On The Association Between Dietary Factors and Mortality Due to Stroke and Diabetes:

A Retrospective Examination of Western and Asian High-Income Countries.

By

Faaiza Castellanos, B.A. (Economics), University of California at Berkeley, 2000 and Jill DeWitt, M.S (Secondary Math Education), Grand Valley State University, 2006

Advisors:
Dr. Jossy Uvah
Dr. Achraf Cohen

A Graduate Capstone Project In Partial Fulfillment of the Degree of Master of Science in Mathematical Sciences University of West Florida

Contents

Li	st of	Figures	iv
Li	st of	Tables	iv
\mathbf{A}	bstra	act	1
1	Inti	roduction	2
	1.1	Statement of Problem	2
	1.2	Relevance of Problem	2
	1.3	Literature Review	2
	1.4	Limitations	4
2	Die	tary Factors and Mortality	5
	2.1	The Data Set	5
	2.2	Principle Components Analysis	6
	2.3	Methodology: Justification, Assumptions and Development of the Model	11
	2.4	Testing and Analysis	13
		2.4.1 Model Formulation for Stroke Rate	13
		2.4.2 Model Formulation for Diabetes Rate	18
3	Cor	nclusions	22
	3.1	Interpretation and Summary	22
	3.2	Suggestions for Further Study	22
\mathbf{R}	efere	nces	24
\mathbf{A}	ppen	dixes	25
	App	endix A: Full Data Set	25
	Ann	endix B: B Code	28

List of Figures

1	PCA Biplot for Year 1990	8
2	PCA Biplot for Year 2000	9
3	PCA Biplot for Year 2010	10
4	Histogram for Stroke rate	14
5	95% Confidence Interval for Lamda, Stroke Rate	14
6	Diagnostic Plot for Log Model, Response Variable = Stroke	15
7	Histogram for Stroke Rate after log transformation	15
8	Histogram for diabetes rate	18
9	95% Confidence Interval for Lamda, Diabetes Rate	19
10	Diagnostic Plot for Box-Cox Model, Response Variable = Diabetes	19
11	Histogram for Diabetes Rate after Box-Cox transformation	20
	List of Tables	
1	Summary Statistics for Stroke Rate	13
2	Linear Regression results for Stroke Rate with Log transformation	16
3	Comparison of full and final models for Stroke Rate	16
4	Final linear regression model results for Stroke Rate, with Log Transformation $$.	17
5	Summary Statistics for diabetes Rate	18
6	Linear Regression results for Diabetes Rate with Box-Cox transformation $\ \ldots \ \ldots$	20
7	Comparison of full and final models for Diabetes Rate	20
8	Final linear regression model results for Diabetes Rate, with Box-Cox Transformation	21

Abstract

A Retrospective Look at the Association Between Dietary Factors and Mortality due to Stroke and Diabetes in Western and Asian High-Income Countries

The purpose of this research paper was to explore the relationship between diet, specifically the consumption of fruit, non-starchy vegetables, bean and legumes, nuts and seeds, unprocessed red meats, sugar-sweetened beverages, fruit juices, milk, protein, calcium, and protein, and rate of stroke and diabetes in Western, high-income countries. Our study took a retrospective approach by looking at data from three years-1990, 2000, and 2010. We used Principal Component Analysis and Multiple Linear Regression to determine which foods or nutrients had a significant effect on stroke or diabetes rates. Gender and year were also considered as factors in the study. The study found that in general, only a few nutrients had a positive or negative impact on disease rates.

1 Introduction

1.1 Statement of Problem

Diet and how it pertains to the development and progression of chronic disease is a growing topic of interest. The increase in prevalence of deadly diseases, such as diabetes and stroke, has led many to examine their diet in a more profound manner, searching for clues that can prevent, halt the progression of, or even reverse such conditions. In response to the escalating interest in diet and disease, many doctors and self-help gurus are promoting a variety of diets, such as keto, paleo, low-fat, low-carb, and many more. Many ancient civilizations have also promoted a variety of foods based on culture and environment. It seems that nutritional intake and its impact on diseases has been a major concern for most people for thousands of years.

With the variety of recommendations and guidelines circulating, the question remains-What foods or nutrients are linked with some of the main killers in our society, namely diabetes and stroke? In order to study diet, it is important to examine a variety of foods and food groups, rather than just focusing on one or a few. In this paper, we investigate a variety of nutrients intake-fruit, non-starchy vegetables, bean and legumes, nuts and seeds, unprocessed red meats, sugar-sweetened beverages, fruit juices, milk, protein, calcium, and protein, and their link to stroke and diabetes/kidney disease. In addition, we restrict our study to Western and Asian high-income countries in an effort to control factors such as high poverty and low healthcare that contribute to lack of resources and education to buy and choose healthy foods.

1.2 Relevance of Problem

The study of food and its link to diabetes and stroke is of paramount importance to the World Health Organization (WHO) [2]. Every year societies spend a large amount of money on healthcare in order to treat diabetes and stroke. Unfortunately, many individuals also die from these diseases, leading to familial emotional turmoil as well as loss of economic contribution from the deceased individual. If such diseases can be prevented, reversed, and even treated with diet, individuals could decrease healthcare cost, benefit the economy, and be socially and emotionally healthier. Diet is a daily part of everyone's lives, so modifications to diet are available to almost all at a much lower price than medication or surgery.

1.3 Literature Review

The importance of diet in preventing death due to diabetes is a major concern of the World Health Organization (WHO) [2], with valid concern. According to the WHO, the number of people with diabetes rose from 108 million in 1980 to 422 million in 2014 and between 2000 and 2016, there was a 5% increase in premature mortality from diabetes. Diabetes is a disease that then can lead to a myriad of other deadly diseases, such as kidney failure, heart attacks, and stroke. Overall, the WHO estimates that diabetes was the seventh leading cause of death in 2016. However, they do stress the importance of a healthy diet in order to prevent and even reverse diabetes, recommending avoiding sugar and saturated fats. The American

Diabetes Association [3] recommends to fill half your plate with non-starchy vegetables and to also include fruits, lean meats and plant-based sources of protein, less added sugar, and less processed foods.

Strokes are another major cause of deaths worldwide, and the WHO [8] estimates that 15 million people suffer from a stroke worldwide annually, of which 5 million die and another 5 million remain permanently disabled. The major cause of stroke is high blood pressure, which is often associated with diet. The Centers for Disease Control and Prevention (CDC) [7] states that up to 80% of strokes can be prevented through healthy lifestyle changes, and diet is a key component of those changes. According to the CDC, eating plenty of fresh fruits and vegetables, foods low in saturated fats, trans fat, and cholesterol, and foods high in fiber can prevent high cholesterol, which in turn lowers your chances of suffering a stroke. Also, limiting sodium can lower blood pressure, which as indicated before, increases the likelihood of a stroke.

The WHO in Europe [1] provides even further research on the connection between diet and disease and offers recommendations for prevention. The importance of preventing disease with diet is a major issue of study and increasing prevalence of these diseases places a huge stress on the healthcare system, impacting national economies and health service budgets negatively. With correct knowledge and programs to education and support populations in implementing healthy dietary guidelines, many countries can increase longevity, improve their economies, de-stress their healthcare systems, and improve mental health. [15]

To study our data in terms of descriptive statistics, we used Principal Component Analysis, or PCA. PCA [9] is a technique for feature extraction, meaning that it combines our variables in a way that we can omit the least influential variables while retaining their most valuable parts. There are three key assumptions that are behind PCA [16] and can indicate when PCA would not provide a strong analysis. The assumptions are linearity, large variances have important structure, and the principal components are orthogonal. The linearity assumption organizes the analysis as a change of basis. As for variances, principal components with larger variances are indicative of structure, while those with smaller variances just represent noise, which is irrelevant to the strength of the signal being analyzed. The assumption that principal components are orthogonal is suggestive that PCA can be solved using linear algebra decomposition techniques.

We then used Multiple Linear Regression with a Box-Cox transformation to model the data. Halinski and Feldt [11] provide a framework for choosing the best procedure to pursue while keeping two goals in mind. First, the best model should produce an equation that yields the best predictions for the population. Second, the best model should contain an optimal number of explanatory variables. We began our model-building using a general linear model using a Gamma, then Poisson distribution, but ultimately settled on a normal multiple linear regression model with a transformation of the response variable. In general, the multiple regression models in this study strive to balance accuracy and parsimony, that is, the models should accurately describe both the systematic and random components and be as simple as possible. There are four basic assumptions that must be met to use multiple linear regression analysis [12]:

1. There must be a linear relationship between the response variable and the explanatory variables.

- 2. The model error term (referred to as the model "residuals") must be normally distributed. Residuals can be thought of as the information not explained by the model. A residual plot and QQ plot can be used to determine if this assumption is met.
- 3. There should not be any multicollinearity. This means that explanatory variables are not correlated with each other. This assumption can be tested by examining the Variance Inflation Factor.
- 4. Homoscedasticity—This means that the variance of the error terms are consistent across the explanatory variables. A Scale-Location plot can be used to determine if this assumption is met.

The regression analysis in this study addresses each of these assumptions for each proposed multiple linear regression model. Nested models were compared using the Akaike's Information Criterion (AIC) and Bayesian Information Criterion (BIC). Our goal was to consider these two measures together and to choose models that were were favored by both criterion, as argued by Kuha [13].

1.4 Limitations

There were many limitations of the various methods of data collection and analysis. We collected data from the Global Dietary Database, or GDD, and they state that they collect data on dietary habits via surveys, which then are based on the volunteer's bias. Some volunteers might not remember their exact diet, others might overstate or understate their food consumption, and yet others might just not wish to disclose honest information. Of course, the GDD uses a complex food description and classification system to address the issue of variation of description of diet, but this does not address the issue of volunteer dishonesty or lack of memory recall.

