Homework 4 (100 Points)

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Exercise 1 (40pts)

In this exercise we will implement a Natural Language Processing (NLP) system using binary logistic regression.

The data you will be working with comes from the Yelp Dataset. Each line is a review that consists of a label (0 for negative reviews and 1 for positive reviews) and a set of words.

```
1 i will never forget this single breakfast experience in mad...
0 the search for decent chinese takeout in madison continues ...
0 sorry but me julio fell way below the standard even for med...
1 so this is the kind of food that will kill you so there s t...
```

In order to transform the set of words into vectors, we will rely on a method of feature engineering called word embeddings. Rather than simply indicating which words are present, word embeddings represent each word by "embedding" it in a low-dimensional vector space which may carry more information about the semantic meaning of the word. (for example in this space, the words "King" and "Queen" would be close).

word2vec.txt contains the word2vec embeddings for about 15 thousand words. Not every word in each review is present in the provided word2vec.txt file. We can treat these words as being "out of vocabulary" and ignore them.

Example

Let x_i denote the sentence "a hot dog is not a sandwich because it is not square" and let a toy word2vec dictionary be as follows:

```
hot 0.1 0.2 0.3
not -0.1 0.2 -0.3
sandwich 0.0 -0.2 0.4
square 0.2 -0.1 0.5
```

we would first trim the sentence to only contain words in our vocabulary: "hot not sandwich not square" then embed x_i into the feature space:

```
 \$\$ \ \phi 2(x_i)) = \frac{1}{5} \ (word2vec(\text{hot}) + 2 \cdot word2vec(\text{hot})) + word2vec(\text{text}\{not\}) + word2vec(\text{text}\{sandwich\}) + word2vec(\text{text}\{square\})) = \left[0.02 \cdot hspace\{2mm\} \ 0.06 \cdot hspace\{2mm\} \ 0.12 \cdot hspace\{2mm\} \ T \ \$ \right]
```

```
Part 1 (20pts)
```

a) Implement a function to trim out-of-vocabulary words from the reviews. Your function should return an nd array of the same dimension and dtype as the original loaded dataset. (10pts)

```
import csv
import numpy as np
VECTOR LEN = 300 # Length of word2vec vector
MAX WORD LEN = 64 # Max word length in dict.txt and word2vec.txt
#########
# We have provided you the functions for loading the tsv and txt
files. Feel #
# free to use them! No need to change them at all.
#########
def load tsv dataset(file):
   Loads raw data and returns a tuple containing the reviews and
their ratings.
   Parameters:
       file (str): File path to the dataset tsv file.
   Returns:
      An np.ndarray of shape N. N is the number of data points in
the tsv file.
      Each element dataset[i] is a tuple (label, review), where the
label is
      an integer (0 or 1) and the review is a string.
   dataset = np.loadtxt(file, delimiter='\t', comments=None,
encoding='utf-8',
                     dtype='l,0')
   return dataset
def load tsv dataset for embeddings(file):
   Loads raw data and returns a tuple containing the reviews and
their ratings.
   Parameters:
       file (str): File path to the dataset tsv file.
```

```
Returns:
        An np.ndarray of shape N. N is the number of data points in
the tsv file.
        Each element dataset[i] is a tuple (label, review), where the
label is
        an integer (0 or 1) and the review is a string.
    dataset = np.loadtxt(file, delimiter='\t', comments=None,
encoding='utf-8',
                     dtype=[('id', 'f8'), ('text', '0')])
    return dataset
def load feature dictionary(file):
    Creates a map of words to vectors using the file that has the
word2vec
    embeddings.
    Parameters:
        file (str): File path to the word2vec embedding file.
    Returns:
       A dictionary indexed by words, returning the corresponding
word2vec
        embedding np.ndarray.
    word2vec map = dict()
    with open(file) as f:
        read file = csv.reader(f, delimiter='\t')
        for row in read file:
            word, embedding = row[0], row[1:]
            word2vec map[word] = np.array(embedding, dtype=float)
    return word2vec map
def trim reviews(path to dataset):
    word2vec = load feature dictionary("./data/word2vec.txt")
    data = load tsv dataset(path to dataset)
    data cleaned=[]
    for text in data:
        row=[]
        tokens = text[1].split(" ")
        matching tokens = [token for token in tokens if token in
word2vec.keys()]
        row.append(text[0])
        row.append(" ".join(matching_tokens))
        data cleaned.append(row)
```

```
return data_cleaned

trim_train = trim_reviews("./data/train_small.tsv")
trim_test = trim_reviews("./data/test_small.tsv")
```

b) Implement the embedding and store it to a .tsv file where the first column is the label and the rest are the features from the embedding. Round all numbers to 6 decimal places. embedded_train_small.tsv and embedded_test_small.tsv contain the expected output of your function. (10pts)

```
def embed reviews(trimmed dataset):
    embedded dataset=[]
    word2vec = load feature dictionary("./data/word2vec.txt")
    for text in trimmed dataset:
        row=[]
        words = text[1].split()
        word vectors = [word2vec[word] for word in words if word in
word2vec1
        feature vector = np.round(np.mean(word vectors, axis=0),6)
        row.append(text[0])
        row.append(feature vector)
        embedded dataset.append(row)
    return embedded dataset
def save as tsv(dataset, filename):
    with open(filename, 'w+') as f:
        for row in dataset:
            f.write(str(row[0]) + '\t' + '\t'.join(map(str, row[1])) +
'\n')
    return
embedded train = embed reviews(trim train)
embedded test = embed reviews(trim test)
#et o=load tsv dataset for embeddings("./data/output/embedded train sm
all.tsv")
save as tsv(embedded train, "./data/output/embedded train small.tsv")
save_as_tsv(embedded_test, "./data/output/embedded_test_small.tsv")
Part 2 (20pts)
```

In this part we'll be implementing Gradient Descent for binary Logistic Regression Classifier.

