

A decorative graphic on the left side of the slide, consisting of a network of white lines and small circles on a blue gradient background, resembling a circuit board or neural network connections.

PERFORMANCE OF MULTILAYER PERCEPTRONS ON ECG ANALYSIS

CS615: DEEP LEARNING

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BACKGROUND

- US Healthcare System substantially behind in basic safety₁
- Annually, 98,000 deaths due to medical errors in US hospitals₁
- IOM encouraged use of IT₂
 - Simplify workflow processes
 - Reduce waste
 - Improve healthcare access

1 (INSTITUTE OF MEDICINE [IOM], 2000, P.5)

2 (IOM, 2009)

BACKGROUND

- Effective treatment of Acute Coronary Syndrome (ACS) is time sensitive
- Current guidelines identify ACS on ECG by:
 - ST-segment elevation (STEMI)
 - Elevated biomarker detection (NSTEMI)
- 24-35% of those with NSTEMI need emergent catheterization
 - Delayed while awaiting biomarker elevation due to inability to detect on ECG

PREVIOUS WORK

- Literature Review 2022
 - 59 Deep Learning studies analyzing ECG data
 - Convolutional Neural Networks and ResNet best performing
 - Reported over 97% accuracy
 - ECG data is noisy
 - May be more suited to DL than ML

PREVIOUS WORK

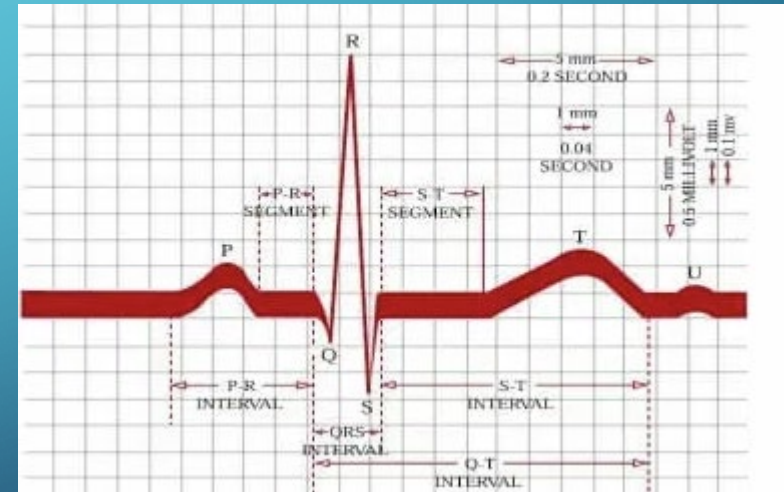
- 2023 prospective, observational
 - 2023 prospective, observational cohort
 - 12-Lead ECG Data
 - 10 different ML classifiers
 - Random Forest Model achieved AUROC 0.91 (95% CI = 0.87-0.96)
 - Outperformed practicing clinicians and commercial ECG systems

PURPOSE

- To analyze the performance of MLP classification models on identifying abnormal heart rhythms and myocardial ischemia on ECG data at various degrees of complexity

QUESTION

- Can a MLP classification model accurately identify cardiac dysrhythmias and myocardial ischemia from ECG tracings?



DATA

- ECG 200¹
 - N = 200
 - 1D time series of ECG tracing, single cycle
 - Classes: Normal vs Abnormal
- MIT-BIH³
 - N = 47 subjects
 - 2 lead ECG tracing, 30 minutes long
 - Classes: Normal vs Ischemia
- ECG 5000²
 - N = 5000
 - 1D time series of ECG tracing, single cycle
 - Classes:
 - Normal
 - R-on-T PVC
 - PVC
 - Escape Beat
 - Ventricular Fusion

1 (DAU, ET AL., 2018)

2 (GOLDBERGER ET AL., 2000)

3 (MOODY & MARK, 2001)

DATA

- Compare our model against Al-Zaiti's 2023 RF Model
 - 12-Lead ECG Data
 - Random Forest Model achieved AUROC 0.91 (95% CI = 0.87-0.96)
 - AUROC 0.91 = Benchmark for our Model

PRE-PROCESSING

- ECG 200 and ECG5000 data
 - Available from author cleaned and split 50/50 training/validation sets
 - Data were re-combined, shuffled, and split 75/25 training/validation
 - ECG5000 class outcomes were one-hot-encoded
 - Significant oversampling of normal tracings

PRE-PROCESSING

- MIT-BIH data
 - Much more complicated; 2 lead ECG tracing over time
 - Sliding window was created to take 6 second snapshot of tracing
 - Any observation with missing data was omitted
 - Each lead data within a window was concatenated together to form single observation
 - 3000 features per observation
 - Subsample of 1725 observations; 861 normal and 864 ischemia

PRE-PROCESSING

- Full ECG data (Al-Zaiti)
 - Data features taken AS-IS
 - No randomization or shuffling. We used their training/validation data, as published
 - $N = 7313 \rightarrow \text{Training} = 4026, \text{Validation} = 3287$
 - 74 features, including patient demographics
 - BUT very imbalanced distribution, very high majority of class normal
 - Approximately 370 of each class in both training and validation set
 - Missingness is a problem
 - Extreme outliers ($> 5 \text{ SD}$) were removed
 - All NaN values were replaced by column mean after removing extreme outliers

METHODS

- On each dataset, compare performance of
 - Shallow MLP (5 Layers); INPUT \rightarrow FCC \rightarrow TANH \rightarrow LOGSIG \rightarrow LOG LOSS
 - Deep MLP (22 Layers); INPUT \rightarrow (FCC \rightarrow TANH) \times 9 \rightarrow FCC \rightarrow LOGSIG \rightarrow LOGLOSS
 - Deep MLP with Skip residuals
 - Same 22 Layers as Deep but each FCC \rightarrow TANH sequence is wrapped with skip residuals and batch normalization layer
 - FCC use hidden state size funneling ; dimension size -2 every other skip layer block
 - TANH is prone to dead activations, skip layer should attenuate that problem
 - TANH proved too unstable for Full ECG data, so we switched all TANH to ReLU
 - BATCHNORM layer just before final LOGSIG for added stability

METHODS

- Hyperparameter Tuning
 - Hidden State (FCC output dimension) and learning rate
 - Optimized by painful trial and error for each model
 - Iterative breadth-search
 - All permutations of $\text{Eta} = [10^{-3}, 10^{-4}, 10^{-5}]$ and Hidden State Size $[8, 16, 32, 64]$
 - ECG200 -----> $\text{Eta} = .00001$, Hidden State Size = 32
 - ECG5000 -----> $\text{Eta} = .0001$, Hidden State Size = 32
 - ECG5000 Multiclass -----> $\text{Eta} = .0001$, Hidden State Size = 32
 - ECG5000 Subsampled -----> $\text{Eta} = .00001$, Hidden State Size = 32
 - MIT-BIH -----> $\text{Eta} = .00001$, Hidden State Size = 32
 - FULL ECG -----> $\text{Eta} = .00032$, Hidden State Size = 16
 - All FCC use ADAM except the FULL ECG pipeline
 - Removed due to instability



