

Pre-conference Workshop: Introduction to Causal Inference for Epidemiologists (II). XXXVII-SEE, Oviedo, Spain 2019

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PART II: ESTIMATION and INFERENCE

ESTIMANDS, ESTIMATORS, and STATISTICAL INFERENCE

ESTIMANDS

Most common ESTIMANDS

<https://migariane.github.io/DeltaMethodEpi.nb.html>

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- $RD = EY(1) - EY(0)$

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Estimands in Causal Inference

- $RD = EY(1) - EY(0)$
- $RR = EY(1) / EY(0)$
- $MOR = EY(1) \times E(1 - Y(0)) / EY(0) \times E(1 - Y(1))$

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G-Formula, (Robins, 1986)

G-Formula for the **identification** of the ATE with observational data

$$\begin{aligned} E(Y^a) &= \sum_y E(Y^a \mid W = w)P(W = w) \\ &= \sum_y E(Y^a \mid A = a, W = w)P(W = w) \text{ by consistency} \\ &= \sum_y E(Y = y \mid A = a, W = w)P(W = w) \text{ by ignorability} \end{aligned}$$

The **ATE**=

$$\sum_w \left[\sum_y P(Y = y \mid A = 1, W = w) - \sum_y P(Y = y \mid A = 0, W = w) \right] P(W = w)$$

$$P(W = w) = \sum_{y,a} P(W = w, A = a, Y = y)$$

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G-Formula

- The sums is generic notation. In reality, likely involves sums and integrals (we are just integrating out the W 's).
- The **g-formula** is a **generalization of standardization** and allow to estimate unbiased treatment effect estimates.

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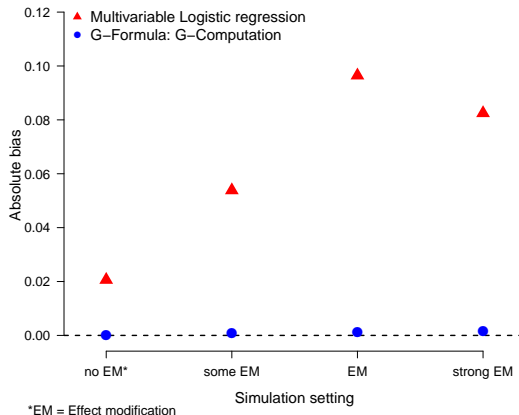
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G-Formula

- In the presence of **treatment or exposure effect heterogeneity** (i.e. effect modification) simple conditional estimate is biased.
- The **g-formula** allows to estimate the **marginal average treatment effect** (i.e., effect for the population averaged across the different level of covariates) allowing to estimate unbiased treatment effect estimates.

Treatment heterogeneity



Simulation treatment heterogeneity and estimation using OR and MOR, $n = 1,000$.

Luque-Fernandez et al.(2019) Effect modification and collapsibility when estimating the effect of public health interventions: A Monte Carlo simulation comparing classical multivariable regression adjustment versus the G-Formula, based on a cancer epidemiology illustration. AJPH <https://maluque.netlify.com/project/hetmor/>

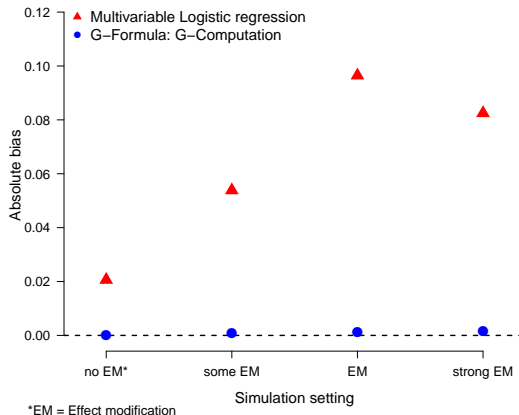
COR

$$COR = \frac{\frac{P(Y=1|A=1,W)}{(1 - P(Y=1|A=1,W))}}{\frac{P(Y=1|A=0,W)}{(1 - P(Y=1|A=0,W))}}$$