Some rules:

- 1. Include an intercept term in your model. You must consider the bias term as part of the weight vector and not a separate term to keep track of.
- 2. Initialize all model parameters to 0
- 3. Use vector and matrix multiplication

```
The expected metrics.txt from the dataset with 500 epochs and 0.001 learning rate is:
error(train): 0.000000
error(test): 0.625000
We will be testing your code on other, larger datasets.
import numpy as np
def load tsv dataset(file):
    return np.loadtxt(file, delimiter='\t', encoding='utf-8')
def sigmoid(x):
    e = np.exp(x)
    return e / (1 + e)
def gd(theta, X, y, learning_rate):
    # TODO: implement in vector form
    grad = (1/len(y)) * np.dot(X.T, (sigmoid(np.dot(X, theta)) - y)) +
(theta/len(y))*theta
    theta -= learning rate * grad
    return theta
def train(theta, X, y, num_epoch, learning_rate):
    for epoch in range(num epoch):
        theta = gd(theta, X, y, learning rate)
    return theta
def predict(theta, X):
    # TODO: implement in vector form
    y pred = sigmoid(np.dot(X, theta))
    y pred[y pred >= 0.5] = 1
    y_pred[y_pred < 0.5] = 0
    return y_pred
def compute error(y pred, y):
    # TODO: implement in vector form
    return np.mean(np.abs(y - y pred))
def write_metrics(train_err, test_err, metrics_out):
    with open(metrics_out, 'w+') as f:
        w = "error(train): " + "{:.6f}".format(train_err) + "\n"
        w += "error(test): " + "{:.6f}".format(test err) + "\n"
```

```
f.write(w)
    return
def logistic_reg(formatted_train, formatted test, metrics out,
num epochs, learning rate):
    X = load_tsv_dataset(formatted_train)
    y = X[:, 0]
    X = np.insert(X[:, 1:], 0, 1, axis=1)
    theta = np.zeros(X.shape[1])
    learned theta = train(theta, X, y, num epochs, learning rate)
    train pred = predict(learned theta, X)
    train err = compute error(train pred, y)
    X test = load tsv dataset(formatted test)
    y test = X test[:, 0]
    X_{\text{test}} = \text{np.insert}(X_{\text{test}}[:, 1:], 0, 1, axis=1)
    test pred = predict(learned theta, X test)
    test err = compute error(test pred, y test)
    write metrics(train err, test err, metrics out)
    return
logistic reg("./data/output/embedded train small.tsv",
"./data/output/embedded_test_small.tsv", "./data/output/metrics.txt",
500, 0.01)
with open('./data/output/metrics.txt', 'r') as f:
    print(f.read())
error(train): 0.000000
error(test): 0.625000
```

Exercise 2 (60 Points)

For this exercise, recall the lab's kaggle competition where you predicted if a cookie was defective or not (assuming you attend labs). For more information on the data and details, click here to view the competition and data details

However, what we will do differently here is check if two of the features were correlated. This is important because if two features are correlated, then we can remove one of the features and still get the same information. This is part of the feature selection process.

Loading in the Data

For this exercise, we are interested in the two features Texture Hardness and Texture Chewiness and want to see if they are related. To do this, we will load in the data and create a scatter plot of the two features.

a) (1 Point) Download the kaggle data from the link above and read in the csv files cookie train.csv and cookie test.csv

```
#Code in this cell
import pandas as pd
cookie_test = pd.read_csv("data/cookie/cookie_test.csv")
cookie train = pd.read csv("data/cookie/cookie train.csv")
```

Combining Training and Testing Data

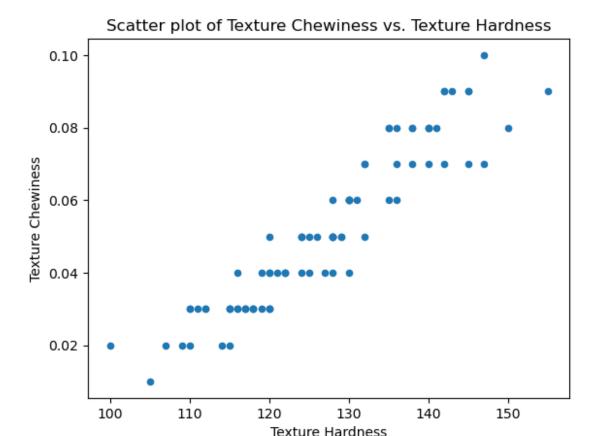
The lab did not have a lot of cookies for the training set, so what we will do is combine the training and testing data together without any loss of generality.

b) (3 Point) Combine the training and testing data together into a single dataframe called cookie_data and plot the scatter plot for Texture Chewiness (our y or our response variable) and Texture Hardness (our x or our explanatory variable). Make sure to label your axes and give your plot a title.

```
#Code in this cell
import matplotlib.pyplot as plt

cookie_data = pd.concat([cookie_train, cookie_test],
ignore_index=True)

cookie_data.plot.scatter(x='Texture Hardness', y='Texture Chewiness')
plt.xlabel('Texture Hardness')
plt.ylabel('Texture Chewiness')
plt.title('Scatter plot of Texture Chewiness vs. Texture Hardness')
plt.show()
```



Checking If We Can Do Linear Regression

It is important to check all requirements/assumptions for linear regression are met before we do linear regression. One of the requirements is that the explanatory variable and response variable are linearly related, which we checked by noticing the scatter plot was linear. However, there is much more to check!

Checking for Normality

It is important to check that the response variable is normally distributed. This is important for hypothesis testing and relates to the idea of linear regression being an unbiased, minimum variance estimator (if you're interested in this, check out the course CAS MA 582).

