

# Methods of Applied Stats, Non-parametric and semi-parametric models

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Sept to Dec 2020

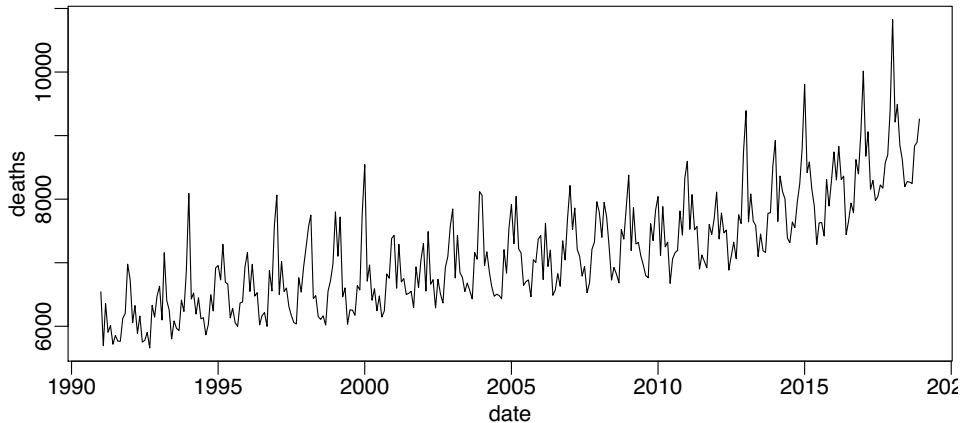
# Non-parametric modelling

also known as fitting wiggly lines through points

- Penalized least squares
- Bayesian semi-parametrics with INLA

## Deaths in Ontario (monthly)

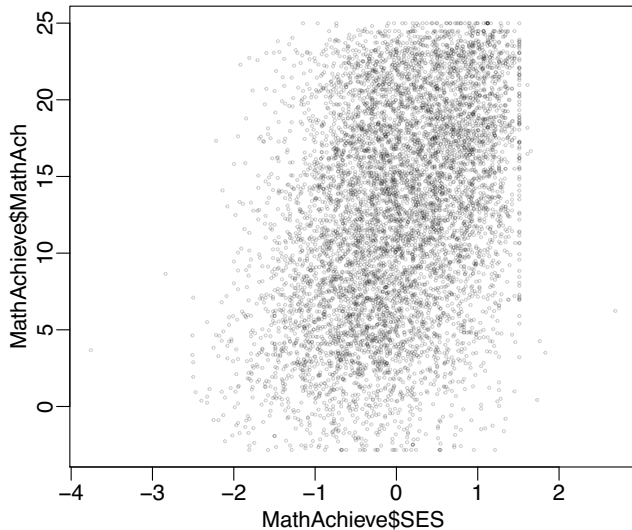
```
> plot(oDeaths$date, oDeaths$Value, xlab='date', ylab='deaths', type='l')
```



see [pbrown.ca/teaching/appliedstats/data/oDeaths.R](http://pbrown.ca/teaching/appliedstats/data/oDeaths.R)

## Math scores

```
> data('MathAchieve',  
+   package='MEMSS')  
> plot(MathAchieve$SES,  
+   MathAchieve$MathAch,  
+   cex=0.3, col='#00000030')
```



## Infant mortality rate

scraped from Wikipedia

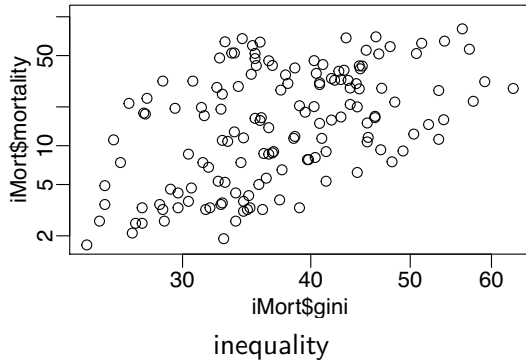
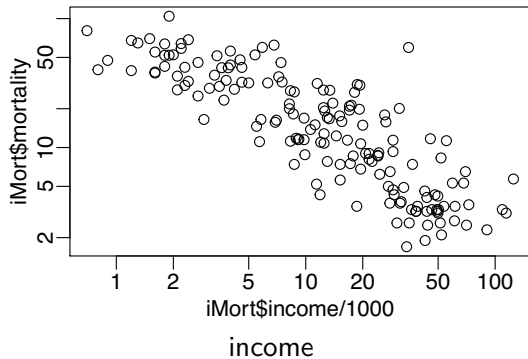
Warning: NAs introduced by coercion

```
> knitr::kable(iMort[grep("United|Japan|Egypt|Yemen|Canada",  
+   iMort$Country), c("Country", "income", "gini",  
+   "mortality")]])
```

	Country	income	gini	mortality
26	Canada	48100	33.8	4.3
42	Egypt	13000	31.5	17.1
72	Japan	42700	32.9	1.9
153	United Arab Emirates	68200	32.5	5.3
154	United Kingdom	43600	34.8	4.1
155	United States	59500	41.4	5.3
160	Yemen	2300	36.7	41.9

# Infant Mortality

infant mortality



# Generalized additive models

Trevor J Hastie and Robert J Tibshirani

(1990). *Generalized additive models*.

Vol. 43. CRC Press

Simon Wood (2006). *Generalized additive models: an introduction with R*. CRC press

$$Y_i \sim (\lambda_i, \theta)$$
$$g(\lambda_i) = X_i\beta + f_1(W_{1i}) + \dots + f_k(W_{ki})$$

- $Y_i$  are responses
- $\theta$  is the response distribution
- $X_i$  and  $W_i$  are covariates
- $f(w)$  is some sort of wiggly line
- If we put no restrictions or assumptions on  $f$ , the estimate  $\hat{f}(w)$  will interpolate the data perfectly
- ...which isn't very interesting

## Penalized likelihood

$$\ell_p(\beta, f, \alpha; \mathbf{Y}) = \log[\text{pr}(\mathbf{Y}; \beta, f)] - \alpha \int \left[ \frac{\partial^2 f(x)}{\partial x^2} \Big|_u \right]^2 du$$

$$Y_i \sim (\lambda_i, \theta)$$
$$g(\lambda_i) = X_i \beta + f(W_i)$$

Estimates, given  $\alpha$

$$\hat{\beta}(\alpha), \hat{f}(\alpha) = \underset{\beta, f}{\text{argmin}} \ell_p(\beta, f, \alpha; \mathbf{Y})$$

- The last term is the integrated squared second derivative
- $\alpha$  is a penalty parameter
- A smooth  $f(x)$  should have small  $f''(x)$
- The second derivative is interpretable as 'energy'
- a good  $\hat{f}$  is a compromise between fitting the data and being smooth.



