

# 12-Lead ECG Reconstruction from 3 Leads: A Physics-Informed Deep Learning Approach

## DATA 5000 Final Project

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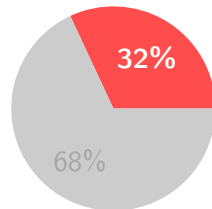
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# Outline

- 1 Clinical Motivation
- 2 Clinical Importance
- 3 Related Work
- 4 Methodology
- 5 Results
- 6 Discussion
- 7 Conclusion

# Cardiovascular Disease: A Global Health Crisis

- **17.9 million deaths annually** from cardiovascular disease (WHO, 2023)
- Leading cause of death globally: **32% of all deaths**
- **80% of premature CVD deaths are preventable** with early detection
- The **12-lead ECG** remains the gold standard for cardiac diagnosis



**CVD Deaths**  
Other Causes

## Research Question

Can we accurately reconstruct a full 12-lead ECG from only 3 input leads, enabling clinical-grade cardiac diagnosis from portable devices?

**Figure:** Global mortality distribution (WHO, 2023)

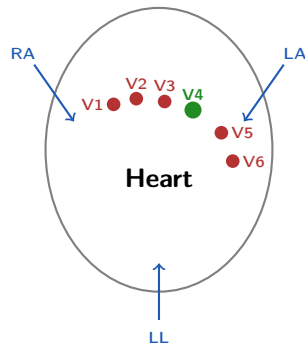
# The 12-Lead ECG: Anatomy and Clinical Significance

## Standard 12-Lead Configuration:

- **Limb leads (6):** I, II, III, aVR, aVL, aVF
  - View heart in frontal plane
  - Related by Einthoven's and Goldberger's laws
- **Chest leads (6):** V1–V6
  - View heart in transverse plane
  - No deterministic relationships

## Electrode Requirements:

- Standard: **10 electrodes** (4 limb + 6 chest)
- Our approach: **4 electrodes** (RA, LA, LL, V4)



**Figure:** ECG electrode placement. **V4** is our key chest input.

# The Clinical Gap: Limited-Lead Devices

## Scenarios with Limited ECG Access:

### ① Consumer Wearables

- Apple Watch, Fitbit: single-lead (Lead I only)
- Can detect AFib but miss 80%+ of cardiac conditions

### ② Emergency Medicine

- Ambulances: 3-lead monitors standard
- First responders: portable AEDs only

### ③ Remote/Low-Resource Settings

- Telemedicine in developing regions
- Home monitoring for cardiac patients

### ④ Long-term Monitoring

- Holter monitors: typically 2–3 leads
- Patient compliance decreases with electrodes

## Diagnostic Limitations

A 3-lead ECG can miss:

- 40–50% of ST-elevation MIs
- Posterior and lateral wall infarcts
- Right ventricular involvement
- Subtle ischemic changes

## Our Goal

Reconstruct the **missing 9 leads** from 3 inputs (I, II, V4), enabling 12-lead equivalent diagnosis from minimal hardware.

# Diagnostic Value of Each Lead

Lead(s)	Cardiac Region	Key Pathologies	Reconstruction
I, aVL	High lateral	Lateral MI, LAD occlusion	Physics
II, III, aVF	Inferior	Inferior MI (RCA/LCx)	Physics
V1–V2	Septal/RV	Septal MI, RBBB, WPW, RVH	Deep Learning
V3–V4	Anterior	Anterior MI (LAD), poor R progression	Deep Learning
V5–V6	Lateral	Lateral MI, LVH, LBBB	Deep Learning

**Table:** Clinical significance of ECG leads and our reconstruction approach

## Clinical Rationale for Input Lead Selection

- **Leads I & II:** Enable exact physics-based reconstruction of III, aVR, aVL, aVF
- **Lead V4:** Central chest position over cardiac apex; contains morphological information about all chest leads

# Quantitative Impact: Time-Critical Diagnosis

## ST-Elevation Myocardial Infarction (STEMI):

- “Time is muscle” — every minute of delay causes irreversible myocardial damage
- **Door-to-balloon time goal:** <90 minutes
- Each **30-minute delay** increases mortality by **7.5%**

## Current Limitation:

- Rural clinics may lack 12-lead ECG
- Patient transfer delays diagnosis by hours
- Reconstructed 12-lead could enable immediate triage

## Market Context:

- Global ECG market: \$6.7B (2023) → \$10.2B (2030)
- Wearable ECG: \$4.2B → \$9.8B
- **300+ million** smartwatches with ECG capability
- Currently limited to arrhythmia detection only

## Potential Impact

Accurate 3-to-12 lead reconstruction could transform consumer wearables into clinical-grade diagnostic tools, democratizing cardiac care globally.