MOR

$$MOR \text{ (G-Formula)} = \frac{\frac{\sum_w P(Y=1|A=1,W=w) P(W=w)}{(1 - \sum_w P(Y=1|A=1,W=w) P(W=w))}}{\frac{\sum_w P(Y=1|A=0,W=w) P(W=w)}{(1 - \sum_w P(Y=1|A=0,W=w) P(W=w))}}$$

Treatment heterogeneity: simulations



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RA: Regression-adjustment

$$\widehat{ATE}_{RA} = N^{-1} \sum_{i=1}^N [E(Y_i | A = 1, W_i) - E(Y_i | A = 0, W_i)]$$

$$m_A(w_i) = E(Y_i | A_i = A, W_i)$$

$$\widehat{ATE}_{RA} = N^{-1} \sum_{i=1}^N [\hat{m}_1(w_i) - \hat{m}_0(w_i)]$$

IPTW (Inverse probability treatment weighting)

Survey theory (Horvitz-Thompson)

$$\hat{P}_i = E(A_i | W_i) ; \text{ So , } \frac{1}{\hat{p}_i} , \text{ if } A = 1 \text{ and , } \frac{1}{(1 - \hat{p}_i)} , \text{ if } A = 0$$

Average over the total number of individuals

$$\widehat{ATE}_{IPTW} = N^{-1} \sum_{i=1}^N \frac{A_i Y_i}{\hat{p}_i} - N^{-1} \sum_{i=1}^N \frac{(1 - A_i) Y_i}{(1 - \hat{p}_i)}$$

AIPTW (Augmented Inverse probability treatment weighting)

Solving Estimating Equations

$$\widehat{ATE}_{AIPTW} =$$

$$N^{-1} \sum_{i=1}^N [(Y(1) | A_i = 1, W_i) - (Y(0) | A_i = 0, W_i)] +$$

$$N^{-1} \sum_{i=1}^N \left(\frac{(A_i = 1)}{P(A_i = 1 | W_i)} - \frac{(A_i = 0)}{P(A_i = 0 | W_i)} \right) [Y_i - E(Y | A_i, W_i)]$$

ATE estimators: drawbacks

Nonparametric

- Curse of dimensionality (sparsity: zero empty cell)

Parametric

- Parametric models are **misspecified** (all models are wrong but some are useful, Box, 1976), and **break down** for high-dimensional data.
- **(RA)** Issue: extrapolation and biased if misspecification, no information about treatment mechanism.
- **(IPTW)** Issue: sensitive to curse of dimensionality, inefficient in case of extreme weights and biased if misspecification. Non information about the outcome.

Double-robust (DR) estimators

Prons: Semi-parametric Double-Robust Methods

- DR methods give **two chances at consistency** if any of two nuisance parameters is consistently estimated.
- DR methods are **less sensitive to course of dimensionality**.

Cons: Semi-parametric Double-Robust Methods

- DR methods are unstable and inefficient if the propensity score (PS) is small (**violation of positivity assumption**) (vand der Laan, 2007).
- AIPW and IPTW-RA do not respect the **limits of the boundary space of Y**.
- **Poor performance if dual misspecification** (Benkeser, 2016).

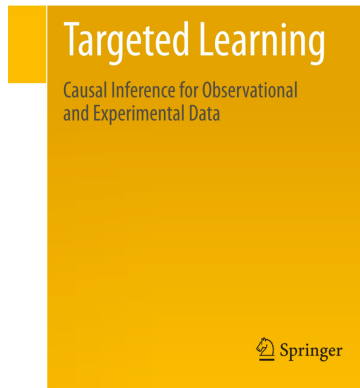
Targeted Maximum Likelihood Estimation (TMLE)

Pros: TMLE

- (TMLE) is a general algorithm for the construction of **double-robust**, **semiparametric** MLE, efficient **substitution** estimator (Van der Laan, 2011)
- **Better performance** than competitors has been largely documented (Porter, et. al., 2011).
- (TMLE) **Respect bounds on Y**, **less sensitive** to **misspecification** and to **near-positivity** violations (Benkeser, 2016).
- (TMLE) **Reduces bias** through **ensemble learning** if misspecification, even dual misspecification.
- For the ATE, **Inference** is based on the **Efficient Influence Curve**. Hence, the **CLT** applies, making inference easier.

