

# Diabetes Factors Analysis

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SDSC2102 STATISTICAL METHOD AND DATA ANALYSIS

# Background and problem formulation

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## What is diabetes

- Diabetes mellitus (DM)
- Group of metabolic disorders characterized by a high blood sugar level over a prolonged period

## Goal

- Assess the relationship between diabetic and its factors
- Finding out the which risk factors are more related to having diabetes
- Significance of that factor causing diabetes

# Data processing method and Justification

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## LOGISTIC REGRESSION

- It is a non-linear model to predict binary class
- Use to predict the odds of occurrence
- Assumes that class attributes is linear in the coefficients of the predictive attributes

We use classification model instead of regression model because:

- Having diabetes or not is a binary variable
- 1 means the person has diabetes
- 0 means the person does not has diabetes
- If we use regression model instead of a classification model
  - Accuracy will be relatively lower than classification model

# Data selected

- Source: Centers for Disease Control and Prevention of U.S. Department of Health & Human Services
- Duration: 2017-March 2020 Pre-Pandemic
- Survey Type:
  - Questionnaire Data
  - Examination Data

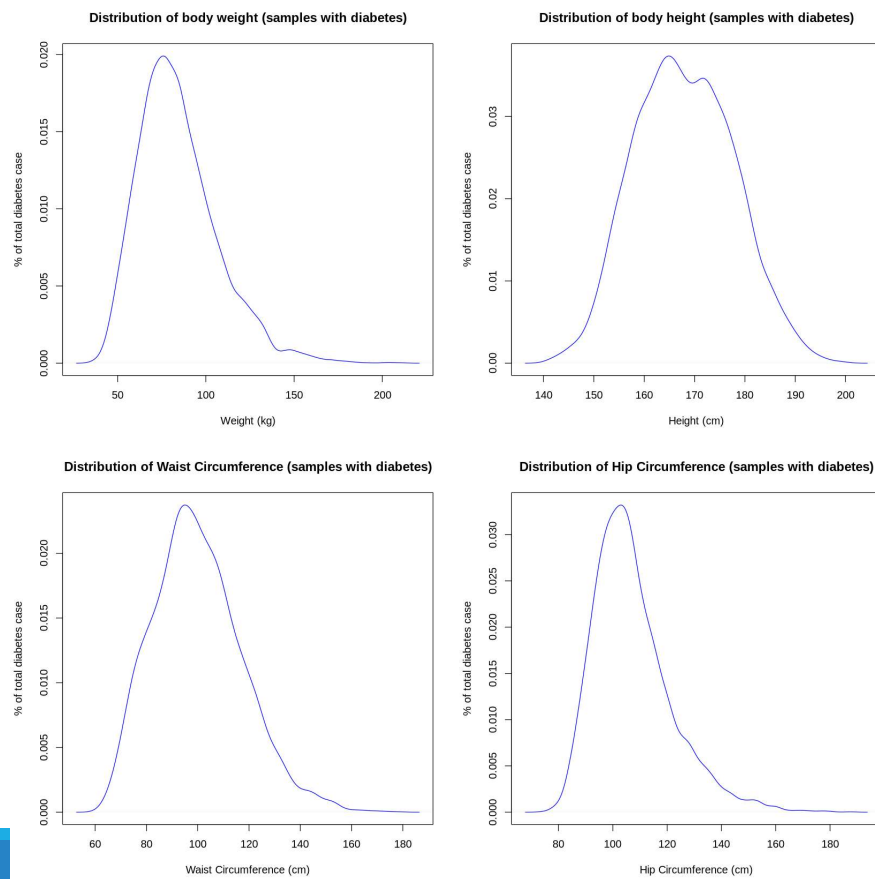
| Data set    |   | # of data           | Data Code |                               | Data Code |  |
|-------------|---|---------------------|-----------|-------------------------------|-----------|--|
| P_ALQ       | 2017-March 2020 Pre-Pandemic Alcohol Use                  | 8965                | SEQN      | Respondent sequence number    | BMXBMI    | Body Mass Index (kg/m**2)                |
| P_BMX       | 2017-March 2020 Pre-Pandemic Body Measures                | 14300               | DIQ010    | Doctor told you have diabetes | BPQ020    | Doctor told you - high blood pressure    |
| P_BPQ       | 2017-March 2020 Pre-Pandemic Blood Pressure & Cholesterol | 10195               | BMXWT     | Weight (kg)                   | BPQ080    | Doctor told you - high cholesterol level |
| P_DIQ       | 2017-March 2020 Pre-Pandemic Diabetes                     | 14986               | BMXHT     | Standing Height (cm)          | ALQ121    | Past 12 mo how often drink alcoholic bev |
|             |   |                     | BMXWAIST  | Waist Circumference (cm)      | ALQ130    | Avg # alcoholic drinks/day - past 12 mos |
|             |   |                     | BMXHIP    | Hip Circumference (cm)        |           |  |
|             |   | Number of Instances |           | Number of Attributes          |           |  |
| After merge |   | 5428                |           | 12                            |           |  |

