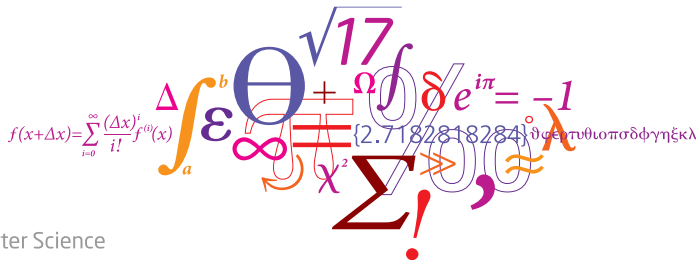


Automated and Early Detection of Disease Outbreaks

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DTU Compute

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Outline

- Data
 - Focusing on STEC
- Novel outbreak detection algorithm
- Model formulas
 - Agegroup
 - Agegroup and seasonality
- Hierarchical Poisson Normal model
 - Implementation
 - Results
- Hierarchical Poisson Gamma model
 - Implementation
 - Results

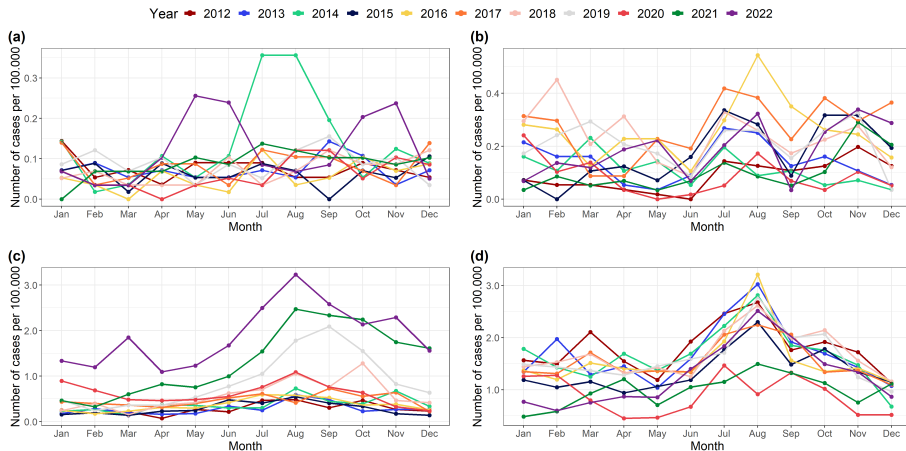
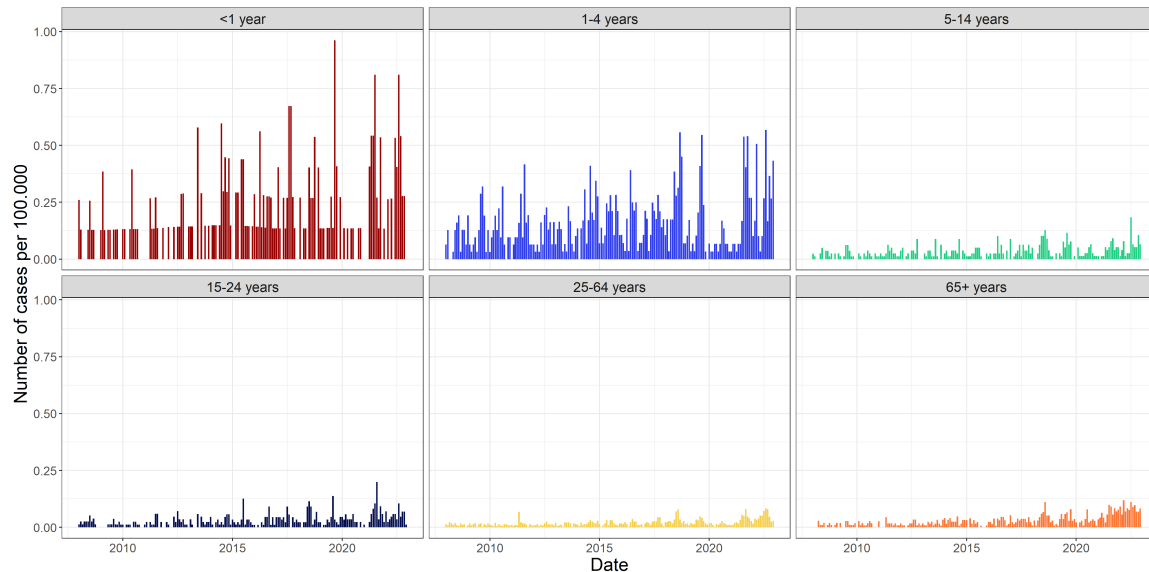


Figure: Epidemic curve showing the incidence per 100,000 in Denmark, 2012-2022, for the subset of diseases considered in this master thesis. (a) Listeriosis, (b) Shigellosis, (c) STEC, and (d) Salmonellosis.

