# MVKT-ECG: Efficient Single-lead ECG Classification on Multi-Label Arrhythmia by Multi-View Knowledge Transferring

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### **Abstract**

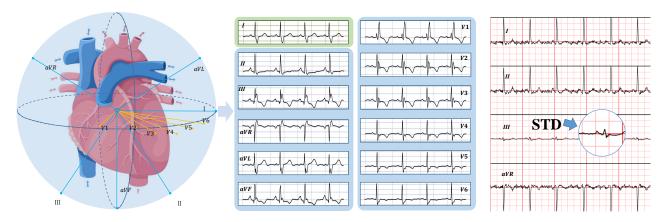
The widespread emergence of smart devices for ECG has sparked demand for intelligent singlelead ECG-based diagnostic systems. However, it is challenging to develop a single-lead-based ECG interpretation model for multiple diseases diagnosis due to the lack of some key disease information. In this work, we propose inter-lead Multi-View Knowledge Transferring of ECG (MVKT-ECG) to boost single-lead ECG's ability for multilabel disease diagnosis. This training strategy can transfer superior disease knowledge from multiple different views of ECG (e.g. 12-lead ECG) to single-lead-based ECG interpretation model to mine details in single-lead ECG signals that are easily overlooked by neural networks. MVKT-ECG allows this lead variety as a supervision signal within a teacher-student paradigm, where the teacher observes multi-lead ECG educates a student who observes only single-lead ECG. Since the mutual disease information between the single-lead ECG and muli-lead ECG plays a key role in knowledge transferring, we present a new disease-aware Contrastive Lead-information Transferring(CLT) to improve the mutual disease information between the single-lead ECG and muli-lead ECG. Moreover, We modify traditional Knowledge Distillation to multi-label disease Knowledge Distillation (MKD) to make it applicable for multi-label disease diagnosis. The comprehensive experiments verify that MVKT-ECG has an excellent performance in improving the diagnostic effect of single-lead ECG.

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### 1. Introduction

Electrocardiogram (ECG) is a commonly used, non-invasive, and convenient diagnostic method for detecting Cardiac arrhythmia and other cardiovascular conditions (Holst et al., 1999). It can be divided into two categories: Multi-lead (12lead system standard in clinical) ECG and single-lead ECG. Multi-lead ECG is acquired by several electrodes arranged on the patient's chest wall and limbs, which can be seen as visualizing the heartbeat signals in multiple different views (Figure 1(a)) (Chen et al., 2021b; 2022). Under the strong ability of deep learning, many multi-lead-based deep ECG interpretation models are widely used in practice. Previous models focused on extracting discriminative features (Kiranyaz et al., 2016; Ribeiro et al., 2020; Wang et al., 2019b). Subsequently, the residual connection is widely used in deeper convolutional network structures to increase the expressive power of the model(Ribeiro et al., 2020). To exploit the properties of different leads, a series of deep neural networks are proposed to establish an attention mechanism between leads (Xia et al., 2019; Liu et al., 2021). More recently, faced with the shortage of high-quality ECG datasets, many self-supervised approaches focus on mining effective information from unlabeled data (Kiyasseh et al., 2021; Chen et al., 2021a; Sarkar & Etemad, 2020). However, this kind of high-quality multi-view ECG data is provided by an environment with strong observation equipment (e.g. hospital) (Hong et al., 2018).

Single-lead electrocardiograms have attracted much attention recently due to the emerging smart ECG devices, such as Apple Watch and Alivecore, which bring great convenience for people. In recent years, many single-lead ECG signals have been produced with the popularization of intelligent ECG devices, raising high expectations for robust intelligent multi-label disease single-lead ECG diagnostic systems. However, the poor observation equipment always provides low-quality and single-view ECG signals (Hong et al., 2018). Due to the lack of views, research on the intelligent diagnosis of single-lead ECG still focuses on a few diseases(Rizas et al., 2022; Hong et al., 2019) (such as atrial fibrillation). Single-lead ECG records the heart's electrical activity from only one perspective, which can be seen as one partial view of a 12-lead ECG. A large gap in



(a) 12-lead ECG from 12 different viewpoints

(b) A ECG waveform with arrhythmia STD (ST-segment depression)

Figure 1. (a): A standard 12-lead ECG signal by depicting heartbeat from 12 different views. Mul-view ECG can record the heart's activity more comprehensively, as it includes rich disease information from various perspective. (b): Lead III of this ECG signal shows the disease STD, while other leads are all normal (only four leads are shown), so judgment based on lead I or other lead (except lead III) alone is likely to lead to missed detection.

arrhythmia diagnosis still subsists between single-lead and multi-lead, indicating that the number of leads plays a key role in achieving better results (see Table. 2).

