Feature Engineering and Missing Value Imputation for Classification of Depression with Deep Neural Networks

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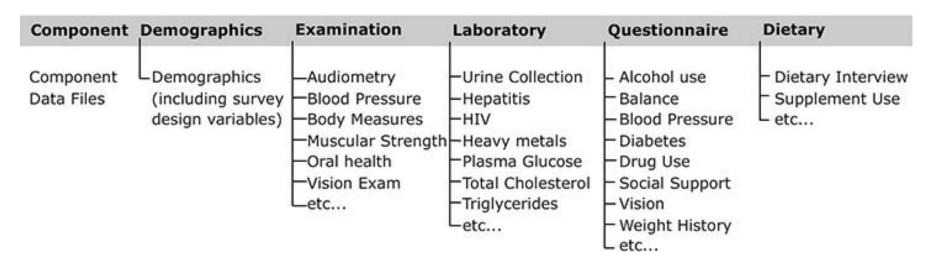
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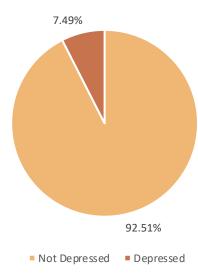
Background

- 1 in every 6 adults in the US will experience depression during their lifetime, which is about 16 million adults each year
- Depression is characterized as a long-term affect in mood caused by a complex combination of genetic, physiological, environmental, and social factors.
- Depression frequently goes undiagnosed and untreated, contributing to the overall burden of disease

National Health & Nutrition Examination Survey (NHANES)



- Sample of the US population amounting to about 5,000 participants each two-year period
- Patient Health Questionnaire nine-item (PHQ-9) depression screening instrument assesses the frequency of depression symptoms over the past 2 weeks
 - Response categories include "not at all," "several days," "more than half the days," and
 "nearly every day" given a point ranging from 0 to 3. A total score of 10 or higher marked as
 'depressed'
- 31,357 complete depression observations between 2005-2016
- 7.49% of the observations are classified as "depressed"



Features

| Survey Component | Number of Features | Binary | Multi-class | Continuous |
|---------------------|-----------------------|--------|-------------|------------|
| Demographics | 7 | 1 | 2 | 4 |
| Dietary | 16 | 1 | 0 | 15 |
| Examination | 7 | 0 | 1 | 6 |
| Laboratory | 15 | 2 | 0 | 13 |
| Questionnaire | 67 | 43 | 14 | 10 |
| Total | 112 | 47 | 17 | 48 |

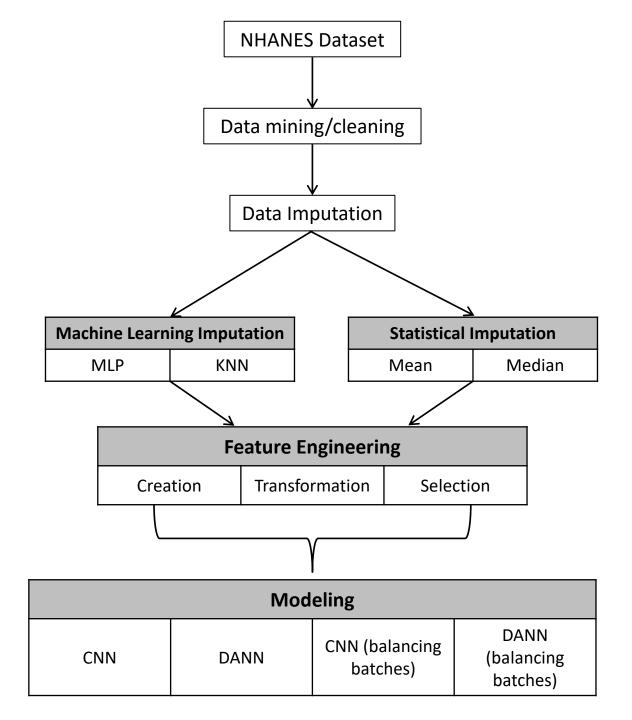
- Originally only 53 NHANES features imported (used in preliminary analyses)
- Features were chosen based on prior associations found in the literature
- Continuous features are utilized for transformations in feature engineering

Problem Statement

Depression is a widely undiagnosed mental health condition that greatly impacts the burden of disease. NHANES data is highly imbalanced (7.49% positive) and contains many missing values. The primary goal of this project is to identify the best imputation and feature engineering methods to optimize depression classification ability of deep neural networks.

