

State-Space Models

State-Space Models....

- future state depends on current state (dynamic model)

$$X_{t+1} \sim X_t$$

State-Space Models....

- future state depends on current state (dynamic model)

$$X_{t+1} \sim X_t$$

- Observe variable Y but want to interpret response X
[when specific to time – Hidden **Markov** Model]

State-Space Models....

For when you observe this state (at time t) :

Y_t



But want to estimate this:

X_t



State-Space Model

$$X_t = f(X_{t-1}) + \varepsilon_t$$

Process Model

$$Y_t = g(X_t) + \omega_t$$

Data Model

- X = latent time series
- Y = observed data
- ε = process error
- ω = observation error

Random Walk State-Space Model

$$X_t = X_{t-1} + \varepsilon_t$$

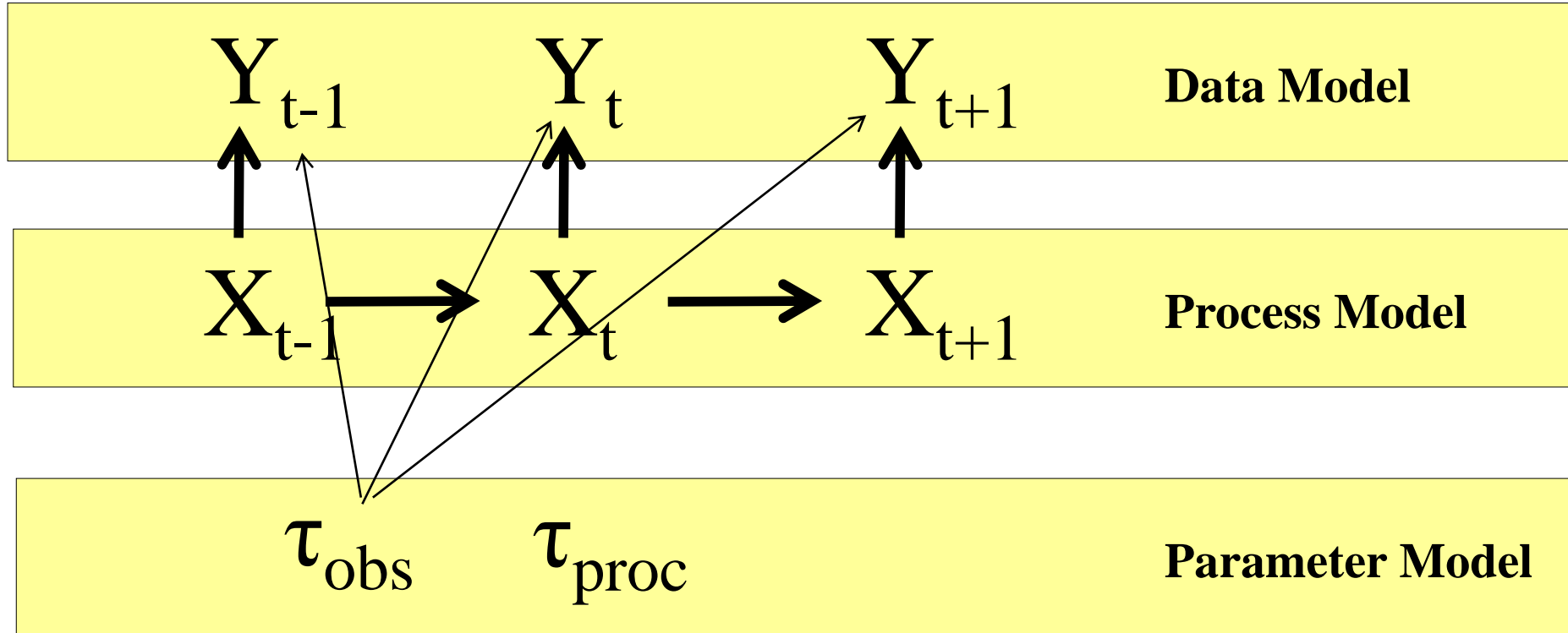
Process Model

$$Y_t = g(X_t) + \omega_t$$

Data Model

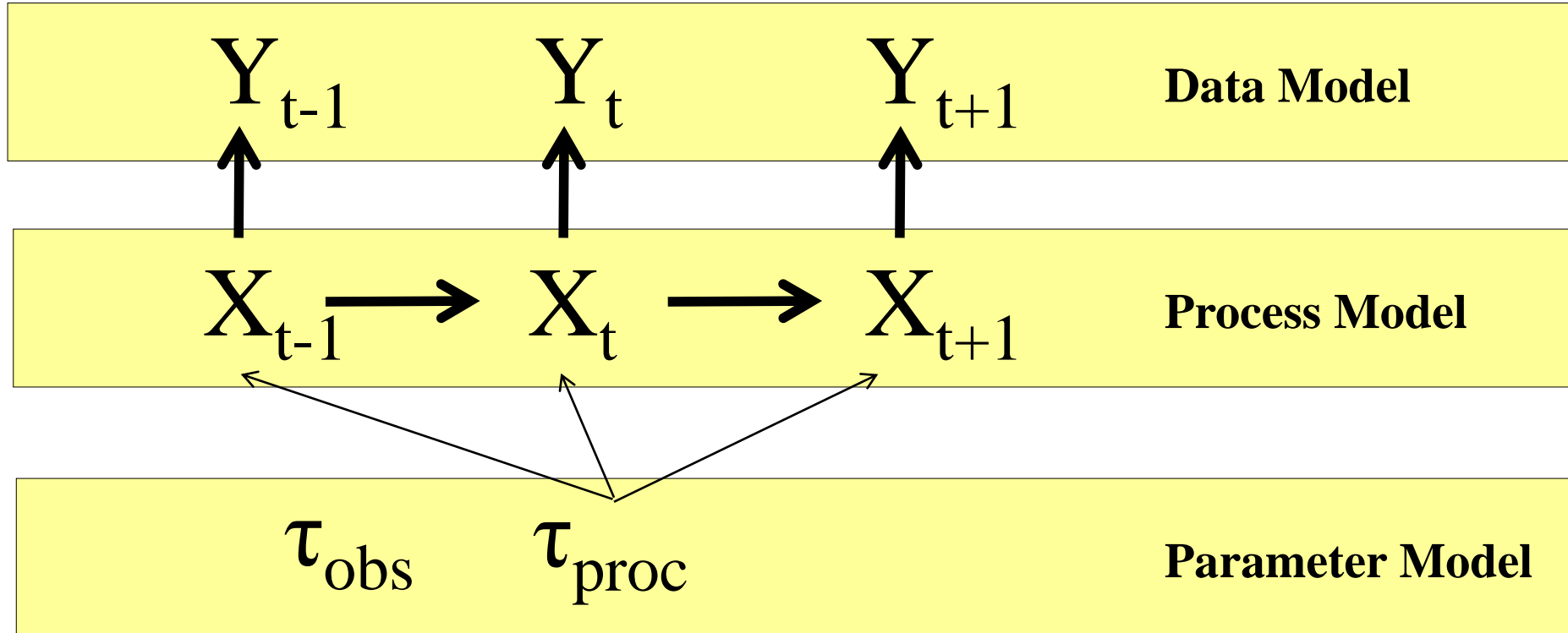
- X = latent time series
- Y = observed data
- e = process error
- w = observation error

Random Walk State-Space Model



Y's are conditionally independent given the X's

Random Walk State-Space Model



X_t estimate depends on X_{t+1} , X_{t-1} and Y_t


```
RandomWalk = "  
model{
```

```
#### Data Model
```

