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Monitoring of Surveillance Time Series by Count Data Regression Charts

Michaela Paul[†] and Michael Höhle^{*‡}



[†]ISPM, University of Zurich, Switzerland, *Department of Statistics, University of Munich, Germany, [‡] Zentrum Mathematik, Technische Universität München, Germany email: michaela paul@ifspm.uzh.ch...hoehle@stat.uni-muenchen.de

email: michaela.paul@ifspm.uzh.ch, hoehle@stat.uni-muenchen.de

Abstract

We perform on-line change point detection in time series of counts arising from infectious disease surveillance using newly developed regression charts for the Poisson and negative binomial distribution. The in-control mean is assumed to be time-varying and linear on the logscale with intercept and seasonal components. Using the generalized likelihood ratio (GLR) statistic a monitoring scheme is formulated to detect on-line whether a shift in the intercept occurred. In case of Poisson the necessary quantities of the GLR detector can be efficiently computed by recursive formulas. Extensions to more general alternatives e.g. containing an autoregressive epidemic component are discussed. A comparison of the Poisson scheme with existing methods is done using Monte Carlo simulations. The practicability of the methods is demonstrated by applying them to the observed number of salmonella hadar cases in Germany.

3 Comparison of time-varying mean Poisson detectors

- The Poisson GLR-detector with multiplicative shift (1) is compared to
- the LR-detector,
- the modified Poisson CUSUM of Rogerson and Yamada (2004),
- the approximate Gaussian CUSUM of Rossi et al. (1999).
- All three methods need an advance specification of which change to detect optimally.
- We considered four scenarios with increasing mean and increasing seasonal amplitude of the monitored disease.



1 Introduction

Assume that the observations x_1, x_2, \ldots originate from some parametric distribution with density f_{θ} such that given the change-point τ the observations are independent realizations

$$x_t | z_t, \tau \sim \begin{cases} f_{\theta_0}(\cdot | z_t) \text{ for } t = 1, \dots, \tau - 1 \text{ (in-control)} \\ f_{\theta_1}(\cdot | z_t) \text{ for } t = \tau, \tau + 1, \dots \text{ (out-of-control)}. \end{cases}$$

- Our interest is to determine τ on-line i.e. new observations are collected until one is convinced that a change has occurred.
- We will assume that f is the negative binomial probability mass function with mean μ_t and a known dispersion parameter.
- A stopping rule is used to determine when enough evidence against $H_0: \mu_t = \mu_{0,t}, t = 1, \dots$ has been collected to stop the sampling.

- The respective thresholds for all methods were selected to achieve an in-control average run length (ARL_0) of 500.
- The obtained ARL_0 using the method of Rossi et al. (1999) are much lower than the anticipated values whereas the other methods are doing a good job in obtaining the desired ARL_0 .
- We estimated out-of-control ARLs for selected shift sizes δ . The ARL_1 for the methods with fixed alternative hypothesis are sensitive to the choice of the shift size Δ . The GLR detector performs best when the actual shift size δ is substantially lower.



Monte-Carlo estimated ARLs (left) and corresponding false alarm probability (right) as a function of c_{γ} for the negative binomial intercept chart with $\hat{\mu}_{0,t}$ estimated from the hadar data. Also shown are point-wise 95% confidence regions.

Analysis of shadar using glrnb: intercept



GLR(n) statistic superimposed on the observed data of 2005-2006. The triangles show all n, where $GLR(n) \ge c_{\gamma}$.

5 Software

The out-of-control situation is characterized by a multiplicative shift

$$\mu_{1,t} = \mu_{0,t} \cdot \exp(\kappa)$$

with $\log(\mu_{0,t})$ specified by cyclic regression.

More involved models for the shift are imaginable, one suggestion is

 $\mu_{1,t}^e = \mu_{0,t} + \lambda x_{t-1}, \quad t > 1,$

(1)

(2)

where $\lambda > 0$ and $\mu_{1,1}^e = \mu_{0,1}$. This auto-regressive model is discussed in Paul et al. (2008) as a model for time series of counts from infectious diseases.

> 2 The Generalized Likelihood Ratio (GLR) detector



 $GLR(n) = \max_{1 \le k \le n} \sup_{\theta \in \Theta} \left[\sum_{t=k}^{n} \log \left\{ \frac{f_{\theta}(x_t | z_t)}{f_{\theta_0}(x_t | z_t)} \right\} \right]$

with stopping rule $N_G = \inf \{n \ge 1 : GLR(n) \ge c_{\gamma} \}.$

• Maximization of the log-likelihood has to be carried out over $\theta \in \Theta$

Monte Carlo estimated out-of-control ARLs for selected values of δ when $X_t \sim \text{Po}(\delta \mu_{0,t})$ and with the charts designed to detect a multiplicative shift of size $\Delta = 2$.

4 Application to German salmonella data

- During the year 2006 the German health authorities noted an increased number of cases due to *salmonella hadar* compared to the previous years.
- We fitted a log-linear negative binomial GLM to the data of 2001 2004 and used this model to predict the expected number of cases $\hat{\mu}_{0,t}$ for the remaining 2 years.
- A window-limited negative binomial intercept chart with $\hat{\mu}_{0,t}$ as in-control mean is applied to the data of 2005 2006.

The presented charts are implemented in the R package **surveillance** available from CRAN. Development versions are available from

http://surveillance.r-forge.r-project.org/

Furthermore, the package provides visualization of routinely collected surveillance data and a test-bench for new surveillance algorithms. Among others **surveillance** contains an implementation of

• the procedure of Farrington et al. (1996),

• the GLR detectors proposed by Höhle and Paul (2008),

• estimation routines for the model described in Paul et al. (2008).

Further information about the package can be obtained from Höhle (2007) or in the accompanying vignette.

6 Acknowledgements

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for each possible change time k between 1 and n.

• Höhle and Paul (2008) show that this computational complexity can be reduced for the multiplicative shift (1) Poisson chart by clever recursive computations of the sums and sups.

• In the more general negative binomial setup or in setups with more flexible out-of-control models such as (2) this reduction is not possible.

• In case of large n an alternative is the so-called *window-limited* GLR scheme where maximization is only performed for a moving window of k values.

• Assuming a fixed out-of-control mean instead of computing the sup over all $\theta \in \Theta$ results in the LR detector.

• To investigate run-length properties of the GLR or LR scheme having time-changing means, Monte-Carlo sampling is necessary. • Direct Monte Carlo simulation was used to obtain a threshold c_{γ} which results in a target ARL of $\gamma = 500$.



Weekly number of *salmonella hadar* cases in Germany taken from the SurvStat@RKI database. The dotted line shows the predicted number of cases $\hat{\mu}_{0,t}$ based on the 2001-2004 data. Höhle, M. (2007). surveillance: An R package for the surveillance of infectious diseases, *Comput. Stat.* **22**(4): 571–582.

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