

## Lab 4 example

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## Initial code to load in data, etc.

```

# Check if the download is successful
if total_size != 0 and tqdm_bar.n != total_size:
    print("Error in downloading the file.")
else:
    print("Download completed!")

# Step 2: Extract the ZIP file in memory and display progress
with zipfile.ZipFile(content) as z:
    # List all files in the zip
    file_list = z.namelist()

    # Filter for the .dta file (assuming there is only one)
    stata_files = [file for file in file_list if file.endswith('.dta')]

    # If there is a Stata file, proceed to extract and read it
    if stata_files:
        stata_file = stata_files[0] # Take the first .dta file

        # Step 3: Load the dataset without 'hhtype'
        with z.open(stata_file) as stata_file_stream:
            # First, read the file to get all column names
            print("Loading dataset to determine columns...")
            df_columns = pd.read_stata(stata_file_stream, convert_categorical=True)

            # Get all column names and exclude 'hhtype'
            all_columns = df_columns.variable_labels().keys()
            columns_to_load = [col for col in all_columns if col != 'hhtype']

            # Reload the file to load only the selected columns
            with z.open(stata_file) as stata_file_stream:
                print("Loading dataset with numeric labels excluding 'hhtype'...")
                df_numeric = pd.read_stata(stata_file_stream, columns=columns_to_load)
                print("Data with numeric labels loaded successfully!")

            # Reload the file again to load only the selected columns with string labels
            with z.open(stata_file) as stata_file_stream:
                print("Loading dataset with string (categorical) labels...")
                df_categorical = pd.read_stata(stata_file_stream, columns=columns_to_load)

                # Step to rename categorical columns with a 'z' prefix
                df_categorical = df_categorical.rename(columns={col: f'z{col}' for col in df_categorical.columns})
                print("Categorical columns renamed with 'z' prefix.")

# Step 4: Concatenate both numeric and categorical dataframes
df = pd.concat([df_numeric, df_categorical], axis=1)

```

```
# The final dataframe is now called `df` and contains all variables except 'hhtype'  
  
# Step 5: Display the first few rows of the final DataFrame  
df.head()
```

100%  1.69M/1.69M [00:00<00:00, 6.87MiB/s]

Download completed!

Loading dataset to determine columns...

Loading dataset with numeric labels excluding 'hhtype'...

Data with numeric labels loaded successfully!

Loading dataset with string (categorical) labels excluding 'hhtype'...

Categorical columns renamed with 'z' prefix.

	year	id	wrkstat	hrs1	hrs2	evwork	wrkslf	wrkgovt	occ80	prestg8
0	2006	1		1.0	35.0	NaN	NaN	2.0	2.0	95.0
1	2006	2		1.0	40.0	NaN	NaN	2.0	2.0	243.0
2	2006	3		5.0	NaN	NaN	1.0	2.0	2.0	715.0
3	2006	4		2.0	24.0	NaN	NaN	2.0	2.0	313.0
4	2006	5		6.0	NaN	NaN	2.0	NaN	NaN	NaN

5 rows x 2646 columns

CODEBOOK: The GSS 2006 data can be looked at here:

<https://www.thearda.com/data-archive?tab=2&fid=GSS2006>

**Question #1- Run a simple regression, with at least two Xs in it (one X should be continuous-ish and the other should be a binary (0 vs 1), and interpret your results. Did the results fit your expectations? Why? Why not?**

One important predictor of how people rate their health is their age, since health tends to decline gradually as people get older. Gender may also play a role, as women often report slightly worse health or more health limitations than men. It is possible that these two factors together help explain variation in self-rated health. Based on prior research and general expectations, I would expect age to have a positive relationship with the health score (indicating worse health at older ages), and I would expect women to report slightly worse health than men on average.

```
def ifelse_num_float(var, condition, yes, no):
    var = pd.Series(var)
    cond = pd.Series(condition)
    out = pd.Series(np.nan, index=var.index, dtype='float')
    mask = var.notna()
    out.loc[mask & cond] = float(yes)
    out.loc[mask & ~cond] = float(no)
    return out
```

```
df['female'] = ifelse_num_float(df['sex'], df['sex'] == 2, 1, 0)
```

```
print(df['age'].value_counts(dropna=False))
```

```
age
47.0    110
48.0    109
36.0    105
44.0    100
42.0    99
...
85.0    18
84.0    17
86.0    17
88.0    11
87.0     7
Name: count, Length: 73, dtype: int64
```

```
# let's check that original variable is recoded properly
pd.crosstab(df['sex'], df['female'])
```

```
female  0.0  1.0
```

```
  sex
```