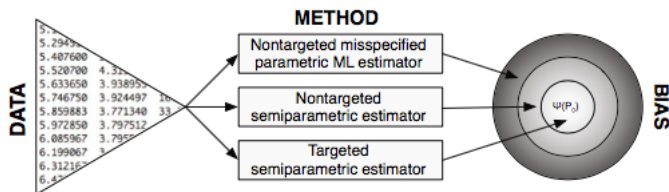
Cons: TMLE

- The procedure is only available in R: **tmle** package (Gruber, 2011).



¹ **Source:** Mark van der Laan and Sherri Rose. Targeted learning: causal inference for observational and experimental data. Springer Series in Statistics, 2011.

Why Targeted learning?



² **Source:** Mark van der Laan and Sherri Rose. Targeted learning: causal inference for observational and experimental data. Springer Series in Statistics, 2011.

MC simulations: Luque-Fernandez et al., 2017 (American Journal of Epidemiology)

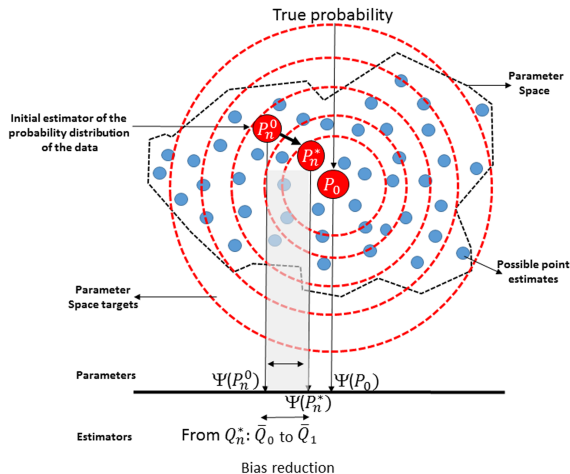
	ATE		BIAS (%)		RMSE		95%CI coverage (%)	
	N=1,000	N=10,000	N=1,000	N=10,000	N=1,000	N=10,000	N=1,000	N=10,000
First scenario* (correctly specified models)								
True ATE	-0.1813							
Naïve	-0.2234	-0.2218	23.2	22.3	0.0575	0.0423	77	89
AIPTW	-0.1843	-0.1848	1.6	1.9	0.0534	0.0180	93	94
IPTW-RA	-0.1831	-0.1838	1.0	1.4	0.0500	0.0174	91	95
TMLE	-0.1832	-0.1821	1.0	0.4	0.0482	0.0158	95	95
Second scenario ** (misspecified models)								
True ATE	-0.1172							
Naïve	-0.0127	-0.0121	89.2	89.7	0.1470	0.1100	0	0
BFit AIPTW	-0.1155	-0.0920	1.5	11.7	0.0928	0.0773	65	65
BFit IPTW-RA	-0.1268	-0.1192	8.2	1.7	0.0442	0.0305	52	73
TMLE	-0.1181	-0.1177	0.8	0.4	0.0281	0.0107	93	95

*First scenario : correctly specified models and near-positivity violation

**Second scenario: misspecification, near-positivity violation and adaptive model selection

³ Data-Adaptive Estimation for Double-Robust Methods in Population-Based Cancer Epidemiology: Risk differences for lung cancer mortality by emergency presentation (2017). AJE.
<https://academic.oup.com/aje/article/doi/10.1093/aje/kwx317/4110407>

TMLE ROAD MAP



⁴ Luque-Fernandez, MA. 2017. TMLE steps adapted from van der Laan MJ, 2011.