METHODS

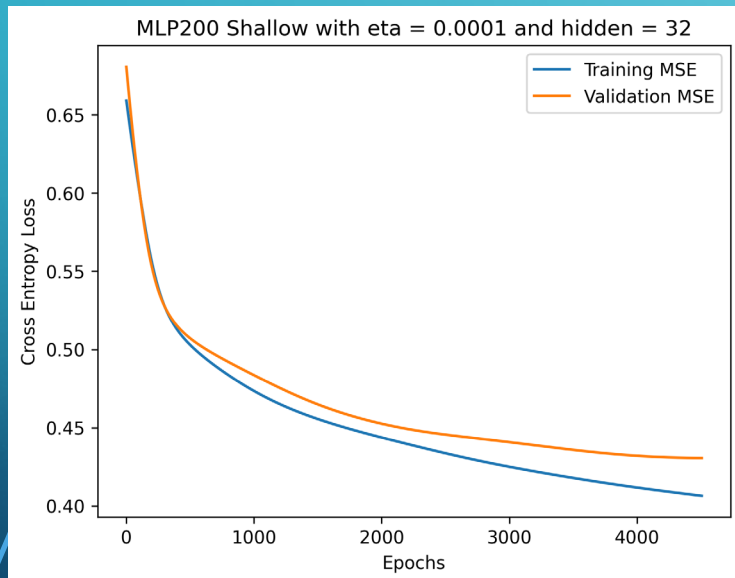
- FCC weight and bias initialization
 - Random, uniformly distributed, Xavier

- $\left[-\sqrt{\frac{6}{h_{in}+h_{out}}}, \sqrt{\frac{6}{h_{in}+h_{out}}} \right]$

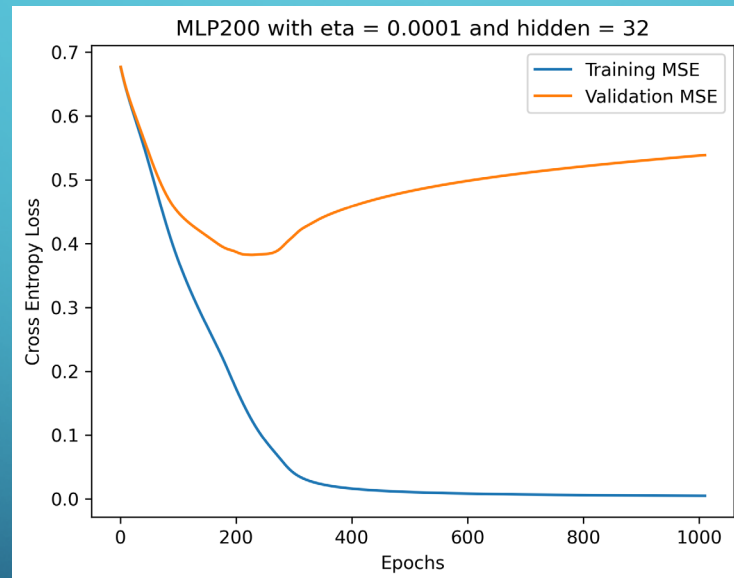


RESULTS-ECG200

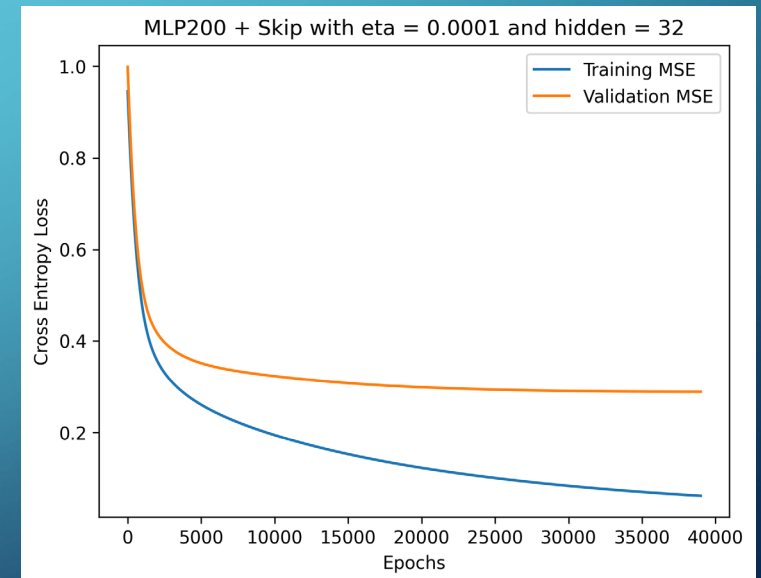
Shallow - Trained Well



Deep – Exploding Gradient



Skip Residuals Fix it!
Still overfit



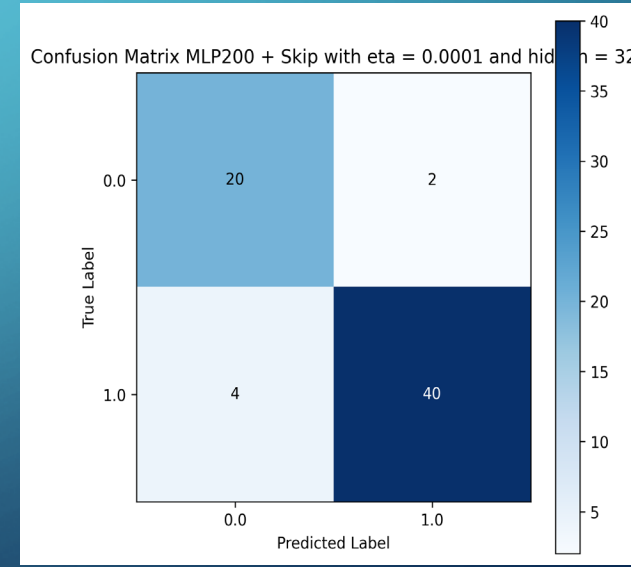
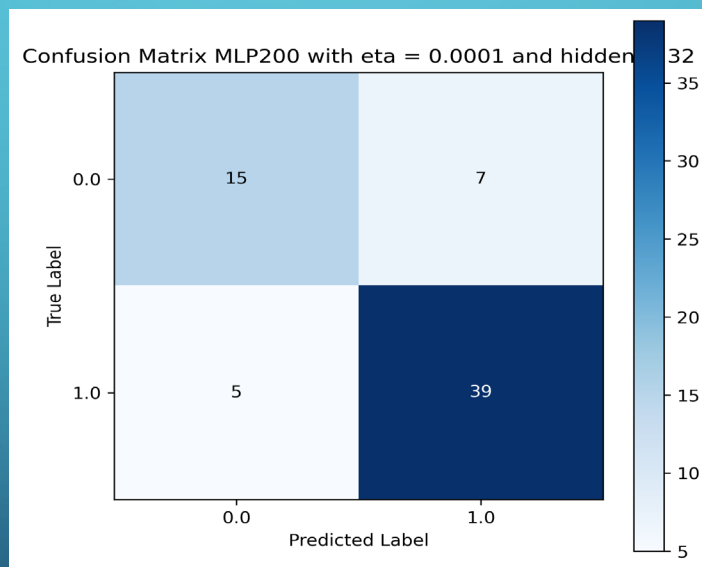
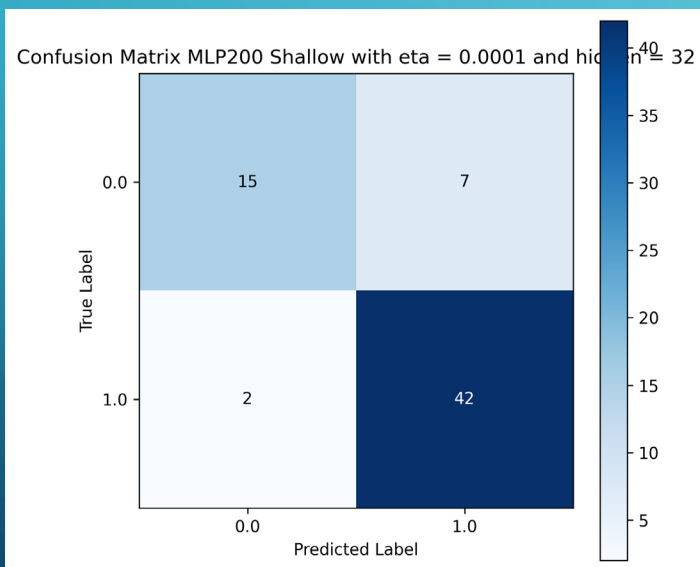
RESULTS-ECG200



