

# regression\_tables\_and\_graphs

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## 1 Python Tutorial Program: Creating Regression Tables and Line Graphs using Pandas and Matplotlib

By Kenneth Burchfiel

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**For more background on this program, please see my corresponding blog post here: [link forthcoming]**

This program demonstrates how Python can be used to determine the statistical significance of an intervention via a Difference-in Differences analysis, then display the effects of that intervention through a series of graphs.

It performs the following tasks:

1. Import a table containing **fictional** beginning-of-year and end-of-year test score data for students in multiple high schools and grades
2. Create individual Pandas DataFrames for each school
3. Perform a regression for each school-grade pair that incorporates 3 variable: (1) time (beginning of year vs. end of year); (2) intervention group; and (3) time-intervention group interaction
4. Append these results to a table for easier comparison/analysis
5. Use this regression data to create and save charts that compare test score improvements for two groups of students
6. Create additional charts that visualize certain regression coefficients and indicate statistical significance
7. Output the table of regression results to an .xlsx worksheet

### 1.1 Scenario:

(Note: both the scenario and the data presented in this program are fictional.)

Orange Valley School District (OVSD) is piloting an after-school math program called Code the Concepts in each of its 8 high schools. This program aims to teach students relevant math skills through programming and computer science applications, and is available in grades 9-12.

Although the district believes that the programming knowledge taught in this course is highly valuable in itself, it also wants to evaluate the impact, if any, this program has on students' math skills. The means of evaluating the math skills will be the Higher Grades Math Assessment (HGMA), a test administered at the beginning and end of each school year.

In [an earlier program](#), I showed how Python could be used to turn a dataset with HGMA scores into pivot tables and charts. Meanwhile, in this program, I will perform regression analyses of this data to determine (1) whether the increase in scores over time for students enrolled in Code the Concepts was significantly higher than the increase for non-enrolled students. These regressions will be followed by line and bar graphs that help visualize the output of the regression analysis.

## 1.2 An explanation of using regressions to run difference-in-differences analyses

If you are already familiar with the use of regression analyses to perform difference-in-difference analyses, feel free to skip over the following text block. If not, please consider reading it, as it will help explain my choices in coding the regression analyses within the program.

Note: A webpage titled “Difference-in-Difference Estimation” ([accessible at publichealth.columbia.edu](http://publichealth.columbia.edu)) was a valuable reference in writing the following summary. It also discusses precautions that I skip over here.

One means of determining whether an intervention had a significant effect is to collect an outcome measure at two different points of time for two different groups: the experimental group (who received the intervention) and the control group (who did not). The magnitude of the experimental group's change can then be compared with the control group's change, producing what is known as a difference-in-differences (or difference-in-difference) analysis.

For example, in this tutorial program, the difference-in-differences analysis will focus on beginning-of-year and end-of-year test scores for (1) students who were enrolled in an optional after-school math class and (2) students who were not. Suppose that the students enrolled in the optional program produced a 10-point score improvement during the school year. This may tempt you to conclude that the optional program was a great success. However, if students who weren't enrolled in the program *also* had a 10-point score improvement, it would seem more likely that other factors (such as students' regular within-school math instruction), rather than the optional program, were responsible for the growth in math scores.

Suppose instead that the enrolled students' scores increased by 15 points, compared to 13 points for non-enrolled students. In this case, enrolled' students scores increased 2 points higher on average than did non-enrolled students' scores. Is this increase statistically significant, or is the chance that the difference was a result of random variation too high?

One way to determine statistical significance is to run a linear regression analysis. Three independent variables (in addition to a constant value) need to be included: (1) time (beginning of year vs. end of year); (2) enrollment status; and (3) an interaction variable (in this case, one specifying whether the data point is from an enrolled student at the end of the year).

The p value of the time coefficient will help determine whether the growth (or decline) in scores across time was statistically significant for the student body as a whole. Meanwhile, the p value of the interaction coefficient provides insight into the intervention's significance. A p value below 0.05 would suggest that students enrolled in the optional program had a significantly larger increase (or decrease) in scores than did non-enrolled students.

With this explanation out of the way, I'll now turn to discussing the actual program.

I will start by importing a number of useful items:

```
[2]: import time
start_time = time.time() # Allows the program's runtime to be measured
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import matplotlib.ticker as mtick
from adjustText import adjust_text
import statsmodels.api as sm
pd.set_option('display.max_rows',100)
pd.set_option('display.max_columns',100)
```

Next, I will import the beginning-of-year and end-of-year HGMA scores for all OVSD high schoolers into a Pandas DataFrame. (This fictional score data was generated within [an earlier tutorial program](#).)

```
[3]: df_scores = pd.read_excel('scores_by_program_enrollment_copy.xlsx')
df_scores.drop('Student_ID',axis=1,inplace=True)
df_scores
```

```
[3]:
```

	School	Grade	Enrolled	Time	Score	Count
0	Bayville	12	Yes	2018_9	39	1
1	Bayville	12	Yes	2019_5	54	1
2	Bayville	12	No	2018_9	77	1
3	Bayville	12	No	2019_5	75	1
4	Eagle	12	No	2018_9	35	1
...	...	...	...	...	...	...
7675	Cardinal	11	No	2018_9	59	1
7676	Eagle	12	Yes	2019_5	52	1
7677	Eagle	12	Yes	2018_9	33	1
7678	Westwood	10	Yes	2018_9	66	1
7679	Westwood	10	Yes	2019_5	81	1

[7680 rows x 6 columns]

Next, I will create a list of all schools within the DataFrame, which will make it easier to produce an individual DataFrame for each school.

```
[4]: school_list = list(pd.unique(df_scores['School'])) # This list will be looped_
      →through in order to create DataFrames for each school
school_list.append('Total')
print(school_list)
```

```
['Bayville', 'Eagle', 'Westwood', 'Fair Lake', 'Central', 'East River', 'Olive',
'Cardinal', 'Total']
```

The following function takes a DataFrame with score data for an individual school, then converts

it into a form that will make later regression analyses easier to perform.

```
[5]: def create_data_source_for_regression(school, df): # The 'school' variable is
    ↪not needed for the function to run, but is included so that the DataFrame
    ↪output can be identified as belonging to a particular school.
        if school == 'Total': # DataFrame will encompass all students, so no need
    ↪to incorporate a query() statement
            df_source = df[['Grade', 'Enrolled', 'Time', 'Score']].copy() # The
    ↪'School' column is not necessary since this DataFrame will encompass all
    ↪schools as a whole
        else:
            df_source = df.query("School == @school")[['Grade', 'Enrolled', 'Time',
    ↪'Score']].copy()
            df_source = pd.get_dummies(df_source.copy(), drop_first=True) # Even if
    ↪'School' had been included as one of the columns earlier, this line would
    ↪end up removing the 'School' column for cases where only one school was
    ↪being queried. This is due to the fact that, with drop_first enabled, the
    ↪first dummy variable for School (School_[school_name]) will get removed. In
    ↪this case, School_[school_name] is the only dummy variable (since all
    ↪students attend that school), so no school column will show up in the final
    ↪output. This is fine, since this column does not provide any information (as
    ↪we already know each student in this DataFrame attends that school.)
            df_source.rename(columns={"Time_2019_5": "Time_EOY"}, inplace=True)
            df_source['EOY_and_Enrolled'] = 0
            for i in range(len(df_source)):
                if (df_source.loc[df_source.index[i], 'Time_EOY'] == 1) and (df_source.
    ↪loc[df_source.index[i], 'Enrolled_Yes'] == 1):
                    df_source.loc[df_source.index[i], 'EOY_and_Enrolled'] = 1
            return school, df_source # Stores the school along with the DataFrame in a
    ↪tuple so that DataFrames can be matched to schools more easily later on
```

```
[6]: school_df_pair_list = [] # Stores tuple outputs of
    ↪create_data_source_for_regression for each school and for the district as a
    ↪whole
    for school in school_list:
        school_df_pair_list.append(create_data_source_for_regression(school,
    ↪df_scores)) # The output, like the input, is a tuple
```

## 2 Here with editing—note how dummy variables, including one for the interaction, were created below

```
[21]: school_df_pair_list[0][0]
```

```
[21]: 'Bayville'
```

```
[7]: school_df_pair_list[0][1]
```

```
[7]:
```

	Grade	Score	Enrolled_Yes	Time_EOY	EOY_and_Enrolled
0	12	39	1	0	0
1	12	54	1	1	1
2	12	77	0	0	0
3	12	75	0	1	0
26	9	61	1	0	0
...	...	...	...	...	...
7619	11	32	0	0	0
7658	9	58	0	1	0
7659	9	62	0	0	0
7672	10	54	1	1	1
7673	10	30	1	0	0