# Data analysis procedure

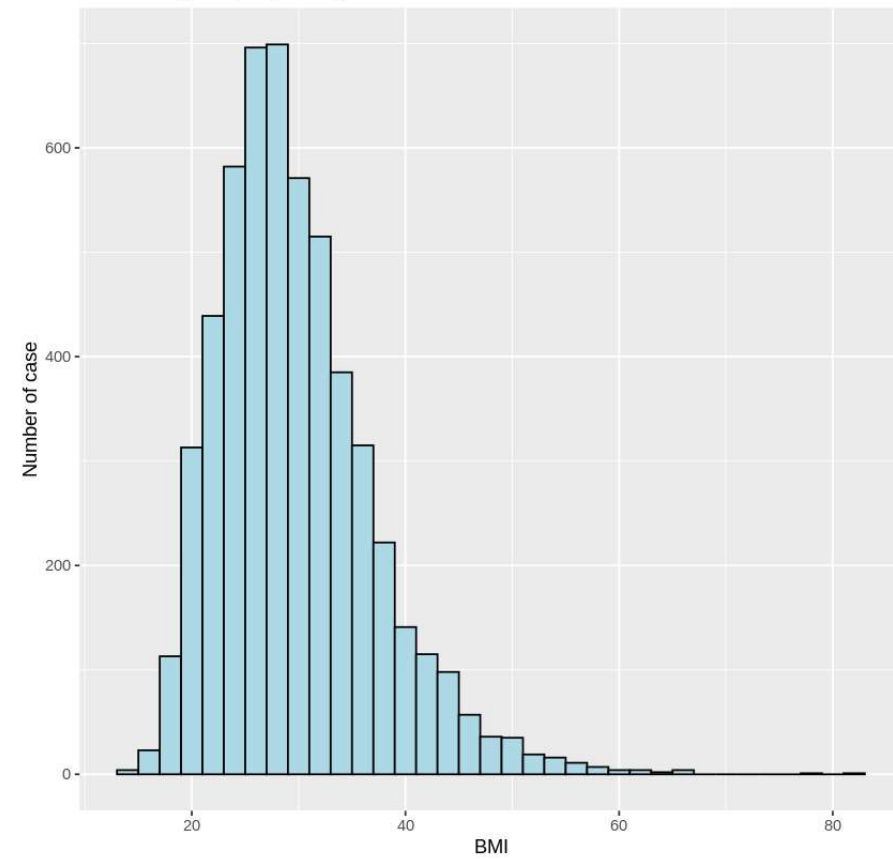
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1. Download data from National Health and Nutrition Examination Survey
2. Merge data into data frame
3. Data transformation
  1. Yes-No answer into binary answers
  2. Data normalization
  3. Data filtering with answers such as “Don’t know”, “Refused to answer”, “Missing” etc.
4. Data pre-processing
  1. Calculate BMI base on weight and height