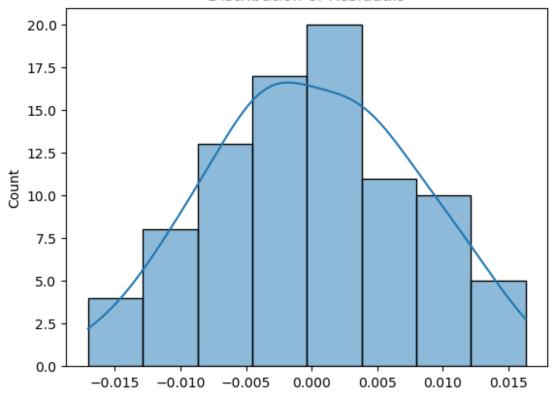
c) (5 Points) Plot an appropriate graph to determine if the response variable Texture Chewiness follows the normality assumption. State whether the normality assumption is met or not. If it is not met, state what can we do to fix this.

```
import seaborn as sns
import matplotlib.pyplot as plt
import statsmodels.api as sm

X = cookie_data['Texture Hardness']
y = cookie_data['Texture Chewiness']
```

```
model = sm.OLS(y, sm.add_constant(X)).fit()
y_pred = model.predict(sm.add_constant(X))
residuals = y - y_pred
sns.histplot(residuals, kde=True)
plt.title('Distribution of Residuals')
plt.show()
```

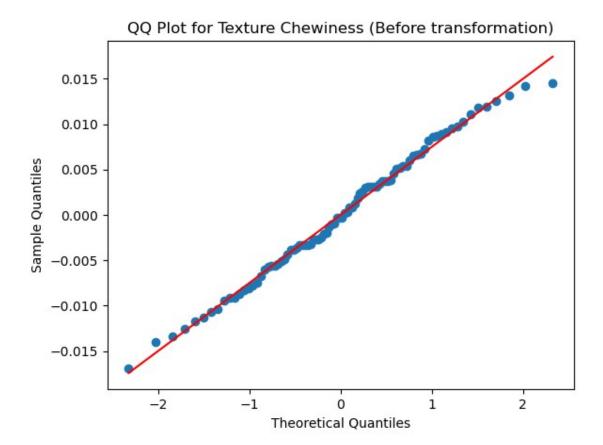
Distribution of Residuals



```
import scipy as sc
theoretical_quantiles = sc.stats.norm.ppf(np.linspace(0.01, 0.99,
num=len(residuals)))
sample_quantiles = np.quantile(residuals, np.linspace(0.01, 0.99,
num=len(residuals)))

model1 = sm.OLS(sample_quantiles,
sm.add_constant(theoretical_quantiles)).fit()

plt.scatter(theoretical_quantiles, sample_quantiles)
plt.plot(theoretical_quantiles, model1.predict(), color='red')
plt.xlabel('Theoretical Quantiles')
plt.ylabel('Sample Quantiles')
plt.title('QQ Plot for Texture Chewiness (Before transformation)')
plt.show()
```



Plotting the histogram with the kde curve shows us that the original data is normally distributed to an extent.

Checking for Homoscedasticity

It is important to check that the variance of the response variable is the same across all values of the explanatory variable. Otherwise, the confidence around a predicted value of the response variable will vary depending on the explanatory variable value.

d) (**5 Points**) Plot an appropriate graph to determine if the response variable Texture Chewiness follows the homoscedasticity assumption. State whether the homoscedasticity assumption is met or not. If it is not met, state what can we do to fix this.

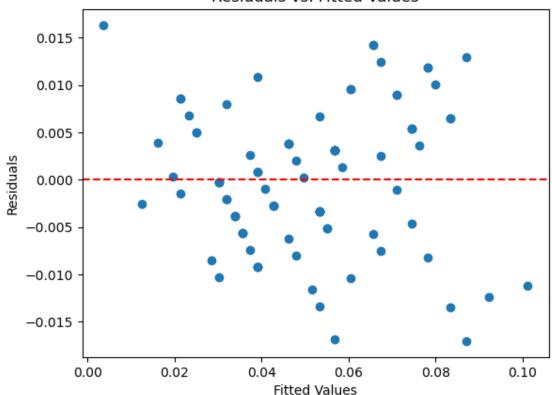
```
X = cookie_data['Texture Hardness']
y = cookie_data['Texture Chewiness']
model = sm.OLS(y, sm.add_constant(X)).fit()

y_pred = model.predict(sm.add_constant(X))
residuals = y - y_pred
plt.scatter(y_pred, residuals)

plt.axhline(y=0, color='red', linestyle='--')
plt.title('Residuals vs. Fitted Values')
```

```
plt.xlabel('Fitted Values')
plt.ylabel('Residuals')
plt.show()
```





In order to find the linear relationship between the predictor and the dependent variables, I used OLS (Ordinary Least Squares) model. The Residuals are the differences between the actual values of the dependent variable and the predicted values whichh I have plotted to find out if the data follows homoscedasticity.

From the above plot, we can see that the residual vs fitted plot is not completely random along the y=0 line. so the assumption of homoscedasticity doesn't hold true.

Checking for Independence

It is important to check that the explanatory variable and response variable are independent. Otherwise, the model isn't really best fit with a line. To do so, we will use the Durbin-Watson test.

e) (**5 Points**) Use the Durbin-Watson test to determine if the explanatory variable and response variable are independent. State whether if the independence assumption is met or not. If it is not met, state what can we do to fix this. You may use any library to do this, and for more information, check out this link.

```
#Code in this cell
X = cookie_data['Texture Hardness']
y = cookie_data['Texture Chewiness']
model = sm.OLS(y, X).fit()

test_result = sm.stats.stattools.durbin_watson(model.resid)
print(test_result)
if test_result < 1.5:
    print("Positive autocorrelation")
elif test_result > 2.5:
    print("Negative autocorrelation")
else:
    print("No autocorrelation")

2.6104905730803654
Negative autocorrelation
```

We see that the Texture Hardness is negatively correlted with Texture Chewiness which means that if Texture Hardness increases, Texture Chewiness decreases

Adjusting the Model

If you think any of the assumptions are not met, then you should respectively fix the model.

f) (**6 Points**) If you think atleast one of the assumptions were not met, then adjust the model how you described you would. Otherwise, proceed to the next step and in the cell below, write "No adjustments were made to the model". However, if you do adjust the model, then in the cell below, justify why your adjustments is appropriate (i.e. plot the new graphs and statistics found above). You may add as many cells as you like.

