECG is a sequence...



... of Data points

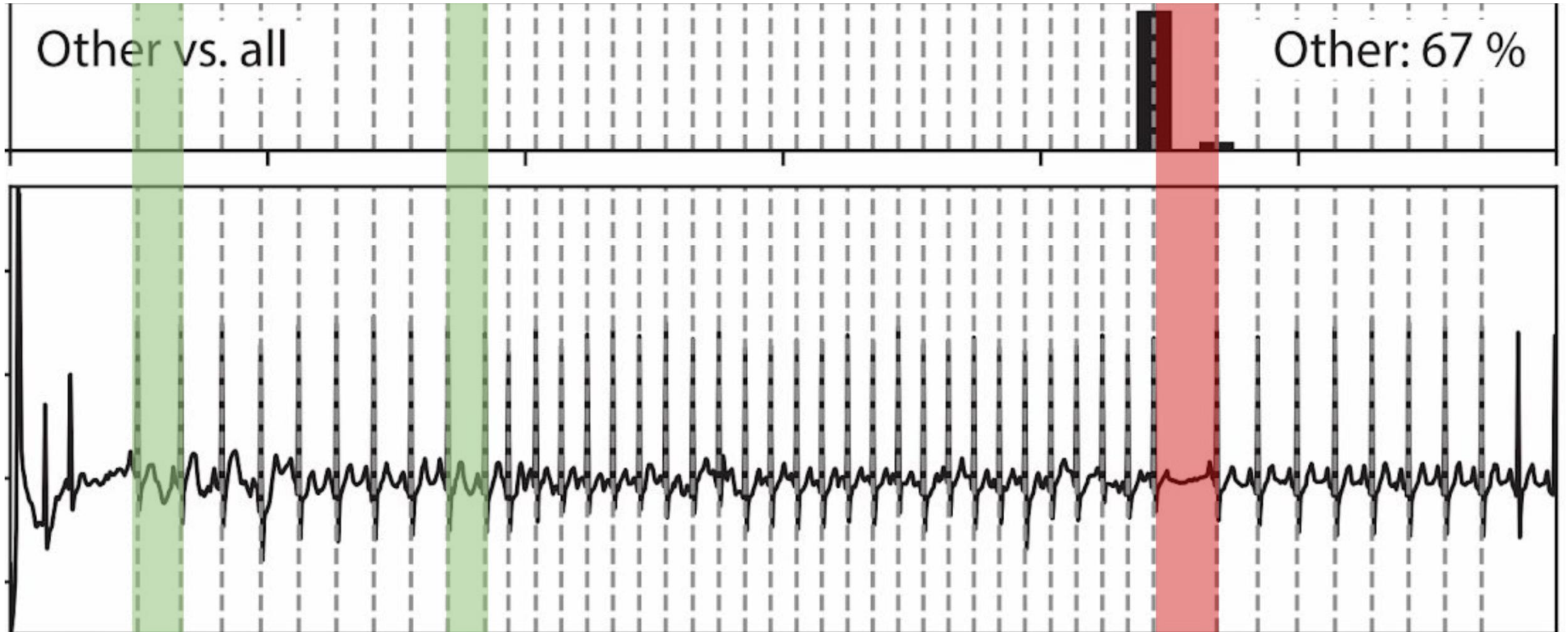


... of Heartbeats



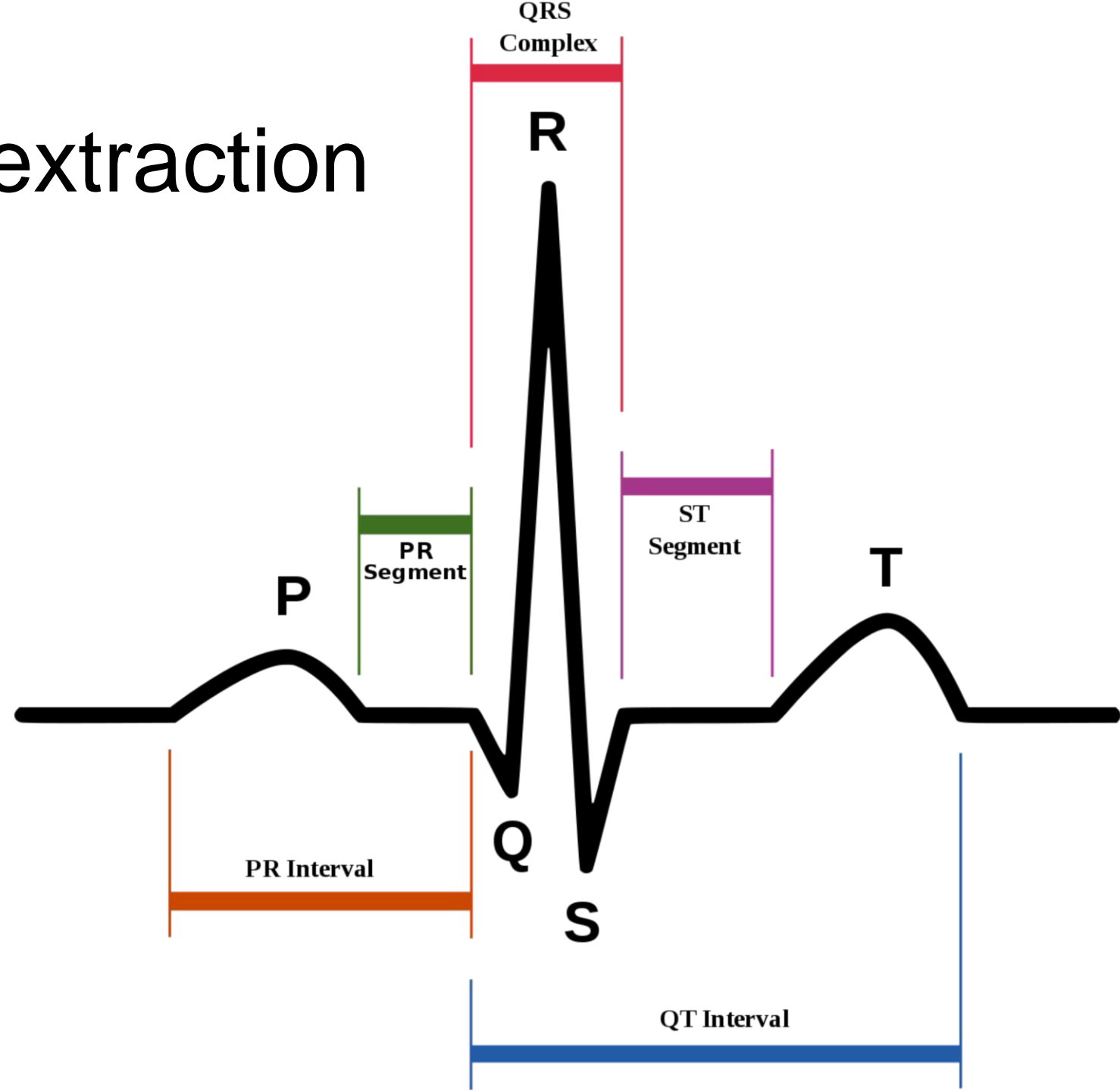


# Splitting into Heartbeats



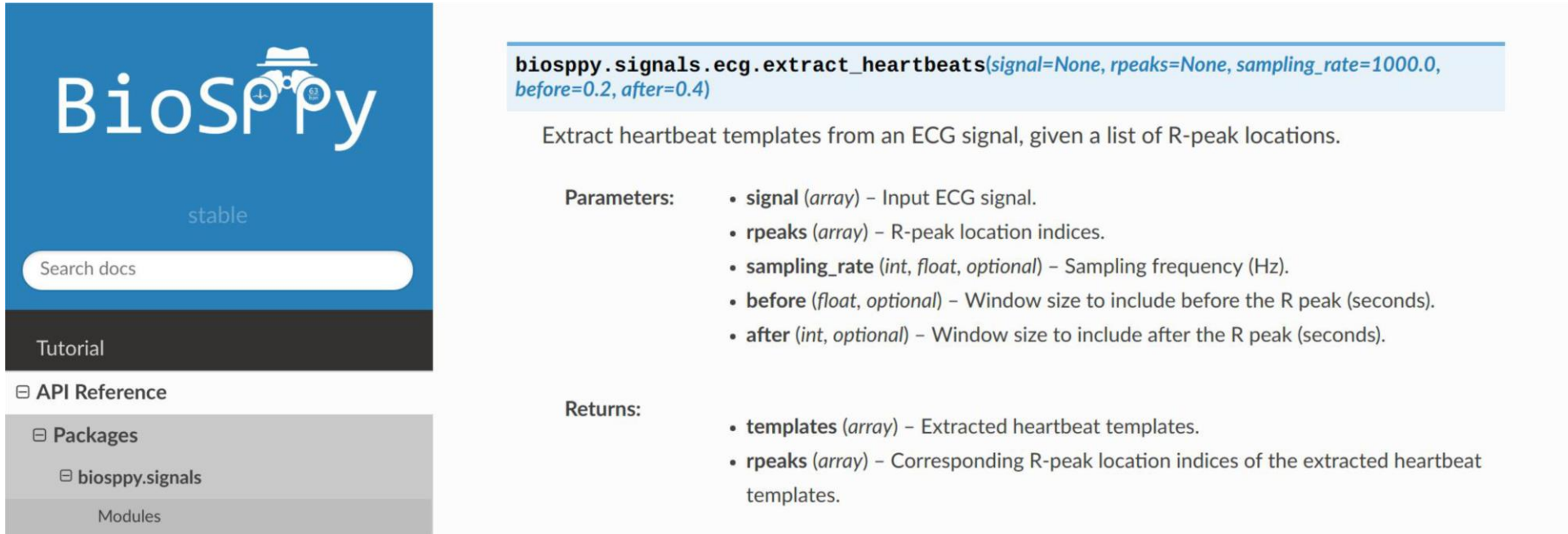
# Manual feature extraction

- RR interval
- R amplitude
- Q amplitude
- QRS duration
- Heart rate variability
- Frequency domain





# import biosppy.signals.ecg as ecg



The screenshot shows the BioSPPy documentation website. On the left is a sidebar with the BioSPPy logo, a search bar, and a navigation menu with links to Tutorial, API Reference, Packages, biosppy.signals, and Modules. The main content area displays the function `biosppy.signals.ecg.extract_heartbeats` with its signature: `(signal=None, rpeaks=None, sampling_rate=1000.0, before=0.2, after=0.4)`. Below the signature is a description: "Extract heartbeat templates from an ECG signal, given a list of R-peak locations." This is followed by a "Parameters:" section with a bulleted list of arguments: `signal` (array) - Input ECG signal, `rpeaks` (array) - R-peak location indices, `sampling_rate` (int, float, optional) - Sampling frequency (Hz), `before` (float, optional) - Window size to include before the R peak (seconds), and `after` (int, optional) - Window size to include after the R peak (seconds). A "Returns:" section follows with a bulleted list: `templates` (array) - Extracted heartbeat templates, and `rpeaks` (array) - Corresponding R-peak location indices of the extracted heartbeat templates.

`biosppy.signals.ecg.extract_heartbeats(signal=None, rpeaks=None, sampling_rate=1000.0, before=0.2, after=0.4)`

Extract heartbeat templates from an ECG signal, given a list of R-peak locations.

**Parameters:**

- `signal` (array) – Input ECG signal.
- `rpeaks` (array) – R-peak location indices.
- `sampling_rate` (int, float, optional) – Sampling frequency (Hz).
- `before` (float, optional) – Window size to include before the R peak (seconds).
- `after` (int, optional) – Window size to include after the R peak (seconds).

**Returns:**

- `templates` (array) – Extracted heartbeat templates.
- `rpeaks` (array) – Corresponding R-peak location indices of the extracted heartbeat templates.