Shallow Accuracy = 86%

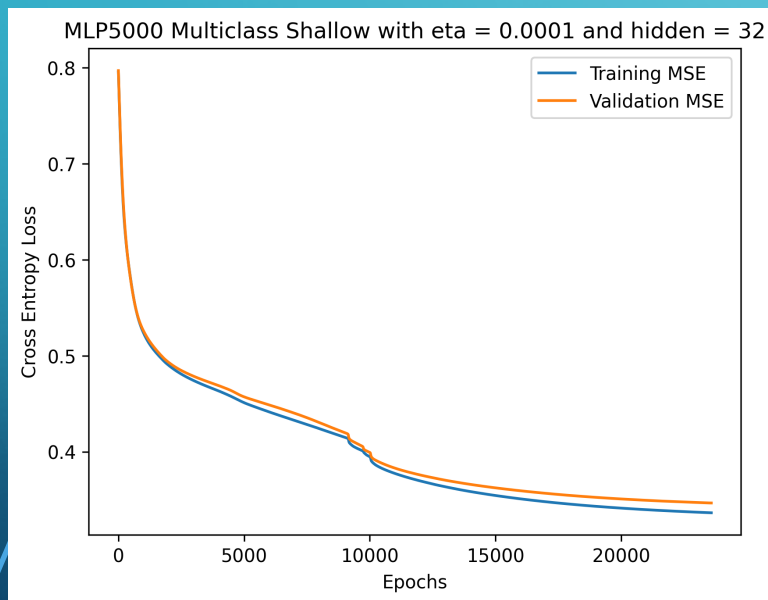
Deep Accuracy = 81%

Deep with Skip Accuracy = 91%

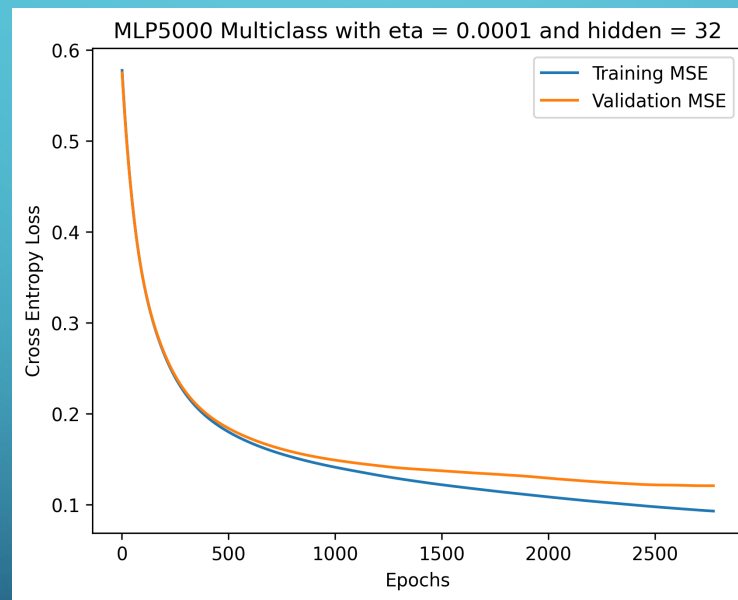


RESULTS-ECG5000 NORMAL VS ABNORMAL

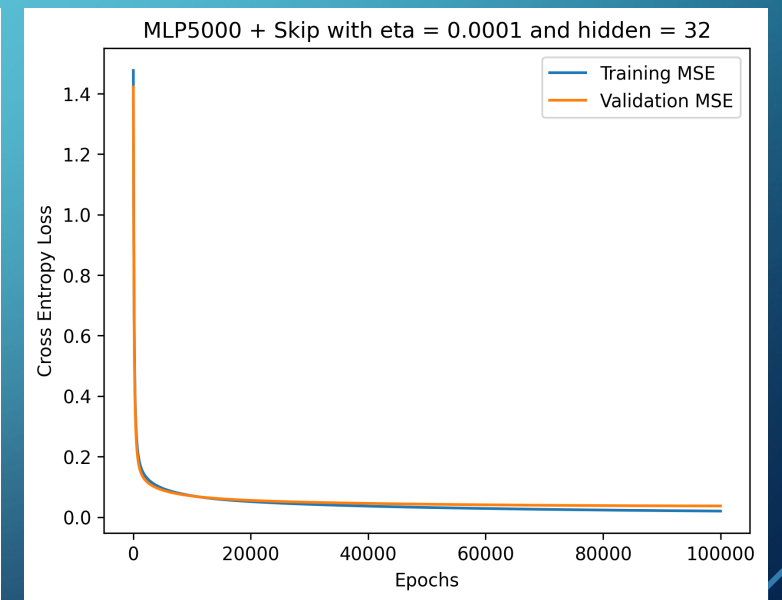
Shallow - Good



Deep - Better

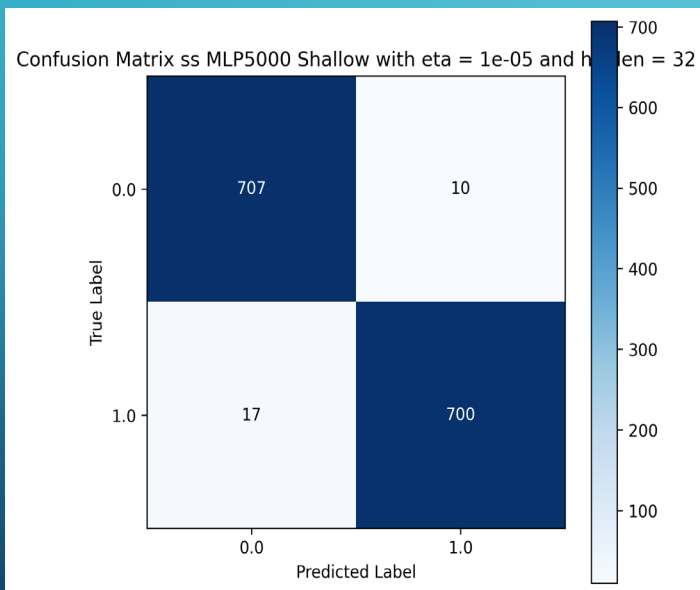


Skip Residuals - Best

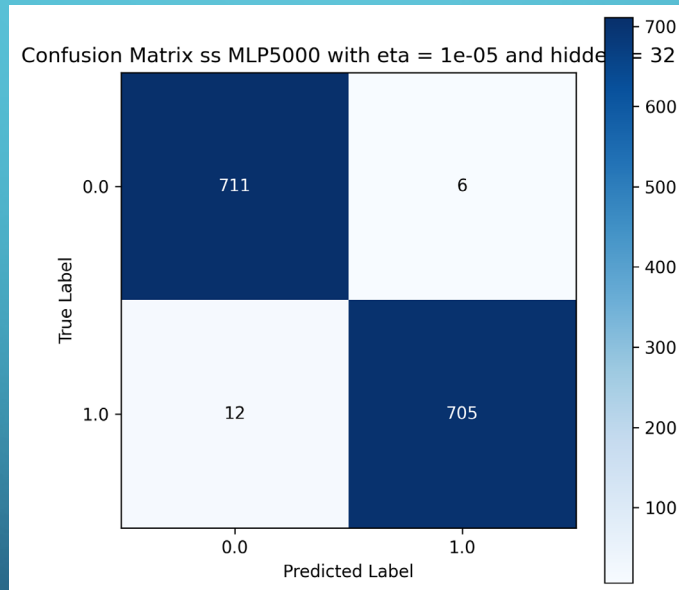


