

Causal Inference in Python: A Vignette

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Installation

CausalInference can be installed using `pip`, and will run provided the necessary dependencies are in place. On Ubuntu systems, the following commands should take care of all the essential steps if you are starting from scratch:

```
$ sudo apt-get update
$ sudo apt-get install python-pip python-numpy python-scipy
$ sudo pip install causalinference
```

Minimal Example

The main object of interest in CausalInference is the class `CausalModel`. It takes as inputs three NumPy arrays: `Y`, an N -vector of observed outcomes; `D`, an N -vector of treatment status indicators; and `X`, an N -by- K matrix of covariates. The following code snippet illustrates how the class can be invoked using random data simulated from NumPy's `random` module.

```
>>> import numpy as np
>>> from causalinference import *
>>> Y = np.random.rand(1000)
>>> D = np.random.randint(0, 2, 1000)
>>> X = np.random.rand(1000, 3)
>>> causal = CausalModel(Y, D, X)
```

Class Attributes and Methods

Once an instance of the class `CausalModel` has been created, it will contain a number of attributes and methods that are relevant for conducting a causal analysis. Tables 1 and 2 contain a brief description of these attributes and methods.

Attribute	Description
<code>N</code> , <code>N_c</code> , <code>N_t</code> , <code>K</code>	Integers indicating sample sizes and number of covariates.
<code>covariates</code>	Dictionary-like object containing summary statistics for the covariate variables.
<code>pscore</code>	Dictionary-like object containing propensity score data, including estimated logistic regression coefficients, predicted propensity score, maximized log-likelihood, and the lists of the linear and quadratic terms that are included in the regression.
<code>cutoff</code>	Floating point number specifying the cutoff point for trimming on propensity score.
<code>blocks</code>	Either an integer indicating the number of equal-sized blocks to stratify the sample into, or a list of ascending numbers specifying the boundaries of the strata.
<code>strata</code>	List-like object containing the list of stratified propensity bins.
<code>est</code>	Dictionary-like object containing treatment effect estimates for each estimator used.

Table 1: Attributes of the class `CausalModel`. Invoking `print` on any of the dictionary- or list-like attribute above yields customized summary tables. Note that some attributes are only created after the relevant methods have been called.

Using CausalInference Interactively

```
>>> print causal.covariates
```

Covariates Summary

Covariate	Controls (N_c=267)		Treated (N_t=233)		Nor-diff
	Mean	S.d.	Mean	S.d.	
X0	-0.507	0.936	0.390	0.907	0.972
X1	-0.295	0.902	0.380	0.905	0.748
X2	-0.514	0.950	0.367	0.871	0.967

Estimate propensity score.

```
>>> causal.propensity_s()
```

Method	Description
<code>restart</code>	Reinitializes data to original inputs, and drop any estimated results.
<code>propensity</code>	Estimates via logit the propensity score using specified linear and quadratic terms.
<code>propensity_s</code>	Estimates via logit the propensity score using the covariate selection algorithm of Imbens and Rubin (2015).
<code>trim</code>	Trims data based on propensity score using the threshold specified by the attribute <code>cutoff</code> .
<code>trim_s</code>	Trims data based on propensity score using the cutoff selected by the procedure of Crump, Hotz, Imbens, and Mitnik (2008).
<code>stratify</code>	Stratifies the sample based on propensity score as specified by the attribute <code>blocks</code> .
<code>stratify_s</code>	Stratifies the sample based on propensity score using the bin selection procedure suggested by Imbens and Rubin (2015).
<code>blocking</code>	Estimates average treatment effects using regression within blocks.
<code>matching</code>	Estimates average treatment effects using matching with replacement.
<code>weighting</code>	Estimates average treatment effects using the Horvitz-Thompson weighting estimator modified to incorporate covariates.
<code>ols</code>	Estimates average treatment effects using least squares.

Table 2: Methods of the class `CausalModel`. Invoke `help` on any of the above methods for more detailed documentation.

```
>>> print causal.pscore
```

```
Estimated Parameters of Propensity Score
```

	Coef.	S.e.	z	P> z	[95% Conf. int.]	
Intercept	-0.018	0.136	-0.133	0.447	-0.284	0.248
X0	1.785	0.192	9.305	0.000	1.409	2.161
X2	1.789	0.195	9.177	0.000	1.407	2.171
X1	1.765	0.203	8.678	0.000	1.366	2.163

Trimming based on propensity score.

```
>>> causal.trim_s()
```

```
>>> causal.N, causal.N_c, causal.N_t
```

```
(281, 143, 138)
```

```
>>> print causal.covariates
```

Covariates Summary

Covariate	Controls (N_c=143)		Treated (N_t=138)		Nor-diff
	Mean	S.d.	Mean	S.d.	
X0	-0.171	0.770	0.163	0.893	0.401
X1	-0.046	0.848	0.066	0.765	0.139
X2	-0.178	0.780	0.125	0.802	0.383

Stratifying sample based on propensity score.

```
>>> causal.stratify_s()
```

```
>>> print causal.strata
```

Stratification Summary

Stratum	Propensity score		N_c	N_t	Ave. p-score		Within Est.
	Min.	Max.			Controls	Treated	
1	0.085	0.513	104	37	0.230	0.267	2.274
2	0.518	0.631	18	17	0.570	0.589	2.996
3	0.632	0.753	7	28	0.678	0.699	3.447
4	0.757	0.915	14	56	0.835	0.842	3.367

Computing the blocking estimator.

```
>>> causal.blocking()
```

```
>>> print causal.est
```

Treatment Effect Estimates

Blocking

	Est.	S.e.	z	P> z	[95% Conf. int.]	
ATE	2.782	0.127	21.941	0.000	2.534	3.031
ATT	3.045	0.152	20.046	0.000	2.747	3.342
ATC	2.530	0.134	18.925	0.000	2.268	2.792