In addition, the data collected was based on observational studies, where the volunteers were not affected in any manner and were just questioned on their diet. The data that was studied could easily indicate correlation, but not necessarily causation. A more promising study would be an experimental study on diet, where volunteers would be divided into an experimental and control group. Each nutrient could be then studied separately to see how it affects development of diabetes and stroke. Of course, such a study would be very time-consuming, as human lifespan can last upwards of 100 years. In addition, to study a specific nutrient in such a manner would require that the experimental group consume that nutrient consistently for years on end. Certain foods being studied, such as sugar-sweetened beverages, are commonly considered to be unhealthy, and to require a group of volunteers to consume the beverages consistently for years and years would be unethical.

The data collected was international, indicating a large variety of genetics, spices, culinary techniques, exercise, and social behavior that play a part in disease. Genetics are a factor that can affect someone's chance of developing diabetes or a stroke, as well as influence how sensitive they are to certain food items. The use of spices may also affect how foods react in the body and lead to the development of disease. Culinary techniques, such as deep frying, grilling,

boiling, and sauteing, can also alter foods and contribute to diabetes and strokes. Other than dietary factors, exercise is a main agent in health, affecting diet metabolism and also disease. Lastly, social behavior can lead to change in emotion and eating habits which can greatly affect disease. There is much research on emotions, neurotransmitters, and disease, and even more to study.

2 Dietary Factors and Mortality

2.1 The Data Set

We obtained the data regarding nutrition from The Global Dietary Database (GDD) [4], which is a project of the Gerald J. and Dorothy R. Friedman School of Nutrition Science and Policy at Tufts University. The program is also supported by the Bill and Melinda Gates Foundation, which is actively engaged in programs to support public policy. The goal of the program is to understand and improve diet through data collection, analysis, and recommendation, thus leading to public policies that aim to prevent disease and improve healthcare. GDD collects its data from [6] items, there is a large variation on their descriptions. In order to address this issue, the GDD applied FoodEx2, which is a complex food description and classification system developed by the European Food Safety Authority. This system allows GDD to standardize the global dietary intake, thus leading to data which is more reliable.

We obtained the data regarding disease from the Global Health Data Exchange (GHDx) [5]. The GHDx is a data catalog created and supported by an independent global health research center at the University of Washington. The population figures used to calculate the disease rates are estimated based on World Population Prospects: 2015 Revision, from the United Nations Population Division and disease mortality figures are obtained from the WHO Human Mortality Database.

We are most interested in discovering which nutrients had a positive or negative impact on disease rates and whether gender or year was a significant factor. The data set consisted of the following variables:

- Country-Each row contained data from one of twenty-eight countries. All countries were Western or Asian high-income countries. A full list of the countries can be found in Appendix A.
- Gender was a coded continuous factor where Female = (1) and Male = (2)
- Nutrient Intake- Eleven nutrients were included in the data set- fruit, non-starchy
 vegetables, bean and legumes, nuts and seeds, unprocessed red meats, sugar-sweetened
 beverages, fruit juices, milk, protein, calcium, and protein. All measurements were in
 average grams per day, except calcium and potassium, which were measured in average
 milligrams per day.
- **Disease Rates** Rates for stroke and diabetes for each country were measured as counts per 100,000 people.

• Year- Data from each country was included for three different years, 1990, 2000, and 2010. The year was a coded continuous factor where 1990= (1), 2000= (2), and 2010= (3).

In summary, each of the 28 countries has 6 rows of data, a row for each gender and for each of the three years. We gathered the data from two online sources. The Global Dietary Database [4] provided the nutrient data, and the Global Health Data Exchange [5] provided the disease rate data. We then used the statistical computing and programming language R to read the data, and for all statistical graphics and regression analysis. The R code can be viewed in Appendix B.

2.2 Principle Components Analysis

In an attempt to factor a variety of aspects of diet into our study, we obtained data for 11 nutrients. This is a substantial amount of data, so we utilized Principal Component Analysis (PCA) [16] to identify which variables impact disease the most. We utilized PCA as a purely descriptive statistics method, thus we used it to identify key foods and nutrients for the years 1990, 2000, and 2010, rather than to omit variables in our analysis. Biplots were created, using R, to help visualize the principle components and to see if any clusters of countries are revealed.

We first examine the PCA biplot for 1990. A PCA biplot [14] shows the PCA score plot and the loading plot, where the PCA score plot displays the PCA scores and the loading plot portrays how strongly each of the variables impacts a principal component. The PCA Biplots for all three years are shown at the end of this section (Figures 1-3).

As seen in Figure 1, the PCA score plots are the individual countries where the data was gathered from and the PCA loading plot vectors are the food variables. Examining the PCA scores, clusters of countries represent countries that impact the two PCAs in a similar manner, meaning they exhibit similar characteristics. The country represented by 43 and 44 is the Republic of Korea (male and female), so this country impacts PCA 1 strongly but PCA 2 negligibly. The countries represented by 15, 16, 23, and 24 are Finland and Iceland. These countries impact both PCA 1 and PCA 2. Both of these countries are high-income European nations, and they have similarities in their food intake. Examining the loading plot, vectors that are close to each other and have small angle between them are positively correlated. Non-starchy vegetables and beans and legumes are positively correlated, and so are sugar-sweetened beverages and fruit juice. If vectors meet at a 90°angle, they are not correlated, such as nuts and seeds and milk, as well as fruits juice and non-starchy vegetables. Lastly, if two vectors diverge at a large angle, such as 180°, they are negatively correlated. Non-starchy vegetables, beans and legumes, and fruit are negatively correlated with milk.

The PCA biplot for 2000 (Figure 2) shows us similar information. Again, the Republic of Korea was a cluster by itself, along with a cluster shown for Finland and Iceland. Non-starchy vegetables and beans and legumes are again positively correlated, and so are sugar-sweetened beverages and fruit juice. Nuts and seeds and milk, as well as fruits and non-starchy vegetables are not correlated. Again, non-starchy vegetables, beans and legumes, and fruit are negatively correlated with milk.

Lastly, the PCA biplot for 2010 (Figure 3) shows similarities to the plots for 1990 and 2000 but does vary. The country represented by 35 and 36, the Netherlands, is a cluster by itself. The countries 43, 44, and 45 are also clustered, referring to the Republic of Korea and Singapore. As for the loading plot, non-starchy vegetables and beans, as well as sugar-sweetened beverages, fruit juice, and nuts and seeds are positively correlated. Milk, protein, and potassium are also positively correlated. Protein and fruit juice exhibit no correlated, as well as fruit juice and non-starchy vegetables and beans. Protein and milk are negatively correlated with beans and non-starchy vegetables. The PCA biplots provide us invaluable information on the relationships between the variables being studied, specifically the countries and the food intake.

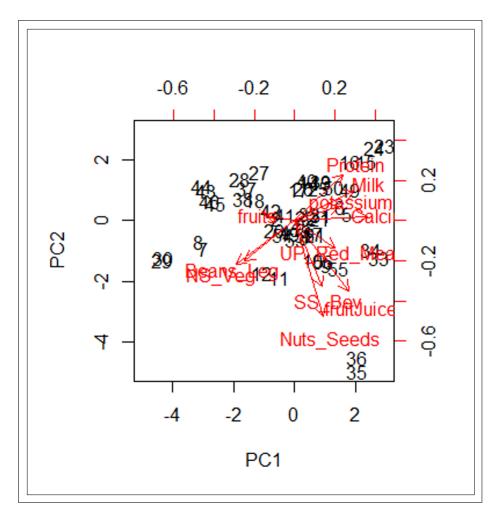


Figure 1: PCA Biplot for Year 1990.

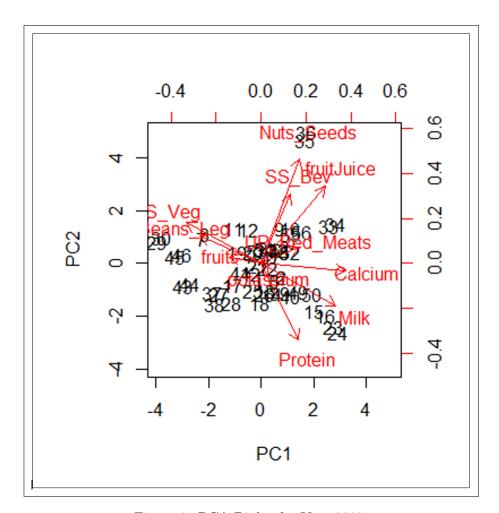


Figure 2: PCA Biplot for Year 2000.

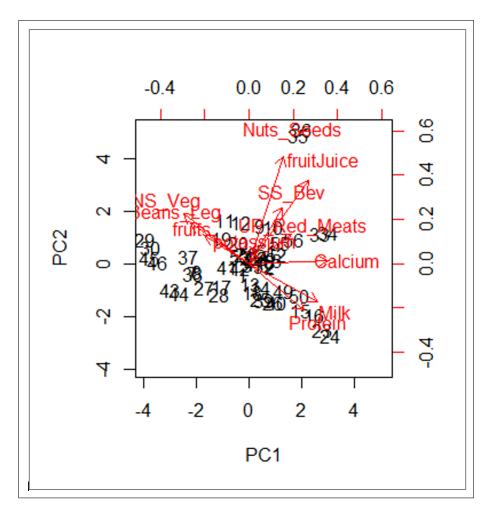


Figure 3: PCA Biplot for Year 2010.