RESULTS-ECG5000 NORMAL VS ABNORMAL

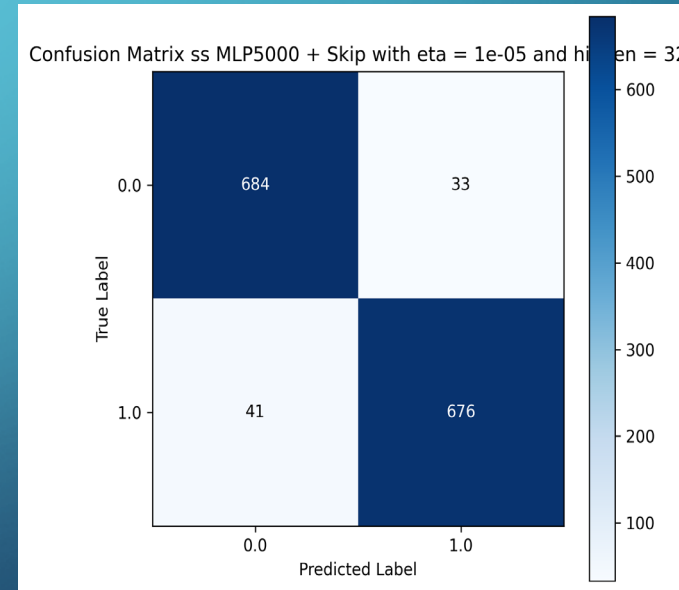
Shallow Accuracy = 98%



Deep Accuracy = 98%

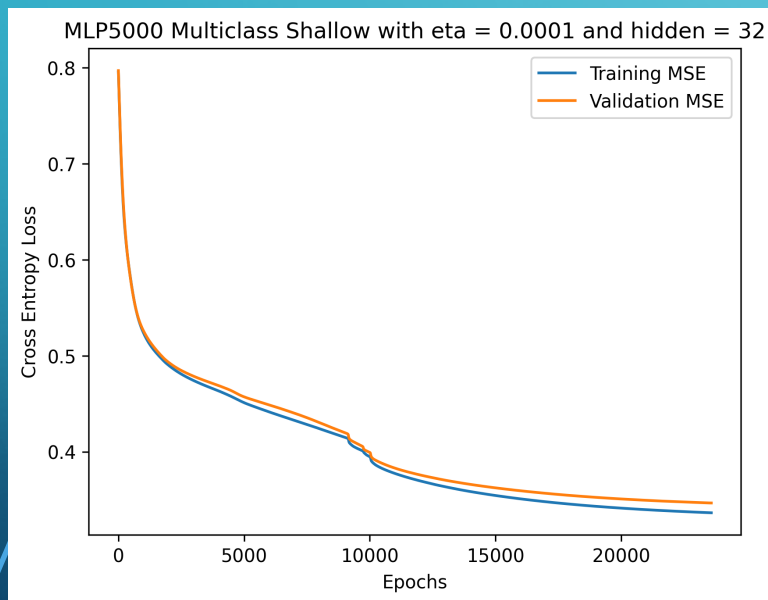


Deep with Skip Accuracy = 98%

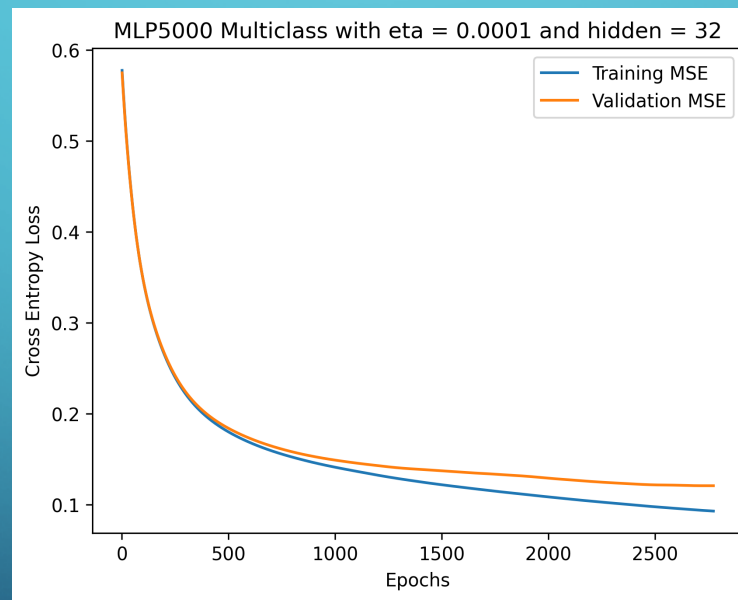


RESULTS-ECG5000 MULTICLASS

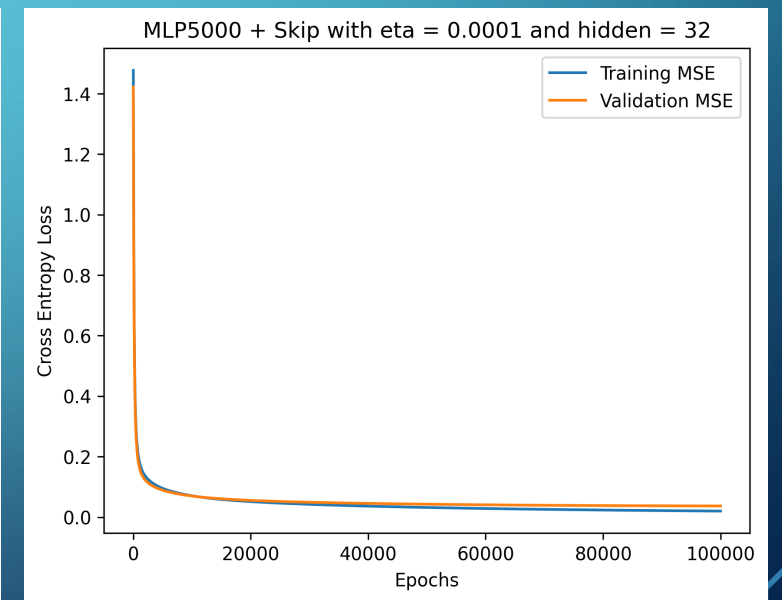
Shallow - Good



Deep - Better



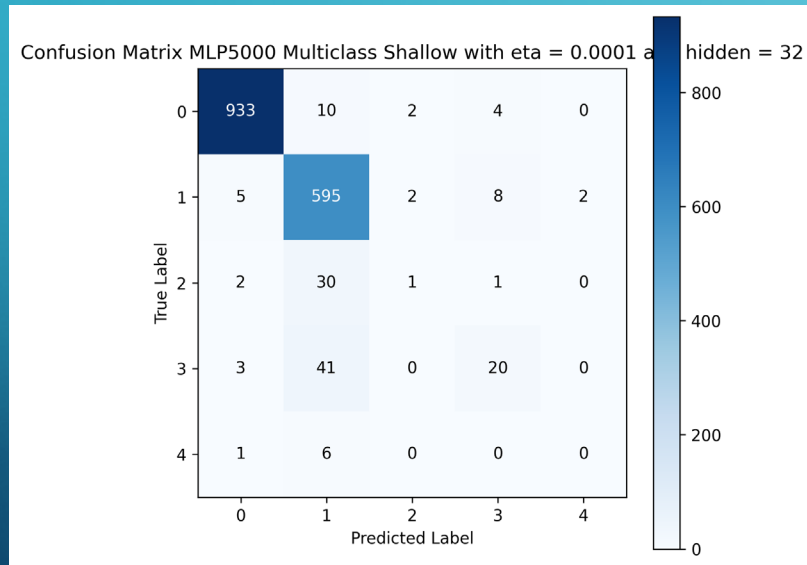
