

Binomial Autoregressive Processes with Density-Dependent Thinning



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Time Series of Counts with a Finite Range

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Motivation & Outline

During last 30 years,
increasing research interest in count data time series.
Initially, research focussed nearly exclusively
on counts with unlimited range $\mathbb{N}_0$.

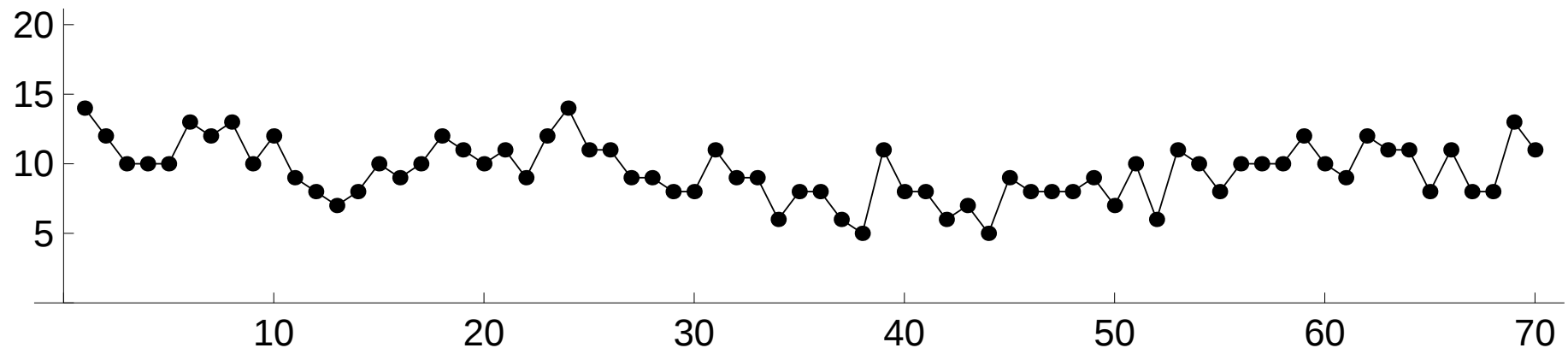
Only during last years, considerable interest also
in time series with **fixed finite range** $\{0, 1, \dots, N\}$.

Examples:

- utilization of computer pools (with N workstations);
- spread of metapopulations (with N patches);
- . . .

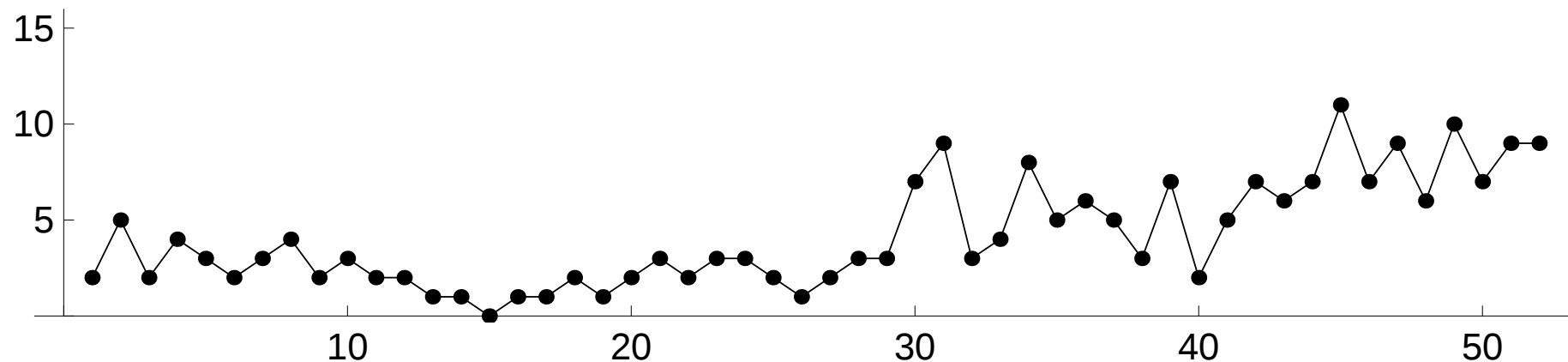
Data example 1: (Weiß & Kim, 2013)

Number of securities companies (among $N = 22$ companies) traded in Korea stock market per 5-min period (Feb. 8, 2011, 09:00–14:50; $T = 70$).



