Heart Disease Detection Using ECG Signals

DATASET UESD

- 1. mit-bih-arrhythmia-database-1.0.0
- 2. mit-bih-artial-fibrillation-database-1.0.0
- 3. mit-bih-malignant-ventricular-ectopy-database-1.0.0
- 4. mit-bih-normal-sinus-rhythm-database-1.0.0
- 5. mit-bih-st-change-database-1.0.0

MIT-BIH-ARRHYTHMIA-DATABASE-1.0.0

- Disease Focus: Cardiac Arrhythmias
- Description: This dataset contains ECG signals from patients with various types of arrhythmias — abnormal heart rhythms that may be too fast, too slow, or irregular. Some common arrhythmias detectable in this dataset include:
 - 1. Premature Ventricular Contractions (PVCs)
 - 2. Bundle Branch Block
 - 3. Atrial Premature Beat
 - 4. Supraventricular Tachycardi
 - 5. Ventricular Tachycardia (VT)

MIT-BIH-ARTIAL-FIBRILLATION-DATABASE-1.0.0

- Disease Focus: Atrial Fibrillation (AF)
- Description: This dataset is specifically curated to help detect Atrial Fibrillation, a common type of arrhythmia where the atria (upper heart chambers) beat chaotically and out of sync with the ventricles. It can lead to:
 - 1. Stroke
 - 2. Heart Failure
 - 3. Fatigue and Palpitations

MIT-BIH-MALIGNANT-VENTRICULAR-ECTOPY-DATABASE-1.0.0

- Disease Focus: Ventricular Arrhythmias (Life-threatening)
- Description: This dataset includes ECG signals from patients who had severe ventricular arrhythmias such as:
 - 1. Ventricular Tachycardia (VT)
 - 2. Ventricular Fibrillation (VF)
 - 3. Multiform Premature Ventricular Complexes (PVCs)
 - 4. Causing sudden Cardiac death

MIT-BIH-NORMAL-SINUS-RHYTHM-DATABASE-1.0.0

- This dataset contains ECG recordings from healthy individuals with normal sinus rhythm. It serves as a baseline to distinguish between:
 - 1. Normal cardiac activity
 - 2. Abnormal or diseased conditions
- Models trained on both normal and abnormal data can learn to detect deviations from healthy heart function.

MIT-BIH-ST-CHANGE-DATABASE-1.0.0

- Disease Focuses: Myocardial Ischemia (ST Elevation or Depression)
- Description: ST-segment deviations are signs of reduced blood flow to the heart (ischemia). This can indicate:
 - 1. Acute coronary syndrome
 - 2. Heart attack (Myocardial Infarction)

Models Used

- 1. 2 Custom CNN Models
- 2. VGG16
- 3. ResNet50
- 4. ResNet101
- 5. Xception
- 6. DenseNet121

First Custom CNN model

CNN Model Architecture (Simplified)

1. Input Layer:

1. Shape: (128, 128, 3)

2. Conv Block 1:

- 1. Conv2D (32 filters, 3×3, ReLU)
- 2. Batch Normalization
- 3. MaxPooling (2×2)
- 4. Dropout (rate = 0.3)

3. Conv Block 2:

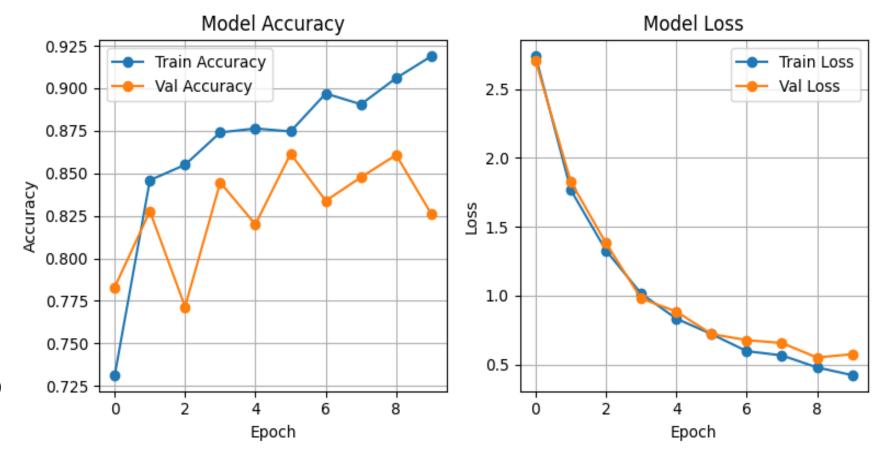
- 1. Conv2D (64 filters, 3×3, ReLU)
- MaxPooling (2×2)

