# **Supplementary Material**

# Semi-supervised learning for automatic atrial fibrillation detection in

## 24-hour Holter monitoring

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### **Supplementary Methods**

### 1, The K-means method combined with supervised learning.

The details of the supervised learning method with K-means clustering are described below:

- Step 1: Two centroids were obtained from the positive labeled samples and negative labeled samples, respectively.
- Step 2: Measuring the Euclidean distance between the unlabeled sample and the two centroids, and the unlabeled sample was clustered to the centroid that had the shorter Euclidean distance.
- Step 3: After all the unlabeled samples were clustered, two new centroids were obtained using the labeled data and the clustered data.
  - Step 4: Repeat Steps 2 and 3 until the clustering results did not change.
  - Step 5: The labeled data and the clustered data were used to train the model.
- Step 6: Apply the trained model to the same testing data as in our semi-supervised learning method were used.

## **Supplementary Tables**

## Supplementary Table I. Overview of training and testing datasets in this study.

Dataset	The number of the patients	The number of AF episodes	AF duration/min	The number of positive samples	The number of negative samples
Labeled training set	10	145±74	4067±585	3815±450	7074±822
Unlabeled training set	790	8088±257	314 457±1777	314 090±2369	549 400±2322
Testing set	200	2058±311	79 631±1516	79 476±2228	139 118±2782

## Supplementary Table II. Parameters of the CNN-LSTM model.

Layer Name	Input size	Output size	Number of parameters
Input		90, 1	0
Conv layer	90, 1	90, 64	384
Conv layer	90, 64	90, 32	6176
Bi-LSTM	90, 32	90, 200	106 400
Global max pooling	90, 200	200,	0
Dropout	200,	200,	0
FC layer	200,	32,	6432
Dropout	32,	32,	0
FC layer	32,	16,	528
Output	13,	2,	34

## Supplementary Table III. The pseudocode of the mixed mean teacher algorithm.

#### Algorithm: Mixed mean teacher

- 1. **Input:** Labeled data and the corresponding labels  $(X, Y) = \{(x_b, y_b); b \in (1, \dots, B_1)\}$ , unlabeled data  $U = \{(u_b); b \in (1, \dots, B_2)\}$ , training epochs es, Initialize model S  $S(\theta_s)$  and model T  $T(\theta_t)$ with the same initial parameters.

```
2. for e = 1 to es do
    for u, x, y in U, X, Y do
       u' = Noise(u) // Apply Gaussian noise as disturbance to u
       l_u = T_{model}(u'; \theta_t) // "l_u" refers to guess labels
       p_u = S_{model}(u; \theta_s)
       l_u = Update(l_u) // Modified guess labels (see eq.4)
       l_u = Sharpen(l_u) // Reduce the entropy of guess labels (see eq.5)
       loss_1 = w_1 * MSE(p_w l_u) // Consistency loss (see eq.2), w_1 is the weight
       loss_2 = w_2 * corssentropy(S_{model}(x; \theta_s), y) / Classification loss (see eq. 1), w_2 is the weight
       optimize S using loss<sub>1</sub> and loss<sub>2</sub>
       optimize T using EMA // See eq.3
   end for
end for
```

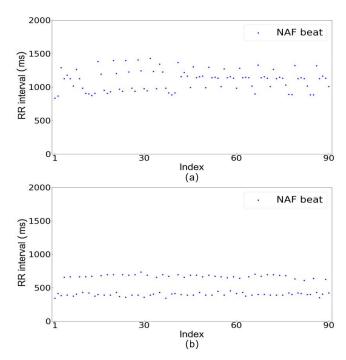
3. **Output:** well-trained model *S*.

# $\label{thm:continuous} \textbf{Supplementary Table IV. Overview of the MIT-BIH\ AF\ database.}$

Dataset	The number of Recordings	The number of AF episodes	AF duration/min	The number of positive samples	The number of negative samples
AFDB	23	291	5492	5870	6655

<sup>\*</sup>Recordings 00735 and 03665 were excluded.

# **Supplementary Figure**



Supplementary Fig. 1. Two examples of the samples that were misclassified by our semi-supervised learning method but correctly detected by the "whole training" method. NAF = non-atrial fibrillation.