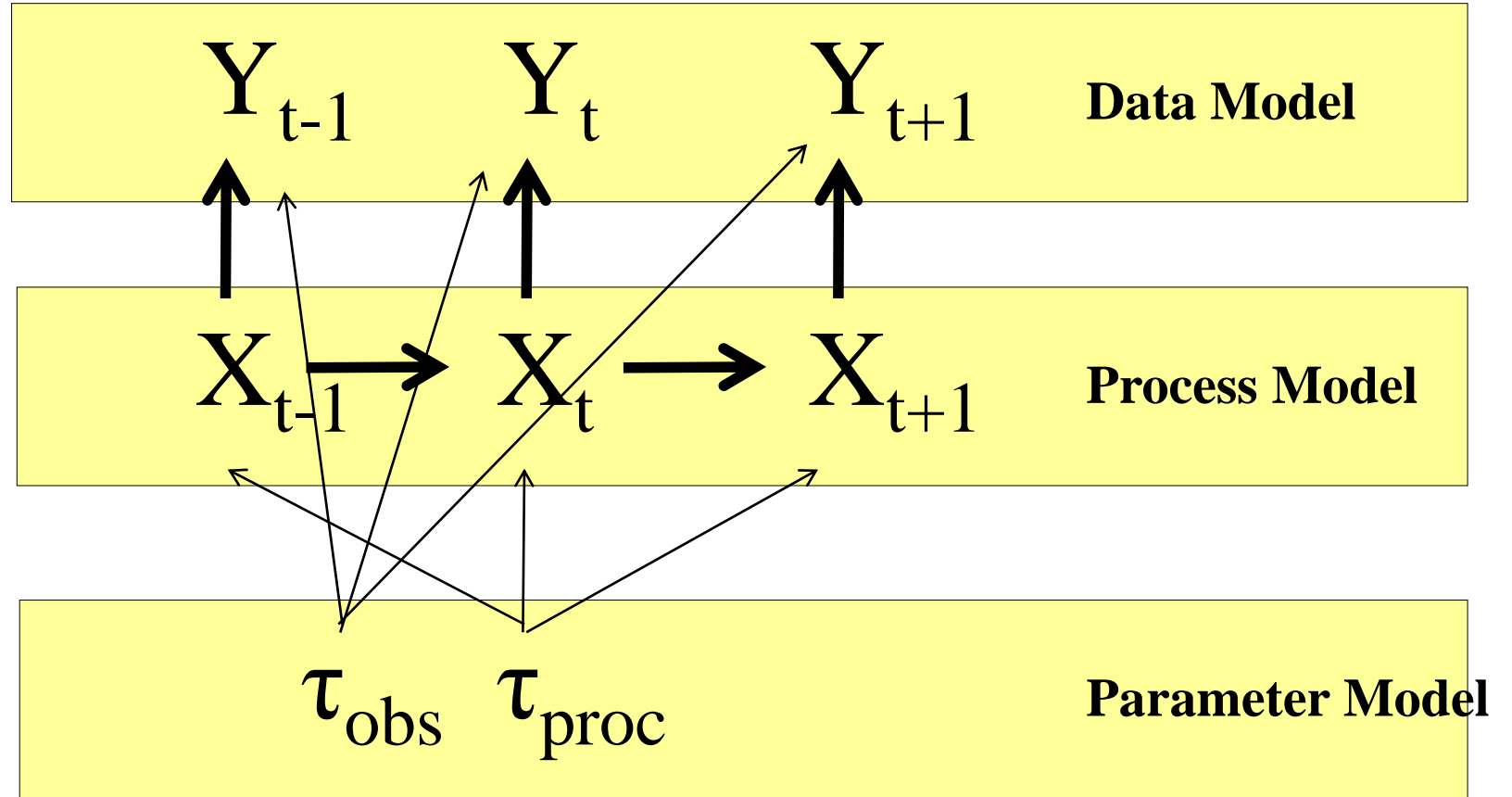
```
for(i in 1:n){  
  y[i] ~ dnorm(x[i],tau_obs)  
}
```

```
#### Process Model
```

```
for(i in 2:n){  
  x[i]~dnorm(x[i-1],tau_proc)  
}
```

```
#### Priors
```

```
x[1] ~ dnorm(x_ic,tau_ic)  
tau_obs ~ dgamma(a_obs,r_obs)  
tau_proc ~ dgamma(a_proc,r_proc)  
}
```



```
RandomWalk = "  
model{
```

```
#### Data Model
```

```
for(i in 1:n){  
  y[i] ~ dnorm(x[i],tau_obs)  
}
```

