

Medical Insurance Regression Analysis

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Introduction

2.7k

Observations

07

Variables

Categorical: smoker, region, sex
Numerical: Cost, age, BMI, children

Analysis Goal

**Our goal is to better understand which factors
affect medical insurance costs**



Data Cleaning

1.3k

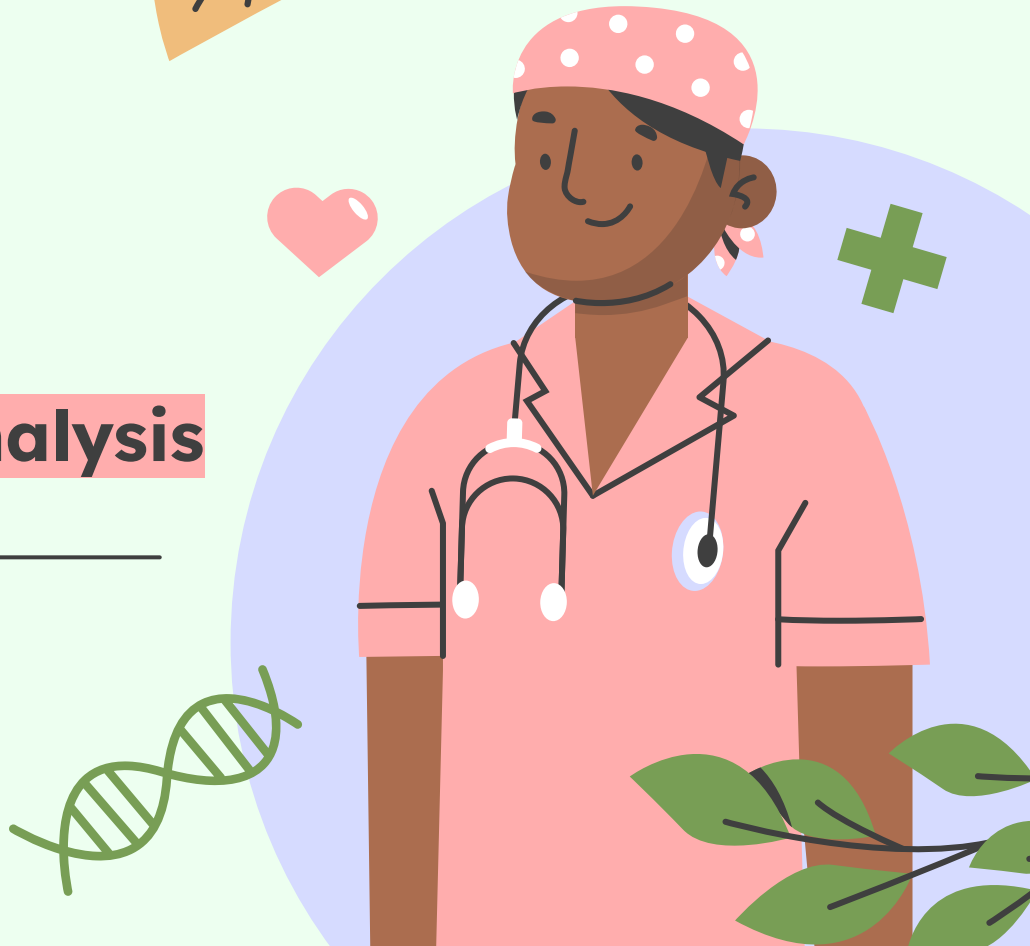
**Observations After Removing
Duplicates and Locating Null Values**

Variables

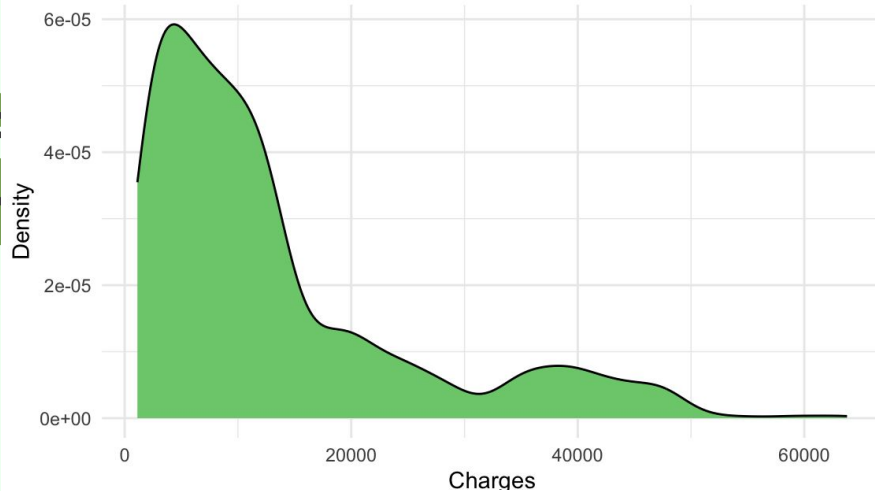
Regressors: Age, Sex, BMI, Children, Smoker, Region
Response: Charges