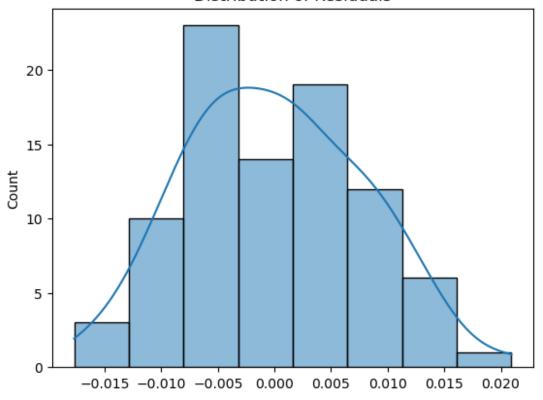
```
#cookie_data['Texture Hardness'] = np.power(cookie_data['Texture
Hardness'], 0.5)
cookie_data['Texture Hardness'] = np.log(cookie_data['Texture
Hardness'])
model = sm.OLS(cookie_data['Texture Chewiness'],
sm.add_constant(cookie_data['Texture Hardness'])).fit()
print(model.summary())
```

OLS Regression Results

```
Dep. Variable: Texture Chewiness R-squared:
0.872
Model: OLS Adj. R-squared:
0.870
Method: Least Squares F-statistic:
584.7
Date: Thu, 13 Apr 2023 Prob (F-statistic):
4.02e-40
```

```
Time:
                     21:47:23 Log-Likelihood:
302.62
                          88 AIC:
No. Observations:
-601.2
Df Residuals:
                             BIC:
                          86
-596.3
Df Model:
                          1
Covariance Type: nonrobust
                coef std err t
                                        P>|t|
[0.025 0.975]
------
              -1.0302 0.045 -23.036 0.000
const
1.119 -0.941
Texture Hardness 0.2235 0.009 24.182 0.000
0.205 0.242
______
_____
                       1.106 Durbin-Watson:
Omnibus:
2.353
Prob(Omnibus):
                       0.575 Jarque-Bera (JB):
1.125
Skew:
                       0.162 Prob(JB):
0.570
Kurtosis:
                       2.550 Cond. No.
269.
______
[1] Standard Errors assume that the covariance matrix of the errors is
correctly specified.
X = cookie_data['Texture Hardness']
y = cookie_data['Texture Chewiness']
model = sm.OLS(y, sm.add constant(X)).fit()
y pred = model.predict(sm.add constant(X))
residuals = y - y_pred
sns.histplot(residuals, kde=True)
plt.title('Distribution of Residuals')
plt.show()
```

Distribution of Residuals

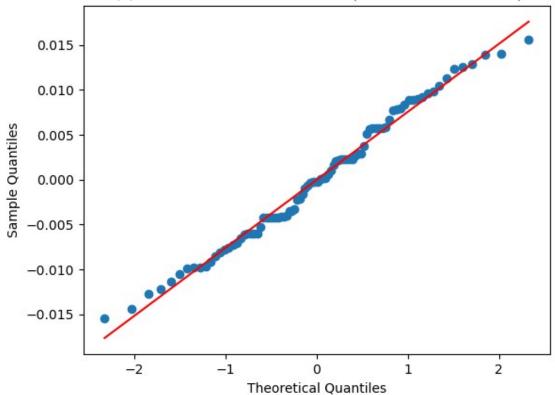


```
theoretical_quantiles = sc.stats.norm.ppf(np.linspace(0.01, 0.99,
num=len(residuals)))
sample_quantiles = np.quantile(residuals, np.linspace(0.01, 0.99,
num=len(residuals)))

model1 = sm.OLS(sample_quantiles,
sm.add_constant(theoretical_quantiles)).fit()

plt.scatter(theoretical_quantiles, sample_quantiles)
plt.plot(theoretical_quantiles, model1.predict(), color='red')
plt.xlabel('Theoretical Quantiles')
plt.ylabel('Sample Quantiles')
plt.title('QQ Plot for Texture Chewiness (After transformation)')
plt.show()
```

QQ Plot for Texture Chewiness (After transformation)



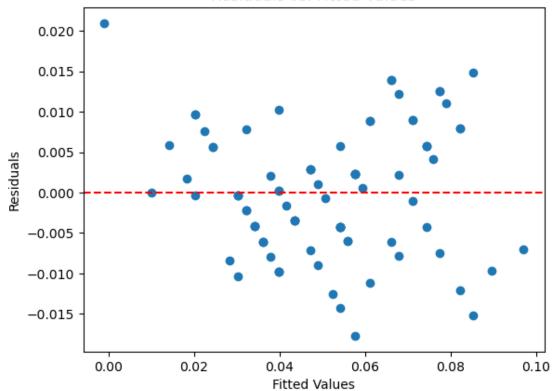
```
X = cookie_data['Texture Hardness']
y = cookie_data['Texture Chewiness']
model = sm.OLS(y, sm.add_constant(X)).fit()

y_pred = model.predict(sm.add_constant(X))
residuals = y - y_pred
plt.scatter(y_pred, residuals)

plt.axhline(y=0, color='red', linestyle='--')

plt.title('Residuals vs. Fitted Values')
plt.xlabel('Fitted Values')
plt.ylabel('Residuals')
```

Residuals vs. Fitted Values



After using power log transformation, as it worked better than the other methods I tried, like sqrt,log,boxcox transformations, we can see from the histogram chart of the residuals, that the outliers have been reduced. And from the model summary, we can see that the Durbin-Watson score is 2.353, which is greater than 1.5 and lesser than 2.5, so there is no autocorrelation

Hypothesis Testing

Now that we have a model that is appropriate, we can do hypothesis testing. We will be using the Linear Regression t-test to test if the explanatory variable is a significant predictor of the response variable.