## Cubic splines

The  $f$  which maximizes the penalized likelihood must be a cubic spline polynomial

- ...a cubic polynomial where the third derivative changes at each data point  $W_i$
- but the 0, 1, and 2 derivatives are continuous

$$\int \left[ \frac{\partial^2 f(x)}{\partial x^2} \Big|_u \right]^2 du = F^T D^T R^{-1} D F$$

$$f'_t \approx f'_t - f'_{t-1}$$

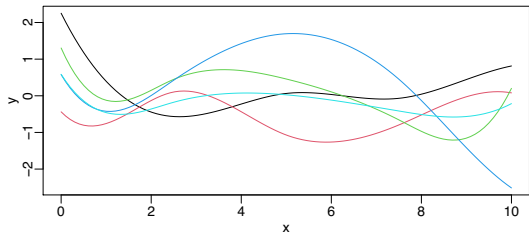
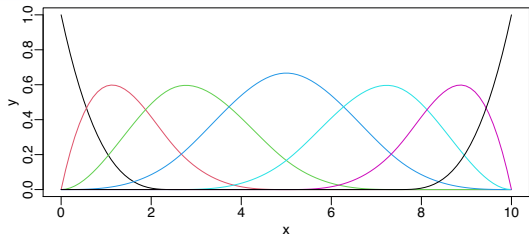
$$f''_t \approx f''_t - f''_{t-1}$$

$$\approx f''_t - 2f''_{t-1} + f''_{t-2}$$

$$F = \begin{pmatrix} f(W_1) \\ \vdots \\ f(W_N) \end{pmatrix} \quad D = \begin{pmatrix} 1 & -2 & 1 & 0 & 0 & 0 \\ 0 & 1 & -2 & 1 & 0 & 0 \\ 0 & 0 & 1 & -2 & 1 & 0 \\ 0 & 0 & 0 & 1 & -2 & 1 \end{pmatrix} \quad R = \begin{pmatrix} 2/3 & 1/6 & 0 & 0 \\ 1/6 & 2/3 & 1/6 & 0 \\ 0 & 1/6 & 2/3 & 1/6 \\ 0 & 0 & 1/6 & 2/3 \end{pmatrix}$$

Peter J Green and Bernard W Silverman (1993). *Nonparametric regression and generalized linear models: a roughness penalty approach*. CRC Press

## B-Splines



$$F = B^T \gamma$$

$$\ell_p = \ln(\mathcal{L}(\beta, \gamma | Y)) - \alpha \gamma^T B D^T R^{-1} D B^T \gamma$$

- Basis functions which approximate cubic polynomials
- There are efficient ways of maximizing the joint probability if  $U$  is restricted to B-splines

1	2	3	4	5	6	7
2.3	0.2	-1.2	0.5	-0.5	0.6	0.8
-0.4	-1.5	1.3	-1.7	-1.0	0.3	0.1
1.3	-1.0	1.0	0.5	-0.5	-1.9	0.2
0.6	-1.2	0.6	2.2	0.9	-1.8	-2.5
0.6	-1.0	0.1	0.1	-0.4	-0.7	-0.2

Paul HC Eilers and Brian D Marx (1996). "Flexible smoothing with B-splines and penalties". In: *Statistical science*, pp. 89–102

Reduced rank approximation

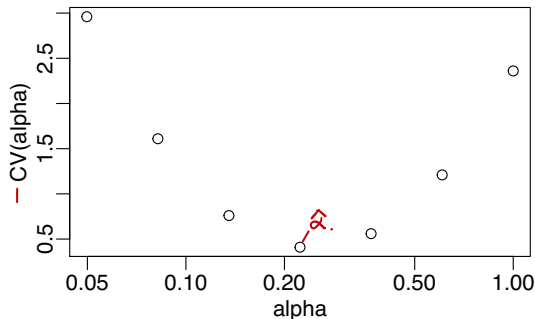
## Estimating $\alpha$

- Big  $\alpha \Rightarrow$  smooth curve
- Small  $\alpha \Rightarrow$  close fit to the data

### Cross validation

- Find  $\hat{\lambda}^{(-i)}$  by excluding observation  $i$
- compute  $pr(Y_i|\hat{\lambda}^{(-i)})$
- repeat for  $i = 1 \dots N$
- $CV(\alpha) = \sum_i \log[pr(Y_i|\hat{\lambda}^{(-i)})]$

$$\hat{\alpha} = \operatorname{argmax}_{\alpha} CV(\alpha)$$



## B-splines and Generalized cross-validation

$$Q_{ii} = g'(\lambda_i)^2 / \text{var}(Y_i | \lambda_i)$$

$$B_{ij} = B_j(W_i)$$

$$U = Ba$$

$$B^T Q (y - \lambda) + B^T Q B a = (B^T Q B + \alpha D^T D) a$$

$$H = B (B^T Q B + \alpha D^T D)^{-1} B^T Q$$

$$GCV(\alpha) = \sum_i \{ (y_i - \lambda_i)^2 / [N - \text{trace}(H)] \}$$

- $Q$  depends on  $\lambda$
- at iteration  $r$ ,  $Q^{(r)}$  is computed using  $\lambda^{(r-1)}$ .
- It suffices to know that the B-spline GCV algorithm is fast

*Actually, the name nonparametric is not always well chosen. It might apply to kernel smoothers and running statistics, but spline smoothers are described by parameters, although their number can be large. It might be better to talk about “overparametric” techniques or “anonymous” models; the parameters have no scientific interpretation.*

Paul HC Eilers and Brian D Marx (1996). “Flexible smoothing with B-splines and penalties”. In: *Statistical science*, pp. 89–102