[https://biosppy.readthedocs.io/en/stable/biosppy.signals.html#biosppy.signals.ecg.extract\\_heartbeats](https://biosppy.readthedocs.io/en/stable/biosppy.signals.html#biosppy.signals.ecg.extract_heartbeats)

***We provide you an example Jupyter notebook using biosppy!  
(you will download it with the data from Kaggle)***

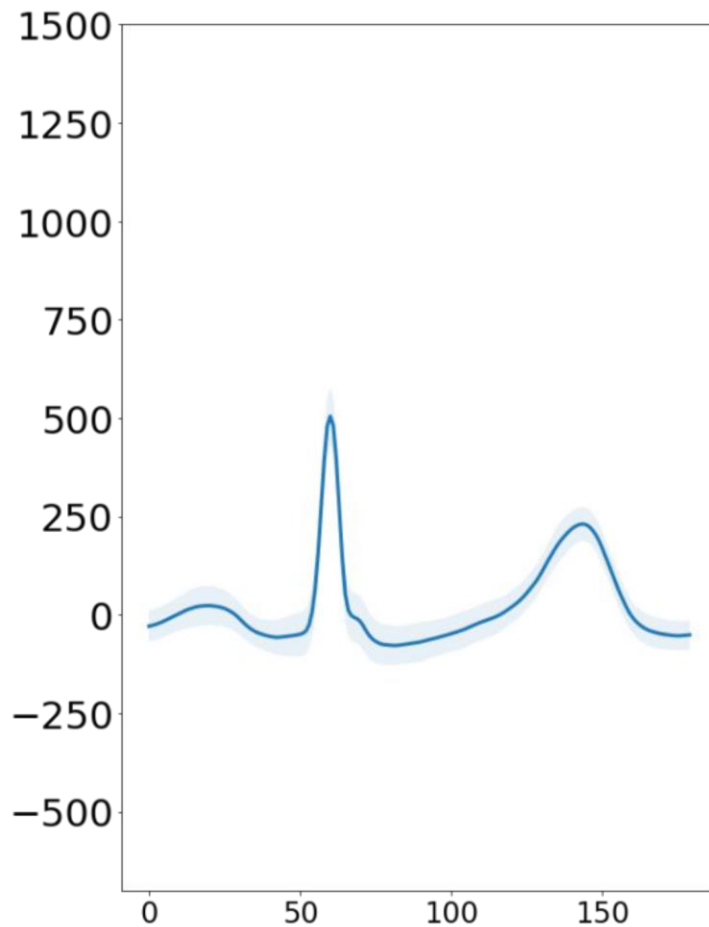
But feel free to experiment with other libraries of your choice:

Some other Python libraries: neurokit, pyhrv, hrv, heartpy, etc...

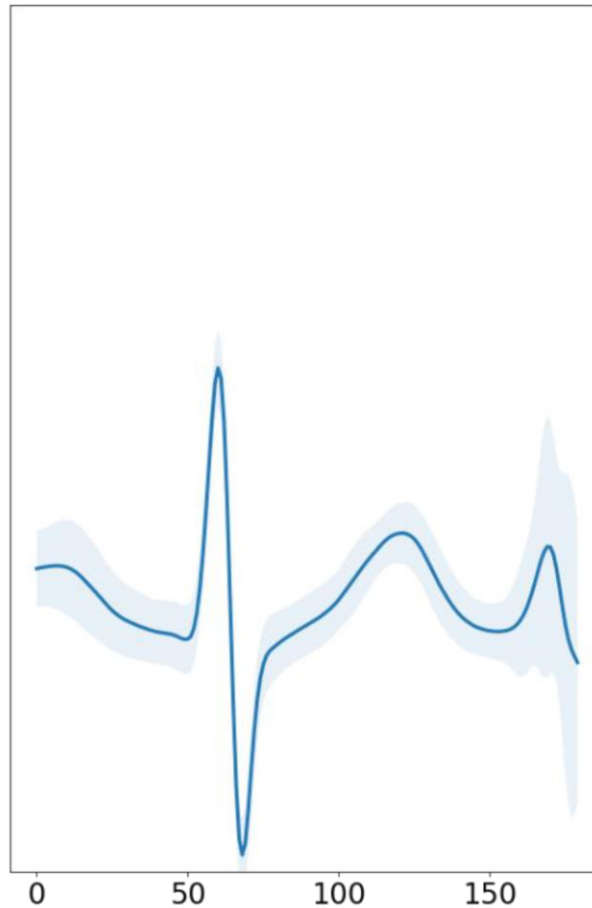
Wave extraction: ecg-kit (Matlab/Octave), ecgpuwave (Fortran, partially ported to Matlab/Octave)