Data example 2: (Robert-Koch-Institut, survstat.rki.de)

Number of districts in Germany (among $N = 38$ districts)
with new case of hantavirus infection
(weekly data, 2011; $T = 52$).



Modeling of time series with fixed finite range $\{0, 1, \dots, N\}$?

Popular **basic approach** (counterpart to AR(1) model):

binomial AR(1) model (BAR(1)) by McKenzie (1985),

which uses **binomial thinning** operation

$\alpha \circ X \sim \text{Bin}(X, \alpha)$ by Steutel & van Harn (1979).

Parameters $\pi \in (0; 1)$, $\rho \in \left(\max \left\{ -\frac{\pi}{1-\pi}, -\frac{1-\pi}{\pi} \right\} ; 1 \right)$,

define thinning probabilities $\beta := \pi (1 - \rho)$ and $\alpha := \beta + \rho$.

BAR(1) recursion

$$X_{t+1} = \underbrace{\alpha \circ X_t}_{\text{survivors}} + \underbrace{\beta \circ (N - X_t)}_{\text{newly occupied}},$$

thinnings performed independently, independent of $(X_s)_{s \leq t}$.

A few well-known properties:

Ergodic Markov chain, transition probabilities

$$P(X_{t+1} = k \mid X_t = l) =$$

$$\sum_{m=\max\{0, k+l-N\}}^{\min\{k, l\}} \binom{l}{m} \binom{N-l}{k-m} \alpha^m (1-\alpha)^{l-m} \beta^{k-m} (1-\beta)^{N-l+m-k},$$

uniquely determined stationary distribution: $\text{Bin}(N, \pi)$.

Autocorrelation function: $\rho_X(k) = \rho^k$ for $k \geq 0$.

Regression properties:

$$E[X_{t+1} \mid X_t] = \rho \cdot X_t + N\beta,$$

$$V[X_{t+1} \mid X_t] = \rho(1-\rho)(1-2\pi) \cdot X_t + N\beta(1-\beta).$$

Limitations:

- Binomial marginal distribution,
- exponentially decaying ACF (AR(1)-like),
- thinning probabilities at time t
do not depend on process up to that time, ...

Approaches for generalization:

- p th order dependence (AR(p)-like): Weiß (2009).
- State dependence of parameters π, ρ resp. α, β at time $t + 1$
 - as function of X_t/N : “density dependence”;
(e.g., proportion of infectives or proportion of occupied patches)
 - acc. to X_t in which regime: “SET approach” (Tong).



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Density-Dependent Binomial $AR(1)$ Processes

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Definition & Properties

Definition: (Weiß & Pollett, 2014)

Let $\pi : [0; 1] \rightarrow (0; 1)$ and $\rho : [0; 1] \rightarrow (0; 1)$, so functions $\beta(y) := \pi(y)(1 - \rho(y))$ and $\alpha(y) := \beta(y) + \rho(y)$ also range $(0; 1)$.

Write $\pi_{t+1} := \pi(X_t/N)$, $\rho_{t+1} := \rho(X_t/N)$,

and $\alpha_{t+1} := \alpha(X_t/N)$, $\beta_{t+1} := \beta(X_t/N)$.

DD-BAR(1) recursion

$$X_{t+1} = \alpha_{t+1} \circ X_t + \beta_{t+1} \circ (N - X_t),$$

thinnings performed independently, independent of $(X_s)_{s \leq t}$.

General properties: (Weiß & Pollett, 2014)

- time-homogeneous finite-state Markov chain
with truly positive transition probabilities

$$P(k|l) := P(X_t = k \mid X_{t-1} = l) = \sum_{m=\max\{0, k+l-N\}}^{\min\{k, l\}} \binom{l}{m} \binom{N-l}{k-m} (\alpha(l/N))^m (1 - \alpha(l/N))^{l-m} (\beta(l/N))^{k-m} (1 - \beta(l/N))^{N-l+m-k};$$

- regression properties

$$E[X_t \mid X_{t-1}] = \rho_t X_{t-1} + N \beta_t,$$

$$V[X_t \mid X_{t-1}] = \rho_t(1 - \rho_t)(1 - 2\pi_t) X_{t-1} + N \beta_t(1 - \beta_t);$$

- . . .