Literature Review

- Linear associations with depression:
 - Demographics such as age, race, gender, and income level [1]
 - Health characteristics including physical activity [2], metabolic syndrome, BMI [3], and other chronic conditions such as asthma, kidney disease [4], and osteoporosis
 - Dietary practices including caffeine consumption [5], household food insecurities [6], and Vitamin B levels [7]
- Jerez et al. 2020 [8] used **imputation methods** based on statistical techniques (mean, hot-deck and multiple imputation) and machine learning techniques (multi-layer perceptron (MLP), self-organization maps (SOM) and k-nearest neighbor (KNN))
 - Machine learning imputation performed best for ANN classification of breast cancer
- Heaton et. al. 2016 [9] reviewed feature engineering techniques for machine learning and found a DANN able to handle addition, summation and multiplication of features. Additionally, feature counts, differences, power transformations, and rational polynomial transformations were all synthesized by the DANN relatively easily
- ANNs have demonstrated accurate classifications of many diseases including heart disease [10], caries [11], and osteoporosis [12, 13]
- Shallow CNN to predict Coronary Heart Disease with NHANES data [14]
- Support vector machines (SVM) and ensemble modeling to predict risk of Diabetes with NHANES data [15]
- Repeated random sub-sampling & Random Forests (RF) to predict the risk of eight chronic diseases [16]



Methods

Data Mining & Cleaning

- Data was downloaded and converted to CSV files using python script found on GitHub [17]
- Remaining preprocessing including importing files, extracting and renaming columns, and merging data for each year was done in Python Jupyter Notebook
- The target column of depression was calculated using the sum of the Patient Health Questionnaire (PHQ-9) nine-item survey ranging from 0 to 27, with a score of 10 or higher marked as 'depressed'. Only data with complete PHQ-9 questionnaires were used in this project
- Responses of '1 yes' and '2 no' were converted to binary 1/0, respectively
- All responses of '7 don't know' or '9 refused' were marked as N/A
- Prior to the implementation of any algorithm predicting depression, all multiclass features are encoded, and continuous features are standardized to a mean of 0 and standard deviation of 1

Data Imputation

- Statistical methods:
 - Simple mean, median, and mode missing value imputation
 - Implemented after dropping columns with a proportion of missing values exceeding a given input threshold (0.75)
- Machine Learning methods:
 - Aim to use available information within the dataset to estimate and substitute the missing values
 - First, separated the data frame into complete data and incomplete data
 - Complete data (excluding the ultimate target variable 'depressed') was used to predict values of the next-mostcomplete column. These predicted values were then imputed into that column and used with the complete data to predict the missing values of the subsequent column
 - Thus this method progressively predicts missing values with the imputed data
 - Used both Multi-layer perceptron (MLP) k-nearest neighbors (KNN) – regression or classification depending on target column