2.3 Methodology: Justification, Assumptions and Development of the Model

The regression analysis focused on two response variables—Stroke rate and diabetes rate. Each response variable is recorded as a count per 100,000. Each model included thirteen potential explanatory variables—eleven nutrients, gender, and year. Several linearized and general linearized models were investigated, analyzed and compared to determine which model had the best fit. Since the response data was count data, two general linearized models were considered. A general linearized model is linear in its parameters. There are three components of a general linearized model. The systematic component is of the form

$$\eta_i = \beta_0 + \beta_1 x_{1i} + \beta_2 x_{2i} + \dots + \beta_3 x_{ni} \tag{1}$$

The random component follows one of the distributions in the family of Exponential Dispersion Models (EDMs). The final component is the link function component, g, which links the mean, μ , to the linear predictor (1), such that

$$g(\mu_i) = \eta_i = \beta_0 + \beta_1 x_{1i} + \beta_2 x_{2i} + \dots + \beta_3 x_{ni}$$

The general linearized model assumes that the response comes from an EDM. Initially, for this study two EDM's were considered-Poisson and Gamma. The Poisson distribution [10] was considered because the data was recorded as counts (per 100,000), which could be considered a rate, but since population data was not available to be used as a meaningful offset it was determined that the Poisson distribution was not appropriate. Next, the Gamma distribution was considered. The Gamma distribution can be considered when the response data is positive and continuous which typically means that the data is skewed to the right. Although, the responses in the data under study was positive and skewed right, it was determined that a transformation of the response did a fairly good job of normalizing the response variable. Ultimately, a general linearized model using the Normal distribution as the EDM was used. This can be justified when the response comes from a normal distribution or a transformation of the response makes it approximately normal. A multiple linear regression model is a special case of the general linearized model. Multiple linear regression models with a response variable y, and p explanatory variables, $x_1, x_2, ... x_p$, consist of two components—a systematic component and a random component. The systematic component is the expected value of the response variable which is linearly related to the explanatory variables x_i such as:

$$\mu_i = \beta_0 + \sum_{i=1}^p \beta_j x_{ji} \tag{2}$$

In equation (2), $\mu_i = E[y_i]$ is the expected value of the response variable and the intercept is β_0 . The regression parameters, β_j, x_{ji} represent each of the p explanatory variables. The response variable, y_i , is the random component in the model. it is assumed to have a constant

variance, σ^2 , and to be normally distributed, that is, $y_i \sim N(\mu_i, \sigma^2)$. It should be noted that linear models are a special case of general linearized models. The goal of the statistical model is to mathematically represent the most important systematic and random components of the data. With a good model, one can understand how variables are related to each other and how the explanatory variables significantly affect the response variable. All model assumptions where checked and analyzed using R. The "Residuals vs. Fitted" plot is used to determine constant variance. Ideally, the plot should show a random pattern around the horizontal line at zero. The "Normal Q-Q plot" can be used to show whether the random component is normally distributed. The "Residuals vs. Leverage" plot can be used to determine if there are outliers or influential observations in the data. The R code for all statistical analysis is located in Appendix B.

Model appropriateness was determined using an ANOVA F-test. This is important because it should be determined whether the explanatory variables are useful predictors of the response variable. This can be determined by testing whether the regression sum of squares (SSReg) is larger than the residual sum of squares (RSS). The ANOVA F-test produces the following results:

• F Statistic:

$$F = \frac{SSReg/(p)}{RSS/(n-p-1)} = \frac{MSReg}{MS_E} \sim F_{(p,n-p-1)}$$

• The Coefficient of Determination, R^2 . This is the proportion of the total variation explained by regression:

$$R^2 = 1 - \frac{RSS}{SS_T}T$$

https://www.overleaf.com/project/60541973b5a9aa428084ba62

• Adjusted R^2 . This is the proportion of the total variation explained by the regression adjusted for the number of explanatory variables:

$$R_{adjusted}^2 = 1 - \frac{RSS/(n-p-1)}{SS_T/(n-1)}$$

In the spirit of parsimony, several nested models were considered. Also, a transformations of the response variables was considered. The goal was to get the simplest models with the best predictive and interpretive value. To compare nested models, Akaike's Information Criterion (AIC) was calculated:

$$AIC = nlog(RSS/n) + 2(p+1)$$
(3)

Smaller AIC values (closer to $-\infty$) represent better models. In equation (3), the term, 2(p+1), is called the penalty. The Bayesian Information Criterion (BIC) was also calculated:

$$BIC = nlog(RSS/n) + log(n)(p+1)$$
(4)

The BIC is inclined to select more parsimonious models than AIC. Smaller BIC values (closer to $-\infty$) represent better models. In equation (4), the term, log(n)(p+1), is called the penalty.

If the assumption of normality of the random variable y_i is not satisfied, a transformation of y_i can be considered. Common transformations include a logarithmic or square root transformation. Another useful method is the Box-Cox transformation. The Box-Cox transformation is used to determine the best lambda, λ , that should be used to transform the response variable, y, where the transformed model becomes $y^{\lambda} = \beta x + \epsilon$. Often, a Box-Cox transformation can improve linearity and homoscedasticity so it can be a very useful tool. The Box-Cox transformation uses a maximum likelihood estimator for λ . If λ is equal to 1, then no transformation is needed. If λ is 0, then a \log transformation should be used, that is $y^{\lambda} = \log(y)$. If $\lambda \neq 0$, then

$$y^{\lambda} = \frac{y^{\lambda} - 1}{\lambda}.$$

Lambda should be relatively small, usually between -3 and 3. R was used to determine the best lambda for the Box-Cox transformation of the data.

2.4 Testing and Analysis

The data was regressed using the thirteen explanatory variables. All model assumptions were checked and addressed if necessary. After an acceptable model was produced, the significant explanatory variables were kept in the model and model adequacy and diagnostics were reevaluated. Ultimately, two models were developed for the response variables stroke rate and diabetes rate. All of the details for model development follow, starting with the model for stroke rate.

2.4.1 Model Formulation for Stroke Rate

Let's begin by examining some descriptive statistics for the response variable Stroke rate. From Table 1, we can see that the mean is greater than the median, so it appears that this data is skewed to the right. The histogram of the variable stroke rate also shows that the data is right skewed (see Figure 4). In order to satisfy the linearity assumption, a transformation of

Minimum	1st Quartile	Median	Mean	3rd Quartile	Maximum
16.88	84.32	123.44	163.55	249.60	451.71

Table 1: Summary Statistics for Stroke Rate

the response variable stroke rate was necessary. The Box-Cox method was used to determine the best transformation. R was used to determine the best lambda, λ , for the Box-Cox transformation. Figure 5 shows the 95% confidence interval for the maximum likelihood estimator, λ .

Using the boxcox function in R, the best lambda was calculated to be $\lambda = \frac{10}{99}$. Note that zero is withing the 95% confidence interval for Lambda in Figure 5. Since zero is within the 95%

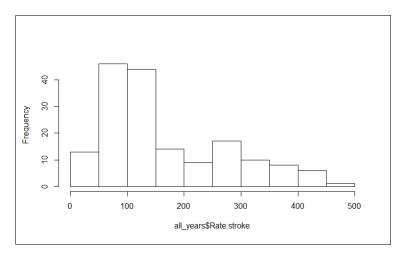


Figure 4: Histogram for Stroke rate

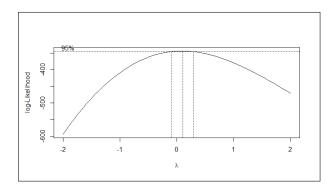


Figure 5: 95% Confidence Interval for Lamda, Stroke Rate.

confindence interval, A log transformation was used. The model diagnostics for the log model are shown on the next page in Figure 6. A visual analysis of the diagnostic plot in Figure 6 demonstrates that there are no major violations of the model assumptions for the transformed model. The residuals vs fitted plot shows a generally random pattern, the normal QQ plot shows some slight deviations from normality near the tails of the data, the scale-location plot shows that the data has homoscedasticity, and the residuals vs. leverage plot show that there are no outliers or influential data points.

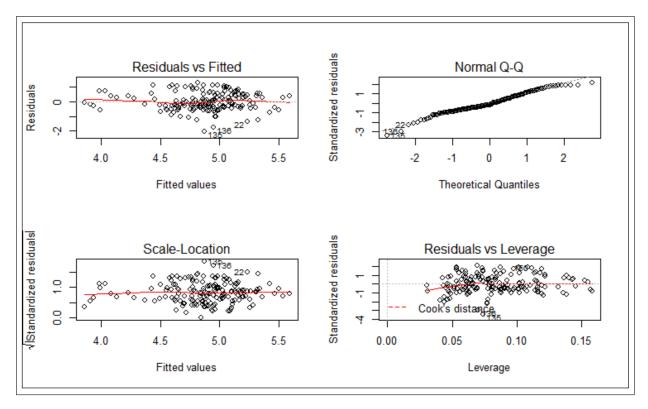


Figure 6: Diagnostic Plot for Log Model, Response Variable = Stroke.

To visualize the effect that a log transformation has on the response variable, we created a histogram of the transformed data. Figure 7 is a histogram of the stroke rate after a log transformation. We can see that the transformed data appears to have more normal distribution.

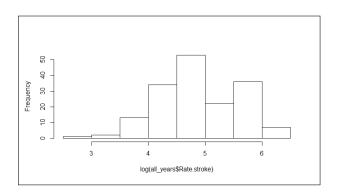


Figure 7: Histogram for Stroke Rate after log transformation.

Table 2 includes a full summary of the multiple linear regression model with a log transformation and all thirteen explanatory variables included in the model. The F-statistic for the model was 3.261, with a corresponding p-value of 0.0002194. The value of R^2 was 0.2155. Using a level of significance of 0.05, the regression model shows that the significant explanatory

	Estimate	Std. Error	t value	$\Pr(> t)$
(Intercept)	1.4537	0.1146	12.69	0.0000
Gender	0.0023	0.0193	0.12	0.9072
year	-0.0130	0.0122	-1.07	0.2860
fruits	-0.0004	0.0003	-1.29	0.2006
$NS_{-}Veg$	0.0004	0.0003	1.26	0.2101
$Beans_Leg$	0.0011	0.0006	1.79	0.0752
$Nuts_Seeds$	-0.0109	0.0043	-2.55	0.0116
UP_Red_Meats	0.0008	0.0004	2.15	0.0334
SS_Bev	-0.0000	0.0002	-0.10	0.9210
fruitJuice	-0.0001	0.0004	-0.21	0.8369
Protein	-0.0010	0.0011	-0.96	0.3395
Calcium	0.0001	0.0001	1.72	0.0879
potassium	0.0000	0.0000	1.86	0.0641
Milk	0.0002	0.0002	1.03	0.3044

Table 2: Linear Regression results for Stroke Rate with Log transformation

variables were nuts & seeds, and unprocessed red meat. Beans & legumes, calcium, and potassium were mildly significant predictors with p-values of 0.0752, 0.0879, and 0.0641 respectively. Next, the model was recalculated using these five explanatory variables. Potassium was no longer significant and was dropped from the model. The final model had an AIC of -757.22 and a BIC of -259.7149, which is evidence that this model is better than the full model containing all thirteen explanatory variables. Comparison of the full and final model are summarized in Table 3.