g) (5 Points) Use the Linear Regression t-test to determine if the explanatory variable Texture Hardness is a significant predictor of the response variable Texture Chewiness. State whether if the explanatory variable is a significant predictor of the response variable or not using an alpha significance level of $\alpha = 0.05$, the null hypothesis $\beta_{hardness} = 0$ and alternative hypothesis $\beta_{hardness} \neq 0$. If it is not, state what can we do to fix this. You may use any library to do this.

```
#Your code in this cell
X = sm.add_constant(cookie_data['Texture Hardness'])
y = cookie_data['Texture Chewiness']
model = sm.OLS(y, X).fit()
```

```
t value = model.tvalues[1]
p value = model.pvalues[1]
p gt abs t value = float(model.summary().tables[1].data[1][4])
print("t-value:" +str(t_value))
print("p-value:" +str(p value))
print("p>|t|:" +str(p_gt_abs_t_value))
alpha = 0.05
if p_gt_abs_t_value < alpha:</pre>
    print("The feature variable 'Texture Hardness' is a significant
predictor of the predictor variable 'Texture Chewiness'.")
else:
    print("The feature variable 'Texture Hardness' is not a
significant predictor of the predictor variable 'Texture Chewiness'.")
t-value:24.18151553521705
p-value: 4.017398526064704e-40
p>|t|:0.0
The feature variable 'Texture Hardness' is a significant predictor of
the predictor variable 'Texture Chewiness'.
```

Looking at the summary above, we see that the p-value (P>|t|) of Texture Hardness using the Texture Chewiness as the dependent variable is 0.000.

This means that the p-value is very small, smaller than the alpha significance level of $\alpha = 0.05$.

This is strong evidence against the null hypothesis and shows that the variable Texture Hardness is a significant predictor of the response variable Texture Chewiness. Hence, the null hypothesis $\beta_{hardness}$ =0 can be rejected.

Adding a New Feature

Now that we have determined that the explanatory variable is a significant predictor of the response variable, you noticed the fit isn't 100% perfect still (e.g. your model is not capturing all the variation in the data). You think that there is another feature that can help improve the model. You think that the feature Taste Sweetness can help improve your model. You want to test if this is true. You may assume that all the assumptions are met for this new model to conduct linear regression.

h) (10 Points) Now that you have two features to predict your response variable, you will have to use multiple linear regression. Use the statsmodels library to do multiple linear regression. Then, determine if the explanatory variables Texture Hardness and Taste Sweetness are significant predictors of the response variable Texture Chewiness using an alpha significance level of α =0.001, the null hypothesis $\beta_{hardness}$ =0 and $\beta_{sweetness}$ =0 and alternative hypothesis $\beta_{hardness}$ ≠0 and $\beta_{sweetness}$ ≠0. If it is not, state what can we do to fix this. You may use any library to do this.

Moreover, justify whether the overall model is significant or not using the F-test with the same α given above. If it is not, state what can we do to fix this.

You may use the transformed model made (if one was made) from the previous parts and add this new feature to it.

```
#Your code in this cell
X = sm.add constant(cookie data[['Texture Hardness', 'Taste
Sweetness'11)
y = cookie_data['Texture Chewiness']
model = sm.OLS(y, X).fit()
alpha = 0.001
print(model.pvalues[1])
p_gt_abs_t_value_hardness = float(model.summary().tables[1].data[1]
[4])
print("p>|t| for hardness:" +str(p qt abs t value hardness))
if p gt abs t value hardness < alpha:</pre>
    print("The feature variable 'Texture Hardness' is a significant
predictor of the predictor variable 'Texture Chewiness'.")
    print("The feature variable 'Texture Hardness' is not a
significant predictor of the predictor variable 'Texture Chewiness'.")
p gt abs t value sweetness = float(model.summary().tables[1].data[2]
[4])
print("p>|t| for sweetness:" +str(p gt abs t value sweetness))
if p_gt_abs_t_value_sweetness < alpha:</pre>
    print("The feature variable 'Taste Sweetness' is a significant
predictor of the predictor variable 'Texture Chewiness'.")
    print("The feature variable 'Taste Sweetness' is not a significant
predictor of the predictor variable 'Texture Chewiness'.")
if p qt abs t value hardness < alpha and p qt abs t value sweetness <</pre>
alpha:
    print("Both feature variables are significant predictors of the
predictor variable.")
else:
    print("At least one of the explanatory variables is not a
significant predictor of the predictor variable.")
p value f = model.f pvalue
if p value f < alpha:</pre>
    print("The overall model is significant.")
    print("The overall model is not significant.")
```

```
3.3802313909162803e-15
p>|t| for hardness:0.0
```

The feature variable 'Texture Hardness' is a significant predictor of the predictor variable 'Texture Chewiness'.

p>|t| for sweetness:0.0

The feature variable 'Taste Sweetness' is a significant predictor of the predictor variable 'Texture Chewiness'.

Both feature variables are significant predictors of the predictor variable.

The overall model is significant.

The p-value of the model is 7.312011412914803e-07, which is smaller than the alpha significance level of $\alpha = 0.001$. This is strong evidence that the overall model is significant.

Interpretation

Covariance Type:

It is important to be able to interpret the results of the model. This is important because it allows us to understand the model and how it works. It also allows us to understand the relationship between the predictors and the response variable, and this applies to not just linear regression, but all learning models.

i) (10 Points) Interpret the coefficients of the model made in part h. What do they mean? What do they represent? What do they tell us about the relationship between the predictors and the response variable? If your code output in part h does not provide these results, you are welcome to use any library here to do so.

```
#Your code in this cell if you need it
print(model.summary())
```

OLS Regression Results

```
Dep. Variable:
                   Texture Chewiness
                                       R-squared:
0.928
Model:
                                 0LS
                                       Adj. R-squared:
0.926
Method:
                       Least Squares F-statistic:
547.4
                    Thu, 13 Apr 2023
Date:
                                       Prob (F-statistic):
2.81e-49
Time:
                            21:47:24
                                       Log-Likelihood:
327.98
No. Observations:
                                       AIC:
                                  88
-650.0
                                       BIC:
Df Residuals:
                                  85
-642.5
Df Model:
                                   2
```

nonrobust

==========	=========	=======	=========		
========	coef	std err	t	P> t	
[0.025 0.		stu err	Ĺ	P> L	
const	-0.6036	0.062	-9.687	0.000	-
0.727 -0. Texture Hardne		0.013	9.595	0.000	
0.103 0.		0.013	3.333	0.000	
Taste Sweetnes		0.001	8.140	0.000	
0.008 0.	014 =======	:=======			======
Omnibus:		1.231	Durbin-Watso	on:	
1.920 Prob(Omnibus):		0.540	Jarque-Bera	(1D).	
0.800		0.340	Jai que-bei a	(JB):	
Skew:		-0.218	Prob(JB):		
0.670 Kurtosis:		3.169	Cond. No.		
564.		2.103	Cond. No.		
=========		=======	========		
=======					

Notes:

[1] Standard Errors assume that the covariance matrix of the errors is correctly specified.