01

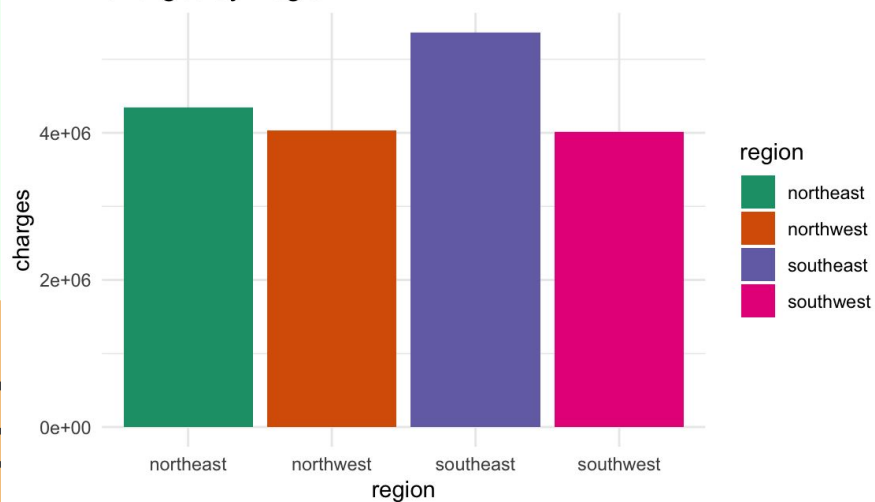
Exploratory Data Analysis



Distribution of Charges

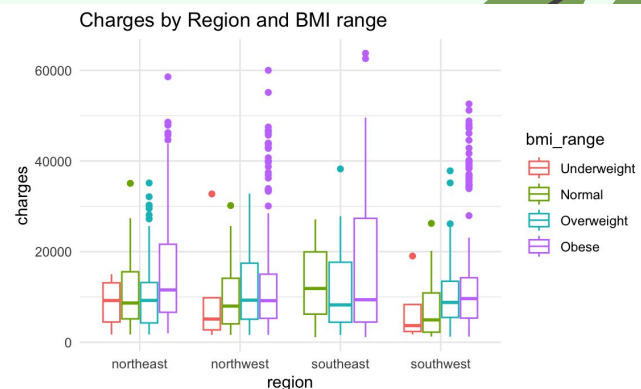
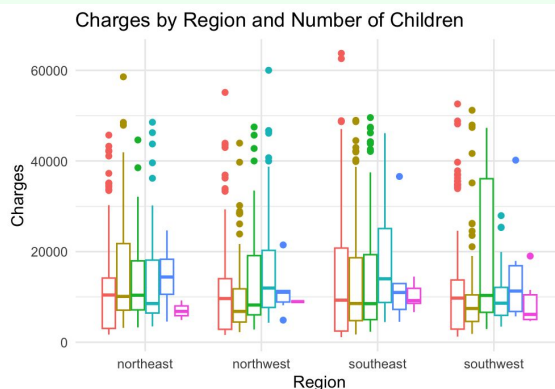
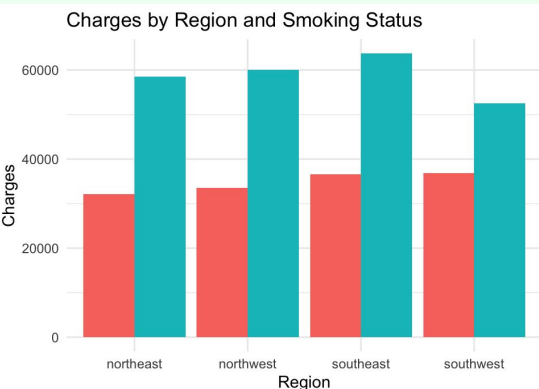


Charges by Region



- ❑ Right skewed distribution
- ❑ Majority of charges are under \$20,000
- ❑ Range from \$1.12k to \$63.8k on the high end
- ❑ Southeast region has highest insurance charges
- ❑ Other regions have about the same