Model	AIC	BIC	R^2	F-Statistic	p-value
Full Model	-144.9	380.7262	0.2155	3.261	0.0002194
Final Model	-752.22	-259.7149	0.1631	7.939	0.000007153

Table 3: Comparison of full and final models for Stroke Rate

Table 4 summarizes the new multiple regression model with parameter estimates, with a log transformation of the response variable, stroke rate, with the following explanatory variables: nuts seeds, unprocessed red meat, beans legumes, calcium. The sign of the coefficient on the explanatory variable indicates a positive or negative effect on Stroke rate. A positive effect would indicate stroke rate decreased so the sign on the explanatory variable would be negative, likewise, a negative effect would indicate that stroke rate increased so the sign on the explanatory variable would be positive. Nuts & seeds have a positive effect on stroke rate, but unprocessed red meat, beans & legumes, and calcium have a negative effect on stroke rate. According to the CDC [7], foods low in saturated fats can prevent stroke, so nuts and seeds can have high saturated fats thus contributing to a higher risk of stroke, which our study supports. In addition, the CDC states that foods high in in fiber can prevent high cholesterol which can lead to a stroke, and beans and legumes are high in fiber, which our study shows has a negative effect on stroke. Unprocessed red meats can actually have high levels of saturated fat

and cholesterol, which would imply that they would increase the risk of stroke according to the CDC, but we did not find that in our results.

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	1.4458	0.0493	29.32	0.0000
$Nuts_Seeds$	-0.0115	0.0029	-3.88	0.0002
Calcium	0.0002	0.0001	3.18	0.0018
Beans_Leg	0.0015	0.0005	3.07	0.0025
UP_Red_Meats	0.0009	0.0003	2.64	0.0090

Table 4: Final linear regression model results for Stroke Rate, with Log Transformation

2.4.2 Model Formulation for Diabetes Rate

Next, we investigated the analysis of the response variable, diabetes rate. The analysis followed the same process as the analysis for stroke rate, starting with a look at some descriptive statistics for the response variable, diabetes rate (Table 5).

Minimum	1st Quartile	Median	Mean	3rd Quartile	Maximum
17.99	46.75	72.05	116.67	149.87	473.51

Table 5: Summary Statistics for diabetes Rate

From Figure 8, we can see that, like stroke rate, the variable diabetes rate is skewed to the right. Similar to stroke rate, a transformation of the response variable, diabetes rate, was necessary.

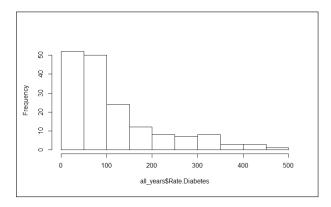


Figure 8: Histogram for diabetes rate.

We used R to determine the best lambda (λ) for the Box-Cox transformation. Figure 9 shows the 95% confidence interval for the maximum likelihood estimator, λ . Using the Box-Cox function in R, the best lambda was calculated to be $\lambda = -\frac{2}{9}$. The 95% confidence interval did not contain zero or another convenient estimate for Lambda, so $\lambda = -\frac{2}{9}$ was used for the transformation.

The model diagnostics for the Box-Cox model are shown in Figure 10. A visual analysis of the diagnostic plot shows that there are no major violations of the model assumptions for either model. The residuals vs fitted plot shows a generally random pattern, the normal QQ plot shows some slight deviations from normality near the tails of the data, the scale-location plot shows that the data has homoscedasticity, and the residuals vs. leverage plot shows that there are no outliers or influential data points.

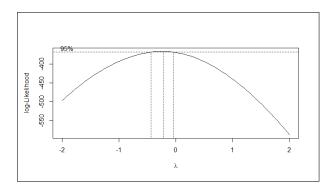


Figure 9: 95% Confidence Interval for Lamda, Diabetes Rate.

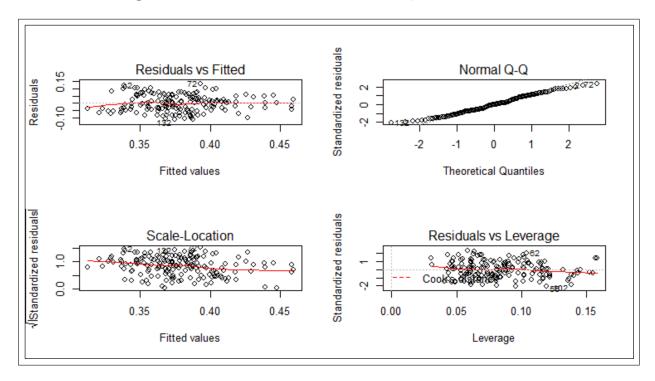


Figure 10: Diagnostic Plot for Box-Cox Model, Response Variable = Diabetes.

To visualize the effect that the Box-Cox transformation has on the response variable, we created a histogram of the transformed data. Figure 11 is histogram of the diabetes rate after a Box-Cox transformation (with $\lambda = -\frac{2}{9}$). We can see that the Box-Cox transformation does a good job "normalizing" the response variable.

Table 6 includes a full summary of the multiple linear regression model with a Box-Cox transformation and all thirteen explanatory variables included in the model. Using a level of significance of 0.05, the regression model shows that the significant explanatory variables were unprocessed red meat, protein, and milk. Year, beans & legumes, and nuts & seeds were mildly significant predictors with p-values of 0.07011, 0.09395, and 0.08111 respectively.

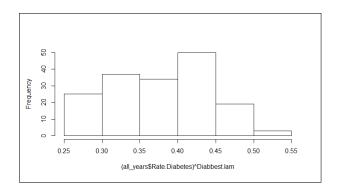


Figure 11: Histogram for Diabetes Rate after Box-Cox transformation.

	Estimate	Std. Error	t value	$\Pr(> t)$
(Intercept)	0.4098	0.0657	6.23	0.0000
Gender	-0.0175	0.0111	-1.58	0.1161
year	0.0127	0.0070	1.82	0.0701
fruits	0.0003	0.0002	1.48	0.1404
$NS_{-}Veg$	-0.0003	0.0002	-1.30	0.1951
Beans_Leg	0.0006	0.0004	1.69	0.0940
$Nuts_Seeds$	0.0043	0.0024	1.76	0.0811
UP_Red_Meats	-0.0007	0.0002	-2.91	0.0042
SS_Bev	0.0001	0.0001	0.52	0.6006
fruitJuice	0.0000	0.0002	0.16	0.8762
Protein	-0.0014	0.0006	-2.32	0.0217
Calcium	0.0000	0.0000	0.38	0.7060
potassium	0.0000	0.0000	0.57	0.5676
Milk	0.0003	0.0001	2.10	0.0372

Table 6: Linear Regression results for Diabetes Rate with Box-Cox transformation

Next, the model was recalculated using these six explanatory variables. Unprocessed red meat and protein were no longer significant and were dropped from the final model. Table 7 summarizes other model statistics that can be used to determine which model is the best. A comparison of the AIC and BIC for both transformations shows that the model using a Box-Cox transformation is best. Based on the model diagnostics, histogram, AIC/BIC, and Coefficient of Determination \mathbb{R}^2 , the linearized model using a Box-Cox transformation was determined to be the best and was used to develop the final parsimonious model.

Model	AIC	BIC	R^2	F-Statistic	p-value
Full Model	-457.7402	-410.8807	0.2019	2.996	0.0006074
Final Model	-937.5627	-440.0555	0.1172	5.41	0.0004082

Table 7: Comparison of full and final models for Diabetes Rate

The final model had an AIC of -937.56 and a BIC of -440.0555, which is evidence that the

final model is better than the full model containing all thirteen explanatory variables. Table 8 summarizes the new multiple regression model, including parameter estimates, with a Box-Cox transformation with the following explanatory variables: year, nuts & seeds, milk, beans & legumes. Nuts & seeds, milk, and beans & legumes all had a negative effect on the response variable, diabetes rate. The explanatory variable "year" was a coded numerical factor where year 1990 was coded as "1", 2000 was coded as "2", and 2010 was coded as "3". In this model that means that for every decade, from 1990 to 2010, the rate of diabetes increased by about 0.28. According to the WHO [2], saturated fats increase the risk of diabetes and non-starchy vegetables, plant-based proteins, and lean meats decrease the risk of diabetes. Our study supports these recommendations as beans and legumes had a negative effect on diabetes and they are plant-based protein sources that the WHO recommends. Nuts and seeds are also also plant based protein sources, yet they can also have high levels of saturated fats. Our study shows that nuts and seeds have a negative relationship. In addition, the WHO estimates that the number of diabetes cases rose from 108 million in 1980 to 422 million in 2014, and our study agrees with the increase in diabetes rate from 1990 to 2010.

