



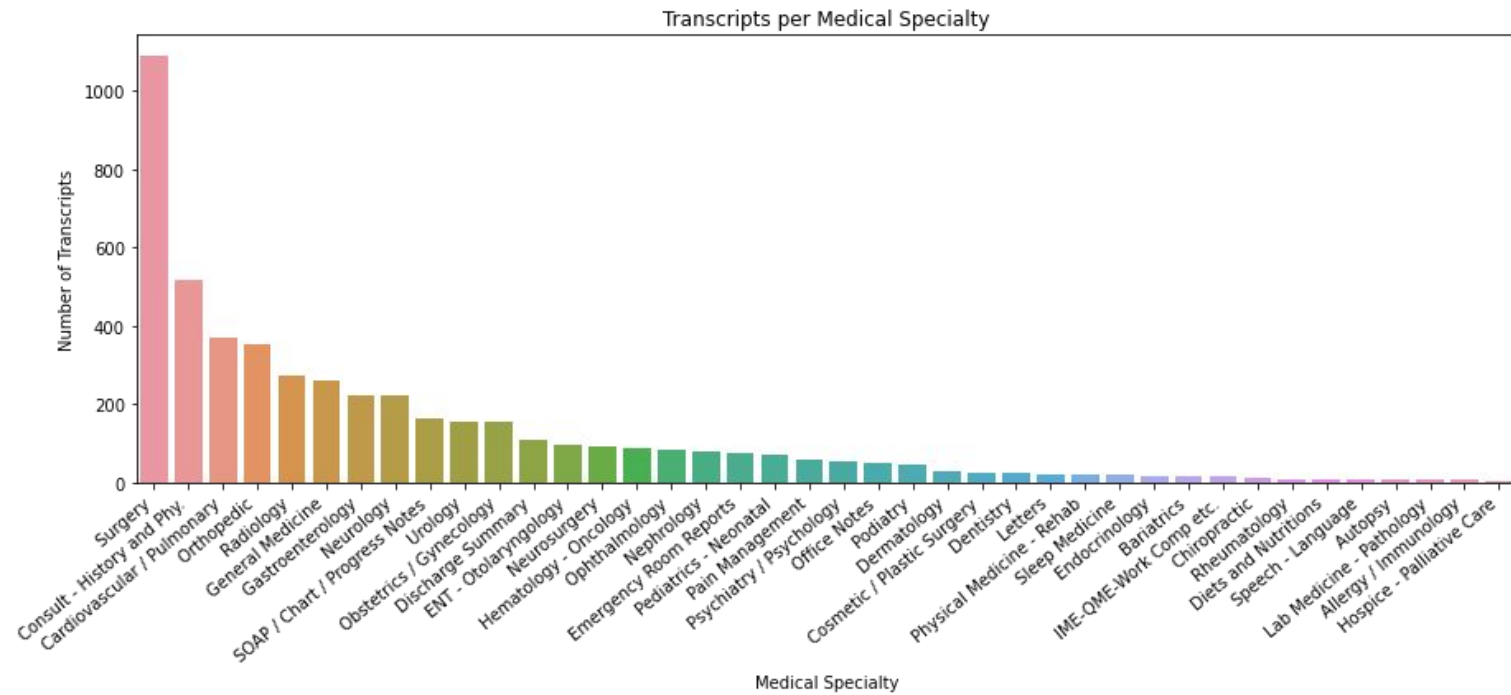
# Medical Transcription Classification

Freya Gray  
CS39AA

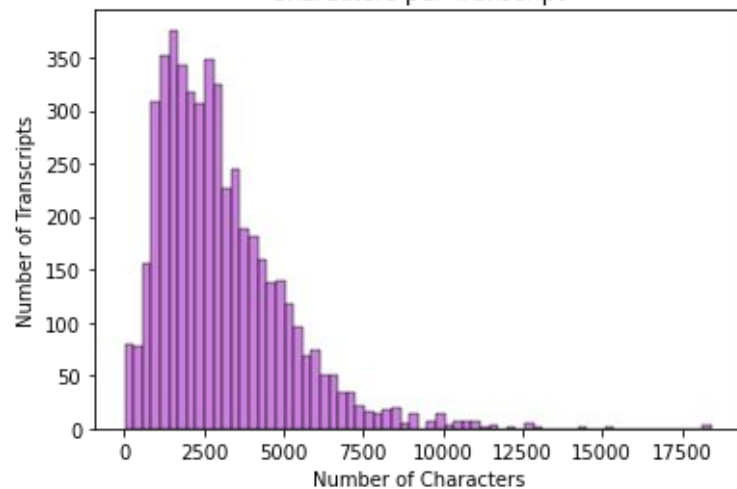


# Dataset

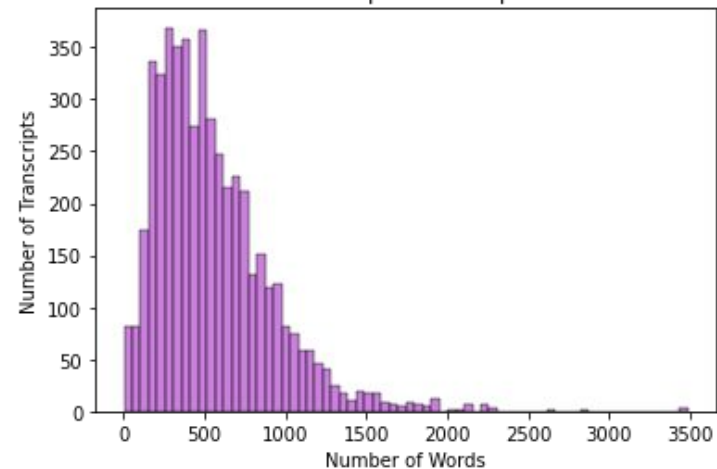
- Medical transcripts from [mhsamples.com](https://mhsamples.com)
- Transcripts from medical professionals after seeing a patient
- All names and dates have been changed or removed
- 40 different medical specialties
- 4966 transcripts
- Transcripts come in a wide range of formats and styles