# Data overview



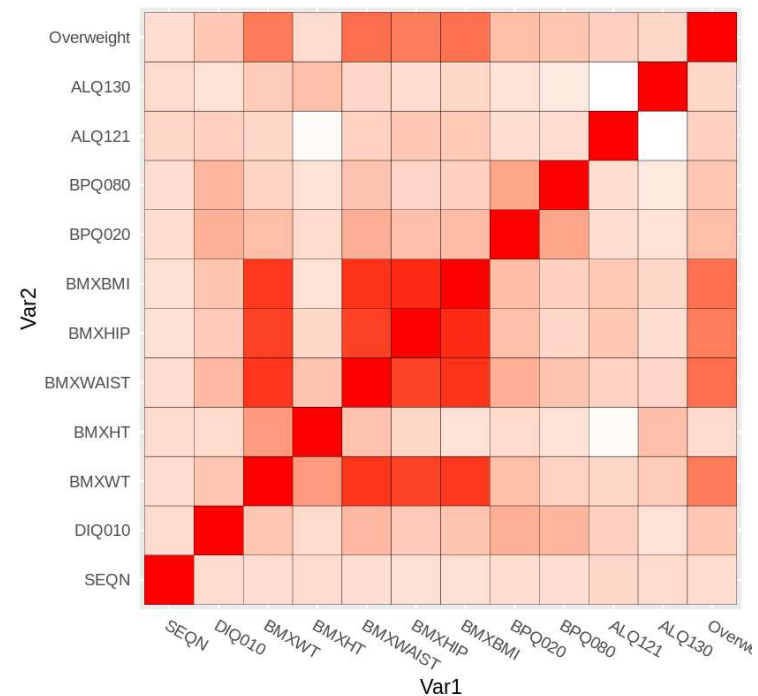
BMI histogram (step as 2)



# Data overview

Highly correlated pair:

- BMI – Hip Circumference
- BMI – Waist Circumference
- BMI – Weight
- BMI – Height
- Diabetes – Waist Circumference
- Diabetes – BMI
- Diabetes – high blood pressure
- Diabetes – high cholesterol level



Correlation heatmap

# Results - Logistic regression model

```
Call:
glm(formula = DIQ010 ~ BMDXBMI + BMDXMAIST, family = binomial(link = "logit"),
    data = OtData_selected)

Deviance Residuals:
    Min       1Q   Median       3Q      Max
-1.7834  -0.5330  -0.4000  -0.2877   2.6825

Coefficients:
            Estimate Std. Error z value Pr(>|z|)
(Intercept) -7.357782   0.314908  -23.365  <2e-16 ***
BMDXBMI      -0.138204   0.014418   -9.586  <2e-16 ***
BMDXMAIST     0.092273   0.006203   14.876  <2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
Call:
glm(formula = DIQ010 ~ Overweight, family = binomial(link = "logit"),
    data = OtData_selected)

Deviance Residuals:
    Min       1Q   Median       3Q      Max
-0.5592  -0.5592  -0.5592  -0.3163   2.4577

Coefficients:
            Estimate Std. Error z value Pr(>|z|)
(Intercept) -2.9701     0.1217  -24.409  <2e-16 ***
Overweight    1.1935     0.1298   9.197  <2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
Call:
glm(formula = DIQ010 ~ BPQ080, family = binomial(link = "logit"),
    data = OtData_selected)

Deviance Residuals:
    Min       1Q   Median       3Q      Max
-0.7279  -0.3721  -0.3721  -0.3721   2.3258

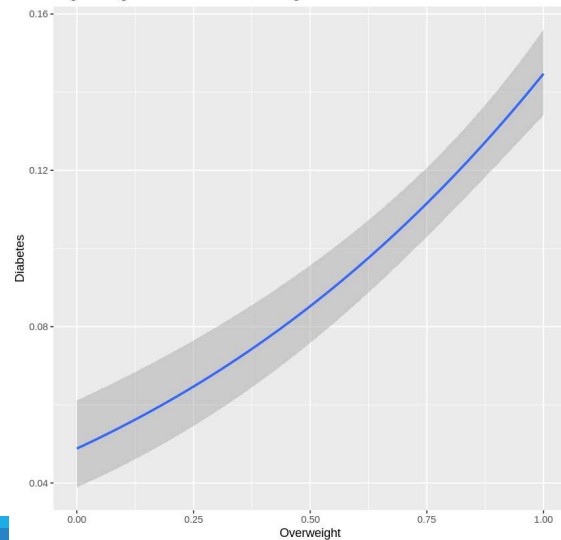
Coefficients:
            Estimate Std. Error z value Pr(>|z|)
(Intercept) -2.6353     0.0656  -40.17  <2e-16 ***
BPQ080        1.4422     0.0871   16.56  <2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
Call:
glm(formula = DIQ010 ~ BPQ020, family = binomial(link = "logit"),
    data = OtData_selected)

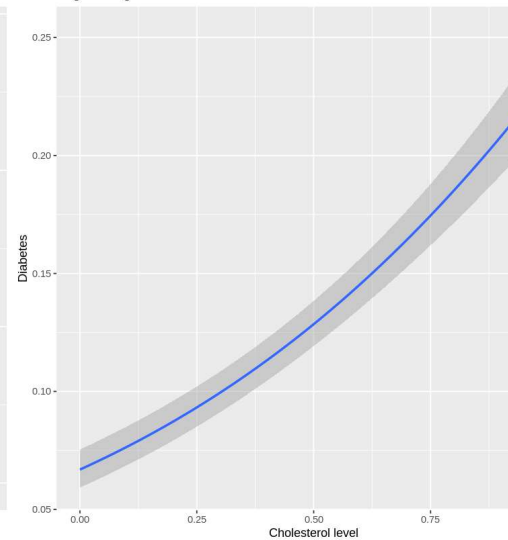
Deviance Residuals:
    Min       1Q   Median       3Q      Max
-0.7493  -0.3457  -0.3457  -0.3457   2.3863

Coefficients:
            Estimate Std. Error z value Pr(>|z|)
(Intercept) -2.78751   0.07076  -39.39  <2e-16 ***
BPQ020        1.66088   0.08977   18.50  <2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

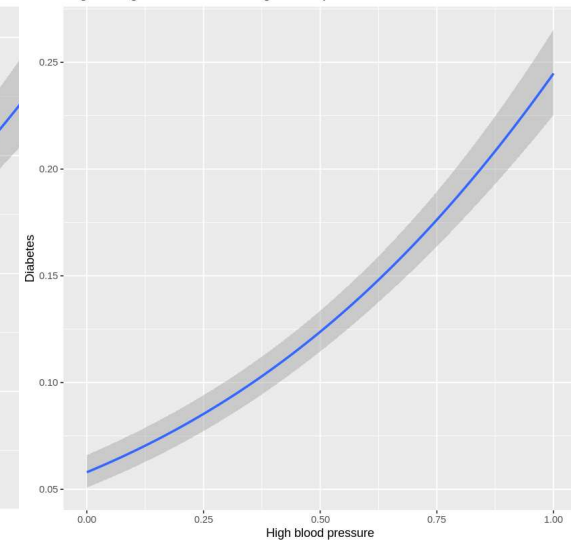
Logistic regression model of Overweight ~ Diabetes



Logistic regression model of Cholesterol level ~ Diabetes



Logistic regression model of High blood pressure ~ Diabetes





# Result – Multiple Logistic regression model

