



Prediction of Age and Biological Sex using Spatial-Temporal Graph CNN on fMRI Scans

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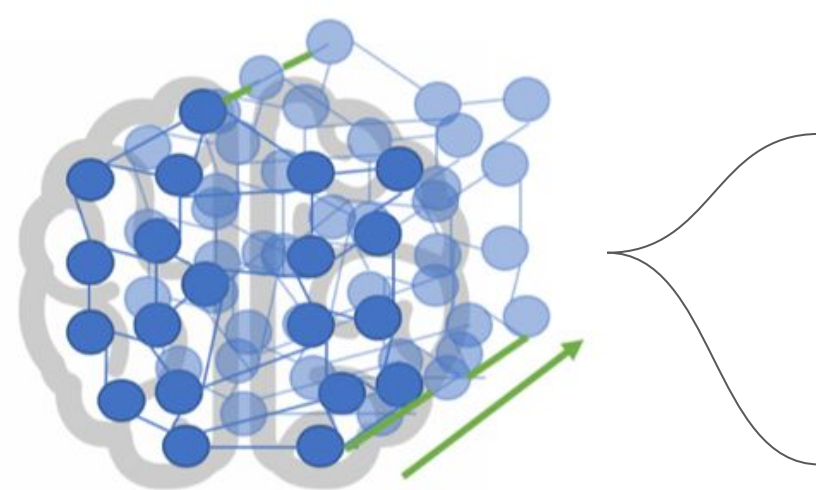
CS 230: Deep Learning
 Stanford University, Fall 2019

Introduction

- The **Human Connectome Project** aims to build a complete functional and structural map of the human brain
- We use their fMRI and structural MRI brain scans to **predict sex and age**
- We use a novel approach by employing **ST-GCN**
- This research project can be used to **discover** and **improve understanding** of sex-linked and age-related neurological diseases

Problem

- Given a **4D brain scan**, classify age and sex of a patient
- Regression on age, binary classification of age (old, young adult) and sex (male, female)



Age

Sex

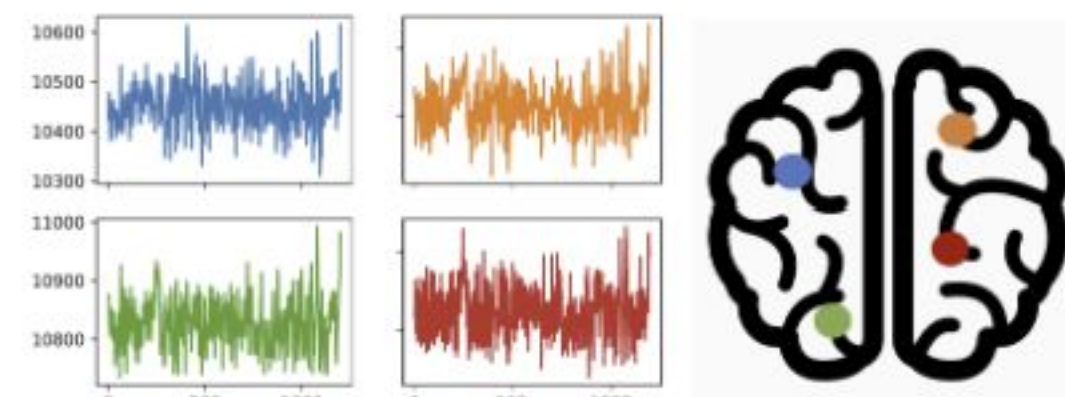
4D Brain Scan

- ST-GCN Test Accuracies**
 - Age : **63%** , Sex: **78%**

Dataset

- Original input data
 - fMRI scans from **1108 healthy individuals**, 603 female, 505 male
 - Each file is of 200 brain regions of 1,200 timesteps
- Data augmentation
 - Each file was **fragmented** into 100 time steps, which resulted in **12 fragments**
 - Coarser brain sampling (**22 brain regions** instead of 200)
 - This resulted in **13,082 samples**