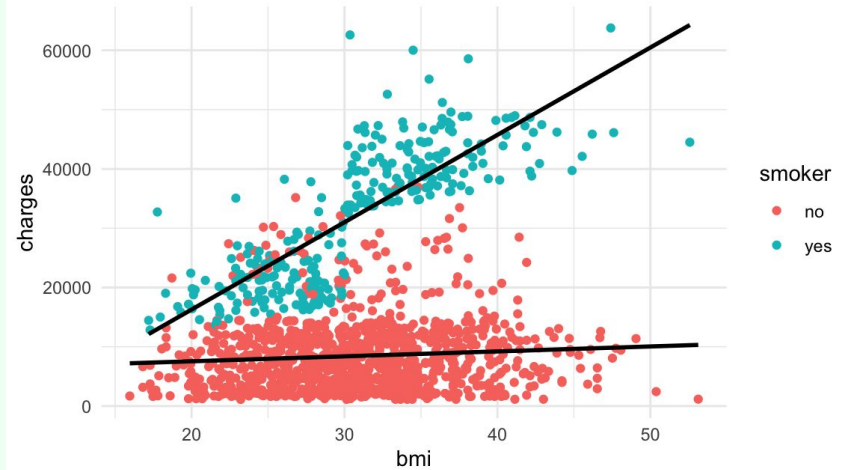
Do the higher charges in the Southeast region affect other variables?



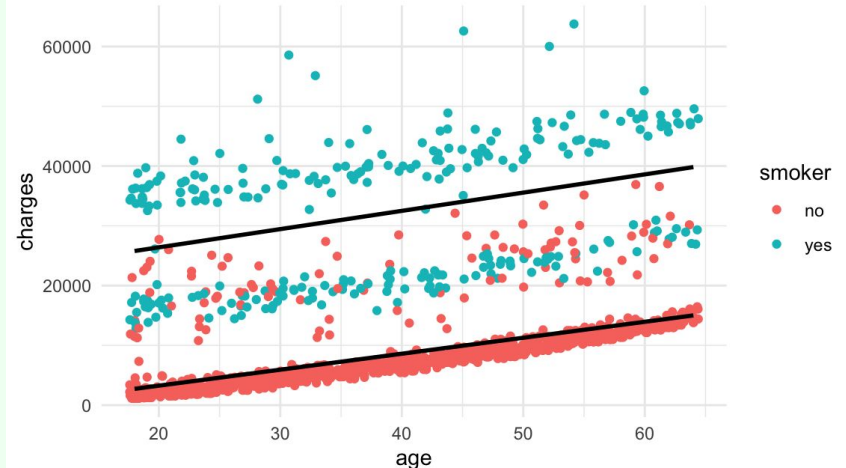
- ❑ Insurance charges are significantly higher for smokers considering they only make 20% of sample population
- ❑ IQR of charges for all counts of children do not have significant variation
- ❑ BMI's that show obesity have the largest ranges of charges generally on the higher end
- ❑ Southeast region does have higher insurance charges in their reasonable range
- ❑ Overall, Southeast region does not show effect from smoking status, number of children, or BMI

- Smokers generally have significantly higher charges than non smokers
- BMI and smokers together show steep linear trend
- Weak but evident linear trend between age and charges

Scatterplot of Charges vs BMI



Scatterplot of Charges vs Age



02

Fitting Our Model



Models

```
summary(full_model)
```

Call:

```
lm(formula = charges ~ age + sex + bmi + children + smoker +  
    region, data = data)
```

Residuals:

Min	1Q	Median	3Q	Max
-11305.1	-2850.3	-979.9	1395.0	29992.8

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	-11936.56	988.23	-12.079	< 2e-16 ***
age	256.76	11.91	21.555	< 2e-16 ***
sexmale	-129.48	333.20	-0.389	0.697630
bmi	339.25	28.61	11.857	< 2e-16 ***
children	474.82	137.90	3.443	0.000593 ***
smokeryes	23847.33	413.35	57.693	< 2e-16 ***
regionnorthwest	-349.23	476.82	-0.732	0.464053
regionsoutheast	-1035.27	478.87	-2.162	0.030804 *
regionsouthwest	-960.08	478.11	-2.008	0.044836 *

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 6064 on 1328 degrees of freedom
Multiple R-squared: 0.7507, Adjusted R-squared: 0.7492
F-statistic: 500 on 8 and 1328 DF, p-value: < 2.2e-16

```
summary(reduced_model)
```

Call:

```
lm(formula = charges ~ age + bmi + children + smoker + region,  
    data = data)
```

Residuals:

Min	1Q	Median	3Q	Max
-11366.5	-2841.4	-976.9	1364.0	29936.4

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	-11987.42	979.21	-12.242	< 2e-16 ***
age	256.88	11.91	21.577	< 2e-16 ***
bmi	338.73	28.57	11.856	< 2e-16 ***
children	473.86	137.83	3.438	0.000604 ***
smokeryes	23835.21	412.04	57.847	< 2e-16 ***
regionnorthwest	-348.25	476.66	-0.731	0.465152
regionsoutheast	-1034.63	478.71	-2.161	0.030852 *
regionsouthwest	-959.42	477.95	-2.007	0.044914 *