[904 rows x 5 columns]

```
[8]: grade_test = 'Grade 9'
grade_int = int(grade_test.replace('Grade ', ''))
school_df_pair_list[0][1].query("Grade == @grade_int")['Score']
```

```
[8]: 26      61
      27      69
      208     36
      209     38
      282     43
      ..
      7351    63
      7358    57
      7359    48
      7658    58
      7659    62
Name: Score, Length: 204, dtype: int64
```

```
[9]: def create_regression_table(school, data_source_for_regression):
      regression_table = pd.DataFrame(index=['Non_Enrolled_BOY_Mean',
      ↪ 'Non_Enrolled_EOY_Mean', 'Enrolled_BOY_Mean', 'Enrolled_EOY_Mean',
      ↪ 'Time_Coeff', 'Time_Pval', 'Time_Significant', 'EOY_and_Enrolled_Coeff',
      ↪ 'EOY_and_Enrolled_Pval', 'EOY_and_Enrolled_Significant', 'R_Squared',
      ↪ 'Adj_R_Squared'], columns = ['Grade 9', 'Grade 10', 'Grade 11', 'Grade 12',
      ↪ 'Total'])
      for i in range(len(regression_table.columns)):
          if regression_table.columns[i] == 'Total':
              filtered_df = data_source_for_regression.copy()
          else:
              grade_int = int(regression_table.columns[i].replace('Grade ', ''))
              filtered_df = data_source_for_regression.query("Grade ==
      ↪ @grade_int")
              y = filtered_df['Score']
```

```

x_vars = filtered_df[['Time_EOY', 'Enrolled_Yes', 'EOY_and_Enrolled']]
x_vars = sm.add_constant(x_vars)
model = sm.OLS(y,x_vars)
results = model.fit()

regression_table.loc['Time_Coeff'][regression_table.columns[i]] =
↳results.params['Time_EOY']
regression_table.loc['Time_Pval'][regression_table.columns[i]] =
↳results.pvalues['Time_EOY']
    if results.pvalues['Time_EOY'] < 0.05:
        regression_table.loc['Time_Significant'][regression_table.
↳columns[i]] = 1
    else:
        regression_table.loc['Time_Significant'][regression_table.
↳columns[i]] = 0

regression_table.loc['EOY_and_Enrolled_Coeff'][regression_table.
↳columns[i]] = results.params['EOY_and_Enrolled']
regression_table.loc['EOY_and_Enrolled_Pval'][regression_table.
↳columns[i]] = results.pvalues['EOY_and_Enrolled']
    if results.pvalues['EOY_and_Enrolled'] < 0.05:
        regression_table.
↳loc['EOY_and_Enrolled_Significant'][regression_table.columns[i]] = 1
    else:
        regression_table.
↳loc['EOY_and_Enrolled_Significant'][regression_table.columns[i]] = 0

regression_table.loc['Non_Enrolled_BOY_Mean'][regression_table.
↳columns[i]] = results.params['const'] # These methods of calculating means
↳by adding different regression coefficients together produced the same
↳results as calculating them by querying specific parts of the DataFrame and
↳taking the means of those parts.

regression_table.loc['Non_Enrolled_EOY_Mean'][regression_table.
↳columns[i]] = results.params['const'] + results.params['Time_EOY']

regression_table.loc['Enrolled_BOY_Mean'][regression_table.columns[i]]
↳= results.params['const'] + results.params['Enrolled_Yes']

regression_table.loc['Enrolled_EOY_Mean'][regression_table.columns[i]]
↳= results.params['const'] + results.params['Time_EOY'] + results.
↳params['Enrolled_Yes'] + results.params['EOY_and_Enrolled']

regression_table.loc['R_Squared'][regression_table.columns[i]] =
↳results.rsquared

```

```

        regression_table.loc['Adj_R_Squared'][regression_table.columns[i]] =
↪results.rsquared_adj

    return school, regression_table # the school is included in the output to
↪help with identification.

```

```

[10]: regression_table_list = []

for i in range(len(school_df_pair_list)): # school_df_pair_list contains both
↪school names and dataframes for those schools. These names and DataFrames
↪can then be used as arguments for the create_regression_table function when
↪it loops through regression_table_list.
    regression_table_list.
    ↪append(create_regression_table(school_df_pair_list[i][0],
    ↪school_df_pair_list[i][1])) # i refers to the tuple in the list; [0] refers
    ↪to the school name; and [1] refers to the DataFrame returned by
    ↪create_data_source_for_regression.

```

```

[11]: len(regression_table_list) # Determines how many sheets need to be added to the
↪Google Sheet that will store the DataFrame outputs

```

```
[11]: 9
```

```
[12]: regression_table_list[0][0]
```

```
[12]: 'Bayville'
```

```
[20]: regression_table_list[0][1]
```

```

[20]:

```

	Grade 9	Grade 10	Grade 11	Grade 12	\
Non_Enrolled_BOY_Mean	54.66	52.050847	56.785714	52.95	
Non_Enrolled_EOY_Mean	58.04	53.542373	58.5	53.916667	
Enrolled_BOY_Mean	56.096154	53.232143	55.157895	54.693548	
Enrolled_EOY_Mean	64.038462	70.125	63.982456	70.967742	
Time_Coeff	3.38	1.491525	1.714286	0.966667	
Time_Pval	0.247258	0.590873	0.552521	0.722588	
Time_Significant	0	0	0	0	
EOY_and_Enrolled_Coeff	4.562308	15.401332	7.110276	15.307527	
EOY_and_Enrolled_Pval	0.264748	0.000138	0.081081	0.00008	
EOY_and_Enrolled_Significant	0	1	0	1	
R_Squared	0.058384	0.197423	0.046398	0.203692	
Adj_R_Squared	0.044259	0.186769	0.033512	0.193738	
Total					
Non_Enrolled_BOY_Mean	54.048889				
Non_Enrolled_EOY_Mean	55.875556				
Enrolled_BOY_Mean	54.770925				

Enrolled_EOY_Mean	67.418502
Time_Coeff	1.826667
Time_Pval	0.197455
Time_Significant	0
EOY_and_Enrolled_Coeff	10.82091
EOY_and_Enrolled_Pval	0.0
EOY_and_Enrolled_Significant	1
R_Squared	0.117464
Adj_R_Squared	0.114522

```
[13]: def plot_line(xpoints, ypoints, color=('000000'), line_label='_',
    ↪point_label_1 = '_', point_label_2 = '_'): # X and Y points need to be in a
    ↪list format; if they derive from a DataFrame, apply list() to them
    ↪beforehand. Each list should have only two points.
        # Matplotlib documentation notes that "Specific lines can be excluded from
    ↪the automatic legend element selection by defining a label starting with an
    ↪underscore" (https://matplotlib.org/stable/api/\_as\_gen/matplotlib.pyplot.legend.html), which is why '_' is used as the default label.
        # print("x values:",xpoints) # for debugging
        # print("y values:",ypoints)
        plt.plot(xpoints, ypoints, linestyle='solid',label=line_label,
    ↪linewidth=2,color=color)
        plt.plot(xpoints[0], ypoints[0], 'om', label=point_label_1) # Plots a data
    ↪point on the left of the line. See the 'notes' section within https://matplotlib.org/stable/api/\_as\_gen/matplotlib.pyplot.plot.html
        plt.plot(xpoints[1], ypoints[1], 'og',label=point_label_2) # Plots a data
    ↪point on the right of the line.
```

```
[14]: for t in range(len(regression_table_list)):
    t_min = min(regression_table_list[t][1].iloc[0:4,0:5].min()) # First finds
    ↪the minimum value for each column, then determines the lowest value out of
    ↪those values
    t_max = max(regression_table_list[t][1].iloc[0:4,0:5].max())
    if t == 0:
        all_schools_min = t_min
        all_schools_max = t_max
    else:
        if t_min < all_schools_min:
            all_schools_min = t_min
        if t_max > all_schools_max:
            all_schools_max = t_max

    print(all_schools_min, all_schools_max)
```