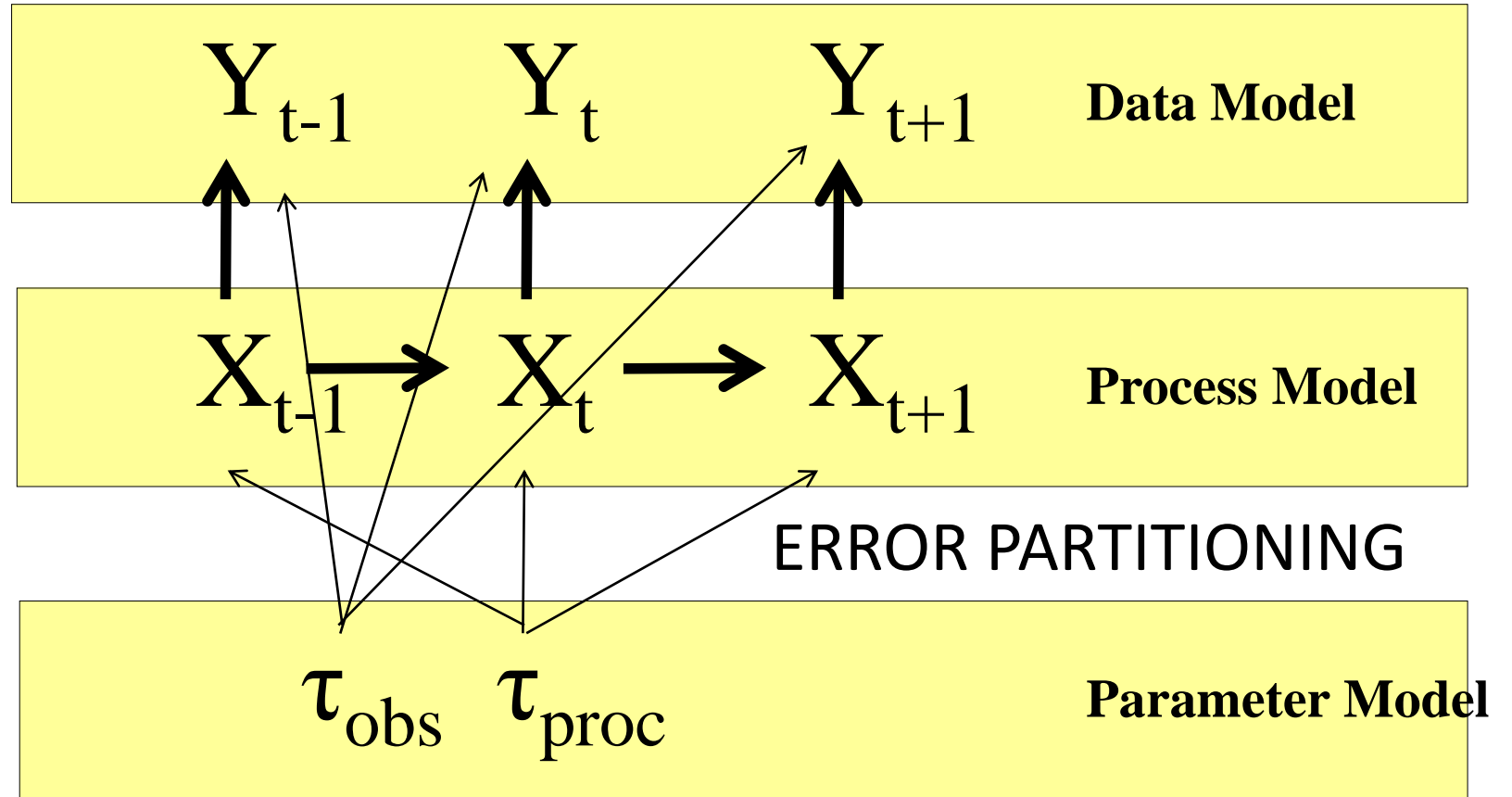
```
#### Process Model
```

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for(i in 2:n){  
  x[i]~dnorm(x[i-1],tau_proc)  
}
```

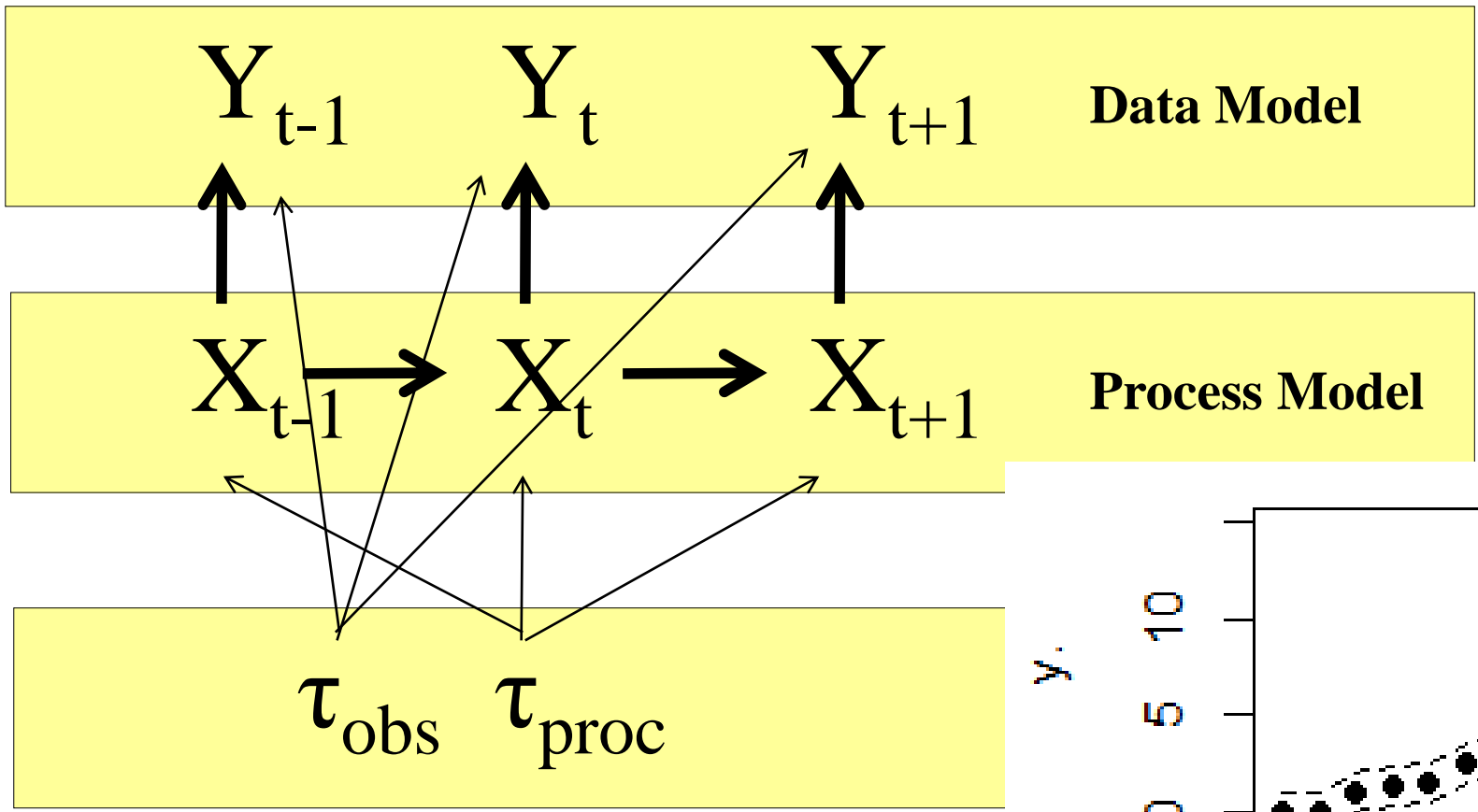
```
#### Priors
```

```
x[1] ~ dnorm(x_ic,tau_ic)  
tau_obs ~ dgamma(a_obs,r_obs)  
tau_proc ~ dgamma(a_proc,r_proc)
```

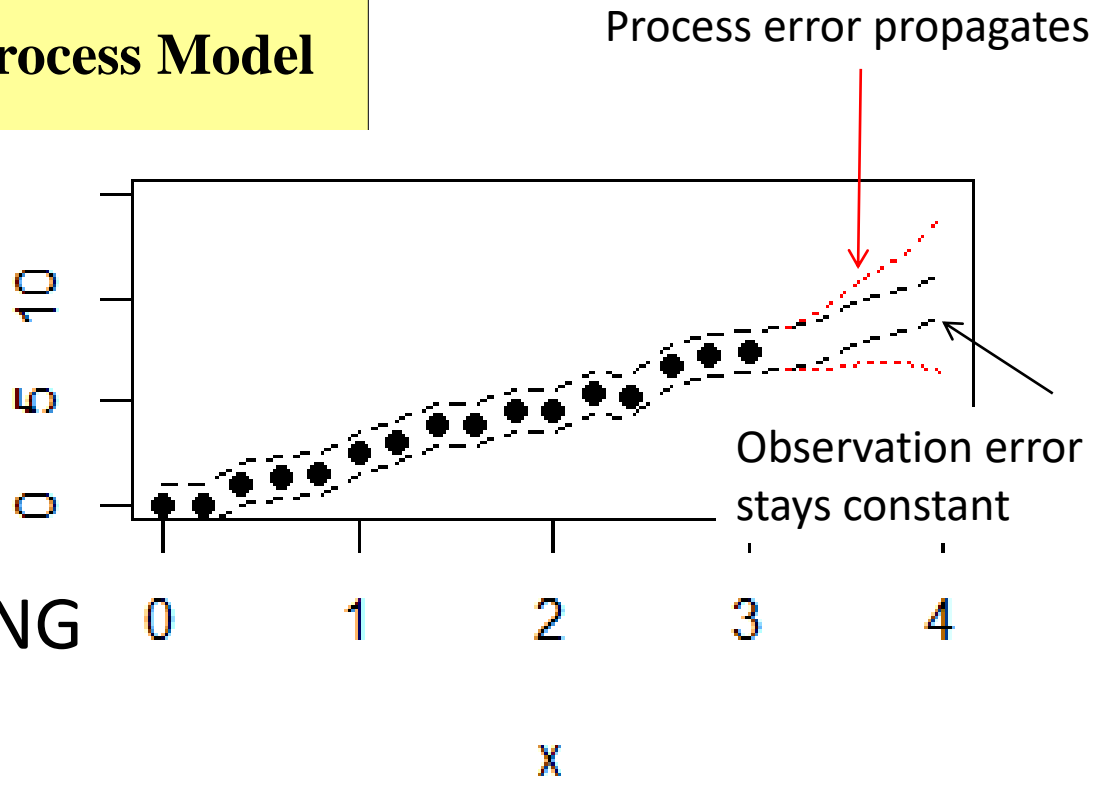
```
}
```



X_t depends on X_{t+1} and X_{t-1} and their uncertainties (τ_{proc})



ERROR PARTITIONING



