

ProtoECGNet: Case-Based Interpretable Deep Learning for Multi-Label ECG Classification with Contrastive Learning



Sahil Sethi^{1,2}, David Chen², Thomas Statchen^{1,2}, Michael C. Burkhart², Nipun Bhandari³, Bashar Ramadan⁴, Brett Beaulieu-Jones²

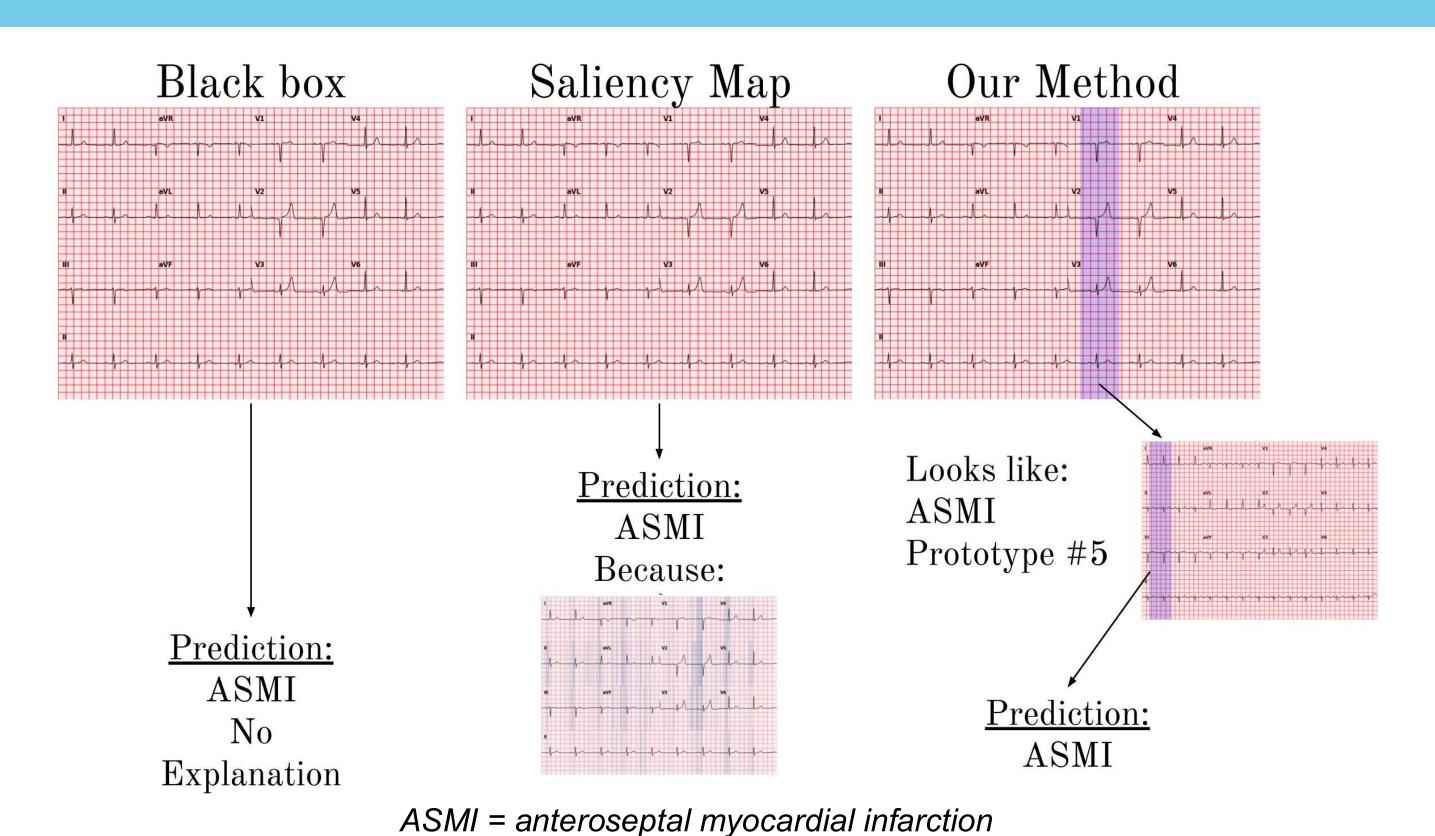


¹UChicago Pritzker School of Medicine, ²UChicago Center for Computational Medicine & Clinical AI, ³UC Davis Division of Cardiovascular Medicine, ⁴UChicago Medicine Section of Hospital Medicine