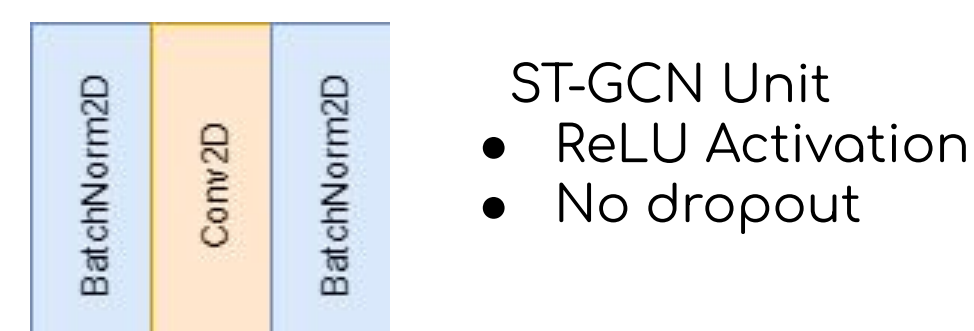
Feature Mapping



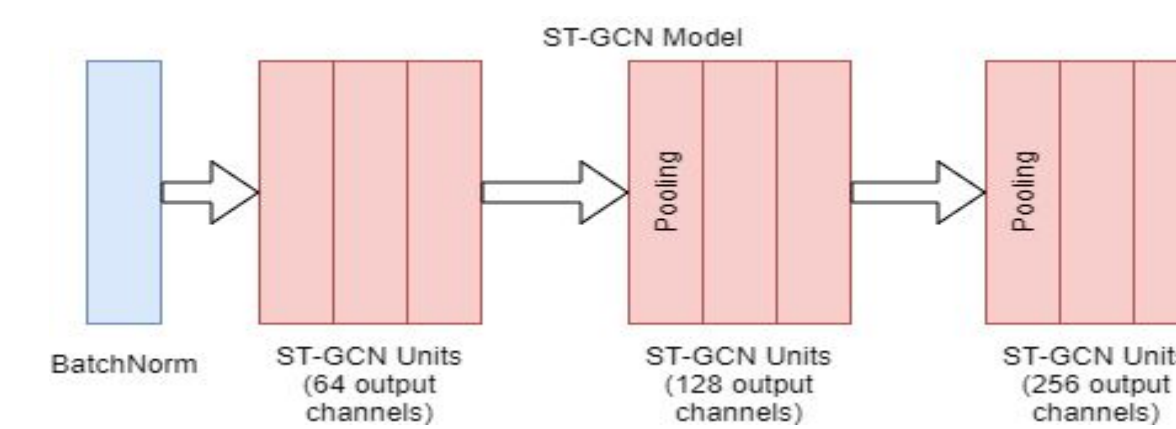
- Brain regions act as **nodes** in the graph. Correlation between regions act as **edges**
- Adjacency matrix** captures **correlations** of brain region **across timesteps**
- Feature mapping is computed by using