48.777777777777786 74.8360655737705



```

[15]: for t in range(len(regression_table_list)): # This loop runs through each
    ↪ database stored in the regression_table_list. i is used as an iterator below
    ↪ and I prefer not to use the same letter for two different iterators within a
    ↪ nested for loop, so I instead used t (for 'tuple') here.
        data_source = regression_table_list[t][1] # Giving this data_table a
    ↪ shorter variable name makes the code more readable and, more importantly,
    ↪ makes it easy to replace this DataFrame with a different one.
        # regression_table_list[t][2] returns a dataframe.
        school = regression_table_list[t][0]
        y_min = min(data_source.iloc[0:4,0:5].min())
        y_max = max(data_source.iloc[0:4,0:5].max())

        dl_1 = [0, 1.5, 3, 4.5, 6] # dl = data_labels. These points, along with
    ↪ those in dl_2, help provide correct spacing for the graphs and axis labels.
    ↪ Each line has a length of 1, and there is a gap of 0.5 in between each line.
        # dl_1 stores the x axis start points for each graph, whereas dl_2 stores
    ↪ the x axis end points.
        dl_2 = [1, 2.5, 4, 5.5, 7]

        col_labels = list(data_source.columns.copy()) # Creates a list of all the
    ↪ grades and the 'Total' category in the same order that they appear within
    ↪ the table. The list() operation is necessary in the event that changes to
    ↪ the column names need to be made.
        fig, ax = plt.subplots(figsize=[9,5])
        fig.set_facecolor('white')
        plt.ylim(all_schools_min-5,all_schools_max+5)
        plt.grid(b=True,which='both',axis = 'y')
        # plt.grid(axis = 'y', which='minor', linewidth=1)

        xtick_list = []

        x_avoid_list = []
        y_avoid_list = [] # These two lists will store a series of coordinates that
    ↪ adjust_text will move data labels away from.
        data_label_list = []

        for i in range(len(data_source.columns)):
            if i == 0: # This if statement, along with the parameter values in the
    ↪ two plot_line statements below, cause only the first set of labels to be
    ↪ graphed, not the subsequent ones (in order to avoid duplicate labels).
                current_non_enrolled_line_label = 'Not Enrolled'
                current_enrolled_line_label = 'Enrolled'
                current_point_label_1 = 'Beginning of Year'
                current_point_label_2 = 'End of Year'
            else:
                current_non_enrolled_line_label = '_'

```

```

        current_enrolled_line_label = '_'
        current_point_label_1 = '_'
        current_point_label_2 = '_'

    non_enrolled_yvals = [data_source.
↳loc['Non_Enrolled_BOY_Mean'][data_source.columns[i]], data_source.
↳loc['Non_Enrolled_EOY_Mean'][data_source.columns[i]]]
    enrolled_yvals = [data_source.loc['Enrolled_BOY_Mean'][data_source.
↳columns[i]], data_source.loc['Enrolled_EOY_Mean'][data_source.columns[i]]]

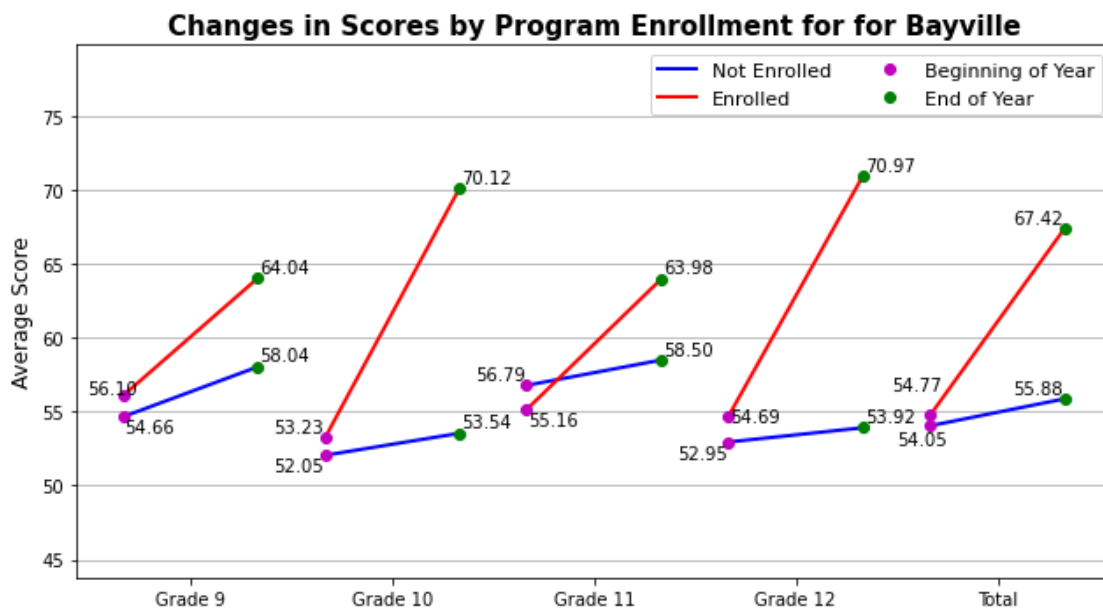
    □
    plot_line([dl_1[i],dl_2[i]],non_enrolled_yvals,color=('blue'),line_label=□
↳current_non_enrolled_line_label)
    plot_line([dl_1[i],dl_2[i]],enrolled_yvals,color=('red'), line_label =□
↳current_enrolled_line_label, point_label_1 = current_point_label_1,□
↳point_label_2 = current_point_label_2)
    # Creating data labels for both sets of lines
    for j in range (2): # Loop will start with the non-enrolled students□
↳graph and then conclude with the enrolled students graph
        if j == 0:
            yval_source = non_enrolled_yvals # Contains points for the□
↳leftmost part of each graph
            if j == 1:
                yval_source = enrolled_yvals # Contanins points for the□
↳rightmost part of each graph
            left_data_label = plt.text(dl_1[i],yval_source[0],'{:.2f}'.
↳format(yval_source[0]),ha='center') # Plots a percentage label next to the□
↳left end of the line using the points in dl_1 for x values and the□
↳percentage values from the DataFrame for y values.
            data_label_list.append(left_data_label)
            right_data_label = plt.text(dl_2[i],yval_source[1],'{:.2f}'.
↳format(yval_source[1]),ha='center')
            data_label_list.append(right_data_label)
            x_avoid = np.linspace(0,1,21) # Creates 20 points between (and□
↳including) 0 and 1, as each line has an x length of 1
            y_avoid = yval_source[0] + x_avoid*(yval_source[1] -□
↳yval_source[0]) # Creates a series of y values that line up with the lines□
↳created earlier in this for loop. Each value equals the left y□
↳value(yval_source[0]) plus the x value (from 0 to 1) * the total rise of the□
↳line from left to right.
            x_avoid += dl_1[i] # Shifts the points in x_avoid so that they line□
↳up with their corresponding line. (Only the first line actually starts at 0;□
↳the other lines are offset by dl_1[i].
            x_avoid_list.extend(x_avoid)
            y_avoid_list.extend(y_avoid)
    adjust_text(data_label_list, x=x_avoid_list, y=y_avoid_list)

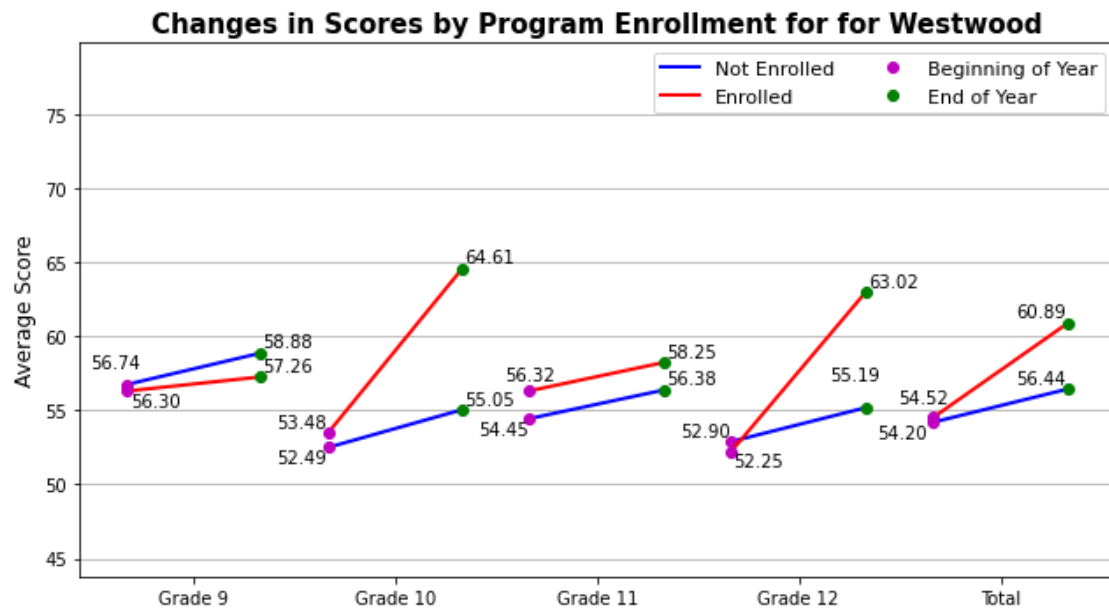
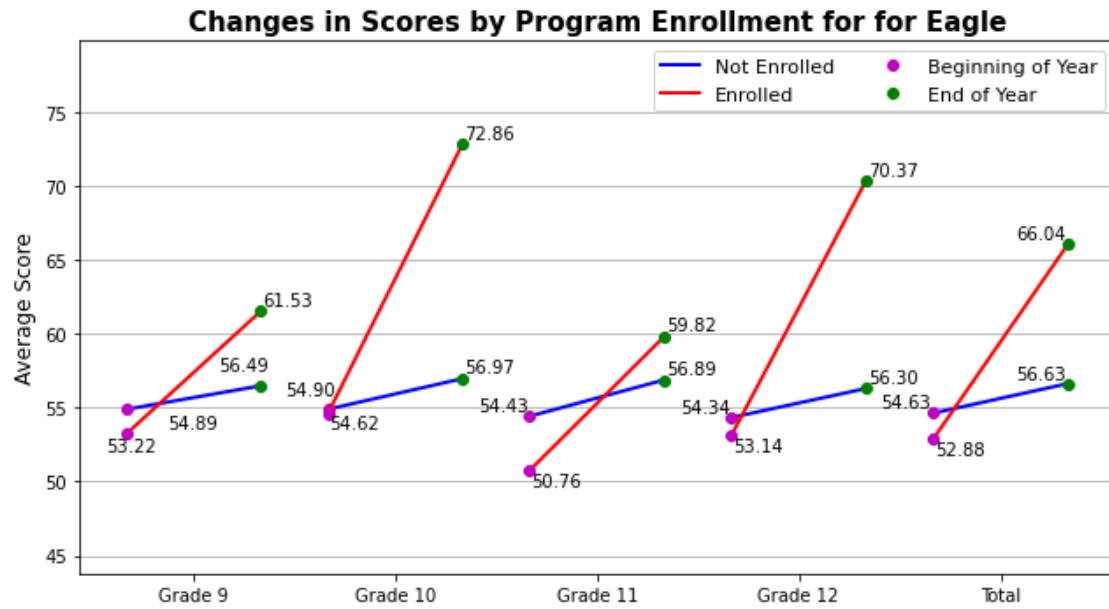
```