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 6062 on 1329 degrees of freedom
Multiple R-squared: 0.7507, Adjusted R-squared: 0.7494
F-statistic: 571.8 on 7 and 1329 DF, p-value: < 2.2e-16

Anova

Analysis of Variance Table

Model 1: charges ~ age + bmi + children + smoker + region

Model 2: charges ~ age + sex + bmi + children + smoker + region

	Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)
1	1329	4.8844e+10				
2	1328	4.8838e+10	1	5553651	0.151	0.6976

- ❑ Since the p value is > 0.05 , then that leads us to be able to assume that we can reduce our model to not utilize the “sex” variable

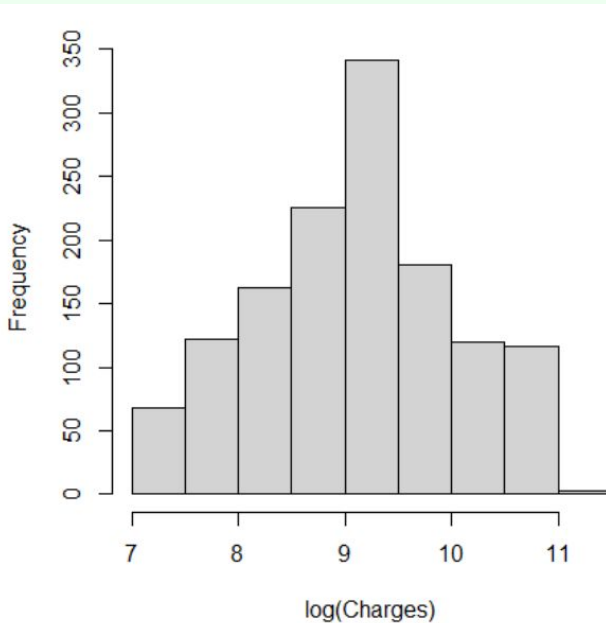
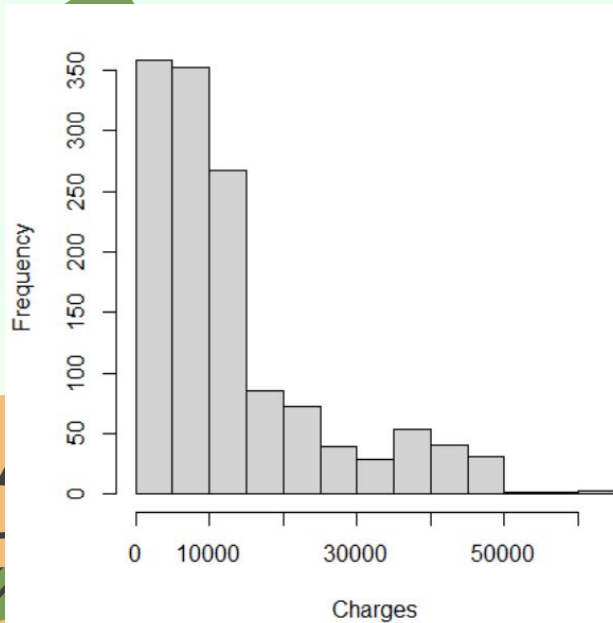
03