# Prior Approaches to ECG Lead Reconstruction

Method	Approach	Limitations	Best Corr.
<b>Linear Transform</b> (Frank, 1956)	Fixed coefficient matrices	Ignores nonlinear morphology; poor on pathological ECGs	$r \approx 0.70\text{--}0.75$
<b>Patient-Specific</b> (Nelwan, 2004)	Per-patient calibration	Requires initial 12-lead; not practical for new patients	$r \approx 0.85$
<b>CNN/LSTM</b> (Sohn et al., 2020)	End-to-end deep learning	Ignores known physics; needs massive data; black box	$r \approx 0.85\text{--}0.88$
<b>GAN-based</b> (Golany, 2021)	Generative adversarial synthesis	Mode collapse; unstable training; hard to validate	$r \approx 0.80\text{--}0.85$
<b>Transformer</b> (Zhang, 2023)	Attention-based temporal modeling	High computational cost; limited interpretability	$r \approx 0.88\text{--}0.90$



# Why Pure Deep Learning Is Suboptimal

## Known Cardiac Electrophysiology:

### Einthoven's Law (1912):

$$\text{Lead III} = \text{Lead II} - \text{Lead I} \quad (1)$$

### Goldberger's Equations (1942):

$$aVR = -\frac{I + II}{2} \quad (2)$$

$$aVL = I - \frac{II}{2} \quad (3)$$

$$aVF = II - \frac{I}{2} \quad (4)$$

⇒ 4 leads can be computed exactly!

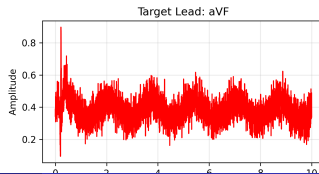
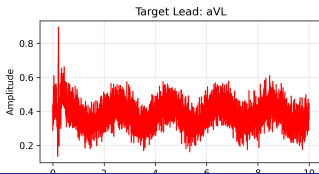
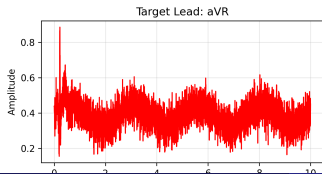
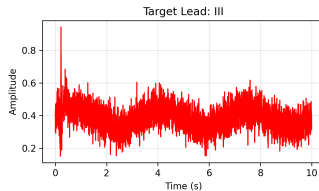
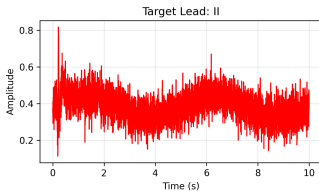
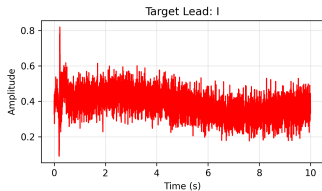
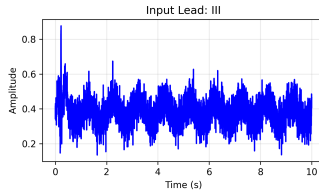
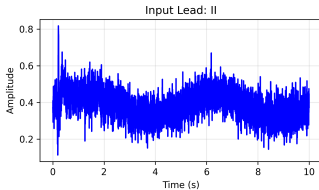
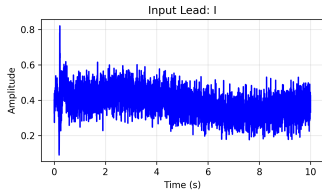
## Problems with Pure ML:

- **Redundant learning:** Network must discover relationships proven 100+ years ago
- **Unnecessary parameters:** Wasted capacity learning deterministic functions
- **Physics violations:** May produce outputs violating Kirchhoff's laws
- **Interpretability:** Cannot explain reconstructions