# Penalized likelihood and random effects models

## Random effects

$$Y_i|U_i \sim (\lambda_i)$$

$$g(\lambda_i) = X_i\beta + U_i$$

$$\mathbf{U} \sim \text{MVN} \left[ 0, \frac{2}{\alpha} (D^\top R^{-1} D)^{-1} \right]$$

Frequentist inference:

$$\hat{\mathbf{U}} = E(\mathbf{U}|\mathbf{Y}; \hat{\beta}, \hat{\alpha})$$

Bayesian inference:

$$\hat{\mathbf{U}} = E(\mathbf{U}|\mathbf{Y})$$

## Penalized likelihood

Write  $U_i = f(W_i)$

$$\begin{aligned} \ell_p(\beta, f, \alpha; \mathbf{Y}) &= \log[\text{pr}(\mathbf{Y}; \beta, f)] - \alpha \int f''(u)^2 du \\ &= \log[\text{pr}(\mathbf{Y}; \beta, f)] - \alpha \mathbf{U}^\top D^\top R^{-1} D \mathbf{U} \\ &= \log[\text{pr}(\mathbf{Y}, \mathbf{U}; \beta, \alpha)] + C_1 \\ &= \log[\text{pr}(\mathbf{U}|\mathbf{Y}; \beta, \alpha)] + C_2 \end{aligned}$$

Inference using

$$\hat{\mathbf{U}} = \text{argmax}_{\mathbf{U}} \text{pr}(\mathbf{U}|\mathbf{Y}; \hat{\beta}, \hat{\alpha})$$

the conditional mode

## Wading into a philosophical minefield

- Frequentist:
  - plug-in MLE's of parameters  $\alpha$ ,  $\beta$
  - effect  $f$  by conditional mean, or median, or mode
- Bayesian:
  - put priors on parameters  $\alpha$ ,  $\beta$
  - effect  $f$  by posterior mean, or median, or mode
- Non-parametric
  - plug-in CV estimate  $\alpha$  by CV, MLE  $\beta$
  - effect  $f$  by conditional mode
  - Is this *Empirical Bayes*?
  - Restricting inference to the conditional mode is not a small detail!
  - ...the tails of the distribution are ignored

# How to write a GAM

## Non-parametric

$$Y_i \sim (\lambda_i, \theta)$$
$$g(\lambda_i) = X_i\beta + f(W_i; \nu)$$

where  $f(w; \nu)$  is a smoothly-varying function with roughness parameter  $\nu$ .

## Parametric

$$Y_i \sim (\lambda_i, \theta)$$
$$g(\lambda_i) = X_i\beta + U(W_i)$$
$$U(w) \sim \text{ARIMA}_{0,2,1}(\sigma^2, 2 - \sqrt{3})$$

$U(w)$  is <sup>almost</sup> a random walk of order 2 with variance  $\sigma^2$ .

Patrick E Brown and P de Jong (2001). "Nonparametric smoothing using state space techniques". In: *Canadian Journal of Statistics* 29.1, pp. 37–50. DOI: [10.2307/3316049](https://doi.org/10.2307/3316049)



## Math

$$Y_i \sim N(\mu_i, \sigma^2)$$

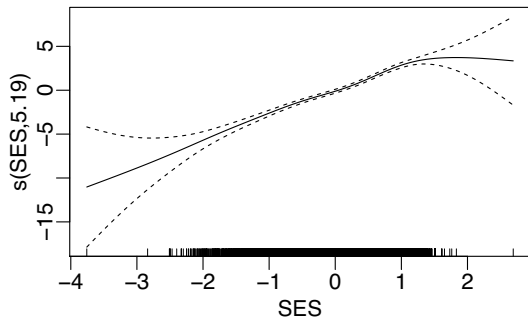
$$\mu_i = X_i\beta + f(W_i)$$

SES.

```
> library('mgcv')
> mathGam = gam(
+   MathAch ~ s(SES) + Minority*Sex,
+   data=MathAchieve)
> knitr::kable(
+   summary(mathGam)$p.table[,1:2],
+   digits=1)
```

	Estimate	Std. Error
(Intercept)	12.9	0.1
MinorityYes	-2.7	0.2
SexMale	1.4	0.2
MinorityYes:SexMale	-0.2	0.3

```
> plot(mathGam)
```



```
> mathGam$sp —  $\hat{\sigma}^2$ 
s(SES)
0.8254378
```

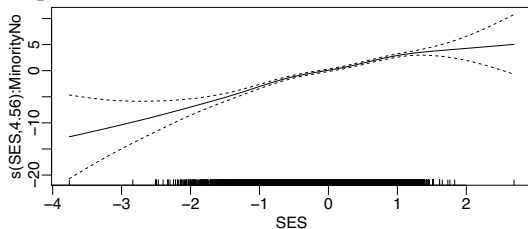
$$\mu_i = X_i\beta + \begin{cases} f_1(W_i), & X_{ii}=1 \\ f_2(W_i), & X_{ii}=0 \end{cases}$$

## Math, SES-minority interaction

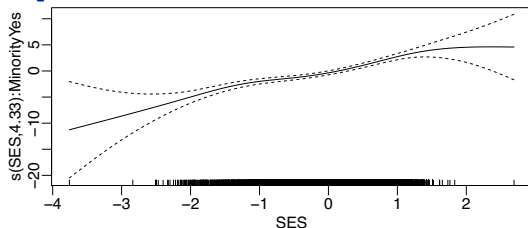
```
> mathGamInt = gam(
+   MathAch ~ s(SES, by=Minority) +
+   Minority*Sex,
+   data=MathAchieve)
> mathGamInt$sp
s(SES):MinorityNo s(SES):MinorityYes
      0.820158      0.614983
```

	Estimate	Std. Error
(Intercept)	12.8	0.1
MinorityYes	-2.9	0.2
SexMale	1.4	0.2
MinorityYes:SexMale	-0.1	0.3

```
> plot(mathGamInt, select = 1)
```



```
> plot(mathGamInt, select = 2)
```



## A common smoothing parameter?

$$Y_{ij} \sim N(\lambda_{ij}, \tau^2)$$

$$\lambda_{ij} = X_{ij}\beta + f_i(W_{ij}; \nu)$$

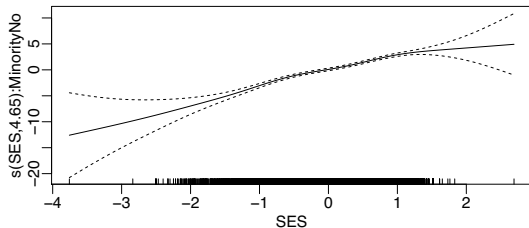
- $Y_{ij}$  is the observation for individual  $j$  in ethnic group  $i$
- $X_{ij}$  is a vector of covariates (ethnic group, sex, interaction)
- $f_i(w; \nu)$  is the smoothly-varying function of SES
  - for ethnic group  $i$
  - with roughness parameter  $\nu$ .