G-computation RA: LAB 1

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Ways to implement the G-Formula: RA

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IPTW: LAB 2

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ESTIMATION

G-Computation and IPTW in STATA

Classical regression adjustment

```
use http://www.stata-press.com/data/r14/cattaneo2.dta
reg bweight i.mbsmoke mage
```

Source	SS	df	MS	Number of obs	=	60
				F(2, 57)	=	83.71
Model	1341718.7	2	670859.348	Prob > F	=	0.0000
Residual	456829.236	57	8014.54801	R-squared	=	0.7460
				Adj R-squared	=	0.7371
Total	1798547.93	59	30483.8633	Root MSE	=	89.524

bweight	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
mbsmoke						
smoker	-368.1796	29.8213	-12.35	0.000	-427.8957	-308.4635
mage	15.9611	3.324991	4.80	0.000	9.30292	22.61928
_cons	3191.268	73.22399	43.58	0.000	3044.64	3337.897

Stata v.14 new module for Tx effects

```
teffects ra (bweight mage) (mb smoke)
```

```
Treatment-effects estimation          Number of obs    =          60
Estimator      : regression adjustment
Outcome model  : linear
Treatment model: none
```

		Robust					
bweight		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	

ATE							
mbsmoke							
(smoker vs nonsmoker)		-375.5157	29.62863	-12.67	0.000	-433.5867	-317.4446

POmean							
mbsmoke							
nonsmoker		3598.81	24.22443	148.56	0.000	3551.331	3646.289

Computing the RA by "hand"

```
reg bweight mage if mbsmoke==1  
predict double y1hat
```

```
reg bweight mage if mbsmoke==0  
predict double y0hat
```

```
mean y1hat y0hat
```

```
lincom _b[y1hat] - _b[y0hat]
```

Mean	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
(1)	-375.5157	6.601436	-56.88	0.000	-388.7251	-362.3062

Non-parametric estimation: conditional means saturated model

```
reg bweight ibn.mbsmoke ibn.mbsmoke#c.mage, noconstant vce(robust) vsquish
```

```
//Potential outcomes
```

```
margins mbsmoke, vce(unconditional)
```

```
Predictive margins                                Number of obs      =           60  
Expression   : Linear prediction, predict()
```

		Unconditional		t	P> t	[95% Conf. Interval]	
	Margin	Std. Err.					
mbsmoke							
nonsmoker	3598.81	25.07467	143.52	0.000	3548.579	3649.04	
smoker	3223.294	21.43679	150.36	0.000	3180.351	3266.237	

Non-parametric estimation: conditional means saturated model

```
reg bweight ibn.mbsmoke ibn.mbsmoke#c.mage, noconstant vce(robust) vsquish  
//ATE  
margins r.mbsmoke, contrast(nowald)
```

Contrasts of predictive margins

Model VCE : Robust

Expression : Linear prediction, predict()

		Delta-method		
		Contrast	Std. Err.	[95% Conf. Interval]

	mbsmoke			
(smoker vs nonsmoker)		-375.5157	30.87175	-437.3592 -313.6721

Estimation of the CRR

```
teffects ra (bweight mage) (mb smoke), coeflegend
```

```
nlcom 100*_b[ATE:r1vs0.mb smoke]/_b[POmean:r0.mb smoke]
```

bweight	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
_nl_1	-10.43444	.7741387	-13.48	0.000	-11.95172	-8.917157

Estimation of the IPTW

```
. teffects ipw (bweight) (mb smoke mage, logit), nolog vsquish
```

```
Treatment-effects estimation          Number of obs    =      4,642
Estimator      : inverse-probability weights
Outcome model  : weighted mean
Treatment model: logit
```

	bweight	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATE							
	mb smoke						
(smoker vs nonsmoker)		-275.5595	22.63192	-12.18	0.000	-319.9173	-231.2018
POmean							
	mb smoke						
	nonsmoker	3408.938	9.308235	366.23	0.000	3390.695	3427.182