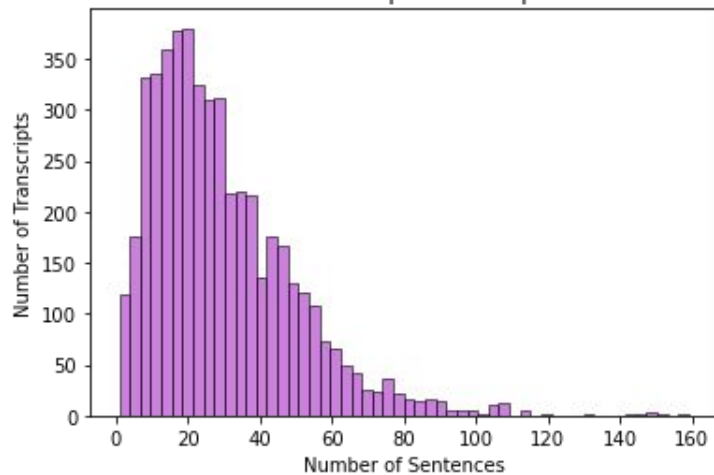
Characters per Transcript



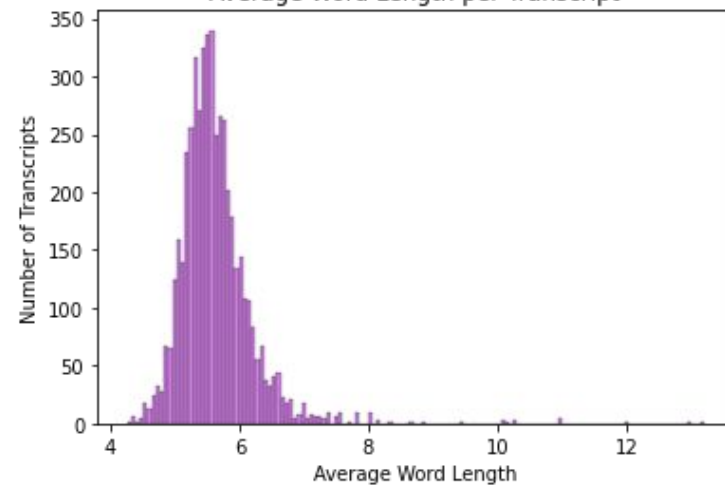
Words per Transcript