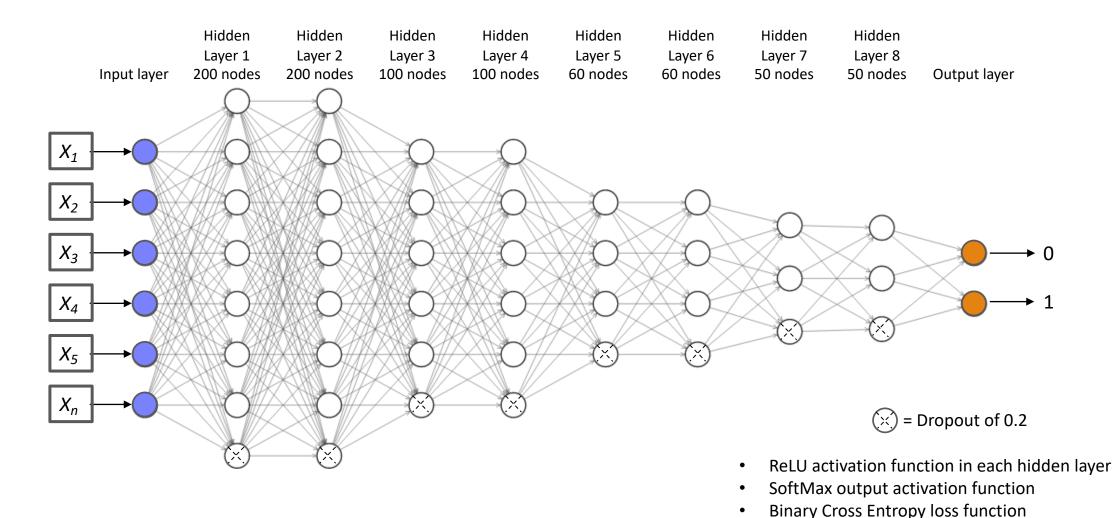
Preliminary Analyses: Results

- Original 53 columns of NHANES data
- With such low F1-scores, and accuracies averaging at 92%, the challenge with imbalanced data is highlighted in these results
- These results lead to the addition of features and construction of deep neural networks (DNNs)