The R-squared value of 0.928 indicates that the model explains 92.8% of the variance in the dependent variable. This is a high value, suggesting that the model is a good fit for the data.

The coefficients for the predictor variables indicate the strength and direction of their relationship with the dependent variable. The coefficient for Texture Hardness is 0.1295, indicating that for every one unit increase in Texture Hardness, Texture Chewiness increases by 0.1295 units, holding all other variables constant. The coefficient for Taste Sweetness is 0.0111, indicating that for every one unit increase in Taste Sweetness, Texture Chewiness increases by 0.0111 units, holding all other variables constant.

The p-values for both predictor variables are less than 0.05, indicating that they are statistically significant in predicting Texture Chewiness.

The Durbin-Watson value of 1.92 suggests that there is no significant autocorrelation in the residuals of the model.

The equation to predict Texture Chewiness based on the above multiple linear regression model is:

Texture Chewiness = -0.6036 + 0.1295(Texture Hardness) + 0.0111(Taste Sweetness)

Using Taste Bitterness over Taste Sweetness

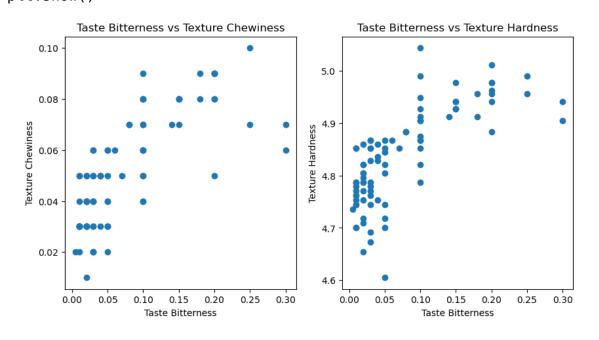
You think that the feature Taste Bitterness can help improve your model since Taste Sweetness wasn't it.

For this section, you will continue to use your fixed/adjusted features for Texture Hardness and Texture Chewiness.

j) (10 Points) Plot 2 scatter plots: Taste Bitterness vs your possibly adjusted Texture Chewiness and Taste Bitterness vs your possibly adjusted Texture Hardness. Given these plots, propose an appropriate model. Justify your answer by fitting it and seeing its accuracy (you can use any metric to prove so), showing appropriate models/graphs, and ensuring assumptions are satisfied (if any). You may use any library to do this.

```
fig, (ax1, ax2) = plt.subplots(1, 2, figsize=(10, 5))
```

```
ax1.scatter(cookie_data["Taste Bitterness"], cookie_data["Texture
Chewiness"])
ax1.set_xlabel("Taste Bitterness")
ax1.set_ylabel("Texture Chewiness")
ax1.set_title("Taste Bitterness vs Texture Chewiness")
ax2.scatter(cookie_data["Taste Bitterness"], cookie_data["Texture
Hardness"])
ax2.set_xlabel("Taste Bitterness")
ax2.set_ylabel("Texture Hardness")
ax2.set_title("Taste Bitterness vs Texture Hardness")
plt.show()
```

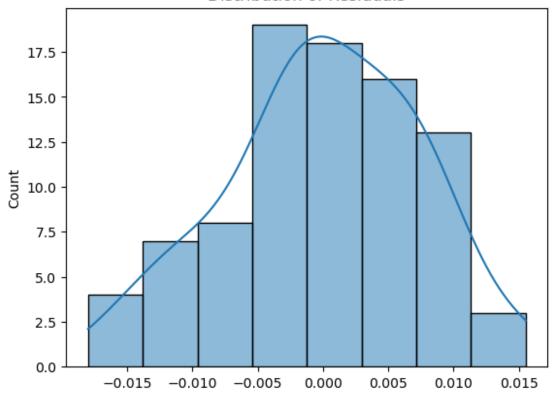


```
X = cookie_data[['Texture Hardness', 'Taste Bitterness']]
y = cookie_data['Texture Chewiness']
model = sm.OLS(y, sm.add_constant(X)).fit()

y_pred = model.predict(sm.add_constant(X))
residuals = y - y_pred

sns.histplot(residuals, kde=True)
plt.title('Distribution of Residuals')
plt.show()
```

Distribution of Residuals



```
cookie_data['Taste Bitterness'] = np.power(cookie_data['Taste
Bitterness'], 0.5)
cookie_data['Taste Bitterness'] = np.log(cookie_data['Taste
Bitterness'])
```

Drop any rows with missing or NaN values
cookie_data.dropna(inplace=True)

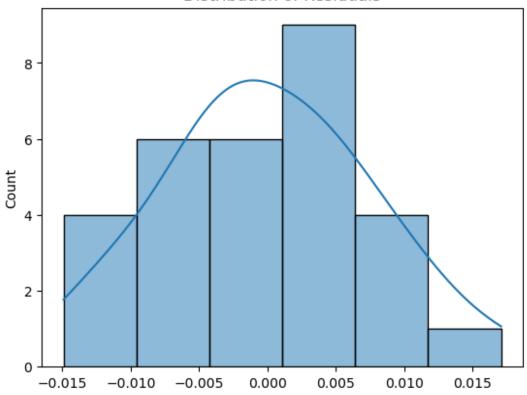
```
X = sm.add_constant(cookie_data[['Texture Hardness', 'Taste
Bitterness']])
y = cookie_data['Texture Chewiness']
model = sm.OLS(y, X).fit()
```

```
X = cookie_data[['Texture Hardness', 'Taste Bitterness']]
y = cookie_data['Texture Chewiness']
model = sm.OLS(y, sm.add_constant(X)).fit()

y_pred = model.predict(sm.add_constant(X))
residuals = y - y_pred

sns.histplot(residuals, kde=True)
plt.title('Distribution of Residuals')
plt.show()
```