	Estimate	Std. Error	t value	$\Pr(> t)$
(Intercept)	0.2873	0.0219	13.14	0.0000
year	0.0185	0.0061	3.03	0.0029
$Nuts_Seeds$	0.0046	0.0016	2.86	0.0047
Milk	0.0002	0.0001	2.30	0.0227
Beans_Leg	0.0005	0.0003	1.89	0.0602

Table 8: Final linear regression model results for Diabetes Rate, with Box-Cox Transformation

3 Conclusions

3.1 Interpretation and Summary

The aim of this project was to study if there is a relationship between diet and death by strokes and diabetes. We started by utilizing Principal Component Analysis to describe the data. We noticed that there were clusters of countries laying geographically close to each other who had similar food intake, such as Finland and Iceland. We also noticed that non-starchy vegetables and legumes were similar in characteristics, and so were sugar-sweetened beverages and fruit juice. After the descriptive statistics, we progressed to create a model for our variables by utilizing regression analysis. Multiple linear regression results provided parameter estimates for the multiple linear regression equations. Following are the final regression equations for the response variable stroke rate and diabetes rate, respectively

$$log(y_{Stroke}) = 1.4458 - 0.0115(NutsSeeds) + 0.0002(calcium) + 0.0015(BeansLegumes) + 0.0009(UPRedMeats)$$

$$(y_{Diabetes})^{\lambda} = 0.2873 - 0.0185(Year) + 0.0046(NutsSeeds) + 0.0002(milk) + 0.0005(BeansLegumes)$$

Note that in each case, the response variable has been transformed. For the response variable, stroke rate a \log transformation was used, and for the response variable, diabetes rate, the Box-Cox transformation, with $\lambda = -\frac{2}{9}$, was used. In this context, interpretation of the intercept does not make sense since that would imply that consuming none of the eleven nutrients is within the parameters of the model. Both models show that consumption of beans and legumes is associated with slightly higher rates of both stroke and diabetes. Nuts and seeds have a positive effect on the rate of diabetes, but a negative effect on the rate of stroke. Consumption of unprocessed red meat has a slightly negative effect on the rate of stroke. Finally, the variable year is a significant factor in the model for diabetes. Because this effect is negative, we can conclude that over time, specifically each decade from 1990 to 2010, the rate of diabetes has decreased. Overall, these models shows some interesting associations between certain nutrients and disease rates for stroke and diabetes. These correlations do not imply causation and further research and experimentation is needed to state definitively that nutrition and consumption of certain foods causes higher or lower rates of stroke or diabetes.

3.2 Suggestions for Further Study

The conclusions made by the model presented demonstrate correlation, but in order to study if causation does in fact exist, an experimental study would need to be conducted. In addition, nuts and seeds, unprocessed red meat, and beans and legumes were determined to influence diabetes and stroke, but the method in how these foods are prepared and seasoned can play a huge role on their effect on disease. It would be useful to study how such methods could affect the correlation, as well as how the addition of other food items, such as spices and condiments, not mentioned in our study could further influence the incidence of disease or stroke. Also, the food categories studied were extremely general, and studying more specific foods could uncover more accurate information on the link of food and disease. For example, when studying unprocessed red meat, specifying how the animals were raised, such as factory-farmed versus grass-fed, and the type of red meat, such as lamb, cattle, or bison, would be extremely relevant to study.

References

- [1] Cancer: Overview, prevention, and management. https://www.who.int/health-topics/cancertab=tab₂, March2021.
- [2] Diabetes: Key facts. https://www.who.int/news-room/fact-sheets/detail/diabetes, March 2021.
- [3] Eat good to feel good. https://www.diabetes.org/nutrition/healthy-food-choices-made-easy, March 2021.
- [4] The gdd 2015 beta-version. https://www.globaldietarydatabase.org/gdd-2015-beta-version, March 2021.
- [5] Global health data exchange. http://ghdx.healthdata.org/, March 2021.
- [6] Scope of current data collection. https://www.globaldietarydatabase.org/methods/scope-current-data-collection, March 2021.
- [7] Stroke. https://www.cdc.gov/stroke/, March 2021.
- [8] Stroke, cerebrovascular accident. https://www.emro.who.int/health-topics/stroke-cerebrovascular-accident/index.html, March 2021.
- [9] M. Brems. A one-stop shop or principle components analysis. Towards Data Science, 2017.
- [10] E. L. Frome. The analysis of rates using poisson regression models. *Biomettics*, Volume 39, No. 3, 1983.
- [11] R. Halinski. The selection of variables in multiple regression analysis. *Journal of Educational Measurement*, Volume 7:151-157, 1970.
- [12] Osborne Jason. Four assumptions of multiple linear regression that researchers should always test. *Practical Assessment, Research, and Evaluation*, Volume 8, 2003.
- [13] J. Kuha. Aic and bic: Comparisons of assumptions and performance. *Sociological Methods* and Research, Volume 33(2):188-229, 2004.
- [14] Linh Ngo. Principal component analysis explained simply. https://blog.bioturing.com/2018/06/14/principal-component-analysis-explained-simply/, June 2021.
- [15] A. Robertson. Diet and disease. WHO Regional Publications, European Series, 2021.
- [16] J. Shlens. A tutorial on principal component analysis. Google Research, 2014.