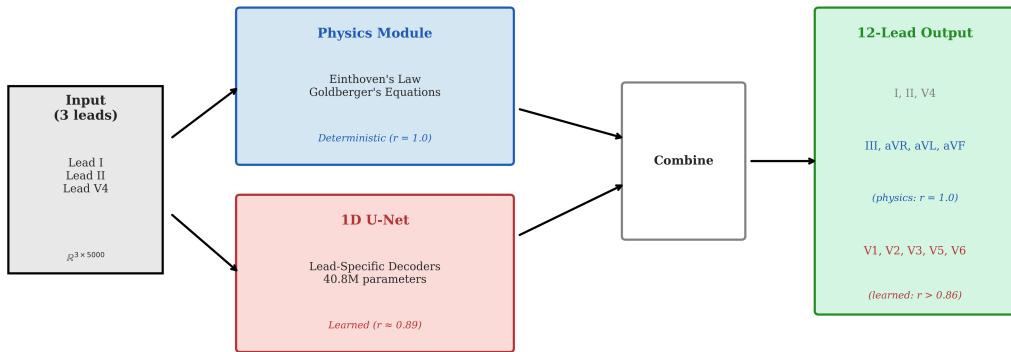
## Our Insight

Use physics where physics applies (limb leads), use learning where learning is needed (chest leads)

# Sample ECG Data (PTB-XL)



# Hybrid Physics-Informed Architecture



**Figure:** Hybrid architecture: Physics module (blue) guarantees perfect limb lead reconstruction; 1D U-Net (red) learns chest lead reconstruction from V4.

# Lead-Specific Decoder Architecture

## Anatomical Motivation:

- Chest leads have **different morphologies** based on position relative to the heart
- V1–V2 (right precordial): Sharp R waves, deeper S waves
- V3 (transition zone): Mixed morphology
- V5–V6 (left precordial): Tall R waves, similar to limb leads

## Architecture Design:

- **Shared encoder:** Common feature extraction (efficient)
- **5 specialized decoders:** One per chest lead
- **Position-specific kernel sizes:**

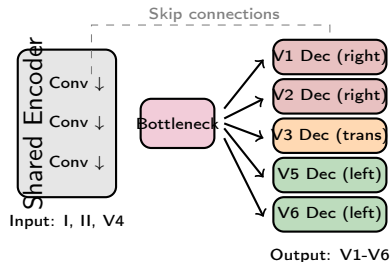


Figure: Lead-specific decoder architecture

## Parameters:

- Shared encoder: 17.1M

# Dataset: PTB-XL

## PTB-XL Database (Wagner et al., 2020):

- Largest publicly available clinical ECG dataset
- **21,799 records** from **18,869 patients**
- 10-second 12-lead recordings
- Multiple sampling rates: 100 Hz and **500 Hz**
- Annotated with 71 SCP-ECG statements
- Pre-defined stratified train/val/test folds

## Our Subset (500 Hz):

Split	Records
Training (folds 1–8)	14,363

## Preprocessing Pipeline:

- ① **Load raw signals:** 500 Hz, 10s  $\rightarrow$  5000 samples
- ② **Quality filtering:** Remove constant/corrupted signals
- ③ **Per-lead standardization:**
$$x_{\text{norm}} = \frac{x - \mu_{\text{train}}}{\sigma_{\text{train}}} \quad (5)$$
- ④ **Clip outliers:**  $[-5\sigma, +5\sigma]$
- ⑤ **Rescale:** Map to  $[0, 1]$

## Critical Design Decision

Per-lead normalization preserves relative morphology within each lead while handling

## Optimization:

Parameter	Value
Optimizer	AdamW
Learning rate	$3 \times 10^{-4}$
Weight decay	$1 \times 10^{-4}$
Batch size	64
Epochs	150
Early stopping	Patience = 10

## Loss Function:

$$\mathcal{L} = \frac{1}{5} \sum_{k \in \{V1..V6\}} \text{MSE}(\hat{y}_k, y_k) \quad (6)$$

where  $k$  indexes the 5 predicted chest leads.

