

Statistical methods for observational studies

Nina Masson

Supervised by Alexia Letierce and Simon Hitier

Ensaï

massonnina@yahoo.fr

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Overview

- 1 Introduction
- 2 Observational studies
- 3 Standard methods for non-randomized studies
- 4 Marginal Structural Models
- 5 Conclusion

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- Clinical trials are the gold standard to demonstrate efficacy in clinical research
- Participants receive specific interventions according to the protocol
- 4 phases
 - Phase I: Evaluate safety, tolerance threshold, side effects on healthy volunteers
 - Phase II: Test efficacy (and continue safety): comparative studies
 - Phase III: Gather more information on safety and efficacy
 - Phase IV: After approval, detect possible rare undesirable side effects
- However, limitations need to be acknowledged (less appropriate for rare conditions, restricted generalization...)
- Observational studies can complete missing information of clinical trials

Any intervention studied is determined by clinical practice and not the protocol.

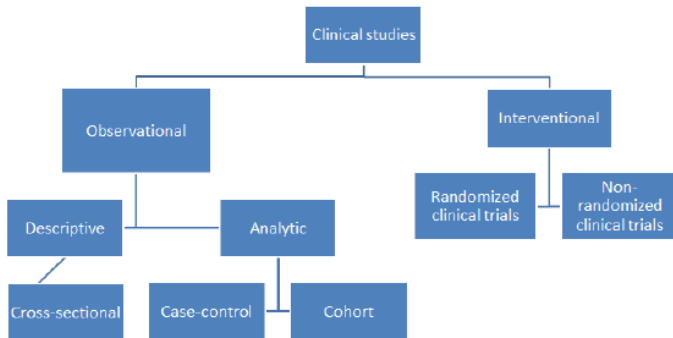
Useful to analyze

- Effectiveness: efficacy in real life
- Possible safety concerns and long-term complications
- Patients' compliance to treatment in real life
- Understudied populations
- Use of treatments (concomitant medication, off-label use,...)
- Rare conditions

Introduction

Observational studies

- Descriptive epidemiology: Monitoring the occurrence of a disease
- Analytic epidemiology: Studying the risk factors of a disease depending on an exposure



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Observational studies

The designs

Cross-sectional studies: Exposure and outcome determined simultaneously



Cohort studies: Inclusion of exposed and unexposed and analysis of whether or not they develop the outcome



Case-control studies: Inclusion of subjects with and without the outcome and collection of information on the past exposure



Observational studies

Potential biases

Selection bias:

- Distortion in estimating the association between the exposure and the outcome
- Consequence of how subjects are selected in the study

Example: Case-control study: influence of NSAIDs on colon cancer
Selection bias if the control chosen are patients hospitalized for arthritis

⇒ Both cases and controls will be exposed to NSAIDs

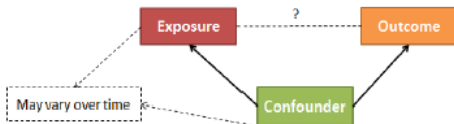
Information bias:

- Distortion in estimating which is the consequence of measurement errors or bad classifications of patients

Observational studies

Confounding

Confounders: factors associated with the exposure and with the outcome studied



Example:



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Adjusting for confounding with multivariate regression

- Include treatment and confounder in the model

Stratification with the Cochran-Mantel-Haenszel method (cmh)

- Comparing crude and stratum-specific associations

$$H_0 : OR_1 = OR_2 = \dots = OR_K = 1$$

Matching

- Controlling confounding by forming homogeneous pairs regarding the confounder
- Statistical tests for matched data (ex: paired t-test for a continuous outcome)

Standard approaches

Propensity score methods

Definition

Conditional probability for a patient to receive the treatment given his/her observed covariates

$$e_i = P(A_i = a_i | L_i)$$

where

- e_i : propensity score for subject i
- A_i : treatment of subject i
- L_i : covariate for subject i



→ Estimate e_i with logistic regression

Standard approaches

Propensity score methods

Methods using the propensity score:

- **Adjustment:** propensity score \hat{e}_i is included in the regression model with the treatment A

$$E[Y_i] = \beta_0 + \beta_1 A_i + \beta_2 \hat{e}_i$$

- **Stratification:** propensity score is cut in percentiles and the association is observed in each stratum
- **Matching:** a treated patient is matched with an untreated patient with the closest propensity score
- Inverse Probability of Treatment Weighted (IPTW)

