

Problem Set #3

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CreationDate: [Feb_10](#)

Title: [Problem Sets #3 experiments and regressions](#)

Notes: [Colab](#)

In-class notebook exercises

1.1 Replicate and simulate a real study

1.1.1

```
print(r_col3.summary())
print(r_col4.summary())
```

1.1.2

- *Column 4* has more control variables
- to calculate the *R Square*

```
print(r_col3.rsquared) ## --> 0.0671339542940943
print(r_col4.rsquared) ## --> 0.10819457890249662
```

Column 4 has a higher R-squared, because it contains more control variables hence the model has more explanatory power;

- to calculate the *Std.ev*

```
print(r_col3.bse.rem_any) ## --> 0.009153114318622113
print(r_col4.bse.rem_any) ## --> 0.008949743622819572
```

Column 4 has a smaller a standard error on the estimated *ATE*;

1.1.3

```

=====
                        Model 1   Model 2
-----
d                        0.0317*** 0.0324***
                        (0.0082) (0.0082)
grad_high_school                0.1043***
                                (0.0080)
R-squared                   0.0011   0.0135
No. coefficients 2           3
=====
Standard errors in parentheses.
* p<.1, ** p<.05, ***p<.01

```

- for the estimation of the effect of `rem_any` in the experiments;
the estimated coefficients and the standard error are listed as follow

```

print(r_col3.params.rem_any) ## --> 0.03185505049068642
print(r_col4.params.rem_any) ## --> 0.03186131518614749
print(r_sim_biv.params.d)     ## --> 0.031709816056386286
print(r_sim_control.params.d) ## --> 0.03240621068111807

```

- the estimated coefficients and the standard error from the simulation are pretty close to real data;

1.1.4

- set the `beta_hs = 0.35` for the simulation
- set `B = 1000`

similar to the `compare_lpm_prop_test` function, we define a new function

```

def new_simulation_function(
    N = 10000,
    beta_hs = 0.35,
    ATE = 0.032
):
    grad_high_school = np.random.binomial(n=1, p=0.5, size=N)
    D = np.random.binomial(n=1, p=0.61, size=N)
    baseline_probability = 0.25 + beta_hs * grad_high_school
    Y0 = np.random.binomial(n=1, p=baseline_probability)
    Y1 = np.random.binomial(n=1, p=baseline_probability + ATE * D)
    dff = pd.DataFrame({
        'grad_high_school': grad_high_school,
        'd': D,
        'y0': Y0,
        'y1': Y1
    })
    dff['y'] = dff.eval("y1 * d + y0 * (1 - d)")

```

```
regr = sfa.ols("y ~ d", dff).fit()
a = regr.params['d']
return a
```

and we loop it;

```
B = 1000
res = pd.DataFrame([new_simulation_function() for i in np.arange(0, B)])
```

and we see the results;

	0	1	2	3	4	5	6	7
index	count	mean	std	min	25%	50%	75%	max
0	1000	0.0319545	0.0099911	-0.000497493	0.0254244	0.0324918	0.0388504	0.07187

1.1.5

for this question, we replicate what we did from the previous simulation;

and we plot the `params['d']` to show the estimated treatment effects;