Recently, Knowledge Distillation methods are proposed to improve single-lead ECG's performance (Sepahvand & Abdali-Mohammadi, 2022; Hong et al., 2018), where the teacher model was already developed through multi-lead ECG signals and the student model was developed through single-lead ECG signals. Since the student is educated under the supervision of the teacher to mimic the output space and inner feature or attention maps of its teacher, the internal feature maps to be processed must be strictly consistent. And this method can only diagnose ECG with single-label. It is noted that mutual information plays a key role in improving single-lead ECG's potential. So limiting consideration on the information relationship between single-lead ECG and multi-lead ECG, just forcing the same output of student with the teacher does not guarantee the robustness of the student.

In this paper, in order to efficiently stimulate the single-lead ECG signals' potential, we propose a novel and efficient architecture to transfer multi-view information of ECG for multi-disease label classification — Multi-View Knowledge Transferring of ECG (MVKT-ECG). We adopt the teacher-student paradigm and extend it to a more general inter-lead knowledge transfer approach. Since mutual information is the key to improving single-lead ECG's ability, we formulate a new objective to maximize the mutual information of the feature representations between teacher and student — Contrastive Lead-information Transferring (CLT) Loss. The CLT can maximize a lower bound to the mutual information between the teacher and student representations.

Further, We modify traditional Knowledge Distillation to multi-label Knowledge Distillation to make it applicable to the context of multi-disease labels. To verify the effectiveness of our method, in this paper, we conduct extensive experiments on two commonly used public datasets, PTB-XL and ICBEB2018. Experimental results demonstrate the effectiveness and robustness of multi-view knowledge transferring in the single-lead ECG classification task. To sum up, the contributions of this paper include the following:

- We propose a more general and efficient framework MVKT-ECG to transfer multi-view information from multi-lead ECG signals to single-lead-based models.
- We explore the nature of inter-lead knowledge transfer and design a novel inter-lead information transferring objective—Contrastive Lead Transferring(CLT) Loss.
- We formulate multi-label Knowledge Distillation (MKD) and apply it to to fit the context of more general multi-disease label detection.

### 2. Related Work

(Hannun et al., 2019) firstly proposed a comprehensive evaluation of an end-to-end deep neural network for ECG analysis across various diagnostic classes, demonstrating that an end-to-end deep learning approach can classify a broad range of distinct arrhythmias from single-lead ECGs with high diagnostic performance similar to that of cardiologists. Due to the limited information provided by single lead ECGs, all kinds of intelligent ECG equipment can only support a few types of detection at present. For example, the

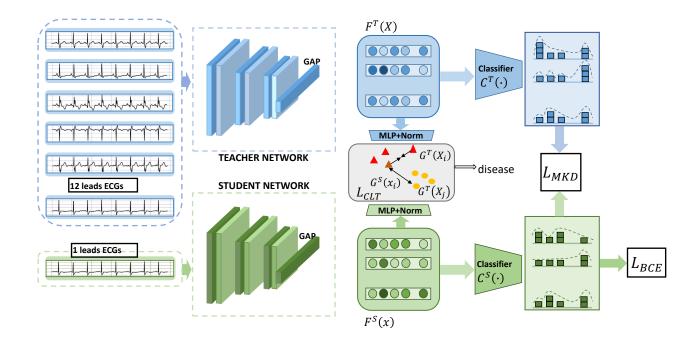


Figure 2. An Overview of Multi-View Knowledge Transferring: the network using single-lead ECG signals is optimized by the network using 12-lead ECG signals. The student network receives the multi-view Knowledge in two ways: (a) minimizing  $\mathcal{L}_{CLT}$  to improve the inter-lead disease mutual information (b) matching the logits of the teacher by minimizing the  $\mathcal{L}_{MKD}$ ;

Apple Watch can only detect sinus rhythm(SR) and atrial fibrillation(AF) (Rizas et al., 2022). And most academic research has also focused on that. Existing studies have achieved high detection accuracy in the detection of AF (Clifford et al., 2017). However, limited by input information, the single-lead model performs poorly in classifying other arrhythmias, such as STD and PAC (Relevant experimental data will be presented in the Appendix).

To address this problem of lacking ECG views, people proposed the method of restruction and synthesis of ECG. (Zhang & Frick, 2019) restored another nine leads based on noisy three-lead signals through LSTM. (Golany et al., 2021) transformed the reconstruction of missing leads into solving the least squares problem by mapping the known ECG leads to the corresponding Koopman space and using the theory that the Koopman operator is linear. (Chen et al., 2021b) proposed the concept of ECG panorama and the corresponding generation network Nef-Net and further proposed disease-aware synthesis method—ME-GAN, which attains panoptic electrocardio representations conditioned on heart diseases. These methods can restructure or synthesize multi-view ECG signals, but lacking evaluation on downstream classification tasks makes it difficult to guarantee the quality of the reconstructed signals to detect diseases.