Inverse Probability of Treatment Weighted (IPTW)

- Calculate a weight from the propensity score for each subject:

$$IPTW_i = \frac{1}{P(A_i = a_i | L_i = l_i)} = \frac{1}{\hat{e}_i}$$

For a binary treatment: $IPTW_i = \frac{A_i}{\hat{e}_i} + \frac{1-A_i}{1-\hat{e}_i}$ (demo)

→ In the weighted sample (pseudo-population): no confounding \Rightarrow a usual regression model can be applied (demo)

- Stabilized weights to increase statistical efficiency

$$sw_i = \frac{P(A_i = a_i)}{P(A_i = a_i | L_i = l_i)} = P(A_i = a_i) \times IPTW_i$$

Methodology

Comparisons

Methods	Strengths	Weaknesses
Multivariate adjustment	<ul style="list-style-type: none">- Simultaneously adjustment for multiple confounders- Use of all information in continuous variables	<ul style="list-style-type: none">- Adequacy and assumptions of the model
Stratification	<ul style="list-style-type: none">- Simple- Estimation of the effects by stratum	<ul style="list-style-type: none">- Difficult to interpret if multiple confounders with multiple levels- Can cause loss of information or not remove enough confounding according to how the stratification is done
Matching	<ul style="list-style-type: none">- Elimination of the influence of strong constitutional confounders- Elimination of the influence of confounders that are hard to measure	<ul style="list-style-type: none">- Difficult to acquire an adequate sample size- Overmatching
Propensity score	<ul style="list-style-type: none">- Simultaneously control for multiple confounders- Ability to directly see confounding through distribution of the propensity score	<ul style="list-style-type: none">- Difficult to understand the results- Measurement of all relevant covariates

Standard approaches

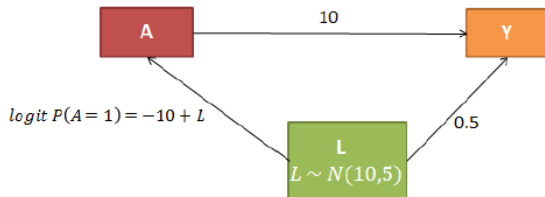
Applications: simulated data

- Point-treatment study

- $L \sim \mathcal{N}(10, 5)$: a continuous confounder
- $\text{logit } P(A = 1) = -10 + L$: a dichotomous treatment
- $Y = 10A + 0.5L + \mathcal{N}(-10, 5)$: a continuous outcome
- $i = 1, \dots, n$ subjects with $n = 1000$
- 200 simulations

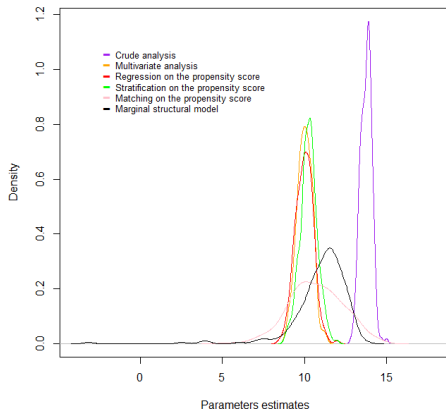
- Objective: estimate the causal effect of A on Y

- True value: 10



Standard approaches

Applications: simulated data



- Crude analysis overestimates the effect
- Adjusting on the confounder, adjusting on the propensity score, stratification on the propensity score: similar results with values centered around 10
- Matching on the propensity score and IPTW: more scattered estimates (Truncating method for the weights)

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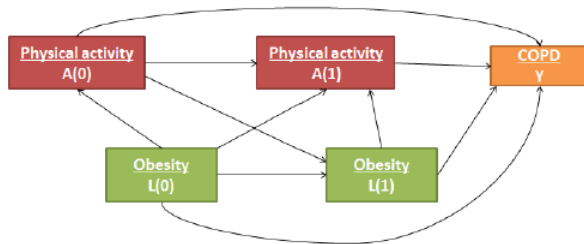
Marginal Structural models

Introduction

■ Literature search:

- "Marginal structural models and causal inference in epidemiology", Robins JM, Hernan MA, Brumback B (2000), *Epidemiology*
- "Marginal structural models to estimate the joint causal effect of nonrandomized treatments", Hernan MA, Brumback B, Robins JM (2001), *Journal of American Statistical Association*

■ Marginal structural models (MSMs) are useful when:



=> Relationship between treatment and outcome can be confounded

Marginal Structural Models

Notations

Notations

- $A_i(t)$: treatment at time t for subject i
- $\bar{A}_i(t) = (A_i(0), \dots, A_i(t))$ for subject i , treatment history until t
- $L_i(t)$: confounder value at time t for subject i
- $\bar{L}_i(t) = (L_i(0), \dots, L_i(t))$ for subject i , covariate history until t
- V_i : subset of time-fixed covariates ($V_i = L_i(0)$)
- $C_i(t)$: censoring indicator for subject i at time t
- K : study duration
- Outcome: Y_i for binary or continuous variables and T_i for survival analysis

Potential outcome

- $Y_{\bar{a}}$: a subject's binary or continuous outcome had he/she been treated with \bar{a} rather than his/her observed treatment \bar{A}
- $T_{\bar{a}}$: a subject's time-to-event if he/she had followed treatment history \bar{a} from the start of follow-up rather than his/her observed treatment history

Marginal Structural Models

Weights

Marginal Structural Models

Weights

- Computing the Inverse Probability of Treatment Weights **IPTW**
 $t = 1, \dots, K$ (implementation)

$$sw_i(t) = \prod_{k=0}^t \frac{f(A_i(k) | \bar{A}_i(k-1), V_i)}{f(A_i(k) | \bar{A}_i(k-1), \bar{L}_i(k))}$$

Marginal Structural Models

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- Computing the Inverse Probability of Censoring Weights **IPCW**

$$sw_i^\dagger(t) = \prod_{k=0}^t \frac{P[C_i(k) = 0 | \bar{C}_i(k-1) = 0, \bar{A}_i(k), V_i]}{P[C_i(k) = 0 | \bar{C}_i(k-1) = 0, \bar{A}_i(k), \bar{L}_i(k)]}$$

Marginal Structural Models

Weights

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- Computing the final weights

$$sw_i(t) \times sw_i^\dagger(t)$$

Marginal Structural Models

Weights

- Computing the Inverse Probability of Treatment Weights **IPTW**
 $t = 1, \dots, K$ (implementation)

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- Computing the final weights

$$sw_i(t) \times sw_i^\dagger(t)$$

\Rightarrow 1 weight per person and per time (sw)

Marginal Structural Models

Modeling

- **Binary outcome:** $\boxed{\text{logit } P(Y_{\bar{a}} = 1) = \beta_0 + \beta_1 \text{cum}(\bar{a})}$ ¹
measured at end of follow-up

$$sw_i(K) \times sw_i^\dagger(K)$$

- **Continuous outcome:** $\boxed{E[Y_{\bar{a}}] = \beta_0 + \beta_1 \text{cum}[\bar{a}]}$
→ Fit a standard GEE linear model giving the time-specific weight $sw_i(t) \times sw_i^\dagger(t)$ to each subject

- **Survival:** $\boxed{\lambda_{T_{\bar{a}}}(t) = \lambda_0(t) \exp(\gamma_1 a(t) + \gamma_2 V)}$
⇒ Weighted pooled logistic regression treating each person-time as an observation
⇒ Use weights $sw_i(t) \times sw_i^\dagger(t)$ for
 $\text{logit } P[D(t) = 1 | D(t-1) = 0, \bar{A}(t-1)] = \theta_0(t) + \theta_1 A(t-1) + \theta_2 V$

¹ $\text{cum}(\bar{a}) = \sum_{k=0}^K a_k$

Marginal Structural Model

Assumptions

■ Exchangeability:

- No unmeasured confounding assumption
- Measure enough joint predictors of exposure and outcome
- Sensitivity analyses (asses the impact of adding further potential confounders)

■ Consistency:

- Links the counterfactual data $Y_{\bar{a}}$ to the observed data (Y, A)
- A subject's counterfactual outcome under his or her observed exposure history is his or her observed outcome

$$Y_{\bar{A}} = Y$$

■ Positivity:

- The experimental treatment assumption
- Both treated and untreated patients at every level of the confounder

■ Specifications of the models:

- Models for initiation of treatment and censoring, given past covariate and treatment history need to be correctly specified

Marginal Structural Models

Applications: real-life data

The CREDIT study: a long-term international non-interventional study in patients with type 2 diabetes, treated with insulin (Initial protocol)

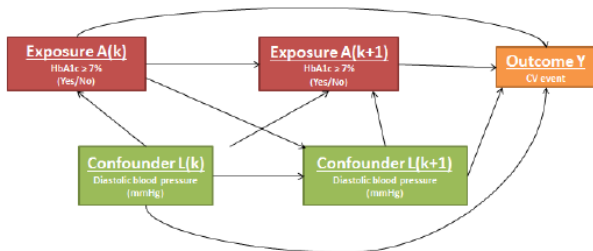
Clinical context:

- Diabetes:
 - Chronic disease that can be controlled but for which there is no cure
 - Triggered by a shortage or a deficiency of insulin (hormone normally produced by the pancreas to help control blood glucose level)
- When glucose lacks of insulin
 - ⇒ glucose does not give energy to cells and is not absorbed by them
 - ⇒ its accumulation causes damages to the organs (retina, kidney,...)

Marginal Structural Models

Application: real-life data

A binary analysis to study the relationship between CV events and glycemic control, in presence of a time-dependent confounder, the diastolic blood pressure



- 2,524 patients followed up to 4 visits
- Dataset:
 - one row of data per patient for each visit until their time to CV event
 - if no CV event: one row of data per visit until censoring

Marginal Structural Models

Application: real-life data

	HbA1c < 7% at baseline N=143	HbA1c ≥ 7% at baseline N=2,381	Total N=2,524
CV event <u>N (%)</u>	9 (6.3%)	106 (4.5%)	115 (4.6%)
Age (years) mean (SD)	63.44 (10.01)	61.30 (10.04)	61.42 (10.05)
Sex Men	78 (54.5%)	1,172 (49.2%)	1,250 (49.5%)
Women	65 (45.5%)	1,209 (50.8%)	1,274 (50.5%)
Diastolic blood pressure mean (SD)	79.81 (10.73)	80.97 (11.53)	80.90 (11.49)

- Only 4.6% of the patients enrolled present a CV event during follow-up
- Diastolic blood pressure: time-dependent confounder
 - Relationship between diastolic blood pressure and cv event
 - Relationship between diastolic blood pressure and glycemic control

Marginal Structural Models

Application: real-life data

Implementation:

Marginal Structural Models

Application: real-life data

Implementation:

- 1 Computing IPTW

Marginal Structural Models

Application: real-life data

Implementation:

1 Computing IPTW

- Numerator: PROC LOGISTIC

- outcome: exposure $A_i(k)$
- covariates: previous exposure $A_i(k - 1)$ and baseline variables

Marginal Structural Models

Application: real-life data

Implementation:

1 Computing IPTW

- Numerator: PROC LOGISTIC
 - outcome: exposure $A_i(k)$
 - covariates: previous exposure $A_i(k-1)$ and baseline variables
- Denominator: PROC LOGISTIC
 - same method, adding the confounder history as explanatory variable

Marginal Structural Models

Application: real-life data

Implementation:

1 Computing IPTW

- Numerator: PROC LOGISTIC

- outcome: exposure $A_i(k)$

- covariates: previous exposure $A_i(k-1)$ and baseline variables

- Denominator: PROC LOGISTIC

- same method, adding the confounder history as explanatory variable

2 Computing IPCW

Marginal Structural Models

Application: real-life data

Implementation:

1 Computing IPTW

- Numerator: PROC LOGISTIC

- outcome: exposure $A_i(k)$

- covariates: previous exposure $A_i(k-1)$ and baseline variables

- Denominator: PROC LOGISTIC

- same method, adding the confounder history as explanatory variable

2 Computing IPCW

- same method but outcome: $C_i(k)$

Marginal Structural Models

Application: real-life data

Implementation:

1 Computing IPTW

- Numerator: PROC LOGISTIC

- outcome: exposure $A_i(k)$

- covariates: previous exposure $A_i(k-1)$ and baseline variables

- Denominator: PROC LOGISTIC

- same method, adding the confounder history as explanatory variable

2 Computing IPCW

- same method but outcome: $C_i(k)$

3 Computing the final stabilized weights

Marginal Structural Models

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- Denominator: PROC LOGISTIC

- same method, adding the confounder history as explanatory variable

2 Computing IPCW

- same method but outcome: $C_i(k)$

3 Computing the final stabilized weights

- One weight per patient and per visit

- Binary analysis \Rightarrow only the weight at last visit for each patient is retrieved (formula)

Marginal Structural Models

Application: real-life data

Implementation:

1 Computing IPTW

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4 Fitting final marginal structural model: PROC LOGISTIC with the

weight option: $\text{logit } P(Y_{\bar{a}} = 1) = \beta_0 + \beta_1 \sum_{k=0}^K a_k$

- $\sum_{k=0}^K a_k$: duration of exposure to an HbA1c value $\geq 7\%$

Marginal Structural Models

Application: real-life-data

Parameter		Estimate	Standard Error	Pr > ChiSq
Intercept		-3.7579	1.0118	0.0002
Visit	2	-0.6392	0.2826	0.0237
Visit	3	-1.2237	0.3384	0.0003
Visit	4	-4.5909	0.4643	<.0001
HbA1c		0.2838	0.1233	0.0214
Sex	M	1.5312	1.3011	0.2392
Age		0.0387	0.0151	0.0102
Age*Sex	M	-0.0241	0.0199	0.2264

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- $OR = 1.32 [1.01-1.69]$

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- $OR = 1.32$ [1.01-1.69]
- Significant difference between the exposed and the unexposed

Marginal Structural Models

Application: real-life data

Comparison with an unweighted model:

Parameter		Estimate	Standard Error	95% Confidence Limits		Pr > ChiSq
Intercept		-9.7361	1.3187	-12.3207	-7.1516	<.0001
Visit	2	-0.2105	0.2374	-0.6758	0.2549	0.3754
Visit	3	-0.2183	0.2442	-0.6970	0.2604	0.3714
Visit	4	-1.0216	0.3303	-1.6690	-0.3742	0.0020
Sex	M	2.2480	1.3572	-0.4120	4.9080	0.0976
Age		0.0579	0.0155	0.0276	0.0883	0.0002
Age*Sex	M	-0.0289	0.0207	-0.0694	0.0115	0.1611
HbA1c	≥ 7%	0.3626	0.2317	-0.0915	0.8168	0.1176
DBP		0.0193	0.0089	0.0019	0.0367	0.0293

Marginal Structural Models

Application: real-life data

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Visit	2	-0.2105	0.2374	-0.6758	0.2549	0.3754
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Visit	4	-1.0216	0.3303	-1.6690	-0.3742	0.0020
Sex	M	2.2480	1.3572	-0.4120	4.9080	0.0976
Age		0.0579	0.0155	0.0276	0.0883	0.0002
Age*Sex	M	-0.0289	0.0207	-0.0694	0.0115	0.1611
HbA1c	≥ 7%	0.3626	0.2317	-0.0915	0.8168	0.1176
DBP		0.0193	0.0089	0.0019	0.0367	0.0293

Marginal Structural Models

Discussion and perspectives

- Only a few patients with CV event

Marginal Structural Models

Discussion and perspectives

- Only a few patients with CV event
- Relevant time-dependent confounder

Marginal Structural Models

Discussion and perspectives

- Only a few patients with CV event
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- Assumptions:
 - Exchangeability \Rightarrow sensitivity analysis
 - Positivity
 - Specification of the models

Marginal Structural Models

Discussion and perspectives

- Only a few patients with CV event
- Relevant time-dependent confounder
- Assumptions:
 - Exchangeability \Rightarrow sensitivity analysis
 - Positivity
 - Specification of the models
- Other application: marginal structural Cox model vs unweighted method \Rightarrow Similar results

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Conclusion

- Objective: Statistical methods for observational studies

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- Application on real-life data:
 - Results going in the same direction but,
 - Non-significant results for the unweighted method

Conclusion

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- Looking through the literature, the propensity score methods rapidly directed us towards the MSM with IPTW estimator
- MSMs enable time-dependent confounders to be taken into account
- Point treatment study: difference between results obtained
- Application on real-life data:
 - Results going in the same direction but,
 - Non-significant results for the unweighted method
- In literature: important difference between marginal structural models and unweighted methods

THANK YOU FOR YOUR ATTENTION !

