

Likelihood estimation for the INAR(p) model by saddlepoint approximation

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Recent developments in statistics for complex dependent data

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Outline

- 1 Saddlepoint techniques
- 2 INAR(p) processes
- 3 SPA for INAR(p) processes
- 4 Simulations
- 5 Application

Past research

- Introduced into statistics in a seminal paper by Daniels (1954).
- Studied extensively during the last two decades, see e.g. Field and Ronchetti (1990), Jensen (1995) and Butler (2007).
- Quite complex theory but fairly straightforward use in practice.
- **Main advantage:** The high accuracy with which they can approximate intractable densities and tail probabilities, even for extremely small sample sizes (Davison, 2003).

Basic idea: From Taylor series to Saddlepoints

(Goutis and Casella, 1999)

- Consider a positive function $f(x)$ and suppose that our aim is to approximate its value at some point x_0 .
- Let $h(x) = \log f(x) \Rightarrow f(x) = \exp h(x)$
- **Taylor series expansion:**

$$f(x) \approx \exp \left\{ h(x_0) + (x - x_0)h'(x_0) + \frac{(x - x_0)^2}{2}h''(x_0) \right\}.$$

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- Equation (1) is exact if $h(x)$ is a quadratic equation.

Basic idea: From Taylor series to Saddlepoints (con't)

Laplace approximation

- Equation (1) can be useful for computing integrals of positive functions:

$$\int f(x)dx \approx \int \exp \left\{ h(\hat{x}) + \frac{(x - \hat{x})^2}{2} h''(\hat{x}) \right\} dx \quad (2)$$

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- If \hat{x} is a maximum, $h''(\hat{x}) < 0$ and (2) is equivalent to

$$\int f(x)dx \approx \exp\{h(\hat{x})\} \left(-\frac{2\pi}{h''(\hat{x})} \right)^{1/2} \quad (3)$$

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- Hence, for any fixed x ,

$$\begin{aligned} f(x) &\approx \int \exp \left\{ k(x, \hat{u}(x)) + \frac{(u - \hat{u}(x))^2}{2} \frac{\partial^2 k(x, u)}{\partial u^2} \Big|_{\hat{u}(x)} \right\} du \\ &= \exp \{k(x, \hat{u}(x))\} \left(-\frac{2\pi}{\frac{\partial^2 k(x, u)}{\partial u^2} \Big|_{\hat{u}(x)}} \right)^{1/2} \end{aligned} \quad (4)$$

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- Equation (4) is a **saddlepoint approximation** of $f(x)$ and $\hat{u}(x)$ is the **saddlepoint** that maximizes $k(x, u)$.

Basic idea: From Taylor series to Saddlepoints (con't)

- For a **density** $f(x)$, using the inversion formula, it can be proved that

$$k(x, u) = K_X(u) - ux$$

where $K_X(u)$ is the **cumulant generating function**, i.e.

$$K_X(u) = \log M_X(u) = \log \int_{-\infty}^{+\infty} \exp(ux) f(x) dx,$$

whith $M_X(u)$ denoting the moment generating function.

- The **saddlepoint** is the point $\hat{u}(x)$ that satisfies

$$K'_X(u) = x.$$

- After some tedious calculations, the **saddlepoint approximation to $f(x)$** proves to be

$$f_X(x) \approx \left(\frac{1}{2\pi K''_X(\hat{u}(x))} \right)^{1/2} \exp \{ K_X(\hat{u}(x)) - \hat{u}(x)x \}.$$

Basic idea: The saddlepoint method in a time series framework

Let $\{X_t, t \in \mathbb{Z}\}$ be a discrete-valued time series with available conditional pmf $f_{X_t|X_{t-1}, \dots, X_{t-p}}(x_t)$ and cumulant generating function (cgf)

$$K_t(u) = \log E(\exp(uX_t)|X_{t-1}, \dots, X_{t-p}),$$

for some $p \geq 1$. Then, the **saddlepoint approximation (SPA)** to the **conditional pmf** of X_t given X_{t-1}, \dots, X_{t-p} , is given by

$$\tilde{f}_{X_t|X_{t-1}, \dots, X_{t-p}}(x_t) = \left\{ \frac{1}{2\pi K_t''(\tilde{u})} \right\}^{1/2} \exp\{K_t(\tilde{u}) - \tilde{u}x_t\}, \quad (5)$$

where $K_t'(u)$ and $K_t''(u)$ denote the first and second order derivatives of $K_t(u)$ with respect to u , and \tilde{u}_t is the unique value of u satisfying $K_t'(\tilde{u}) = x_t$, for $t = p+1, p+2, \dots$