## Math, common smoothing parameter

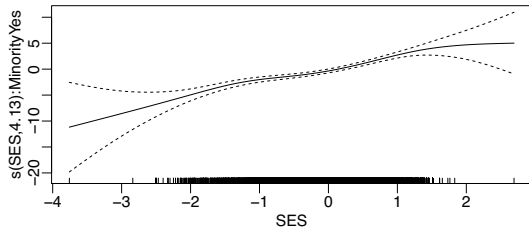
```
> mathGamIntC = gam(MathAch ~  
+   s(SES, by=Minority, id=1) +  
+   Minority*Sex,  
+   data=MathAchieve)  
> mathGamIntC$sp  
s(SES):MinorityNo  
0.7492505
```

	Estimate	Std. Error
(Intercept)	12.8	0.1
MinorityYes	-2.9	0.2
SexMale	1.4	0.2
MinorityYes:SexMale	-0.1	0.3

```
> plot(mathGamIntC, select = 1)
```



```
> plot(mathGamIntC, select = 2)
```



## Math 2d

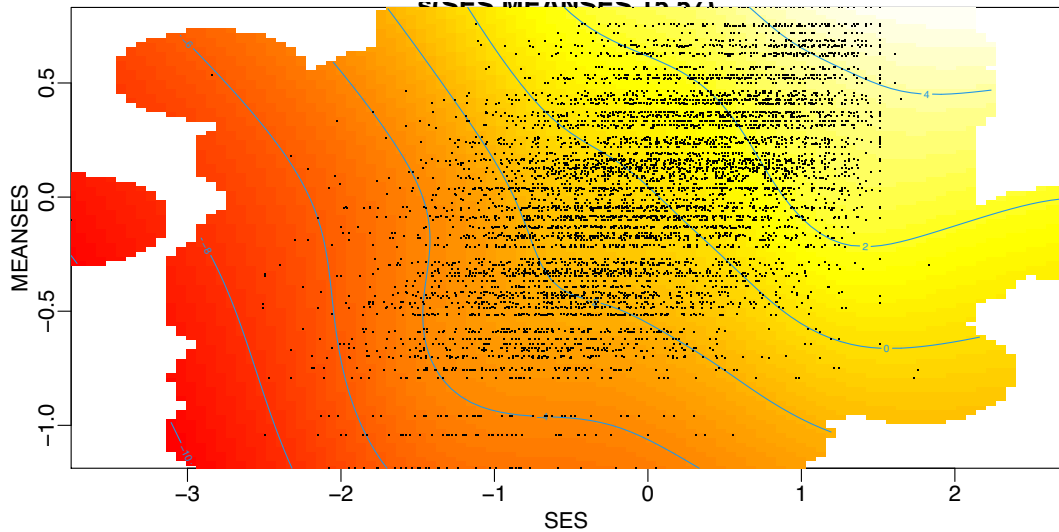
- two dimensional smoothing
- penalize curvature
- 2d b-splines

*school average*



```
> mathGam2 = gam(MathAch ~ s(SES, MEANSES) + Minority *  
+   Sex, data = MathAchieve)  
> plot(mathGam2, scheme = 2, n2 = 100)
```

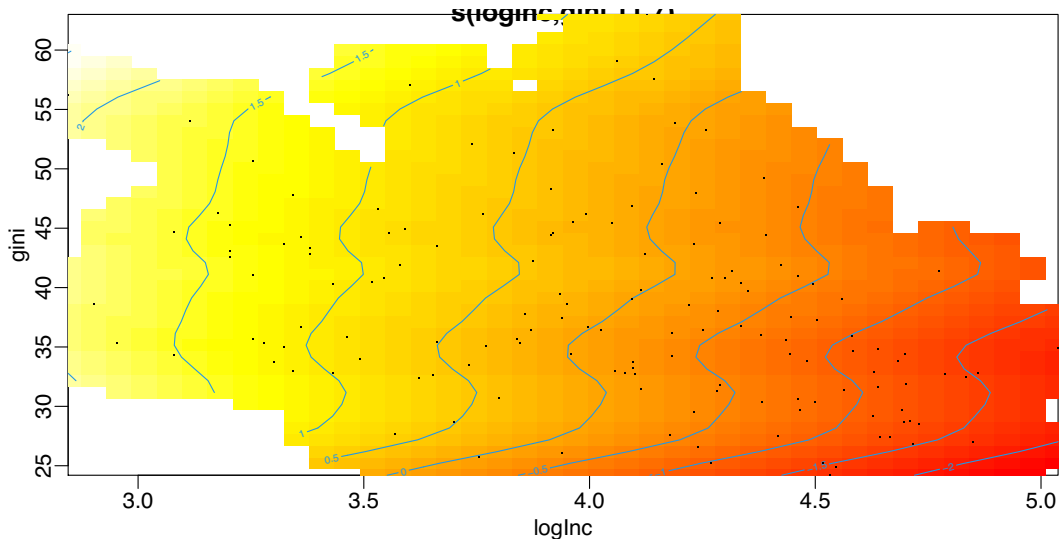
## Math 2d



## Infant mortality

```
> library("mgcv")
> iMort$logInc = log10(iMort$income)
> iMort$logMort = log(iMort$mortality)
> mortGam = gam(logMort ~ s(logInc, gini), data = iMort)
> plot(mortGam, scheme = 2)
```