4. Fully Connected Layers:

- 1. Flatten
- 2. Dense (128 units, ReLU, L2 Regularization)
- 3. Dense (32 units, ReLU)

5. Output Layer:

1. Dense (5 units, Softmax)



Validation Set Accuracy: 82.62%

Second Custom CNN model

CNN Model Architecture (Simplified)

1. Input Layer:

1. Shape: (128, 128, 3)

2. Conv Block 1:

- 1. Conv2D (32 filters, 3×3, ReLU)
- 2. Batch Normalization
- 3. MaxPooling (2×2)
- 4. Dropout (rate = 0.3)

1. Data Augmentation:

- 1. Random Flip
- 2. Random Rotation (±10%)
- 3. Random Zoom (±10%)

2. Conv Block 2:

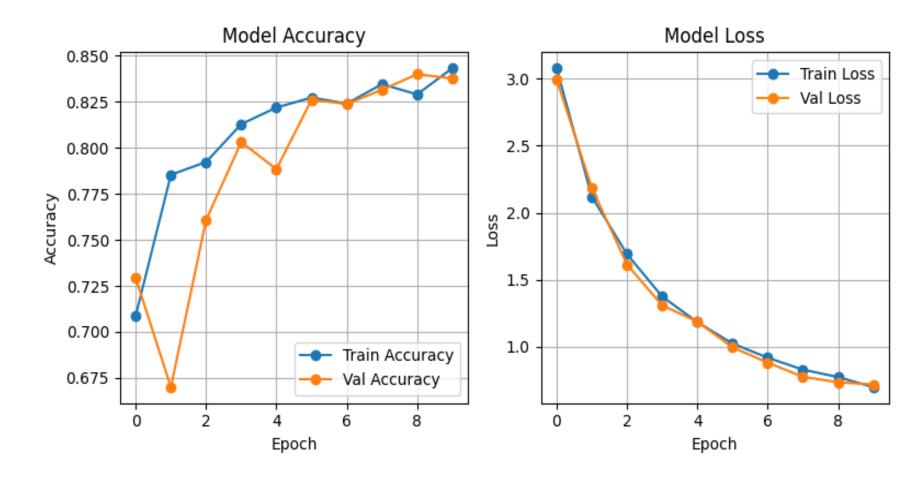
- 1. Conv2D (64 filters, 3×3, ReLU)
- 2. MaxPooling (2×2)

3. Fully Connected Layers:

- Flatten
- 2. Dense (128 units, ReLU, L2 Regularization)
- 3. Dense (32 units, ReLU, L2 Regularization)

4. Output Layer:

1. Dense (5 units, Softmax)



Validation Set Accuracy: 83.77%

VGG16 (transfer learning)

Transfer Learning Model Architecture

1. Base Model:

1. Pretrained VGG16

2. Weights: ImageNet

3. Top layers removed (include_top=False)

4. Input shape: (128, 128, 3)

5. All layers frozen

2. Custom Head:

Flatten

2. Dense (64 units, ReLU)

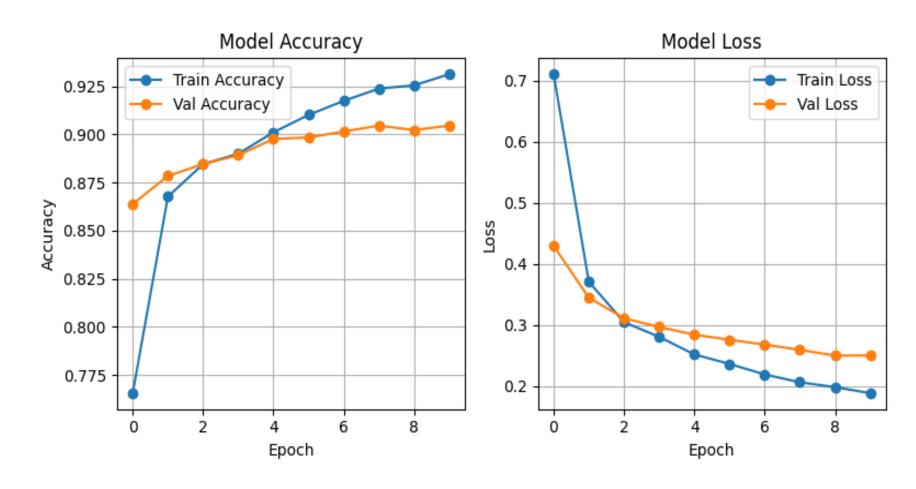
3. Output: Dense (5 units, Softmax)