General properties: (cont.)

- due to primitive transition matrix $\mathbf{P} = (P(k|l))_{k,l=0,\dots,N}$, ergodic with unique stationary distribution, obtained numerically from equation

$$\mathbf{P} p = p \quad (\text{eigenvalue problem})$$

(also allows to evaluate stationary moments);

- second-order moments with time-lag h
from $P(X_t = k, X_{t-h} = l)$ as entries of $\mathbf{P}^h \text{diag}(p)$, etc.;
- . . .

Large- N Approximations: Define

$$f(x) := \alpha(x)x + \beta(x)(1-x) = \rho(x)x + \beta(x),$$

$$v(x) := \alpha(x)(1-\alpha(x))x + \beta(x)(1-\beta(x))(1-x),$$

so $E[X_t/N | X_{t-1}/N] = f(X_{t-1}/N)$, $N V[X_t/N | X_{t-1}/N] = v(X_{t-1}/N)$.

Several large- N results, among others

- law of large numbers $X_t/N \xrightarrow[N \rightarrow \infty]{P} y_t$, where $y_{t+1} = f(y_t)$;
- central limit law for scaled fluctuations $\sqrt{N}(X_t/N - y_t)$;
- if y^* stable fixed point of f , and $\kappa = f'(y^*)$, then

$$\mu_X \approx N y^*, \quad \sigma_X^2 \approx N \frac{v(y^*)}{1 - \kappa^2}, \quad \rho_X(k) \approx \kappa^k.$$

(Buckley & Pollett, 2010; Weiß & Pollett, 2014)



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Density-Dependent Binomial AR(1) Processes

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Special Cases

Binomial index of dispersion for r.v. with support $\{0, \dots, N\}$, and with mean μ and variance σ^2 :

$$I_d := \frac{N\sigma^2}{\mu(N - \mu)} \in (0; \infty).$$

For $\text{Bin}(N, \pi)$ -distributed r.v., we have $I_d = 1$ for any $\pi \in (0; 1)$.

If $I_d > 1$, then **overdispersion** w.r.t. binomial distribution (“extra-binomial variation”),

while **underdispersion** refers to $I_d < 1$.

Definition: Let $\rho(y) = \rho \in (0; 1)$ be constant,
and $\pi(y) = a + by$ be linear, where $a, a + b \in (0; 1)$. So

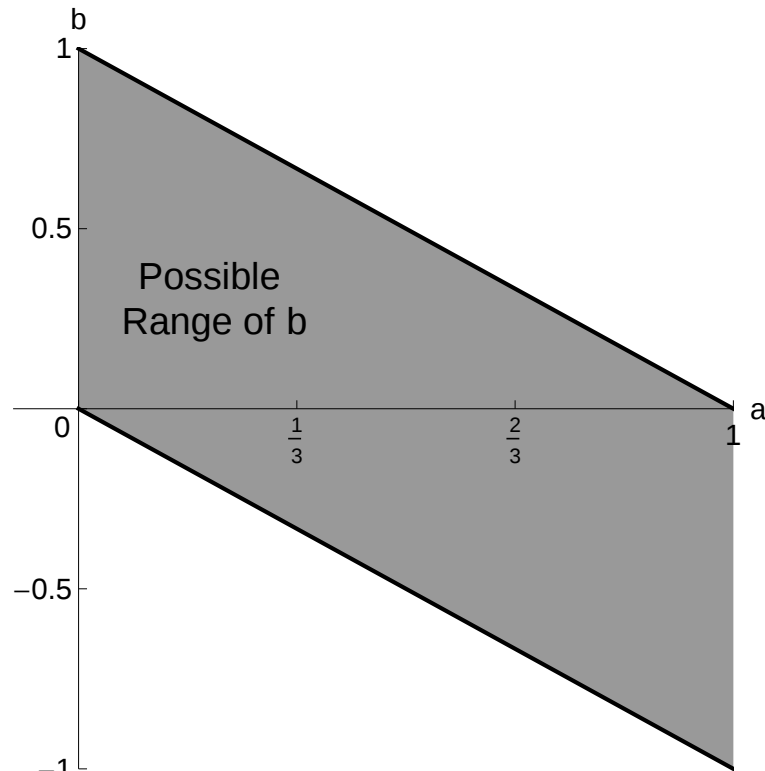
$$\beta_t = (1 - \rho) (a + b X_{t-1}/N), \quad \alpha_t = (1 - \rho) (a + b X_{t-1}/N) + \rho.$$