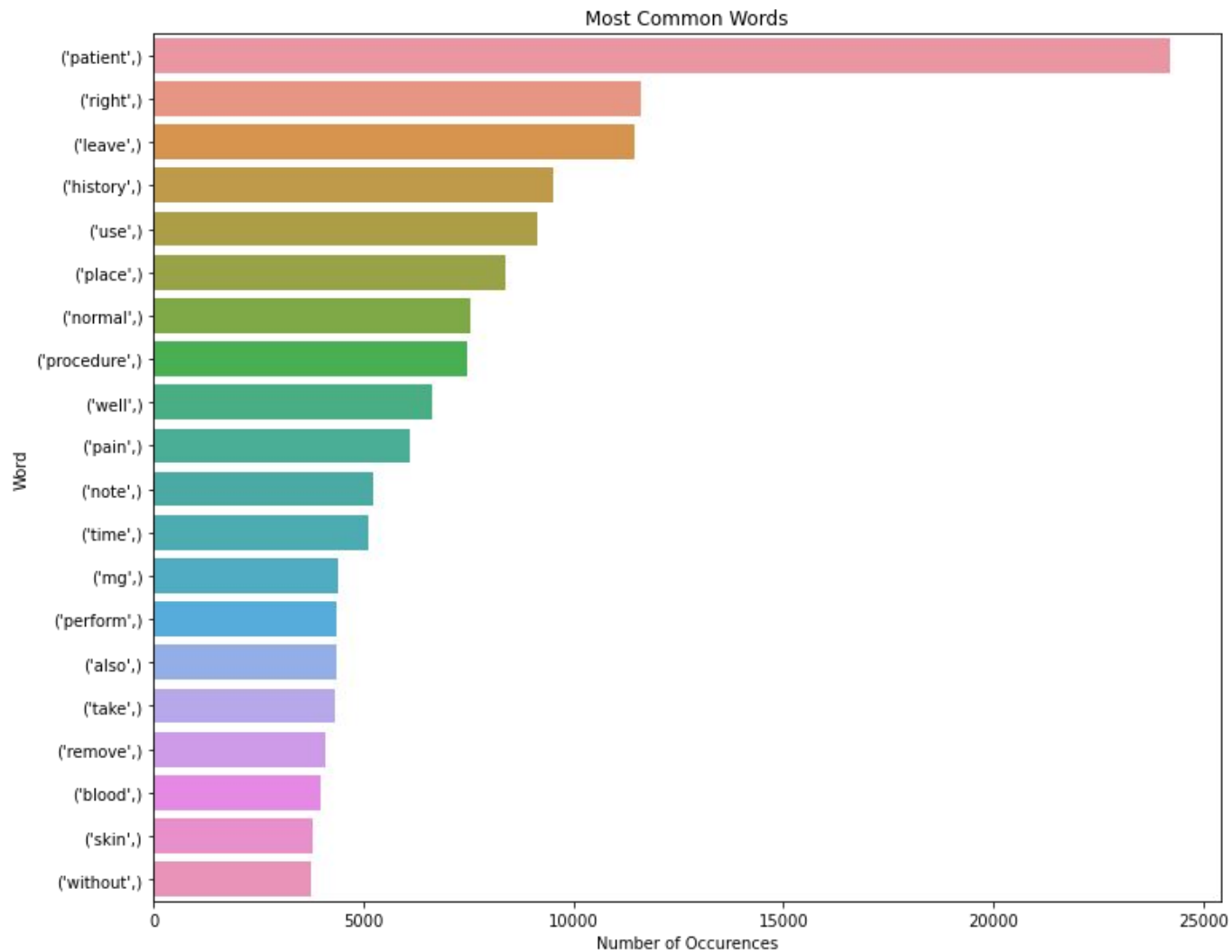


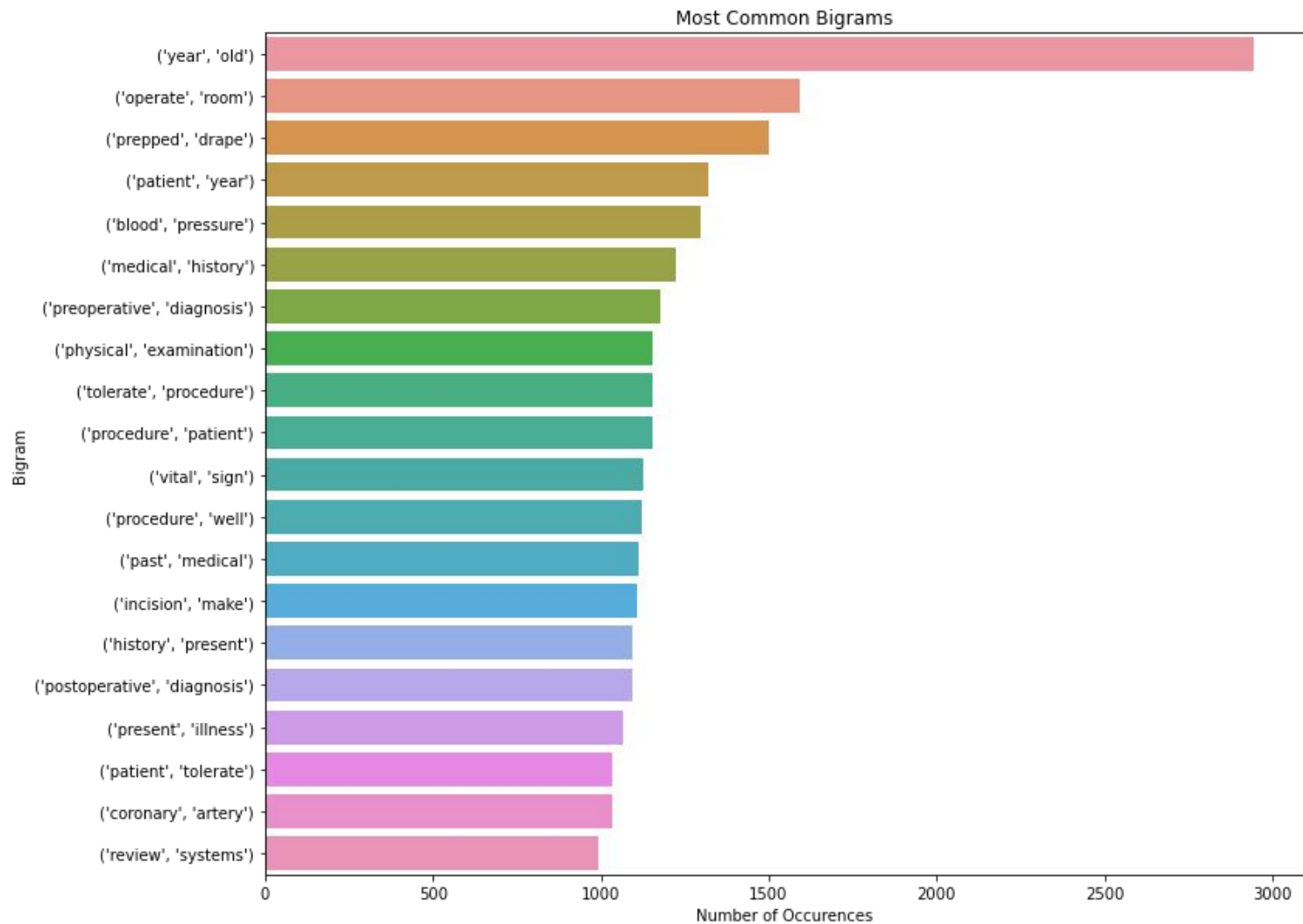
Sentences per Transcript



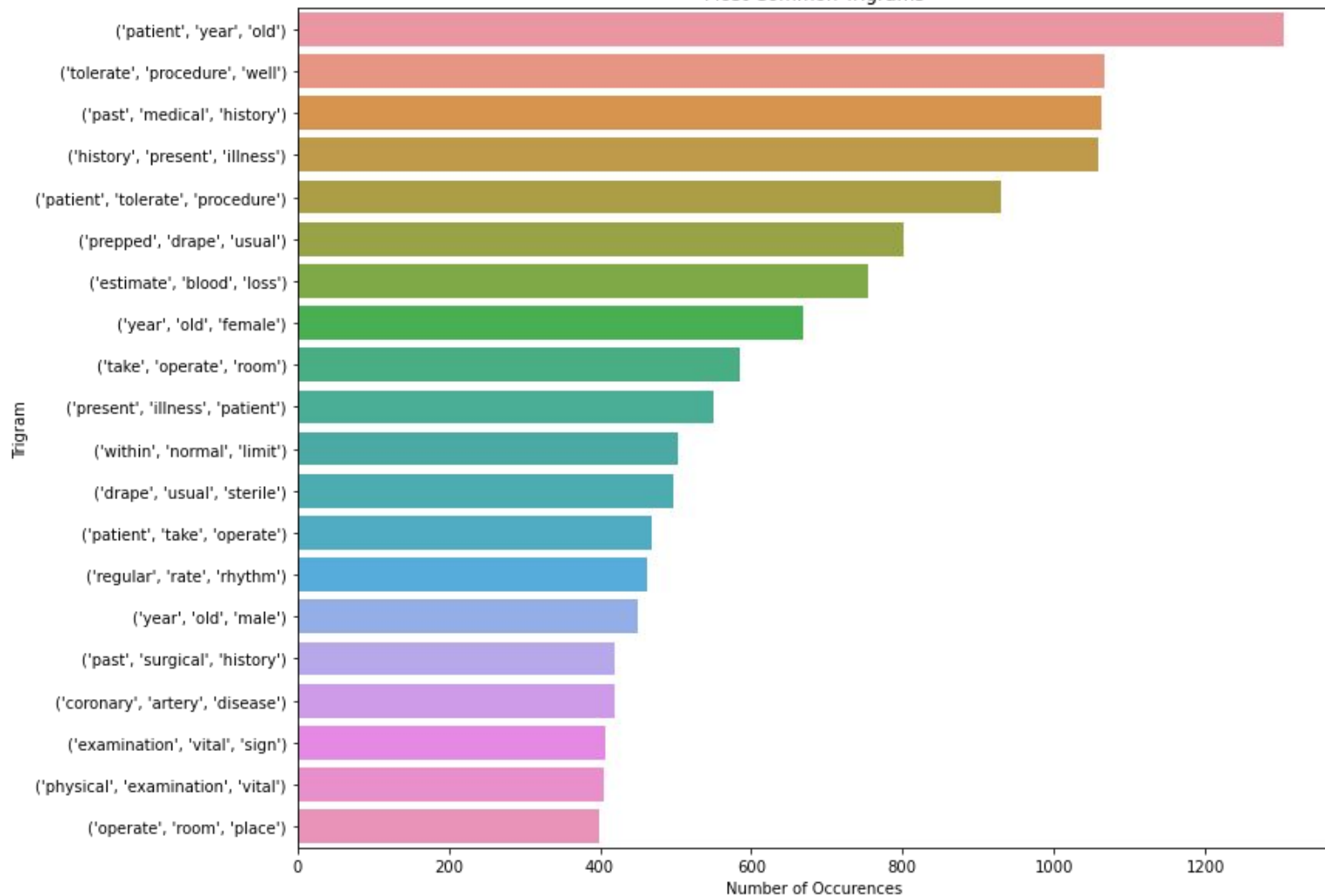