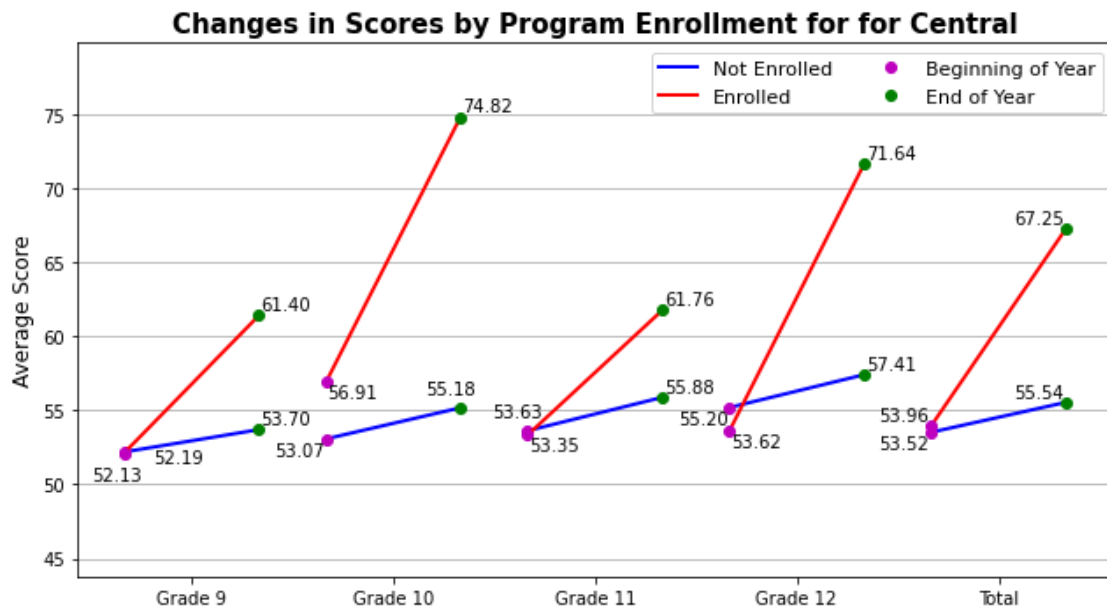
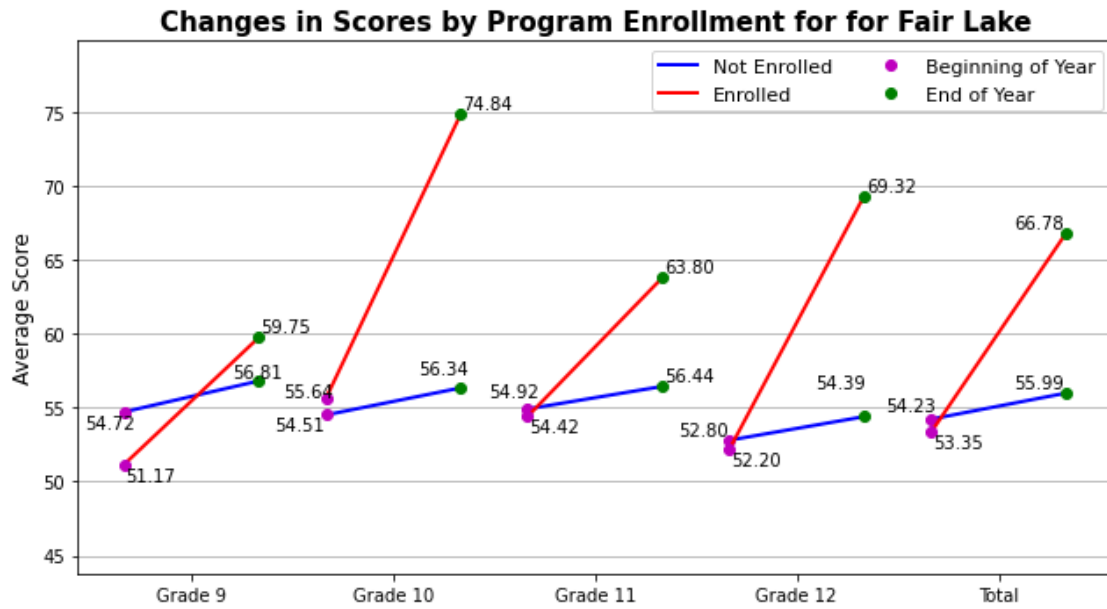
```

if school == 'Total':
    plt.title("Changes in Scores by Program Enrollment for All_
→Schools",fontweight='bold',fontsize=15)
else:
    plt.title("Changes in Scores by Program Enrollment for for_
→"+school,fontweight='bold',fontsize=15)
    for m in range(len(dl_1)):
        xtick_list.append(np.mean([dl_1[m], dl_2[m]])) # Calculates the_
→midpoint between each pair of dl_1 points and dl_2 points.
        ax.set_xticks(xtick_list) # These ticks are placed in the middle of each_
→graph in order to make the graph labels align correctly.
    ax.set_xticklabels(col_labels)
    ax.set_ylabel('Average Score',fontsize=12)
    plt.tight_layout()
    plt.legend(ncol=2,fontsize=11)
    file_string = "score_graphs\\"+school + '_score_changes.png'
    plt.savefig(file_string,dpi=400)
    plt.show()
    #plt.close()

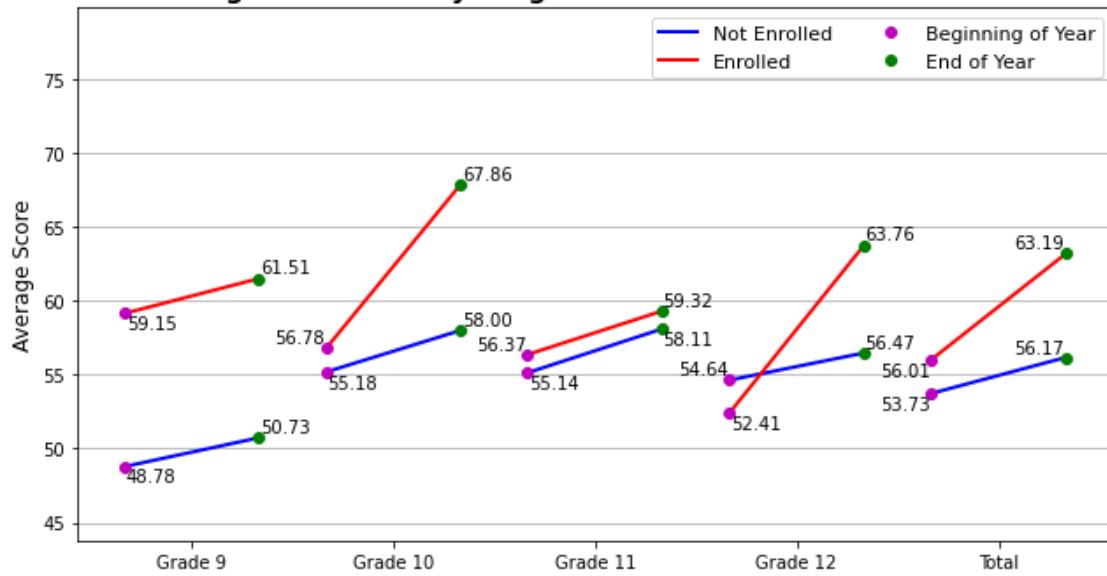
```