Estimation of the IPTW by hand

```
. clear
. use http://www.stata-press.com/data/r14/cattaneo2.dta
. global Y bweight // Outcome
. global A mbsmoke // Exposure or treatment
. global W mmarried prenatal1 fbaby medu mage mrace // Confounders.
. teffects ipw ($Y) ($A $W, logit), nolog vsquish
```

```
Treatment-effects estimation      Number of obs      =      4,642
Estimator      : inverse-probability weights
Outcome model  : weighted mean
Treatment model: logit
```

	bweight	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATE							
	mbsmoke						
(smoker vs nonsmoker)		-240.2212	24.82773	-9.68	0.000	-288.8826	-191.5597
POmean							
	mbsmoke						
nonsmoker		3408.235	9.440857	361.01	0.000	3389.732	3426.739

Estimation of the IPTW by hand

```
. logit $A $W, vce(robust) nolog
. predict double ps
. generate double ipw1 = ($A==1)/ps
. regress $Y [pw=ipw1]
```

```
Linear regression      Number of obs      =      864
```

		Robust				
bweight	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
_cons	3168.014	23.35919	135.62	0.000	3122.167	3213.862

```
. generate double ipw0 = ($A==0)/(1-ps)
. regress $Y [pw=ipw0]
```

Linear regression Number of obs = 3,778

bweight	Coef.	Robust Std. Err.	t	P> t	[95% Conf. Interval]	
cons	3408.235	9.523029	357.89	0.000	3389.565	3426.906

Estimation of the IPTW by hand: bootstrapping for SE

```
. program drop ATE
. program define ATE, rclass
.   capture drop y1
.   capture drop y0
.   regress $Y [pw=ipw1]
.   matrix y1 = e(b)
.   gen double y1 = y1[1,1]
.   regress $Y [pw=ipw0]
.   matrix y0 = e(b)
.   gen double y0 = y0[1,1]
.   mean y1 y0
.   lincom _b[y1]-_b[y0]
.   return scalar ace = `r(estimate)´
. end
```

Estimation of the IPTW by hand

```
. qui bootstrap r(ace), reps(1000): ATE  
. estat boot, all
```

```
Bootstrap results                                Number of obs    =      4,642  
                                                Replications      =      1000
```

```
command: ATE  
_bs_1: r(ace)
```

	Observed Coef.	Bias	Bootstrap Std. Err.	[95% Conf. Interval]		
_bs_1	-240.22116	.2588013	24.989583	-289.1998	-191.2425	(N)
				-287.8047	-192.4459	(P)
				-287.6467	-192.4318	(BC)

```
(N) normal confidence interval  
(P) percentile confidence interval  
(BC) bias-corrected confidence interval
```

Estimation of the IPTW by hand

```
. // Test balance after adjustment
. qui teffects ipw ($Y) ($A $W)
. tebalance summarize
```

Covariate balance summary

	Raw	Weighted
Number of obs =	4,642	4,642.0
Treated obs =	864	2,282.2
Control obs =	3,778	2,359.8

	Standardized differences		Variance ratio	
	Raw	Weighted	Raw	Weighted
mmarried	-.5953009	-.020219	1.335944	1.017057
prenatal1	-.3242695	-.0186635	1.496155	1.0278
fbaby	-.1663271	.0217106	.9430944	1.005001
medu	-.5474357	-.1156001	.7315846	.5112843
mage	-.300179	-.0740731	.8818025	.7979432
foreign	-.1706164	-.034976	.4416089	.8643349
mrace	-.1029446	-.028183	1.198452	1.052116

Survival-time treatment effects example

The time to a second heart attack among women aged 45 to 55 years. The treatment, smoking, is stored in the 0, 1 indicator smoke. These data also contain each woman's age at the time of her first heart attack (age), and indices of her exercise level (exercise), diet quality (diet), and education attainment (education) prior to her first heart attack.