2000 HIC	HO	HIC	Asia	2000 HIC 2000 Asia	2000 HIC	HC	품	2000 HIC	픙	픙	HC	픙	1990 HIC	등등	픙	HC	1990 HIC	Asia	Asia	1990 Asia	등등	HC	1990 HIC	등등	HC	1990 HIC	등등	HC	HC I	1990 Asia	HC	1990 HIC	1990 HIC	HC	1990 HIC	HC	HC 8	1990 HIC	등등		픙	1990 HIC	HIC	품 등	1990 HIC	Asia	1990 Asia	등	HC E	1990 HIC	등	rign income countries including HIC in Year Asia
CYP	CAN	CAN	BRN	BRN F		AUT	TUA	AUS	USA	USA	GBR	GBR	2 5	SWE	SWE	ESP	ESP 4	SGP	KOR	KOR Z	PRT	NOR	NOR F	NZ KZ	NLD	N E	MLT	Ex	Ę.	D P	ATI	Ā	<u> </u>	IST.	ISL SE	GRC	DEU		FRA	Ī	FE S	DNK	CYP	CYP	CAN	BRN	BRN	品	AUT	AUT	AUS	Country Label
Cyprus	Canada	Canada	Brunei Darussalam	Brunei Darussalam	Belgium	Austria	Austria	Australia	USA, Puerto Rico and US Virgin Islands	USA, Puerto Rico and US Virgin Islands	United Kingdom	United Kingdom	Switzerland, Liechtenstein	Sweden Linchtanetain	Sweden	Spain	Spain	Singapore	Republic of Korea	Republic of Korea	Portugal	Norway, Svalbard and Jan Mayen	Norway, Svalbard and Jan Mayen	New Zealand	Netherlands	Netherlands	Malta	Luxembourg	Luxembourg	Japan	Italy	Italy	Ireland	loeland	Iceland	Greece	Germany	Germany	France, Monaco	Finland	Finland	Denmark	Cyprus	Cyprus	Canada	Brunei Darussalam	Brunei Darussalam	Belgium	Austria	Australia	Australia	Country Name
	0	1		_ 0		0		0 -	. 0		0		0 -	4 0		0			0			0	1		0	_ 0		0			0			0												0			0 -			Gender (female 1, Male 0)
176.8679				93.255791				139 05083					130 14381				130.89755			146.37234			115.48391			168.65582				108 78735		222.10255		_	105.65095		167.60831	205.88083	114.96436	108.36874	135.48993	132.31978	162.84715	193.49896	125 86967		92.637444			140.58208	159.31151	Fruits, grams per day
201.18304				162.02965	139.57938			131 08856		ľ			124 05106			_	129.3286			164.59535			115.7535	_		172.63239				264.8472		166.62062			90.401009			221.08168				129.69385		2	173.51277		166.19374		-	139.63774	_	Non-starchy vegetables, grams per day
55.898224		9.0262394	7.2445016	7.9055419	4.1109142	9.1030989	8.6576595	25.394457	29.514435	24.638857	47.046387	45.199345	6.3736196	13.146868	11.793897	38.108578	35.975891	48.366718	64.499657	72.050262	44.418266	3.7233541	3.485769	83.292503	53.699947	46.96714	25.925381	13.032359	12.215356	81.330879	23.790369	22.692709	39.341621	5.5842638	5.192903	25.325388		11.432217	18.402555	8.3972712	7.8366876	31.080357	74.188736	69.882217	14.888444	7.1396027	7.8864026	4.978188	9.5277596	18.12635	_	Beans and legumes, grams per day
5.8970523	6.8948326	5.9419432	4.2414904	4.2786779	3.3174531	4.2216549	3.6854455	3,4587562	4.2735281	3.6891048	5.1456985	4.4890604	4.4161673	2.8985648	2.5711465	9.1152754	7.9604626	3.1226339	0.5511151	0.57572073	2.9519796	2.7910926	2.3564482	1.5869818	14.155262	12,439658	8.1420155	4.0825024	3.5702846	2.2926466	0.90611005	0.81436813	1.2548673	0.55479509	0.4955461	5.7366557	2.8755314	2.4357324	2.6559777	2.7361574	2,4079967	2.4302604	6.9270687	6.1379318	5.814899	4.1945276	4.3079109	3.0052705	4.6677866	3.6225703	3.1044965	Nuts and seeds, grams per day
105.23022	61.286419	49.979919	17.538639	17.416063	71.919029	117.13522	94.041382	97 200378	54.942394	44.533421	46.56868	42.320396	65.442749	105.86221	83.810837	82.47747	73.190536	37.38768	88.043091	81.908554	107.1513	105.40411	97.540077	70.793404	93.261543	85.827393	123.61172	103.8773	95.585258	56.286198	69.745476	62.514748	43.903561	71.198212	63.2145	101.76545	67.213448	52.780968	44.472233	72.645805	55.491825	63.012394	141.64493	128.18497	57.704567	18.750452	18.684093	76.479645	128.04825	106.43897	65.648636	Unprocessed red meats, grams per day
103.84737	162.68036	124.32528	218.08769	184.18207	157.39064	125.10043	94.232254	159 60054	304.28864	255.1413	183.50287	149.96219	123.73541	93.084892	70.250031	128.84766	99.709251	97.072838	41.35413	26.420126	96.472931	96.584694	71.19194	54.38353	197.34187	168.32877	130.75967	136.64531	103.49871	90.136383	61.049908	48.194157	108.63662	127.63717	102.15916	71.497383	132.15805	89.08918	74.039932	75.450768	50.992348	59.524784	129.36247	100.04411	125.40845	218.29222	182.9133	156.85667	124.90359	156.58426	110.39997	Sugar-sweetened beverages, grams per day
54.305317		_		6.5921154	55.795696	49.08712	54.662483	39 966999	89.807762			4	78.3274				25.057211	7.791151		19.495726	9.5310802	36.161125	40.325745	49.510426	175.69302	194.53049	102.49653	60.644188	67.308907	23.148926	17.334097	19.297182			61.972347			78.332245				45.789452			102 07849		6.6887312	60.583244	54.030766	43.886585		Fruit juices, grams per day
80.811172	73.98101	68.692108	73.263397	73.15107	93.424423	72.300339	67.276268	95 085266	87.827232	82.591835	72.19091		75 791641	83.173424	78.412903	82.673058	76.925148	68.793411	74.754173	74.834068	73.558792	85.101128	79.160393	94.773636	67.804375	66.675926	77.823555	72.079422	66.910301	69.524361	84.018501	76.472603	85.864693	99.234215	92.503426	80.343323	78.160034	75.364861	80.656746	91.423538	85.024368	81.217339	87.33876	81.211693	74.112221	68.177971	68.125137	96.847198	72.493256	92.454269	85.916626	Total protein, grams per day
852.57922	821.00854	901.23657	451.89902	490.51385	829.38818	710.97168	761.10437	774.39563	788.9516	804.12561	807.54443	862.82397	942.86267	833.33661	865.55841	765.55975	846.508	474.39111	443.55917	465.50217	837.24664	751.64691	794.15875	715.68317	883.30524	955.84821	1068.3932	814.24664	870.11877	576.47546 470.68033	821.38068	838.22504	833.08337	965.21991	1044.8717	951.56122	949.29346	1021.333	806.68726	1044.5688	1117,4998	941.45563	772.82031	826.2702	876.97925 798 76971	442.32208	480.82251	802.00458	688.99634	763.20398	749.19263	Calcium, milligrams (mg) per day
2497.0767	3198.7422	3167.7688	1864.545	1908.8055	4503.3428	2999.2932	2973.7349	2761.6936	2738.8032	2707.9653	3449.6738	3242.7217	2750.479	2761.4944	2849.2632	3541.4204	3513.2173	2619.7888	2551.3154	2715.2434	3521.0405	2587.7366	2568.8999	2875.283	2722.731	2770.4138	2250.5452	3298.0132	3276.2073	2774.6497	2937.6484	2688.6426	3049.2185	2364.0696	2344.3608	3975.9375	3330.6543	3431.9038	2929.7466	2555.4165	2542.1951	2788.4971	2519.4734	2503.519	2838.0876	1697.9625	1742.6666	4208.9834	2832.2078	2778.8025		Potassium, milligrams (mg) per day
143.65512	161.26653	166.7795	25.672522	26.36484	86.222519	99.398003	101.09904	156 34441	188.2646	175.69659	196.50546	207.31943	141.2681	331.35205	338.36707	182.50151	184.49921	51.106758	99.735306	117.42743	145.51485	179.80382	183.36372	193.0836	158.26497	163.142	253.13715	106.84019	108.67527	97 743011	116.83884	126.7723	219.54808	331.54584	336.1077	104.14078	90.159378	89.988457	158.46062	319.23703	319.85489	159.17484	170.48256	170.62866	193.03419	27.153236	27.901379	98.078117	116.34063	188.52759	170.35217	Total milk, grams per day
