# Web-based Supplementary Materials for "A space-time conditional intensity model for invasive meningococcal disease occurence"

by

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### Web Appendix A: Calculus of the Score Function

Let  $\boldsymbol{\vartheta}$  denote any subvector of  $\boldsymbol{\theta}$ . Then, the partial derivative of the log-likelihood with respect to  $\boldsymbol{\vartheta}$  is

$$\boldsymbol{s}_{\boldsymbol{\vartheta}}(\boldsymbol{\theta}) := \frac{\partial}{\partial \boldsymbol{\vartheta}} l(\boldsymbol{\theta}) = \sum_{i=1}^{n} \frac{\frac{\partial}{\partial \boldsymbol{\vartheta}} \lambda_{\boldsymbol{\theta}}^{*}(t_{i}, \boldsymbol{s}_{i}, \kappa_{i})}{\lambda_{\boldsymbol{\theta}}^{*}(t_{i}, \boldsymbol{s}_{i}, \kappa_{i})} - \int_{0}^{T} \int_{W} \sum_{\kappa \in \mathcal{K}} \frac{\partial}{\partial \boldsymbol{\vartheta}} \lambda_{\boldsymbol{\theta}}^{*}(t, \boldsymbol{s}, \kappa) \, \mathrm{d}t \, \mathrm{d}\boldsymbol{s} , \qquad (1)$$

and the score function is  $s(\theta) = \frac{\partial}{\partial \theta} l(\theta) = \left(s'_{\beta_0}, s'_{\beta}, s'_{\gamma}, s'_{\sigma}, s'_{\alpha}\right)'(\theta)$ . The necessary partial derivatives of the CIF with their respective time-space-mark integrals are given in the following subsections and can be plugged into the equation (1). The analytic derivatives of the interaction function f and g with respect to  $\sigma$  and  $\alpha$ , respectively, have to be determined for the specific model at hand. For instance, a type-specific spatial Gaussian kernel

$$f_{\sigma}(\boldsymbol{s}|\kappa) = \exp\left(-\frac{\|\boldsymbol{s}\|^2}{2\sigma_{\kappa}^2}\right)$$

with  $\boldsymbol{\sigma} = (\sigma_1, \dots, \sigma_K)'$  has partial derivatives

$$\frac{\partial}{\partial \sigma_k} f_{\sigma}(\boldsymbol{s}|\kappa) = \mathbb{1}_{k=\kappa}(\kappa) \cdot \exp\left(-\frac{\|\boldsymbol{s}\|^2}{2\sigma_k^2}\right) \frac{\|\boldsymbol{s}\|^2}{\sigma_k^3}, \quad \text{for any } k \in \mathcal{K}.$$

The type-specific temporal interaction function  $g_{\alpha}(t|\kappa) = e^{-\alpha_{\kappa}t}$  with  $\alpha = (\alpha_1, \dots, \alpha_K)$  has partial derivatives  $\frac{\partial}{\partial \alpha_k} g_{\alpha}(t|\kappa) = \mathbbm{1}_{k=\kappa}(\kappa) \cdot (-t e^{-\alpha_k t})$ , for any  $k \in \mathcal{K}$ . While the integral of  $\frac{\partial}{\partial \sigma_{\kappa}} f_{\sigma}(s|\kappa)$  over the region  $R_j$  will be approximated by numerical integration, the temporal function  $\frac{\partial}{\partial \alpha_{\kappa}} g_{\alpha}(t|\kappa)$  is assumed to permit analytical integration.

