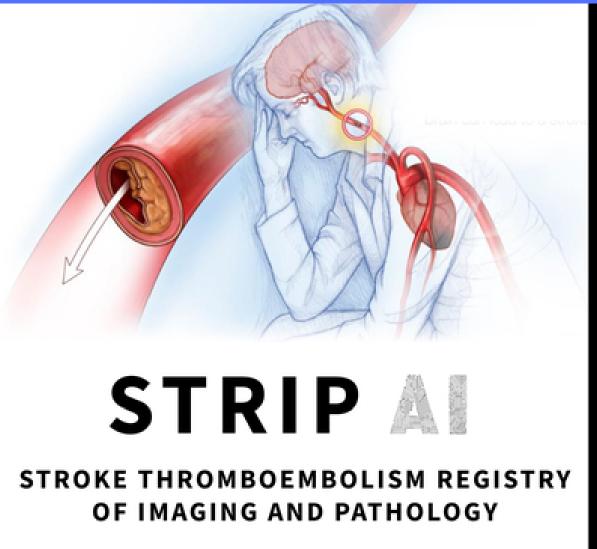
MAYO CLINIC

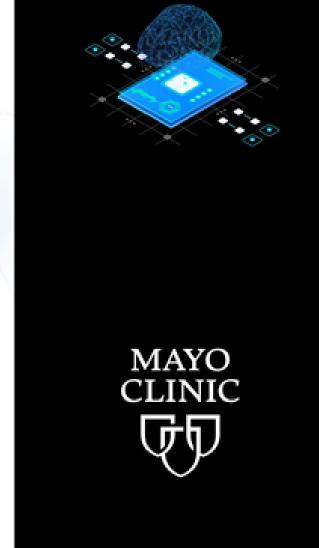


PREDICTING STROKE ETIOLOGY FROM IMAGING OF BLOOD CLOTS

SAM OLIVER - FREELANCE DATA SCIENTIST







- BACKGROUND
- DATA

- METHODS
- MODEL RESULTS
- THE DIAGNOSTIC TOOL

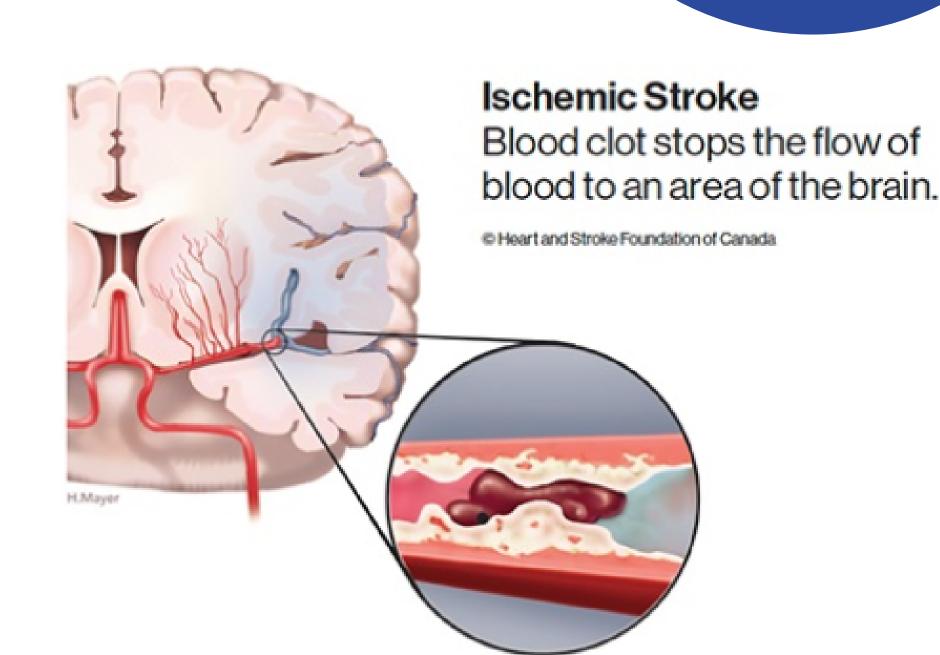
- FURTHER
 RECOMMENDATIONS
- CONTACT DETAILS

BACKGROUND

- 700K STROKES / YR IN US
- MAYO CLINIC + KAGGLE

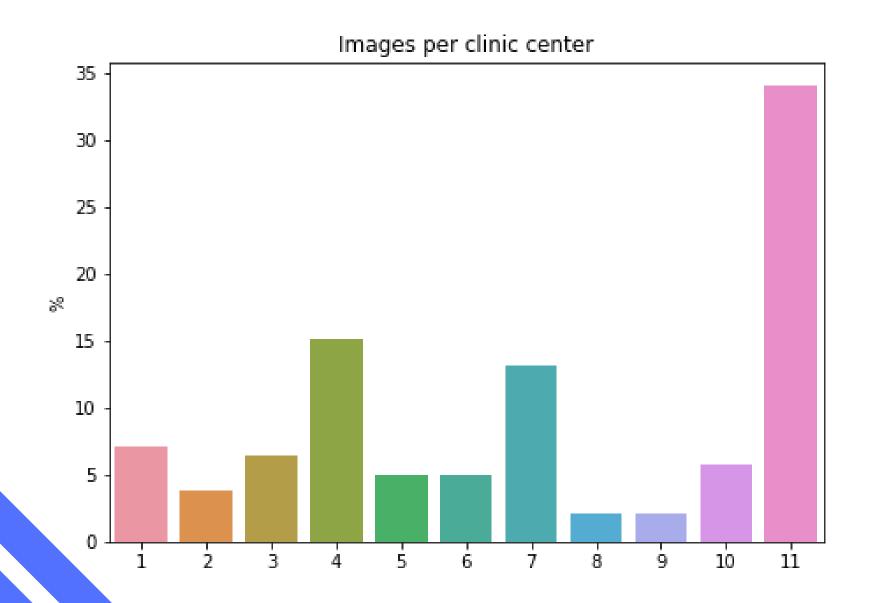
GOALS

- PREDICT TYPE OF STROKE
 FROM EXTRACTED CLOT IMAGE
- CREATE DIAGNOSTIC TOOL TO ASSIST THE MAYO CLINIC WITH DIAGNOSIS OF PATIENTS

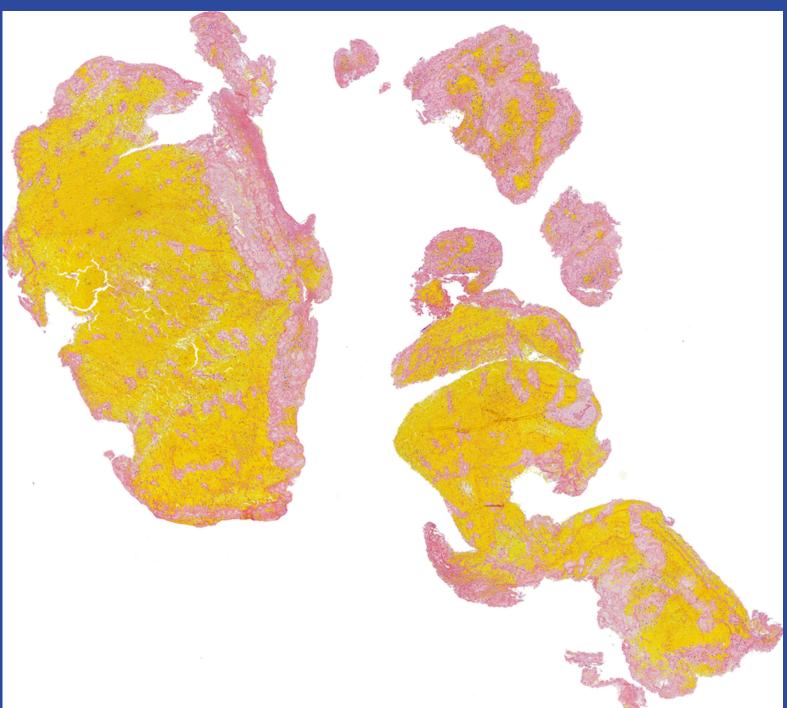


DATA

1,000 WHOLE SLIDE IMAGES - 400 GB



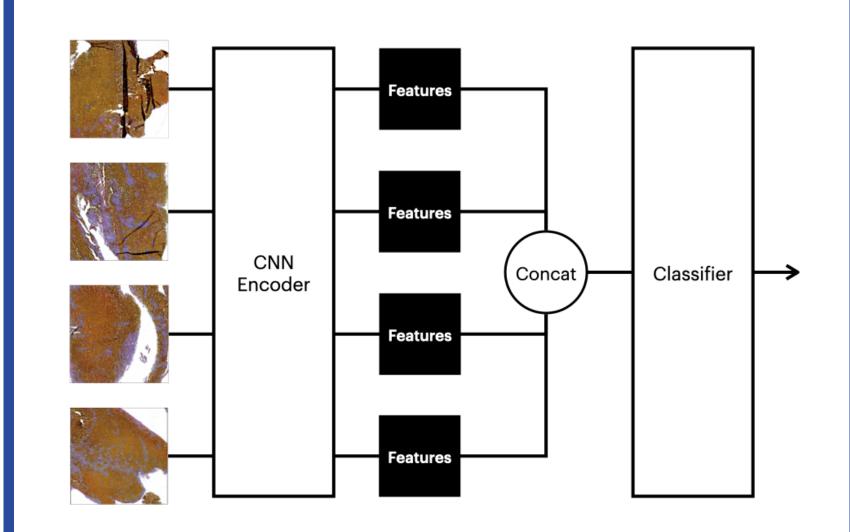
WHOLE SLIDE IMAGE OF CLOT



METHODS

- CLEAN DATASET
- HANDLE SIZE OF DATA
- MODEL THE DATA: NN & MIL
- EVALUATE & COMPARE

MIL MODELING



MODEL RESULTS

LOSS SCORES:

• IMPROVEMENT FROM BASELINE TO BEST

BASELINE	BEST
0.663	0.604



THE DIAGNOSTIC TOOL

1 CREATES DIAGNOSTIC PREDICTION
Automatic diagnosis given

2 READS IN NEW WHOLE SLIDE IMAGES
Processes data and refines algorithm

OPTIMIZED FOR ADDITIONAL DATA

Processes biomarkers and mass spectrometry readings



Further Recommendations

1 EXPAND COMPUTATIONAL RESOURCES

2 MORE DATA: BIOMARKERS + MASS SPECTROMETRY READINGS

2 EXPLORE OTHER SOLUTIONS:
PREVENTATIVE HEALTHCARE
PREDICT RISK + RECOMMEND SCREENING

CONTACT INFO

- samueloliver303@gmail.com
- https://www.linkedin.com/in/samo
 liver3/
- https://github.com/samoliver3