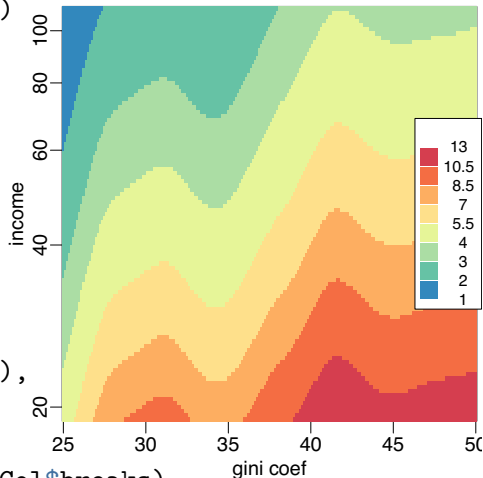
## Infant mortality





## Infant mortality

```
> predList = list(gini = seq(25,50,len=201), logInc = seq(
+   log10(19000), log10(110000), len=101))
> mortPred = exp(predict(mortGam,
+   do.call(expand.grid, predList),
+   type='response'))
> mortCol = mapmisc::colourScale(
+   mortPred, digits=1.5, col='Spectral',
+   style='equal', transform=0.5,
+   breaks=9, rev=TRUE)
> image(predList$gini,
+   10^predList$logInc/1000,
+   matrix(mortPred, length(predList$gini),
+   length(predList$logInc)),
+   xlab = 'gini coef', ylab='income',
+   log='y', col=mortCol$col, breaks=mortCol$breaks)
> mapmisc::legendBreaks("right", mortCol, cex=0.8, inset=0)
```



## Deaths in Ontario

```
> timeOrigin = ISOdate(2000,1,1,0,0,0, tz='UTC')
```

```
> oDeaths$timeNumeric = as.numeric(
```

```
+   difftime(oDeaths$date, timeOrigin,
```

```
+   units='days'))
```

```
> oDeaths[c(1,100,200),
```

```
+   c('date', 'month', 'Value',
```

```
+   'timeNumeric')]
```

	date	month	Value	timeNumeric
159	1991-01-01	January	6542	-3287
3077	1999-04-01	April	6463	-275
6231	2007-08-01	August	6527	2769

- Monthly death count is Value
- The date variable is a fancy data type for times
- strptime creates date objects
- timeNumeric is number of days since 1 Jan 2000

## Ontario GAM

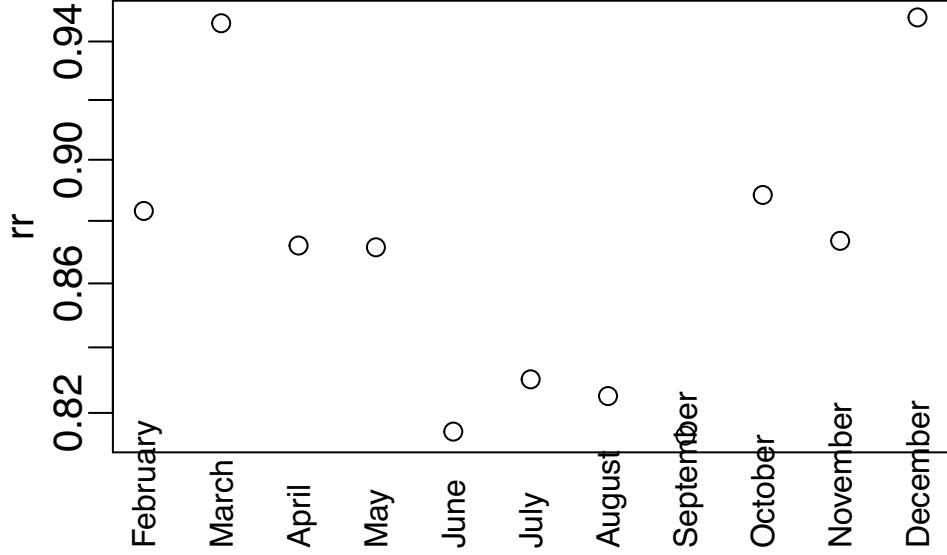
$$Y_i \sim \text{Poisson}(\lambda_i)$$

$$\log(\lambda_i) = X_i\beta + f(\text{time}_i)$$

```
> deathsGam = gam(  
+   Value ~ month + s(timeNumeric),  
+   data=oDeaths, family='poisson'  
+ )  
> knitr::kable(  
+   summary(deathsGam)$p.table[,1:2],  
+   digits=3, col.names=c('est', 'se'))
```

	est	se
(Intercept)	9.001	0.002
monthFebruary	-0.124	0.003
monthMarch	-0.055	0.003
monthApril	-0.137	0.003
monthMay	-0.138	0.003
monthJune	-0.205	0.003
monthJuly	-0.186	0.003
monthAugust	-0.192	0.003
monthSeptember	-0.207	0.003
monthOctober	-0.118	0.003
monthNovember	-0.135	0.003
monthDecember	-0.053	0.003

Relative rate for each month



## Number of days in each month

```
> oDeaths$daysInMonth = Hmisc::monthDays(oDeaths$date)
> oDeaths$nDays = log(oDeaths$daysInMonth)
> oDeaths[c(1, 2, 100, 200), c("date", "month",
+   "daysInMonth", "nDays")]
```

	date	month	daysInMonth	nDays
159	1991-01-01	January	31	3.433987
161	1991-02-01	February	28	3.332205
3077	1999-04-01	April	30	3.401197
6231	2007-08-01	August	31	3.433987

- calculate the number of days in each month
- the log number of days will be an offset
- rates modelled will be number of deaths per day

