Model & likelihood vs. code

Model

ODE modul:

$$\frac{dS}{dt} = -\beta SI$$

where S, I, R are the fractions of the population that are susceptible, infectious, and recovered

Measurment egn

suppose our data is a measure of prevalence, but not every infection is fested/reported.

Then we weasure:

Code

```
SIRode <- function(t, x, params){
    S = x[1]
    I = x[2]
    R = x[3]

    b = params[1]
    g = params[2]

    dSdt = -b*S*I
    dIdt = b*S*I - g*I
    dRdt = g*I

    list(c(dSdt, dIdt, dRdt))
}</pre>
```

= the reported # of infections

but, to simplify things, we'll combine k and N into one parameter, K

Initial Conditions

We can use the measurement equation to roughly estimate the initial condition for I:

I(0) = data(0)/K

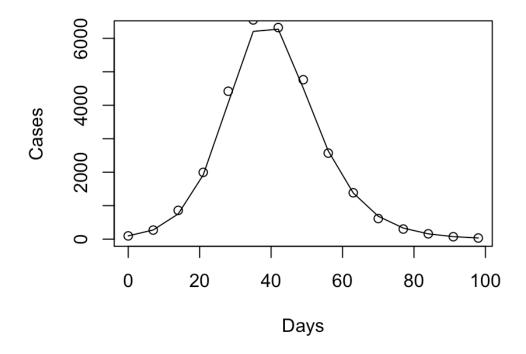
```
yfun = function(odeSim, params){odeSim[,3]/params[3]}
```

Simulate the model

```
times = c(0, 7, 14, 21, 28, 35, 42, 49, 56, 63, 70, 77, 84, 91, 98)
cases = c(97, 271, 860, 1995, 4419, 6549, 6321, 4763, 2571, 1385, 615, 302, 159, 72, 34)
dataset = cbind(times, cases)

params = c('beta'=0.4,'gamma'=0.25, 'kappainv'=1/80000)

xinit <- ode(x0fun(cases,params), times, SIRode, params, method='ode45')
plot(times, yfun(xinit,params), type='l')
points(dataset)</pre>
```



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(negative log likelihood

- LL = Poisson(z,y)

det mean=y

equivalent to:

- LL = Žyt + Žzt logyt

t=1 yt

```
SIRML=function(params, times, data) {
  params = abs(params)

# Simulate model
  xcurr = ode(x0fun(data,params), times, SIRode, params, method='ode45')

# Measurement equation
  y = yfun(xcurr,params)

# Negative Log Likelihood (NLL)
  NLL = sum(y) - sum(data*log(y)) # Poisson ML
}
```