Skip Residuals - Best



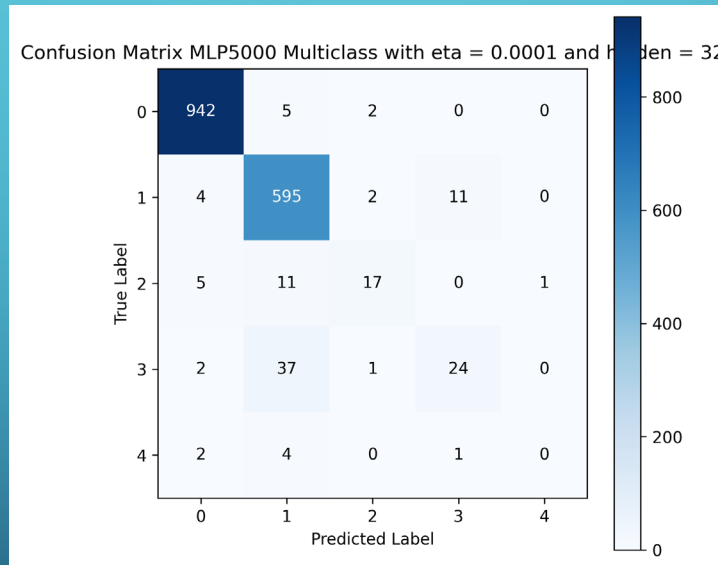
RESULTS-ECG5000 MULTICLASS



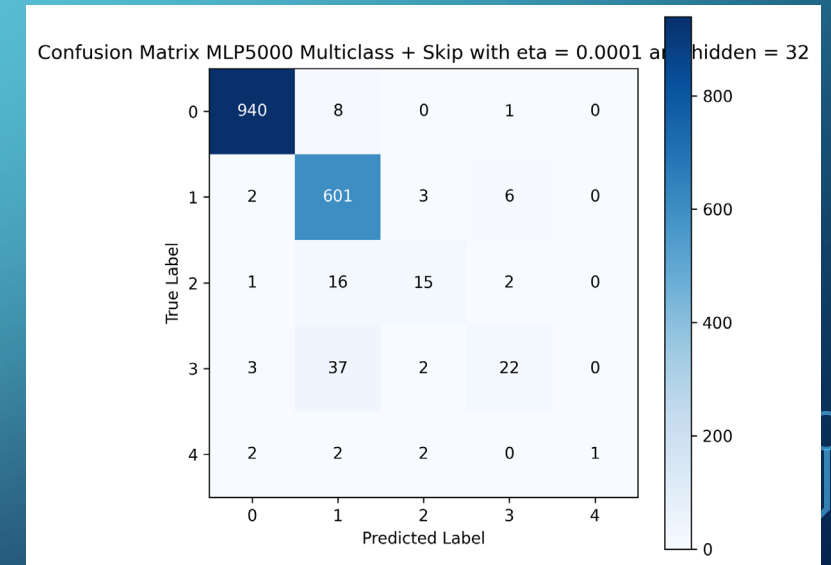
Shallow Accuracy = 92%



Deep Accuracy = 95%



Deep with Skip Accuracy = 95%



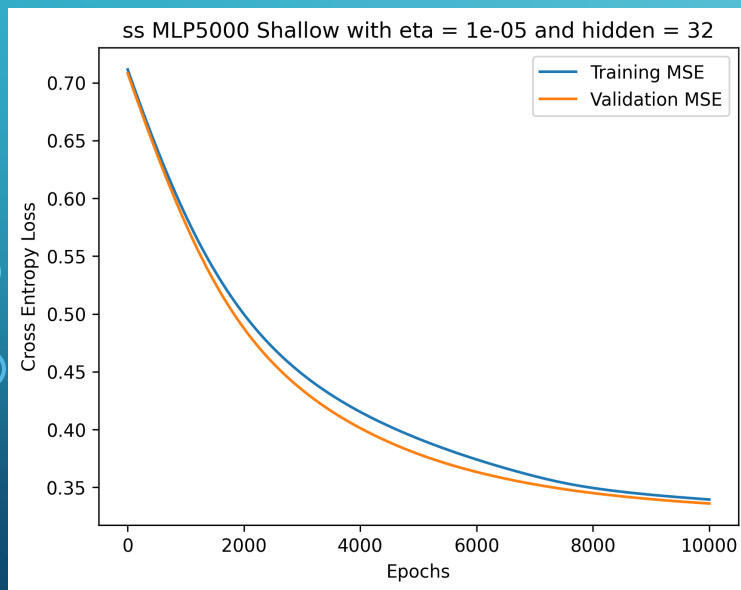
BUT THAT'S A LOT OF NORMAL CASES...

- Does it perform well just because of oversampled normal cases?