Basic idea: The saddlepoint method in a time series framework (con't)

- Expression (5) is the leading term of a saddlepoint density approximation, but further terms can be included.
- For instance, a **second-order approximation** is

$$\tilde{f}_{X_t|X_{t-1},\dots,X_{t-p}}(x_t) = \left\{ \frac{1}{2\pi K_t''(\tilde{u})} \right\}^{1/2} \exp \{K_t(\tilde{u}) - \tilde{u}_t x_t\} \left\{ 1 + \left(\frac{1}{8} \tilde{\kappa}_t^{(4)} - \frac{5}{24} [\tilde{\kappa}_t^{(3)}]^2 \right) \right\},$$

where $\tilde{\kappa}_t^{(r)} = K_t^{(r)}(\tilde{u}) / \{K_t''(\tilde{u})\}^{r/2}$, $r \geq 3$ is the r -th order standardized cumulant of X_t (Butler, 2007).

- In the classical setting, X_t is taken to be an average of m independent observations, and then it may be shown that its true and approximate densities f and \tilde{f} are related by $f(x_t) = \tilde{f}(x_t) \{1 + \mathcal{O}(m^{-1})\}$.

Remarks

- $\tilde{f}_{X_t|X_{t-1},\dots,X_{t-p}}(x_t)$ is always positive, but it does not sum to one exactly. Hence, it is often renormalized, a practice which also improves the relative order of the approximation, see e.g. Field and Ronchetti (1990) and Kolassa (2006). Renormalization is not possible when considering conditional probabilities.

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- Approximation (5) is based on the normal distribution but alternative approximations based on other distributions, such as the Gamma and inverse Gaussian distributions, are also available (Wood et al., 1993).

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- The saddlepoint approximation requires that the cgf exists.

Model

Definition

A sequence of random variables $\{X_t, t \in \mathbb{Z}\}$ is an **integer-valued autoregressive**, INAR(p), **process** if it satisfies a difference equation of the form

$$X_t = \sum_{i=1}^p \alpha_i \circ X_{t-i} + \epsilon_t, \quad (6)$$

where $\alpha_i \in [0, 1)$ are fixed constants and $\{\epsilon_t\}$ is a sequence of uncorrelated non-negative integer-valued random variables having mean μ_ϵ and finite variance σ_ϵ^2 . The sequence $\{\epsilon_t\}$ is called the innovation process (McKenzie, 1985; Al-Osh and Alzaid, 1987).

Model (con't)

- Definition (6) is based on the notion of **binomial thinning**.
- The binomial thinning operator ' \circ ' is defined by

$$\alpha \circ X = \sum_{j=1}^X Y_j$$

where Y_j are i.i.d. Bernoulli random variables with $P(Y_j = 1) = 1 - P(Y_j = 0) = \alpha$.

- The binomial thinning operations involved in (6) are independent, so the INAR(p) process has the classical AR(p) correlation structure.
- A unique strictly stationary and ergodic solution of (6) exists if

$$\sum_{i=1}^p \alpha_i < 1.$$

Example: $p = 1$

Let $\alpha \in [0, 1)$ and let $\{\epsilon_t\}$ be a sequence of i.i.d. non-negative integer valued random variables with $E[\epsilon_t] = \mu_\epsilon$ and $\text{Var}[\epsilon_t] = \sigma_\epsilon^2$. The integer-valued autoregressive process of order 1, **INAR(1)**, $\{X_t, t \in \mathbb{Z}\}$ is defined by the equation

$$X_t = \alpha \circ X_{t-1} + \epsilon_t,$$

where $\alpha \circ X_{t-1}$ is the sum of X_{t-1} Bernoulli random variables all of which are independent of X_{t-1} .