#### Endemic intercept(s) $\beta_0$ :

Let  $\beta_{0,k}$ ,  $k \in \{1, ..., K\}$  be one of the type-specific intercepts in  $\beta_0$ . Then,

$$\frac{\partial}{\partial \beta_{0,k}} \lambda_{\boldsymbol{\theta}}^*(t, \boldsymbol{s}, \kappa) = \mathbb{1}_{k=\kappa}(\kappa) \cdot \exp\left(\beta_{0,k} + o_{\xi(\boldsymbol{s})} + \boldsymbol{\beta}' \boldsymbol{z}_{\tau(t), \xi(\boldsymbol{s})}\right)$$

since the parameter  $\beta_{0,k}$  appears in the endemic component  $h_{\theta}(t, \mathbf{s}, \kappa)$  if and only if  $\kappa = k$ . The corresponding integrated value is

$$\int_0^T \int_W \sum_{\kappa \in \mathcal{K}} \frac{\partial}{\partial \beta_{0,k}} \lambda_{\boldsymbol{\theta}}^*(t, \boldsymbol{s}, \kappa) \, \mathrm{d}t \, \mathrm{d}\boldsymbol{s} = e^{\beta_{0,k}} \cdot \sum_{\tau=1}^D \sum_{\xi=1}^M |C_{\tau}| |A_{\xi}| \exp\left(o_{\xi} + \boldsymbol{\beta}' \boldsymbol{z}_{\tau, \xi}\right) ,$$

cf. the integral of the endemic component in equation (6) of the paper. If the model assumes a type-invariant endemic intercept  $\beta_0 = \beta_0$ , then

$$\frac{\partial}{\partial \beta_0} \lambda_{\boldsymbol{\theta}}^*(t, \boldsymbol{s}, \kappa) = \exp\left(\beta_0 + o_{\xi(\boldsymbol{s})} + \boldsymbol{\beta}' \boldsymbol{z}_{\tau(t), \xi(\boldsymbol{s})}\right)$$

with integrated value

$$\int_0^T \int_W \sum_{\kappa \in \mathcal{K}} \frac{\partial}{\partial \beta_0} \lambda_{\boldsymbol{\theta}}^*(t, \boldsymbol{s}, \kappa) \, \mathrm{d}t \, \mathrm{d}\boldsymbol{s} = K \, e^{\beta_0} \cdot \sum_{\tau=1}^D \sum_{\xi=1}^M |C_\tau| |A_\xi| \exp\left(o_\xi + \boldsymbol{\beta}' \boldsymbol{z}_{\tau, \xi}\right) \, .$$

#### Endemic covariate effects $\beta$ :

$$\frac{\partial}{\partial \boldsymbol{\beta}} \lambda_{\boldsymbol{\theta}}^*(t, \boldsymbol{s}, \kappa) = \exp\left(\beta_0^{(\kappa)} + o_{\xi(s)} + \boldsymbol{\beta}' \boldsymbol{z}_{\tau(t), \xi(s)}\right) \cdot \boldsymbol{z}_{\tau(t), \xi(s)}$$

with corresponding integral vector (element-wise integral values)

$$\left(\sum_{\kappa \in \mathcal{K}} \exp\left(\beta_0(\kappa)\right)\right) \cdot \sum_{\tau=1}^{D} \sum_{\xi=1}^{M} |C_{\tau}| |A_{\xi}| \exp(o_{\xi} + \boldsymbol{\beta}' \boldsymbol{z}_{\tau,\xi}) \boldsymbol{z}_{\tau,\xi}.$$

#### Epidemic effects $\gamma$ :

$$\frac{\partial}{\partial \boldsymbol{\gamma}} \lambda_{\boldsymbol{\theta}}^*(t, \boldsymbol{s}, \kappa) = \sum_{j \in I^*(t, \boldsymbol{s}, \kappa)} e^{\boldsymbol{\gamma}' \boldsymbol{m}_j} g_{\boldsymbol{\alpha}}(t - t_j | \kappa_j) f_{\boldsymbol{\sigma}}(\boldsymbol{s} - \boldsymbol{s}_j | \kappa_j) \boldsymbol{m}_j,$$

and the corresponding integral can be deduced similar to equation (7) of the paper as

$$\sum_{j=1}^{n} q_{\kappa_j,\bullet} e^{\gamma' m_j} \left[ \int_0^{\min\{T-t_j;\varepsilon\}} g_{\alpha}(t|\kappa_j) dt \right] \left[ \int_{R_j} f_{\sigma}(\boldsymbol{s}|\kappa_j) d\boldsymbol{s} \right] \boldsymbol{m}_j.$$