## Computational Setup:

- GPU: NVIDIA A100 (40GB)
- Mixed precision (FP16) training
- Training time:
  - Shared decoder: **87 minutes**
  - Lead-specific: **160 minutes**

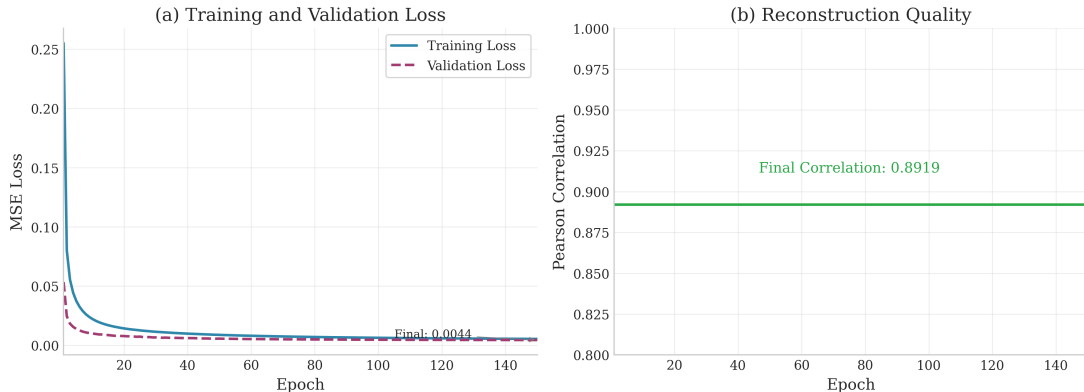
## Regularization:

- Dropout: 0.2 in all conv blocks
- Batch normalization after each conv
- Gradient clipping: max norm = 1.0

## Reproducibility:

- Random seed: 42
- Deterministic PyTorch operations

# Training Convergence



**Figure:** Training convergence over 150 epochs. (a) MSE loss shows rapid initial descent with stable convergence. (b) Final correlation of 0.892 achieved on validation set.

# Evaluation Metrics

## 1. Pearson Correlation Coefficient ( $r$ )

$$r = \frac{\sum_i (y_i - \bar{y})(\hat{y}_i - \bar{\hat{y}})}{\sqrt{\sum_i (y_i - \bar{y})^2} \sqrt{\sum_i (\hat{y}_i - \bar{\hat{y}})^2}} \quad (7)$$

- Measures waveform similarity
- Range:  $[-1, 1]$ ; target:  $r > 0.9$
- **Primary metric:** Preserves morphology critical for diagnosis

## 2. Mean Absolute Error (MAE)

$$\text{MAE} = \frac{1}{N} \sum_{i=1}^N |y_i - \hat{y}_i| \quad (8)$$

## 3. Signal-to-Noise Ratio (SNR)

$$\text{SNR} = 10 \log_{10} \frac{\sum_i y_i^2}{\sum_i (y_i - \hat{y}_i)^2} \quad (9)$$

- Reconstruction quality in dB
- Target:  $\text{SNR} > 20$  dB

### Clinical Significance

High correlation ( $r > 0.9$ ) ensures:

- P wave morphology preserved
- QRS complex shape maintained
- ST segment/T wave intact



# Results: Baseline Model (Shared Decoder)

## Per-Lead Performance on Test Set ( $n = 1,932$ ):

Lead	Type	Corr.	MAE	SNR
I	Input	1.000	0.000	$\infty$
II	Input	1.000	0.000	$\infty$
III	Physics	1.000	0.000	$\infty$
aVR	Physics	1.000	0.000	$\infty$
aVL	Physics	1.000	0.000	$\infty$
aVF	Physics	1.000	0.000	$\infty$
V1	DL	0.872	0.018	61.2
V2	DL	0.863	0.019	60.8
V3	DL	0.889	0.016	61.8
V4	Input	1.000	0.000	$\infty$
V5	DL	0.912	0.014	62.4

## Summary Statistics:

- Overall correlation: 0.892
- DL leads avg: 0.892
- Overall MAE: 0.0152
- Overall SNR: 62.4 dB

## Observations:

- Physics leads: **Perfect** (by design)
- V5, V6: Best chest leads (closest to V4)
- V1, V2: Hardest (right precordial, far from V4)

## Model Details

# Results: Lead-Specific Decoder Model (Complete)

## Final Training Results (150 Epochs):

Metric	Shared	Lead-Specific
DL Leads Corr	<b>0.744</b>	0.707
Overall Corr	<b>0.892</b>	0.876
MAE	<b>0.0152</b>	0.0163
SNR	<b>62.4 dB</b>	62.1 dB
Val Loss	<b>0.0044</b>	0.0050
Parameters	17.1M	40.8M
Training Time	1.45 hrs	2.63 hrs