To achieve model compression of single-lead-based model

and minimize the performance gap between the arrhythmia classification model with multi-lead ECG signals and the arrhythmia classification model with single-lead ECG signals, (Sepahvand & Abdali-Mohammadi, 2022) proposed the teacher-student architecture. In order to deploy models in the poor-data environment without requiring direct access to multi-modal data acquired from a rich-data environment, (Hong et al., 2018) proposed a knowledge distillation (KD) method (RDPD) to enhance a predictive model trained on poor data using knowledge distilled from a high-complexity model trained on rich, private data. These methods use the traditional method of KD, forcing the student to imitate the output and inner features or attention of the teacher. It realizes the compression of the model through mature knowledge distillation loss and reduces the gap between single-lead ECG and 12-lead ECG. But this method must require the same dimension of the feature map between teacher and student, which makes the network structure lack flexibility. Different from this work, we explore the nature of inter-lead information transferring and formulate two new losses. In addition, our architecture can also deal with the muli-label-disease task, which is more general and efficient.

### 3. Method

### 3.1. Architecture

Here we briefly describe the framework of our method and introduce the notations used in this paper. Given a 12 leads ECG signal  $X = [x^{(1)}, x^{(2)}, ..., x^{(12)}]$  or a single lead ECG signal  $\boldsymbol{x} = x^{(i)}$ , where  $\boldsymbol{X} \in \mathbb{R}^{12 \times L}$  and  $\boldsymbol{x} \in \mathbb{R}^{L}$ . Our models are aimed to learn a function  $\mathcal{F}_{\theta}(X)$ , mapping a set of ECGs  $[X_1, X_2, ..., X_N]$  to a representative features space. Then the classifier  $\mathcal{C}_{\theta}(\cdot)$  will project the above features into category space  $\hat{Y} \in \mathbb{R}^C$ . We consider the medical reality that sometimes patients with heart disease often have complex conditions, often multiple arrhythmias. So we consider the multi-label disease scenario, so the sigmoid function was used as the final activation function. Here, we hope that this kind of representation can 1)minimize the information gap between single-lead ECGs and multi-lead ECGs signals and 2)be robust to the lead reduction in the single-lead ECGs. To achieve this, as shown in Figure 2, our proposed method is a two-stage procedure, as follows.

First step: we train the teacher network using standard 12 leads ECGs  $\boldsymbol{X} = \begin{bmatrix} x^{(1)}, x^{(2)}, ..., x^{(12)} \end{bmatrix}$ . The module  $\mathcal{F}_{\theta}^T(\boldsymbol{X}): \mathbb{R}^{12 \times L} \to \mathbb{R}^D$  will map each ECG signal  $X_i$  to a fixed-size representation  $d_i$  (in xresnet1d101 (Strodthoff et al., 2021) D=4096). We train the teacher network-which will observe 12-lead ECG using the multi-label classification term — Binary Cross Entropy Loss ( $\mathcal{L}_{BCE}$ ), which can be formulated as:

$$\mathcal{L}_{BCE} = -\left[ \boldsymbol{y} \log \hat{\boldsymbol{y}} + (1 - \boldsymbol{y}) \log(1 - \hat{\boldsymbol{y}}) \right] \tag{1}$$

where y represents the one-hot labels and  $\hat{y}$  represents the predicted classes probability.

**Second step**: we use the teacher network trained in the first step to instantiate the new student network, which only observes single-lead ECG, and the teacher's parameters are frozen. When we have trained the teacher model, we hope the representation ability of the network, which only observes single-lead ECGs, can be improved. In order to achieve this, we aim at the information knowledge we can gather from different leads, depicting the same heart condition under different views. When facing a 12-lead ECG classification task, one can often exploit lead viewpoints to provide a variety of appearances for a target disease. We want to teach a network to recover as much full-lead information just from a single-lead ECG. Although some diseases can not be inferred from single-lead ECG signals, what we want to do is to shorten the information gap between single-lead ECG and 12-lead ECG as far as possible, encouraging the student to focus more on key details to some particular diseases and maximum the mutual disease information between single-lead and 12-lead, further to reduce the incidence of misdiagnosis.

Typically, due to the information gap between 12-lead ECG

and single-lead ECG, the teacher network is always more powerful than the student network. This kind of asymmetry between the teacher and the student can produce a distillation objective different from the one due to the differences in model complexity. MVKT-ECG improves the mutual information between the feature representations of teacher and student by the Contrastive Lead-information Transferring (CLT) Loss. In the process of doing so, we also allow the student to imitate the teacher's output from the single-lead, which is a part of the 12-lead. To do it, we refine the Knowledge Distillation loss (Hinton et al., 2015) to a new knowledge distillation objective that can process multi-label disease problems.

### 3.2. Contrastive Lead-information Transferring(CLT)

To improve the inter-lead mutual information, we propose the disease-aware Contrast Lead-information Transferring(CLT) Loss, which can transfer useful disease information by maximizing the mutual information between single-lead ECG and multi-lead ECG.

