```
In [1]:
         import pandas as pd
         from scipy.stats import ttest rel, wilcoxon
         from scipv.stats import pearsonr
         import numpy as np
         import statsmodels.api as sm
         from statsmodels.formula.api import ols
         # Reading in the data
         gzero_data = 'Gatorade_Zero_Taste_Test_Survey_Responses_2.csv'
         rawdata = pd.read_csv(gzero_data)
         # Separating the data for each product
         original = rawdata[rawdata['Product'] == 'Original']
         new formulation = rawdata[rawdata['Product'] == 'New Formulation']
         # Ensuring the data is aligned by Panelist ID
         original = original.sort values('Panelist ID')
         new formulation = new formulation.sort values('Panelist ID')
         # Performing paired t-test for each liking attribute
         liking attributes = [col for col in original.columns if 'Liking' in col]
         t test results = {}
         wilcoxon test results = {}
         for attribute in liking_attributes:
             t stat, p value = ttest rel(original[attribute], new formulation[attribute])
             t test results[attribute] = (t stat, p value)
         # Performing Wilcoxon signed-rank test to doublecheck results
             w stat, p value = wilcoxon(original[attribute], new formulation[attribute])
             wilcoxon test results[attribute] = (w stat, p value)
         # Displaying results
         t_test_results_df = pd.DataFrame.from_dict(t_test_results, orient='index', colum
         wilcoxon test results df = pd.DataFrame.from dict(wilcoxon test results, orient=
         print("Paired t-test Results:")
         print(t_test_results_df)
         print("\nWilcoxon Signed-Rank Test Results:")
         print(wilcoxon test results df)
         #Printing mean scores
         liking means = rawdata.groupby('Product')[liking attributes].mean().round(2)
         # Transposing the dataframe for better readability
         liking_means_transposed = liking_means.T
         liking means transposed.reset index(inplace=True)
         liking means transposed.columns = ['Liking Attribute', 'Original', 'New Formulat
         # Displaying the table
         print("\nLiking Means:")
         print(liking_means_transposed)
```

```
Paired t-test Results:
```

t_stat p_value Overall Liking 0.000000 1.000000

```
Flavor_Liking
                              -1.238540 0.219184
        Aroma Liking
                               0.454048 0.651039
        Color Liking
                              -0.136003 0.892165
        Orange Flavor Liking 0.474017 0.636795
                              -0.671139 0.504090
        Sweetness Liking
        Mouthfeel Liking
                              -0.056902
                                          0.954767
        Aftertaste Liking
                              -1.547787 0.125670
        Wilcoxon Signed-Rank Test Results:
                                        p value
                               w stat
        Overall Liking
                               1240.5
                                       0.990636
        Flavor_Liking
                               1086.0 0.199374
        Aroma Liking
                               1266.5
                                       0.643467
        Color Liking
                               1285.5
                                       0.872598
        Orange_Flavor_Liking
                               1151.5 0.593412
        Sweetness_Liking
                               1151.5 0.467149
        Mouthfeel Liking
                               1253.5 0.888011
        Aftertaste_Liking
                               1138.0 0.128262
        Liking Means:
                Liking Attribute Original New Formulation
        0
                  Overall_Liking
                                       5.29
                                                        5.29
        1
                  Flavor_Liking
                                       5.56
                                                        4.96
        2
                                                        5.36
                    Aroma_Liking
                                       5.14
        3
                    Color Liking
                                       5.42
                                                        5.36
           Orange Flavor Liking
                                       5.05
                                                        5.28
        5
                Sweetness Liking
                                       5.69
                                                        5.39
        6
               Mouthfeel Liking
                                       5.31
                                                        5.29
        7
              Aftertaste Liking
                                       5.60
                                                        4.94
        # Now working on JAR scores
In [2]:
         # Define the list of JAR attributes
         jar attributes = [col for col in rawdata.columns if 'JAR' in col]
         # Function to calculate the JAR category percentages
         def calculate jar percentages(df, attributes):
             jar_percentages = pd.DataFrame(index=attributes, columns=['Not Enough', 'JAF
             for attribute in attributes:
                 not enough = df[attribute].isin([1, 2]).sum()
                 just about right = df[attribute].isin([3]).sum()
                 too_much = df[attribute].isin([4, 5]).sum()
                 total = len(df)
                 jar_percentages.at[attribute, 'Not Enough'] = (not_enough / total) * 100
                 jar_percentages.at[attribute, 'JAR'] = (just_about_right / total) * 100
jar_percentages.at[attribute, 'Too Much'] = (too_much / total) * 100
             return jar percentages.round(2)
         # Separate the data for each product
         original_data = rawdata[rawdata['Product'] == 'Original']
         new formulation data = rawdata[rawdata['Product'] == 'New Formulation']
         # Calculate the JAR category percentages for each product
         original_jar_percentages = calculate_jar_percentages(original_data, jar_attribut
         new formulation jar percentages = calculate jar percentages(new formulation data
         original_jar_percentages, new_formulation_jar_percentages
```