Focusing on STEC



Algorithm 1: Test

Data: This text

Result: Hello

for $i = 1$ **to** T **do**

 | Do something

end

$$\log(\lambda_{it}) = \mathbf{x}_{it}^T \boldsymbol{\beta} + \log(n_{it}), \quad i = 1, \dots, k, \quad t = 1, \dots, T \quad (1)$$

$$\log(\lambda_{it}) = \beta(\text{ageGroup}_i) + \log(n_{it}) \quad (2)$$

$$\log(\lambda_{it}) = \beta(\text{ageGroup}_i)\beta_i + \sin\left(\frac{\pi \cdot \text{monthInYear}_t}{6}\right)\beta_{\sin} + \cos\left(\frac{\pi \cdot \text{monthInYear}_t}{6}\right)\beta_{\cos} + \log(n_{it})$$

(3)

Hierarchical Poisson Normal model

$$\mathbf{Y}|\mathbf{u} \sim \text{Pois}(\boldsymbol{\lambda} \exp(\mathbf{u})) \quad (4a)$$

$$\mathbf{u} \sim \text{N}(\mathbf{0}, I\sigma^2) \quad (4b)$$

Hierarchical Poisson Normal model

Implementation

```
#include <TMB.hpp>           // Links in the TMB libraries

template<class Type>
Type objective_function<Type>::operator() ()
{
    DATA_VECTOR(y);          // Data vector transmitted from R
    DATA_VECTOR(x);          // Data vector transmitted from R
    DATA_MATRIX(X);          // Design matrix transmitted from R

    PARAMETER_VECTOR(u);      // Random effects

    // Parameters
    PARAMETER_VECTOR(beta);    // Parameter value transmitted from R
    PARAMETER(log_sigma_u);    // Parameter value transmitted from R

    vector<Type> lambda = exp(X*beta-log(x)+u);
    Type sigma_u = exp(log_sigma_u);

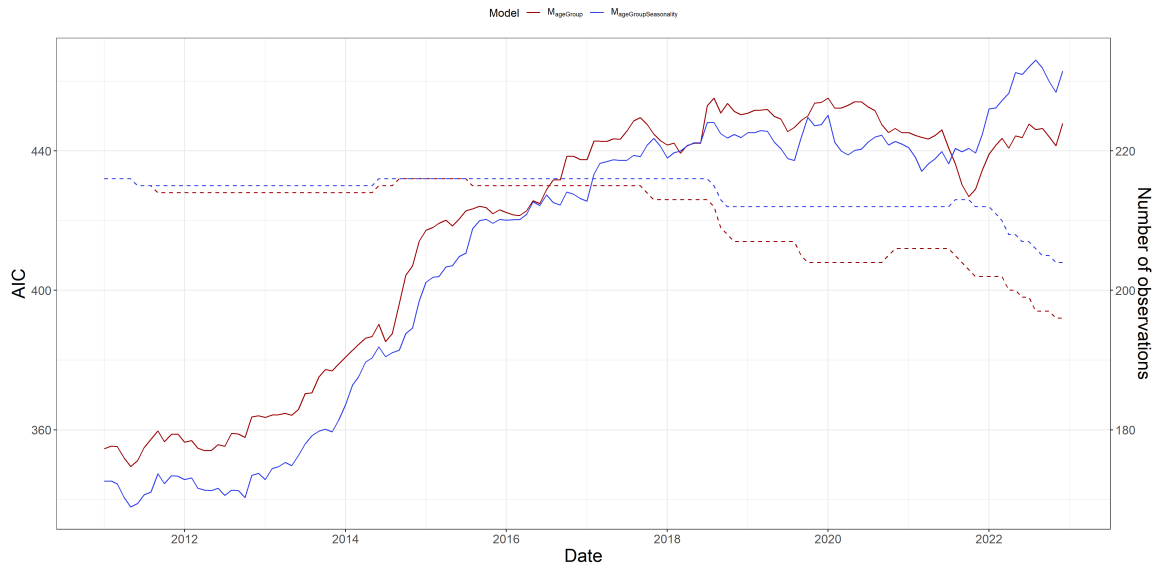
    int nobs = y.size();
    Type mean_ran = Type(0);

    Type f = 0;               // Declare the "objective function"
    for(int t=0; t < nobs; t++){
        f -= dnorm(u[t],mean_ran,sigma_u,true);
        f -= dpois(y[t],lambda[t],true);
    }

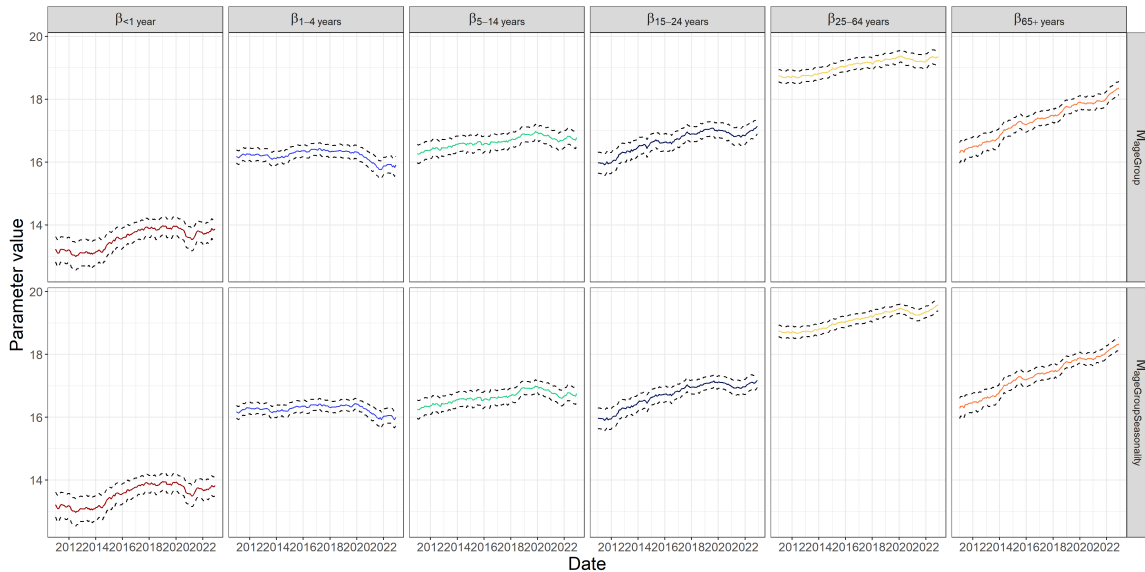
    return f;
}
```