As we all know, due to the lack of data dimensions, the disease information in the feature extracted by a neural network from single-lead ECG is less compared with 12-lead ECG, but it still contains part disease information. We refer to this information as the mutual information between single-lead ECG and 12-lead ECG. In another aspect, the absence of available data for single-lead ECG signals also tends to trigger a false diagnosis. For example, abnormal T wave morphology in any lead would be clinically diagnosed as a T wave change. In the case of abnormal T waves in other leads, judgment based on lead I alone is likely to lead to missed detection Fig 1(b). We refer to this information that triggers a tendency to misdiagnosis as misleading information.

We know the lower limit of mutual information (Oord et al., 2018):

$$I(G^{T}(X), G^{S}(X))$$

$$= \sum_{i,j} p(G^{T}(X_{i}), G^{S}(X_{j})) \log \frac{p(G^{T}(X_{i}), G^{S}(X_{j}))}{p(G^{T}(X_{i}))(G^{S}(X_{j}))}$$

$$\geq \log(N) + \mathbb{E} \left[ \log \frac{f(G^{T}(X_{i}), G^{S}(X_{j}))}{f(G^{T}(X_{i}))(G^{S}(X_{j}))} \right]$$

$$= \log(N) + \mathbb{E} \left[ \log \frac{exp\left[ (G^{T}(X_{i})G^{S}(X_{j}))/\tau \right]}{\sum exp\left[ (G^{T}(X_{i})G^{S}(X_{j}))/\tau \right]} \right]$$

$$= \log(N) - \mathcal{L}_{CLT}$$
(2)

The key idea of contrastive learning is learning a representation that is close for "positive" pairs and pushing apart the representation between "negative" pairs. In detail, for samples  $X_i$  and  $X_j$ , MVKT-ECG shortens the student model and teacher model's representation of the same sample  $G^S(x_i)$  and  $G^T(X_i)$  and further the feature representation

of the two models for different samples, such as  $G^S(X_i)$  and  $G^T(X_j)$ . This can be achieved by minimizing:

$$\mathcal{L}_{\text{CLT}} = -\mathbb{E}\left[\log\frac{exp\left[\left(G^{T}(X_{i})G^{S}(X_{j})\right)/\tau\right]}{\sum exp\left[\left(G^{T}(X_{i})G^{S}(X_{j})\right)/\tau\right]}\right]$$
(3)

N is the number of positive and negative sample pairs; the larger the better. Then a memory buffer is used to store the features of all training samples in this study. Since the teacher has observed multi-view ECGs, we believe that the different distances between "positive" and "negative" pairs yield a strong description of corresponding disease identities. During the training, N samples are randomly selected to calculate the loss so that it can meet the demand of the model for the number of negative samples. The expression of each sample in the memory will carry on momentum updates according to the latest feature extracted from the network.

### 3.3. Multi-label diseases Knowledge Distillation(MKD)

The traditional KD focuses on single-label classification problems, so we propose MKD, a new knowledge distillation objective that can process multi-label disease problems. Especially, we separate the output probability  $p_i$  with  $1-p_i$ , and for each sample, we calculate the fore-and-aft softmax by temperature coefficient  $\tau_{kd}$ .

$$q_i = \frac{e^{p_i/\tau_{kd}}}{e^{p_i/\tau_{kd}} + e^{(1-p_i)/\tau_{kd}}} \tag{4}$$

$$\mathcal{L}_{\text{MKD}} = \tau_{kd}^2 \sum_{i} \text{KL}\left(q_i^T || q_i^S\right)$$
 (5)

where  $\tau_{kd}$  represents the temperature coefficient of multilabel disease knowledge distillation loss.

### 3.4. Student Optimization

In summary, the MVKT-ECG overall objective combines the distillation terms ( $\mathcal{L}_{\mathrm{MKD}}$ ), the Contrastive Leadinformation Transferring terms( $\mathcal{L}_{\mathrm{CLT}}$ ), and the ones optimized by hard label- $\mathcal{L}_{\mathrm{BCE}}$ , which provide a higher conditional likelihood w.r.t. ground truth labels. To sum up, MVKT aims to boost single lead ECG's representation ability by the following optimization problem:

$$\operatorname{argmin}_{\theta_{S}} \mathcal{L}_{\text{MVKT}} = \mathcal{L}_{\text{BCE}}^{S} + \alpha \mathcal{L}_{\text{MKD}} + \beta \mathcal{L}_{\text{CLT}}$$
 (6)

where  $\alpha$ ,  $\beta$  are hyperparameters balancing the contributions of  $\mathcal{L}_{\mathrm{MKD}}$  and  $\mathcal{L}_{\mathrm{CLT}}$  to the total loss  $\mathcal{L}_{\mathrm{MVKT}}$ . Regarding the student initialization, we found that the CLECG's self-supervised strategy (Chen et al., 2021a) is beneficial, which will be mentioned in Sect. 5.5.