```
Call:
glm(formula = DIQ010 ~ ALQ130, family = binomial(link = "logit"),
    data = OtData_selected)

Deviance Residuals:
    Min       1Q   Median       3Q      Max
-0.5265 -0.5265 -0.5103 -0.4794  2.4057

Coefficients:
            Estimate Std. Error z value Pr(>|z|)
(Intercept) -1.83975    0.06819  -26.980  < 2e-16 ***
ALQ130      -0.06647    0.02297   -2.894  0.00381 **
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

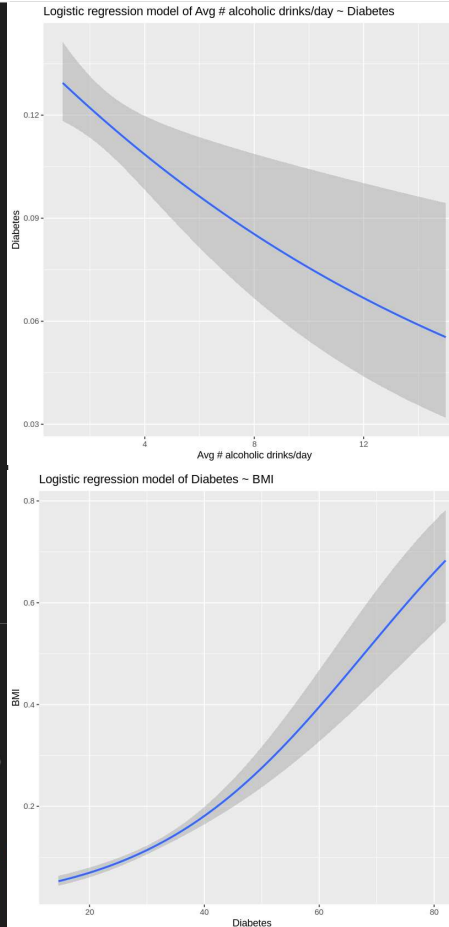
(Dispersion parameter for binomial family taken to be 1)