#### Parameters $\sigma$ and $\alpha$ of the interaction functions:

For a general spatial kernel  $f_{\sigma}(s|\kappa)$ ,

$$\frac{\partial}{\partial \boldsymbol{\sigma}} \lambda_{\boldsymbol{\theta}}^*(t, \boldsymbol{s}, \kappa) = \sum_{j \in I^*(t, \boldsymbol{s}, \kappa)} e^{\eta_j} g_{\boldsymbol{\alpha}}(t - t_j | \kappa_j) \left[ \frac{\partial}{\partial \boldsymbol{\sigma}} f_{\boldsymbol{\sigma}}(\boldsymbol{s} - \boldsymbol{s}_j | \kappa_j) \right]$$

with corresponding integral

$$\sum_{j=1}^{n} q_{\kappa_{j},\bullet} e^{\eta_{j}} \left[ \int_{0}^{\min\{T-t_{j};\varepsilon\}} g_{\alpha}(t|\kappa_{j}) dt \right] \left[ \int_{R_{j}} \frac{\partial}{\partial \boldsymbol{\sigma}} f_{\boldsymbol{\sigma}}(\boldsymbol{s}|\kappa_{j}) d\boldsymbol{s} \right].$$

Similarly, for a general temporal kernel  $g_{\alpha}(t|\kappa)$ ,

$$\frac{\partial}{\partial \boldsymbol{\alpha}} \lambda_{\boldsymbol{\theta}}^*(t, \boldsymbol{s}, \kappa) = \sum_{j \in I^*(t, \boldsymbol{s}, \kappa)} e^{\eta_j} \left[ \frac{\partial}{\partial \boldsymbol{\alpha}} g_{\boldsymbol{\alpha}}(t - t_j | \kappa_j) \right] f_{\boldsymbol{\sigma}}(\boldsymbol{s} - \boldsymbol{s}_j | \kappa_j)$$

with corresponding integral

$$\sum_{j=1}^{n} q_{\kappa_{j},\bullet} e^{\eta_{j}} \left[ \int_{0}^{\min\{T-t_{j};\varepsilon\}} \frac{\partial}{\partial \boldsymbol{\alpha}} g_{\boldsymbol{\alpha}}(t|\kappa_{j}) dt \right] \left[ \int_{R_{j}} f_{\boldsymbol{\sigma}}(\boldsymbol{s}|\kappa_{j}) d\boldsymbol{s} \right].$$

# Web Appendix B: Fisher Information Matrix

The inverse of the Fisher information matrix at the maximum likelihood estimate (MLE)  $\hat{\boldsymbol{\theta}}_{ML}$  is in general likelihood theory used as an estimate of the variance of  $\hat{\boldsymbol{\theta}}_{ML}$ . The precise conditions under which asymptotic properties of MLEs hold for spatio-temporal point processes have been established by Rathbun (1996). Specifically, the conditions for existence, consistence and asymptotic normality of a local maximum  $\hat{\boldsymbol{\theta}}_{ML}$  as  $T \to \infty$  for a fixed observation region W are discussed in Meyer (2009, Section 4.2.3).

The expected Fisher information  $\mathcal{I}(\boldsymbol{\theta})$  can be estimated by the "optional variation process" adapted to the marked spatio-temporal setting (Rathbun, 1996, equation (4.7))

$$\int_0^T \int_W \int_{\mathcal{K}} \left( \frac{\partial}{\partial \boldsymbol{\theta}} \log \lambda_{\boldsymbol{\theta}}^*(t, \boldsymbol{s}, \kappa) \right)^{\otimes 2} \, \mathrm{d}N(t, \boldsymbol{s}, \kappa)$$