### 4. Dataset

To build robust and efficient single-lead ECG interpretation models in a multi-label disease context. So we conduct experiments on two multi-label freely accessible datasets. ICBEB2018 dataset(Liu et al., 2018) contains 6,877 12lead ECG recordings, each ranging in length from 6 to 60 seconds. The dataset considered one normal and eight abnormal arrhythmia categories, including AF, I-AVB, LBBB, PAC, PVC, RBBB, STD, and STE. Each record may have more than one label. This paper follows the processing method in (Liu et al., 2018) to divide the dataset into ten folds by stratified sampling, and the original label distribution is maintained in each fold. Among them, the first eight were used as training sets, while the 9th and 10th were used as verification and test sets, respectively. Then, we preprocess them to get equal-length data, and the length n = 10000.

PTB-XL dataset(Wagner et al., 2020) is the to-date largest freely accessible clinical 12-lead ECG-waveform dataset comprising 21,837 records from 18,885 patients of 10 seconds long. This work uses all signals down-sampled at 100 Hz as the labeled data source. This dataset provides rich multi-level annotations, which, in terms of diagnosis, include superclasses of 5 classes and subclasses of 24 classes. We followed the Settings in (Wagner et al., 2020) to divide the dataset into ten folds of class-balanced, with the first eight folds as the training set and the 9th and 10th folds as the verification set and test set, respectively. The signal length of each sample is 1,000 points.

### 5. Experiments

**Evaluation Metrics.** In the followings, we report performance in terms of Area under the Receiver Operating Characteristic (ROC-AUC) and the F1 score.

### 5.1. Experimental setups

We implemented the MVKT algorithm in the framework of Pytorch. During the training, the model was trained according to the two-stage teacher-student training process, and the 12-lead data was first used to train the teacher model. All the teacher networks are trained for 100 epochs using Adam (Kingma & Ba, 2014). During the information transferring period, the parameters of the teacher model are frozen, and only the student model is updated. We feed 12-lead ECG signals to the teacher and single-lead ECGs to the student. In the process of information transferring, the temperature coefficient of knowledge distillation  $\tau_{KD}$  is set as 1.5 because the dataset we used contains not many categories and the network prediction scores for each category are not concentrated. The temperature coefficient in the loss of comparative representation  $\tau$  is set as 0.07, and the di-

Table 1. Comparision MVKT's performance with State-of-the-Art single-lead ECG interpretation methods on public datasets. Since KD
loss is only applicable to single-label classification, we use the original output KL-div when computing KD loss on these two data sets.

Method	ICBEB2018			PTB	XL.subd	iagnostic	PTBXL.superdiagostic			
Wichiod	AUC	ACC	F1-score	AUC	ACC	F1-score	AUC	ACC	F1-score	
resnet1d_Liu	90.2	92.2	65.6	84.9	88.5	48.8	82.6	80.2	60.9	
DenseCNN_wang	88.7	90.6	61.7	84.5	88.3	47.9	82.7	81.1	61.7	
SEresnet18	93.5	94.4	72.5	85.0	88.8	48.5	82.7	81.3	61.6	
SEresnet34	93.7	94.4	73.1	84.4	89.3	47.7	82.9	81.8	62.3	
CNN_Hannun	94.0	94.7	74.6	84.9	89.2	48.9	83.1	80.8	61.4	
KD+FitNet	94.0	94.8	74.8	83.6	87.2	46.4	83.8	76.5	60.1	
KD+AT	92.8	93.8	71.4	82.0	86.1	44.2	80.0	78.5	59.8	
MVKT-ECG	95.7	95.7	78.0	86.1	88.3	58.3	84.3	82.2	62.6	

mension of teacher and student representation d is 128 in all structures. The size of the training batch is 32. In each training round, 1024 samples are randomly selected from the memory storage area to calculate the loss- $\mathcal{L}_{\text{CLT}}$ . In the concrete realization, the negative samples are extracted with the student output and the teacher output as the anchor points, respectively. The loss is composed of two symmetric parts. In the concrete realization, the negative samples are extracted with the student output and the teacher output as the anchor points, respectively. The loss is composed of two symmetric parts.

# 5.2. Comparision with State-Of-The-Art Single Lead ECG's Classification Methods

Table 1 reports a thorough comparison with current state-ofthe-art (SOTA) methods across datasets. In the Physionet Challenge 2017 competition (Clifford et al., 2017), Hannun's CNN based model (Hannun et al., 2019) achieved the best score. The DenseCNN\_wang (Wang et al., 2019a), and resnet1d\_Liu (Liu et al., 2019) achieved the top performance in China ECG AI Contest 2019 competition. (Zhao et al., 2020) proposed a large kernel size model SEresnet\_Zhao based on SE-block (Hu et al., 2018), achieving second place in the PhysioNet 2020 competition (Alday et al., 2020). (Sepahvand & Abdali-Mohammadi, 2022) (KD+FitNet) and (Hong et al., 2018) (KD + attention) also use the distillation idea to bridge the gap between the model with multilead ECG signals and single-lead ECG signals by trying to make the student's output and inner features the same as the teacher's. Compared with KD, Our proposed method performs better and outperforms other SOTA single-lead models. This result is fully consistent with our goal of robustness when providing only a single lead ECG as a query.