```
RandomWalk = "  
model{
```

```
#### Data Model
```

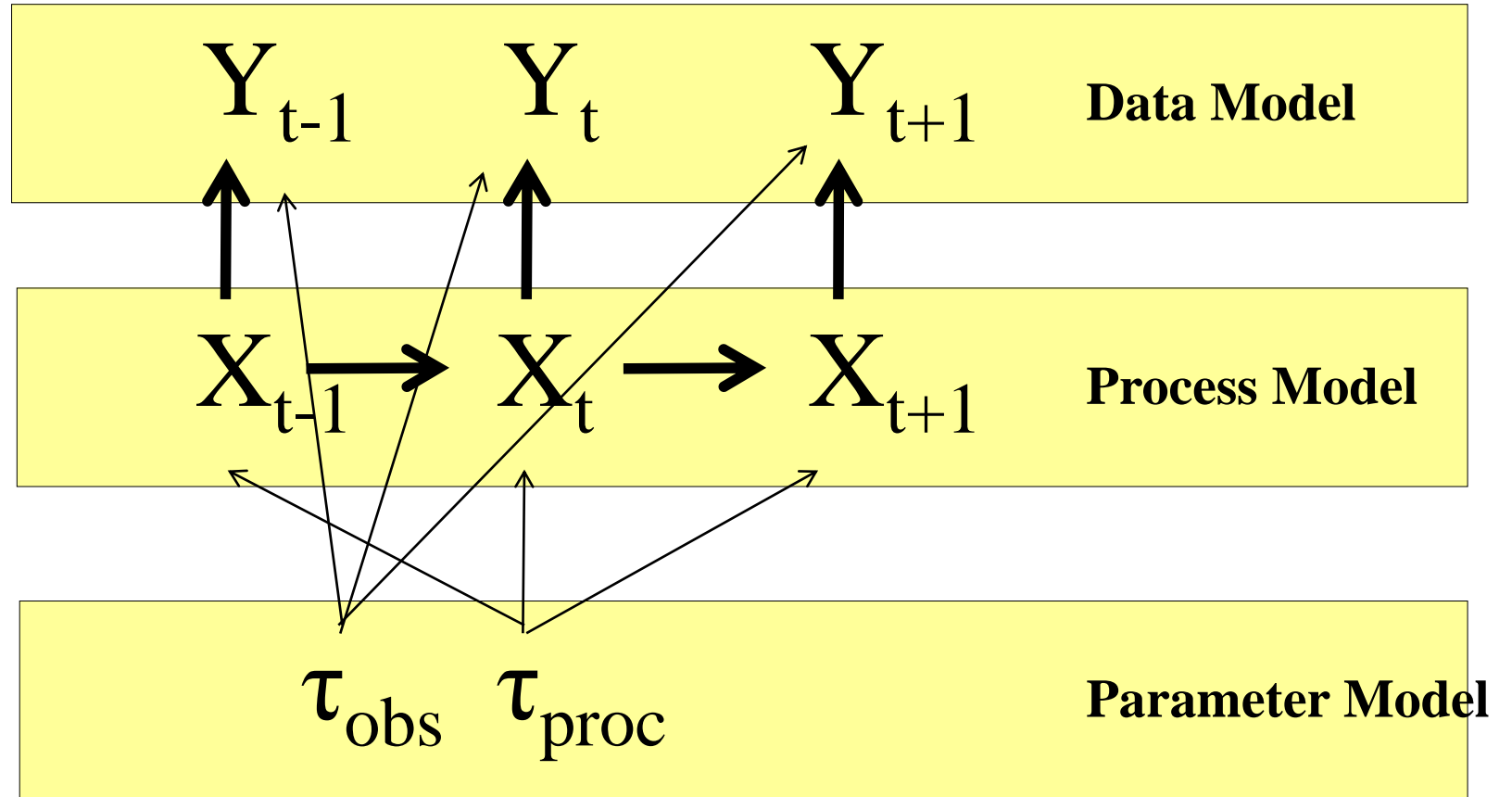
```
for(i in 1:n){  
  y[i] ~ dnorm(x[i],tau_obs)  
}
```

```
#### Process Model
```

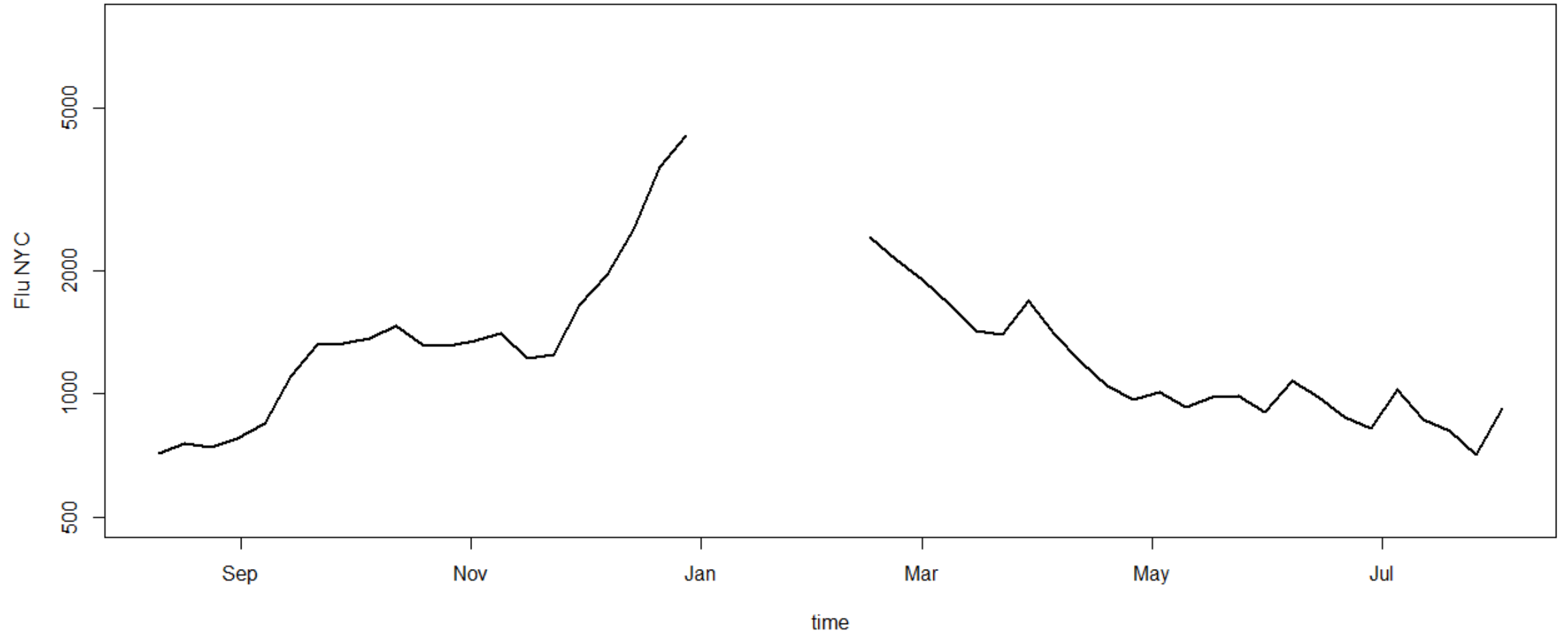
```
for(i in 2:n){  
  x[i]~dnorm(x[i-1],tau_proc)  
}
```