Motivation

- Transparency in model predictions is essential for clinical adoption
- Post-hoc explainability methods are often not faithful to a model's reasoning
- For ECG model explanations to be useful, they should be aligned with how clinicians reason, and cover the full diagnostic spectrum of ECG interpretation

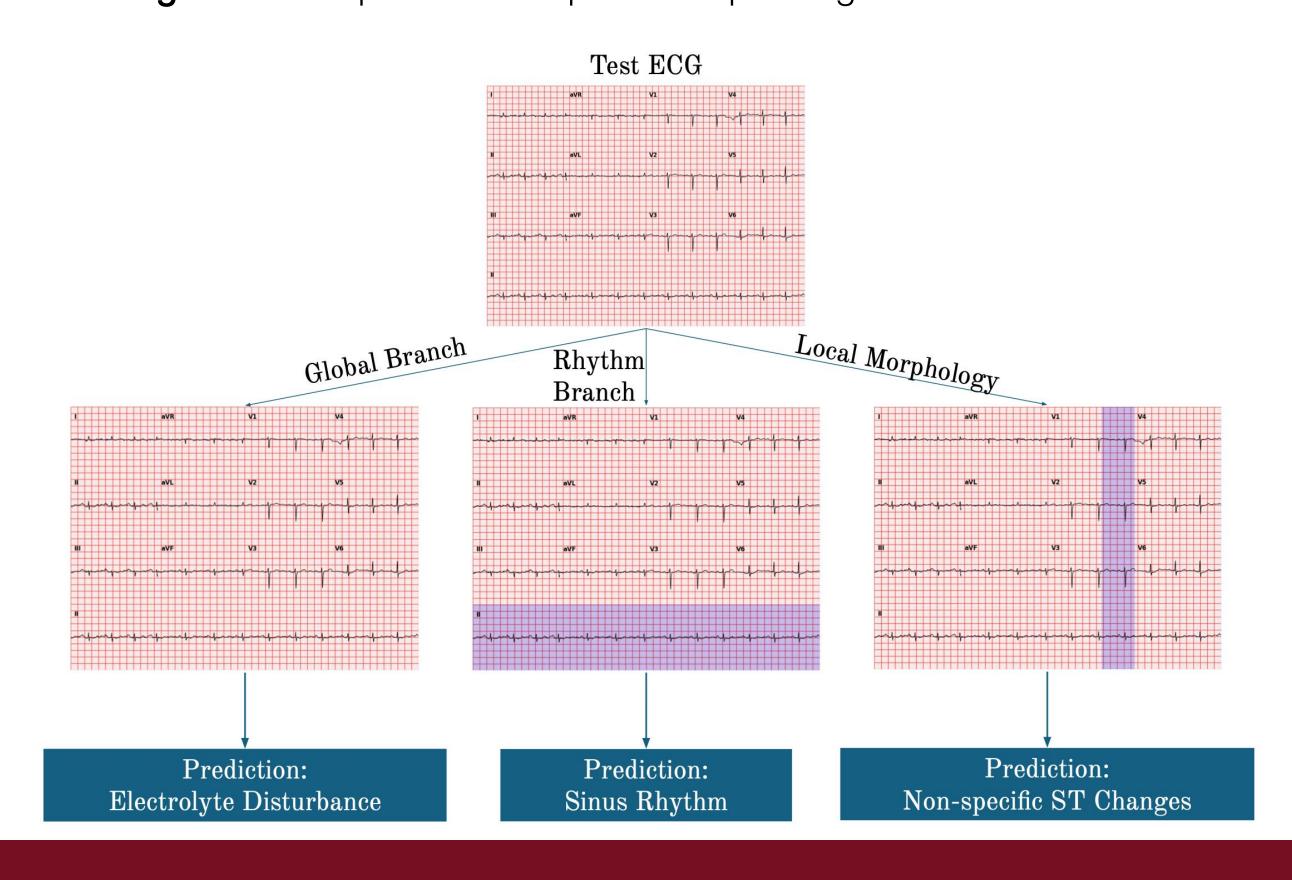
ProtoECGNet provides case-based explanations, tailored to the type of visual reasoning used by clinicians for each diagnosis, that are faithful to its internal reasoning process