### 5.3. MVKT-ECG on different backbones

We indicate the baseline and teacher model with the name of the backbone and append "MVKT" for the model after the MVKT (e.g. ResMVKT34). We first benchmark 5 SOTA models' performance when they observe 12-lead ECGs and single-lead ECGs, respectively. The 5 models: CNN\_Hannun (Hannun et al., 2019), ResNet1d34 (He et al., 2016), ResNet1d\_wang (Wang et al., 2017), inception1d (Ismail Fawaz et al., 2020), XresNet1d101 (Strodthoff et al., 2020) are set as our backbones. The backbones' performance on single-lead ECGs is set as our baseline. Table 2 reports the comparison for different backbones. Table 4 and Table 5 demonstrates detail promotion of each disease. In detail, the student XresMVKT1d101 outperforms its baseline observably (3.2 % on ICBEB2018, 1.3 % on PTB.super, and 1.4 % on PTB.sub). Based on this, we draw the following conclusions: 1) according to the objective which the student seeks to optimize, our method can gain significant improvement when single-lead ECGs are available; 2) compared with the baseline, the students do gain improvement, but it is hard for them to exceed the teacher's performance. As far as we can tell, the single-lead ECGs contain only a fraction of the 12-lead ECGs information that the MVKT can elicit to the greatest extent, but some pieces of information which single-lead ECG drops can never be recovered.

As additional proof, plots from Figure 3 draw a comparison between models before and after information transferring. MVKT-ECG improves the performance considerably on ICBEB2018 dataset for each selected lead. Surprisingly, the lead II, AVR and limb lead (always v3–v6) always perform better. Surprisingly, MVKT-ECG can make lighter single-lead networks superior to complex models several times deeper: as an example, ResMVKT34 scores better than

Table 2. MVKT's results on different datasets, settings, and architectures.

	backbone	AUC (%)					
	backbone	PTB.sub	PTB.super	ICBEB2018			
Teacher	CNN_Hannun	92.3	92.0	97.0			
1 lead	CNN_Hannun	84.8	83.3	94.0			
1 leau	CNN_Hannun_MVKT	85.6	83.4	95.7			
Teacher	ResNet1d34	92.3	92.8	94.5			
1 lead	ResNet1d34	84.5	82.6	90.7			
1 lead	ResMVKT34	85.2	83.8	93.9			
Teacher	ResNet1d_wang	92.7	92.1	91.0			
1 lead	ResNet1d_wang	84.9	83.1	87.3			
	ResMVKT_wang	85.8	84.0	89.3			
Teacher	Inception1d	93.2	92.4	93.8			
1 lead	Inception1d	85.2	83.2	88.1			
1 Icau	InceptionMVKT	86.1	84.3	90.7			
Teacher	XresNet1d101	92.1	93.7	95.3			
1 lead	XresNet1d101	83.9	82.7	92.6			
1 icau	XresMVKT101	85.3	84.0	94.6			

even XresNet101 on ICBEB2018, regardless of the selected lead of ECGs.

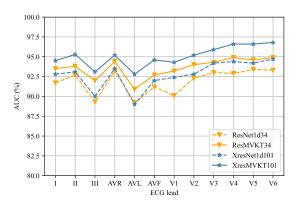


Figure 3. Performance (AUC) at evaluation time when changing the lead of ECG in ICBEB2018 dataset.

### 5.4. Analysis on MVKT-ECG

### The impact of Loss Terms.

We conducted a thorough ablation study for the loss terms. Without loss of generality, we focus our analysis on Xres-MVKT101 on the ICBEB2018 dataset. Table 3 reports the ablation's result. In the three losses,  $\mathcal{L}_{\mathrm{CLT}}$  plays an important role. Although we only use the hard label and  $\mathcal{L}_{\mathrm{CLT}}$ , it can also achieve a significant result, which also demonstrates our proposed role of information transferring between leads. And as expected, the greatest performance of AUC is obtained with all the losses.

#### Visualization of the classes distribution.

Table 3. Ablation study in terms of the impact of each loss term.

	$\mathcal{L}_{ ext{BCE}}$	$\mathcal{L}_{ ext{MKD}}$	$\mathcal{L}_{ ext{CLT}}$	AUC (%)
XresNet101(Teacher)				95.3
	✓			88.2
XresMVKT101	$\checkmark$	$\checkmark$		93.7
(Student)	$\checkmark$		$\checkmark$	93.9
	$\checkmark$	$\checkmark$	$\checkmark$	94.4

To visually assess the differences between the baseline and student, we use the T-SNE graphs to highlight the feature distribution. Figure 4 depicts the impact of MVKT between different backbones on ICBEB2018. As we can see, the distribution of features of normal (Norm), atrial premature beat (PAC), ST-segment depression (STD), and ST-segment elevation (STE) in the baseline network are confused to some extent. After MVKT, although the four categories are still relatively concentrated in the same area, the feature distribution of the distilled model in all nine categories became denser. As for the issue of category overlap, the status is greatly improved, especially on CNN\_Hannun, ResNet1d34, and XresNet1d101 backbones. (We can see the improvement of each disease on ResNet1d\_wang and Inception1d from Table 4). This suggests that the information conveyed by other leads can make up for the lack of single leads for various arrhythmias.