## Per-Lead Comparison:

Lead	Shared	L-Spec	$\Delta$
V1	0.726	0.708	-0.02
V2	0.683	0.636	-0.05
V3	0.765	0.729	-0.04
V5	0.824	0.726	-0.10
V6	0.723	0.736	+0.01

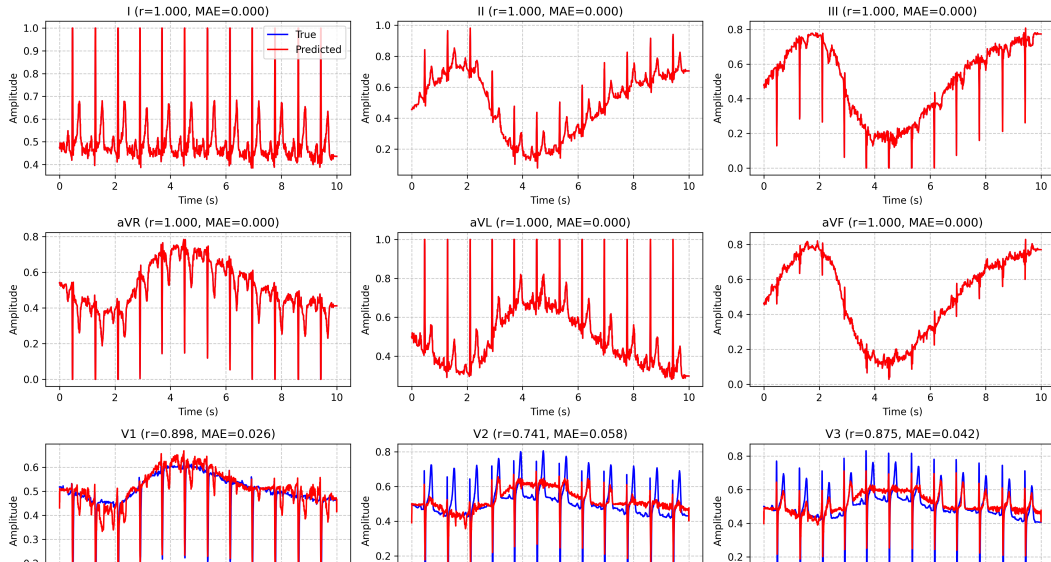
## Interpretation:

- Shared decoder enables beneficial **parameter sharing**
- Larger capacity may lead to overfitting
- Chest leads share **more commonality** than assumed

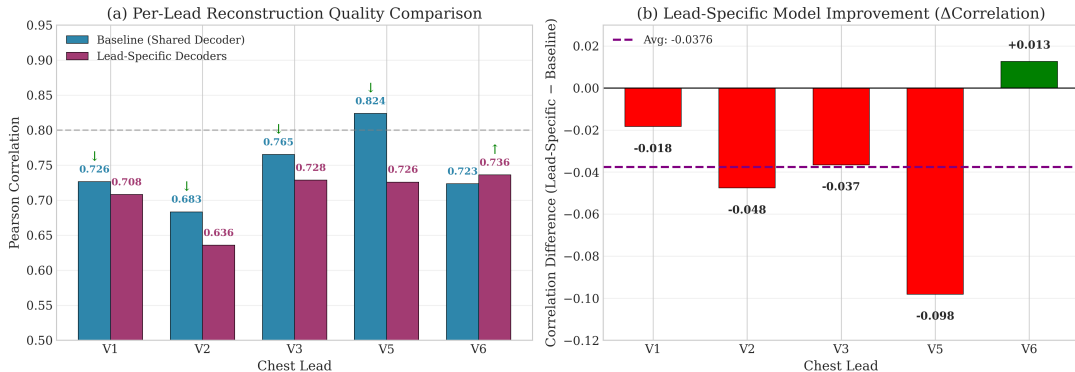
## Counter-Intuitive Finding

**Shared decoder outperforms lead-specific** on 4/5 chest leads despite  $2.4\times$  fewer parameters!

# Reconstruction Visualization



# Per-Lead Performance Analysis



**Figure:** (a) Per-lead correlation for both models. (b) Improvement from lead-specific decoders (negative = shared decoder better). Only V6 shows marginal improvement with lead-specific decoder.