JAR Too Much

Not Enough

```
Out[2]: (
         Flavor_JAR
                                31.25
                                        17.5
                                                51.25
         Aroma_JAR
                                41.25 21.25
                                                 37.5
         Color JAR
                                41.25
                                      18.75
                                                   40
         Orange_Flavor_JAR
                                46.25
                                       17.5
                                                36.25
         Sweetness_JAR
                                  40
                                          25
                                                   35
         Mouthfeel JAR
                                43.75 13.75
                                                 42.5
         Aftertaste JAR
                                41.25
                                          20
                                                38.75,
                           Not Enough
                                         JAR Too Much
         Flavor JAR
                                 42.5
                                          15
                                                 42.5
         Aroma JAR
                                 37.5 18.75
                                                43.75
         Color JAR
                                 37.5
                                          15
                                                 47.5
         Orange_Flavor_JAR
                                 47.5
                                        12.5
                                                   40
         Sweetness JAR
                                41.25 58.75
                                                    0
         Mouthfeel JAR
                                21.25
                                       26.25
                                                38.75
         Aftertaste JAR
                                38.75
                                        17.5
                                                43.75)
In [3]:
         #Finding Correlation for directional JAR Scores to Overall Liking
         # Transform JAR scores into weighted dummy variables for "Not Enough" and "Too M
         for attribute in jar attributes:
             rawdata[f'{attribute}_Not_Enough'] = rawdata[attribute].apply(lambda x: 1 if
             rawdata[f'{attribute}_Too_Much'] = rawdata[attribute].apply(lambda x: 1 if x
         # Create the formula for the regression model
         dummy variables = [f'{attr} Not Enough' for attr in jar attributes] + [f'{attr}
         formula = 'Overall_Liking ~ ' + ' + '.join(dummy_variables)
         # Function to fit the model for a specific product
         def fit_model_for_product(product_data):
             model = ols(formula, data=product data).fit()
             print(f"Model summary for {product data['Product'].iloc[0]}:")
             print(model.summary())
             print("\n")
             return model
         # Separate the data for each product
         original data = rawdata[rawdata['Product'] == 'Original']
         new formulation data = rawdata[rawdata['Product'] == 'New Formulation']
         # Fit the models for each product
         original_model = fit_model_for_product(original_data)
         new_formulation_model = fit_model_for_product(new_formulation_data)
         # Save the model summaries to text files
         with open('Original_Product_Multiple_Regression_Summary.txt', 'w') as f:
             f.write(original_model.summary().as_text())
         with open('New Formulation Product Multiple Regression Summary.txt', 'w') as f:
             f.write(new formulation model.summary().as text())
        Model summary for Original:
                                    OLS Regression Results
        Dep. Variable:
                               Overall_Liking
                                                R-squared:
                                                                                  0.143
        Model:
                                          0LS
                                                Adj. R-squared:
                                                                                 -0.042
        Method:
                                Least Squares
                                                F-statistic:
                                                                                 0.7739
        Date:
                             Fri, 31 May 2024
                                                Prob (F-statistic):
                                                                                  0.693
        Time:
                                     23:01:31
                                                Log-Likelihood:
                                                                               -179.22
        No. Observations:
                                           80
                                                AIC:
                                                                                 388.4
                                           65
                                                                                  424.2
        Df Residuals:
                                                BIC:
                                           14
        Df Model:
```

