

Models

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DTU Compute

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Outline



- Data exploration
 - VTEC / STEC
- Model formulation
 - Poisson-Lognormal
 - Poisson-Gamma

Data exploration VTEC / STEC



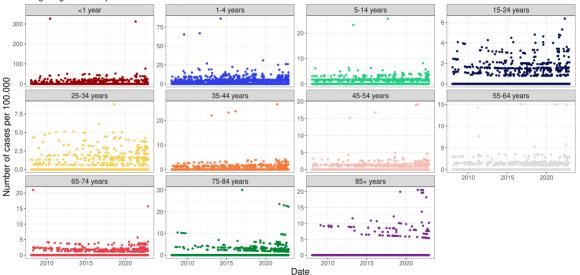
Date	ageGroup	у
2008-01-01 2008-01-01	<1 year 1-4 years	2
2008-01-01	5-14 years	2
2008-01-01	15-24 years	1
2022-12-01	55-64 years	2
2022-12-01	65-74 years	5
2022-12-01	75-84 years	4
2022-12-01	85+ years	3

Data exploration

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VTEC / STEC

Shiga- og veratoxin producerende E. coli.



Poisson-Lognormal



$$Y_i \sim \operatorname{Pois}(\lambda_i \exp(u_i))$$
 (1a)

$$u_i \sim N(0, \sigma^2)$$
 (1b)



Implementation - Objective function in C++

```
#include <TMB.hpp>
                               // Links in the TMB libraries
template<class Type>
Type objective function Type :: operator() ()
 DATA VECTOR(v):
                                      // Data vector transmitted from R
 DATA_FACTOR(ageGroup); // Data factor transmitted from R
 PARAMETER VECTOR(u):
                                   // Random effects
 // Parameters
 PARAMETER VECTOR(lambda):
                             // Parameter value transmitted from R
 PARAMETER(log_sigma_u);
                                      // Parameter value transmitted from R
 Type sigma_u = exp(log_sigma_u);
 int nobs = v.size();
 Type mean ran = Type(0):
 int j;
 Type f = 0;
                          // Declare the "objective function" (neg. log. likelihood)
 for(int i=0; i < nobs; i++){
   f -= dnorm(u[i],mean_ran,sigma_u,true);
   j = ageGroup[i];
   f -= dpois(v[i],lambda[j]*exp(u[i]),true);
 return f;
```

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Implementation - Call from R

```
# Import libraries
library(readr)
library(dplvr)
library(TMB)
# Import the data
dat <- read rds(file = "../../data/processed/dat.rds")</pre>
# Only consider some of the data
v <- dat %>%
  filter(caseDef == "Shiga- og veratoxin producerende E. coli.") %>%
  group_by(Date, ageGroup) %>%
  summarize(v = sum(cases))
compile(file = "PoissonLognormal.cpp") # Compile the C++ file
dvn.load(dvnlib("PoissonLognormal")) # Dunamically link the C++ code
# Function and derivative
PoisLN <- MakeADFun(
 data = list(y = y$y, ageGroup = y$ageGroup),
 parameters = list(u = rep(1, length(y$y)),
                    lambda = rep(1, nlevels(y$ageGroup)),
                    log_sigma_u = log(1)),
  random = "u",
  DLL = "PoissonLognormal"
opt <- nlminb(start = PoisLN$par, PoisLN$fn, PoisLN$gr, lower = c(0.01, 0.01))
```

Results

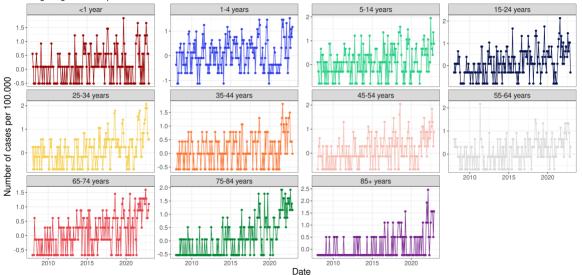


Parameter	Estimate	Std. Error
$\lambda_{<1year}$	0.83	0.09
$\lambda_{1-4years}$	3.39	0.29
$\lambda_{5-14 years}$	1.73	0.16
$\lambda_{15-24years}$	1.78	0.17
$\lambda_{25-34years}$	1.38	0.14
$\lambda_{35-44years}$	1.02	0.11
$\lambda_{45-54years}$	1.29	0.13
$\lambda_{55-64 years}$	1.25	0.12
$\lambda_{65-74 years}$	1.31	0.13
$\lambda_{75-84 years}$	0.90	0.10
$\lambda_{85+years}$	0.31	0.04
$\log(\sigma_u)$	0.01	0.03



Results

Shiga- og veratoxin producerende E. coli.



Poisson-Gamma



$$Y_i \sim \operatorname{Pois}(\lambda_i u_i)$$
 (2a)
 $u_i \sim \operatorname{G}(1, \phi)$ (2b)

$$u_i \sim \mathrm{G}(1,\phi)$$
 (2b)