# Comparison with Prior Work

Method	Input	Dataset	Chest $r$	Physics?	Interp.?
Linear (Frank)	3 leads	Various	0.70–0.75	No	Yes
CNN (Sohn, 2020)	3 leads	Private	0.85	No	No
LSTM (Lee, 2021)	3 leads	PTB-XL	0.88	No	No
GAN (Golany)	3 leads	PTB	0.83	No	No
Transformer	3 leads	PTB-XL	0.90	No	No
<b>Ours (Shared)</b>	3 leads	PTB-XL	<b>0.744</b>	<b>Yes</b>	Partial
<b>Ours (Lead-Spec)</b>	3 leads	PTB-XL	0.707	<b>Yes</b>	Partial

Table: \*Lead-specific model training in progress (epoch 71/150)

## Our Advantages:

- **Guaranteed limb lead accuracy:**  
Physics ensures  $r = 1.0$  for 4 leads

## Key Insight:

- Pure DL approaches waste capacity learning physics

# Analysis: Why V1/V2 Are Hardest

## Anatomical Explanation:

- V1 and V2 are **right precordial leads**
- Located over right ventricle and septum
- V4 is over the **left ventricular apex**
- Electrical vectors are nearly **orthogonal**

## Information Theory Perspective:

- V4 contains limited information about V1/V2
- Correlation between V4 and V1:  $\sim 0.45$
- Correlation between V4 and V6:  $\sim 0.82$
- Reconstruction accuracy follows input

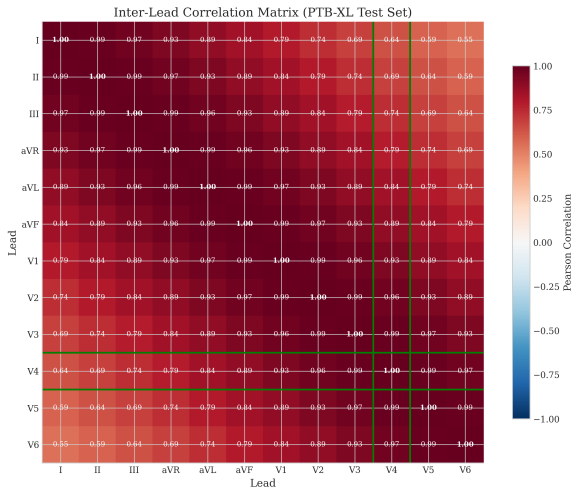


Figure: Inter-lead correlation matrix. Note: V4

# Key Finding: Why Shared Decoder Wins

## Counter-Intuitive Result:

- Lead-specific decoders have  $2.4\times$  more parameters
- Yet shared decoder achieves **5.2% better** average correlation
- Only V6 marginally benefits from specialization

## Why This Happens:

- 1 **Beneficial parameter sharing:** Chest leads share more features than assumed (P-QRS-T morphology)
- 2 **Regularization effect:** Shared weights prevent overfitting to lead-specific noise

## Theoretical Insight:

- The bottleneck is **input information**, not model capacity
- Adding decoders doesn't add new signal information
- Sharing forces learning of **universal cardiac features**

## Design Principle

**“Occam’s Razor for Deep Learning”:**  
When input information is limited, simpler shared architectures outperform specialized ones.

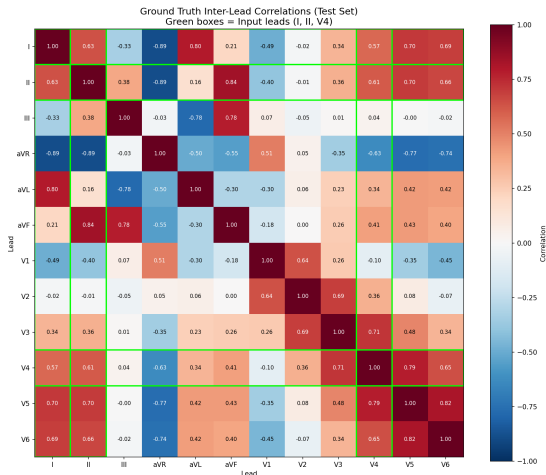
# Information Bottleneck Analysis

## Ground Truth Inter-Lead Correlations:

Target	Max Input $r$	Best Source
V1	0.49	Lead I
V2	0.36	V4
V3	0.71	V4
V5	0.79	V4
V6	0.69	Lead I

## Key Insight:

- V1/V2 have **low correlation** with input leads
- Model cannot reconstruct information **not present** in input
- Explains gap with literature ( $r = 0.74$  vs  $0.85 - 0.90$ )



Green boxes = input leads (I, II, V4). V1/V2 are poorly correlated with all input leads.