```
#### Priors
```

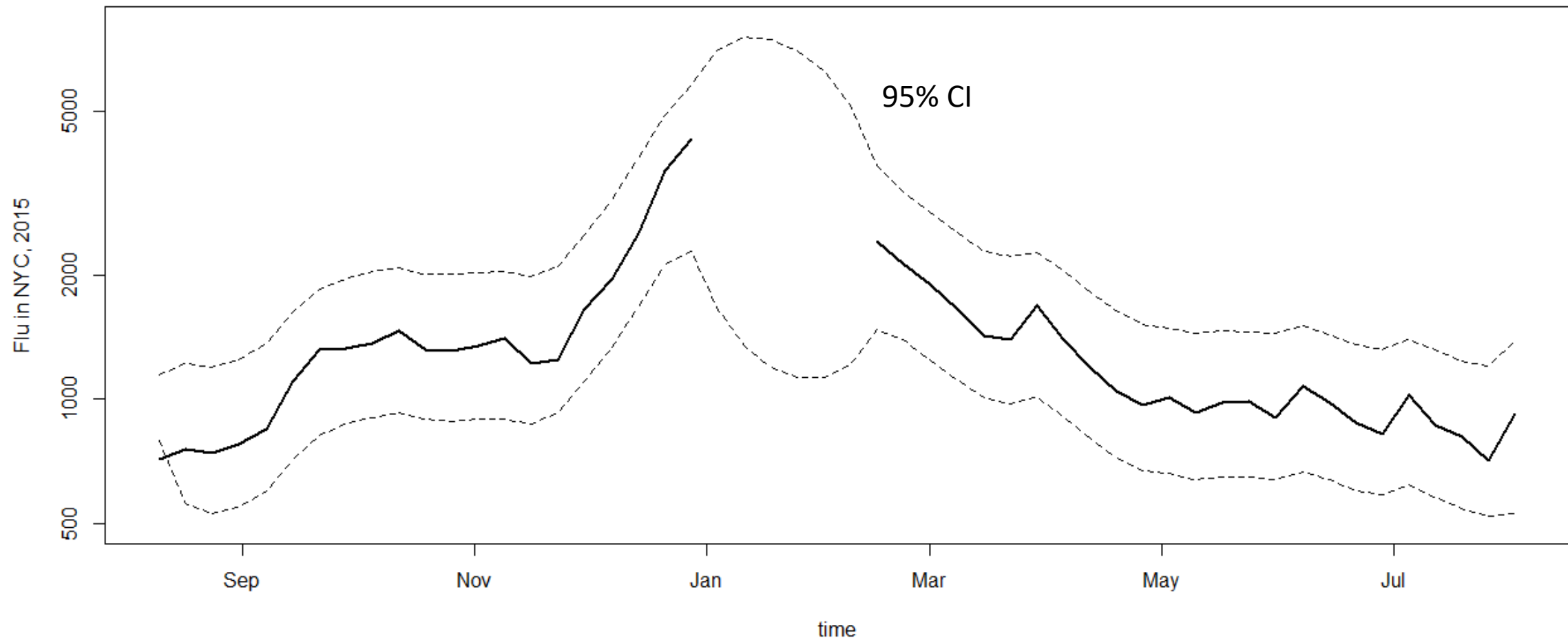
```
x[1] ~ dnorm(x_ic,tau_ic)  
tau_obs ~ dgamma(a_obs,r_obs)  
tau_proc ~ dgamma(a_proc,r_proc)  
}
```



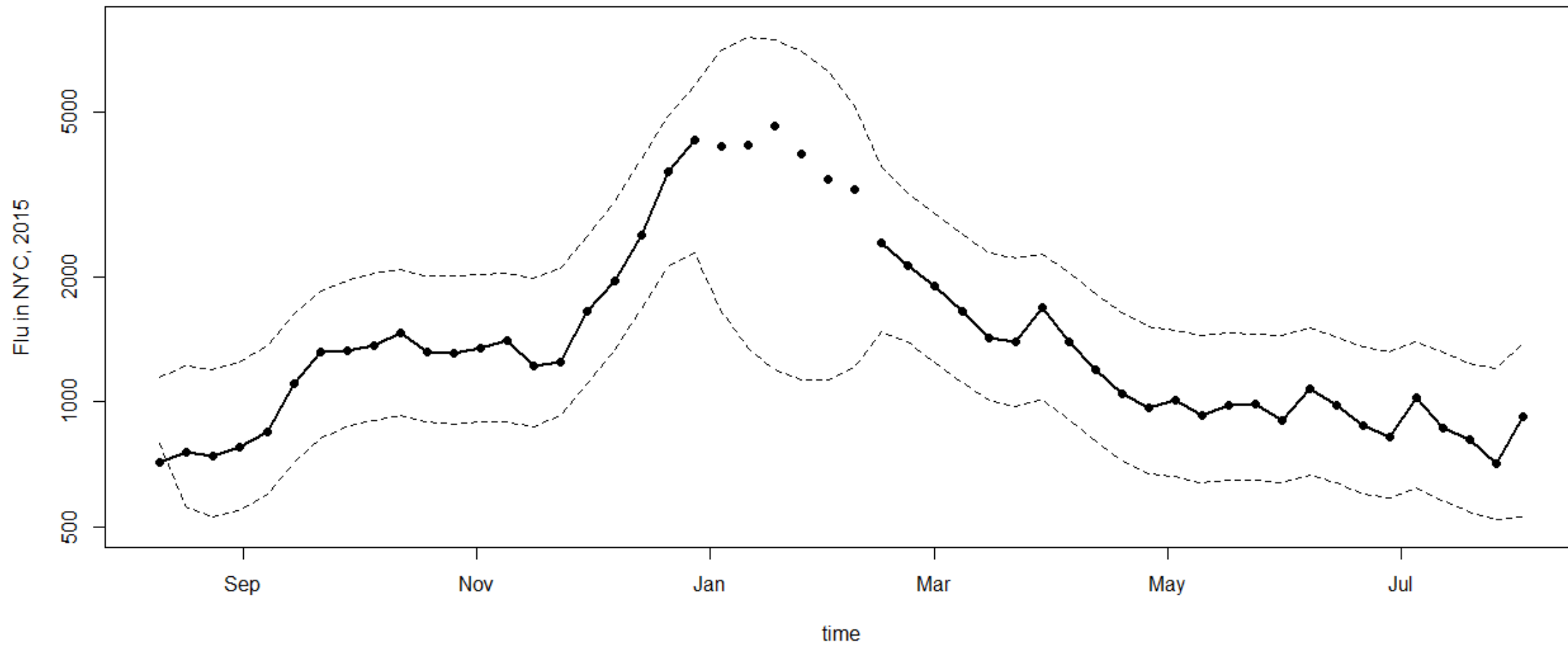
Random Walk State Space Model



Random Walk State Space Model



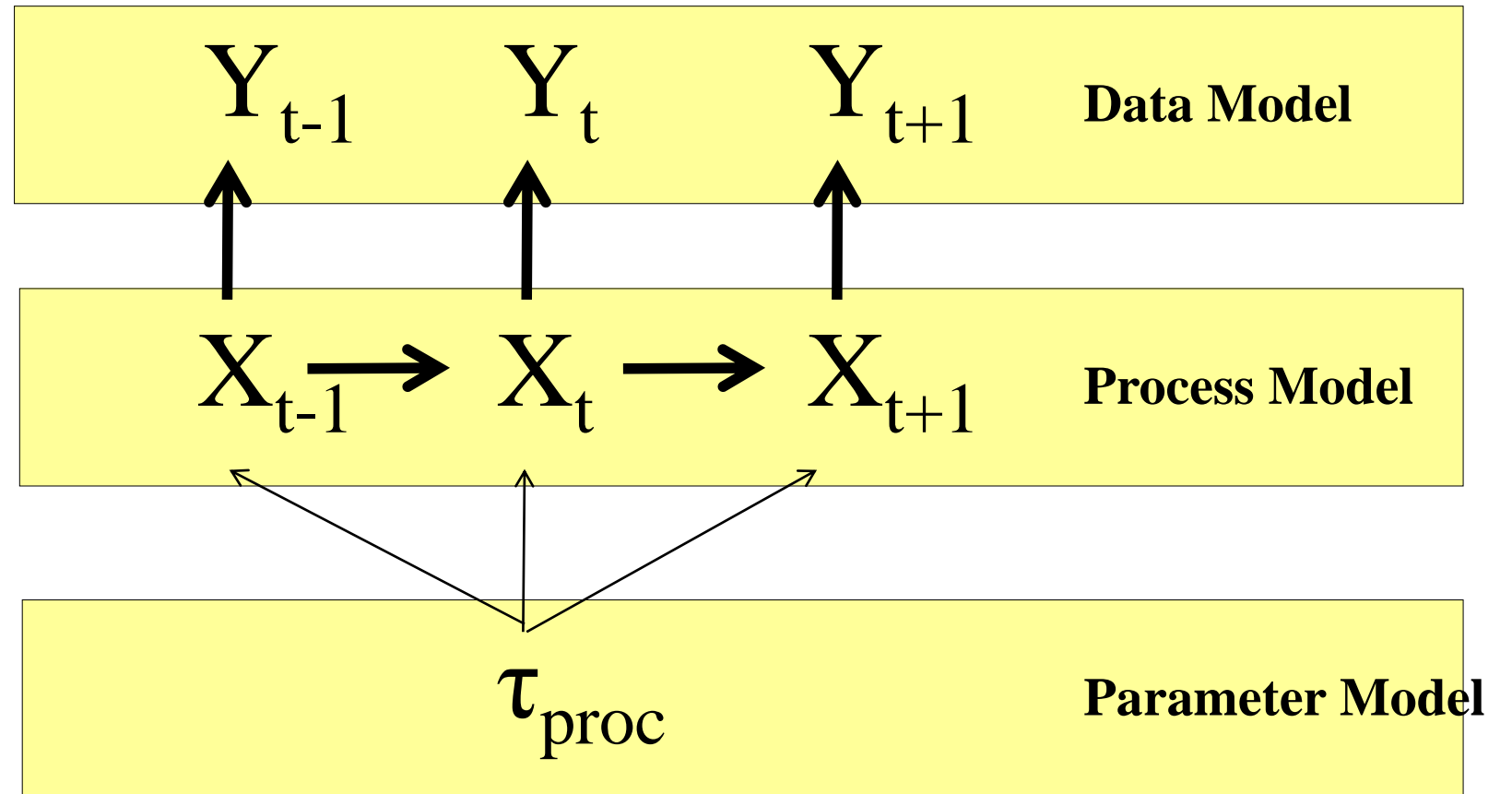
Random Walk State Space Model



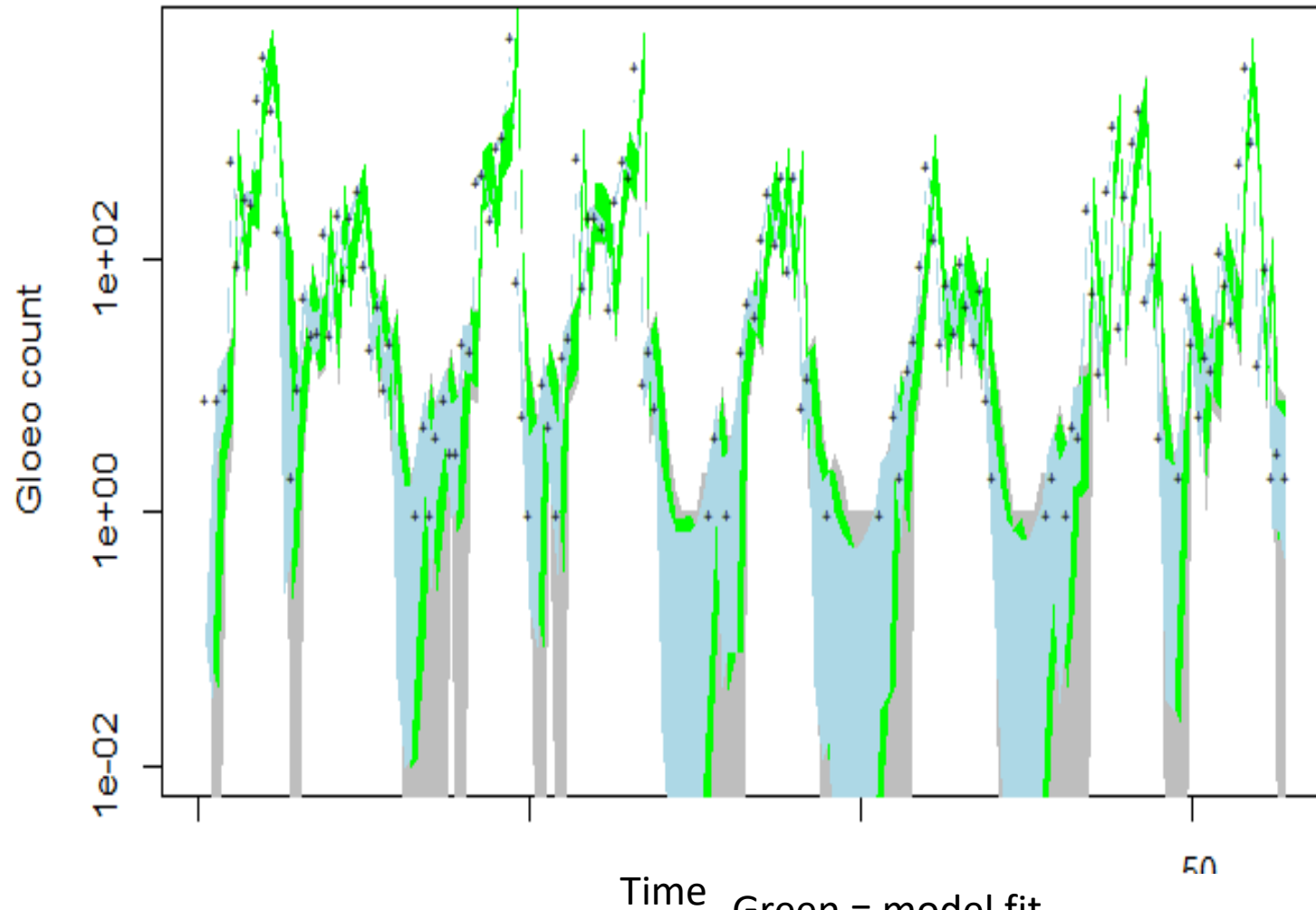
Beyond the Gaussian Random Walk

- Non-Gaussian data model
- Covariates
- Detection Models

Algal Bloom Example – Poisson RW



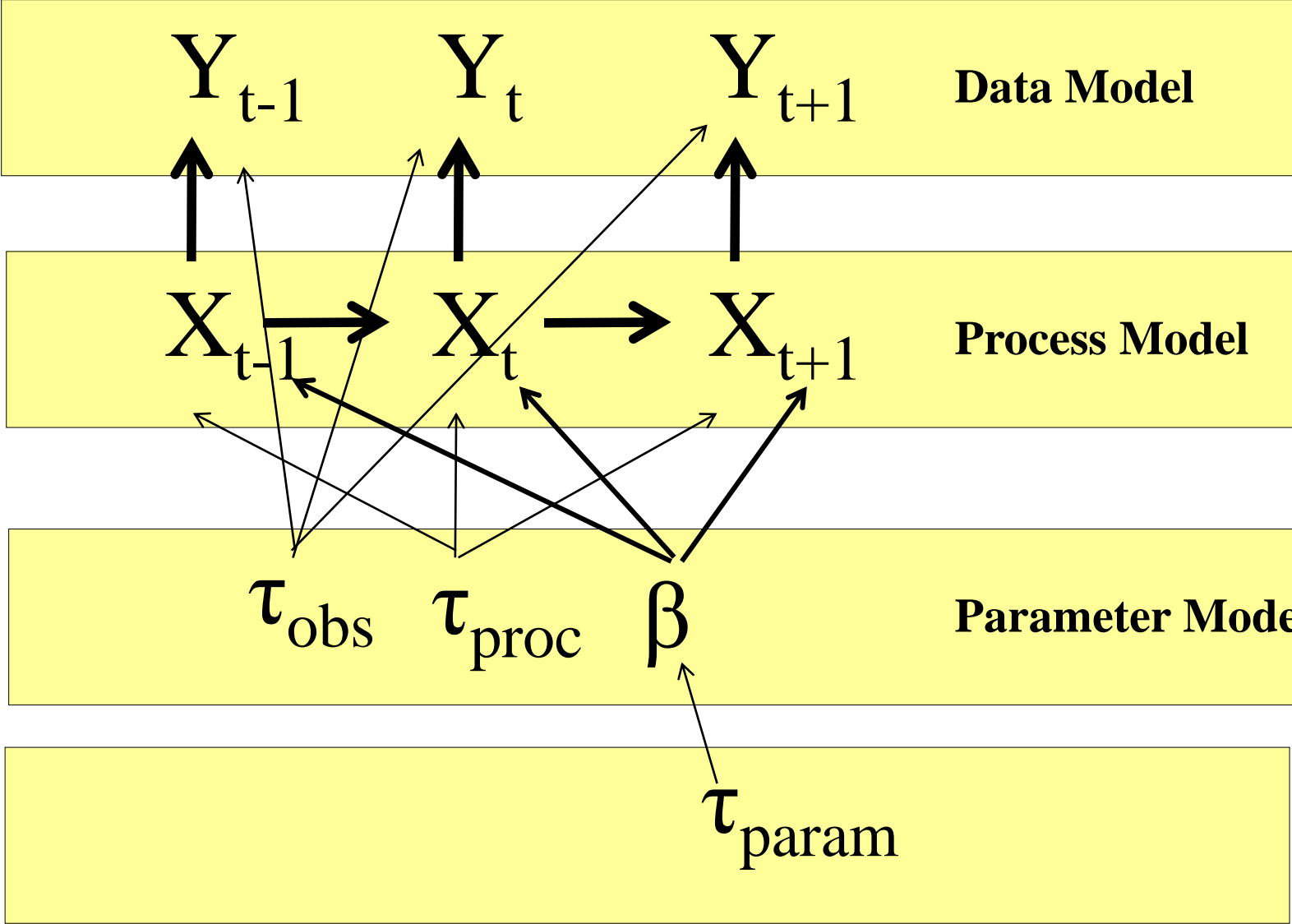
Algal Bloom Example – Poisson RW



GLEON: Lofton, Brentrup, Beck,
Zwart, Bhattacharya, McCullough

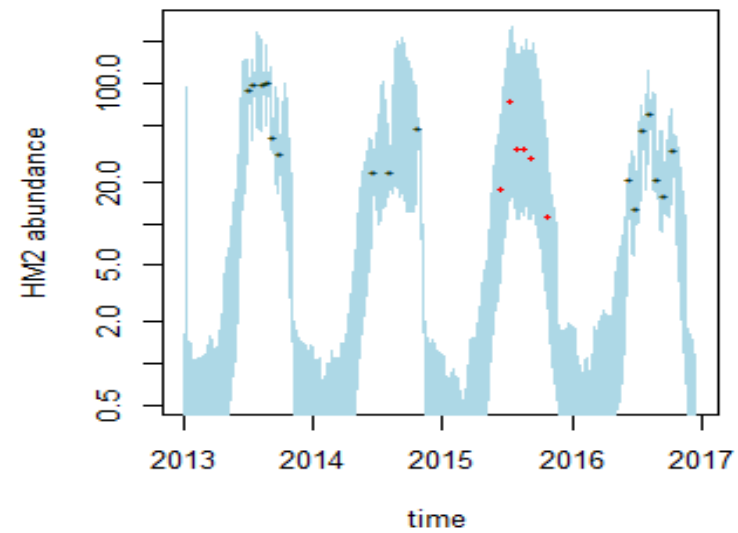