Residual and Influential Analysis



Transforming Our Data

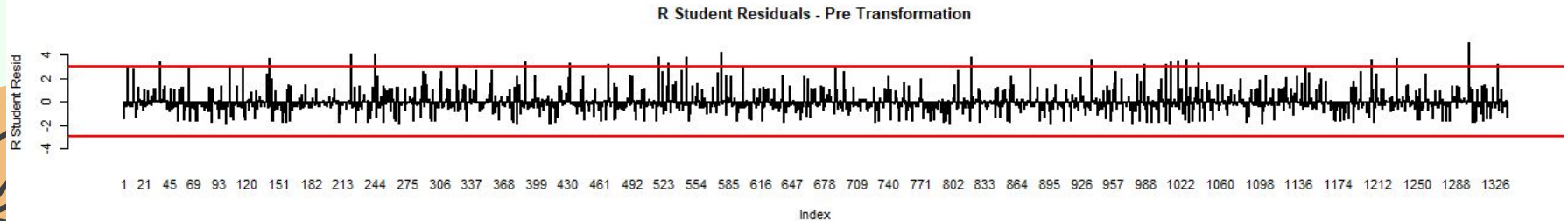
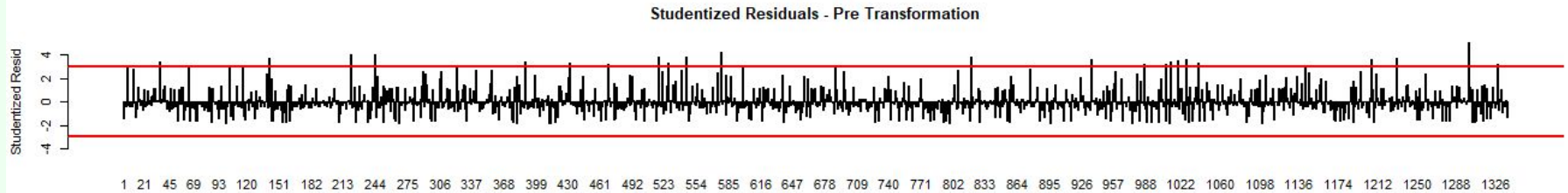
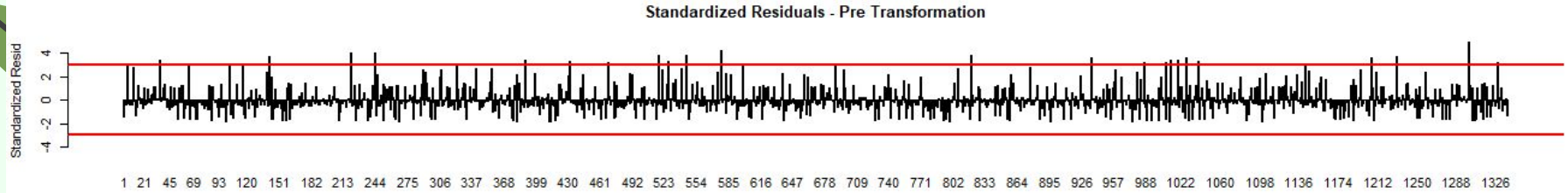


```
> AIC(fit, fit_log)
      df      AIC
fit     10 27096.2154
fit_log 10 -594.9611
> BIC(fit, fit_log)
      df      BIC
fit     10 27148.1973
fit_log 10 -542.9792
```

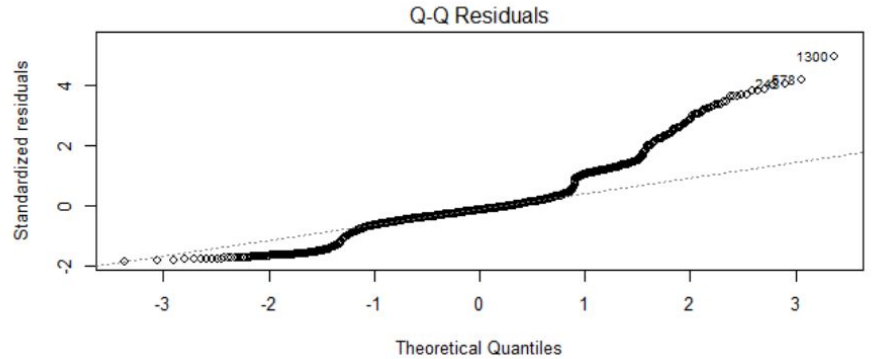
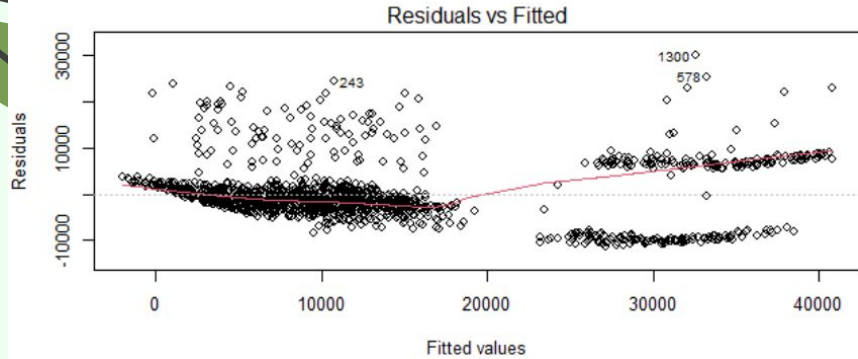
Applying the log function to our charges gave us an approximately normal distribution