Covariance Type:	nonrobust				
[0.025 0.975]	coef	std err	t	P> t	
Intercept	4.1625	1.623	2.565	0.013	
0.921 7.404					
Flavor_JAR_Not_Enough	0.2190	1.086	0.202	0.841	-
1.950 2.388	0.0054	0.054	4 004	0 001	
Aroma_JAR_Not_Enough	0.9054	0.851	1.064	0.291	_
0.795 2.605 Color_JAR_Not_Enough	-1.1877	0.995	-1.194	0.237	
3.174 0.799	-1.10//	0.993	-1.194	0.237	_
Orange_Flavor_JAR_Not_Enough	h -0.5474	0.917	-0.597	0.553	_
2.380 1.285	0.017	0.517	01337	0.555	
Sweetness_JAR_Not_Enough	0.9587	0.916	1.046	0.299	_
0.871 2.788					
Mouthfeel_JAR_Not_Enough	0.1133	1.010	0.112	0.911	_
1.903 2.130					
Aftertaste_JAR_Not_Enough	1.0976	0.929	1.181	0.242	_
0.758 2.953	0.7001	0.004	0.700	0 426	
Flavor_JAR_Too_Much 1.098 2.514	0.7081	0.904	0.783	0.436	_
1.098 2.514 Aroma_JAR_Too_Much	0.1422	0.934	0.152	0.880	_
1.723 2.008	0.1422	0.354	0.132	0.000	
Color_JAR_Too_Much	-0.8374	0.890	-0.941	0.350	_
2.614 0.940					
Orange_Flavor_JAR_Too_Much	0.0656	0.962	0.068	0.946	_
1.855 1.986					
Sweetness_JAR_Too_Much	1.5840	0.910	1.741	0.086	_
0.233 3.401					
Mouthfeel_JAR_Too_Much	0.2151	0.980	0.220	0.827	_
1.741 2.171 Aftertaste_JAR_Too_Much	0.2335	1.041	0.224	0.823	
1.846 2.313	0.2333	1.041	0.224	0.023	_
					=====
Omnibus:	4.809	Durbin-Wats	on:		2.082
<pre>Prob(Omnibus):</pre>	0.090	Jarque-Bera			2.886
Skew:	-0.261				0.236
Kurtosis:	2.229	Cond. No.			11.8

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

Model summary for New Formulation: ${\tt OLS~Regression~Results}$

Dep. Variable: Model: Method: Date: Time:	Overall_Liking OLS Least Squares Fri, 31 May 2024 23:01:31	R-squared: Adj. R-squared: F-statistic: Prob (F-statistic): Log-Likelihood:	====	0.428 0.315 3.799 0.000152 -171.66
No. Observations: Df Residuals: Df Model:	80 66 13	AIC: BIC:		371.3 404.7
Covariance Type:	nonrobust			
	=======================================	=======================================	=====	=======================================
	coe	f std err	t	P> t

[0.025 0.975]