$$f_{out} = D^{-1/2}(A + I)D^{-1/2}f_{in}W$$

ST-GCN Unit and Model



- ST-GCN Unit
 - ReLU Activation
 - No dropout

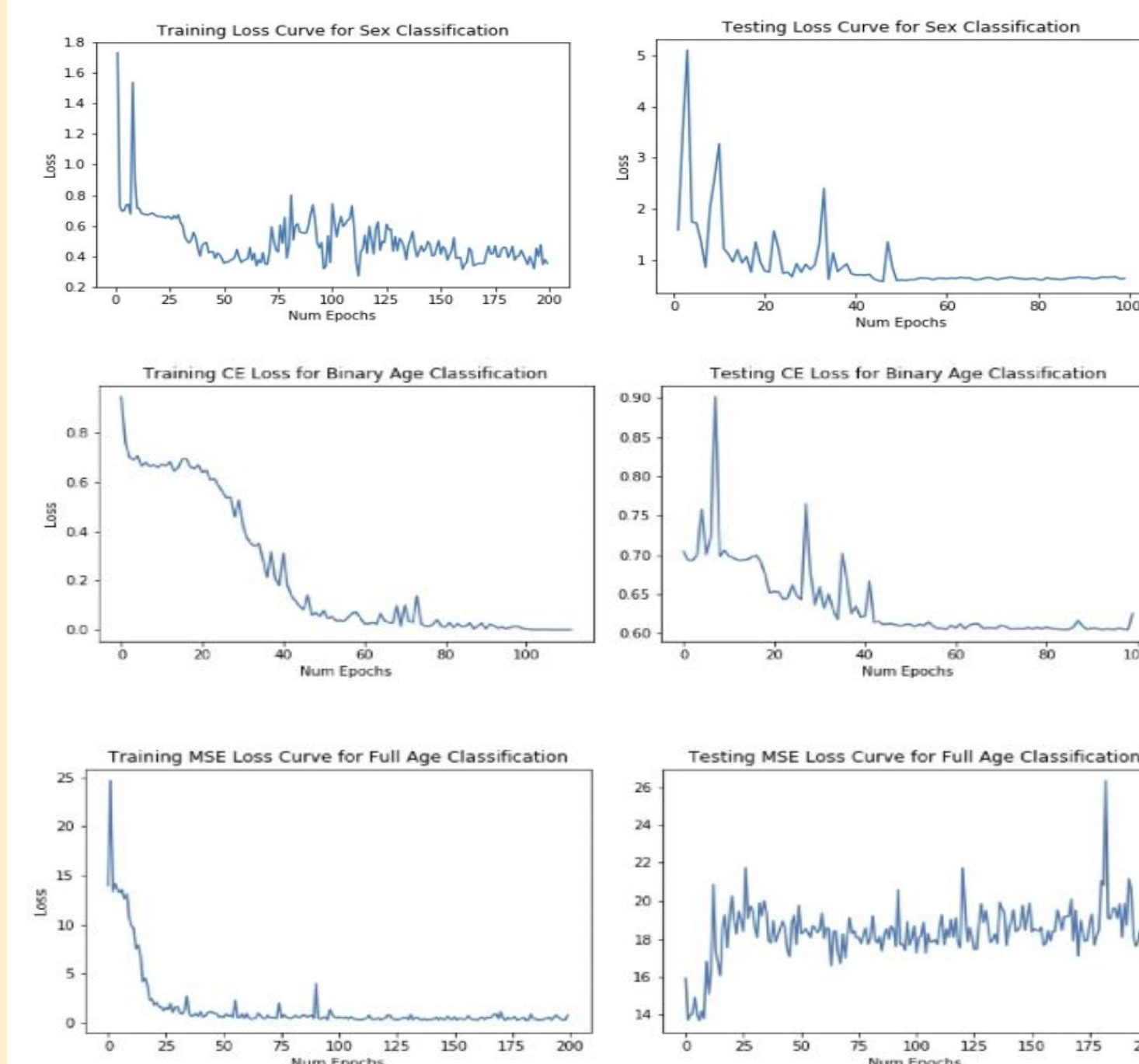


- Final layer predictions computed with and without Softmax for **binary** and **real-number** outputs

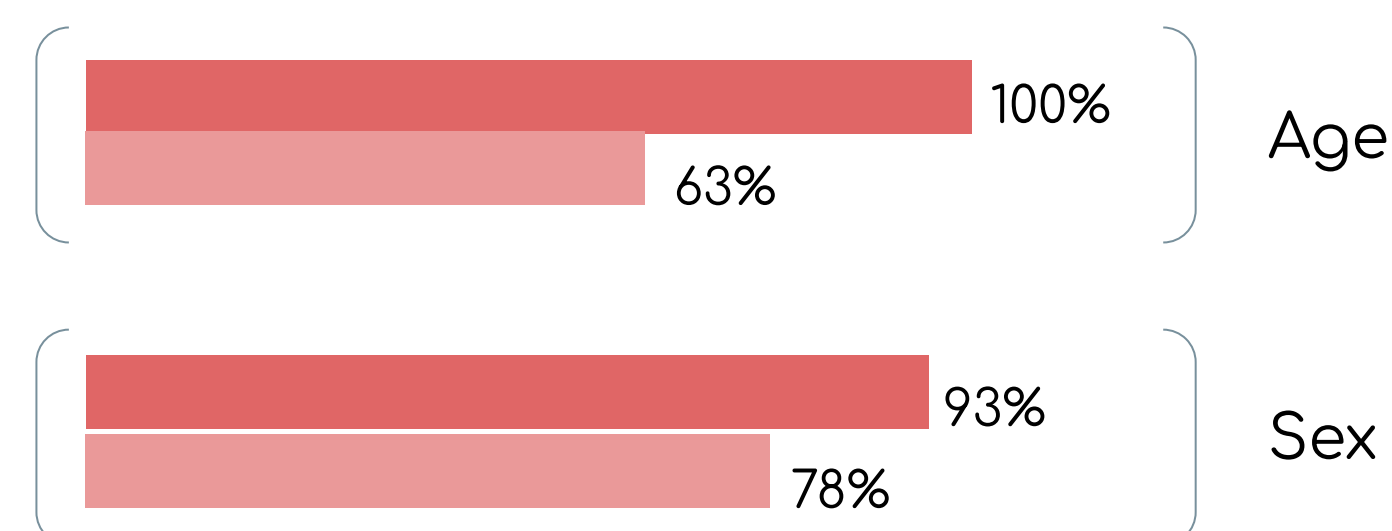
$$\mathcal{L}(y_i, \hat{y}_i) = -\frac{1}{m} \sum_{i=1}^m y_i \log(\hat{y}_i) + (1 - y_i) \log(1 - \hat{y}_i) \quad \text{CE Loss}$$

$$\mathcal{L}(y_i, \hat{y}_i) = \frac{1}{m} \sum_{i=1}^m (y_i - \hat{y}_i)^2 \quad \text{MSE Loss}$$