How to use Cochran-Mantel-Haenszel Test and Breslow-Day Test

- Calculate crude OR_{Y-A}
- Calculate stratum-specific OR's: $OR_{Y-A|L}$
- If $OR_{Y-A} = OR_k \Rightarrow L$ is unlikely to be a confounder
- If $OR_{Y-A} \neq$:
 - If $OR_1 = OR_2 = \dots = OR_K \Rightarrow L$ may be a confounder
 - Apply CMH test of conditional independence: H_0 : A and Y are conditionnally independent

$$CMH = \frac{(\sum_k n_{11k} - \mu_{11k})^2}{\sum_k \sigma_{11k}^2}$$

- Calculate OR_{MH} (if H_0 is rejected, OR_{MH} should be $\neq 1$)
- If stratum-specific Or's differ from each oether \Rightarrow there might be an effect modification (interaction)
 - Apply Breslow-Day test of homogeneity of the OR's: H_0 : stratum-specific OR's are equal (homogeneous)

$$Q_{BD} = \sum_k \sum_i \sum_j \frac{(n_{ijk} - m_{ijk})^2}{m_{ijk}} \sim K\chi^2(K-1)$$

Stratification

	$L = k$		
	$Y = 1$	$Y = 0$	
$A = 1$	n_{11k}	n_{10k}	n_{1+k}
$A = 0$	n_{01k}	n_{00k}	n_{0+k}
	n_{+1k}	n_{+0k}	n_k

- CMH test statistic: $M^2 = \frac{[\sum_k (n_{11k} - \mu_{11k})]^2}{\sum_k \text{Var}(n_{11k})} \sim \chi^2(1)$, where
 - $\mu_{11k} = \frac{n_{1+k}n_{+1k}}{n_k}$ is the expected frequency of $A = 1$ and $Y = 1$ for $L = k$ assuming the conditional independence holds
 - $\text{Var}(n_{11k}) = \frac{n_{1+k}n_{0+k}n_{+1k}n_{+0k}}{n_k^2(n_k - 1)}$
- Common odds ratio:

$$\widehat{OR}_{MH} = \frac{\sum_k (n_{11k}n_{00k})/n_k}{\sum_k (n_{10k}n_{01k})/n_k}$$

Paired t-test

- Comparing 2 measures of a quantitative variable realized on the paired subjects
- Treating the 2 matched samples as one dataset on which the difference between both measures would have been computed

■ H_0 : *the mean difference between both measures is null*

■ Test statistic: $t = \frac{\frac{1}{n} \sum_{i=1}^n \text{Diff}_i}{\sqrt{\frac{SCE_d}{n(n-1)}}} \sim St(n-1)$

(where $SCE_d = \sum_{i=1}^n \text{Diff}_i^2 - \frac{(\sum_{i=1}^n \text{Diff}_i)^2}{n}$)

H_0 rejected \Rightarrow significant difference between both outcomes

back

Computation of the unstabilized weights

Proof in a point treatment study

Actual population: n_{ijk} is the number of patients with the outcome i , under treatment j and with value covariate k in the actual population

	$L = 1$		$L = 0$		Total
	$A = 1$	$A = 0$	$A = 1$	$A = 0$	
$Y = 1$	n_{111}	n_{101}	n_{110}	n_{100}	n_{1++}
$Y = 0$	n_{011}	n_{001}	n_{010}	n_{000}	n_{0++}
Total	n_{+11}	n_{+01}	n_{+10}	n_{+00}	n_{+++}

Pseudo-population: N_{ijk} is the number of patients with the outcome i , under treatment j and with value covariate k in the pseudo-population

	$L = 1$		$L = 0$		Total
	$A = 1$	$A = 0$	$A = 1$	$A = 0$	
$Y = 1$	N_{111}	N_{101}	N_{110}	N_{100}	N_{1++}
$Y = 0$	N_{011}	N_{001}	N_{010}	N_{000}	N_{0++}
Total	N_{+11}	N_{+01}	N_{+10}	N_{+00}	N_{+++}