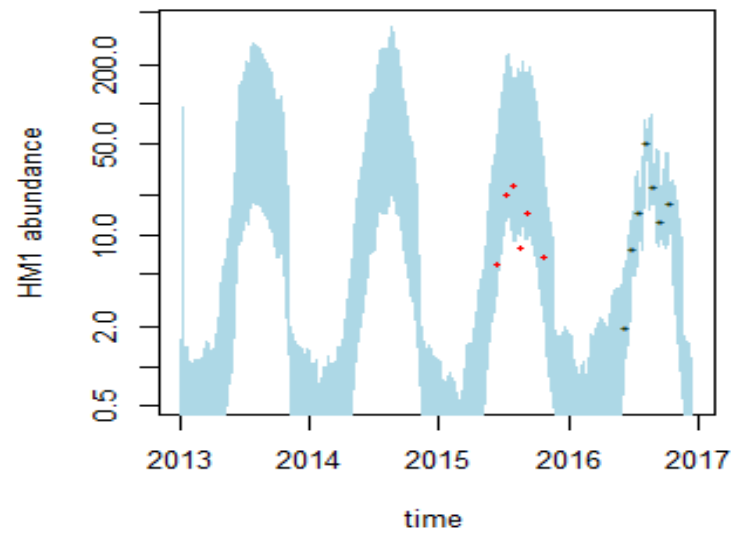
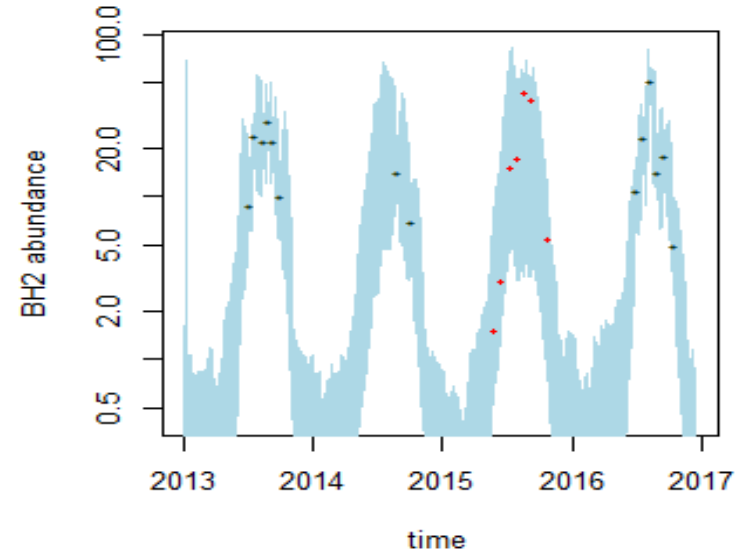
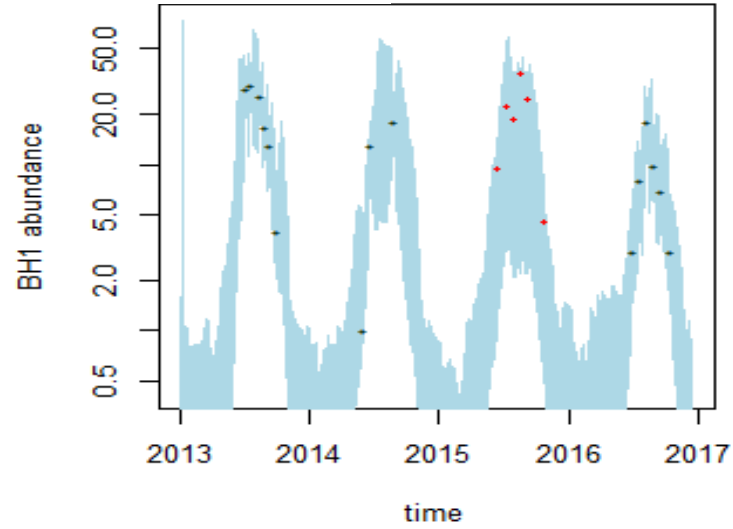
Green = model fit
Blue = predictive interval
Gray = predictive interval + data model

Jake Zwart – GIF showing 1-step ahead forecast RW

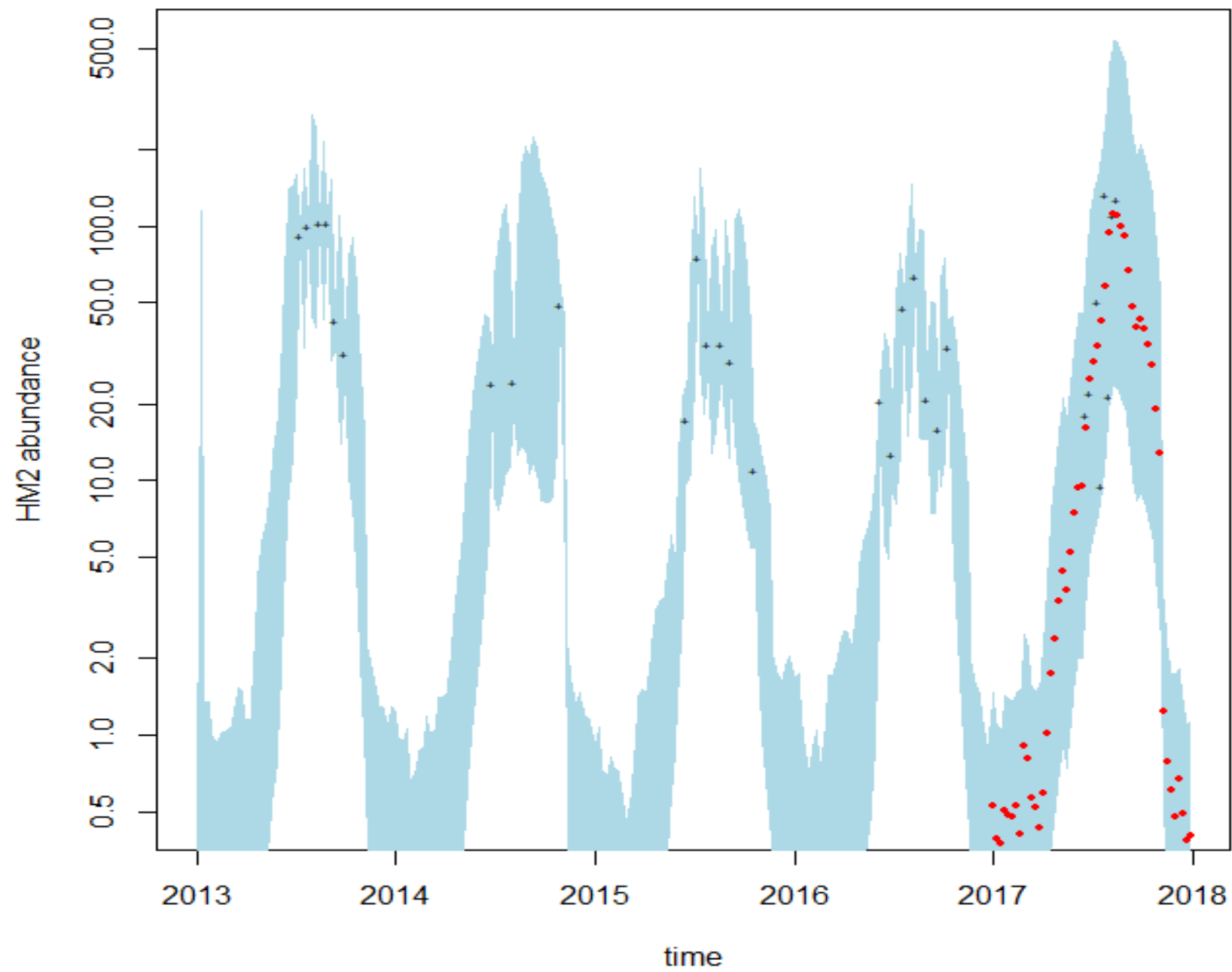


Process Model

$$X_{it} = f(X_{i,t-1} + \text{weathert} + \text{juv habitati}) + \varepsilon_{it}$$



Data Model $Y_{it} = g(X_{it})$

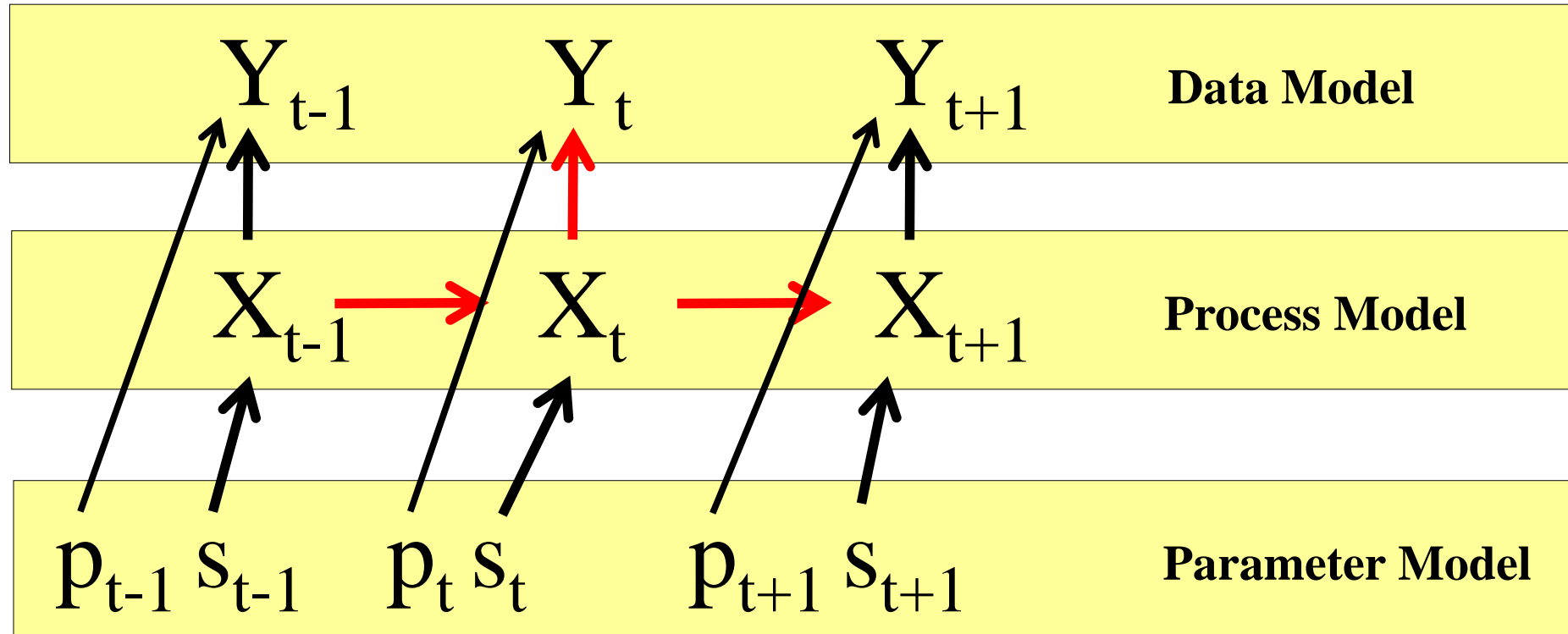


Capture-Recapture

- Individuals captured, marked, and released with goal of estimating population size.
- Over repeated censuses will recapture some fraction of the population
- Assume recapture is random