Example: $p = 1$ (con't)

It can be shown that

- $E[X_t] = \mu_\epsilon / (1 - \alpha)$,
- $\text{Var}[X_t] = (\alpha\mu_\epsilon + \sigma_\epsilon^2) / (1 - \alpha^2)$
- the autocovariance function evaluated at lag k , $c(k)$, is given by

$$c(k) \equiv \text{Cov}[X_t, X_{t-k}] = \alpha^k c(0). \quad (7)$$

- Consequently, the autocorrelation function, $\rho(k)$, is

$$\rho(k) = \frac{c(k)}{c(0)} = \alpha^k, \quad (8)$$

so that $\rho(k)$ decays exponentially with the lag k as in AR(1), but unlike the autocorrelation of a stationary AR(1) process, it is always positive for $\alpha \in (0, 1)$.

Saddlepoint Approximation for INAR processes

Proposition

Let $\{X_t\}$ be the INAR(p) process defined by (6). For large values of x_{t-i} and if $0 < \sum_{i=1}^p \alpha_i < 1$, the exact and approximate densities are related by

$$\begin{aligned}
 f_{X_t|X_{t-1},\dots,X_{t-p}}(x_t) &= \tilde{f}_{X_t|X_{t-1},\dots,X_{t-p}}(x_t) \left\{ 1 + \mathcal{O} \left(\frac{1}{\sum_{i=1}^p x_{t-i} + 1} \right) \right\} \\
 &= \left\{ \frac{1}{2\pi K_t''(\tilde{u}_t)} \right\}^{1/2} \exp \{ K_t(\tilde{u}_t) - \tilde{u}_t x_t \} \\
 &\times \left\{ 1 + \mathcal{O} \left(\frac{1}{\sum_{i=1}^p x_{t-i} + 1} \right) \right\}.
 \end{aligned}
 \tag{9}$$

Inference

Maximum likelihood estimation (MLE)

Let $\boldsymbol{\theta} = (\alpha_1, \alpha_2, \dots, \alpha_p, \mu_\epsilon)^T$ be the unknown parameter vector. The maximum likelihood estimator of $\boldsymbol{\theta}$ is calculated by maximizing the conditional log likelihood function

$$\ell(\boldsymbol{\theta}) = \sum_{t=p+1}^n \log P(X_t = x \mid X_{t-1} = x_{t-1}, \dots, X_{t-p} = x_{t-p}),$$

that is $\hat{\boldsymbol{\theta}}_{\text{MLE}} = \arg \max_{\boldsymbol{\theta}} \ell(\boldsymbol{\theta})$.

Inference (con't)

Example

An important special case is that of **Poisson INAR(1) process**

$X_t = \alpha \circ X_{t-1} + \epsilon_t$, with $\{\epsilon_t\}$ a sequence of i.i.d. Poisson random variables with mean μ_ϵ . The conditional distribution of X_t given X_{t-1} is

$$P(X_t = x | X_{t-1} = x_{t-1}) = x_{t-1}! \exp(-\mu_\epsilon) \sum_{i=1}^m \frac{\alpha^i (1 - \alpha)^{x_{t-1}-i} \mu_\epsilon^{x-i}}{i! (x_{t-1} - i)! (x - i)!},$$

where $m = \min(x_{t-1}, x)$.

Inference (con't)

Example

Under a **Poisson INAR(p)** process,

$$\begin{aligned}
 P(X_t = x | X_{t-1} = x_{t-1}, \dots, X_{t-p} = x_{t-p}) = \\
 \sum_{i_1=0}^{\min(x_{t-1}, x)} \binom{x_{t-1}}{i_1} \alpha_1^{i_1} (1 - \alpha_1)^{x_{t-1} - i_1} \\
 \times \sum_{i_2=0}^{\min(x_{t-2}, x - i_1)} \binom{x_{t-2}}{i_2} \alpha_2^{i_2} (1 - \alpha_2)^{x_{t-2} - i_2} \\
 \dots \sum_{i_p=0}^{\min(x_{t-p}, x - (i_1 + \dots + i_{p-1}))} \binom{x_{t-p}}{i_p} \alpha_p^{i_p} (1 - \alpha_p)^{x_{t-p} - i_p} \\
 \times \frac{\exp(-\mu_\epsilon) \mu_\epsilon^{x - (i_1 + \dots + i_p)}}{(x - (i_1 + \dots + i_p))!}.
 \end{aligned}$$