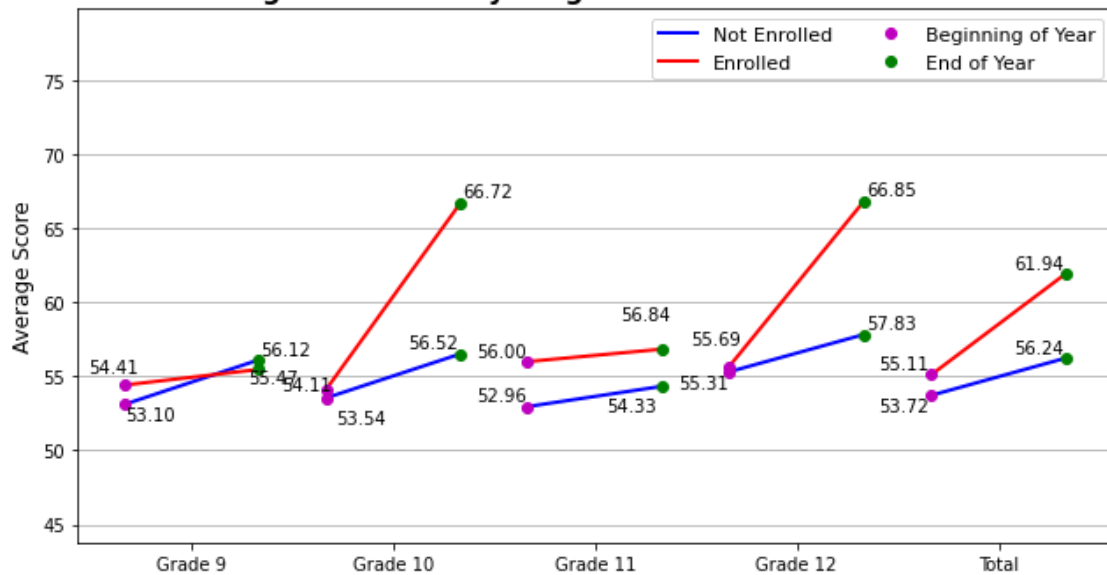


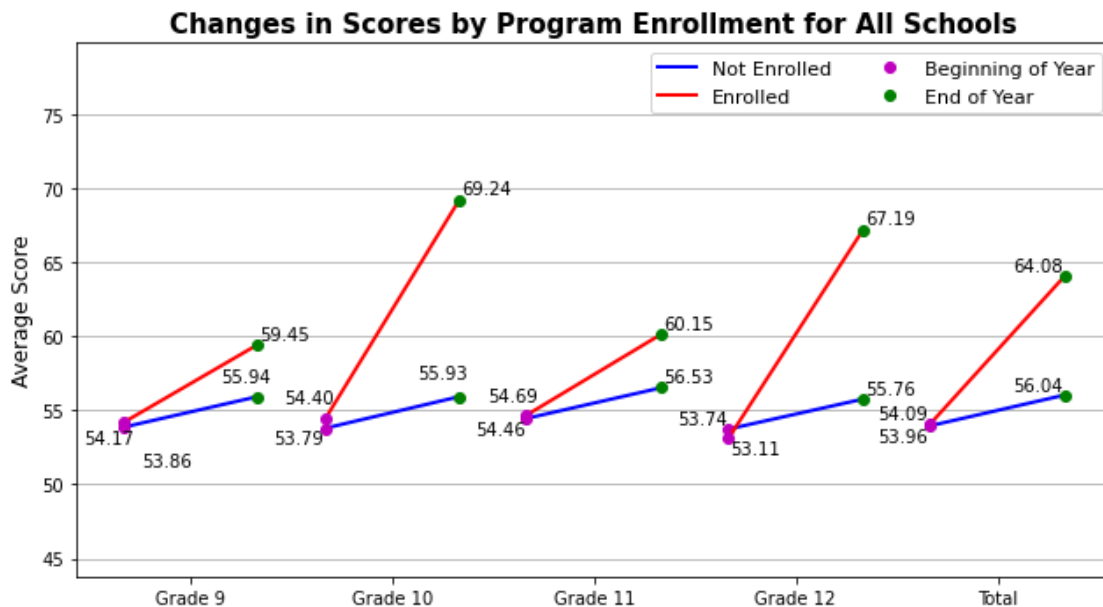
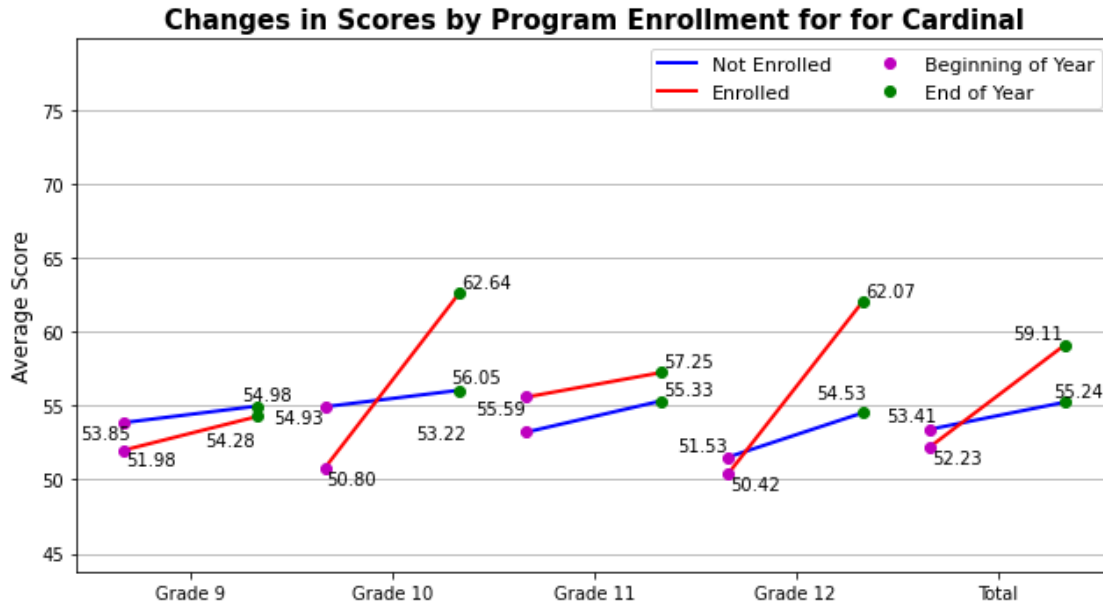


**Changes in Scores by Program Enrollment for for East River**



**Changes in Scores by Program Enrollment for for Olive**





A virtually identical set of graphs was created in [an earlier program](#). Although these graphs were based off regression outputs, whereas the graphs in the earlier program were based off pivot tables, the numbers shown in both graphs are identical.

The following block of code creates charts that show the relative magnitude of the different coefficients within the DataFrames `regression_table_list`.

```

[17]: for t in range(len(regression_table_list)): # This loop runs through each
    ↪ database stored in the regression_table_list. i is used as an iterator below
    ↪ and I prefer not to use the same letter for two different iterators within a
    ↪ nested for loop, so I instead used t (for 'tuple') here.

    data_source = regression_table_list[t][1] # Giving this data_table a
    ↪ shorter variable name makes the code more readable and, more importantly,
    ↪ makes it easy to replace this DataFrame with a different one.
    # regression_table_list[t][2] returns a dataframe.
    school = regression_table_list[t][0]

    fig, ax = plt.subplots(figsize=[12.8,7.5]) # Widening the graph gives the x
    ↪ axis labels more space
    col_labels = list(data_source.columns.copy())
    print(type(col_labels))
    fig.set_facecolor('white')

    x_values = np.linspace(0, len(col_labels)-1, len(col_labels)) # Creates x
    ↪ values for plotting bars and other features. The list of x values iterates
    ↪ by 1 and is equal to the length of the number of column labels.
    # It's useful to set these manually so that other points on the graph (such
    ↪ as significance labels) can be plotted at the right positions.

    bar_width = 0.4
    ax.set_xticks(x_values)
    plt.grid(axis = 'y') # https://matplotlib.org/stable/api/_as_gen/matplotlib.
    ↪ artist.Artist.set_zorder.html#matplotlib.artist.Artist.set_zorder
    ax.set_xticklabels(col_labels, rotation = 0)
    ax.set_axisbelow(True) # https://matplotlib.org/stable/api/_as_gen/
    ↪ matplotlib.axes.Axes.set_axisbelow.html#matplotlib.axes.Axes.set_axisbelow
    time_coeff_bars = plt.bar(x_values - bar_width/2, data_source.
    ↪ loc['Time_Coeff'][:, bar_width, label='Time
    ↪ Coefficient', color='red', edgecolor='white', linewidth=1) # [:] is used
    ↪ because all columns are being included; however, this could be adjusted to
    ↪ only include a range of columns if needed. Converting the output to a list
    ↪ avoids an error message.
    time_enrolled_interaction_coeff_bars = plt.bar(x_values + bar_width/2,
    ↪ data_source.loc['EOY_and_Enrolled_Coeff'][:, bar_width,
    ↪ label='Time-Enrolled
    ↪ Interaction\nCoefficient', color='blue', edgecolor='white', linewidth=1)
    ax.bar_label(time_coeff_bars, label_type =
    ↪ 'edge', color='red', fontweight='bold', fmt='%.3f')
    ax.bar_label(time_enrolled_interaction_coeff_bars, label_type =
    ↪ 'edge', color='blue', fontweight='bold', fmt='%.3f')