126.0473428	62.73937182	86.35602805	113.6835738	120.1687452	150.5538975	97.90480565	156.0734155	99.08980825	70.31567222	98.92616685	135.4848063	204.4979975	108 2680061	355,4465049	308.1679258	145.5112908	192.1619232	81.83478081	155.3351278	178.6540091	386.6025086	156.6131725	202.410189	124.3430179	97.48312181	138.1856437	138.3912947	405.410232	300.5039534	132 8325032	50.84434268	69.70774718	286.7809788	269.5963559	241.0430157	58.69082204	137.8756664	213.4856576	148.2218127	296.3244102	250.75526	363.2394776	109.5757082	161.6577128	85.77922094	85.81213626	81.36804483	186.2897654	218.5355215	82.99883236	110.5469533	Cause of death, Stroke, per 100k
158.6225631	50.53897904	49.54854092	53.2538607	61.8143391	54.18887139	36.10563245	48.99565223	38.92229661 41 28216489	326.9580503	265.5123857	28.09494174	31.54492021	34 65011637	136.1715584	174.7321513	51.95833946	72.38189698	130.5069454	202.8179374	130.3108224	64.6913029	363.9308139	293.1489594	31.28807023	37.37624852	58.09927991	65.6080854	170.9741857	232.6309167	204.091117	153.3633403	187.2215708	155.249822	90.37161784	110.2412137	312.9250049	36.03715234	61.71786397	260.1729965	125.9928919	191.3584389	182.9027015	101.8639551	137.8075959	36.48229886	141.5343469	141.3815499	332.151185	322.3100979	31.61263907	35.14046125	Cause of death, Diabetes, per 100k
115.9413065	310.874087	252.9996763	67.34412945	66.34622216	308.4196382	351.5719924	295.3756573	306 2768703	41.62657791	47.21529154	430.5239469	360.5623333	338.6819592	33.64092733	31.99576986	379.0985148	217.7680699	36.42483614	31.24411562	26.29604963	249.8211236	29.26598459	27.23773228	262.3652613	382.769191	267,4459986	204.3869992	30.89844745	44.39328323	32.59231802	425.6637881	271.546507	37.0675936	17.31555298	16.34069072	236.3690614	343.0851244	298.2117361	38.01627802	16.77726931	25.31529482	31.09857235	193.9368954	163.1952326	233.952078	80.59348883	82.03344863	53.80870776	47.26288964	304.9853905	223.9384147	Cause of death, Cancer, per 100k

2010 HIC	2010 HIC			2010 HIC			2010 HIC	2010 HIC		2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 Asia	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC		2000 Asia	2000 Asia	2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC			2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC	2000 HIC
Ā	쿈	쿈	Ē	is exc	GRC	DEU		FRA	Ξ	ŦΝ	DNK	CYP	CYP	CAN	CAN	R R	B E	BEL	AUT A	AUS.	AUS	USA	GBR	GBR F	유유	SWE	SWE	ESP	SGP	SGP R	KOR	PRT Z	NOR NOR	NOR	Z Z	- N	NLD	MCT I	E X	Ex	JPN :	.ip i	I TA	쿈	P P	2 2	GRC	GRC		FRA	F I	ΞΞ	DNK	DNK
Italy	Ireland	Ireland	lceland	Iceland	Greece	Germany	Germany	France, Monaco	Finland	Finland	Denmark	Cyprus	Cyprus	Canada	Canada	Brunei Darussalam	Belgium	Belgium	Austria	Australia	Australia	USA, Puerto Rico and US Virgin Islands	United Kingdom	United Kingdom	Switzerland, Liechtenstein	Sweden	Sweden	Spain	Singapore	Republic of Korea Singapore	Republic of Korea	Portugal	Norway, Svalbard and Jan Mayen	Norway, Svalbard and Jan Mayen	New Zealand	Netherlands	Netherlands	Malta	Luxembourg	Luxembourg	Japan	Italy	Italy	Ireland	Ireland	Iceland	Greece	Greece	Germany	France, Monaco	France, Monaco	Finland	Denmark	Denmark
. 0		0	_	0 -	. 0	_	0 -	. 0	_	0	_ <		. 0	_	0 -	_ 0	, _	0	_ 0		0	0 -	. 0	_ 0	o -1	0	_ 0		0	1 0	_	0 -	. 0	_	0 1	. 0	_	0 -	0	_	0 -	4 0		0	_ 0		0	_ (0 1	0	_ 0	o -1	0	_
204.8378	_			92.403442			151,77888				133.47656					99 911453			134.07495		_	90.856201		120.68939			137.94211			208.34708		126.97682			272.64822			124.51641				144 44579			100.32801			156.94447			118.25168			
136.36047	Ť			74,490395				169.24455		9	112.9466				139.78021				117.62232			132.38176		139.30351	_		132.68839	_		153,89455		129.95859			161.52966			98.586006				248 51274			149.22032			153.26363			190.27296			
15.71965	_			17.407503				13.125037	Ė	Ì	6 17.705496					7 7 686 539			6.2503548	Ť		6 18.141495 3 21.805296		34.943378			9 9.9513531			36 890083		38.768681			58.157494 1 57.224464			18.673586	Ť	П		18.414806 14 71 121567			25.93013				8 8.8725176		6 14.073345			Γ
0.92574114	Г	П		7 0.64352256			3,0301588				2.1877427			П		4,6099582		Ì	18 3.5690949			3.7196779 6 4.3188114	П	8 4.3475399			51 8.1687603 51 2.3461249	Ì		0.6232236		3.3407730		N	1.5501611 1.686506			6 9.3423347		П		0.9002974			3 1.1429518	T		5.4906931			5 2.7509842			
57.455776			47	6 54,0448				49.775745		cn	7 56.10442		0,		on on	2 14.834725			9 87.993935	T		4 47 584606		9 36.446835			9 70.147667			6 52.948009 5 19.432501		7 89.300606			66.783241			7 101.62215			4	9 45 46479			8 34.537098			79.633827			2 39.275307			65.019264
59.037506	120.15189	151.4272	100.51331	124,99861	96.712852	83.018646	119.38448	97.037376	50.704315	73.933449	61.610359	106.4399	137.17429	120.29539	156.87065	175 26044	160.17274	204.43196	95.449593	120.16269	169.54768	264.21091	185.81911	153.44957	98.249626	93.741737	70.988266	99.919197	111.94642	42.220142 89.782166	27.05831	127.98326	102.30985	75.855202	63.855618	201.24881	172.63936	178.52522	144.0769	110.22777	117.56184	89 806671	48.001633	152.44305	119.6334	101.85502	98.269821	74.905106	126 41478	98.419083	71.635826	50.244759	83.519646	60.344112
15.169219	Ī			1 49.656445			8 62,023224	_			9 38.407429	Ī			5 89.625923				3 50.372704			89.290665		7 36,51432			6 65.023178			2 16.574518 6 7.6196752		6 7.921093			9 45.151615 8 40.474728			2 77.943062			4 21.21431			ω	4 41.89539			6 25.502964	7		6 22.135496			
84.550903				5 104.88444				96.989616			9 83.950027					68 923637	Ī		63.252792			81.013123		2 68.516869	Ī		82.012474			8 72.970657		80.59671			83,190598			90.797775				83,995117		П	98.738945			4 82.284889	Ī		5 79.237938		Ì	
863.24982				1010.6261		Ì	1017.9942				1003.3715				848.00806				777.98779			825.93347		887.56586			888.84308	00		472.26044	Ì	798.82837	~		548.36914			1020.1349			487.07666	_			887.65271			973.43542			835.85052			
3022.2859				2971,3704		П	3326,1897				3410.5505		N		3270.7629		4	Ì	3084.0129		N	2995.9187		3534.6196			3237.252			2886.0369		3401.4138			3092.6438			3145.1926			2379.3999			Ì	3570.6689			4343.5977			2855.0383			Ī
93.904793				261.75064			75,722328		2		118.02145			_	143.9976				82,628593			149.42509		174.43863			298.54892			80.429169	9	129.8493			132.28484			196.69159				102.46165			164.86142	T	Ĺ	92,874718		ĺ	139.41356			136.37762
71.8183208	cn			16.87892068	N		113,5270951			_	439.351859	Ť		_	69.7841577				343.5863986			70 29134253		172.0273409	١.		2 321.3963728		4	115.0388961		256.2272459			253.1818323			92.78423219				123 73 11 899			264.6181251	-	N	284.8685702		П	270.3462026	Т		
143.6714626		П		8 221.4814037	Т	П	344,5457844	Т	П		94.1391337	_	Т	П	46.75667394	Т			62,73309779			5 65.66429682	П	26.04674614			5 51.30999434 3 179.3799129		_	1 45.7171956		9 73.58777775		П	3 113.5167618	Т		9 51.4539371	Т			34 28639075			1 116.2718072	Т	П	2 46.58440255	_	П	5 113.6834122	N)	П	Т
302.2160063		N		65.26339031		П	78.78684544	Т			62.24757492				269.7434081	Ť			62.67347708		-	277.9705239	П	345.7956347	-		40.07808654	-	П	253.770251	П	383.4590324			39.32462781		2	267.575776				126.0305243		Ť	32.24011579			260.2242339			2 44.6769527			387.839504