Mark Recapture State Space



Capture-Recapture

- Suppose an individual record consists of capture data

$$Y_i = [1,0,1,0,0]$$

- This is compatible with the following survival

$$X_i = [1,1,1,0,0]$$

$$X_i = [1,1,1,1,0]$$

$$X_i = [1,1,1,1,1]$$

Basic Mark-Recapture State Space

- Process model

$$P(X_t = 1 \mid X_{t-1} = 1) = s_t$$

$$P(X_t = 1 \mid X_{t-1} = 0) = 0$$

$$P(X_t = 0 \mid X_{t-1} = 1) = 1 - s_t$$

$$P(X_t = 0 \mid X_{t-1} = 0) = 1$$

Bernoulli Survival Probability

Basic Mark-Recapture State Space

- Process model

$$P(X_t = 1 \mid X_{t-1} = 1) = s_t$$

$$P(X_t = 1 \mid X_{t-1} = 0) = 0$$

$$P(X_t = 0 \mid X_{t-1} = 1) = 1 - s_t$$

$$P(X_t = 0 \mid X_{t-1} = 0) = 1$$

- Observation model

$$P(Y_t = 1 \mid X_t = 1) = p_t$$

$$P(Y_t = 1 \mid X_t = 0) = 0$$

$$P(Y_t = 0 \mid X_t = 0) = 1$$

$$P(Y_t = 0 \mid X_t = 1) = 1 - p_t$$

- Priors on p and s (e.g. Beta)

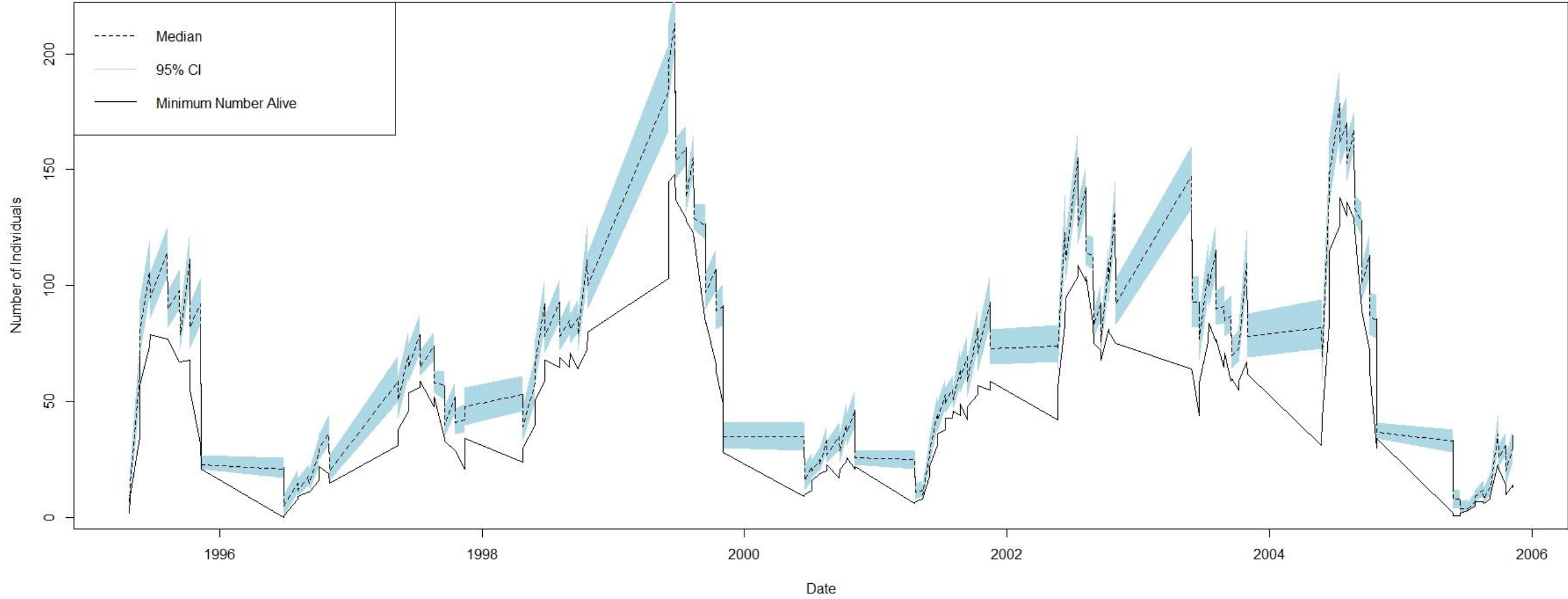
Bernoulli Survival Probability

$$s^x (1-s)^{1-x}$$

Bernoulli Detection Probability

$$p^x (1-p)^{1-x}$$

Estimated Mouse Abundance



John Foster

Generality of the State Space Model

- Neither X nor Y need be Normal
- X and Y don't need to be the same type of data
- X and Y don't need to have the same time scale
- Easily handles missing data (gaps)
- Can accommodate multiple data sources (Y 's) – and they can be different data types