


Cardiovascular Disease Prediction Exploratory Data Analysis



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DSCI 303



Overview

- Cardiovascular diseases (CVD's) have been the #1 cause of death globally for the past few years. According to the World Health Org, close to 18 million people die annually from CVD's.
- That being said, many studies show that almost 80% of CVD's can indeed be prevented, including heart disease and stroke.
- Our Goal: Create a tool to help doctors with predicting a patient's chances of getting CVD's with high accuracy, so that they can be prescribed necessary treatments/ medication early-on.

Previous Research

Machine Learning–Driven Models to Predict Prognostic Outcomes in Patients Hospitalized With Heart Failure Using Electronic Health Records: Retrospective Study

- **Goal:** Predict 1-year in-hospital mortality, use of positive inotropic agents, and 1-year all-cause readmission rate
- **Method:** Decision tree of mortality risk (after consideration of logistic regression, support vector machine, artificial neural network, random forest, and extreme gradient boosting models)
- **Data:** real-world electronic health records

Using Deep Learning to Identify High-Risk Patients with Heart Failure with Reduced Ejection Fraction | Published in Journal of Health Economics and Outcomes Research

- **Goal:** Predict hospitalizations, worsening HF events, and 30-day and 90-day readmissions in patients with heart failure with reduced ejection fraction (HFrEF)
- **Data:** Adult HFrEF patients from IBM® MarketScan® Commercial and Medicare Supplement databases (2015-2017)
- **Method:** Sequential model architecture based on bi-directional long short-term memory (Bi-LSTM) layers (also tested traditional ML models such as logistic regression, random forest, and eXtreme Gradient Boosting (XGBoost))
- **Results:** For all outcomes assessed, the DL approach outperformed traditional machine learning models

Data Overview

Link to data: <https://www.kaggle.com/sulianova/cardiovascular-disease-dataset>

Snapshot of data:

12 Features

	id	age	gender	height	weight	ap_hi	ap_lo	cholesterol	gluc	smoke	alco	active	cardio
0	0	18393	2	168	62.0	110	80	1	1	0	0	1	0
1	1	20228	1	156	85.0	140	90	3	1	0	0	1	1
2	2	18857	1	165	64.0	130	70	3	1	0	0	0	1
3	3	17623	2	169	82.0	150	100	1	1	0	0	1	1
4	4	17474	1	156	56.0	100	60	1	1	0	0	0	0

Number of observations: 70000

Target (0=does not have CVD; 1= has CVD)

Feature Overview

```
<class 'pandas.core.frame.DataFrame'>
```

```
RangeIndex: 70000 entries, 0 to 69999
```

```
Data columns (total 13 columns):
```

#	Column	Non-Null Count	Dtype
0	id	70000 non-null	int64
1	age	70000 non-null	float64
2	gender	70000 non-null	int64
3	height	70000 non-null	int64
4	weight	70000 non-null	float64
5	ap_hi	70000 non-null	int64
6	ap_lo	70000 non-null	int64
7	cholesterol	70000 non-null	int64
8	gluc	70000 non-null	int64
9	smoke	70000 non-null	int64
10	alco	70000 non-null	int64
11	active	70000 non-null	int64
12	cardio	70000 non-null	int64

```
dtypes: float64(2), int64(11)
```

```
memory usage: 6.9 MB
```

```
None
```

Objective (measured) Features

- **Age** (in days) [NUMERIC]
- **Gender** (1=female, 2=male) [BINARY]
- **Height** (in cm) [NUMERIC]
- **Weight** (in kg) [NUMERIC]

Examination Features

- **Systolic blood pressure/ ap_hi** (mm of mercury - mmHg) [NUMERIC]
- **Diastolic blood pressure/ ap_lo** (mm of mercury - mmHg) [NUMERIC]
- **Cholesterol** (1=norm; 2=above norm; 3=well above norm) [TERNARY]
- **Glucose** (1=norm; 2=above norm; 3=well above norm) [TERNARY]

Subjective Features

- **Smoking** (0=does not smoke; 1=smokes) [BINARY]
- **Alcohol Intake** (0=does not drink; 1=frequent drinker) [BINARY]
- **Physical Activity** (0=not very active; 1=active) [BINARY]

Ideally, there shouldn't be too many subjective features. We may need to give slightly less weight to these features in our final model to decrease variance.