Intercept	7.0844	1.388	5.106	0.000		
4.314 9.855						
Flavor_JAR_Not_Enough	-0.0867	0.847	-0.102	0.919	-	
1.779 1.605						
Aroma_JAR_Not_Enough	0.3469	0.870	0.399	0.691	_	
1.390 2.084						
Color_JAR_Not_Enough	0.6205	0.833	0.745	0.459	_	
1.042 2.283						
Orange_Flavor_JAR_Not_Enough	-0.8550	0.867	-0.986	0.328	_	
2.587 0.877						
Sweetness_JAR_Not_Enough	-4.9018	0.820	-5.977	0.000	_	
6.539 -3.264						
Mouthfeel_JAR_Not_Enough	0.1296	1.455	0.089	0.929	_	
2.775 3.034	01-200		0.000	0.10_0		
Aftertaste_JAR_Not_Enough	-0.0426	0.819	-0.052	0.959	_	
1.677 1.592	010420	01013	01032	01333		
Flavor_JAR_Too_Much	-0.0380	0.832	-0.046	0.964	_	
1.699 1.623	0.0300	01032	01040	0.304		
Aroma_JAR_Too_Much	0.8392	0.876	0.958	0.342		
0.910 2.589	0.0392	0.070	0.930	0.342	_	
	0 6530	0 053	-0.686	0.495		
Color_JAR_Too_Much	-0 . 6538	0.953	-0.080	0.495	_	
2.557 1.249	0.0041	0.000	0 770	0 444		
Orange_Flavor_JAR_Too_Much	-0.6641	0.862	-0.770	0.444	_	
2.385 1.057	2 002 45	2 02 45	4 007		2 67	
Sweetness_JAR_Too_Much	3.963e-15	3.82e-15	1.037	0.304	-3.67	
e-15 1.16e-14						
Mouthfeel_JAR_Too_Much	-0.0712	0.768	-0.093	0.926	_	
1.604 1.462						
Aftertaste_JAR_Too_Much	-0.7135	0.907	-0.786	0.434	_	
2.525 1.098						
=======================================				========	=====	
Omnibus:	5.303	Durbin-Watso	on:		1.922	
<pre>Prob(Omnibus):</pre>	0.071	Jarque-Bera (JB):			2.424	
Skew:	-0.068	Prob(JB):			0.298	
Kurtosis:	2.158	Cond. No.		5.2	7e+17	
=======================================		=========		========	=====	

Notes:

- [1] Standard Errors assume that the covariance matrix of the errors is correctly specified.
- [2] The smallest eigenvalue is 6.37e-34. This might indicate that there are strong multicollinearity problems or that the design matrix is singular.

In [4]: #Intensity score analysis intensity_attributes = [col for col in rawdata.columns if 'Intensity' in col] intensity_means = rawdata.groupby('Product')[intensity_attributes].mean().round(# Separating the data for each product original_data = rawdata[rawdata['Product'] == 'Original'] new_formulation_data = rawdata[rawdata['Product'] == 'New Formulation'] # Performing paired t-test and Wilcoxon signed-rank test for each intensity attr t_test_results = {} wilcoxon_test_results = {} for attribute in intensity_attributes: t_stat, t_p_value = ttest_rel(original_data[attribute], new_formulation_data t_test_results[attribute] = (t_stat, t_p_value)

```
w_stat, w_p_value = wilcoxon(original_data[attribute], new_formulation_data[
             wilcoxon_test_results[attribute] = (w_stat, w_p_value)
         # Creating dataframes for the results
         t_test_results_df = pd.DataFrame.from_dict(t_test_results, orient='index', colum
         wilcoxon test results df = pd.DataFrame.from dict(wilcoxon test results, orient=
         intensity_means, t_test_results_df, wilcoxon_test_results_df
         #
                          Sweetness Intensity Orange Flavor Intensity \
Out[4]:
         Product
         New Formulation
                                         4.94
                                                                  4.95
                                         4.99
         Original
                                                                  5.24
                          Aftertaste_Intensity
         Product
         New Formulation
                                          5.22
                                          5.75
         Original
                                             p_value
                                    t stat
         Sweetness_Intensity
                                  0.177445
                                            0.859613
         Orange_Flavor_Intensity
                                  0.811947
                                            0.419262
         Aftertaste_Intensity
                                  1.588946 0.116067,
                                           p_value
                                  w stat
         Sweetness_Intensity
                                  1062.0 0.779423
         Orange_Flavor_Intensity 1221.0 0.366285
         Aftertaste_Intensity
                                   731.5 0.122381)
In []:
In []:
```