Distribution of Residuals



```
#Code in this cell
X = cookie_data[['Texture Hardness', 'Taste Bitterness']]
y = cookie_data['Texture Chewiness']
model = sm.OLS(y, X).fit()

test_result = sm.stats.stattools.durbin_watson(model.resid)
print(test_result)
if test_result < 1.5:
    print("Positive autocorrelation")
elif test_result > 2.5:
    print("Negative autocorrelation")
else:
    print("No autocorrelation")
```

```
2.0136453560752363
No autocorrelation

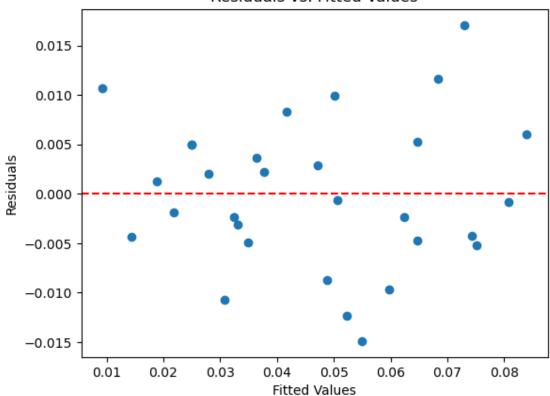
model = sm.OLS(y, sm.add_constant(X)).fit()

y_pred = model.predict(sm.add_constant(X))
residuals = y - y_pred
plt.scatter(y_pred, residuals)

plt.axhline(y=0, color='red', linestyle='--')

plt.title('Residuals vs. Fitted Values')
plt.xlabel('Fitted Values')
plt.ylabel('Residuals')
```

Residuals vs. Fitted Values



```
X = cookie_data[['Texture Hardness', 'Taste Bitterness']]
X = sm.add_constant(X)
X['Texture Hardness^2'] = X['Texture Hardness'] ** 2
X['Taste Bitterness^2'] = X['Taste Bitterness'] ** 2
y = cookie_data['Texture Chewiness']
model = sm.OLS(y, X).fit()
```

```
print(model.summary())
y pred = model.predict(X)
mse = np.mean((y - y pred)**2)
print(f"Mean Squared Error: {mse}")
                      OLS Regression Results
Dep. Variable: Texture Chewiness R-squared:
0.896
Model:
                           OLS Adj. R-squared:
0.879
Method:
                  Least Squares F-statistic:
53.81
Date:
                Thu, 13 Apr 2023 Prob (F-statistic):
6.34e-12
Time:
                       21:47:24 Log-Likelihood:
106.14
No. Observations:
                               AIC:
                           30
-202.3
Df Residuals:
                           25
                               BIC:
-195.3
Df Model:
Covariance Type:
             nonrobust
_____
                 coef std err t P>|t|
[0.025 0.975]
------
                  3.2808 2.462
                                   1.333 0.195
const
1.789 8.351
               -1.5057 1.021 -1.475 0.153
Texture Hardness
3.608
     0.596
                 -0.0056
                           0.027 -0.204
Taste Bitterness
                                             0.840
0.062
        0.051
Texture Hardness^2
                 0.1731
                           0.106 1.637
                                             0.114
0.045 0.391
Taste Bitterness^2
                 -0.0043
                           0.010 -0.432 0.669
0.025
         0.016
Omnibus:
                         0.667 Durbin-Watson:
2.049
```

0.717 Jarque-Bera (JB):

Prob(Omnibus):

```
0.647
                                0.316 Prob(JB):
Skew:
0.724
Kurtosis:
                                2.655
                                        Cond. No.
4.54e+04
Notes:
[1] Standard Errors assume that the covariance matrix of the errors is
correctly specified.
[2] The condition number is large, 4.54e+04. This might indicate that
there are
strong multicollinearity or other numerical problems.
Mean Squared Error: 4.948531009180381e-05
X = cookie data[['Texture Hardness', 'Taste Sweetness', 'Taste
Bitterness'll
X = sm.add constant(X)
X['Texture Hardness^2'] = X['Texture Hardness'] ** 2
X['Taste Bitterness^2'] = X['Taste Bitterness'] ** 2
X['Taste Sweetness^2'] = X['Taste Sweetness'] ** 2
y = cookie_data['Texture Chewiness']
model = sm.OLS(y, X).fit()
print(model.summary())
y pred = model.predict(X)
mse = np.mean((y - y_pred)**2)
print(f"Mean Squared Error: {mse}")
                            OLS Regression Results
Dep. Variable:
                   Texture Chewiness R-squared:
0.920
Model:
                                  0LS
                                        Adj. R-squared:
0.899
                        Least Squares F-statistic:
Method:
44.05
                     Thu, 13 Apr 2023 Prob (F-statistic):
Date:
1.77e-11
Time:
                             21:53:29
                                        Log-Likelihood:
110.07
No. Observations:
                                   30
                                        AIC:
-206.1
Df Residuals:
                                   23
                                        BIC:
-196.3
```

Df Model: 6

Covariance Type: no	nrobust
---------------------	---------

[0.025	0.975]	coef	std er	r t	P> t	
const 4.254	6 277	1.0117	2.54	5 0.397	0.695	-
Texture Hardness 2.721		-0.5287	1.06	-0.499	0.623	-
		-0.0020	0.01	-0.147	0.884	-
		-0.0036	0.02	-0.143	0.887	-
		0.0679	0.11	0.617	0.544	-
		-0.0033	0.00	-0.349	0.730	-
Taste Swee		0.0019	0.00	0.746	0.463	-
========		========		:========	========	=====
Omnibus: 1.761			0.490	Durbin-Watso	on:	
Prob(Omnibus): 0.617		0.783	Jarque-Bera (JB):			
Skew: 0.735		0.164	Prob(JB):			
Kurtosis: 5.33e+04		2.379	Cond. No.			
=======	=======			=========		=====

Notes:

=======

- [1] Standard Errors assume that the covariance matrix of the errors is correctly specified.
- [2] The condition number is large, 5.33e+04. This might indicate that there are

strong multicollinearity or other numerical problems.