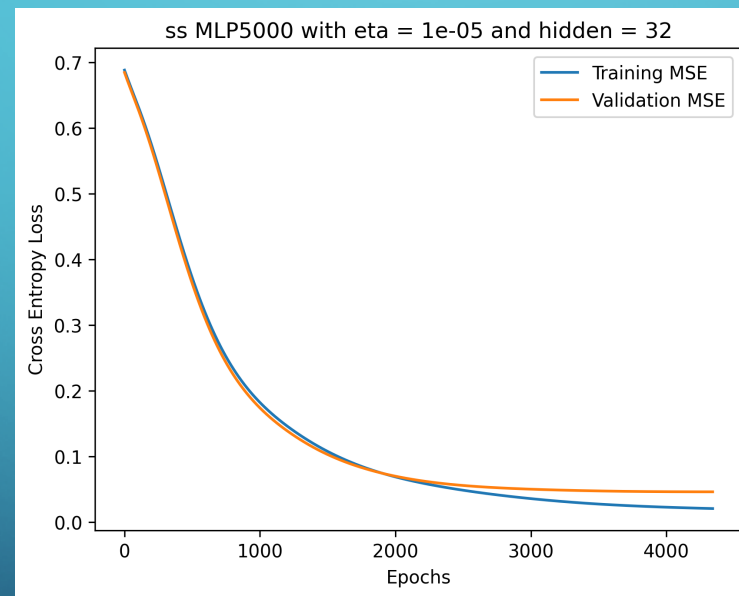


RESULTS-ECG5000 SUBSAMPLED NORMAL VS ABNORMAL

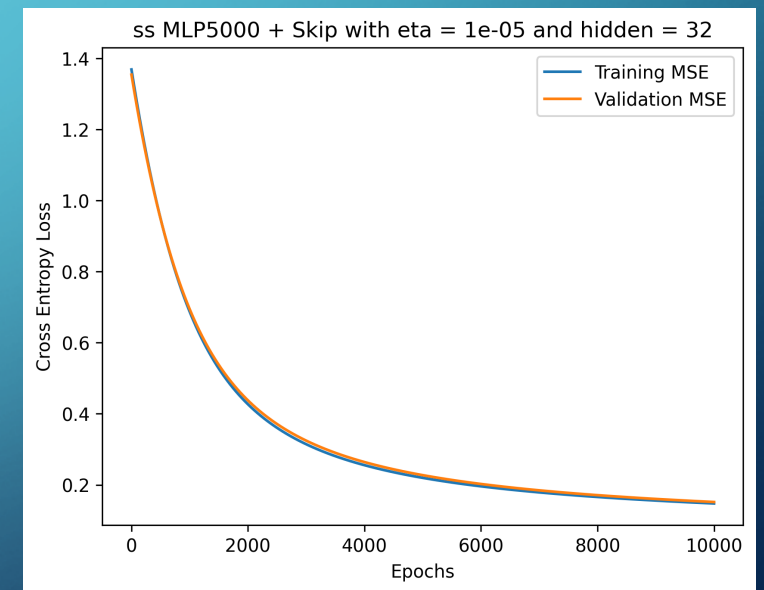
Shallow



Deep



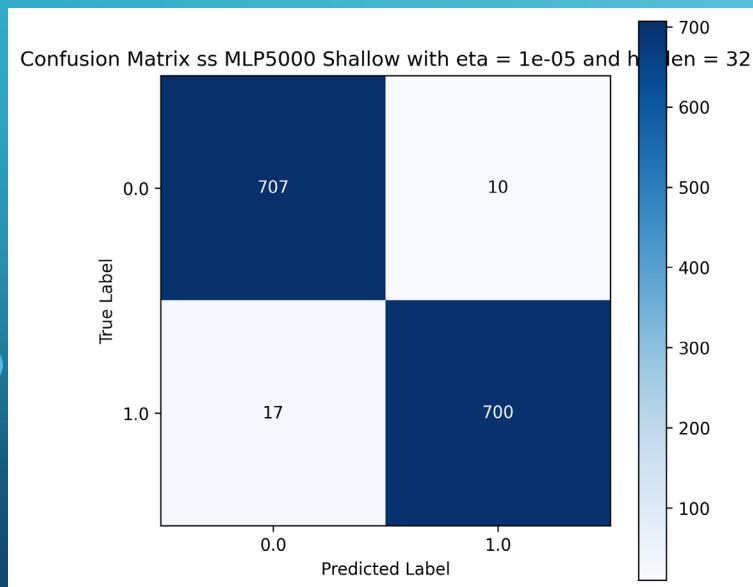
Deep with Skip



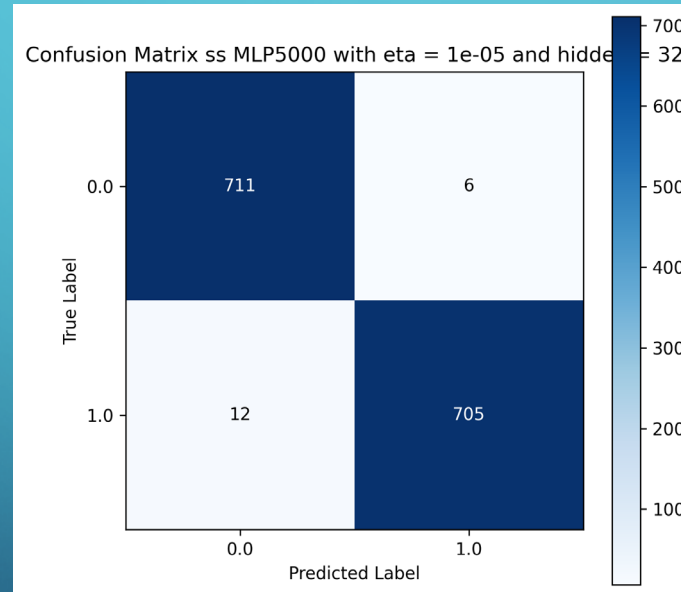
RESULTS-ECG5000 SUBSAMPLED NORMAL VS ABNORMAL



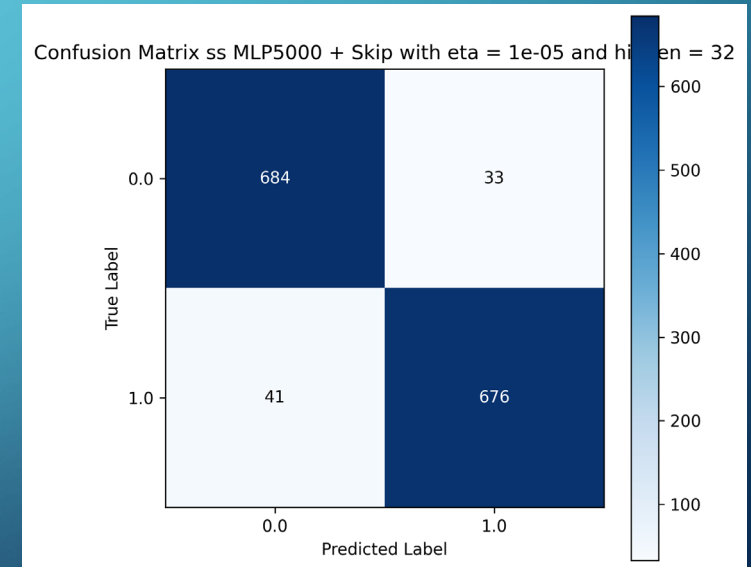
Shallow Accuracy = 98%



Deep Accuracy = 99%

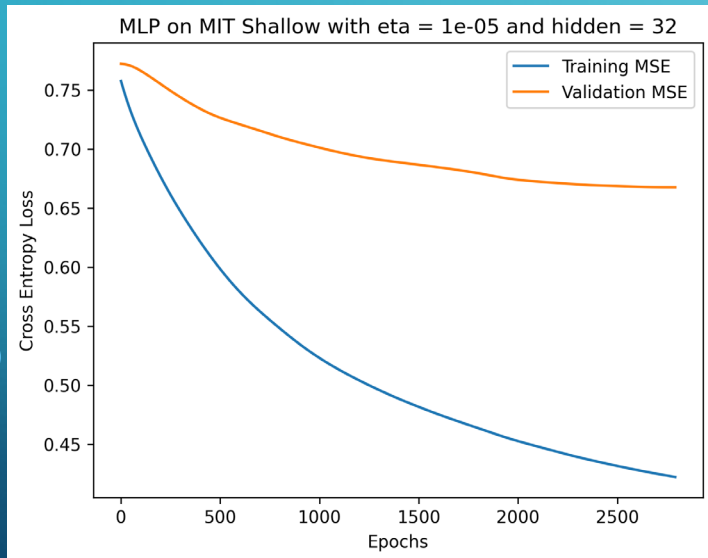


Deep with Skip Accuracy = 95%

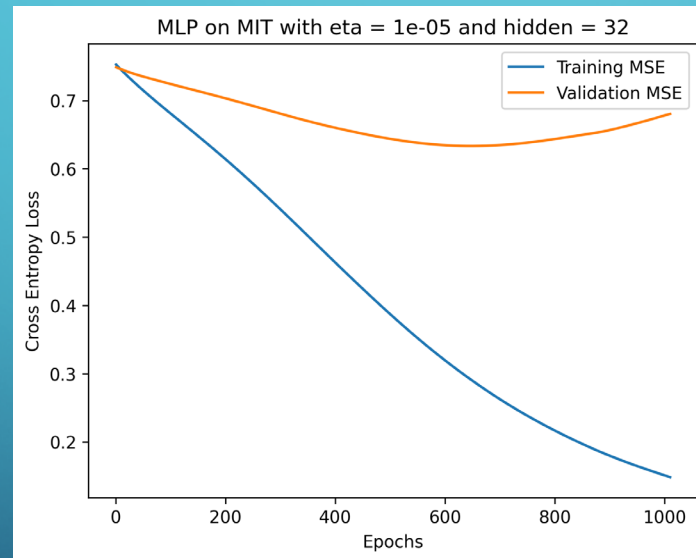


RESULTS MIT-BIH

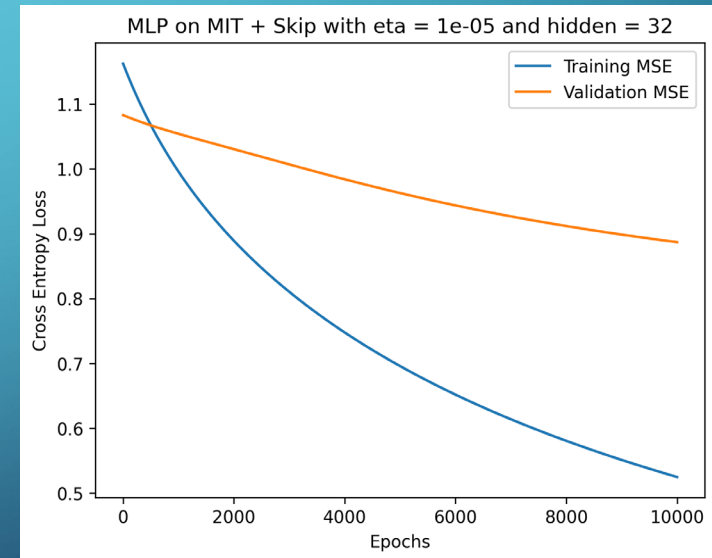
Shallow – Very overfit



Deep – Exploding Gradient



Skip Residuals... at least the line is pointing in the right direction

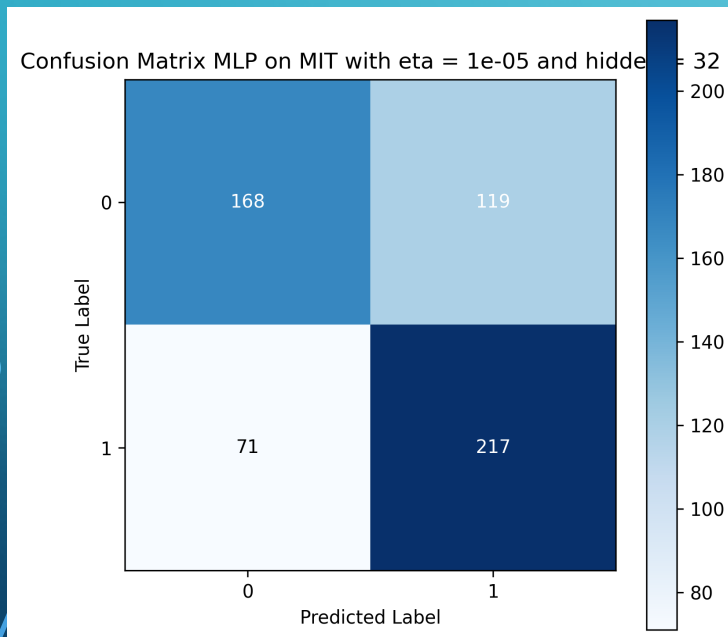


I RAN THIS FOR ANOTHER 10000 EPOCHS..
IT DOES NOT IMPROVE

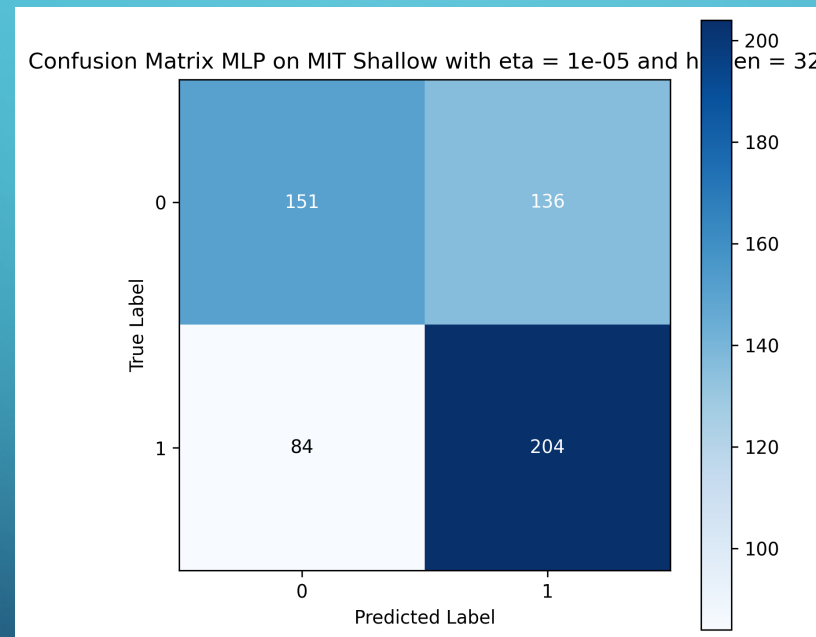
RESULT MIT-BIH



