## **Exercise 6: Linear models and machine learning**

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March 23, 2020

## Problem 6.1.1 - Wrapping in a function

This first question should be straight forward to complete, we simply wrap the code in a def and return statement

```
def simulate(n = 1000, m = 1500, k = 10, plot = False):
    # [function code]
    return (d, z, Pi, gamma, nu), (y, x, delta, alpha, u)
```

with an if-statement ensuring that the plotting code is only run when plot = True. Notice that the function returns two tuples of data. This is simply to make unpacking results more intuitive, as we can call

```
aux, main = simulate()
d, z, Pi, gamma, nu = aux
y, X, delta, alpha, u = main
```

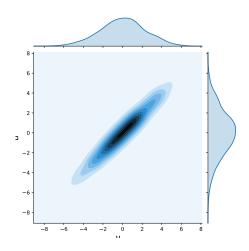
Now let us quickly discuss what the code is actually doing. The main part of the code is straight forward, it simulates data according to the two equations

$$\begin{aligned} d_i &= x_i' \gamma_0 + z_i' \delta_0 + u_i \\ y_i &= \alpha_0 d_i + x_i' \beta_0 + \epsilon_i \end{aligned} \tag{Auxilliary}$$

The important part happens in one of the first lines, namely

```
errors = np.random.multivariate_normal(mean = [0,0], cov = [[5,5],[5,5]], size = n)
```

which constructs  $u_i$ ,  $v_i$  as correlated random normals, as evident from figure 1. The consequence of this is that whenever  $u_i$  is high we also expect  $\epsilon_i$  to be high and vice versa. As we will see shortly this becomes an issue if we want to estimate the actual impact of  $d_i$  on  $y_i$ . i.e.  $\alpha_0$ .



**Fig 1.** Correlation structure of  $\nu_i$  and  $u_i$ .

## Problem 6.1.2 - The naive regression

In this problem we will see what quantity we can expect to estimate by running the regression

$$y_i \sim \beta_0 + \beta_1 d_i + \eta_i \tag{1}$$

If this regression provides unbiased estimates on  $d_i$  we will expect  $\hat{\beta}_1 \approx \alpha_0$ . Repeating the regression many times should reveal that  $\hat{\beta}_1$  falls evenly on either side of the true value  $\alpha_0$ . However, this is not what we will find, and the reason lies in the correlated errors.

Because  $u_i$  and  $\epsilon_i$  are positively correlated  $u_i > 0 \Rightarrow \mathbb{E}[\epsilon_i|u_i] > 0$  - i.e. positive errors in the main equation are associated with positive errors in the auxilliary equation. Assume for a moment that  $u_i$  and  $\epsilon_i$  are correlated such that  $\partial \epsilon_i/\partial u_i = \sigma$ . Then increasing  $u_i$  dicretly increases  $\epsilon_i$  by a proportional amount. In this case what happens when  $u_i = 1$ ?  $d_i$  increases by 1 above  $\mathbb{E}[d_i|z_i,x_i]$ , and  $y_i$  increase by  $\alpha_0 + \sigma$  above  $\mathbb{E}[y_i|x_i]$ . From observing only  $y_i$  and  $d_i$  we cannot separate out what changes in  $y_i$  were due to changes in  $d_i$  ( $\alpha_0$ ), and what changes were due to changes in  $u_i$  affecting  $\epsilon_i$  ( $\sigma$ ). For this reason we should expect a bias in our naive regression results. The code for this problem starts off with setting

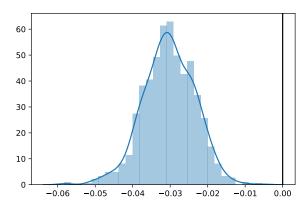


Fig 2. Distribution of estimated values for  $\hat{\beta}_1$  's deviation from truth in the naive regression.

up two lists and a for-loop. The first few lines of the loop simply simulates a new dataset and unpacks all of the return values.

```
A,B = [], []
for _ in range(1000):
    aux, main = simulate()
    d, z, Pi, gamma, nu = aux
    y, X, delta, alpha, u = main
```

Next we fit the naive regression

```
model = OLS(y, add_constant(d))
result = model.fit()
```

and store the true  $\alpha_0$  alongside  $\hat{\alpha}_0$  in the two lists.

```
A.append(alpha)
B.append(result.params[1])
```

The histogram can easily be drawn with seaborn,

```
sns.distplot([(a - b) for a,b in zip(A,B)])
plt.axvline(o,color = 'black')
```