On martingales, causality, identifiability and model selection

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PhD defense, 14 February, 2014

Agenda

- A motivating problem
- 2 Exponential martingales
- Simplified proofs in the general theory of processes
- A notion of causality for SDEs
- Identifiability in ICA
- 6 Degrees of freedom in nonlinear regression

Consider the following situation:

- A patient has a disease
- The disease may lead to an event (such as death)
- A treatment can be given, depending on health condition
- The patient may at any time be censored (e.g., recover)

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We wish to model this scenario and estimate the causal effect of the treatment on the time to the event.

We may use counting processes to set up a modeling framework (Røysland 2010). Assume:

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- N^L is a multivariate counting process

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$$T_A = \inf\{t \ge 0 \mid N_t^A = 1\}$$
, similarly for T_C and T_D

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$$A_t = \int_0^t \mathbf{1}_{(s \le T_A)} dN_s^A$$
, similarly for C and D

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Here:

- A is the counting process for initiation of treatment
- C is the counting process for censoring
- *D* is the counting process for the event
- L measures the patients multivariate health condition

Illustration of the model setup:



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We put restrictions on the dependencies, corresponding to the local independence graph:



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Røysland shows that this facilitates estimation of the causal effect of A on D in the marginal model where L is unobserved.

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- The estimation methodology outlined above requires the existence of randomized trial measures. What are sufficient criteria ensuring this existence?
- Our modeling discussion involved notions of causality. How do we formalize such notions in a continuous-time framework?

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From a more abstract perspective, our problem is thus:

Problem I. Assume given processes λ and μ , and a multidimensional counting process N with intensity λ . When is it possible to construct from this a multidimensional counting process with intensity μ ?

To give a solution to this problem, we assume given:

- A filtered probability space $(\Omega, \mathcal{F}, (\mathcal{F}_t), P)$
- Predictable *d*-dimensional processes λ and μ on the probability space
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And we define:

• $M_t^i = N_t^i - \int_0^t \lambda_s^i \, \mathrm{d}s$ • $\gamma_t^i = \mu_t^i / \lambda_t^i$ • $H_t^i = \gamma_t^i - 1$

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All integrals are vector integrals, meaning that

$$(H \cdot M)_t = \sum_{i=1}^d \int_0^t H_s^i \mathrm{d} M_s^i.$$

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The following lemma shows that the martingale property of $\mathcal{E}(H \cdot M)$ can be used to construct a counting process with intensity μ by a change of measure.

Lemma. Assume that $\mathcal{E}(H \cdot M)$ is a martingale. Let $t \ge 0$ and let Q_t have Radon-Nikodym derivative $\mathcal{E}(H \cdot M)_t$. Then N has intensity μ on [0, t] under Q_t .

Thus, by the lemma, it suffices to consider:

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We obtained the following sufficient criterion:

Theorem. Assume that there is $\varepsilon > 0$ such that for $0 \le u \le t$ with $t - u \le \varepsilon$, it holds that one of the following two conditions are satisfied:

$$\begin{split} &E\exp\left(\sum_{i=1}^d\int_u^t(\gamma_s^i\log\gamma_s^i-(\gamma_s^i-1))\lambda_s^i\,\mathrm{d}s\right)<\infty\quad\text{or}\\ &E\exp\left(\sum_{i=1}^d\int_u^t\lambda_s^i\,\mathrm{d}s+\int_u^t\log_+\gamma_s^i\,\mathrm{d}N_s^i\right)<\infty. \end{split}$$

Then $\mathcal{E}(H \cdot M)$ is a martingale.

For the case of a homogeneous Poisson process, meaning that $\lambda = 1$, we get the following corollary.

Corollary. Assume that there is $\varepsilon > 0$ such that for $0 \le u \le t$ with $t - u \le \varepsilon$, it holds that one of the following two conditions are satisfied:

$$\begin{split} & E \exp\left(\sum_{i=1}^{d} \int_{u}^{t} \mu_{s}^{i} \log_{+} \mu_{s}^{i} \, \mathrm{d}s\right) < \infty \quad \text{or} \\ & E \exp\left(\sum_{i=1}^{d} \int_{u}^{t} \log_{+} \mu_{s}^{i} \, \mathrm{d}N_{s}^{i}\right) < \infty. \end{split}$$

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This yields criteria for the existence of:

• Randomized trial measures

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- Counting processes with intensity given as transformations of SDEs
- Some self-exciting counting processes

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Theorem (Protter & Shimbo, 2008). If $E \exp(\frac{1}{2} \langle M^c \rangle_{\infty} + \langle M^d \rangle_{\infty})$ is finite, then $\mathcal{E}(M)$ is a uniformly integrable martingale.