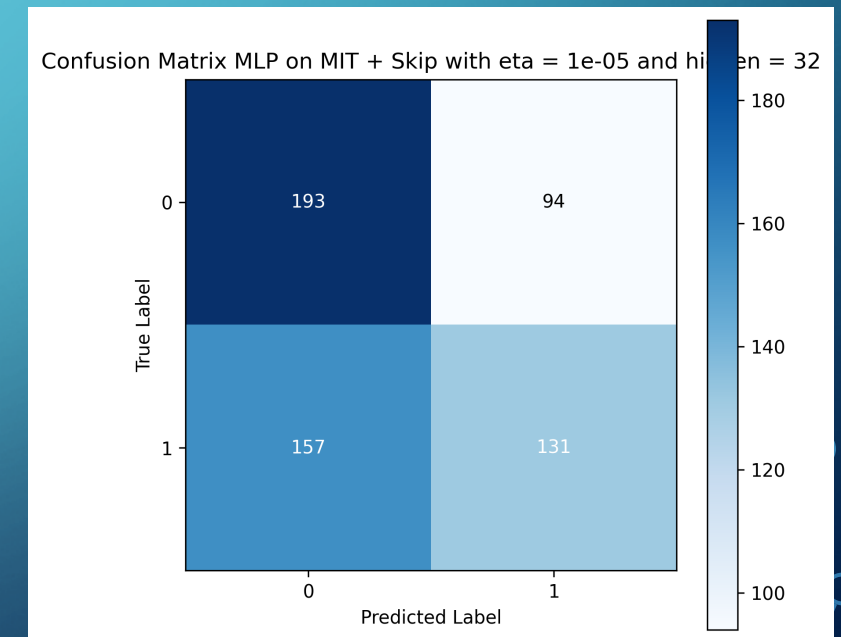
Shallow Accuracy = 67%



Deep Accuracy = 61%

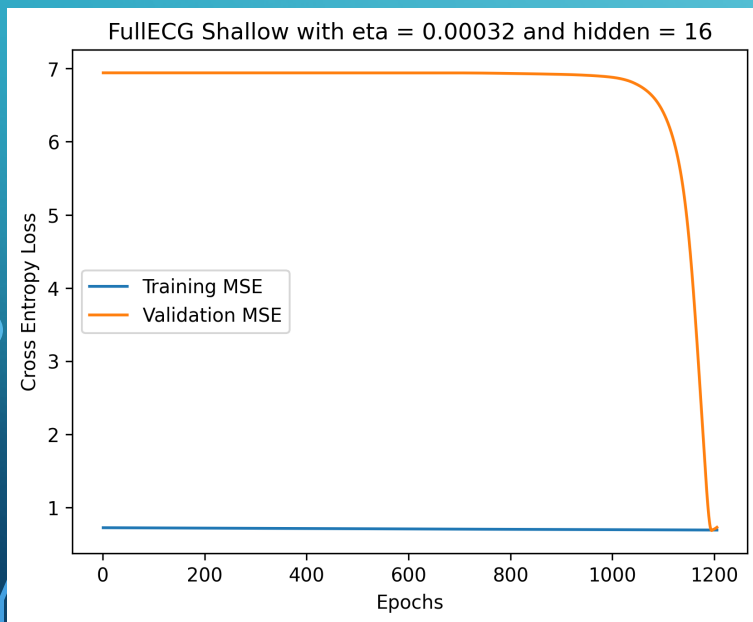


Deep with Skip Accuracy = 56%

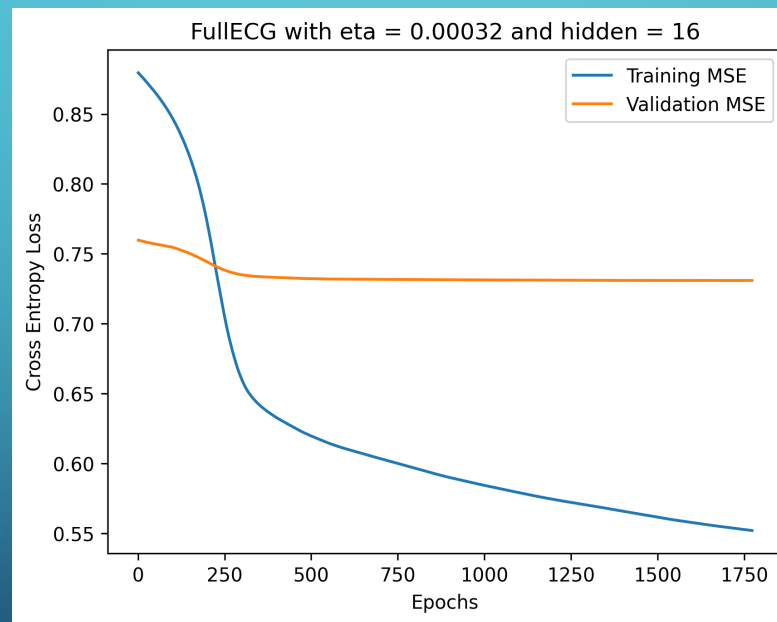


RESULT-FULL ECG

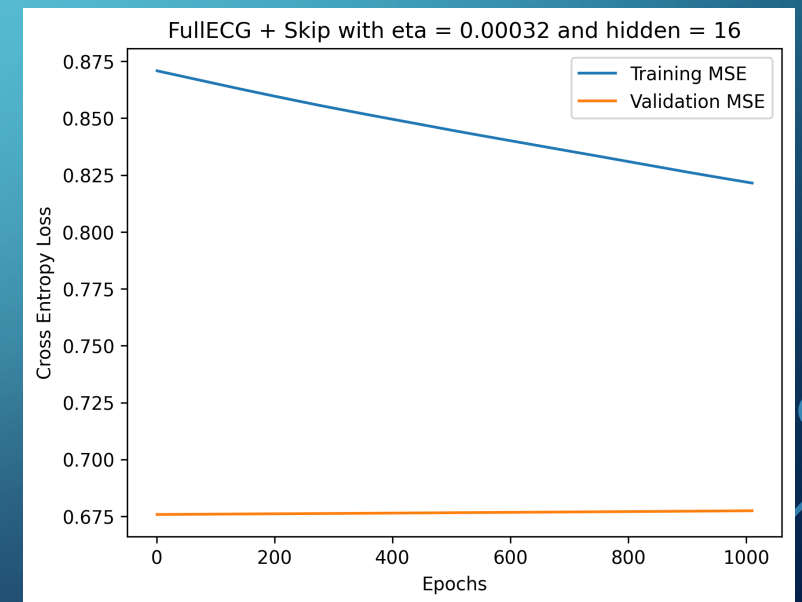
Shallow – Learned Something?



Deep – Overfit



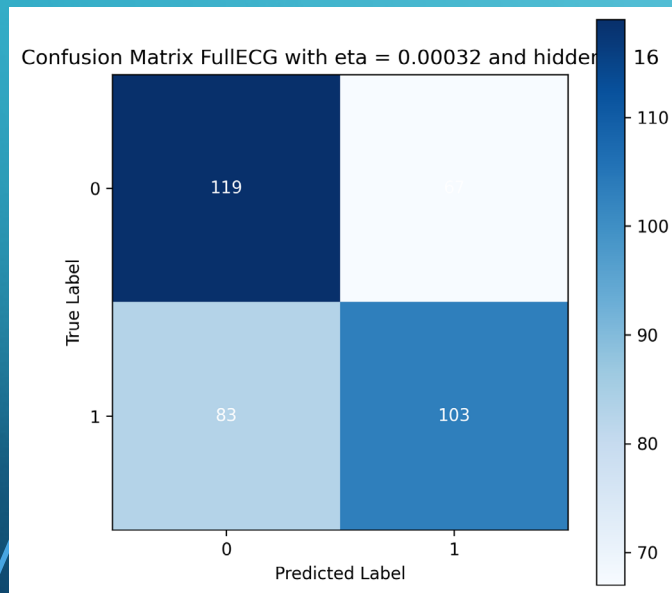
Deep with Skip - Guesses



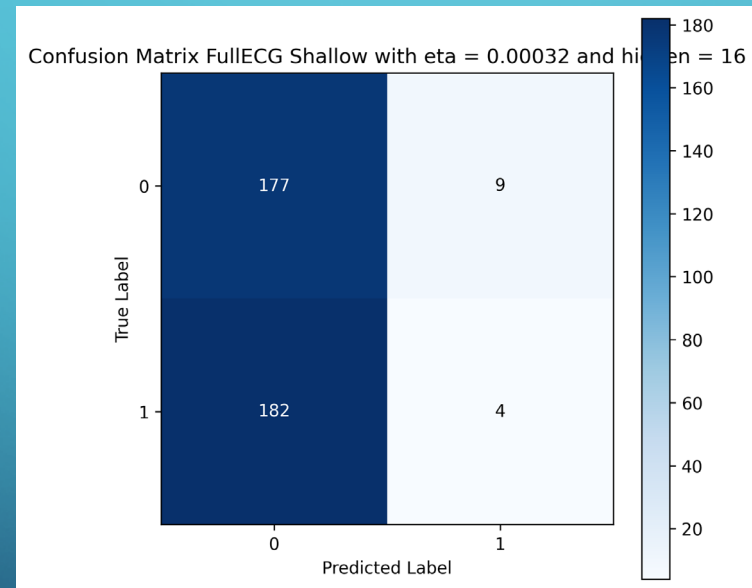
RESULTS-FULL ECG



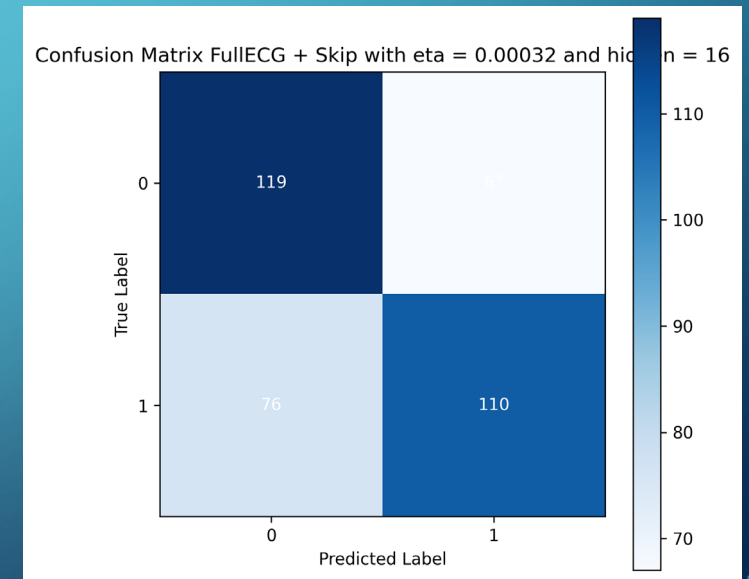
Shallow Accuracy = 72%



Deep Accuracy = 48%



Deep with Skip Accuracy = 75%



ANALYSIS – THE GOOD!

- Our MLP was effectively trained to identify normal vs ischemic single-cycle ECG tracing with an accuracy of 91%, made possible with the addition of skip residuals
- Our MLP was effective at identifying normal vs abnormal single-cycle ECG tracings. The shallow model performed with an accuracy of ***, leaving little motivation for more complex models or skip residuals

ANALYSIS – THE NOT SO GOOD...

- More complex datasets were not effectively analyzed by our MLP, despite numerous architecture modifications and optimizations, including a breadth search for optimal hyperparameters and attenuation of data missingness

DISCUSSION

- While MLP may be suitable for many classification problems, feature rich data may still prove too complex for this method.
- No globally-optimal model exists
- Multi-lead ECG data may be better classified using more complex models, such as Convolutional Neural Networks, Transformers, or Random Forest Models
- Future work is recommended to evaluate these datasets using more complex models

REFERENCES

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