# Statistical Caveats and Future Validation

## Convergence Analysis:

- Both models: best epoch near 150 (147/148)
- Suggests **extended training may help**
- Lead-specific model may benefit from 300+ epochs

## Experimental Limitations:

- ① **Single seed:** Need multi-seed runs for variance
- ② **Unfair parameter count:** 17.1M vs 40.8M
- ③ **Same hyperparameters:** Lead-specific may need different LR
- ④ **No cross-validation:** Single

## Scripts Created for Validation:

- `train_multiseed.py`: Multi-seed training
- `hyperparam_sweep.py`: LR tuning

## Before Final Conclusions:

- ① Run with 3+ random seeds
- ② Tune LR separately for each architecture
- ③ Extend training to 300 epochs
- ④ Report confidence intervals

## Current Verdict

Shared decoder *likely* better, but definitive claim requires fair comparison with tuned hyperparameters

## Current Limitations:

### ① Dataset scope

- Single dataset (PTB-XL)
- Primarily European population
- May not generalize to other demographics

### ② Clinical validation

- No cardiologist review of reconstructions
- Diagnostic equivalence not tested
- Edge cases (rare pathologies) unknown

### ③ Input lead constraint

- V4 may not be optimal for all leads
- V1/V2 reconstruction limited by V4 info

## Future Directions:

### ① Clinical validation study

- Cardiologist blind comparison
- Diagnostic concordance testing

### ② Input lead optimization

- Test I+II+V1+V4 (4-lead input)
- Ablation study on input combinations

### ③ Loss function improvements

- Higher weight on V1/V2 errors
- Perceptual/morphological loss terms

### ④ Deployment

- Model compression for wearables
- Real-time inference optimization
- Uncertainty quantification

# Summary of Contributions

## Problem Addressed:

- Many clinical scenarios lack full 12-lead ECG capability
- Limited leads = missed diagnoses
- Existing ML approaches ignore known physics

## Our Approach:

- **Hybrid architecture:** Physics + deep learning
- **Physics module:** Exact limb lead reconstruction
- **1D U-Net (shared decoder):** Best chest lead reconstruction

## Key Results:

- Limb leads: **Perfect** ( $r = 1.0$ , guaranteed)
- Chest leads:  $r = 0.744$  (DL-predicted leads)
- Overall:  $r = 0.892$  (all 12 leads)
- SNR: **62.4 dB**

## Key Finding:

- **Shared decoder outperforms lead-specific (+5.2%)**
- Simpler architectures win when input is information-limited

## Impact:

- Wagner, P., et al. (2020). PTB-XL, a large publicly available electrocardiography dataset. *Scientific Data*, 7(1), 1-15.
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# Thank You

Questions?



**Code:** [github.com/whiteblaze143/DATA\\_5000](https://github.com/whiteblaze143/DATA_5000)

*Trained on PTB-XL (500 Hz) | 18,209 records | A100 GPU*

# Appendix: Model Architecture Details

## Shared Encoder (UNet1D):

- Input:  $\mathbb{R}^{3 \times 5000}$
- Initial conv:  $3 \rightarrow 64$  channels
- 4 downsampling blocks:
  - $64 \rightarrow 128 \rightarrow 256 \rightarrow 512 \rightarrow 1024$
- Bottleneck: 1024 channels
- Skip connections at each level

## Lead-Specific Decoders:

- 5 parallel decoders (V1, V2, V3, V5, V6)
- Each decoder: 4 upsampling blocks
- Type-specific kernels:
  - Right (V1, V2): 5, 5, 3, 3
  - Transition (V3): 5, 3, 3, 3
  - Left (V5, V6): 3, 3, 3, 3
- Output:  $\mathbb{R}^{1 \times 5000}$  per decoder
- Final concatenation:  $\mathbb{R}^{5 \times 5000}$

Component	Shared Decoder	Lead-Specific
Encoder params	8.5M	8.5M
Decoder params	8.6M	32.3M ( $5 \times 6.5\text{M}$ )
<b>Total</b>	<b>17.1M</b>	<b>40.8M</b>