Motivation

- Maximization of $\ell(\boldsymbol{\theta})$ is quite cumbersome, owing to the nested summations involved in its computation, and the numerical difficulties that can arise when summing up many small probabilities.
- **Alternative approach:** Approximate the log-likelihood of the INAR(p) model by the saddlepoint method.
- **The saddlepoint approximation (SPA),**
 - circumvents computational difficulties,
 - provides simple and highly accurate approximations to the conditional maximum likelihood estimators even for complicated INAR models.

Generalities

Key observation

Conditional on $X_{t-p} = x_{t-p}, \dots, X_{t-1} = x_{t-1}$, X_t is a sum of independent binomial variates with parameters (x_{t-i}, α_i) , plus the innovation term ϵ_t . Therefore, its conditional cgf is

$$K_t(u) = \sum_{i=1}^p x_{t-i} \log\{1 - \alpha_i + \alpha_i \exp(u)\} + K_{\epsilon_t}(u),$$

where $K_{\epsilon_t}(u)$ denotes the cumulant-generating function of ϵ_t .

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Remarks:

- The domain of $K_t(u)$ is the same set as the domain of $K_{\epsilon_t}(u)$. This is the entire real line for Poisson innovations, and an interval of form $(-\infty, c)$ for negative binomial innovations.
- $K_t(u)$ is a sum of convex functions and thus itself convex.

Generalities (con't)

Approximate log-likelihood function

$$\tilde{\ell}(\boldsymbol{\theta}) = \sum_{t=p+1}^n \log \tilde{f}_{X_t|X_{t-1}, \dots, X_{t-p}}(x_t),$$

where $\tilde{f}_{X_t|X_{t-1}, \dots, X_{t-p}}(x_t) = \left\{ \frac{1}{2\pi K_t''(\tilde{u})} \right\}^{1/2} \exp \{K_t(\tilde{u}) - \tilde{u}_t x_t\}$, is the SPA to the conditional pmf of X_t (Daniels, 1954). The value $\tilde{\boldsymbol{\theta}}$ maximizing $\tilde{\ell}(\boldsymbol{\theta})$ is the saddlepoint maximum likelihood estimator (SPMLE).

Generalities (con't)

Evaluation of $\tilde{\ell}(\theta)$

- It requires the computation of $n - p$ values of \tilde{u} , one for each term in the sum.
- For $x_t = 0$ the pmf need not to be approximated, since

$$P(X_t = 0 \mid X_{t-1} = x_{t-1}, \dots, X_{t-p} = x_{t-p}) = f_{\epsilon_t}(0) \prod_{i=1}^p (1 - \alpha_i)^{x_{t-i}}.$$

- If the saddlepoint equation $K'_t(u) = x_t$ cannot be solved analytically, then \tilde{u}_t can be approximated numerically, e.g. using the Newton-Raphson method of Lieberman (1994): linear expansion of the saddlepoint equation in the neighborhood of an initial value u_0 , i.e. $K'_t(u_0) + K''_t(u_0)(\tilde{u}_t - u_0) \approx x_t$, to provide the update

$$\tilde{u}_t = \frac{x_t - K'_t(u_0)}{K''_t(u_0)} + u_0,$$

which is iterated until the approximation is adequate.

Asymptotic properties of the SPMLE

- Established result (Bu et al., 2008):

$$\sqrt{n}(\hat{\boldsymbol{\theta}} - \boldsymbol{\theta}_0) \xrightarrow{d} N\{0, I^{-1}(\boldsymbol{\theta}_0)\}, \quad n \rightarrow \infty, \quad (10)$$

where $\hat{\boldsymbol{\theta}}$ is the MLE of $\boldsymbol{\theta}_0$ based on data X_1, \dots, X_n from a stationary INAR(p) process, and $I(\boldsymbol{\theta}_0)$ is the Fisher information in a single observation.