## Ontario GAM with offset

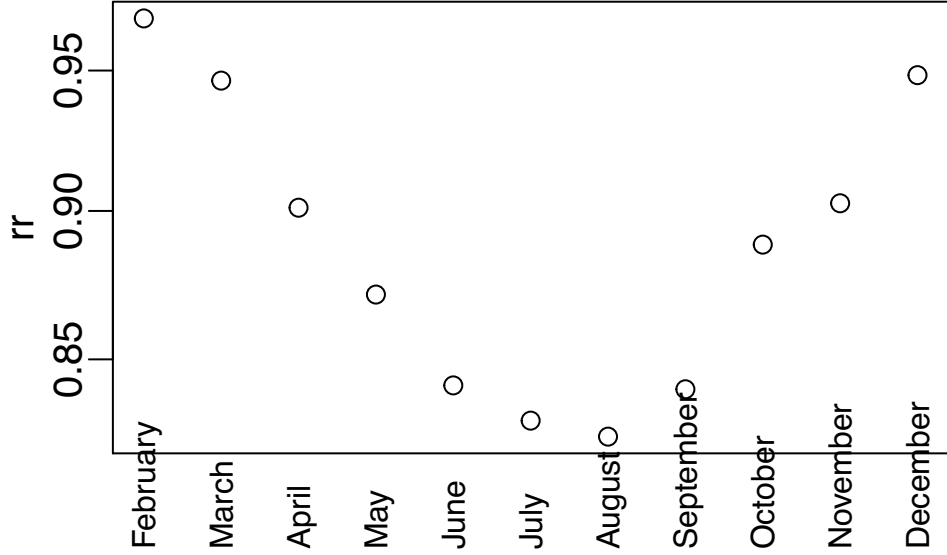
$$Y_i \sim \text{Poisson}(O_i \lambda_i)$$
$$\log(\lambda_i) = X_i \beta + f(\text{time}_i)$$

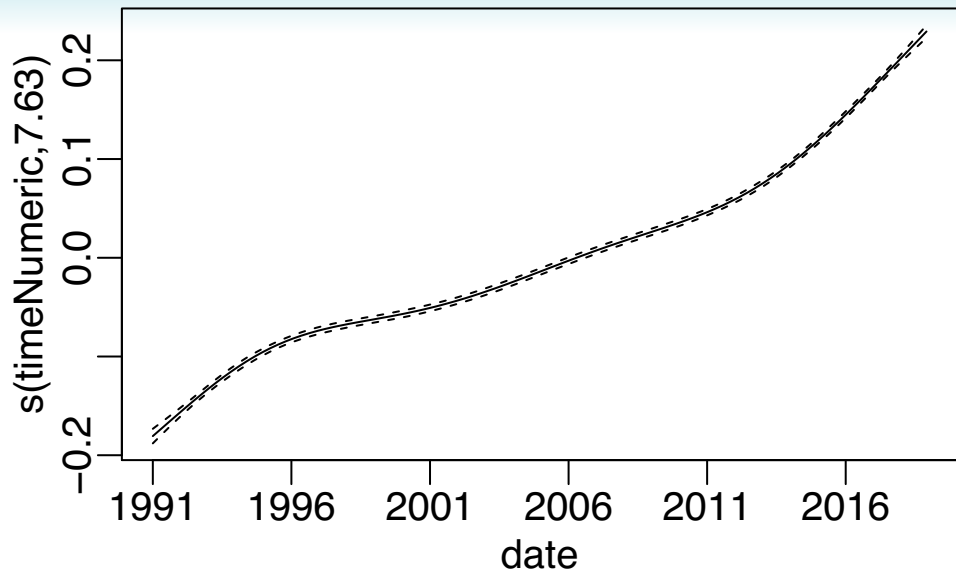
*O<sub>i</sub>: days in month*  
*λ<sub>i</sub>: deaths / day*

```
> deathsGam = gam(  
+   Value ~ month + s(timeNumeric) +  
+   offset(nDays), data=oDeaths,  
+   family='poisson')  
> knitr::kable(  
+   summary(deathsGam)$p.table[,1:2],  
+   digits=3, col.names=c('est', 'se'))
```

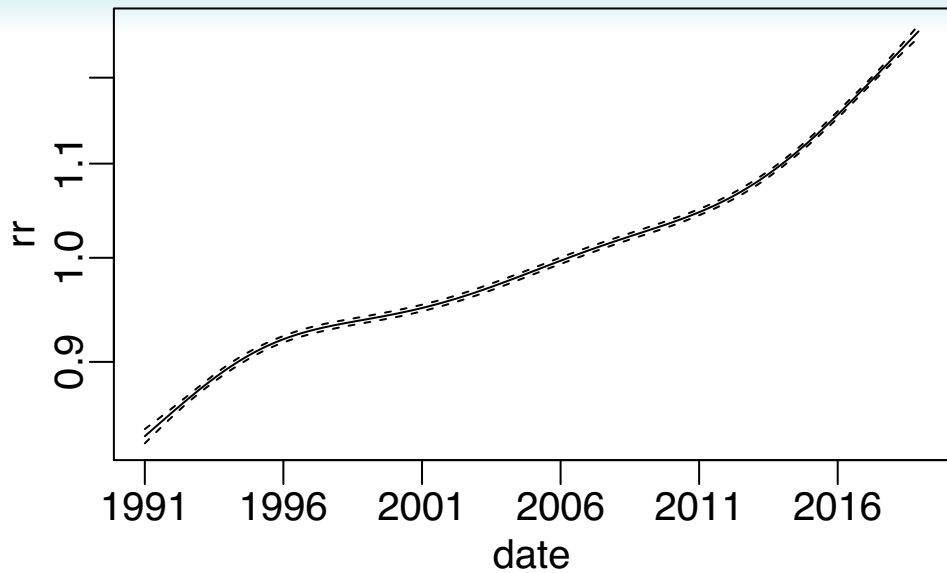
	est	se
(Intercept)	5.567	0.002
monthFebruary	-0.031	0.003
monthMarch	-0.055	0.003
monthApril	-0.104	0.003
monthMay	-0.138	0.003
monthJune	-0.173	0.003
monthJuly	-0.186	0.003
monthAugust	-0.192	0.003
monthSeptember	-0.174	0.003
monthOctober	-0.118	0.003
monthNovember	-0.102	0.003
monthDecember	-0.053	0.003

Relative rate for each month









```
> dSeq = seq(from = min(oDeaths$date), by = "5 years",
+   length.out = 10)
> deathPred = as.matrix(as.data.frame(predict.gam(deathsGam,
+   oDeaths, type = "terms", terms = "s(timeNumeric)",
+   se.fit = TRUE)))
> deathPred = exp(deathPred %*% Pmisc::ciMat())
> matplot(oDeaths$timeNumeric, deathPred, log = "y",
+   xaxt = "n", xlab = "date", type = "l", lty = c(1,
+   2, 2), col = "black", ylab = "rr")
> axis(1, at = difftime(dSeq, timeOrigin, units = "days"),
+   labels = format(dSeq, "%Y"))
```