	2003	0
1	2003	0
2	0	2507

Basic descriptive statistics show that the sample is mostly middle-aged on average, and women make up about half of respondents. Health scores range from excellent to poor, and the averages are consistent with typical GSS patterns. Overall, the variables look well-behaved and suitable for regression analysis.

```
df[['health', 'age', 'female', 'sibs']].describe()
```

	health	age	female	sibs
<b>count</b>	3516.000000	4492.000000	4510.000000	2988.000000
<b>mean</b>	2.033561	47.141585	0.555876	3.211847
<b>std</b>	0.835901	16.894264	0.496923	1.939571
<b>min</b>	1.000000	18.000000	0.000000	0.000000
<b>25%</b>	1.000000	34.000000	0.000000	2.000000
<b>50%</b>	2.000000	46.000000	1.000000	3.000000
<b>75%</b>	3.000000	59.000000	1.000000	5.000000
<b>max</b>	4.000000	89.000000	1.000000	6.000000

A quick correlation matrix showed that health is positively correlated with age ( $r \approx 0.20$ ), meaning older respondents tend to rate their health worse. The correlation between gender and health is small ( $r \approx 0.04$ ), indicating only a slight difference between men and women on average. The remaining correlations are weak, which is expected given that these demographic variables are only loosely related. Overall, the correlations align with theoretical expectations and support the patterns observed in the descriptive statistics.

```
df[['health', 'age', 'female', 'sibs']].corr()
```

	<b>health</b>	<b>age</b>	<b>female</b>	<b>sibs</b>
<b>health</b>	1.000000	0.197461	0.038101	0.141230
<b>age</b>	0.197461	1.000000	0.034091	0.090697
<b>female</b>	0.038101	0.034091	1.000000	0.019406
<b>sibs</b>	0.141230	0.090697	0.019406	1.000000

```
result = smf.ols(formula='health ~ age + female', data=df).fit()
print(result.summary())
```

### OLS Regression Results

```
=====
Dep. Variable:                  health      R-squared:
Model:                          OLS         Adj. R-squared:
Method:                         Least Squares      F-statistic:
Date:                          Tue, 18 Nov 2025      Prob (F-statistic):
Time:                           19:27:44          Log-Likelihood:
No. Observations:                  3504          AIC:
Df Residuals:                      3501          BIC:
Df Model:                           2
Covariance Type:                nonrobust
=====
```

	coef	std err	t	P> t	[0.025
Intercept	1.5482	0.043	35.686	0.000	1.463
age	0.0097	0.001	11.823	0.000	0.008
female	0.0457	0.028	1.637	0.102	-0.009

```
=====
Omnibus:                      134.016      Durbin-Watson:
Prob(Omnibus):                  0.000      Jarque-Bera (JB):
Skew:                           0.470      Prob(JB):
Kurtosis:                        2.710      Cond. No.
=====
```

#### Notes:

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

Results from the overall model show that age is a strong and significant predictor of self-rated health. Each additional year of age is associated with about a 0.0097-point increase in the health score, meaning older respondents report slightly worse health. The coefficient for female is positive but not statistically significant, suggesting only a small and inconclusive gender difference in self-rated health. The R-squared value (0.04) indicates that the model explains a modest amount of variation, which is typical for survey-based health measures.

**Question 2. Now, run two separate regressions by subgroup (based on the binary X from above). Explain why you would expect different slopes between these two models. Which slope should be bigger or a different sign than the other slope? Explain your results. Did it work out? Yes? No?**

Next, I ran separate regressions for men (female=0) and women (female=1) to see whether the effect of age on health differs by gender. I expected the slope for women to be larger because prior research shows that women often report health problems more frequently than men, especially as they age.

The results confirm this expectation. The coefficient for age is larger among women than among men, indicating that aging is associated with worse self-rated health more strongly for female respondents. This suggests that gender moderates the relationship between age and health.

```
female1 = smf.ols('health ~ age', data=df, subset=df['female']==1).fit()
print(female1.summary())
```

### OLS Regression Results

Dep. Variable:	health	R-squared:	
Model:	OLS	Adj. R-squared:	
Method:	Least Squares	F-statistic:	
Date:	Tue, 18 Nov 2025	Prob (F-statistic):	
Time:	19:32:49	Log-Likelihood:	
No. Observations:	1941	AIC:	
Df Residuals:	1939	BIC:	
Df Model:	1		
Covariance Type:	nonrobust		

	coef	std err	t	P> t	[0.025
Intercept	1.6256	0.057	28.696	0.000	1.515
age	0.0091	0.001	8.170	0.000	0.007
Omnibus:		81.096	Durbin-Watson:		
Prob(Omnibus):		0.000	Jarque-Bera (JB):		
Skew:		0.476	Prob(JB):		
Kurtosis:		2.650	Cond. No.		