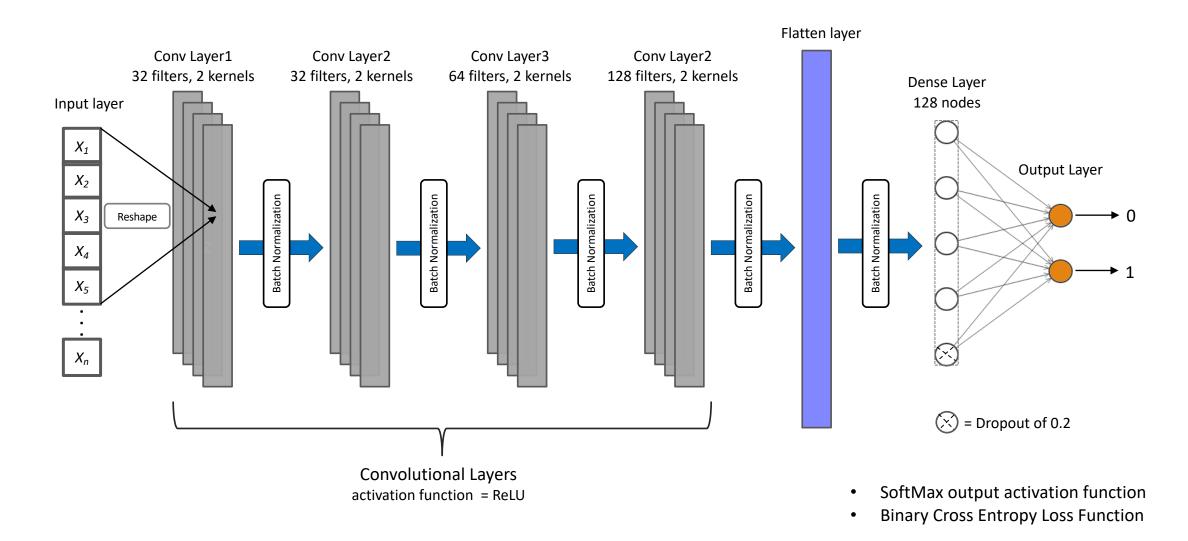
F1-SCORES

| Imputation Type | Logistic Regression | Decision Tree | Random Forest | XGBoost | KNN | Naïve Bayes | Single Perceptron | MLP |
|--------------------|------------------------|------------------|------------------|---------|--------|----------------|----------------------|--------|
| MLP | 0.1575 | 0.0129 | 0.0129 | 0.1839 | 0.0946 | 0.2465 | 0.2822 | 0.1489 |
| KNN | 0.1430 | 0.0077 | 0.0077 | 0.1630 | 0.0943 | 0.2459 | 0.0303 | 0.1675 |
| Mean | 0.1398 | 0.0154 | 0.0154 | 0.1833 | 0.0953 | 0.2473 | 0.1171 | 0.2346 |
| Median | 0.1398 | 0.0154 | 0.0154 | 0.1833 | 0.0953 | 0.2473 | 0.1171 | 0.0832 |

Deep Neural Network



1-D Convolutional Neural Network



Verification of Neural Networks

- Implementation of the neural networks to classify:
 - Different target variable within NHANES dataset 'overweight'
 - Classification of 'diagnosis' with UCI Breast Cancer dataset

| | | | F1-SC | ORES |
|--------------------------------|-----------------|------------------|--------|--------|
| Data | Target Variable | % positive cases | DANN | CNN |
| UCI Wisconsin Breast Cancer | diagnosis | 37.26 | 0.8859 | 0.9385 |
| NHANES MLP imputed | overweight | 33.75 | 0.7143 | 0.7538 |

- Breast Cancer high F1-Scores confirm the architectures and implementations of the DNNs
- NHANES target of 'overweight' F1-Scores are not as impressive
 - Features in the NHANES dataset have less predictive power than those in the Breast Cancer dataset
- Nevertheless, classification of 'overweight' result in much higher F1-scores than in the case of depression
 - Difference in class imbalance proportions (33.75% vs. 7.49%)
 - The nature of the two variables, 'overweight' having more recognizable attributes
- Conclusions:
 - DNNs are implemented and functioning properly
 - The high level of class imbalance with respect to depression is a significant hindrance
 - The NHANES data may, by nature, not be highly discriminatory or indicative for any classification

Feature Engineering



Creation

Addition of feature columns by either counting, summing, subtracting, dividing, or multiplying two or more existing features

15 new features created



Transformation

New features consisting of the square root, log, power, polynomial, and sine functions applied to the original continuous numerical features

Continual augmentation of transformed features to existing features



Selection

Sub-setting of features

Feature association with depression (literature review)

Reduce correlation between features

| Feature Engineering Procedure | Number of Features (after encoding) |
|----------------------------------------------------------------------------------------|-------------------------------------|
| NHANES original | 79 |
| NHANES baseline features | 166 |
| Baseline + Created features (BCF) | 186 |
| BCF + log(cont) | 240 |
| BCF + (cont) ^{1/2} | 240 |
| BCF + (cont) ² | 240 |
| BCF + polynomial | 240 |
| BCF + sin(cont) | 240 |
| BCF + (cont) ² + log(cont) | 293 |
| $BCF + (cont)^{2} + log(cont) + (cont)^{1/2}$ | 346 |
| BCF + (cont) ² + log(cont) + (cont) ^{1/2} + polynomial | 399 |
| BCF + (cont) ² + log(cont) + (cont) ^{1/2} + polynomial + sin(cont) | 452 |
| Subset of CF | 105 |
| Subset + log(cont) | 135 |
| Subset + log(cont) + (cont) 1/2 | 165 |



Balancing Batches (BB)

- For each data imputation method and feature engineering experiment, both the DANN and CNN were trained with standard training data (imbalanced) and with balanced minibatches
- Balanced mini-batches used random under sampling of the majority class without replacement to create a random, balanced batches of size 32
- Implemented to determine whether resampling methods are beneficial in these DNN classifications