Low AIC and BIC values

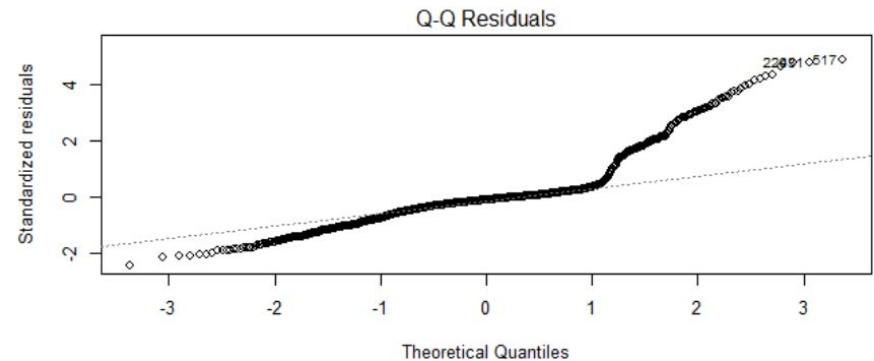
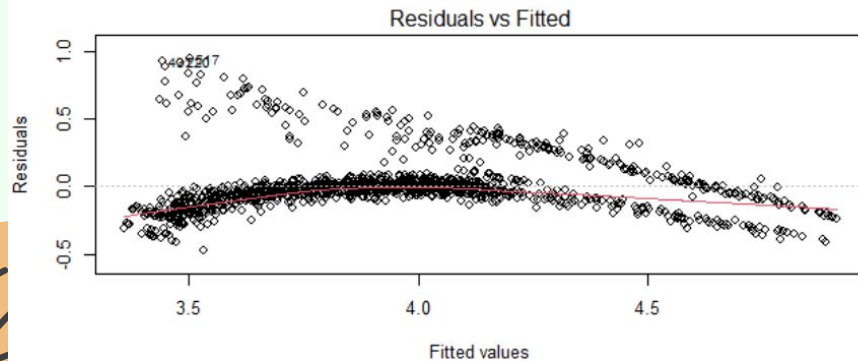
Pre-Transformation Residuals