Hierarchical Poisson Normal model

Model performance

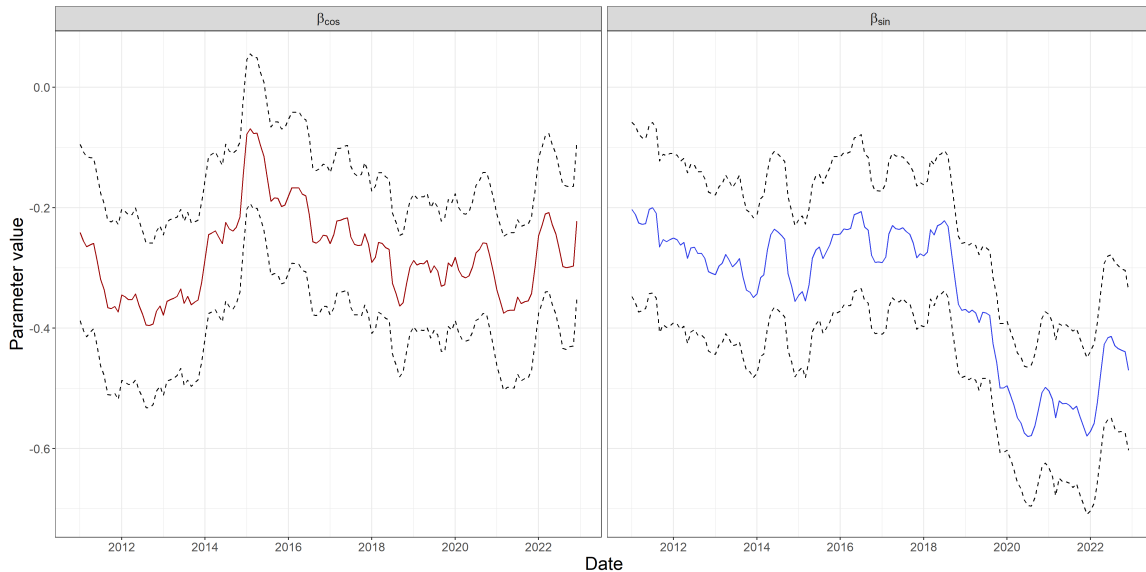


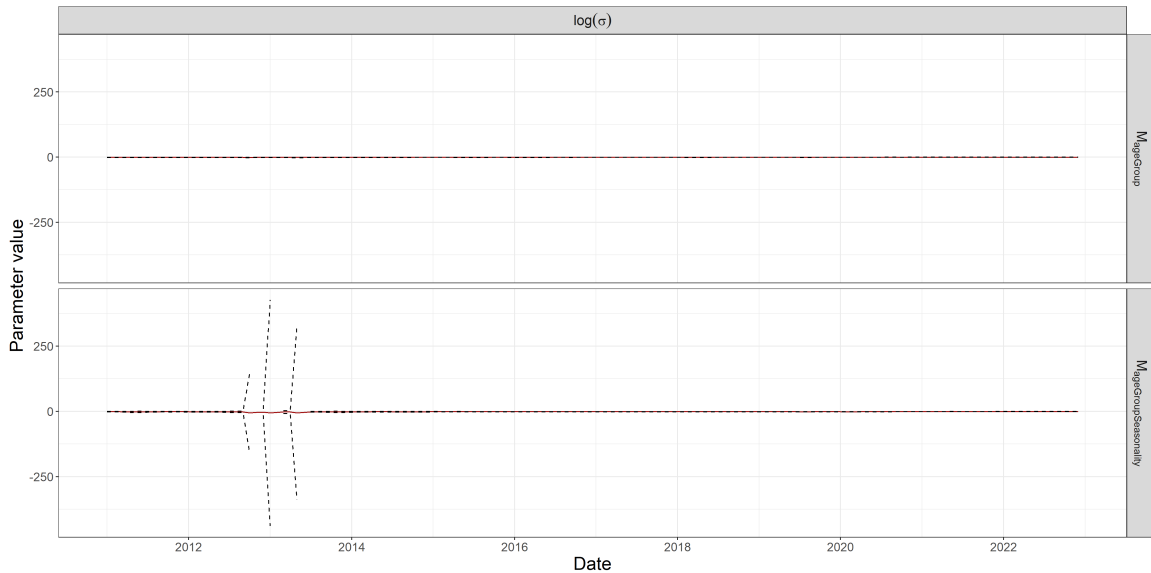
Agegroup parameters



Hierarchical Poisson Normal model

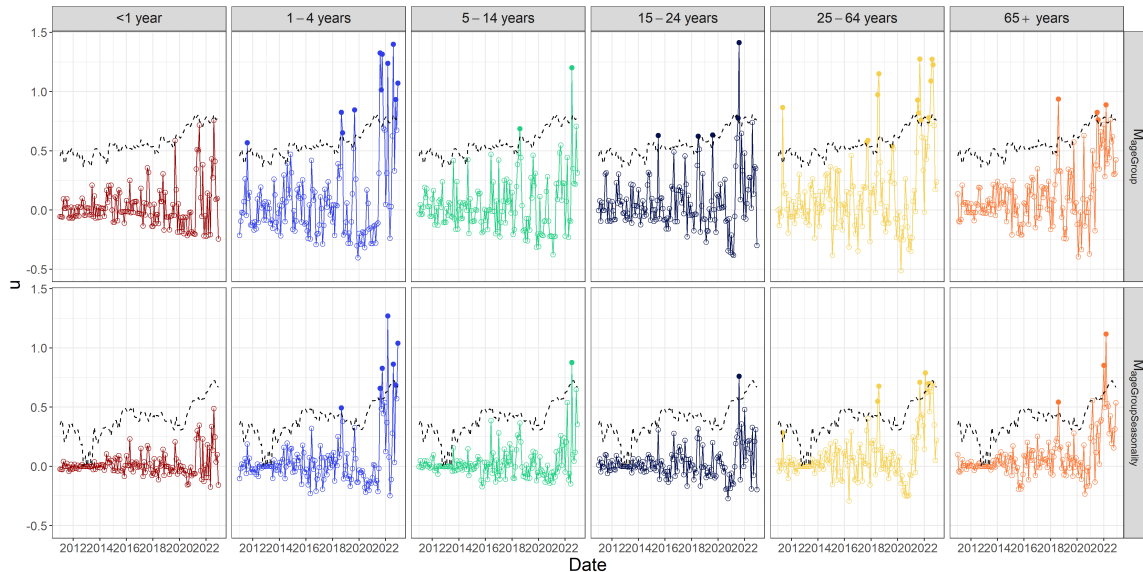
Seasonality parameters



Variance parameter

Hierarchical Poisson Normal model

Outbreak detection



Hierarchical Poisson Gamma model

$$\mathbf{Y}|\mathbf{u} \sim \text{Pois}(\lambda\mathbf{u}) \quad (5a)$$

$$\mathbf{u} \sim \text{G}(\mathbf{1}/\phi, \phi) \quad (5b)$$