Evaluation Metrics

Results are evaluated using F1-Score:

$$2 * \frac{Precision*Recall}{Precision+Recall}$$

Where,

$$Precision = \frac{TP}{TP + FP}$$

$$Recall = \frac{TP}{TP + FN}$$

| | Predicted: 0 | Predicted: 1 |
|-----------|--------------|--------------|
| Actual: 0 | TN | FP |
| Actual: 1 | FN | TP |

MODEL AVERAGE F1-SCORES

| DANN | 0.1458 |
|---------|--------|
| CNN | 0.2640 |
| DANN BB | 0.2513 |
| CNN BB | 0.2488 |

Results: Model Comparison

- Average across all feature engineering techniques
- CNN has highest average

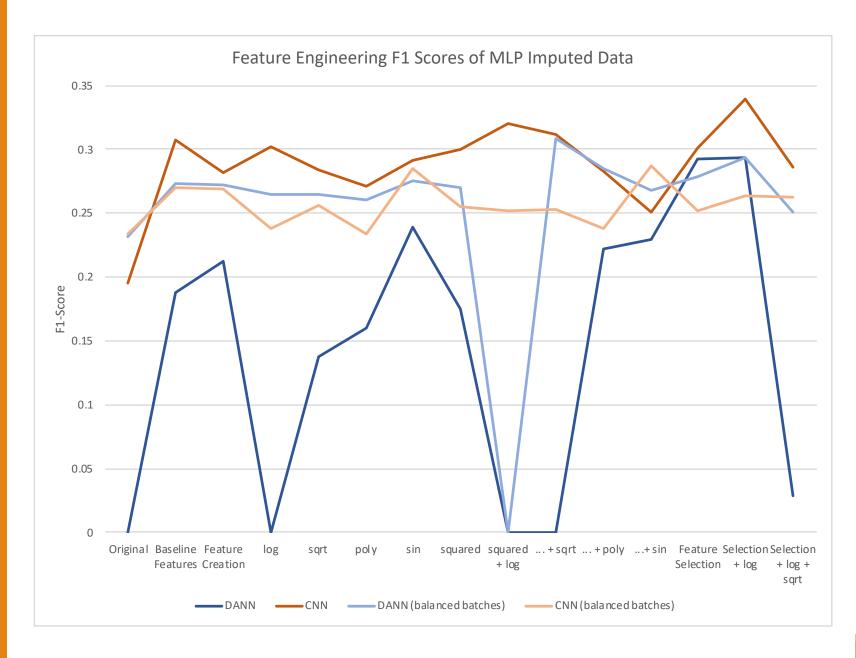
| IMPUTATION METHOD | AVERAGE F1- SCORE (ALL MODELS) | CNN AVERAGE F1- SCORE |
|----------------------|--------------------------------------|--------------------------|
| MLP | 0.2312 | 0.2794 |
| KNN | 0.2429 | 0.2726 |
| Mean | 0.2083 | 0.2521 |
| Median | 0.2274 | 0.2520 |

Results: Imputation Methods

- Results averaged across all feature engineering results
- KNN provides the highest average F1score averaging across all models
- Average F1-score of just the CNN results, MLP imputation has the best F1-score