through its observed realisation

$$\hat{\mathcal{I}}(\boldsymbol{\theta}) = \sum_{i=1}^{n} \left( \frac{\partial}{\partial \tilde{\boldsymbol{\theta}}} \log \lambda_{\tilde{\boldsymbol{\theta}}}^*(t_i, \boldsymbol{s}_i, \kappa_i) \bigg|_{\tilde{\boldsymbol{\theta}} = \boldsymbol{\theta}} \right)^{\otimes 2} = \sum_{i=1}^{n} \left( \frac{\frac{\partial}{\partial \tilde{\boldsymbol{\theta}}} \lambda_{\tilde{\boldsymbol{\theta}}}^*(t_i, \boldsymbol{s}_i, \kappa_i)}{\lambda_{\tilde{\boldsymbol{\theta}}}^*(t_i, \boldsymbol{s}_i, \kappa_i)} \bigg|_{\tilde{\boldsymbol{\theta}} = \boldsymbol{\theta}} \right)^{\otimes 2} ,$$

where  $\mathbf{a}^{\otimes 2} := \mathbf{a}\mathbf{a}'$  for a vector  $\mathbf{a}$ . Uncertainty of the parameter estimates is thus deduced from the diagonal of  $\hat{\mathcal{I}}^{-1/2}(\hat{\boldsymbol{\theta}}_{ML})$ , which contains their standard errors.

#### References

Meyer, S. (2009). Spatio-temporal infectious disease epidemiology based on point processes. Master's thesis, Department of Statistics, Ludwig-Maximilians-Universität, München. Available as http://epub.ub.uni-muenchen.de/11703/.

Rathbun, S. L. (1996). Asymptotic properties of the maximum likelihood estimator for spatiotemporal point processes. *Journal of Statistical Planning and Inference* **51**, 55–74.

# Web Appendix C: Simulation Algorithm

This appendix provides a more implementational view on the simulation algorithm described in Section 4 of the paper. In addition to the notation of the paper, let L(t) be the next time point after time t where any endemic covariate in any tile changes its value, or a previously infected individual stops spreading the disease, i.e.:

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L(t) = \min \{t' > t \mid (\exists \xi \in \{1, \dots, M\} : \mathbf{z}_{\tau(t'), \xi} \neq \mathbf{z}_{\tau(t), \xi}) \lor (\exists j \in \{1, \dots, N_q(t)\} : t' = t_j + \varepsilon)\}.
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An implementational perspective on the simulation algorithm is then:

#### Algorithm 1: Ogata's modified thinning adapted for twinstim

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1 Given current time t, update L(t) and calculate local upper bound \overline{\lambda_a^*}(t);
 2 Generate proposed waiting time \Delta \sim \operatorname{Exp}(\overline{\lambda_a^*}(t));
 3 if t + \Delta > L(t) then
        Let t = L(t);
 5 else
 6
         Let t = t + \Delta;
         Accept t with probability \lambda_q^*(t)/\lambda_q^*(t), otherwise goto step 1;
 7
         Draw the source of infection (infective individual j or endemic) with weights equal to the
 8
         respective components of \lambda_a^*(t) (cf. equation (8) of the paper);
        if endemic source of infection then
 9
              Draw the type \kappa of the new event with weights \exp(\beta_0(\kappa)), \kappa \in \mathcal{K};
10
              Draw the tile A_{\xi} of the new event with weights |A_{\xi}| \rho_{\tau(t),\xi} e^{\beta' z_{\tau(t),\xi}}, \xi \in \{1,\ldots,M\};
11
              Draw the location s of the new event uniform within the sampled tile A_{\xi};
12
        else
13
              Draw the type \kappa of the new event at random out of the types which can be triggered by
14
             the source individual j, i.e. draw from U(\{\kappa \in \mathcal{K} : q_{\kappa_j,\kappa} = 1\});
              Draw the relative location oldsymbol{v} of the new event (relative to the source j) from the density
15
             f(s|\kappa_j)/\int_{R_i} f(s|\kappa_j) \,\mathrm{d}s on R_j, i.e. s=s_j+v;
         Draw additional marks according to the pre-specified distribution;
16
        Update the event history;
17
   Goto step 1;
18
```

An R implementation of the algorithm can be found as the function simEpidataCS in the popular surveillance package.