Hierarchical Poisson Gamma model Implementation

```
#include <TMB.hpp>           // Links in the TMB libraries

template<class Type>
Type objective_function<Type>::operator() ()
{
    DATA_VECTOR(y);           // Data vector transmitted from R
    DATA_VECTOR(x);           // Data vector transmitted from R
    DATA_MATRIX(X);           // Design matrix transmitted from R

    // Parameters
    PARAMETER_VECTOR(beta);     // Parameter value transmitted from R
    PARAMETER(log_phi_u);       // Parameter value transmitted from R

    vector<Type> lambda = exp(X*beta-log(x)); // Construct the model parameters
    Type phi_u = exp(log_phi_u); // ... and the model parameters

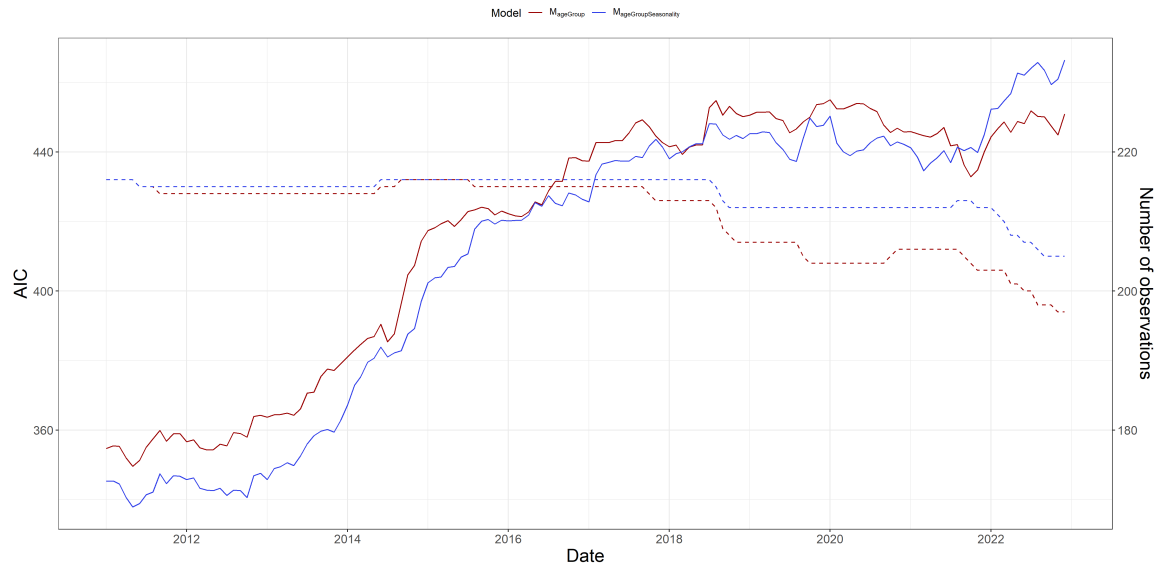
    Type r = 1/phi_u; // Construct the size
    vector<Type> p = 1/(lambda*phi_u+1); // ... and the probability parameter

    Type f = -sum(dnbinom(y, r, p,true)); // Calculate the "objective function"

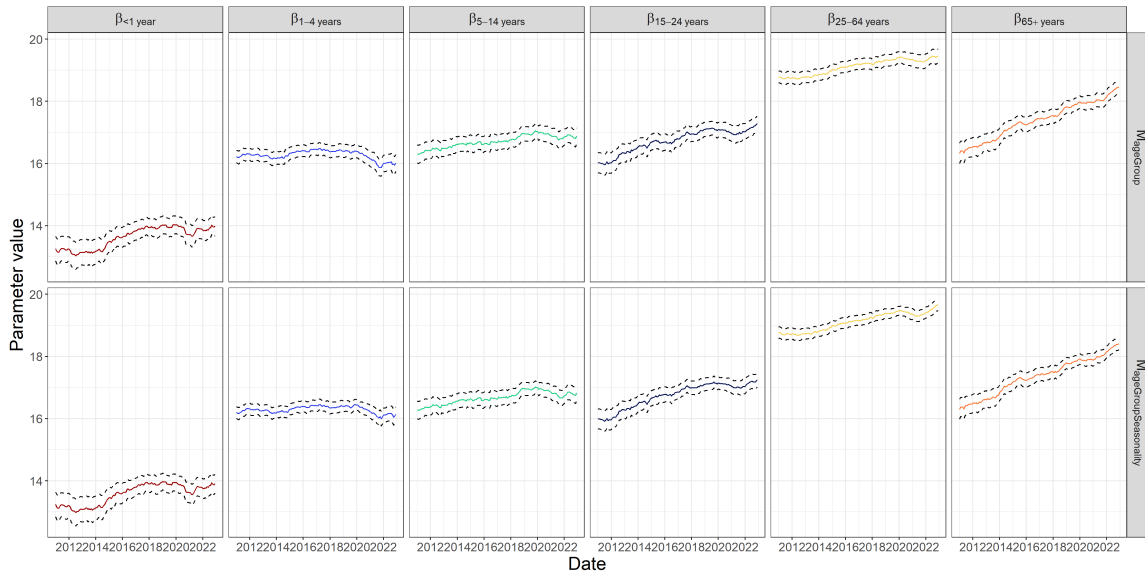
    return f;
}
```