Results: Feature Engineering

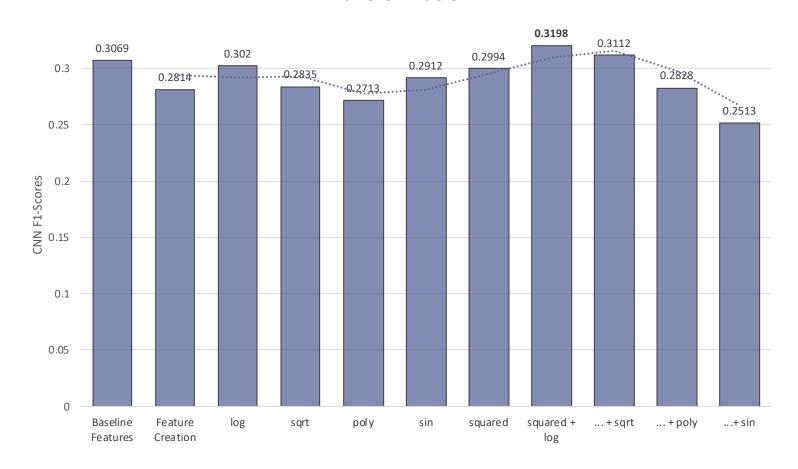
- DANN has the greatest fluctuations and is the least reliable model
- DANN is generally improved by training on balanced batches.
- All models benefit from the addition of features from 'original' to 'baseline features'
- Created features may have no benefit
- Transformations have varying effects
- Most models benefit from some feature selection and subsequent transformations



CNN and MLP Imputation

- Feature creation results in a decrease in the F1-score
- Log, sin, and squared transformations have better outcomes than polynomial and square root
- The best transformation and augmentation combination is the addition of both the square and log, resulting in a higher F1-score than the baseline (0.3198)
- Subsequent augmentations negatively affect performance.

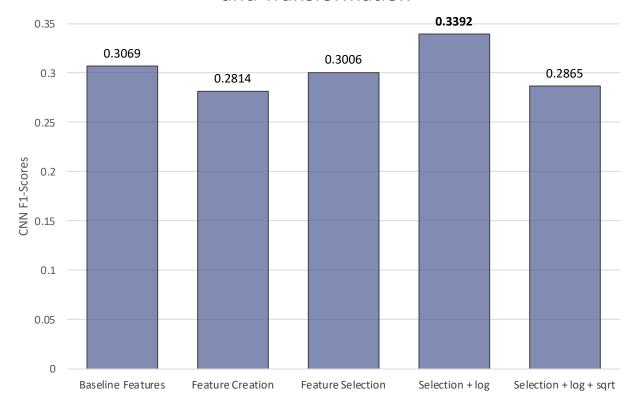
MLP Imputation CNN F1-scores: Feature Creation and Transformation



CNN and MLP Imputation Cont.

- Feature selection alone leads to improvement from feature creation
- The addition of the logtransformed features to this subset provides the best results (0.3392) among all F1-scores of each imputation, model, and feature engineering combination.

MLP Imputation CNN F1-scores: Feature Selection and Transformation



Best Results

This model is achieved using the CNN without balancing batches, MLP imputation, and a subset of the extended number of NHANES features along with the 15 created features, and the log of continuous features

This F1-Score is an 11% increase from that without any feature engineering (0.3069)

| | Predicted: 0 | Predicted: 1 | |
|-----------|--------------|--------------|--|
| Actual: 0 | 9036 | 545 | |
| Actual: 1 | 499 | 268 | |
| ' | | | |
| F1-Score | 0.3392 | | |
| Precision | 0.3296 | | |
| Recall | 0.3494 | | |
| ROC AUC | 0.6463 | | |
| Accuracy | 0.8991 | | |

Summary & Conclusions

- Depression is a complex mental disease with unspecified causes. Despite the unique combination of demographic, dietary, clinical, and questionnaire data from NHANES, I am unable to build a model to accurately classify depression
- The nature of this data with only 7.49% positive cases and many missing values did not provide enough information for DNNs to predict depression in the NHANES population.
- Progressive machine learning imputation out-performed standard statistical imputation techniques.
- The 1D CNN classification results were superior to the more commonly used machine learning algorithms
- Feature engineering techniques including feature selection and augmentation of transformed features afforded slight improvements to each model's performance.
- A combination of feature creation, selection, and transformation boosted the F1-score of the baseline model by 11%
- Future work could benefit from implementation of MLP progressive imputation, 1D CNNs for binary classifications, and experimenting with feature engineering

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Code available at: https://github.com/csklaver/Capstone-Group6

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