- We have proved that

$$\tilde{f}(x_t | x_{t-1}, \dots, x_{t-p}) = \left\{ \frac{1}{2\pi\sigma_t^2(\boldsymbol{\theta})} \right\}^{1/2} \exp\left[-\frac{\{x_t - \mu_t(\boldsymbol{\theta})\}^2}{2\sigma_t^2(\boldsymbol{\theta})}\right], \quad (11)$$

which is the probability density function of a Gaussian random variable with

$$\mu_t(\boldsymbol{\theta}) = E(X_t | X_{t-1}, \dots, X_{t-p}) = \sum_{i=1}^p \alpha_i x_{t-i} + \mu,$$

$$\sigma_t^2(\boldsymbol{\theta}) = \text{Var}(X_t | X_{t-1}, \dots, X_{t-p}) = \sum_{i=1}^p \alpha_i (1 - \alpha_i) x_{t-i} + \sigma_\epsilon^2.$$

Asymptotic properties of the SPMLE (con't)

- The conditional mean and variance can be used to construct a quasilielihood function for the estimation of θ , and the resulting estimator is asymptotically normally distributed; see Theorem 3.2.23 of Taniguchi and Kakizawa (2000).
- Empirical results indicate close agreement between the MLE $\hat{\theta}$ and the SPMLE $\tilde{\theta}$.
- Proposition 2.1 can be used to show that in the usual asymptotic setting, with $n \rightarrow \infty$ and the parameters fixed, the SPMLE is inconsistent. However, when the process assumes large values, i.e., when the mean of process is growing, then it can be shown that that the SPMLE is a consistent estimator of θ .

SPA for Poisson INAR(p) processes

Assume that the error sequence $\{\epsilon_t\}$ consists of i.i.d. Poisson random variables with mean λ . Then,

- $K_t(u) = \sum_{i=1}^p x_{t-i} \log(1 - \alpha_i + \alpha_i e^u) + \lambda(e^u - 1),$
- $K'_t(u) = \sum_{i=1}^p \left[x_{t-i} \frac{\alpha_i e^u}{\alpha_i e^u + (1 - \alpha_i)} \right] + \lambda e^u,$
- $K''_t(u) = \sum_{i=1}^p \left[x_{t-i} \frac{\alpha_i(1 - \alpha_i)e^u}{(\alpha_i e^u + (1 - \alpha_i))^2} \right] + \lambda e^u.$

SPA for Poisson INAR(p) processes (con't)

The SPA to the log-likelihood is given by

$$\begin{aligned}\tilde{\ell}(\boldsymbol{\theta}) = & -\frac{1}{2} \sum_{t=p+1}^n \log \left\{ \sum_{i=1}^p \left[x_{t-i} \frac{\alpha_i (1 - \alpha_i) e^{\tilde{u}_t}}{(\alpha_i e^{\tilde{u}_t} + (1 - \alpha_i))^2} \right] + \lambda e^{\tilde{u}_t} \right\} \\ & + \sum_{t=p+1}^n \left\{ \left[\sum_{i=1}^p [x_{t-i} \log(\alpha_i e^{\tilde{u}_t} + (1 - \alpha_i))] + \lambda (e^{\tilde{u}_t} - 1) \right] - \tilde{u}_t \right\},\end{aligned}$$

evaluated for each time point $t = p + 1, \dots, n$, at $u = \tilde{u}_t$.