Results

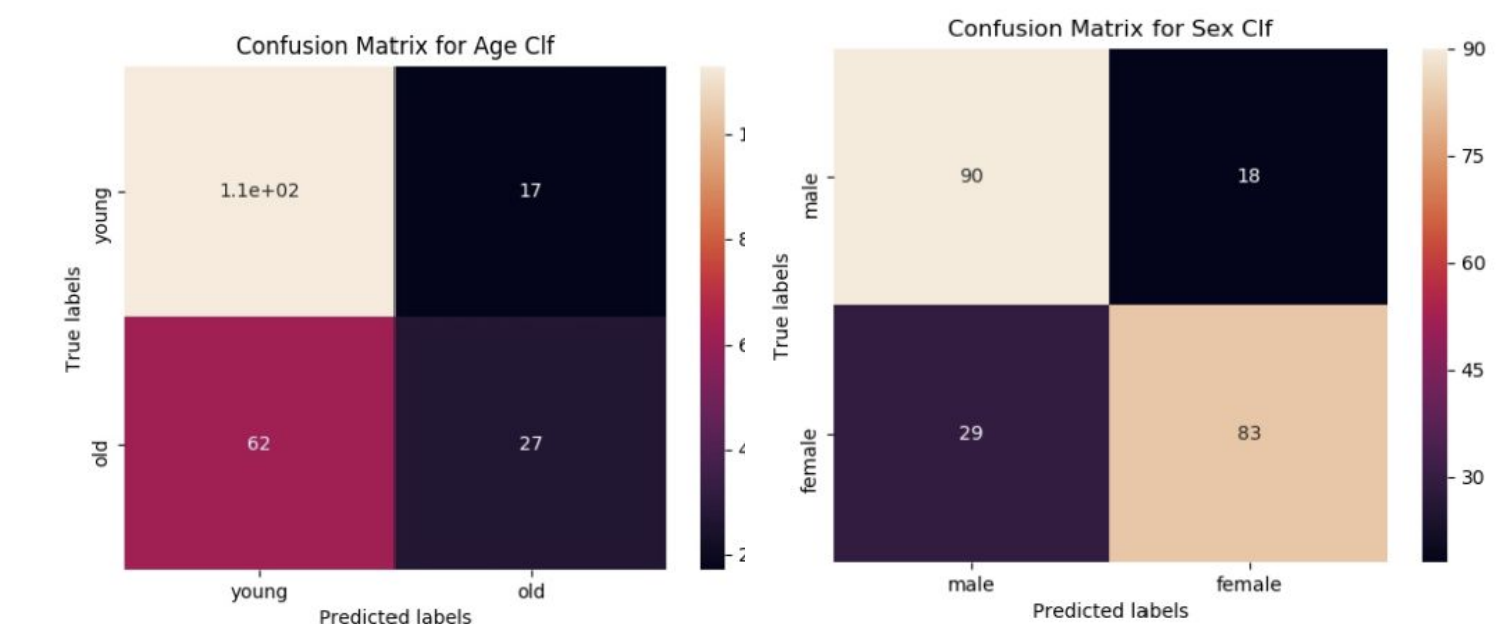


Accuracies (Train/Test)



Discussion

	Precision	Recall	F1 Score
Age	0.6457	0.8692	0.741
Sex	0.7563	0.833	0.793



- Model show similar accuracies for male and female brains but predicted male more often
- Age classification accuracy lower, but model was still able to distinguish age bins

Future Work

- Large gap in train/test accuracies suggest we use dropout and weight regularization
- Analyze subject level prediction instead of sample level prediction for better accuracies

Notable References

For full list please checkout github.com/ericksiavichay/cs230-final-project

- Kipf, T.N., Welling, M.: Semi-Supervised Classification with Graph Convolutional Network <https://arxiv.org/abs/1609.02907>
- Li, X., et al: Graph Neural Network for Interpreting Task-fMRI Biomarkers <https://arxiv.org/abs/1907.01661>
- Yan, S., Xiong, Y., Lin, D.: Spatial Temporal Graph Convolutional Networks (ST-GCN) for Skeleton-Based Action Recognition in PyTorch <https://github.com/yysijie/st-gcn>