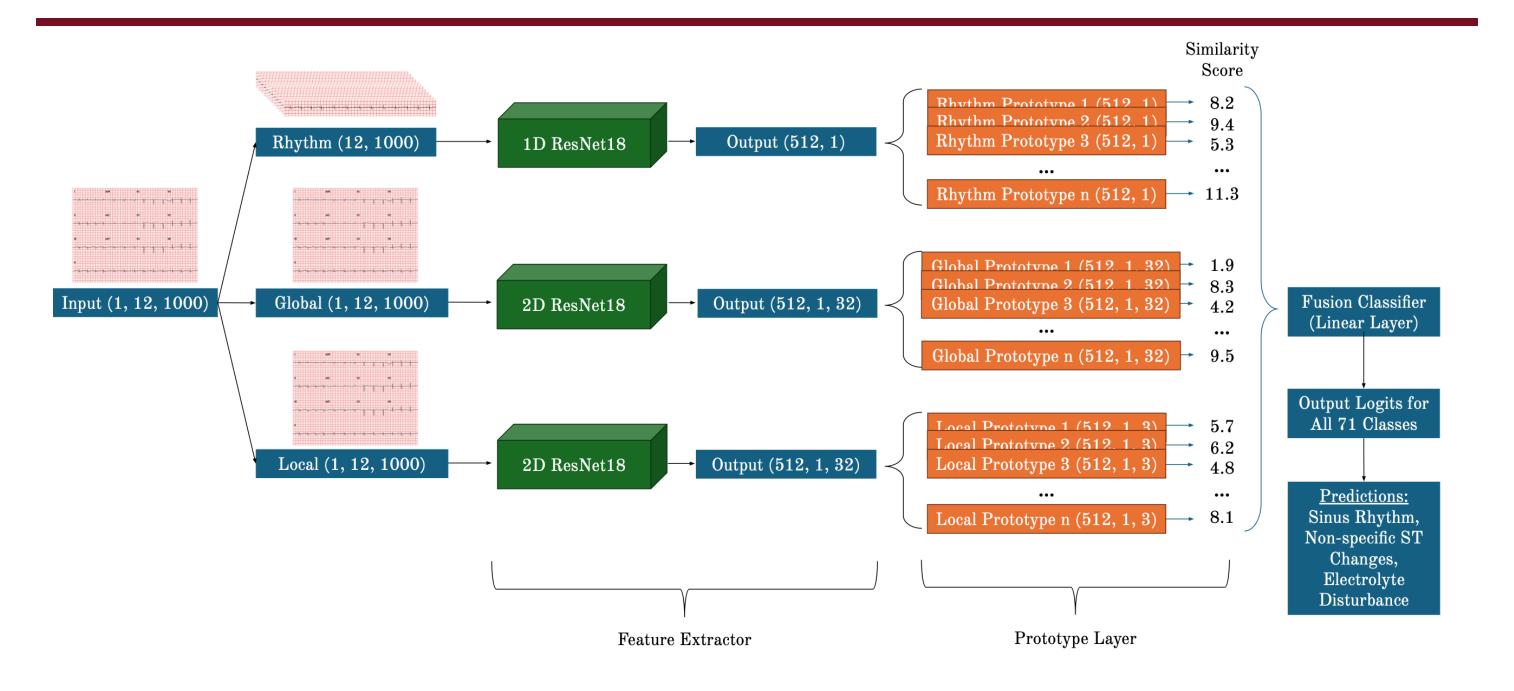
Multi-Branch Approach

We grouped the 71 labels from PTB-XL into three prototype categories based on the type of visual reasoning required for diagnosis:

- 1. Rhythm-based diagnoses—require temporal pattern analysis across full-length ECG signals, often discernible from a single lead
- 2. Morphology-based diagnoses—require localized waveform shape or inter-lead comparisons over short time intervals
- 3. Global diagnoses—require full-lead patterns spanning the full ECG duration