Average Word Length per Transcript







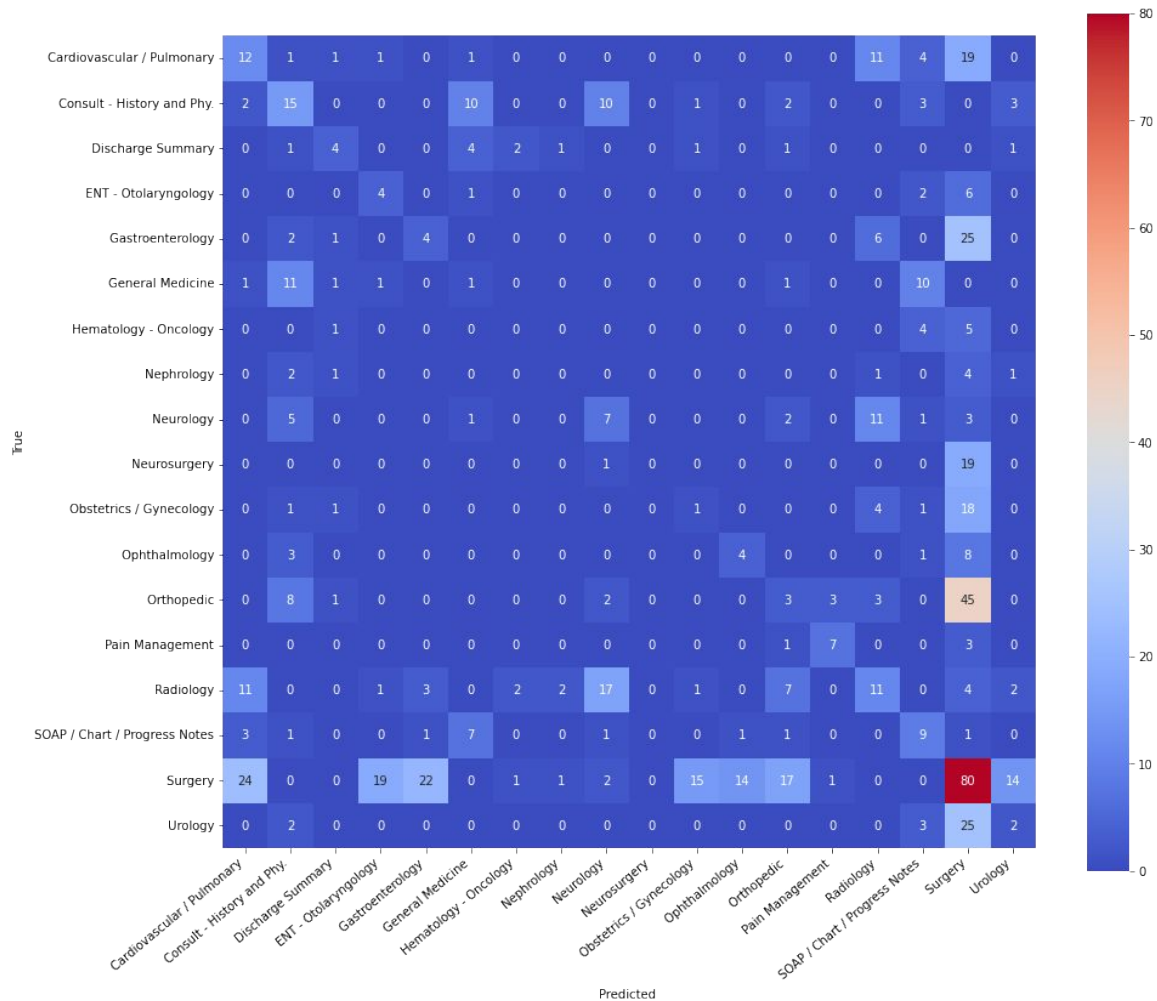
Most Common Trigrams



# Baseline Model

- Reduced medical specialties to 18
- Support Vector Machine
- GridSearchCV to find best parameters
  - Best parameters:  $C = 1$ , kernel = linear
- Accuracy 23%