We showed the following. For a>-1 with $a\neq 0$, define

$$\alpha(a) = \frac{(1+a)\log(1+a) - a}{a^2}$$
$$\beta(a) = \frac{(1+a)\log(1+a) - a}{(1+a)a^2},$$

and extend to $[-1,\infty)$ by continuity.

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Theorem. Let $a \ge -1$ and assume that $\Delta M \mathbb{1}_{(\Delta M \neq 0)} \ge a$. It holds that

$$E \exp(rac{1}{2} \langle M^c \rangle_{\infty} + lpha(a) \langle M^d \rangle_{\infty}) < \infty \Rightarrow \mathcal{E}(M)$$
 is an UI MG
 $E \exp(rac{1}{2} [M^c]_{\infty} + eta(a) [M^d]_{\infty}) < \infty \Rightarrow \mathcal{E}(M)$ is an UI MG.

All constants are optimal. Note that $\beta(-1) = \infty$.

Plot of α and β :



x value

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Corollary. Assume that $\Delta M \ge 0$. If $E \exp(\frac{1}{2} \langle M \rangle_{\infty})$ or $E \exp(\frac{1}{2} [M]_{\infty})$ is finite, then $\mathcal{E}(M)$ is a uniformly integrable martingale.

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Based on this observation, we proved, extending a result by Krylov (2009):

Theorem. Assume that $\Delta M \ge 0$. Fix $0 \le \gamma \le 1$. Assume that

$$\liminf_{\varepsilon \to 0} \varepsilon \log E \exp\left((1-\varepsilon)\frac{1}{2}\left(\gamma[M]_{\infty} + (1-\gamma)\langle M \rangle_{\infty}\right)\right) < \infty,$$

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Beiglböck et al (2012) gave a simplified proof of the Doob-Meyer theorem using the following lemma:

Lemma. Let (X_n) be a uniformly integrable sequence of variables. Then, there exist a limit variable X and convex weights such that

$$\sum_{i=n}^{K_i} \lambda_i^n X_i \xrightarrow{\mathcal{L}^1} X.$$

Simplified proofs in the general theory of processes Applying an \mathcal{L}^2 version of the lemma, we gave simple proofs of:

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- The existence of the quadratic variation process: For a local martingale *M*, there exists an adapted and increasing process [*M*] such that *M*² [*M*] is a local martingale and (Δ*M*)² = Δ[*M*].

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The first proof is a minor variation of the arguments presented in Beiglböck et al (2012).

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Write $M_t^2 = N_t^n + Q_t^n$, where $t_k^n = k2^{-n}$ and

$$N_t^n = 2 \sum_{k:t_{k-1}^n < t} M_{t_{k-1}^n}^t (M_{t_k^n}^t - M_{t_{k-1}^n}^t).$$

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Observe that (N_t^n) is a sequence of local martingales with (N_{∞}^n) bounded in \mathcal{L}^2 . Thus, by the lemma, there exists a limiting martingale N.

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Put $[M] = M^2 - N$ and verify using strong convergence.

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Instead of considering counting processes, where causality already has been discussed through local independence (e.g. Didelez 2008), we considered causality for stochastic differential equations (SDEs) of the type

$$\mathrm{d}X_t = a(X_{t-})\,\mathrm{d}Z_t,$$

where $a : \mathbb{R}^p \to \mathbb{M}(p, d)$ and Z is a d-dimensional semimartingale.

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We begin by considering an example.

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What happens when we intervene by changing the activity level of some gene?

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Increasing the activity of gene D: growth effect.

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Increasing the activity of gene E: no effect. (in spite of positive correlation with growth)

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Increasing the activity of gene C: growth effect. (through gene D)

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Increasing the activity of gene A: large growth effect. (through genes B and D) $% \left({{{\mathbf{D}}_{\mathbf{n}}}_{\mathbf{n}}} \right)$

In reality, expression levels of genes change over time. Consider a network of p genes. One simple model of the activity of these genes over time is through an Ornstein-Uhlenbeck SDE of the type

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If we could endow this model with a notion of causality, we might be able to use it to **predict the effect of interventions in the system.** This is our motivation for introducing a notion of causality for SDEs.

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We consider the SDE

$$\mathrm{d}X_t = a(X_{t-})\,\mathrm{d}Z_t.$$

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In particular, we may use the formula

$$\mathrm{d}X_t = a(X_{t-})\,\mathrm{d}Z_t$$

to make the approximation

$$X_{t+\Delta} = X_t + a(X_t)(Z_{t+\Delta} - Z_t)$$

and consider this as describing how interventions in X_t will influence $X_{t+\Delta}$.