Mean Squared Error: 3.807088056728153e-05

After adding Taste Bitterness instead of Taste Sweetness for predicting Texture Chewiness, I plotted the histogram of the residuals, which showed that the values were not normalized. So, I applied the same power log transformation to Taste Bitterness. Plotting the residuals post which, shows a better normalized curve on the histogram, with no autocorrelation from Durbin-Watson.

The scatter plot for Taste Bitterness against both Texture Chewiness and Texture Hardness show a non linear relationship. hence, in order to capture the non-linearity, the squared terms of the exploratory variables are also needed in the regression model for predicting Texture Chewiness

For the model, I used multiple linear regression using the Ordinary Least Squares (OLS) method. The independent variables used in the regression are Texture Hardness, and Taste Bitterness, and their respective squared terms are also included in the model. The Mean Squared Error is 4.948531009180381e-05 which indicates good prediction by the model.

Adding Taste Sweetness to the independent variables list, to predict Taste Chewiness improved the r-squared score from 0.896 to 0.920, decreased the autocorrelation.

Bonus (20pts)

Lasso Regression is a modification of Regression that adds a regularization penalty to the the parameter β learned by the model. The loss function for Lasso is the following:

$$L(\beta) = \|y - X\beta\|^2 + \lambda \|\beta\| = \beta^T X^T X\beta - 2\beta^T X^T y + y^T y + \lambda \|\beta\|$$

Where λ is a tuning parameter specified by the user.

Using worksheet 17 as a guide:

- 1. create an animation like in part c) for Lasso Regression. [15pts]
- 2. create at least two more animations using different values for λ and the true β you used to generate the dataset (try steeper / less steep curves, and higher or close to zero intercepts). Briefly explain some key takeaways on how to tune λ . [5pts]

```
from PIL import Image as im
import matplotlib.pyplot as plt
TEMPFILE = "temp.png"
def snap(betas, losses):
    # Creating figure
    fig = plt.figure(figsize =(14, 9))
    ax = plt.axes(projection = '3d')
    ax.view init(20, -20)
    ax.set xlim(-5, 4)
    ax.set_xlabel(r'$\beta_0$')
    ax.set ylabel(r'$\beta 1$')
    ax.set ylim(-5, 4)
    ax.set zlim(0, 30000)
    # Creating plot
    ax.plot surface(b0, b1, cost, color='b', alpha=.7)
    ax.plot(np.array(betas)[:,0], np.array(betas)[:,1], losses, 'o-',
c='r', markersize=10, zorder=10)
    fig.savefig(TEMPFILE)
    plt.close()
    return im.fromarray(np.asarray(im.open(TEMPFILE)))
```

```
def snap model(beta):
    xplot = np.linspace(-10,10,50)
    vestplot = beta[0]+beta[1]*xplot
    fig, ax = plt.subplots()
    ax.plot(xplot, yestplot, 'b-', lw=2)
    ax.plot(xlin, y,'ro',markersize=4)
    ax.set xlim(-10, 10)
    ax.set ylim(min(y), max(y))
    fig.savefig(TEMPFILE)
    plt.close()
    return im.fromarray(np.asarray(im.open(TEMPFILE)))
def lasso loss(X, y, beta, lambd):
    y pred = np.dot(X, beta)
    residuals = y - y pred
    rss = np.sum(residuals**2)
    l1 norm = np.sum(np.abs(beta))
    loss = rss + lambd * l1 norm
    return loss
def lasso_gradient(X, y, beta, lambd):
    return 2 * X.T @ X @ beta - 2 * X.T @ y + lambd * np.sign(beta)
def lasso gradient descent(X, y, beta hat, learning rate, lambd,
epochs, images):
    losses = [lasso loss(X, y, beta hat, lambd)]
    betas = [beta hat]
    for in range(epochs):
        images.append(snap model(beta hat))
        beta hat = beta hat - learning rate * lasso gradient(X, y,
beta hat, lambd)
        losses.append(lasso loss(X, y, beta hat, lambd))
        betas.append(beta hat)
    return np.array(betas), np.array(losses)
beta = np.array([1, .5])
xlin = -10.0 + 20.0 * np.random.random(100)
X = np.column stack([np.ones((len(xlin), 1)), xlin])
y = beta[0]+(beta[1]*xlin)+np.random.randn(100)
beta start = np.array([-5, -2])
learning rate = 0.0002
lambd=0.\overline{0}5
images = []
betas, losses = lasso gradient descent(X, y, beta start,
```

```
learning_rate, lambd, 100, images)
images[0].save(
    'lasso model.gif',
    optimize=False,
    save all=True,
    append images=images[1:],
    loop=0,
    duration=200
)
beta = np.array([1, 0.005])
xlin = -10.0 + 20.0 * np.random.random(100)
X = np.column stack([np.ones((len(xlin), 1)), xlin])
y = beta[0]+(beta[1]*xlin)+np.random.randn(100)
beta start = np.array([-5, -2])
learning_rate = 0.0002
lambd=100
images = []
betas, losses = lasso gradient descent(X, y, beta start,
learning rate, lambd, 100, images)
images[0].save(
    'lasso model slope0.005 lambda100.gif',
    optimize=False,
    save all=True,
    append images=images[1:],
    loop=0,
    duration=200
)
beta = np.array([10, 0.5])
xlin = -10.0 + 20.0 * np.random.random(100)
X = np.column stack([np.ones((len(xlin), 1)), xlin])
y = beta[0]+(beta[1]*xlin)+np.random.randn(100)
beta start = np.array([-5, -2])
learning rate = 0.0002
lambd=1000
images = []
betas, losses = lasso_gradient_descent(X, y, beta_start,
learning_rate, lambd, 100, images)
images[0].save(
    'lasso model intercept10 lambda1000.gif',
    optimize=False,
    save all=True,
    append images=images[1:],
    loop=0,
```

```
duration=200
)
```