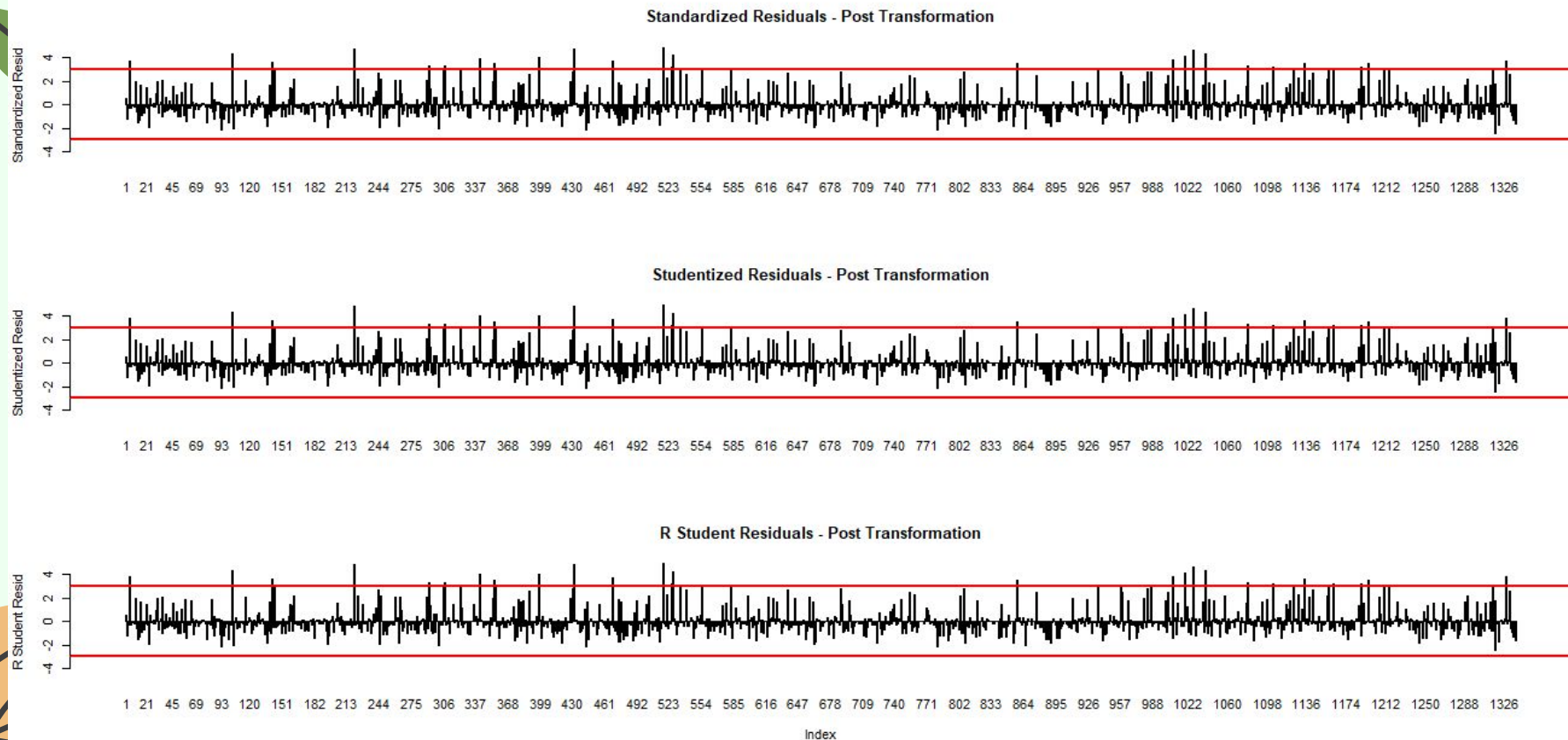
Pre-Transformation



Post-Transformation

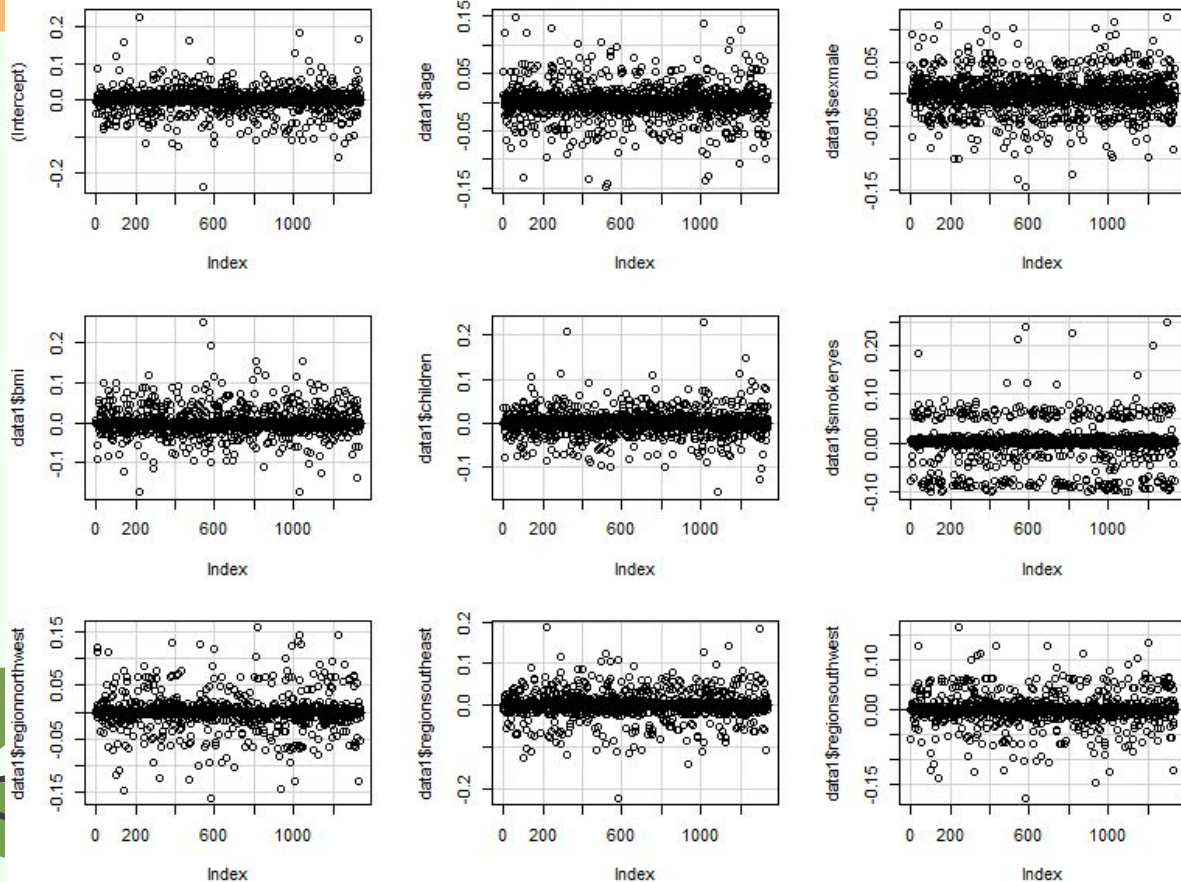


Post-Transformation Residuals



Influential Analysis

dfbetas Plots



A pattern that repeats in most of our graphs is the concentration of observations. This tells us that the majority of points are of comparable influences.

This is the case for most graphs except for the one corresponding to the "smoker" variable. This discrepancy suggests a significant distinction in influence based on whether the user is categorized as a smoker or non-smoker.