### Commonality.

Faced with the shortage of ECG datasets, many self-supervised learning methods are proposed to take advantage of unlabeled data. As for the student network's initialization, (Chen et al., 2021a) introduced a SOTA unsupervised pre-training program — CLECG to mine adequate information from unlabeled data. During the pre-training, CLECG encourages the representations of different augmented views of the same signal to be similar and increases the distance between representations of augmented views from the different signals.

We combine MVKT-ECG with CLECG. As PTB-XL dataset is the to-date largest freely accessible clinical ECG-waveform dataset, we firstly pre-train the network with CLECG on PTB-XL dataset, then using MVKT-ECG to finetune. The different proportions (eighth, quarter, half, all) of ICBEB2018 training set data were used in funtune procedure. AUC indicators of random initialization and self-supervised pre-training were compared and the results are shown in Figure 6.

It can be seen that at different data sizes, MVKT-ECG can improve performance by 1%-2% compared to the baseline model and even surpass the performance of the baseline

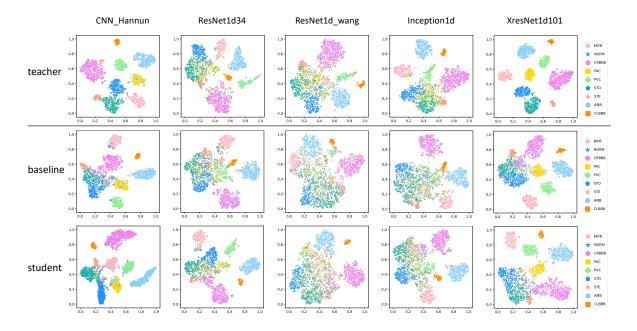


Figure 4. Feature visualization. The first line, second and third lines are the feature distribution of teacher, baseline, and student, respectively.

model trained with total data when using only half of the data. On this basis, using the parameters obtained from external data of CLECG can further improve the performance of the student model, and the AUC metric increases by 0.9% when using all training data, which shows that our method has practical application in combination with other methods. Moreover, this result suggests that the student model can improve performance by accepting knowledge transferred both from unlabeled external data and other leads.

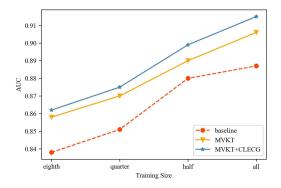


Figure 5. Combine MVKT-ECG with CLECG. The pre-training procedure is based on the PTB-XL dataset, and the finetune procedure is based on the ICBEB2018 dataset.

### 6. Conclusion

An effective ECG smart device recognition algorithm requires higher recognition accuracy and robustness for singlelead ECG signals. To achieve this goal, we propose MVKT-ECG. This is teacher-student information transferring approach in which the student observes only a single lead of the input ECG signals. This strategy encourages students to find better representations and to be closer to the teacher in performance through the knowledge transferring of the 12lead ECG processing network. Notably, MVKT-ECG shows robustness in different datasets and different backbones. Experimental results show that the proposed algorithm is insensitive to the classification granularity and specific categories of the datasets, and the accuracy of the student model in the detection of multiple arrhythmias is greatly improved. The visualization results also show that multi-view knowledge distillation can guide the student model to simulate the expression of the teacher model, which makes up for the deficiency of single-lead ECGs in differentiating some specific categories.

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### A. Detailed experimental results.

Here we report the F1\_score of each disease on different backbones. We also try to use different frameworks for teachers and students. We can see from the Table 4 and Table 5 that MVKT is beneficial in most cases for the diagnosis of any disease except for CNN\_Hannun and Resnet1d\_wang. This suggests that the MVKT operates in an environment with sufficient model capacity.