Internal Architecture



Contrastive Loss

$$\mathcal{L}_{\text{total}} = \mathcal{L}_{\text{BCE}} + \lambda_{\text{clst}} \cdot \mathcal{L}_{\text{clst}} + \lambda_{\text{sep}} \cdot \mathcal{L}_{\text{sep}} + \lambda_{\text{div}} \cdot \mathcal{L}_{\text{div}} + \lambda_{\text{cntrst}} \cdot \mathcal{L}_{\text{cntrst}}$$

$$\mathcal{L}_{\text{cntrst}} = -\frac{1}{\sqrt{P}} \left(\frac{\sum_{i,j} C_{ij} \cdot S(p_i, p_j)}{\sum_{i,j} C_{ij}} - \frac{\sum_{i,j} (1 - C_{ij}) \cdot S(p_i, p_j)}{\sum_{i,j} (1 - C_{ij})} \right)$$

Performance on PTB-XL (71 labels)

The final multi-branch, contrastive model outperforms non-contrastive and single-branch variants, and matches the best black box benchmark

Table 1: Macro-AUROC across branch-specific, single-branch, and multi-branch settings for ProtoECGNet.

| Setting | Model (Label Set) | Black-box | No Contrastive | w/ Contrastive |
|------------------------------|-------------------------------|-----------|----------------|----------------|
| Branch-Specific Labels | Rhythm Branch (16 labels) | 0.9403 | 0.8903 | 0.9064 |
| | Morphology Branch (52 labels) | 0.8872 | 0.8533 | 0.9051 |
| | Global Branch (3 labels) | 0.8649 | 0.8362 | 0.8667 |
| Full 71-Label, Single Branch | 1D Prototype Model | 0.9250 | 0.8646 | 0.8977 |
| | 2D Partial Prototype Model | N/A | 0.8873 | 0.9091 |
| | 2D Global Prototype Model | 0.8990 | 0.8681 | 0.9074 |
| Full 71-Label, Multi-Branch | Macro Aggregation | 0.8982 | 0.8609 | 0.9038 |
| | Fusion Classifier | N/A | 0.8855 | 0.9132 |

Table 2: Weighted AUROC with bootstrapped 95% confidence intervals across branch-specific, single-branch, and multi-branch settings.