## Forecasting

```
> newX = data.frame(date = seq(from = timeOrigin,  
+   by = "months", length.out = 12 * 21))  
> newX$timeNumeric = as.numeric(difftime(newX$date,  
+   timeOrigin, units = "days"))  
> newX$daysInMonth = Hmisc::monthDays(newX$date)  
> newX$nDays = log(newX$daysInMonth)  
> newX$month = months(newX$date)
```

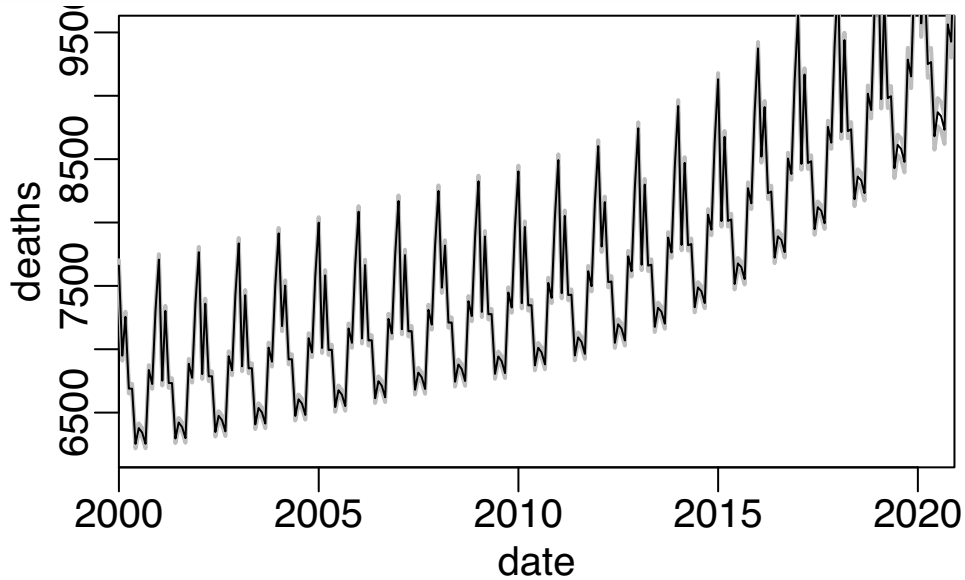
## Predictions

```
> deathsPred = predict(deathsGam, newX, se.fit = TRUE)
> deathsPred = cbind(newX, deathsPred)
> deathsPred$lower = deathsPred$fit - 2 * deathsPred$se.fit
> deathsPred$upper = deathsPred$fit + 2 * deathsPred$se.fit
> for (D in c("fit", "lower", "upper")) {
+   deathsPred[[paste(D, "exp", sep = "")]] = exp(deathsPred[[D]])
+ }
```

## Forecasts

```
> plot(deathsPred$date, deathsPred[, "fitexp"],  
+   type = "n", xlab = "date", ylab = "deaths",  
+   xaxs = "i", ylim = c(6200, 9500))  
>  
> matlines(deathsPred$date, deathsPred[, c("lowerexp",  
+   "upperexp", "fitexp")], lty = 1, col = c("grey",  
+   "grey", "black"), lwd = c(2, 2, 1))  
> points(oDeaths$date, oDeaths$Value, cex = 0.5,  
+   col = "red")
```

## Forecasts



## Favoured models

- Nancy E. Heckman and James O. Ramsay (2000). “Penalized Regression with Model-Based Penalties”. In: *The Canadian Journal of Statistics / La Revue Canadienne de Statistique* 28.2, pp. 241–258. URL: <http://www.jstor.org/stable/3315976>
- what happens when  $\alpha \rightarrow \infty$ ?
- what  $f$  has zero penalty?
- straight lines have  $f''(x) = 0$
- PLS with second derivative penalty encourages straight lines
- connections with Simpson et al. (2017)
- using PLS implicitly assumes  $f$  is almost straight

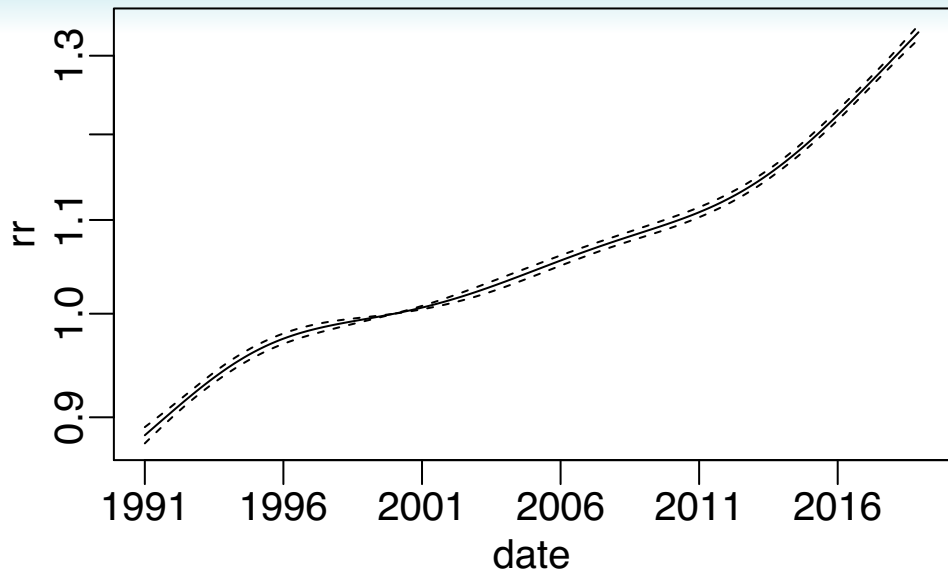
## a different constraint

- adding a constant to  $f(x)$  doesn't change the penalty
- ... any straight line has penalty zero
- can't identify both  $f(x)$  and the intercept without constraints
- a sum-to-zero constraint on  $f(x)$  is the default
- ... alternative is to force  $f(x_0) = 0$

```
> deathsGamC = gam(  
+   Value ~ month + s(timeNumeric, pc=0)  
+   offset(nDays), data=oDeaths,  
+   family='poisson')
```

