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# Chi2 sqaure test for the relationship to be established
import math
# Cramer's Value to measure the chi square to find the association.
def cramers_V(statistical_value, contingency_table, total_observation,
row_count=0, column_count=0):
   # Finding the columns length and rows length
    for rows in contingency_table:
       row_count += 1
    for columns in contingency_table.T:
       column_count += 1
   # Degree Of Freedom
    k = min(row_count, column_count)
    denominator = total_observation * k # Total Observations as well as the
mininmum Degree Of Freedom
    numerator = statistical_value # Chi2 Statistics
   # Cramer's V Formula
    return math.sqrt(numerator / denominator)
comparee = ['Biking Habit', 'Jogging Habit', 'Walking Habit'] # Add the
columns you want to use against
comparer = ['Smoking Habit', 'Drinking Habit'] # Add the columns you want
to use in the first place
def chi_square(healthy_columns, unhealthy_columns):
    try:
       for first_columns in range(len(unhealthy_columns)): # For Unhealthy
Columns
           print()
           print(f'
  ------{unhealthy_columns[fir
st columns]}-----
----')
           print()
           for second_columns in range(len(healthy_columns)): # For
Healthy Columns
               print(f'{unhealthy_columns[first_columns]} to
{healthy_columns[second_columns]}----')
               table = pandas.crosstab(data['Smoking Habit'],
data[healthy_columns[second_columns]])
               chi2_statistics, p_value, degree_of_freedom, expected =
chi2_contingency(table)
               print(
                   f"Cramer's V Score for {
                       unhealthy_columns[first_columns]} and
{healthy_columns[second_columns]
                       cramers V(statistical value=chi2 statistics,
                       contingency_table=table,
                       total_observation=sum(
                           sum(
                               table.values
                   ) # Initiating the Cramer's V to check the accuracy for
the association of Chi2 Test
               if p_value < 0.05:
                   print('There is a strong evidence for the association
to be identified for this relationship')
               else:
                   print('There is not a strong evidence for the
association to be identified for this relationship')
                   print()
    except (IndexError, ValueError, SyntaxError):
       if IndexError:
           print('I want you to perform correct indexing')
       elif ValueError:
           print('I cannot work witht the values you provided, change it!'
)
       elif SyntaxError:
           print('Could you please write clear for me to understand it and
then get you the output')
       else:
           print('Its an error! Please change your code once')
# Final Chi2 Test For Multiple Columns Comparisions
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chi\_square(comparee, comparer)