Weights:

For patients with $L = 1$: When n_{+11} patients are treated, n_{111} have $Y = 1 \Rightarrow$ if all patients with $L = 1$ (n_{++1}) would have been treated, we would have:

$$N_{111} = \frac{n_{++1} \times n_{111}}{n_{+11}} = w_{11} \times n_{111}, \text{ where } w_{11} = \frac{n_{++1}}{n_{+11}} = \frac{1}{P(A=1|L=1)}$$

back

Association between A and L

	$L = 1$	$L = 0$	
$A = 1$	N_{+11}	N_{+10}	N_{+1+}
$A = 0$	N_{+01}	N_{+00}	N_{+0+}
	N_{++1}	N_{++0}	N_{+++}

Objective: show that the proportion of treated who have $L = 1$ is the same as the proportion of untreated who have $L = 1$, i.e. $\frac{N_{+11}}{N_{+1+}} = \frac{N_{+01}}{N_{+0+}}$

$$\frac{N_{+11}}{N_{+1+}} = \frac{w_{11}n_{+11}}{w_{11}n_{+11} + w_{10}n_{+10}} = \frac{\frac{n_{++1}}{n_{+11}} \times n_{+11}}{\frac{n_{++1}}{n_{+11}} \times n_{+11} + \frac{n_{++0}}{n_{+10}} \times n_{+10}} = \frac{n_{++1}}{n_{+++}}$$

$$\frac{N_{+01}}{N_{+0+}} = \frac{w_{01}n_{+01}}{w_{01}n_{+01} + w_{00}n_{+00}} = \frac{\frac{n_{++1}}{n_{+01}} \times n_{+01}}{\frac{n_{++1}}{n_{+01}} \times n_{+01} + \frac{n_{++0}}{n_{+00}} \times n_{+00}} = \frac{n_{++1}}{n_{+++}}$$

Mathematical dimension

- A process is "causally exogenous or ancillary" if

$$Y_{\bar{a}} \perp\!\!\!\perp A(t) | \bar{A}(t-1)$$

$\Rightarrow Y_{\bar{a}}$ is independent of \bar{A}

- A treatment is a "statistically exogenous or ancillary" process if

$$\bar{L}(t) \perp\!\!\!\perp A(t) | \bar{A}(t-1)$$

- $sw_i(t)$ quantifies the degree to which the treatment process is statistically nonexogenous through time t
 - \Rightarrow numerator = denominator for all t with probability 1 if and only if treatment process is statistically exogenous
 - \Rightarrow weighted regression = unweighted analysis only if $A(t)$ is statistically exogenous

Truncation method for the IPTW method

In the simulated study, some weights computed for the MSM are extreme:

- Summary of the weights for the 118th simulated dataset:

Min	1 st qu.	Median	Mean	3 rd qu.	Max
0.46	0.49	0.54	5.59	0.62	4649

⇒ Parameter estimate: -3.15 (true value: 10)

- Summary of the weights for the 1st simulated dataset

Min	1 st qu.	Median	Mean	3 rd qu.	Max
0.48	0.51	0.53	0.91	0.63	74.7

⇒ Parameter estimate: 10.65

(back)

Initial study protocol of the CREDIT study

- Multicenter international non-interventional longitudinal study with a 4-year follow-up per patient carried out on 3,060 subjects
- Visits done according to clinical practice
- Original objectives:
 - Observation of medical practice in the real life, over a 4-year period, in people with type 2 diabetes treated with insulin
 - Evaluation of the evolution and relationship between glycemic control and the risk of cardiovascular events in type 2 diabetic patients treated with insulin, taking into account known cardiovascular risk factors

Initial study protocol of the CREDIT study

Inclusion criteria	Exclusion criteria
<ul style="list-style-type: none">- Male or female, age > 40 years- With type 2 diabetes- Time from diagnosis of diabetes to insulin initiation > 1 year- Treated with insulin (all regimens*) for more than 1 month and less than 6 months prior to study entry- With an HbA1c value within 3 months prior to insulin initiation- Insulin initiated with an intention of a long term treatment- Informed consent must be obtained in writing- Patients able to be followed over a long period of time	<ul style="list-style-type: none">- Type 1 diabetes- Not insulin treated type 2 diabetes- Secondary diabetes (pancreatic history, steroids therapy, endocrine diabetes)- Current temporary insulin therapy (gestational diabetes, pancreas cancer, surgery, clinical trial)- Pregnancy at inclusion