Existence of a finite saddlepoint \tilde{u}_t

Theorem

Let X be a random variable supported on the possibly infinite interval $[-x_1, x_2]$, whose cumulant generating function $K_X(u)$ is defined on an interval with endpoints $-c_1$ and c_2 , and $x_i < \infty$ or $c_i = \infty$ for $i = 1, 2$. Then a solution \tilde{u}_t of the saddlepoint equation exists for all values of $x \in (-x_1, x_2)$ if (Kolassa, 2006)

$$\lim_{u \rightarrow -c_1} K'_X(u) = -x_1, \quad \lim_{u \rightarrow c_2} K'_X(u) = x_2.$$

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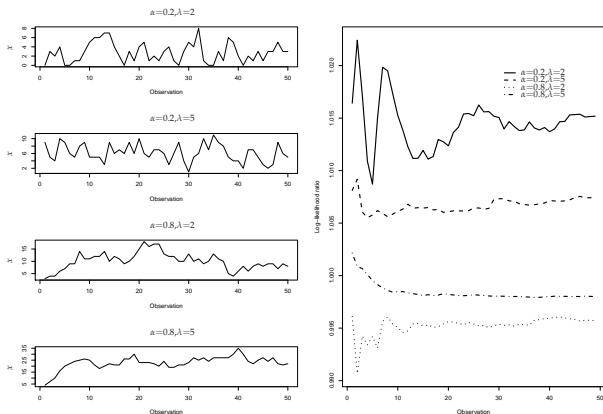
$$\lim_{u \rightarrow -c_1} K'_X(u) = -x_1, \quad \lim_{u \rightarrow c_2} K'_X(u) = x_2.$$

- For an INAR(p) process, $X_t \in [0, \infty)$ and hence $x_1 = 0$ and $x_2 = \infty$.
- For the Poisson INAR(p) model, the existence of a unique solution to the equation $K'_t(u) = x_t$ is easily established, since $K'_t(u)$ is strictly monotone increasing in u , and

$$\lim_{u \rightarrow -\infty} K'_t(u) = 0, \quad \lim_{u \rightarrow \infty} K'_t(u) = \infty.$$

Quality of the approximation

Quality of the proposed approximation for four simulated series of length $n = 50$, shown in the left panel, generated from an INAR(1) process with $\alpha = 0.2$ or 0.8 , $\lambda = 2$ or 5 . The right panel shows the ratio of the true log-likelihood function to the approximate log-likelihood obtained by the saddlepoint method. The relative error is at most $\pm 2.5\%$. All ratios fluctuate around 1 indicating the high quality of the saddlepoint approximation. The accuracy increases for large mean values of the generated process, when the convolutions involved in the transition probabilities become more awkward.



Design

- Simulation experiments under the assumptions of INAR(2) and INAR(4) processes with Poisson and negative binomial innovations.

	Estimation methods			Sample sizes	
	CLS	MLE	SPA	n	
INAR(2)	✓	✓	✓	50	500
INAR(4)	✓		✓	100	500

- Logit and log transformations to avoid inadmissible parameter estimates:

$$\xi = \log(\alpha/(1 - \alpha)) \rightsquigarrow \alpha = \exp(\xi)/(1 + \exp(\xi))$$

$$\eta = \log(\mu) \rightsquigarrow \mu = \exp(\eta)$$

- 1000 replicate simulations in each setting.
- Use of the `optim()` function in R for the minimization of the least squares and negative log likelihoods.

Results for INAR(2) processes

Poisson INAR(2) with $(\alpha_1, \alpha_2, \lambda) = (0.3, 0.4, 5)$

		Bias $\times 100$			MSE $\times 100$		
		CLS	MLE	SPA	CLS	MLE	SPA
$n = 50$	$\hat{\alpha}_1$	-3.0	-1.6	-1.6	1.9	2.4	2.2
	$\hat{\alpha}_2$	-5.4	-3.6	-3.8	1.9	2.2	2.1
	$\hat{\lambda}$	126.7	76.7	79.3	577.5	379.5	390.0
$n = 500$	$\hat{\alpha}_1$	-0.4	-0.2	0.1	0.2	0.2	0.2
	$\hat{\alpha}_2$	-0.6	-0.3	-0.6	0.2	0.1	0.1
	$\hat{\lambda}$	15.8	7.9	10.9	47.8	32.9	34.2