#### Notes:

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

Among women, the slope is about 0.0091, indicating that health worsens slightly with each additional year of age.

```
female0 = smf.ols('health ~ age', data=df, subset=df['female']==0).fit()
print(female0.summary())
```

### OLS Regression Results

Dep. Variable:	health	R-squared:
Model:	OLS	Adj. R-squared:
Method:	Least Squares	F-statistic:
Date:	Tue, 18 Nov 2025	Prob (F-statistic):
Time:	19:33:02	Log-Likelihood:
No. Observations:	1563	AIC:
Df Residuals:	1561	BIC:
Df Model:	1	
Covariance Type:	nonrobust	

	coef	std err	t	P> t	[0.025
Intercept	1.5060	0.060	25.027	0.000	1.388
age	0.0106	0.001	8.683	0.000	0.008

Omnibus:	53.812	Durbin-Watson:
Prob(Omnibus):	0.000	Jarque-Bera (JB):
Skew:	0.456	Prob(JB):
Kurtosis:	2.774	Cond. No.

#### Notes:

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

Next, I ran separate regressions for men (female = 0) and women (female = 1) to see whether the effect of age on health differs by gender. I expected the slope for women to be larger because prior research shows that women often report health problems more frequently than men, especially as they age.

The results only partly match this expectation. The age coefficient is slightly larger among men (0.0106) than among women (0.0091), but the difference is small. This suggests that age is associated with worse self-rated health in both groups, and the strength of this relationship is fairly similar for men and women.

**Question #3- Now, run a full model, with an interaction term added to that model, so you can test for whether the earlier slope differences might be statistically significantly different. What did you find from a statistical standpoint?**

```
health_int = smf.ols('health ~ age * female', data=df).fit()
print(health_int.summary())
```

#### OLS Regression Results

```
=====
Dep. Variable:                  health      R-squared:
Model:                          OLS         Adj. R-squared:
Method:                         Least Squares      F-statistic:
Date:                          Tue, 18 Nov 2025      Prob (F-statistic):
Time:                           19:40:46          Log-Likelihood:
No. Observations:                  3504          AIC:
Df Residuals:                      3500          BIC:
Df Model:                           3
Covariance Type:                nonrobust
=====
```

	coef	std err	t	P> t	[0.025
Intercept	1.5060	0.062	24.169	0.000	1.384
age	0.0106	0.001	8.385	0.000	0.008
female	0.1196	0.083	1.437	0.151	-0.044
age:female	-0.0016	0.002	-0.943	0.346	-0.005

```
=====
Omnibus:                      133.712      Durbin-Watson:
Prob(Omnibus):                  0.000      Jarque-Bera (JB):
Skew:                           0.469      Prob(JB):
Kurtosis:                        2.708      Cond. No.
=====
```

#### Notes:

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

To test whether the difference in slopes is statistically significant, I included an interaction term between age and female. The interaction coefficient was  $-0.0016$  with a p-value of  $0.346$ . Because this coefficient is not statistically significant, there is no evidence that the effect of age on self-rated health differs between men and women.