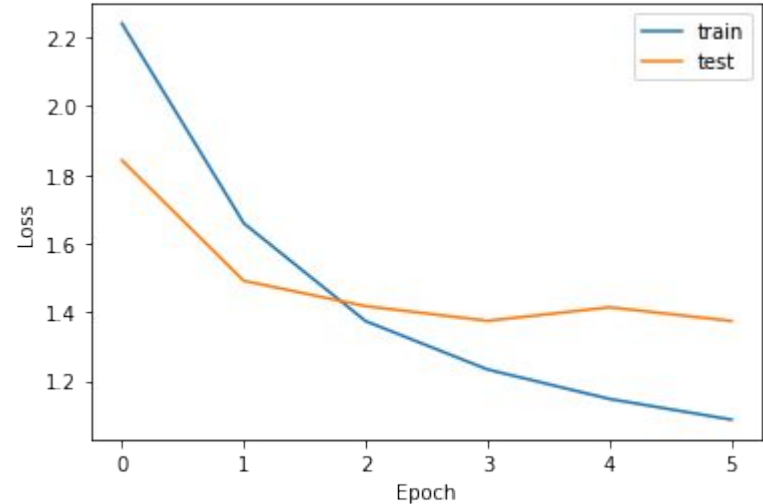


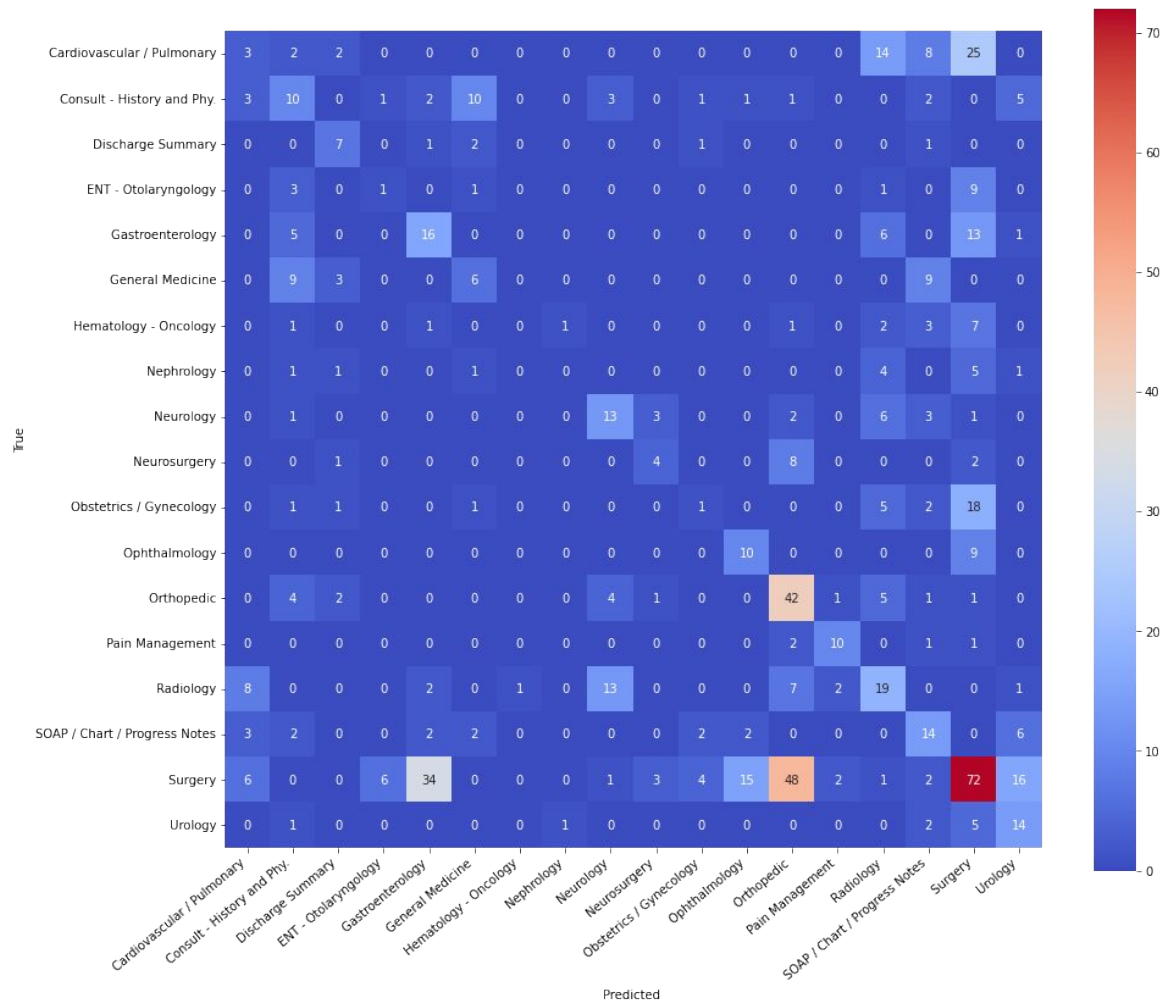


Accuracy: 0.2332859174964438  
 Precision: 0.20922610493758473  
 Recall: 0.2332859174964438  
 F1 score: 0.21742526566778728

# RoBERTa

- Reduced medical specialties to 18
- Model limited max sequence length
- Just transcriptions
- Accuracy: 48.06%
- Loss: 1.0876
- Validation Accuracy: 34.42%
- Validation Loss: 1.3746