# Mean heartbeat with variance

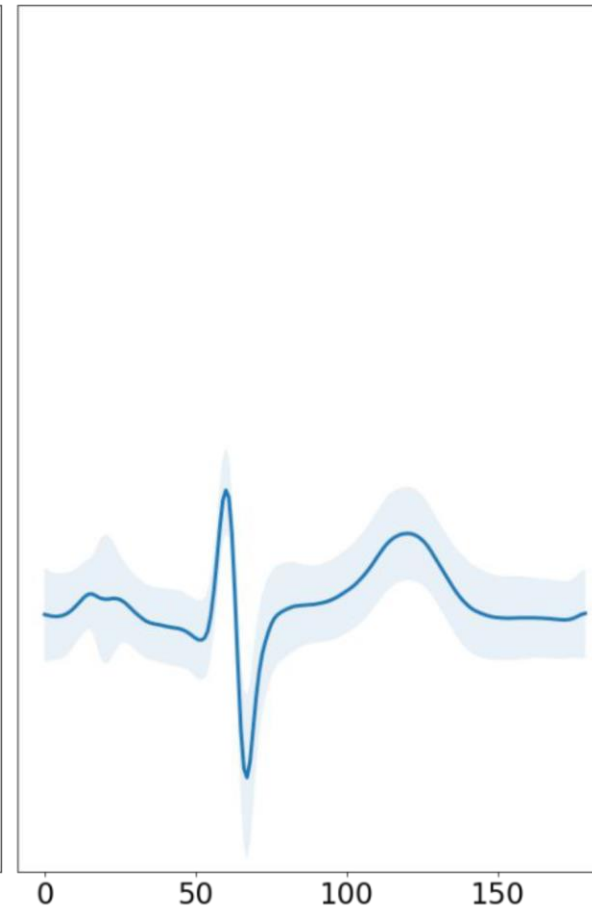
**Class 0**



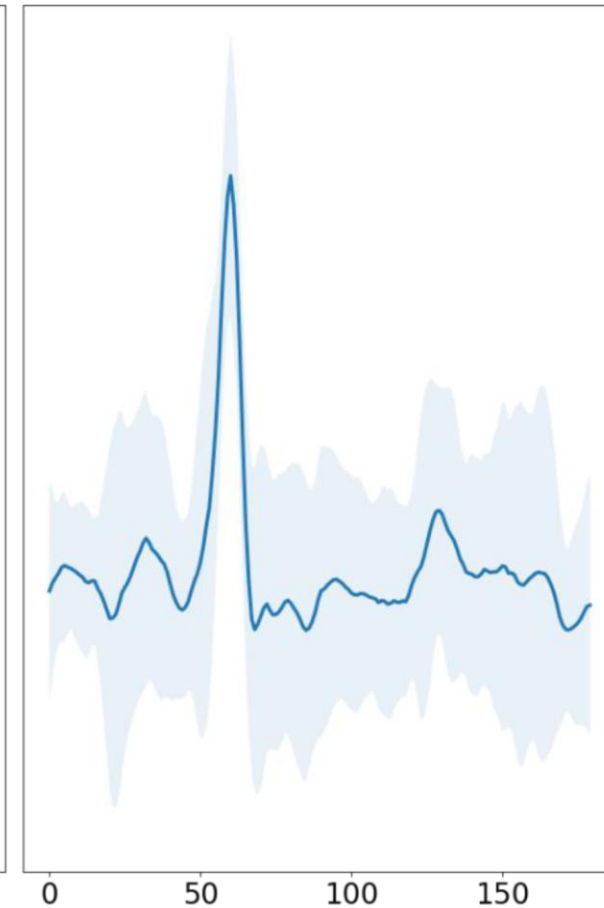
**Class 1**



**Class 2**

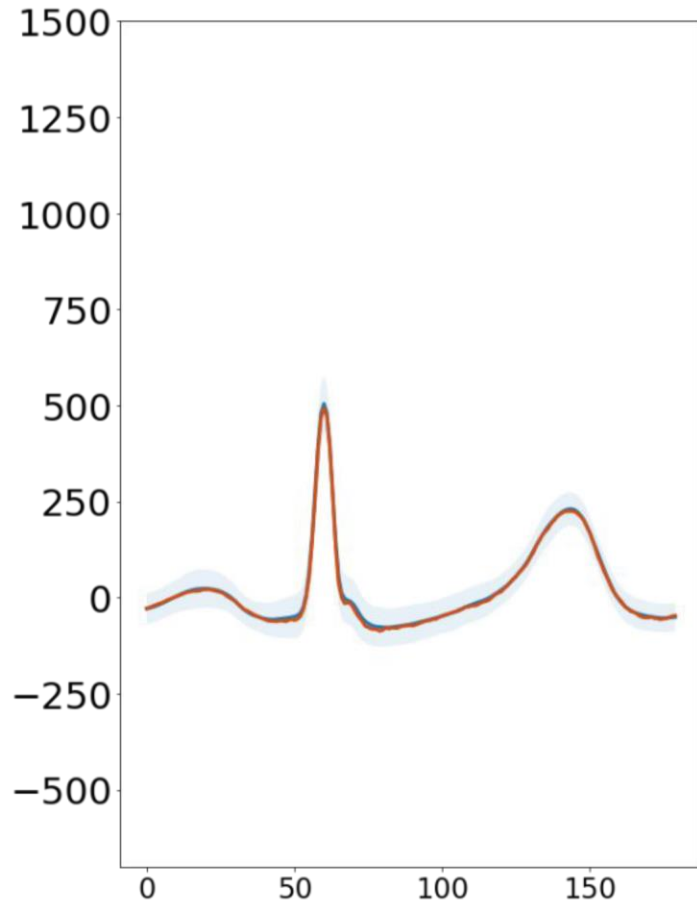


**Class 3**

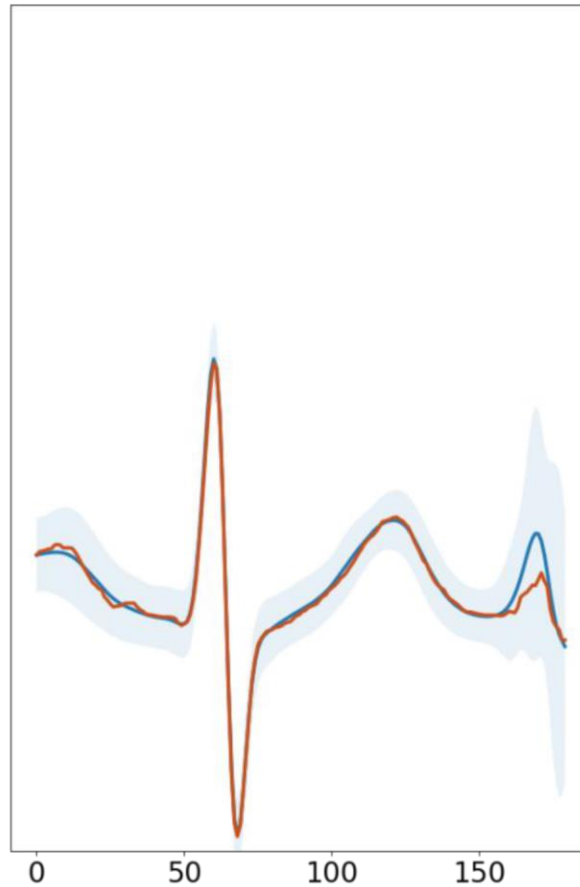


# Mean heartbeat with variance and median

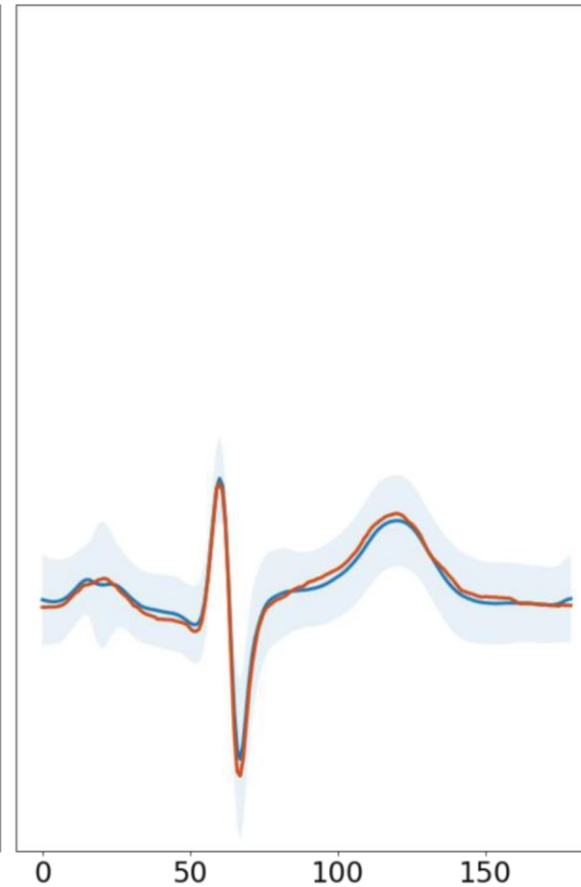
**Class 0**



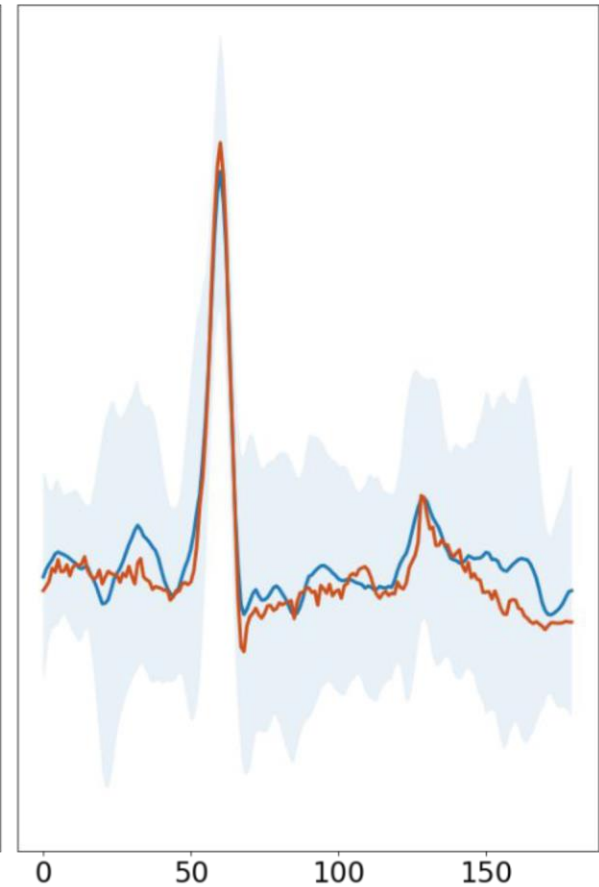
**Class 1**



**Class 2**



**Class 3**



# Submission to Kaggle

- [Kaggle link](#) (you must log in Kaggle with your ETH email)
- Team names must be alphanumeric (A-Z, a-z, 0-9).
- **Each** member of a team must fill out the [Project 2 form](#) in Moodle **separately**.
  - Name of the team in Kaggle, legis, and a description of how you solved the task.
- You only submit a test prediction file to Kaggle, not code. But make sure your results are reproducible. We might ask for your code after the deadline.
- Project period: Nov 11th 3pm – Dec 2nd 2pm
- Public/private leaderboard:
  - Public leaderboard is available during the competition
  - Private leaderboard will open from Dec 2nd 3pm
- Public baseline for passing the project: 0.7
- Check the “Competition Rules” section on the Kaggle competition for more FAQs.

# Moodle project form submission

To obtain points for this project, you have to describe the following in your Moodle form submission:

- You need to select one of your group's submissions for grading. You will only be graded based on this submission.
- You have to write a short description of the approach that you have used. Each student has to write their own description individually, and you **are not allowed to share the description** with your other group members.

If your submission does not include the points above, you will receive zero points for the project.

# Considerations

- Do not use AutoML packages. if in doubt, always ask!
- Be aware of overfitting on the public test set
- Be careful about the submission time
  - Your group has a joint total **10** submissions per day.
- Describe what you did when you hand in the project
- Keep your implementation reproducible for potential review
- Do not wait until the last day to submit something
  - Servers usually get overloaded and crash causing long waiting times



Questions?