also linear, and depending on sign of b , these probabilities increase or decrease with increasing density.

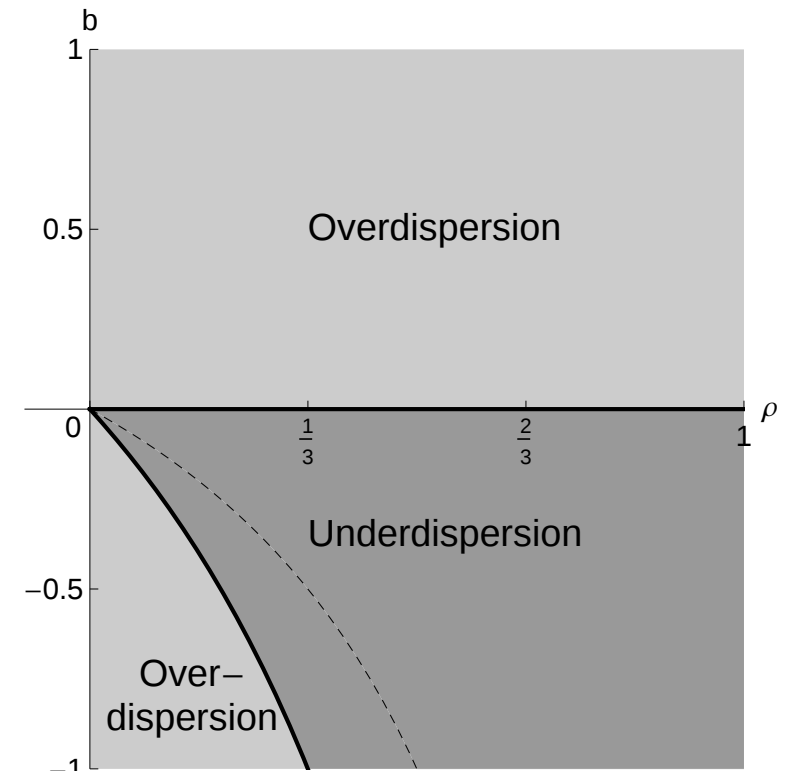
Properties: ACF $\rho_X(k) = (\rho + (1 - \rho)b)^k$, and

$$\mu = \frac{Na}{1 - b}, \quad I_d = \frac{1 - \rho^2}{1 - \frac{\rho^2}{N} - (1 - \frac{1}{N})(\rho + (1 - \rho)b)^2}.$$

Properties:



(a)



(b)

Attainable range of b depending on a in (a)
dispersion determined by b and ρ in (b).

Data example 1: securities counts, $N = 22$.

Analyzing empirical (partial) autocorrelations,
first-order autoregressive dependence structure apparent.

Sample mean and variance are $\bar{x} \approx 9.529$ and $s^2 \approx 4.253$.

Empirical index of dispersion $\hat{I}_d = s^2 / (\bar{x}(1 - \bar{x}/N)) \approx 0.787$
indicates slight degree of binomial underdispersion.

ML-fitted model:

$$\hat{a}_{\text{ML}} \approx 0.693, \quad \hat{b}_{\text{ML}} \approx -0.619, \quad \hat{\rho}_{\text{ML}} \approx 0.630.$$

Boundary case $\rho \rightarrow 0$ leads to

$$X_t \stackrel{D}{=} \text{Bin}(N, a + b X_{t-1}/N).$$

Motivated by analogy to (Poisson) INARCH(1) model (Ferland et al., 2006): **binomial INARCH(1) model** (BINARCH(1)).