Hierarchical Poisson Gamma model

Model performance



Agegroup parameters

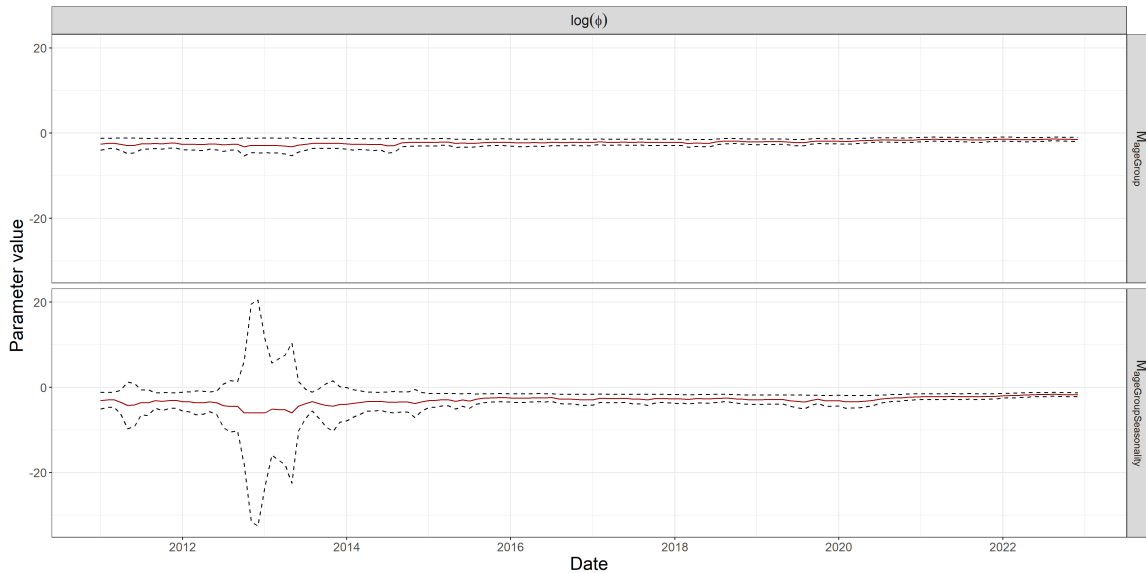


Hierarchical Poisson Gamma model

Seasonality parameters



Variance parameter



Hierarchical Poisson Gamma model

Outbreak detection