3. Compilation:

1. Optimizer: Adam (learning rate = 1e-4)

2. Loss: Categorical Crossentropy

3. Metric: Accuracy



Validation Set Accuracy: 90.46%

ResNet50 (transfer learning)

Transfer Learning Model Architecture

1. Base Model:

Pretrained ResNet50

2. Weights: ImageNet

3. Top layers removed (include_top=False)

4. Input shape: (128, 128, 3)

5. All layers frozen

2. Custom Head:

1. Flatten

2. Dense (64 units, ReLU)

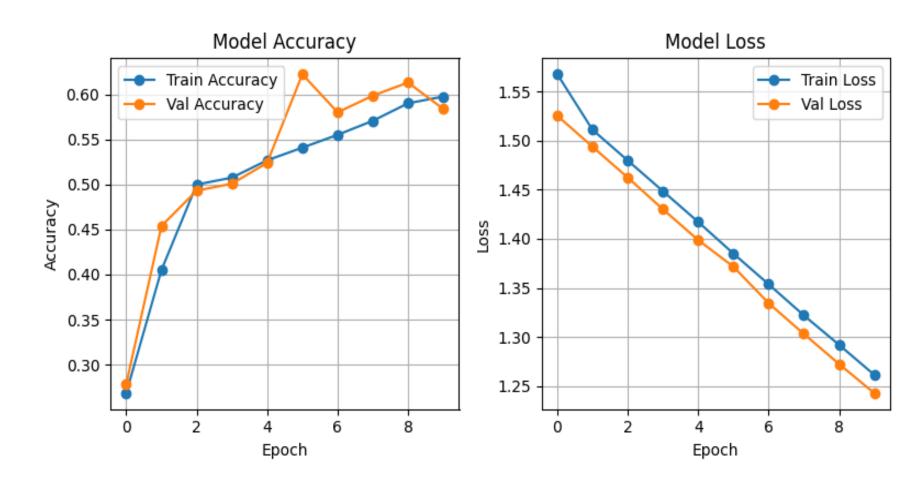
3. Output: Dense (5 units, Softmax)

3. Compilation:

1. Optimizer: Adam (learning rate = 1e-4)

2. Loss: Categorical Crossentropy

3. Metric: Accuracy



Validation Set Accuracy: 58.38%

ResNet101 (transfer learning)

Transfer Learning Model Architecture

1. Base Model:

1. Pretrained ResNet101

2. Weights: ImageNet

3. Top layers removed (include_top=False)

4. Input shape: (128, 128, 3)

5. All layers frozen

2. Custom Head:

1. Flatten

2. Dense (64 units, ReLU)

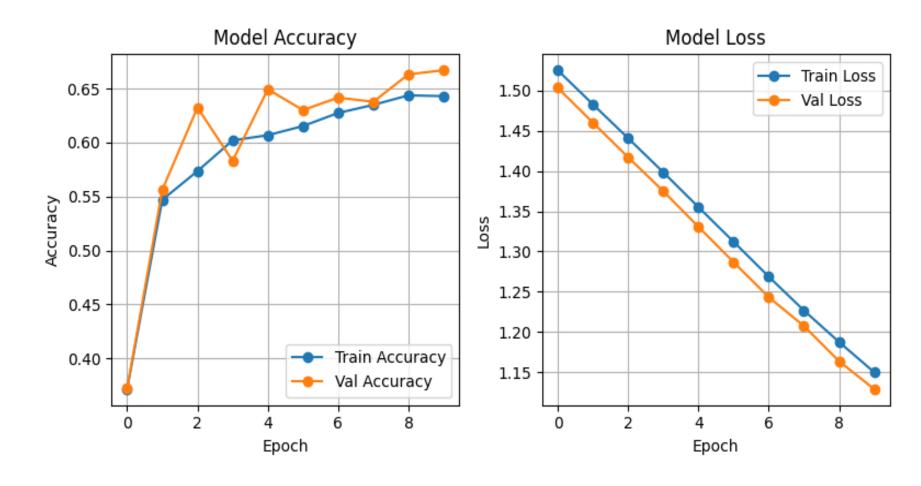
3. Output: Dense (5 units, Softmax)

3. Compilation:

1. Optimizer: Adam (learning rate = 1e-4)

2. Loss: Categorical Crossentropy

3. Metric: Accuracy



Validation Set Accuracy: 66.69%

Why ResNet might be performing poorly?