Properties: ACF $\rho_X(k) = b^k$, and

$$\mu = \frac{Na}{1-b}, \quad I_d = \frac{1}{1 - (1 - \frac{1}{N})b^2} \in [1; N).$$

Consequently, only overdispersion is possible.

Data example 2: hanta counts, $N = 38$.

Sample mean and variance are $\bar{x} \approx 4.173$ and $s^2 \approx 7.793$.

Empirical index of dispersion $\hat{I}_d \approx 2.098$

indicates considerable degree of binomial overdispersion.

	$\hat{\pi}_{\text{ML}}$	$\hat{\rho}_{\text{ML}}$	AIC	BIC	
BAR(1)	0.115 (0.013)	0.535 (0.071)	222.8	226.7	
	$\hat{a}_{\text{ML}}$	$\hat{b}_{\text{ML}}$	$\hat{\rho}_{\text{ML}}$	AIC	BIC
DD-BAR(1)	0.030 (0.016)	0.748 (0.143)	0.000 (0.367)	213.4	219.2
	$\hat{a}_{\text{ML}}$	$\hat{b}_{\text{ML}}$		AIC	BIC
BINARCH(1)	0.030 (0.011)	0.748 (0.108)		211.4	215.3

Definition: Let $\alpha(y) = \alpha \in (0; 1)$ be constant,
and $\beta(y) = \alpha(a + by)$ be linear, where $a, a + b \in (0; 1]$.

Epidemic context: If $b > 0$,
prob. for susceptible becoming infected increases
if number of infectives already large (infection is spreading),
while recovery from infection independent of other infectives.

Properties:

$E[X_t \mid X_{t-1}] = (\alpha - \beta_t) X_{t-1} + N \beta_t$ quadratic in X_{t-1} ,

$V[X_t \mid X_{t-1}]$ cubic polynomial in X_{t-1} .

Explicit large- N approximation for marginal moments and ACF.

Self-Exciting Threshold Binomial $AR(1)$ Processes

(jointly with T.A. Möller, M.E. Silva, M.G. Scotto, I. Pereira)

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“Work in Progress”

Idea: Like in Monteiro et al. (2012), fix threshold value $0 \leq R < N$, which separates range into two regimes:
lower regime $\{0, 1, 2, \dots, R\}$, upper regime $\{R + 1, R + 2, \dots, N\}$.
Depending if previous observation in lower regime ($X_{t-1} \leq R$)
or in upper regime ($X_{t-1} > R$),
parameters of BAR(1) recursion at time t chosen differently.

Note: Possible generalizations include

- more than two regimes;
- increased delay $X_{t-d} \leq R$ vs. $X_{t-d} > R$ with $d > 1$;
- more complex criteria, e.g., $\max \{X_{t-1}, \dots, X_{t-d}\} \leq R$.

Definition: Define $\pi_i \in (0; 1)$, $\rho_i \in \left(\max \left\{ -\frac{\pi_i}{1-\pi_i}, -\frac{1-\pi_i}{\pi_i} \right\}; 1 \right)$, and $\beta_i := \pi_i \cdot (1 - \rho_i) \in (0; 1)$ and $\alpha_i := \beta_i + \rho_i \in (0; 1)$.

SET-BAR(1) process $X_t = \phi_t \circ X_{t-1} + \eta_t \circ (N - X_{t-1})$,

where $\phi_t := \alpha_1 I_{t-1} + \alpha_2 (1 - I_{t-1})$ and $\eta_t := \beta_1 I_{t-1} + \beta_2 (1 - I_{t-1})$ with indicator $I_{t-1} := \mathbb{1}_{\{X_{t-1} \leq R\}}$.

Note: If $R = 0$, then α_1 no influence (can be chosen arbitrarily).

$\Rightarrow \alpha_1$ unidentifiable during parameter estimation.

Same issue with β_2 and $R = N - 1$. Therefore, we set

$\rho_1 = \rho_2$ for $R = 0, N - 1$ ($\rightarrow$ LSET model from below).