Definition. Consider some $m \le p$ and $\zeta \in \mathbb{R}$. The postintervention SDE arising from making the intervention $X^m := \zeta$ in the *p*-dimensional SDE

$$\mathrm{d}X_t = a(X_{t-})\,\mathrm{d}Z_t$$

is the *p*-dimensional SDE

$$\mathrm{d}Y_t = b(Y_{t-})\,\mathrm{d}Z_t$$

where $b : \mathbb{R}^p \to \mathbb{M}(p, d)$, $b_{ij}(y) = a_{ij}(y)$ for $i \neq m$ and $b_{mj}(y) = 0$, $Y_0^i = X_0^i$ for $i \neq m$ and $Y_0^m = \zeta$.

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This **directly defines** interventions in SDEs without specifying the underlying notion of causality.

However, a limiting argument shows that our notion of intervention is in agreement with intervening in the DAG (directed acyclic graph) of the discretized SDE given as below, where arrows corresponds to entries of a(x) independent of particular coordinates of x.



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In our SDE case, the fact that distinct SDEs can have the same distributions (for example, through varying the diffusion matrix) could be a source of complications.

Theorem. Consider the two SDEs

$$\mathrm{d}X_t = a(X_{t-})\,\mathrm{d}Z_t$$

and

$$\mathrm{d}X_t = \tilde{a}(X_{t-})\,\mathrm{d}\tilde{Z}_t,$$

where Z and \tilde{Z} are Lévy processes of dimension d and \tilde{d} , respectively, and a and \tilde{a} are Lipschitz and bounded. Then the results of intervention in the SDEs are the same whenever the Feller semigroups and the initial distributions are the same for the SDEs.

This enables practical inference of postintervention distributions through the following line of inference:



Consider again our previous plant growth example:





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Our results imply that for time-dependent SDE observations with Lévy noise, all intervention effects can be identified.

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In the DAG-based model for causal discovery, we:

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- Assume that the distribution of these variables have conditional independence properties consistent with some DAG.
- Wish to identify the DAG, corresponding to the causal network.

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- Assume that the distribution of these variables have conditional independence properties consistent with some DAG.
- Wish to identify the DAG, corresponding to the causal network.

The central problem is that the DAG is not uniquely identifiable from the distribution.

However, Shimizu et al (2006) showed the following. Assume that the variables X^1, \ldots, X^p are linearly related in the sense that

$$X = CX + \varepsilon$$

where $C \in \mathbb{M}(p, p)$ is acyclic in the sense that PCP^t is strictly lower triangular for some permutation matrix P. Also assume that the error variable ε has independent, non-degenerate and **non-Gaussian** coordinates of mean zero.

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Then, the DAG can be recovered from the distribution of X.

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$$X = A\varepsilon,$$

where ε has independent non-degenerate coordinates of mean zero. Both A and the distribution of ε are assumed unknown and are to be estimated. X is observed.

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From the results of Comon (1994), it follows that:

- **1** If the true distribution of ε has **only Gaussian coordinates**, A is identifiable up to transpose products.
- **2** If the true distribution of ε has **no Gaussian coordinates**, then A is identifiable up to scaling and permutation.

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Nonetheless, we might think that in practice, if the error distribution is close to Gaussian, the behaviour of the model will more closely resemble **the former of the two scenarios.**

Question: What actually happens to identifiability when we sample finitely many times and the error distribution is close to Gaussian while not being Gaussian?

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We set out to understand this phenomenon better.

To obtain results, we considered estimation only of the mixing matrix A, assuming the error distribution fixed. We considered error distributions which are independent and identical contaminated Gaussian distributions.

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Theorem. Consider *n* samples X_1, \ldots, X_n from the *p*-dimensional true distribution. Assume that the common error distribution P_n is $P_n = \beta_n \xi + (1 - \beta_n) \mathcal{N}$, where \mathcal{N} is the standard normal distribution. Letting *n* tend to infinity, it holds that:
Identifiability in ICA

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(Subject to very favorable interpretations)

Recall that one of our motivating examples for the development of a notion of causality for SDEs was an Ornstein-Uhlenbeck SDE for modeling gene expression networks:

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We will take particular interest in sparse networks, meaning that we wish to obtain **sparse estimates of** B (many entries equal to zero).

For simplicity, we consider the Ornstein-Uhlenbeck SDE

$$\mathrm{d}X_t = BX_t\,\mathrm{d}t + \,\mathrm{d}W_t$$

with mean reversion level zero and diffusion matrix I_p .