```



```

ax.set_ylim(min(ax.get_ylim()[0], 0), max(ax.get_ylim()[1]+3,0))
ax_height = ax.get_ylim()[1] - ax.get_ylim()[0]
ax_width = ax.get_xlim()[1] - ax.get_xlim()[0]

add_legend_flag = 0
# Adding labels to indicate which coefficients are statistically significant
for s in range (len(data_source.columns)):
    if data_source.loc['Time_Significant'][s] == 1:
        plt.text(x_values[s] - bar_width/2, data_source.
↪loc['Time_Coeff'][s]/2, 'S', color = (0, 1, 0), fontsize=15,
↪fontweight='bold', ha = 'center') # using [s]/2 places the point in the
↪middle of the bar. Because the y limits of the axis were set so that 0 would
↪always be included, 'S' should never be hidden from view.
        plt.text(x_values[s] - bar_width/2, data_source.
↪loc['Time_Coeff'][s]/2-ax_height/20, 'p=\n'+'{:.3f}'.format(data_source.
↪loc['Time_Pval'][s]), color = 'white', fontsize=10, fontweight='bold', ha =
↪'center')
        add_legend_flag = 1
        # -ax_height/[20] offsets the label by a value that scales with the
↪y axis dimensions so that the p value doesn't overlap with the 'S' metric.
        if data_source.loc['EOY_and_Enrolled_Significant'][s] == 1:
            plt.text(x_values[s] + bar_width/2, data_source.
↪loc['EOY_and_Enrolled_Coeff'][s]/2, 'S', color = (0, 1, 0), fontsize=15,
↪fontweight='bold', ha = 'center')
            plt.text(x_values[s] + bar_width/2, data_source.
↪loc['EOY_and_Enrolled_Coeff'][s]/2-ax_height/20, 'p=\n'+'{:.3f}'.
↪format(data_source.loc['EOY_and_Enrolled_Pval'][s]), color = 'white',
↪fontsize=10, fontweight='bold', ha = 'center')
            add_legend_flag = 1
        if add_legend_flag == 1:
            plt.text(ax.get_xlim()[0]+ax_width/30, ax.get_ylim()[1]-ax_height/20,
↪'S = Statistically significant coefficient', color = (0, 1, 0), fontweight =
↪'bold', fontsize = 11)

plt.legend(loc = 'upper right')

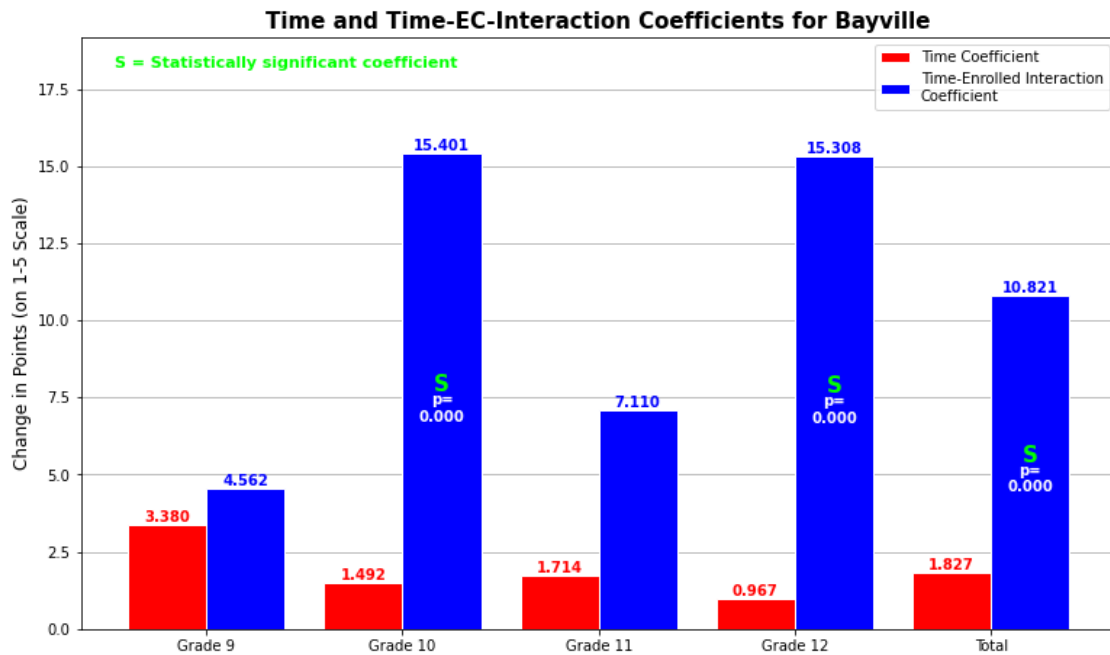
ax.set_ylabel('Change in Points (on 1-5 Scale)',fontsize=12)
if school == 'Total':
    title_string = "Time and Time-Enrolled-Interaction Coefficients for All
↪Schools"
else:
    title_string = "Time and Time-EC-Interaction Coefficients for "+school
plt.title(title_string,fontweight='bold',fontsize=15)

file_string = 'coeff_graphs\\'+school+'_coeffs.png'

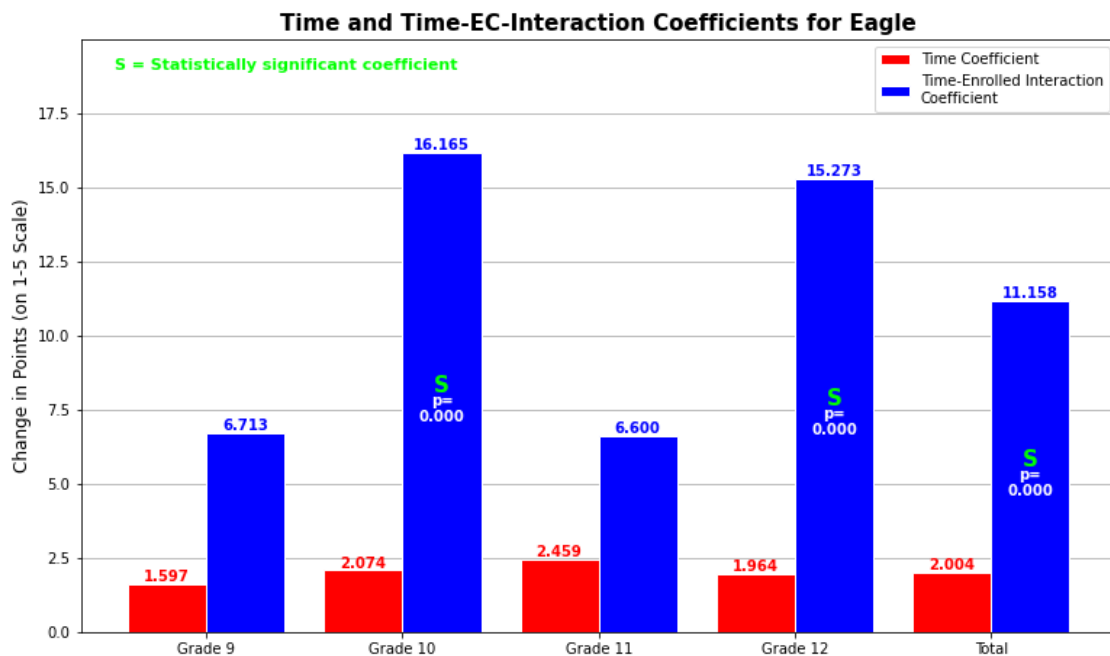
```

```
plt.savefig(file_string,dpi=400)
plt.show()
```