(back)

Marginal Structural Model

Inverse Probability of Treatment Weights (IPTW)

- Binary treatment
- For the denominator of $sw_i(k)$:

$$\text{logit } P(A(k) = 0 | \bar{A}(k-1) = 0, \bar{L}(k)) = \alpha_0(k) + \alpha_1 L(k) + \alpha_2 V$$

$$\begin{cases} \prod_{u=0}^k \hat{p}_i(u) & \text{if subject } i \text{ did not start treatment up to time } k \\ (1 - \hat{p}_i(t)) \prod_{u=0}^{t-1} \hat{p}_i(u) & \text{if subject } i \text{ started treatment at time } t \text{ for } 0 < t \leq k \\ 1 - \hat{p}_i(0) & \text{if subject } i \text{ is treated at time } 0 \end{cases}$$

- For the numerator of $sw_i(k)$: remove $L(k)$ from the logistic model

(back)

Marginal Structural Models

Application: real-life data

A time-to-event analysis

⇒ Application of a marginal structural Cox model

Marginal Structural Models

Application: real-life data

A time-to-event analysis

⇒ Application of a marginal structural Cox model

- 1 Implementing final weights $sw_i(t) \times sw_i^\dagger(t)$: same method as previously

Marginal Structural Models

Application: real-life data

A time-to-event analysis

⇒ Application of a marginal structural Cox model

- 1 Implementing final weights $sw_i(t) \times sw_i^\dagger(t)$: same method as previously
- 2 Fitting final marginal structural Cox model

Marginal Structural Models

Application: real-life data

A time-to-event analysis

⇒ Application of a marginal structural Cox model

- 1 Implementing final weights $sw_i(t) \times sw_i^\dagger(t)$: same method as previously
- 2 Fitting final marginal structural Cox model
 - PROC GENMOD with the *scwgt* option
 - Unstructured correlation matrix
 - *repeated* option

(back)

Applications and results

Real-life data

Parameter		Estimate	Standard Error	95% Confidence Limits		Pr > Z
Intercept		-7.7774	1.0893	-9.9123	-5.6424	<.0001
HbA1c	≥ 7%	0.2180	0.2726	-0.3162	0.7522	0.4238
Visit	2	-0.1487	0.2413	-0.6217	0.3243	0.5378
Visit	3	-0.1768	0.2554	-0.6773	0.3238	0.4888
Visit	4	-0.9356	0.3446	-1.6110	-0.2603	0.0066
Sex	M	2.2918	1.3606	-0.3750	4.9586	0.0921
Age		0.0527	0.0157	0.0219	0.0834	0.0008
Age*Sex	M	-0.0301	0.0207	-0.0706	0.0104	0.1457

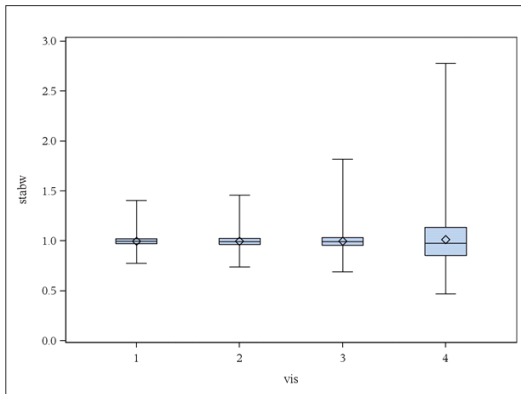
Parameter		Estimate	Standard Error	Pr > ChiSq	Hazard Ratio
Age		0.0414	0.0097	< 0.0001	1.042
Sex	M	0.3641	0.1891	0.0541	1.439
HbA1c	≥ 7%	0.3563	0.2309	0.1228	1.428
DBP		0.0195	0.0089	0.0292	1.020

Table: Unweighted method

- Similar results (weights boxplot)

(back)

Distribution of the final weights



back