Properties:

SET-BAR(1) model instance of

DD-BAR(1) models by Weiß & Pollett (2014)

⇒ adapt results concerning transition probabilities, ergodicity, existence of unique stationary marginal distribution p .

Since finite range, always possible to compute p numerically by solving eigenvalue problem $\mathbf{P} p = p$.

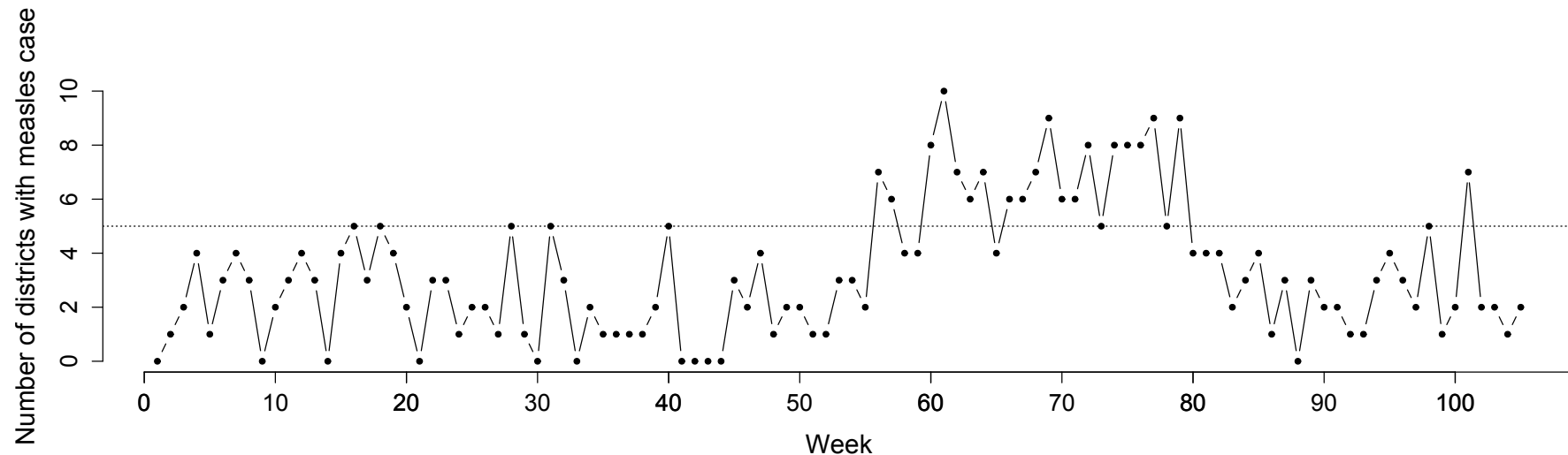
We derived closed-form formulae for mean and variance, but very complex.

Data example 3: (Robert-Koch-Institut)

Weekly No. of districts with new measles case (G., 2004/05).

Finite range $\{0, \dots, N\}$ with $N = 38$ (number of districts).

But ...



... level shift due to measles epidemic!

Above measles time series: level shift in first half of 2005, but no obvious change in serial dependence structure.

Idea: Reduce number of parameters by additional restriction $\rho_1 = \rho_2 =: \rho$.

Definition: Let $\rho \in \left(\max \left\{ -\frac{\pi_1}{1-\pi_1}, -\frac{\pi_2}{1-\pi_2}, -\frac{1-\pi_1}{\pi_1}, -\frac{1-\pi_2}{\pi_2} \right\}; 1 \right)$. A SET-BAR(1) process for which $\rho_1 = \rho_2 =: \rho \neq 0$ holds is called an **LSET-BAR(1) process**.