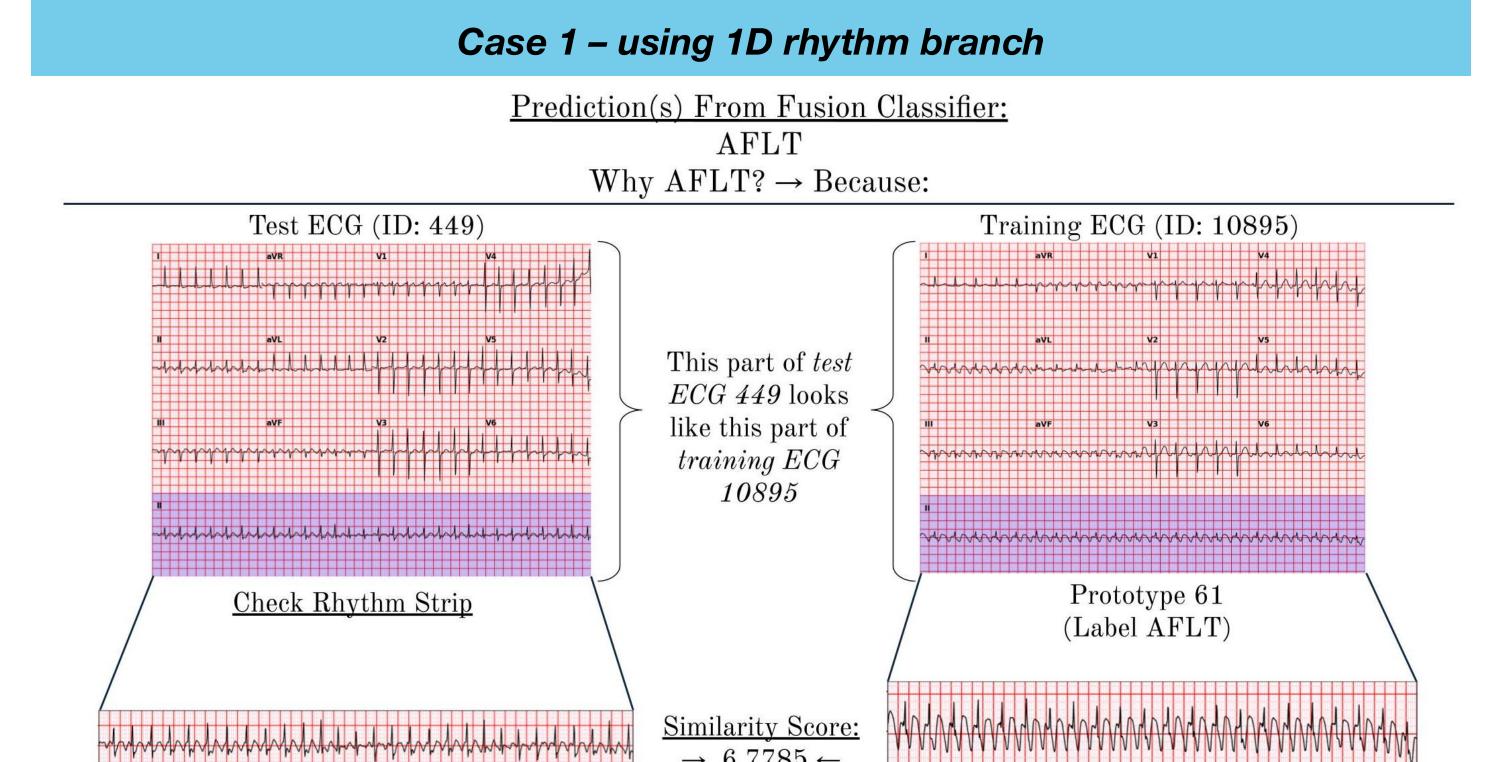
| Setting | Model (Label Set) | Black-box | No Contrastive | w/ Contrastive |
|-----------------|-------------------------------|-------------------------------|-------------------------------|-------------------------------|
| Branch-Specific | Rhythm Branch (16 labels) | 0.8919 (0.8757, 0.9071) | 0.8762 (0.8616, 0.8901) | 0.8853 (0.8708, 0.8991) |
| | Morphology Branch (52 labels) | $0.8996 \ (0.8930, \ 0.9057)$ | $0.8791 \ (0.8727, \ 0.8855)$ | $0.8996 \ (0.8931, \ 0.9059)$ |
| | Global Branch (3 labels) | $0.8307 \ (0.8137, \ 0.8470)$ | 0.6981 (0.6767, 0.7197) | $0.9039 \ (0.8906, \ 0.9164)$ |
| Single-Branch | 1D Prototype Model | 0.9081 (0.9012, 0.9147) | 0.8108 (0.8025, 0.8188) | 0.8857 (0.8782, 0.8930) |
| | 2D Partial Prototype Model | N/A | $0.8605 \ (0.8526, \ 0.8684)$ | $0.8743 \ (0.8666, \ 0.8819)$ |
| | 2D Global Prototype Model | $0.8932\ (0.8859,\ 0.9002)$ | $0.8589 \; (0.8505, 0.8669)$ | $0.8916 \; (0.8841, 0.8992)$ |
| Multi-Branch | Macro Aggregation | 0.8855 (0.8779, 0.8926) | 0.8486 (0.8413, 0.8557) | 0.8950 (0.8882, 0.9016) |
| | Fusion Classifier | N/A | $0.8800 \; (0.8728, 0.8872)$ | 0.9066 (0.9000, 0.9128 |

Clinician Evaluation of Prototypes

Table 3: Average prototype quality scores from structured clinician review (1–5 scale).

| Reviewer | Representativeness (95% CI) | Clarity (95% CI) |
|--------------|-----------------------------|-------------------------|
| Cardiologist | 4.29[4.22,4.35] | $4.48 \ [4.42, \ 4.54]$ |
| Internist | 3.59 [3.52, 3.66] | $4.73 \ [4.69, \ 4.77]$ |