Influential Analysis

Most influential observations:

544 - 54 y/o Female, 47.41BMI, No children, Smoker, Southeast, \$63,770.42

1300 - 45 y/o Male, 30.36BMI, No children, Smoker, Southeast, \$62,592.87

Potential Leverage Points:

439 - 52 y/o Female, 46.75BMI, Five children, Nonsmoker, Southeast, \$12,592.53

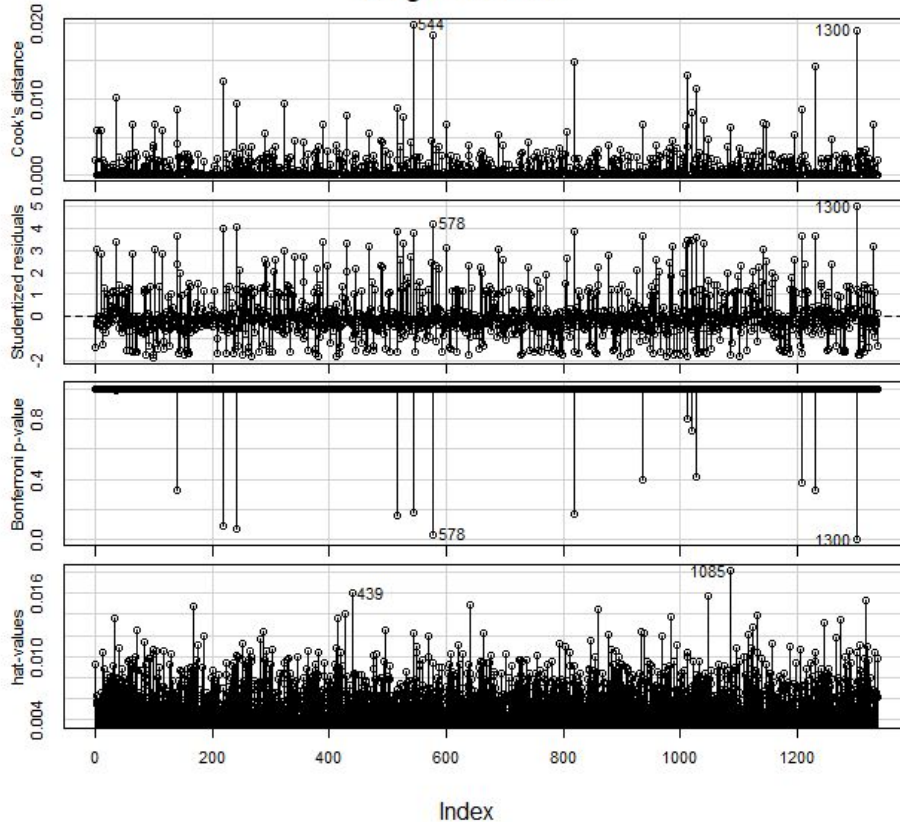
1085 - 39 y/o Female, 18.3BMI, Five children, Smoker, Southwest, \$19,023.26

Our VIFs are small so there is most likely no multicollinearity in our data

```
> vif(fit)
```

	GVIF	Df	$GVIF^{1/(2*Df)}$
data1\$age	1.016794	1	1.008362
data1\$sex	1.008944	1	1.004462
data1\$bmi	1.106742	1	1.052018
data1\$children	1.004017	1	1.002006
data1\$smoker	1.012100	1	1.006032
data1\$region	1.099037	3	1.015864

Diagnostic Plots





Conclusion

Pre-Transformation

Pre-Transformation, our numerical-based model was:

$$\text{Charges} = -12098.82 + 257.77(\mathbf{age}) + 321.87(\mathbf{bmi}) + 472.98(\mathbf{children}) + 23810.40(\mathbf{smoker})$$

Post-Transformation

Post-Transformation, our numerical-based model was:

$$\log(\text{Charges}) = 6.99 + 0.03(\mathbf{age}) + 0.01(\mathbf{bmi}) + 0.1(\mathbf{children}) + 1.54(\mathbf{smoker})$$

Furthermore, we noticed that sex was not a significant predictor for medical insurance costs

Sidenote: we were not able to include the categorical regions in here as R splits that variable up into various ones

The background is a light green color with various medical and nature-themed icons scattered around. In the top left, there is a white circular icon with a black line. Next to it is a red and white capsule. To the right is a green and white capsule. Further right is a white thermometer with a red tip. In the top right corner, there is a blue DNA double helix and a brown leaf with green leaves. On the left side, there is a red and white stethoscope. In the bottom left, there is a green leafy branch. In the bottom center, there is a white pillbox with several blue and red pills. In the bottom right, there is a red bandage, a small blue heart, and a clipboard with a blue clip and a white sheet of paper with black lines.

Thank you!