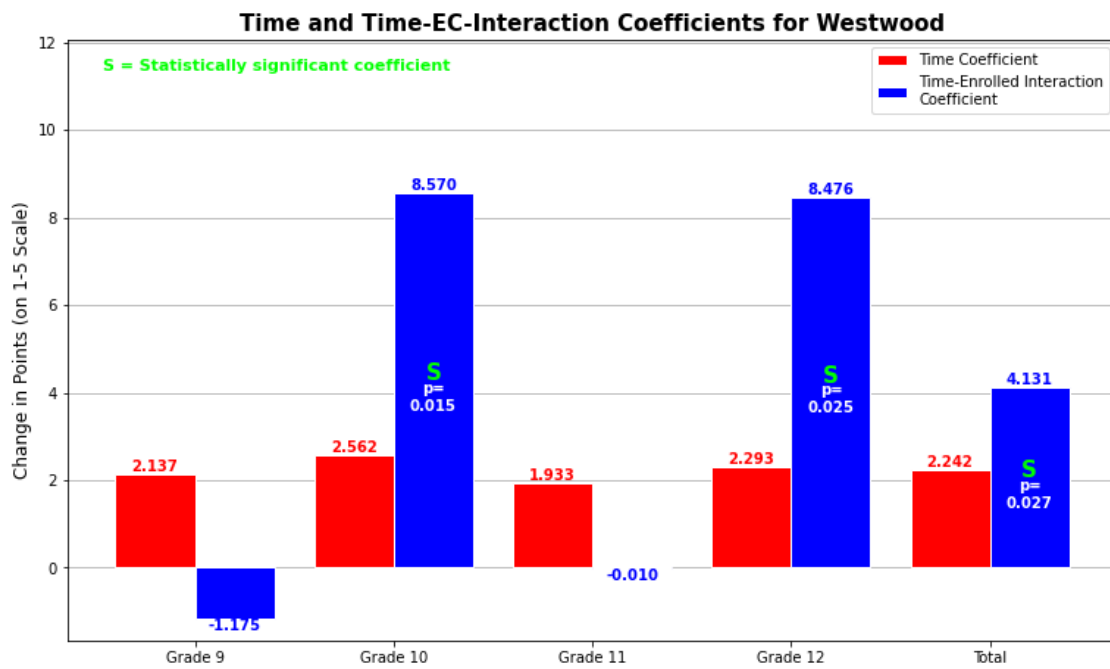
<class 'list'>



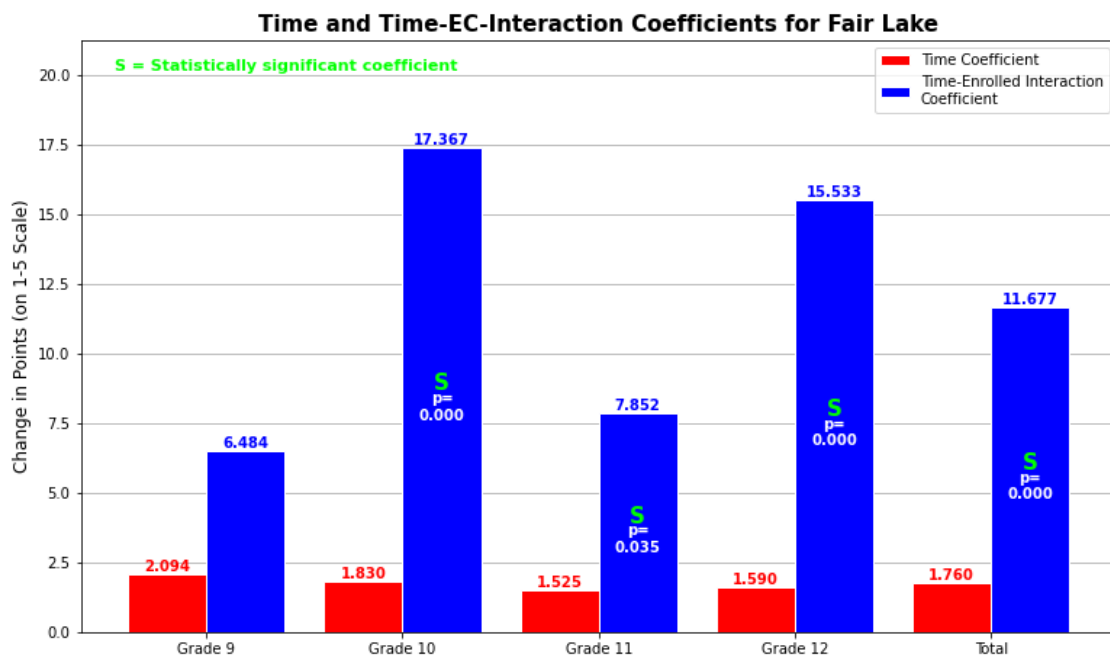
<class 'list'>



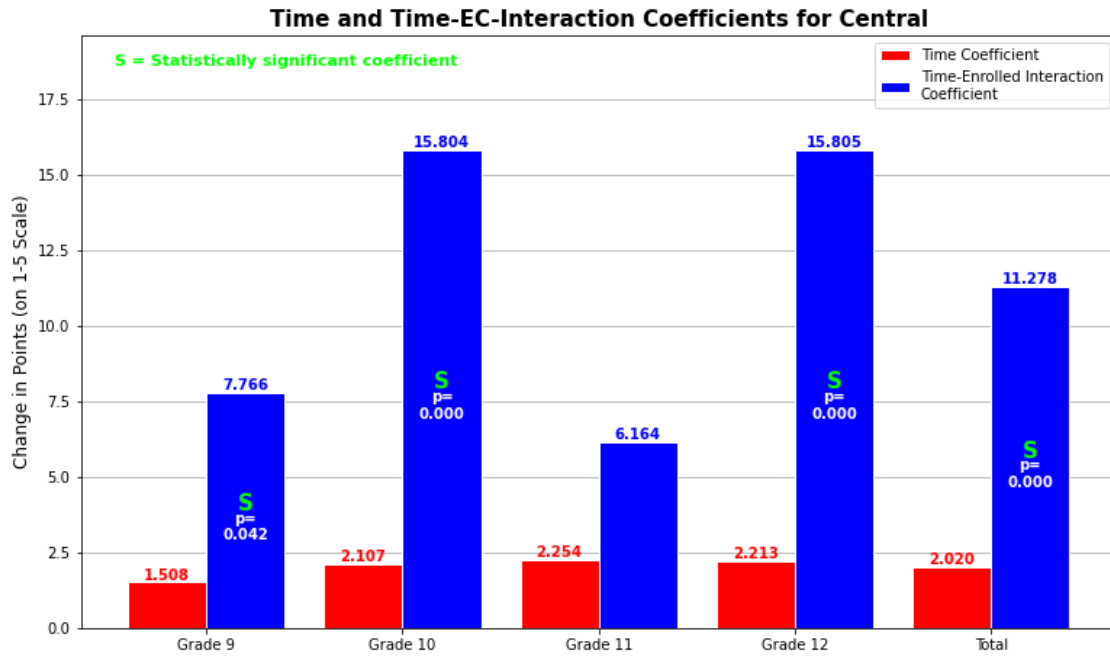
<class 'list'>



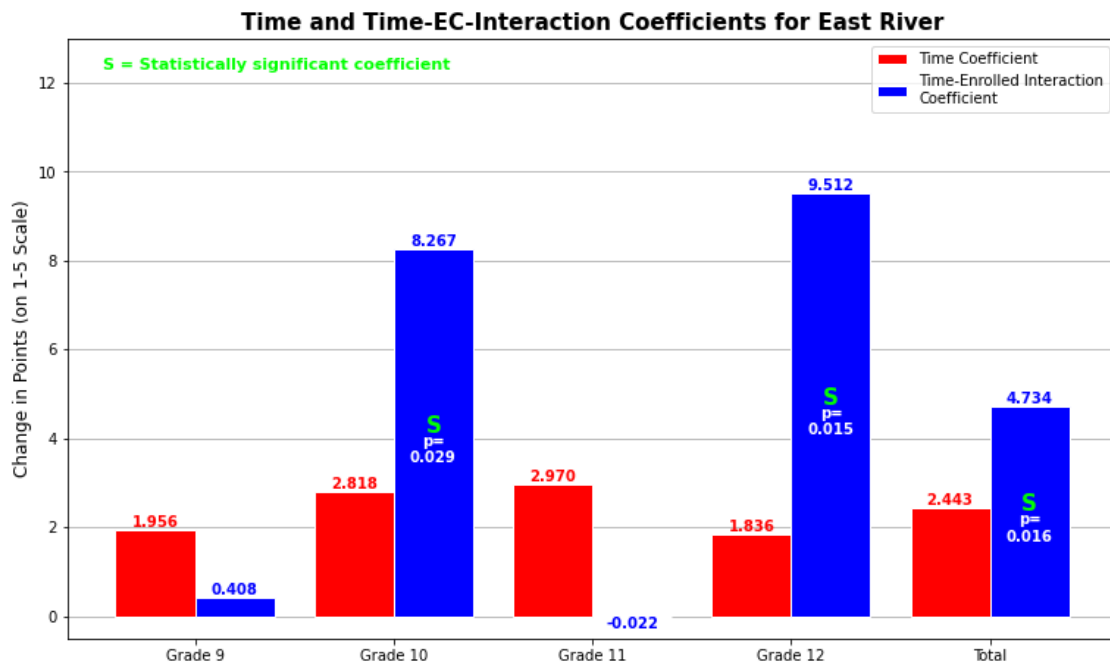
<class 'list'>



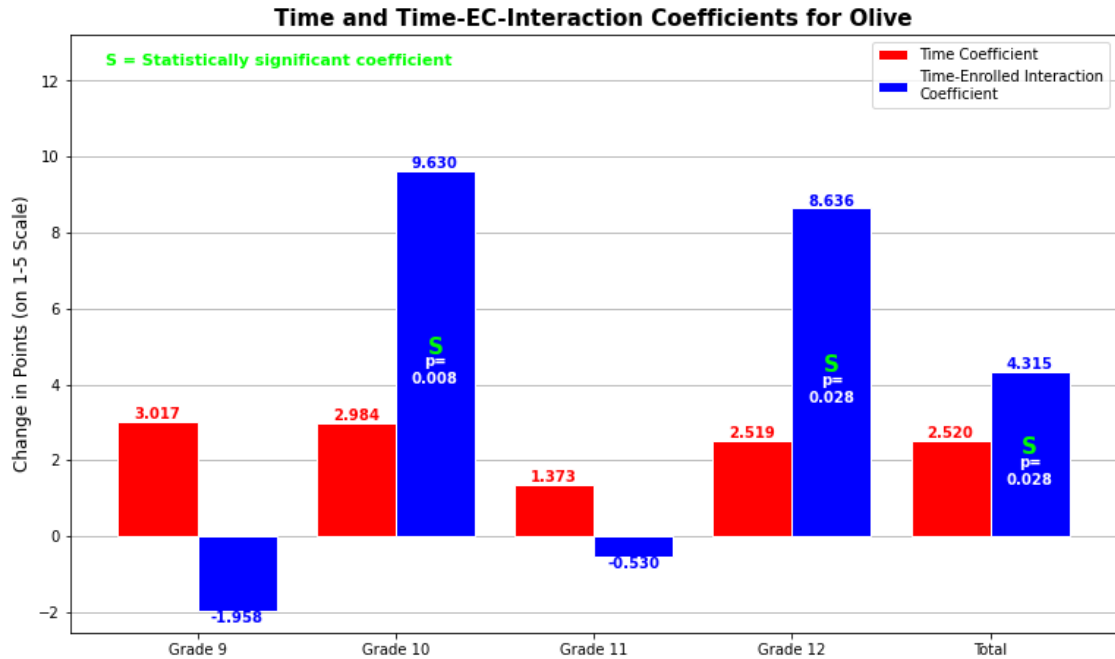
<class 'list'>



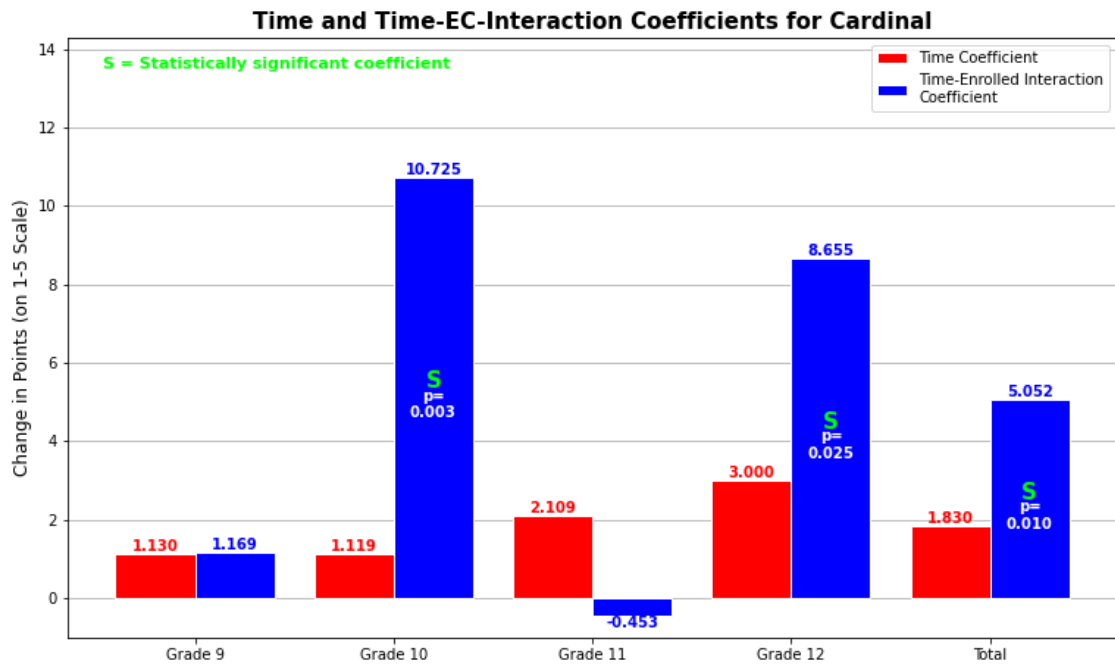
<class 'list'>



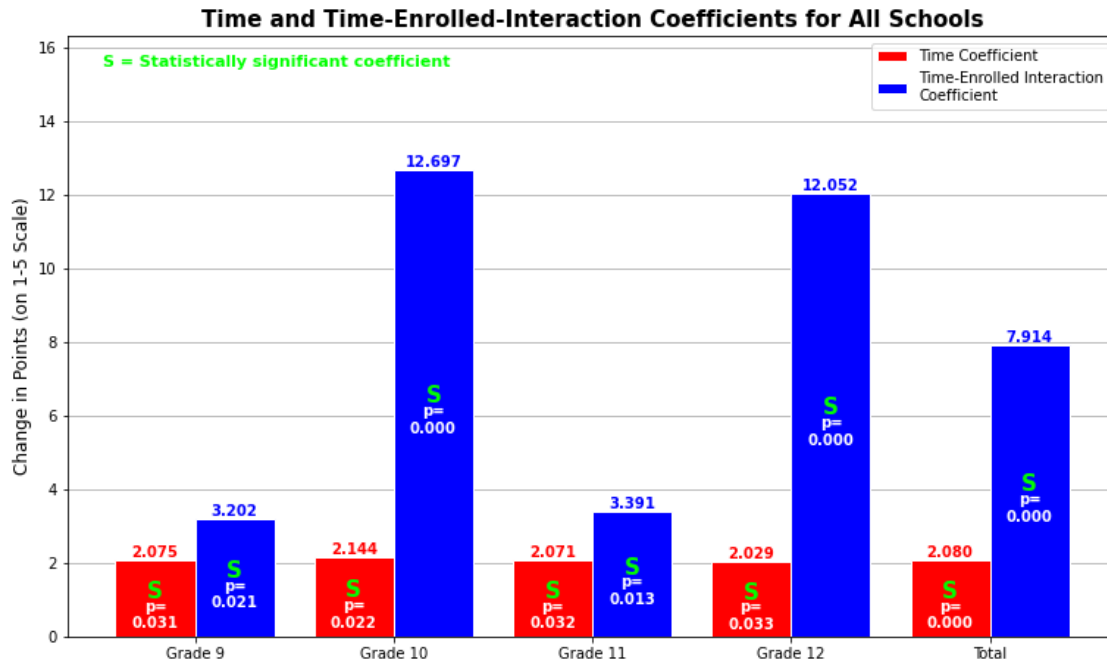
<class 'list'>



<class 'list'>



<class 'list'>



```
[18]: with pd.ExcelWriter('regression_tables.xlsx') as writer:
        for i in range(len(regression_table_list)):
            regression_table_list[i][1].to_excel(writer, sheet_name =
            ↪ regression_table_list[i][0])
```

```
[19]: end_time = time.time()
run_time = end_time - start_time
run_minutes = run_time // 60
run_seconds = run_time % 60
print("Completed run at",time.ctime(end_time),"(local time)")
print("Total run time:",'{:.2f}'.format(run_time),"second(s)"
    ↪ ("+"str(run_minutes),"minute(s) and", '{:.2f}'.
    ↪ format(run_seconds),"second(s))") # Only valid when the program is run
    ↪ nonstop from start to finish
```

Completed run at Mon Jul 5 23:44:46 2021 (local time)  
Total run time: 36.82 second(s) (0.0 minute(s) and 36.82 second(s))