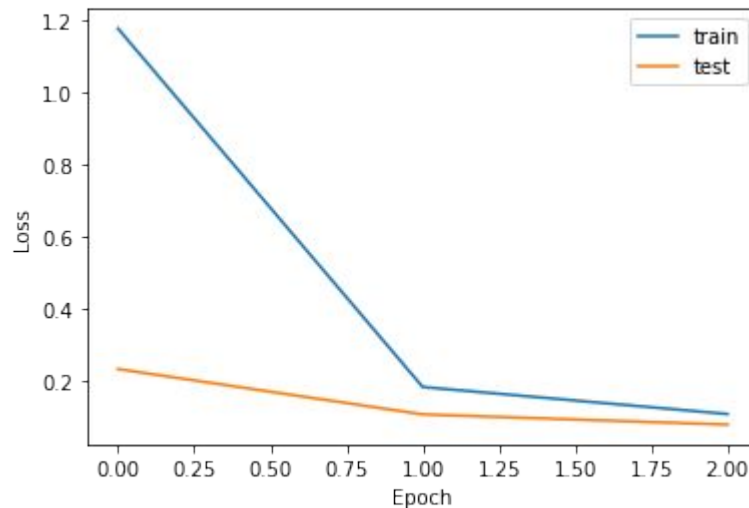


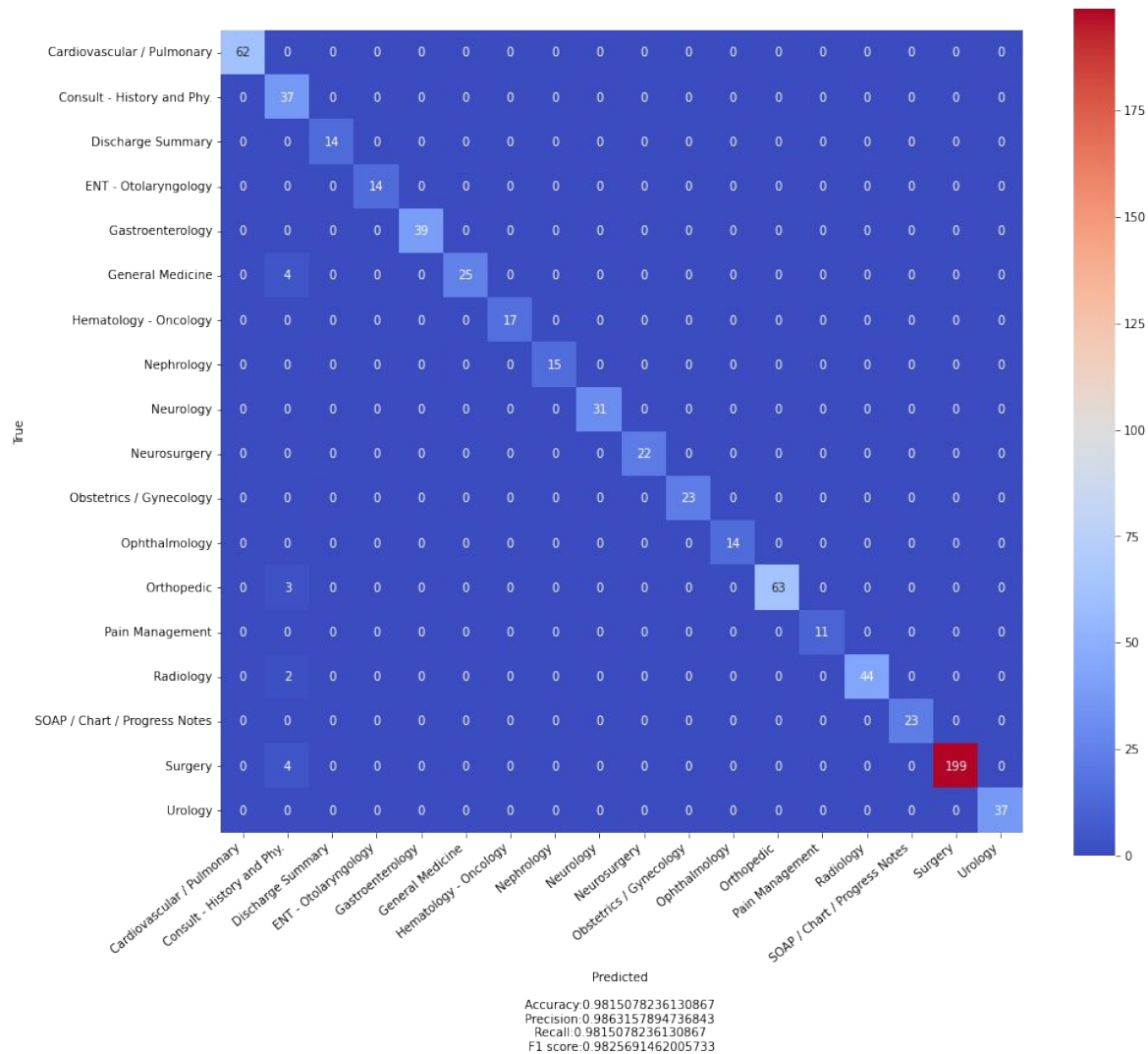


Accuracy: 0.3442389758179232  
Precision: 0.31842947125638177  
Recall: 0.3442389758179232  
F1 score: 0.3199623527339034

# RoBERTa

- Keywords
- Accuracy: 98.29%
- Loss: .1126
- Validation Accuracy: 98.58%
- Validation Loss: .0667





# Conclusions

- Pre-trained model outperformed the baseline model
- Overlap between specialties seems to have increased errors
- Possible Improvements
  - Different Model
  - Better data cleaning
  - Classify transcripts differently

# Sources

- [Dataset](#)
- [Hugging Face Guide](#)
- [Text Cleaning](#)
- Data Exploration: [link 1](#), [link 2](#)