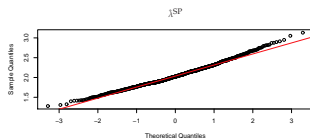
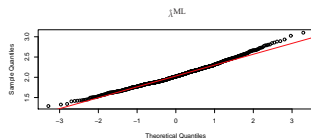
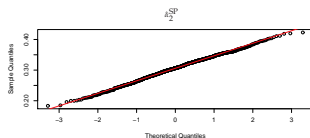
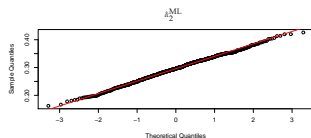
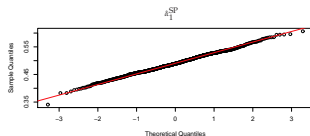
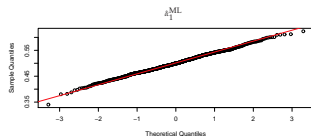
INAR(2)–NegBin innovations with $(\alpha_1, \alpha_2, \mu, r^*) = (0.5, 0.3, 2, 3)$

		Bias $\times 100$			MSE $\times 100$		
		CLS	MLE	SPA	CLS	MLE	SPA
$n = 50$	$\hat{\alpha}_1$	4.1	0.8	1.7	2.3	0.9	1.0
	$\hat{\alpha}_2$	-5.7	-1.2	-0.9	2.1	1.0	1.2
	$\hat{\mu}$	87.7	2.1	-8.7	280.8	37.0	42.2
$n = 500$	$\hat{\alpha}_1$	-0.2	0.1	0.9	0.2	0.1	0.1
	$\hat{\alpha}_2$	-0.9	-0.2	0.6	0.5	0.1	0.1
	$\hat{\mu}$	10.1	0.1	-14.9	15.5	4.0	6.6

* r is kept fixed

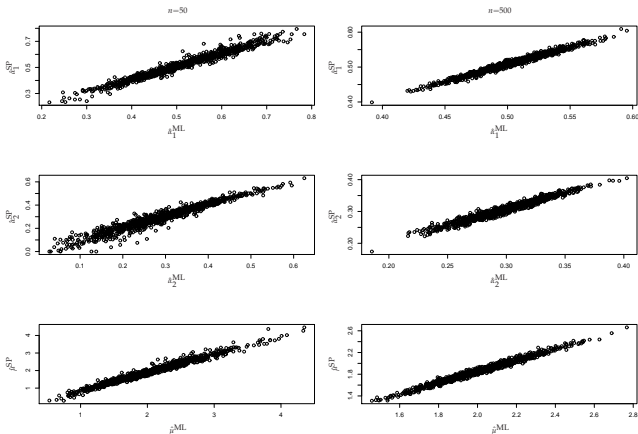
Results for INAR(2) processes (con't)

Normal Q–Q plots for the MLE and SPA estimates obtained from a Poisson INAR(2) model ($n = 500$) with true parameter values $(\alpha_1, \alpha_2, \lambda) = (0.3, 0.4, 5)$. Results have been obtained after 1000 simulation replicates.



Results for INAR(2) processes (con't)

Scatterplots between the MLE and SPA estimates obtained from an INAR(2) model with negative binomial innovations and true parameter values $(\alpha_1, \alpha_2, \mu, r) = (0.5, 0.3, 2, 3)$.

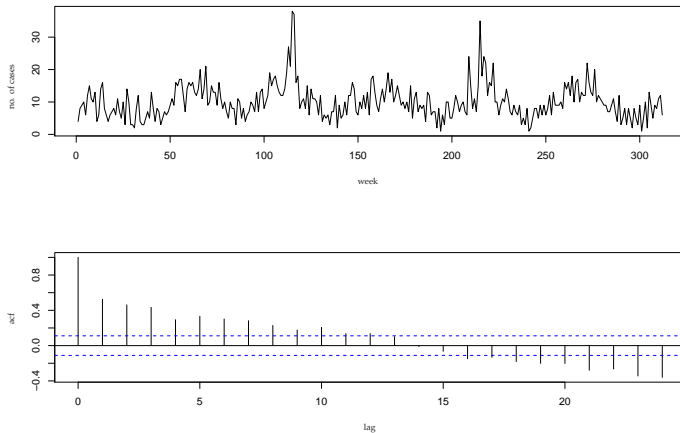


Data

- Weekly number of meningococcal disease cases in Germany for the years 2001–2006 ($n = 312$).
- **Source:** the German national surveillance system for notifiable diseases, administered by the Robert Koch Institute (RKI).
- Significant **overdispersion** present in the data (mean 10.09, variance 27.83) \rightsquigarrow Need for a more flexible distribution than the Poisson distribution, e.g. **negative binomial**.