Table 4. MVKT's results of different diseases on ICBEB2018 dataset

	1 1. 1	MANUET	F1_score (%)									
	backbone	MVKT	NORM	AF	I-AVB	LBBB	RBBB	PAC	PVC	STD	STE	Average
Teacher	CNN_Hannun		75.8	92.0	87.0	88.0	91.7	71.3	86.7	74.0	68.1	81.6
Student	CNN_Hannun	$\checkmark$	65.2 <b>73.9</b>	95.2 <b>95.9</b>	84.8 <b>85.5</b>	91.3 87.5	85.0 <b>86.9</b>	66.7 <b>70.8</b>	86.5 82.2	66.0 <b>72.0</b>	30.8 <b>47.4</b>	74.6 <b>78.0</b>
Teacher	ResNet1d34		73.4	91.7	85.7	88.4	91.0	35.7	82.2	75.5	61.9	76.2
Student	ResNet1d34	$\checkmark$	63.9 <b>65.6</b>	92.1 <b>93.3</b>	84.5 <b>86.3</b>	87.5 <b>90.5</b>	85.5 <b>85.5</b>	33.8 <b>40.3</b>	60.0 <b>79.1</b>	65.1 <b>60.2</b>	26.9 <b>34.3</b>	66.6 <b>70.6</b>
Teacher	Resnet1d_wang		65.9	90.5	55.8	82.6	90.5	22.9	77.2	75.7	57.1	68.7
Student	Resnet1d_wang		60.2 59.4	82.7 <b>85.9</b>	67.7 <b>68.5</b>	84.0 <b>87.0</b>	84.5 84.0	26.3 24.7	36.1 <b>40.7</b>	60.0 <b>61.2</b>	30.4 26.3	59.1 <b>59.8</b>
Teacher	Inception1d		70.6	92.3	85.9	84.0	90.6	39.4	81.8	76.7	61.9	95.9
Student	Inception1d		62.4 <b>67.0</b>	89.9 <b>91.1</b>	81.1 <b>81.3</b>	85.1 <b>85.7</b>	84.07 <b>86.5</b>	27.9 <b>33.7</b>	30.9 <b>56.0</b>	60.0 <b>63.0</b>	26.9 <b>36.7</b>	60.9 <b>66.8</b>
Teacher	XresNet1d101		73.7	95.2	85.3	88.9	92.5	60.2	89.6	77.0	66.7	81.0
	XresNet1d101		65.8 <b>68.1</b>	89.8 <b>95.2</b>	83.8 <b>82.5</b>	87.0 <b>91.3</b>	85.2 <b>86.8</b>	63.1 <b>67.3</b>	80.0 <b>83.0</b>	65.9 <b>68.8</b>	32.3 <b>36.4</b>	72.5 <b>75.5</b>
Student	ResNet1d34	$\checkmark$	63.9 <b>69.1</b>	92.1 <b>93.3</b>	84.5 <b>86.3</b>	87.5 <b>87.5</b>	85.5 <b>87.3</b>	33.8 <b>50.0</b>	60.0 <b>74.5</b>	65.1 <b>66.3</b>	26.9 <b>30.8</b>	66.6 <b>71.7</b>

Table 5. MVKT's results of different diseases on PTB-XL dataset.

	Backbone	MVKT	F1_score						
	Баскоопе		NORM	MI	STTC	CD	HYP	Average	
Teacher	CNN_Ag		74.8	60.5	73.2	86.2	76.0	74.2	
Student	CNN_Ag	$\checkmark$	58.9 <b>60.1</b>	45.8 <b>46.4</b>	56.3 <b>57.4</b>	79.7 <b>80.3</b>	66.5 <b>67.8</b>	61.4 <b>62.4</b>	
Teacher	ResNet1d34		74.5	58.6	75.1	85.8	76.8	74.2	
Student	ResNet1d34		59.2 <b>61.3</b>	45.0 <b>47.3</b>	55.4 <b>56.7</b>	80.1 <b>80.6</b>	66.4 <b>68.7</b>	61.2 <b>62.9</b>	
Teacher	ResNet1d_wang		74.9	58.7	75.9	86.6	76.8	74.6	
Student	ResNet1d_wang		60.2 <b>60.4</b>	46.9 46.4	56.0 <b>57.7</b>	80.8 <b>80.9</b>	67.2 <b>68.5</b>	62.2 <b>62.5</b>	
Teacher	Inception1d		76.6	60.1	76.1	86.4	76.0	75.1	
Student	Inception1d	$\sqrt{}$	58.8 <b>60.2</b>	46.3 <b>47.8</b>	56.2 <b>57.4</b>	80.3 <b>80.6</b>	67.0 <b>66.5</b>	61.7 <b>62.5</b>	
Teacher	XresNet1d101		74.8	53.8	72.7	85.9	75.6	72.6	
Student	XresNet1d101	$\sqrt{}$	58.2 <b>59.8</b>	45.5 <b>47.2</b>	55.6 <b>57.3</b>	80.5 <b>80.4</b>	66.6 <b>67.5</b>	61.3 <b>62.4</b>	

## B. Illustration of Multi-label diseases Knowledge Distillation

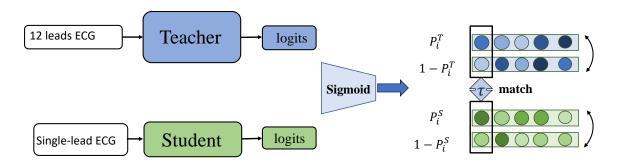


Figure 6. llustration of Multi-label diseases Knowledge Distillation.