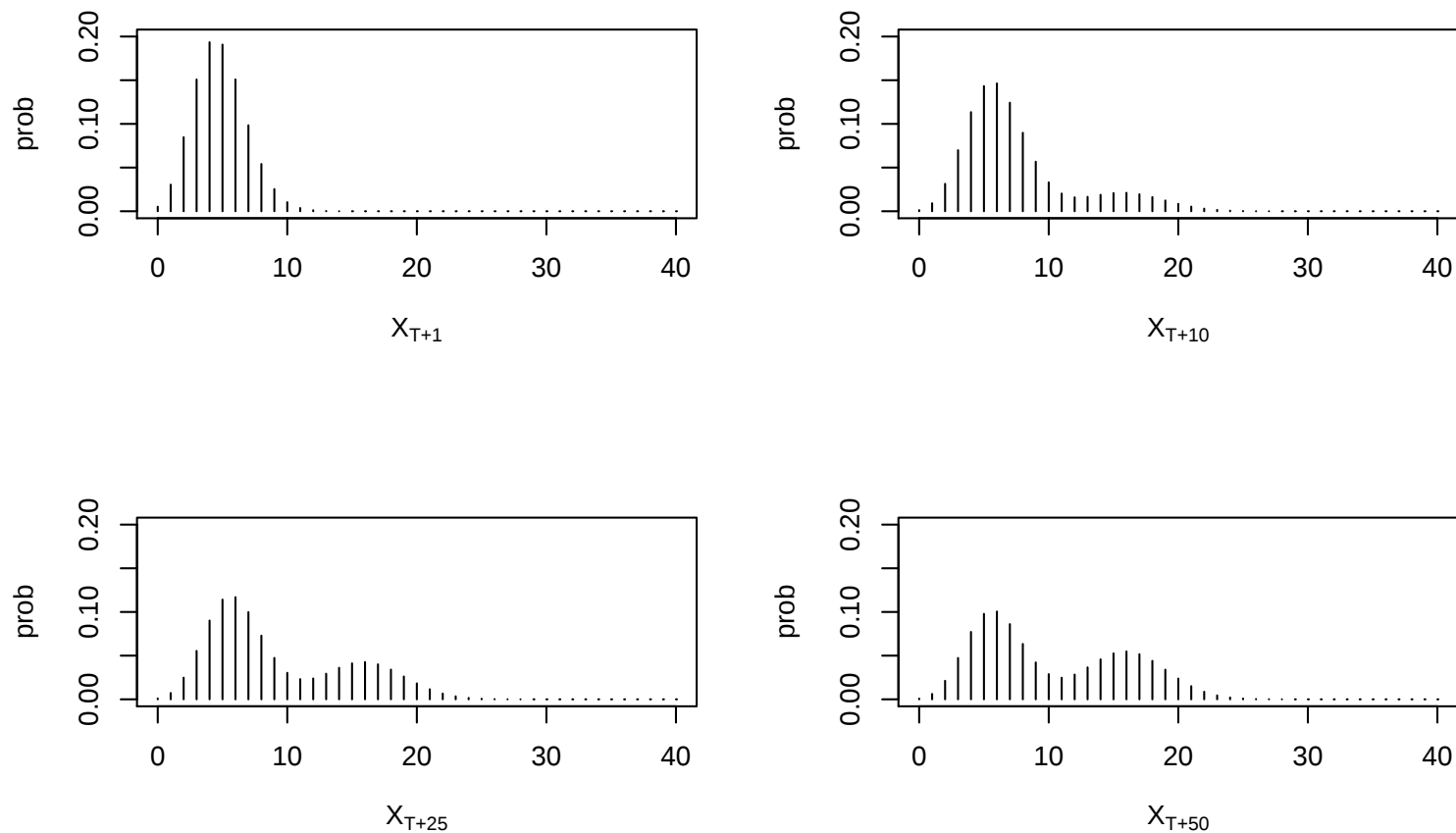
Defining $p := P(X_t \leq R) = E[I_{t-1}]$, $\mu_{IX} := E[I_{t-1}X_{t-1}]$,
we can express unconditional mean and variance as