Observation Overview

Missing Values

```
id          0
age         0
gender      0
height      0
weight      0
ap_hi       0
ap_lo       0
cholesterol 0
gluc        0
smoke       0
alco        0
active      0
cardio      0
dtype: int64
```

No need to impute

Unique Elements

```
id          70000
age         8076
gender       2
height      109
weight      287
ap_hi       153
ap_lo       157
cholesterol  3
gluc         3
smoke        2
alco         2
active       2
cardio       2
dtype: int64
```

Binary and ternary variables
are as expected

Duplicate Rows

```
0          False
1          False
2          False
3          False
4          False
...
69995      False
69996      False
69997      False
69998      False
69999      False
Length: 70000, dtype: bool
Total Number of Duplicates: 0
```

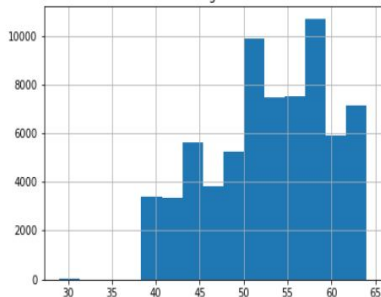
No need to remove tuples

Data is already clean!

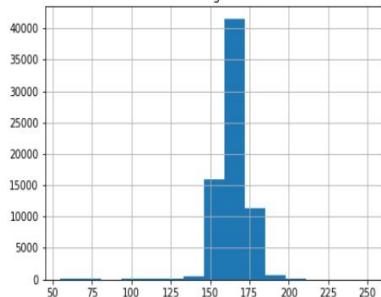
Numerical Feature Distributions

dtype=object)

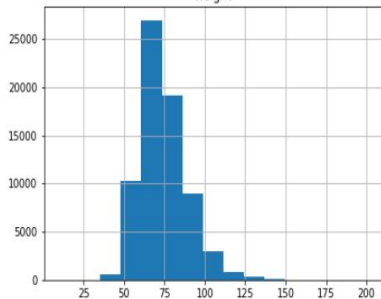
Age



Height



Weight

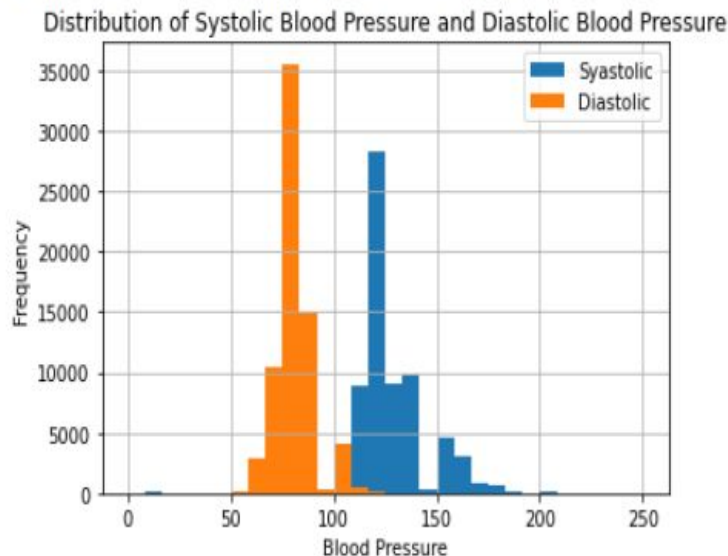


- Age is skewed to the right meaning we have mostly older participants in this dataset.
 - This indicates that our model will not be very applicable to a younger population.
- Height and weight appear to be normally distributed

Numerical Feature Distributions Cont.