2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 Asia	2010 Asia	2010 Asia	2010 Asia	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 HIC	2010 Asia	2010
USA	USA	GBR	GBR	CHE	CHE	SWE	SWE	ESP	ESP	SGP	SGP	KOR	KOR	PRT	PRT	NOR	NOR	NZL	NZL	NLD	NLD	MLT	MLT	LUX	LUX	JPN	-
USA, Puerto Rico and US Virgin Islands	USA, Puerto Rico and US Virgin Islands	United Kingdom	United Kingdom	Switzerland, Liechtenstein	Switzerland, Liechtenstein	Sweden	Sweden	Spain	Spain	Singapore	Singapore	Republic of Korea	Republic of Korea	Portugal	Portugal	Norway, Svalbard and Jan Mayen	Norway, Svalbard and Jan Mayen	New Zealand	New Zealand	Netherlands	Netherlands	Malta	Malta	Luxembourg	Luxembourg	Japan	capain
_	0	_	0	_	0	_	0	_	0	_	0	_	0	_	0	_	0	_	0	_	0	_	0	_	0	_	
85.610626	77.611382	128.05817	106.79031	146.25237	121.26773	140.03665	108.98852	114.5921	101.53709	179.78038	151.5537	147.81392	108.98799	143.55954	123.60711	111.07014	92.75251	258.20181	203.08235	156.00677	136.01802	135.76419	115.87161	118.39726	101.71802	143.60327	
126.75581	117.36357	134.04881	118.46258	127.86882	113.861	124.12965	107.61472	113.21651	102.00794	139.37749	131.36411	127.79362	124.996	137.78847	123.20934	94.128227	83.008415	150.47093	131.3688	151.85458	142.03874	105.42561	92.907547	116.85345	105.29293	235.29352	
14.936326	18.014244	37.272766	39.104225	3.2348995	3.4249625	7.2556591	8.1307516	21.584639	22.848114	25.481451	22.617411	40.311871	36.068375	31.838121	33.584518	0.87635136	0.94205385	43.421249	42.712078	32.883099	37.880409	15.414862	16.629944	10.72389	11.524368	65.309776	001000101
3.8523872	4.4758754	4.5814557	5.2621541	4.4145555	5.0352001	2.6810267	3.0309141	7.4984307	8.5801945	3.9812694	3.9632952	0.69536829	0.67077637	3.0489988	3.473598	2.2352827	2.6522877	1.8943386	2.054219	14.013341	15.996328	7.7011871	8.740593	2.1659908	2.49248	2.5685346	1110011
35.993305	44.456772	32.164341	35.330257	51.507477	56.948917	59.311497	75.007378	56.473423	63.706253	22.458994	23.393072	37.71653	40.552727	70.898842	78.956413	62.921833	68.035492	59.617767	49.662956	70.872177	77.129761	81.192978	87.793488	68.69635	74.674973	41.110466	.0.000.01
262.2959	308.32169	143.84138	172.55777	93.042747	117.40124	73.538139	96.403137	98.242676	126.98962	91.421913	114.15365	25.65185	40.239349	100.88745	130.75531	78.775162	105.28139	58.52277	67.405983	173.45276	200.50316	143.24155	182.77658	113.10076	146.05989	85.407616	
84.853745	76.652885	36.3451	32.57494	78.235161	69.720734	60.30545	54.22554	21.484127	19.332083	6.8724475	6.5692329	14.353333	14.344538	7.978941	7.1638064	31.722685	28.291494	40.62941	36.405117	162.71309	146.56447	78.826424	70.833893	54.232204	48.702518	21.319603	
81.422737	86.656631	67.86953	70.890526	66.961388	70.829819	79.560883	84.408211	76.447685	82.319885	85.890518	86.25061	70.206551	70.160065	74.500534	79.995911	78.052193	83.93605	89.208183	95.942673	80.071655	81.430672	84.887886	90.154648	75.087036	80.890305	68.538383	0110100
845.14313	829.84277	920.23492	862.84344	1056.3242	994.0155	908.10583	874.81073	903.48273	816.99817	531.50629	475.16336	517.81421	493.87964	888.02997	815.50995	854.83472	809.51569	742.83801	562.57703	1019.7216	942.92761	1149.2172	1040.6005	934.83392	875.37512	611.79315	
2841.5598	2880.5049	3533.7256	3770.0779	2726.6533	2761.8665	3223.2559	3127.8467	3083.9258	3113.8892	2721.3833	2647.4302	3037.8237	2859.1116	3377.074	3407.3677	2935.4724	2971.2676	3173.9241	3199.4907	3296.1702	3246.446	3124.4683	3156.5212	3495.3875	3526.5974	2579.1563	1.0000.0.0
135.76813	145.44727	165.94646	158.39336	114.49075	112.69245	261.70264	256.15945	141.09637	138.64375	36.169983	33.101646	73.182602	61.867115	112.91473	112.09917	120.30702	117.7985	121.84873	114.38547	123.96318	120.34307	170.75182	167.79219	76.579056	75.436661	97.622421	0000000000
301.8841784	265.230369	80.91800163	121.5022718	87.99679534	57.85298391	101.1650923	144.8246449	76.36073073	103.3562343	30.69054172	39.25946677	77.74548684	84.76386878	421.1943536	254.1549746	39.60920066	39.69462718	307.1045626	256.0282459	65.24083678	102.6218649	302.331483	213.2543034	74.64477413	112.1511778	451.7099969	
57.2017377	83.04789107	23.16727928	25.40311796	53.81309671	41.49486586	49.18606207	46.93556259	50.33379971	64.58832597	134.7186318	123.5867374	46.93110849	44.74327608	169.7288755	200.4731894	81.98829365	121.5922768	62.77019779	93.37265516	40.4624599	52.30481608	81.84273236	96.20004283	328.8754419	277.9723169	44.58434253	
68.08433031	67.16097462	373.987593	325.8160369	244.6883638	317.2121276	343.5133018	311.320987	368.6210923	218.3745179	15.7089256	20.50034701	271.5005448	157.1828063	84.90402736	95.58113	348.8231187	304.4790117	44.34704244	38.22803639	412.6060599	338.6032818	58.33947441	58.69921152	34.53612412	47.24559132	118.2368252	

```
Appendix B: RCode
setwd("H:/CPC STA6950")
install.packages("xtable")
install.packages("ggplot")
library(ggplot2)
library(xtable)
library(MASS)
library(plyr)
#combine three years into one large data set
y1990<-read.csv("H:/CPC STA6950/STA6950 Project Data Year 1990.csv",
        header = T
v1990$year<-c(1)
y2000<-read.csv("H:/CPC STA6950/STA6950 Project Data Year 2000.csv",
        header = T
v2000$vear<-c(2)
y2010<-read.csv("H:/CPC STA6950/STA6950 Project Data Year 2010.csv",
        header = T)
y2010$year<-c(3)
#rename columns Year 1990
colnames(y1990)[colnames(y1990) %in% c("v01_wt_median","v02_wt_median")]<-c("fruits","NS_Veg")
colnames(y1990)[colnames(y1990) %in% c("v05 wt median","v06 wt median")]<-c("Beans Leg","Nuts Seeds")
colnames(y1990)[colnames(y1990) %in% c("v10_wt_median","v15_wt_median")]<-c("UP_Red_Meats","SS_Bev")
colnames(y1990)[colnames(y1990) %in% c("v16_wt_median","v23_wt_median")]<-c("fruitJuice","Protein")
colnames(y1990)[colnames(y1990) %in% c("v36 wt median","v41 wt median")]<-c("Calcium", "potassium")
colnames(y1990)[colnames(y1990) %in% c("v57_wt_median")]<-c("Milk")
colnames(y1990)[colnames(y1990) %in% c("female.....1.")]<-c("Gender")
colnames(y1990)[colnames(y1990) %in% c("country.name")]<-c("countryname")
#rename columns Year 2000
colnames(y2000)[colnames(y2000) %in% c("v01_wt_median","v02_wt_median")]<-c("fruits","NS_Veg")
colnames(y2000)[colnames(y2000) %in% c("v05_wt_median","v06_wt_median")]<-c("Beans_Leg","Nuts_Seeds")
```

```
colnames(y2000)[colnames(y2000) %in% c("v10_wt_median","v15_wt_median")]<-c("UP_Red_Meats","SS_Bev")
colnames(y2000)[colnames(y2000) %in% c("v16 wt median","v23 wt median")]<-c("fruitJuice","Protein")
colnames(y2000)[colnames(y2000) %in% c("v36_wt_median","v41_wt_median")]<-c("Calcium","potassium")
colnames(y2000)[colnames(y2000) %in% c("v57 wt median")]<-c("Milk")
colnames(y2000)[colnames(y2000) %in% c("female.....1.")]<-c("Gender")
#rename columns Year 2010
colnames(y2010)[colnames(y2010) %in% c("v01_wt_median","v02_wt_median")]<-c("fruits","NS_Veg")
colnames(y2010)[colnames(y2010) %in% c("v05_wt_median","v06_wt_median")]<-c("Beans_Leg","Nuts_Seeds")
colnames(y2010)[colnames(y2010) %in% c("v10_wt_median","v15_wt_median")]<-c("UP_Red_Meats","SS_Bev")
colnames(y2010)[colnames(y2010) %in% c("v16 wt median","v23 wt median")]<-c("fruitJuice","Protein")
colnames(y2010)[colnames(y2010) %in% c("v36_wt_median","v41_wt_median")]<-c("Calcium","potassium")
colnames(y2010)[colnames(y2010) %in% c("v57 wt median")]<-c("Milk")
colnames(v2010)[colnames(v2010) %in% c("female.....1.")]<-c("Gender")
all years<-rbind(v1990,v2000,v2010)
#histogram for response variables
#hist(all years$Rate.Cancer)
hist(all_years$Rate.stroke, main="")
hist(all years$Rate.Diabetes, main="")
##Summary stats for stroke and diabetes rate
s=summary(all years$Rate.stroke)
d=summary(all years$Rate.Diabetes)
#####response var treated as a continuous random var.
####STROKE
##LM model for stroke
modelStr<-lm(formula = Rate.stroke ~ Gender+year+fruits+NS_Veg+Beans_Leg+
        Nuts_Seeds+UP_Red_Meats+SS_Bev+fruitJuice
      +Protein+Calcium+potassium+Milk,
      data = all years)
summary(modelStr)
```

```
par(mfrow=c(2,2))
plot(modelStr)
extractAIC(modelStr)
#BC transformation
bcStr=boxcox(modelStr,lambda = seq(-2,2))
Strbest.lam=bcStr$x[which(bcStr$y==max(bcStr$y))]
##LM model with BC trans
bcmodelstr<-lm((Rate.stroke)^Strbest.lam ~ Gender+year+fruits+NS_Veg+Beans_Leg+
         Nuts_Seeds+UP_Red_Meats+SS_Bev+fruitJuice
       +Protein+Calcium+potassium+Milk,
       data = all_years)
summary(bcmodelstr)
par(mfrow=c(2,2))
plot(bcmodelstr)
extractAIC(bcmodelstr)
hist((all_years$Rate.stroke)^Strbest.lam,main = " ")
BIC(bcmodelstr)
#LM log model
log_modelStr<-lm(formula = log(Rate.stroke) ~ Gender+year+fruits+NS_Veg+Beans_Leg+
          Nuts_Seeds+UP_Red_Meats+SS_Bev+fruitJuice
         +Protein+Calcium+potassium+Milk,
         data = all_years)
summary(log_modelStr)
par(mfrow=c(2,2))
plot(log_modelStr)
extractAIC(log_modelStr)
hist(log(all_years$Rate.stroke),main = " ")
BIC(log_modelStr)
#gamma model link inverse, log trans
```

```
gamma_modelstr=glm(log(Rate.stroke) ~ Gender+year+fruits+NS_Veg+Beans_Leg+
          Nuts_Seeds+UP_Red_Meats+SS_Bev+fruitJuice
         +Protein+Calcium+potassium+Milk,
         data = all years, family = Gamma(link="inverse"))
summary(gamma_modelstr)
par(mfrow=c(2,2))
plot(gamma_modelstr)
extractAIC(gamma_modelstr)
#########End Stroke
####Diabetes
#LM model no trans
modelDiab<-lm(formula = Rate.Diabetes ~ Gender+year+fruits+NS_Veg+Beans_Leg+
        Nuts_Seeds+UP_Red_Meats+SS_Bev+fruitJuice
       +Protein+Calcium+potassium+Milk,
       data = all_years)
summary(modelDiab)
par(mfrow=c(2,2))
plot(modelDiab)
extractAIC(modelDiab)
#BC TRans
bcDiab=boxcox(modelDiab,lambda = seq(-2,2))
Diabbest.lam=bcDiab$x[which(bcDiab$y==max(bcDiab$y))]
#LM model with BC trans
bcmodeldiab<-lm((Rate.Diabetes)^Diabbest.lam ~ Gender+year+fruits+NS_Veg+Beans_Leg+
         Nuts_Seeds+UP_Red_Meats+SS_Bev+fruitJuice
        +Protein+Calcium+potassium+Milk,
        data = all_years)
summary(bcmodeldiab)
par(mfrow=c(1,1))
plot(bcmodeldiab)
```

```
extractAIC(bcmodeldiab)
hist((all_years$Rate.Diabetes)^Diabbest.lam)
BIC(bcmodeldiab)
#LM Log Model
log modeldiab<-lm(formula = log(Rate.Diabetes) ~ Gender+year+fruits+NS Veg+Beans Leg+
          Nuts_Seeds+UP_Red_Meats+SS_Bev+fruitJuice
         +Protein+Calcium+potassium+Milk,
         data = all_years)
summary(log_modeldiab)
par(mfrow=c(2,2))
plot(log_modeldiab)
extractAIC(log_modeldiab)
BIC(log_modeldiab)
#Gamma model link inverse, log trans
gamma_modeldiab=glm(log(Rate.Diabetes) ~ Gender+year+fruits+NS_Veg+Beans_Leg+
           Nuts_Seeds+UP_Red_Meats+SS_Bev+fruitJuice
          +Protein+Calcium+potassium+Milk,
          data = all_years, family = Gamma(link="inverse"))
summary(gamma_modeldiab)
par(mfrow=c(2,2))
plot(gamma_modeldiab)
extractAIC(gamma_modeldiab)
#######END Diabetes
##best models
summary(bcmodelstr)
summary(bcmodeldiab)
summary(log_modelCan)
########mew Models with significant Explanatory Variables only
```

```
####STROKE####
beststroke<-lm((Rate.stroke)^Strbest.lam ~ Nuts_Seeds+Calcium
       +Beans_Leg+UP_Red_Meats,
       data = all_years)
summary(beststroke)
par(mfrow=c(2,2))
plot(beststroke)
extractAIC(beststroke)
BIC(beststroke)
####Diab####
bestdiab<-lm((Rate.Diabetes)^Diabbest.lam ~ year+Nuts_Seeds
           +Milk+Beans_Leg,
       data = all years)
summary(bestdiab)
par(mfrow=c(2,2))
plot(bestdiab)
extractAIC(bestdiab)
BIC(bestdiab)
###export model summaries to Latex###
xtable(beststroke)
xtable(bestdiab)
#PCA for 1990
pr1990<-prcomp(y1990[,5:15],scale=TRUE)
pr1990
summary(pr1990)
par(mfrow=c(1,1))
plot(pr1990,type="l") #Scree plot
biplot(pr1990,scale=0)
```

```
#PCA for 2000
pr2000<-prcomp(y2000[,5:15],scale=TRUE)
pr2000
summary(pr2000)
par(mfrow=c(1,1))
plot(pr2000,type="I") #Scree plot
biplot(pr2000,scale=0)
#PCA for 2010
pr2010<-prcomp(y2010[,5:15],scale=TRUE)
pr2010
summary(pr2010)
par(mfrow=c(1,1))
plot(pr2010,type="l") #Scree plot
biplot(pr2010,scale=0)
plot(modelCan)#PCA for 2010
pr2010<-prcomp(y2010[,5:15],scale=TRUE)
pr2010
summary(pr2010)
par(mfrow=c(1,1))
plot(pr2010,type="l") #Scree plot
biplot(pr2010,scale=0)
##All Years
prAll<-prcomp(all_years[,5:15],scale=TRUE)</pre>
prAll
summary(prAll)
par(mfrow=c(1,1))
plot(prAll,type="l") #Scree plot
biplot(prAll,scale=0)
```