- Possible Reasons
 - The early layers of ResNet (especially ResNet50/101) have large initial kernels and stride, which may lose subtle patterns in highresolution CWT images early on.
 - CWT images are small (i.e, 128×128), deeper models like ResNet50 might not perform optimally due to aggressive downsampling in early layers.

Xception (transfer learning)

Transfer Learning Model Architecture

1. Base Model:

Pretrained Xception

2. Weights: ImageNet

3. Top layers removed (include_top=False)

4. Input shape: (128, 128, 3)

5. All layers frozen

2. Custom Head:

1. Flatten

2. Dense (64 units, ReLU)

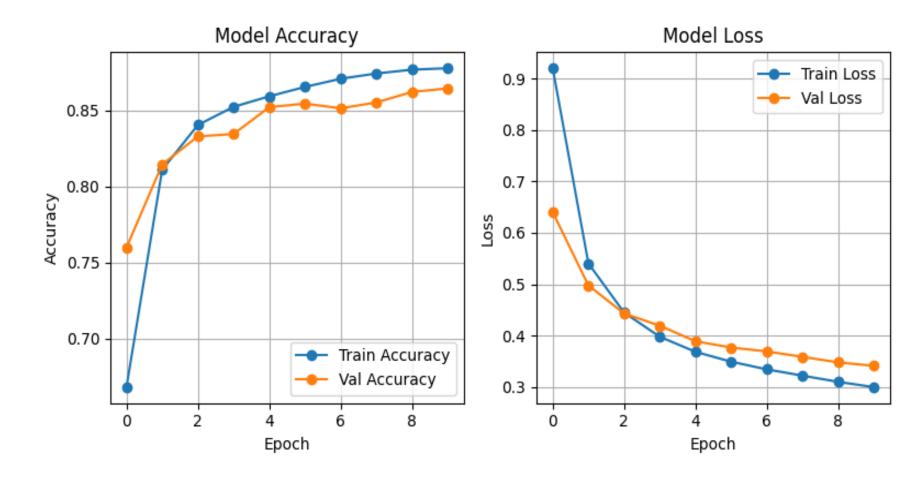
3. Output: Dense (5 units, Softmax)

3. Compilation:

1. Optimizer: Adam (learning rate = 1e-4)

2. Loss: Categorical Crossentropy

3. Metric: Accuracy



Validation Set Accuracy: 86.46%

DenseNet121 (transfer learning)

Transfer Learning Model Architecture

1. Base Model:

1. Pretrained DenseNet121

2. Weights: *ImageNet*

3. Top layers removed (include_top=False)

4. Input shape: (128, 128, 3)

5. All layers frozen

2. Custom Head:

1. Flatten

2. Dense (64 units, ReLU)

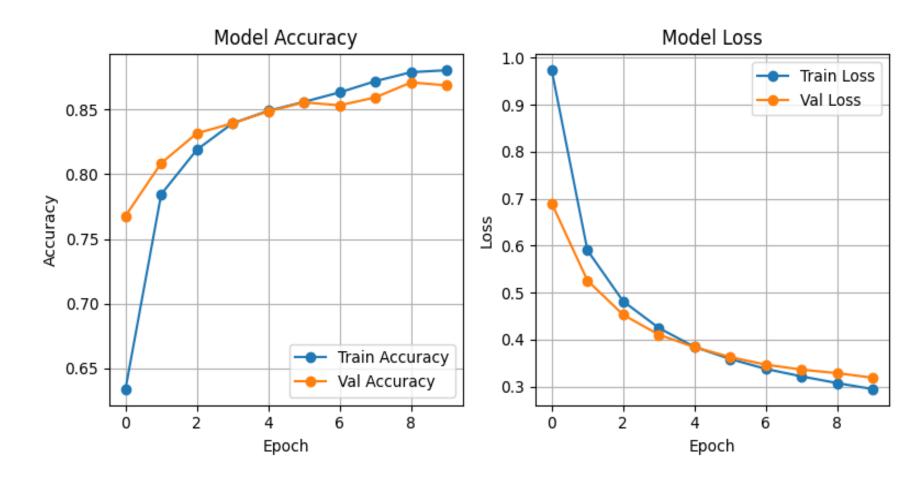
3. Output: Dense (5 units, Softmax)

3. Compilation:

1. Optimizer: Adam (learning rate = 1e-4)

2. Loss: Categorical Crossentropy

3. Metric: Accuracy



Validation Set Accuracy: 86.85%