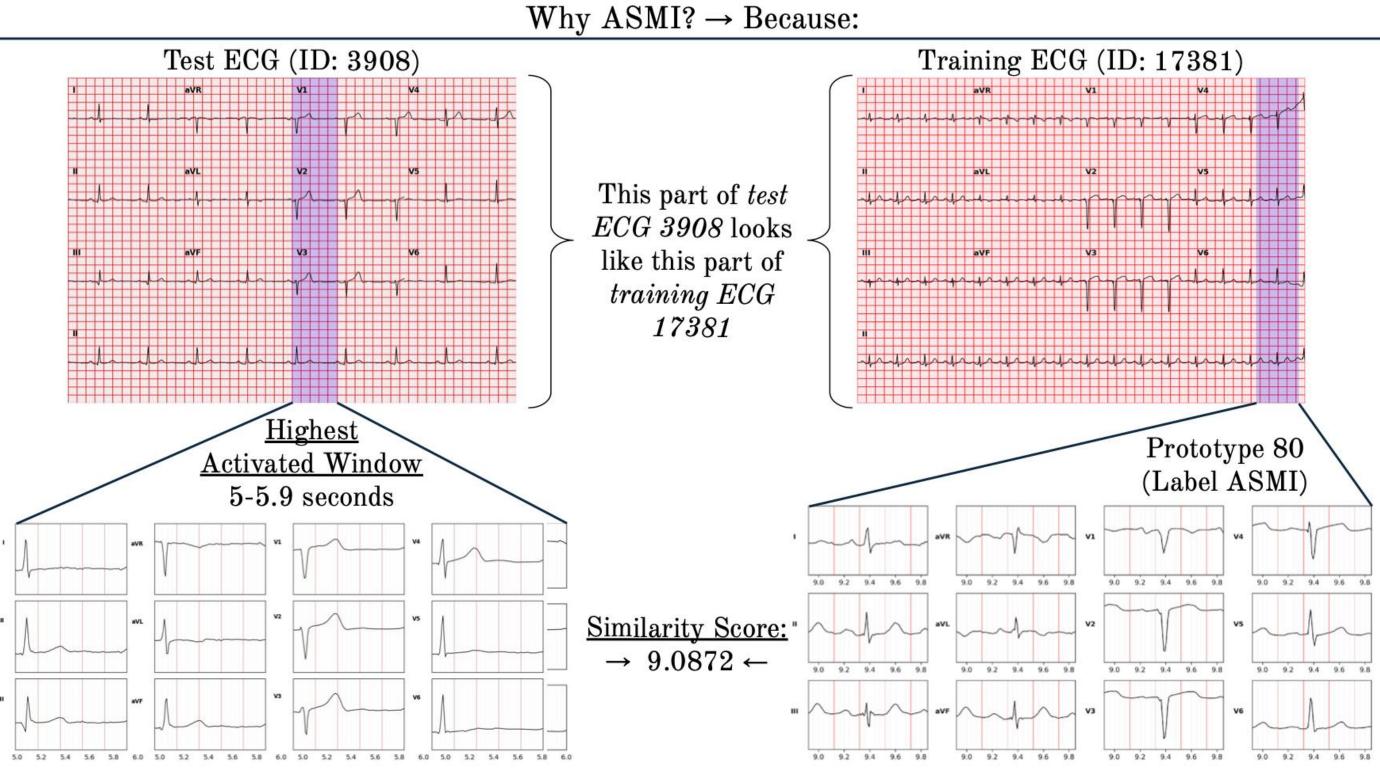
Case-Based Explanations



AFLT = atrial flutter

Case 2 – using 2D partial prototype branch

Prediction(s) From Fusion Classifier: ASMI, SR



ASMI = anteroseptal myocardial infarction, SR = sinus rhythm

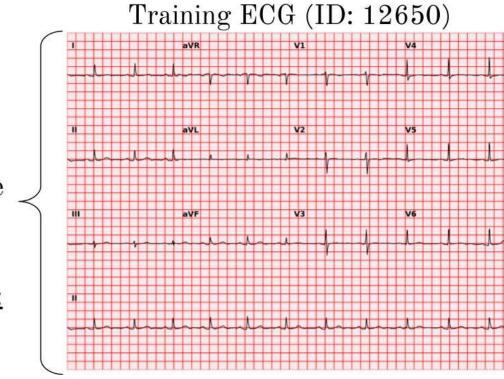
Case 3 – using 2D global prototype branch

Prediction(s) From Fusion Classifier: EL, STD_, SR Why EL? \rightarrow Because:

Test ECG (ID: 12126)

NATION OF THE PROPERTY O

This $test\ ECG$ $12126\ looks\ like$ this training $ECG\ 12650$ Similarity Score: $\rightarrow\ 2.1029 \leftarrow$



EL = electrolyte abnormality, STD_ = ST depression

Paper, Code, and Contact Info