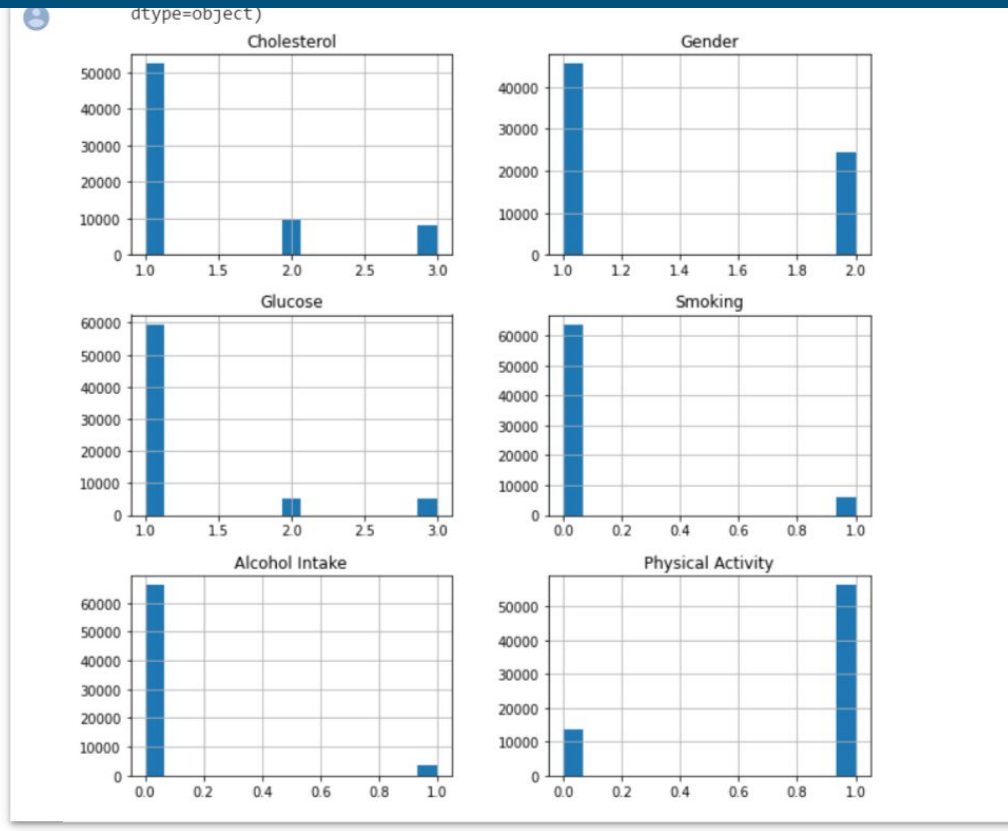
```
6 plt.legend()
```

```
<matplotlib.legend.Legend at 0x7fb60d27e090>
```



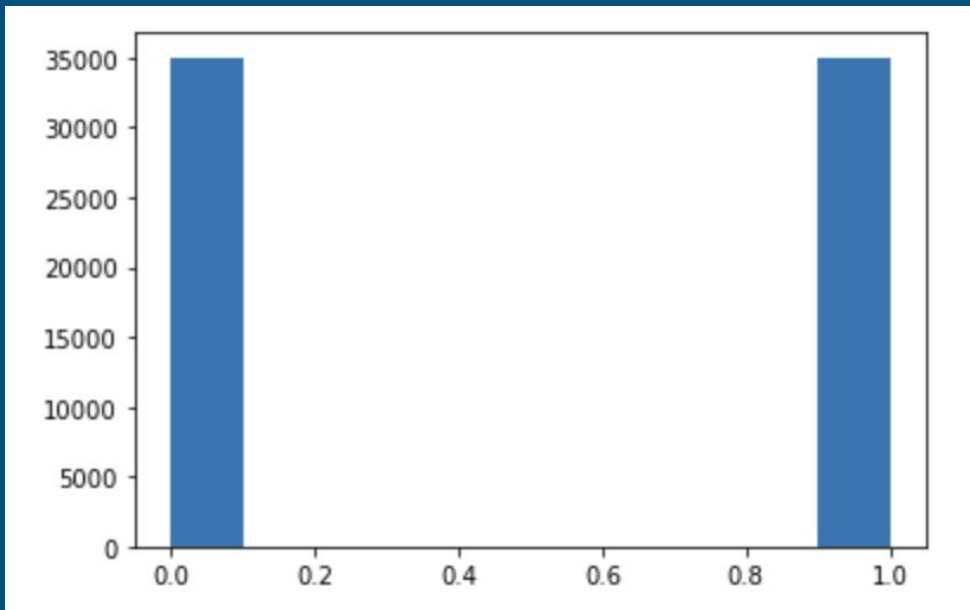
- Systolic Blood Pressure (pressure your heart exerts on your arteries when it pumps) and Diastolic Blood Pressure (pressure on your arteries between pumps of the heart) both appear to be normally distributed.

Categorical Feature Distributions



- Gender shows high bias because our sample deviates from the true distribution of gender in the population.
- Other variables such as cholesterol, glucose, smoking, alcohol intake, and physical activity are also significantly unbalanced.

Target Distribution



- Exactly equal number of observations for patients with cardiovascular disease and without cardiovascular disease.
- Confusion matrix will be a good way for us to measure our model performance.

Feature Correlation

Feature Correlation Matrix



Feature Correlation Continued

Most Correlated Features

- 1) Height and gender: .50
- 2) Cholesterol and glucose: .45
- 3) Alcohol and smoking: .34

Don't have multicollinearity since none of these values exceed .50 (usually considered minimum threshold for high dependence).

Most Correlated with Target

- 1) Age: .24
- 2) Cholesterol: .22
- 3) Weight: .18

Not very linearly correlated (will definitely need to combine multiple features in our models)

Questions?