$$\mu_X = Np\pi_1 + N(1-p)\pi_2,$$

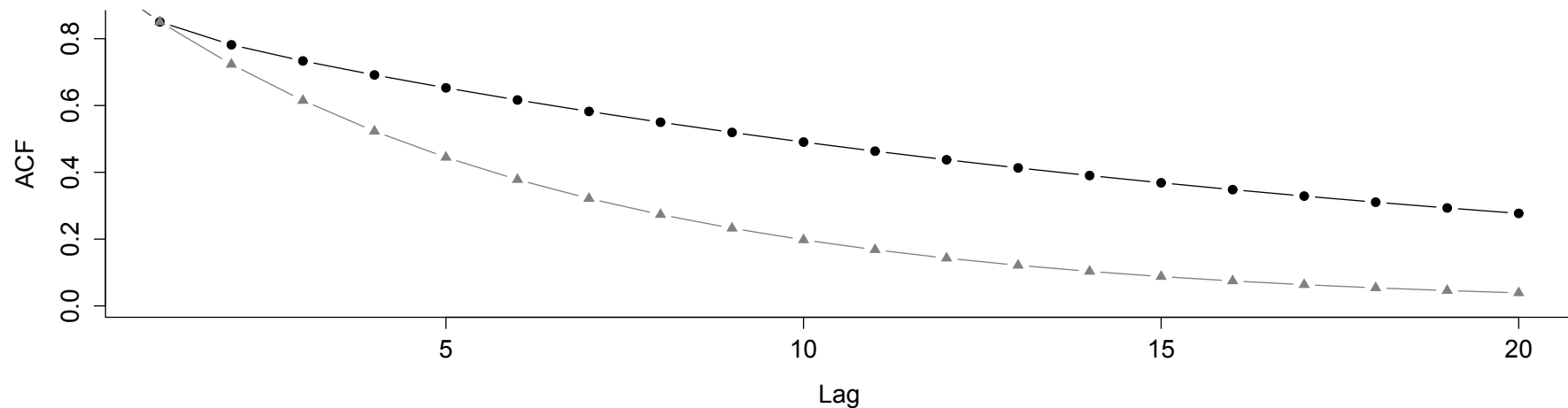
$$\begin{aligned}\sigma_X^2 &= Np\pi_1(1-\pi_1) + N(1-p)\pi_2(1-\pi_2) \\ &\quad + N^2p(1-p)(\pi_2-\pi_1)^2 \\ &\quad + \frac{2\rho}{1+\rho}(N-1)(\pi_2-\pi_1)(Np\pi_1 - \mu_{IX}).\end{aligned}$$

LSET model able to show over- and underdispersion
w.r.t. binomial distribution for appropriate parameter settings.

Forecasting distributions for model $N = 40$, $\rho = 0.3$,
 $\pi_1 = 0.15$, $\pi_2 = 0.4$ and $R = 10$, conditional on $X_T = 2$:



Autocorrelation function for model $N = 40$, $\rho = 0.3$, $\pi_1 = 0.15$, $\pi_2 = 0.4$ and $R = 10$ (black points):



Gray triangles show $f(k) := (\rho_X(1))^k$
 $\Rightarrow$ longer memory than corresponding AR(1)-like model.

Data example 3: measles counts, $N = 38$.

	Par. 1	Par. 2	Par. 3	Par. 4	AIC	BIC
BAR(1) (π, ρ)	0.0882 (0.0070)	0.4158 (0.0550)	-	-	448.2	453.5
DD-BAR(1) (a, b, ρ)	0.0419 (0.0095)	0.5270 (0.1077)	0 (0.1765)	-	436.9	444.9
BINARCH(1) (a, b)	0.0419 (0.0060)	0.5270 (0.0682)	-	-	434.9	440.2
SET-BAR(1) ($\pi_1, \pi_2, \rho_1, \rho_2$)	0.0706 (0.0056)	0.1558 (0.0269)	0.1916 (0.0884)	0.2904 (0.375)	432.5	443.1
LSET-BAR(1) (π_1, π_2, ρ)	0.0707 (0.0057)	0.1604 (0.0169)	0.1947 (0.0876)	-	430.6	438.5

All threshold models include threshold value $R = 5$.

- Time series of counts with finite range $\{0, 1, \dots, N\}$ important topic in practice.
- McKenzie's basic binomial AR(1) model easily generalized in several ways.
- Density-dependent binomial AR(1) model for binomial over- or underdispersion, boundary case of binomial INARCH(1) model.
- Model with density-dependent colonization.
- SET models for counts exhibiting piecewise-type patterns.

Thank You for Your Interest!



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