Null deviance: 3961.9  on 5427  degrees of freedom
Residual deviance: 3952.7  on 5426  degrees of freedom
AIC: 3956.7

Number of Fisher Scoring iterations: 5

Pearson's product-moment correlation

data:  OtData_selected$BMXBMI and OtData_selected$DIQ010
t = 10.935, df = 5426, p-value < 2.2e-16
alternative hypothesis: true correlation is not equal to 0
95 percent confidence interval:
 0.1207105 0.1727718
sample estimates:
cor
0.1468428
```



```
Call:
glm(formula = DIQ010 ~ BMXBMI + BMXWAIST, family = binomial(link = "logit"),
    data = OtData_selected)

Deviance Residuals:
    Min       1Q   Median       3Q      Max
-1.7834 -0.5330 -0.4000 -0.2877  2.6825

Coefficients:
            Estimate Std. Error z value Pr(>|z|)
(Intercept) -7.357782    0.314908  -23.365  <2e-16 ***
BMXBMI      -0.138204    0.014418   -9.586  <2e-16 ***
BMXWAIST      0.092273    0.006203  14.876  <2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 3961.9  on 5427  degrees of freedom
Residual deviance: 3609.4  on 5425  degrees of freedom
AIC: 3615.4

Number of Fisher Scoring iterations: 5
```

# Result – Multiple Logistic regression model

```
Call:
glm(formula = DIQ010 ~ BMXWAIST + BMXHIP + BMXBMI + BPQ020 +
     BPQ080 + ALQ130, family = binomial(link = "logit"), data = OtData_selected)

Deviance Residuals:
    Min       1Q   Median       3Q      Max
-1.6522  -0.4936  -0.3088  -0.1972   2.9010

Coefficients:
              Estimate Std. Error z value Pr(>|z|)
(Intercept) -4.445738    0.525961  -8.453  < 2e-16 ***
BMXWAIST      0.077596    0.006728  11.533  < 2e-16 ***
BMXHIP       -0.057188    0.008747  -6.538  6.24e-11 ***
BMXBMI       -0.004052    0.020098  -0.202   0.84023
BPQ020        1.021170    0.098285  10.390  < 2e-16 ***
BPQ080        0.883811    0.096445   9.164  < 2e-16 ***
ALQ130       -0.077706    0.024492  -3.173   0.00151 **
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

    Null deviance: 3961.9  on 5427  degrees of freedom
Residual deviance: 3248.5  on 5421  degrees of freedom
AIC: 3262.5

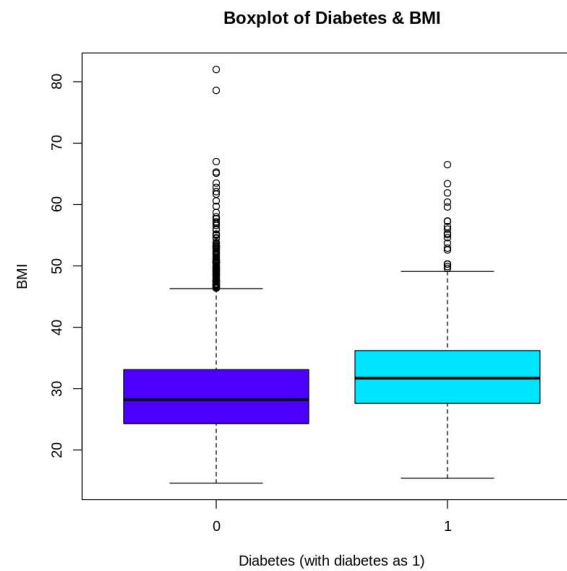
Number of Fisher Scoring iterations: 6
```

Diabetes ~ WAIST+HIP+BMI+BPQ020+BPQ080+ALQ130

# Boxplot analyzing

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## DIABETES & BMI



# Conclusion and Discussion

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- Significant relation with diabetes
  - Waist Circumference
  - Hip Circumference
  - BMI
  - High blood pressure
  - High cholesterol level
  - Heavy alcohol taking
- Having these factors
  - Tended to have higher chance of having diabetes
- Limitation
  - More variables can be used in this project
  - More datasets can be used in this project