Survival-time treatment effects

Does smoking decrease the time to a second heart attack in the population of women aged 45 to 55 who have had one heart attack?

use <http://www.stata-press.com/data/r14/sheart>

```
stset atime, failure(fail)
```

```
    failure event:  fail != 0 & fail < .
```

```
obs. time interval:  (0, atime]
```

```
exit on or before:  failure
```

```
2000  total observations
```

```
    0  exclusions
```

```
2000  observations remaining, representing
```

```
1208  failures in single-record/single-failure data
```

```
3795.226 total analysis time at risk and under observation
```

```
                at risk from t =                0
```

```
                earliest observed entry t =        0
```

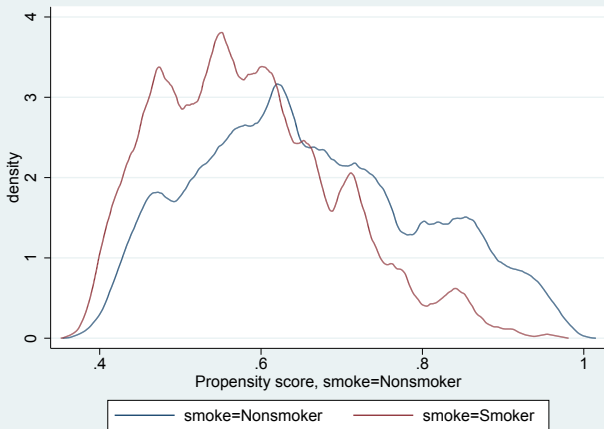
```
                last observed exit t = 34.17743
```


Survival-time treatment effects

```
*Best model for the treatment model
bf fit logit smoke age exercise diet education
display "`r(bvlist)'"
*I am using a double robust method to fit a IPW RA treatment effect model
with weighted adjustment for censoring
stteffects ipwra (age exercise diet education) (smoke age exercise diet education) \\\
(age exercise diet)
      failure _d: fail
      analysis time _t: atime
Iteration 0:   EE criterion = 1.632e-16
Iteration 1:   EE criterion = 9.694e-31
Survival treatment-effects estimation      Number of obs      =      2,000
Estimator      : IPW regression adjustment
Outcome model   : Weibull
Treatment model: logit
Censoring model: Weibull
```

	_t	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Intervall]	
-----+-----							
ATE							
	smoke						
(Smoker vs Nonsmoker)		-2.037944	.6032549	-3.38	0.001	-3.220302	-.855586
-----+-----							
POmean							
	smoke						
Nonsmoker		4.14284	.4811052	8.61	0.000	3.199891	5.085789

Overlap: positivity assumption and SUTVA



Survival-time treatment effects

*IMAI AND RATKOVIC (2014): Over-identification or Test for Balance
(Testing and checking conditional independence or unconfoundedness)
tebalance summ

Covariate balance summary

	Raw	Weighted
Number of obs =	2,000	2,000.0
Treated obs =	738	994.1
Control obs =	1,262	1,005.9

	Standardized differences		Variance ratio	
	Raw	Weighted	Raw	Weighted
age	-.3122094	-.0184574	.8547308	.9370065
exercise	-.4975269	-.0458412	.4966778	.8342339
diet	-.2479756	.0021802	.7937645	1.095347
education	-.4801442	-.0216366	.6015139	.978078

. tebalance over, nolog

Overidentification test for covariate balance

H0: Covariates are balanced:

chi2(5) = 3.28142

Prob > chi2 = 0.6567

TMLE R-markdown 2019

<https://migariane.github.io/TMLE.nb.html>

TMLE Statistics in Medicine 2018

<https://www.ncbi.nlm.nih.gov/pubmed/29687470>

SIM-2018. CODE in R and STATA

<https://github.com/migariane/SIM-TMLE-tutorial>

Practicals

G-Comp. and IPTW using R (RStudio cloud)

Practicals

G-Comp. and IPTW using R (RStudio cloud)

RStudio Cloud link

<https://rstudio.cloud/spaces/19488/project/434105>

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Thank you!



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