Data (con't)

Time series and acf plots for the weekly number of meningococcal disease cases in Germany, 2001–2006.



Modelling

- Let X_t denote the number of meningococcal disease cases at time $t = 1, \dots, 312$.
- X_t is defined as an INAR(p) process of the form,

$$X_t = \sum_{i=1}^p \alpha_i \circ X_{t-i} + \epsilon_t, \quad \epsilon_t \sim \text{NegBin}(\mu_t, r).$$

- To account for seasonality,

$$\log(\mu_t) = \beta_0 + \sum_{s=1}^S (\beta_{1;s} \sin(\omega_s t) + \beta_{2;s} \cos(\omega_s t)),$$

where S is the number of harmonics to include and ω_s are Fourier frequencies of the form $\omega_s = 2\pi s/52$ (see also Paul et al., 2008).

- Challenges:** Model selection & parameter estimation through the application of the SPA technique.

Results

INAR(p) models fitted to the meningococcal disease data. $\log L$ denotes the approximate maximized log-likelihood obtained from the saddlepoint approximation.

Model	p	r	S	$\log L$	AIC
1.	1	0.5	1	-858.11	1724.2
2.	1	0.5	2	-857.73	1727.5
3.	1	1	1	-862.11	1732.2
4.	1	1	2	-861.80	1735.6
5.	2	0.5	1	-851.51	1713.0
6.	2	0.5	2	-851.21	1716.4
7.	2	1	1	-869.01	1748.0
8.	2	1	2	-868.62	1751.2
9.	3	0.5	1	-844.95	1701.9
10.	3	0.5	2	-844.49	1705.0
11.	3	1	1	-859.08	1730.2
12.	3	1	2	-859.09	1734.2
13.	4	0.5	1	-845.46	1704.9
14.	4	0.5	2	-858.95	1731.9
15.	4	1	1	-845.55	1709.1
16.	4	1	2	-862.46	1742.9

Results

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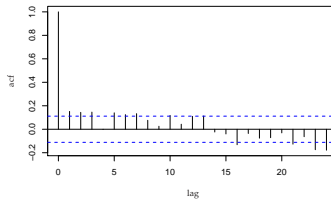
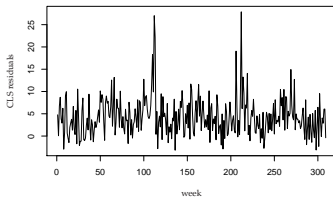
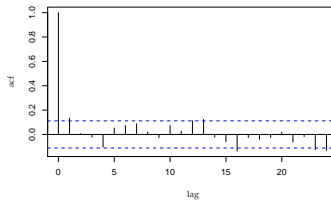
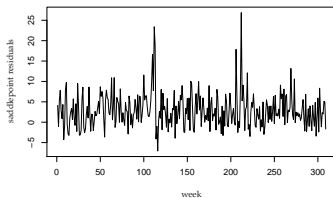
Results (con't)

Parameter estimates and standard errors (s.e.) obtained after fitting model 9 to the meningococcal disease data.

	Saddlepoint approximation		CLS estimation	
	estimate	s.e.	estimate	s.e.
α_1	0.189	0.052	0.216	0.057
α_2	0.196	0.051	0.096	0.059
α_3	0.185	0.049	0.092	0.057
β_0	1.419	0.092	1.749	0.140
$\beta_{1;1}$	0.334	0.083	0.284	0.061
$\beta_{2;1}$	0.324	0.079	0.380	0.058

Results (con't)

Time series and acf plots for the SPA and CLS residuals, $e_t = X_t - \sum_{i=1}^3 \hat{\alpha}_i X_{t-i} - \hat{\mu}_t$, obtained after fitting model 9 to the meningococcal disease data.



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