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We assume that we observe this SDE discretely at times $t_k = \Delta k$ for k = 0, ..., n. A natural loss function for the estimation of B is then $R : \mathbb{M}(p, p) \to [0, \infty)$ given by

$$R(B) = \sum_{k=1}^{n} \|X_{t_k} - \exp(\Delta B) X_{t_{k-1}}\|_2^2$$

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and sparse estimates can be obtained by, for $\lambda \geq$ 0,

$$\hat{B}_{\lambda}(X_{t_0},\ldots,X_{t_n}) = \operatorname*{argmin}_{B\in\mathbb{M}(p,p)} R(B) + \lambda \|B\|_1.$$



(Left matrix: True *B*, right matrix: \hat{B}_{λ} with $\lambda = 0.000$).



(Left matrix: True *B*, right matrix: \hat{B}_{λ} with $\lambda = 0.050$).



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Conclusion. Estimation through L^1 -penalized estimation with loss function R yields reasonable sparse estimates.

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Sad fact. We don't really know how to choose λ .

Workaround. Ignore the hard problem and solve a simpler problem.

Example. Consider the linear regression model

 $Y = X\beta + \varepsilon$

where $X \in \mathbb{M}(n, p)$, $\beta \in \mathbb{R}^p$ and ε is $\mathcal{N}(0, \sigma^2 I_n)$.

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and λ can be chosen by introducing the **generalization error** of \hat{eta}_{λ}

$$\operatorname{Err}_{\beta}(\hat{\beta}_{\lambda}) = E_{\beta} \|Y^* - X\hat{\beta}_{\lambda}\|_2^2,$$

where Y^* is independent of Y and has the same distribution as Y, and choosing λ as the minimizer of an estimate of $\text{Err}_{\beta}(\hat{\beta}_{\lambda})$.

In the linear regression model, Tibshirani and Taylor (2012) showed that

$$E_{\beta} \| Y^* - X \hat{\beta}_{\lambda} \|_2^2 = E_{\beta} \| Y - X \hat{\beta}_{\lambda} \|_2^2 + 2\sigma^2 E_{\beta} \text{ rank } X_{\mathcal{A}_{\lambda}},$$

where $\mathcal{A}_{\lambda} = \{i \leq p \mid \hat{\beta}_{\lambda} \neq 0\}.$

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Let us try this for a more complicated model (though still less complicated than the Ornstein-Uhlenbeck model).

We consider the nonlinear regression model

 $Y = \varphi(\beta) + \varepsilon$

where $\varphi : \mathbb{R}^p \to \mathbb{R}^n$ is continuous and ε is $\mathcal{N}(0, \sigma^2 I_n)$.

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which we call the **degrees of freedom** and the **Steinian degrees of freedom**, respectively, when defined. Here, the divergence can be a weak type of divergence.

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This is important because while $df(\hat{\beta})$ is **hard to estimate**, we can always obtain an unbiased estimate of $df_{\mathcal{S}}(\hat{\beta})$ as

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This allows estimation of the generalization error and therefore allows **model selection**, in our particular case of interest **sparse model selection** through choice of λ .

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Theorem I. For estimators of the type $\hat{\beta} = \operatorname{argmin}_{\beta \in K} ||Y - \varphi(\beta)||_2^2$, with $K \subseteq \mathbb{R}^p$ compact, it holds that $df(\hat{\beta}) \ge df_{\mathcal{S}}(\hat{\beta})$.

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Theorem II. With $\hat{\beta}$ as above and K the centered L^1 -ball of radius s,

$$\widehat{\mathsf{df}}_{\mathcal{S}}(\widehat{\beta}) = \mathsf{tr} \ J_{(\mathcal{A},\mathcal{A})}^{-1} \mathcal{G}_{(\mathcal{A},\mathcal{A})} - \frac{\gamma_{\mathcal{A}}^{t} J_{(\mathcal{A},\mathcal{A})}^{-1} \mathcal{G}_{(\mathcal{A},\mathcal{A})} J_{(\mathcal{A},\mathcal{A})}^{-1} \gamma_{\mathcal{A}}}{\gamma_{\mathcal{A}}^{t} J_{(\mathcal{A},\mathcal{A})}^{-1} \gamma_{\mathcal{A}}}.$$

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Theorem III. For an estimator $\hat{\beta} = \operatorname{argmin}_{\beta \in \mathbb{R}^p} \|Y - \varphi(\beta)\|_2^2 + \lambda \|\beta\|_1$,

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Thanks to:





Niels R. Hansen

My non-advisor co-authors

8

Ernst Hansen



Marloes H. Maathuis



Benjamin Falkeborg

Particular long-suffering teachers



Martin Jacobsen