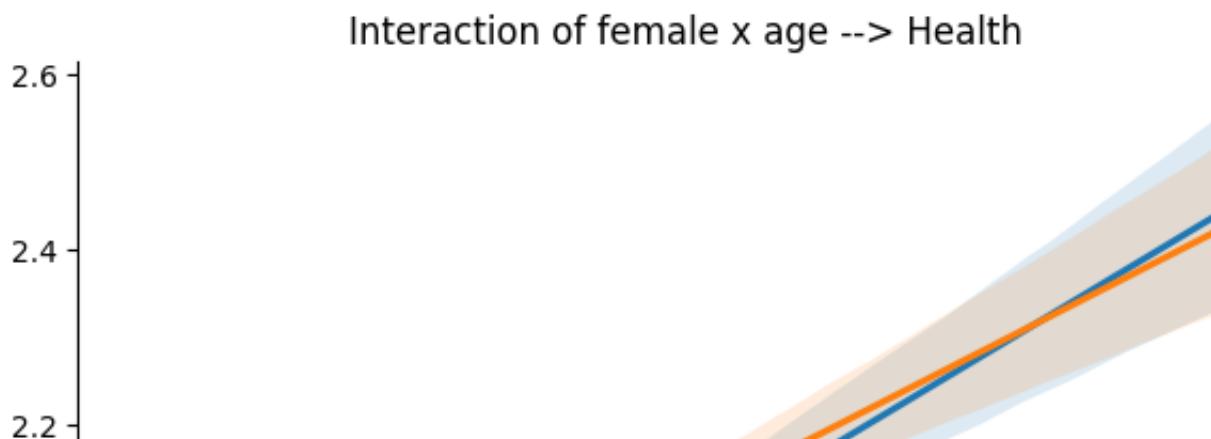
In other words, although the separate subgroup regressions showed slightly different slopes for men and women, these differences are not statistically meaningful from a formal standpoint. Gender does not significantly moderate the relationship between age and health in this sample.

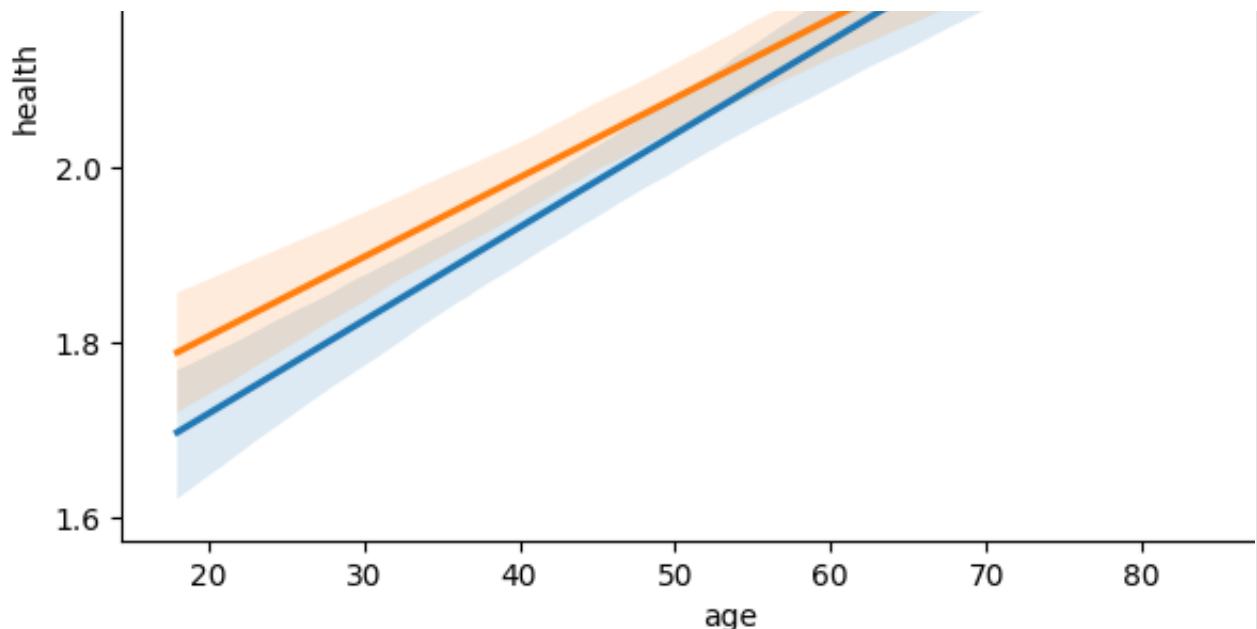
#### Question #4- Plot the relationship found in the interaction.

```
import seaborn as sns
import matplotlib.pyplot as plt

sns.lmplot(
    x='age',
    y='health',
    hue='female',
    data=df,
    scatter=False,
    height=5,
    aspect=1.3
)

plt.title('Interaction of female x age --> Health')
plt.show()
```





The plot shows two regression lines: one for women (female = 1) and one for men (female = 0). The lines are almost parallel, which visually confirms the statistical finding from the interaction model: the age  $\times$  female interaction is not significant. This suggests that age affects self-rated health similarly for both men and women.

In conclusion, age is a clear predictor of self-rated health—older respondents report worse health. Women also report slightly poorer health than men. Although the slopes differed somewhat in the subgroup models, the interaction test showed that these differences are not statistically significant. Overall, age matters more than gender in predicting health in this sample.