	est	se
(Intercept)	5.510	0.003
monthFebruary	-0.031	0.003
monthMarch	-0.055	0.003
monthApril	-0.104	0.003
monthMay	-0.138	0.003
monthJune	-0.173	0.003
monthJuly	-0.186	0.003
monthAugust	-0.192	0.003
monthSeptember	-0.174	0.003
monthOctober	-0.118	0.003
monthNovember	-0.102	0.003
monthDecember	-0.053	0.003





```
> deathPredC = as.matrix(as.data.frame(predict.gam(deathsGamC,  
+   oDeaths, type = "terms", terms = "s(timeNumeric)",  
+   se.fit = TRUE)))  
> deathPredC = exp(deathPredC %*% Pmisc::ciMat())  
> matplot(oDeaths$timeNumeric, deathPredC, log = "y",  
+   xaxt = "n", xlab = "date", type = "l", lty = c(1,  
+   2, 2), col = "black", ylab = "rr")  
> axis(1, at = difftime(dSeq, timeOrigin, units = "days"),  
+   labels = format(dSeq, "%Y"))
```

## About GAM's

- They're easy to fit
- ...but inference is slightly unusual
  - Maximizing the joint probability  $pr(Y, U; \beta, \alpha)$
  - penalty from cross validation

### Bayesian inference

- put priors on  $\alpha, \beta$
- and compute posteriors
- ...with INLA (of course)

## Model for Ontario deaths

$$Y_i \sim \text{Poisson}(O_i \lambda_i)$$

$$\log(\lambda_i) = X_i \beta + U(t_i) + V_i$$

$$[U_1 \dots U_T]^T \sim \text{RW2}(0, \sigma_U^2)$$

$$V_i \sim \text{N}(0, \sigma_V^2) \text{ (independent)}$$

- $U(t)$  is a second-order random walk
- ...second derivatives are  $\text{N}(0, \sigma_U^2)$
- $V_i$  covers independent variation or over-dispersion

## Random Walks

- RW(0), independent

$$V_t \sim \text{iid } N(0, \tau^2)$$

- RW(1), Brownian motion

$$V_{t+1} | V_k, k < t \sim N(V_t, \tau^2)$$

$$V_{t+1} - V_t \sim N(0, \tau^2)$$

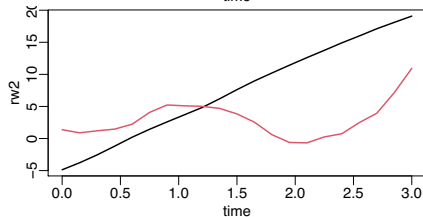
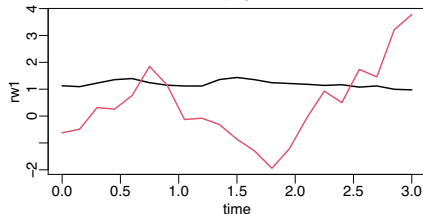
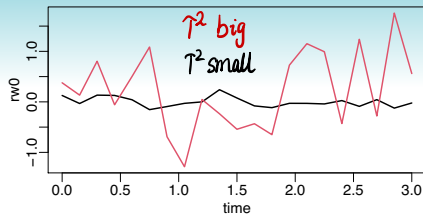
- RW(2), Random slope

$$V_{t+1} | V_k, k < t \sim N(-2V_t + V_{t-1}, \tau^2)$$

$$(V_{t+1} - V_t) - (V_t - V_{t-1}) \sim N(0, \tau^2)$$

$$V_{t+1} - 2V_t + V_{t-1} \sim N(0, \tau^2)$$

$$V_{t+1} - V_t \sim N(V_t - V_{t-1}, \tau^2)$$



## Favoured models and constraints

- $\alpha = \infty$  equivalent to  ~~$\alpha$~~  <sup>$\tau$</sup>  = 0
  - base model, favoured model
- RW(0):  $V_t = 0$  for all  $t$
- RW(1):  $V_t = V_0$ , horizontal line
- RW(2):  $V_t = V_0 + t \cdot (V_1 - V_0)$ , straight line
- RW(1) or RW(2)? favour straight line or horizontal line?
- PC prior: how strongly to encourage favoured model

## Preparing the date variable

- there are many ways to convert dates to numbers
- don't rely on `inla` to do the right thing
- R's Date objects are stored as number of days since 1 January 1970
- convert time from days to years, numerically more stable

```
> class(oDeaths$date)
```

```
[1] "Date"
```

```
> oDeaths$date[1:4]
```

```
[1] "1991-01-01" "1991-02-01" "1991-03-01" "1991-04-01"
```

```
> as.numeric(oDeaths$date)[1:4]
```

```
[1] 7670 7701 7729 7760
```

```
> oDeaths$timeId = oDeaths$timeRw = as.numeric(oDeaths$date)/365.25
```

```
> oDeaths$timeRw[1:4]
```

## Running INLA

- Prior for RW: slope of log rate changes by 0.2 from one year to the next
- `scale.model` defaults to TRUE, which multiplies  $\sigma$  by a constant that some people (not me) find more interpretable.

```
> library("INLA")
> deathsBayes = inla(
+   Value ~ offset(nDays) + month +
+   f(timeRw, model='rw2',
+     prior='pc.prec', param=c(0.2, 0.5),
+     scale.model=FALSE) + (small)
+   f(timeIid, model='iid',
+     prior='pc.prec', param=c(log(1.25), 0.5)),
+   data=oDeaths, family='poisson',
+   control.predictor=list(compute=TRUE))
```

$$Y_i \sim \text{Poisson}(O_i \lambda_i)$$

$$\log(\lambda_i) = X_i \beta + U_i + V_i$$

$$V_i \text{ independent } N(0, \tau^2)$$

$$U_i \sim \text{RW2}(\sigma^2)$$

Priors:

$$\beta \sim \text{MVN}(0, KI)$$

$$\tau \sim \text{Exp}, \text{ median } \log(1.25)$$

$$\sigma \sim \text{Exp}, \text{ median } 0.2$$

↓  
25% change is the prior  
median (a lot).

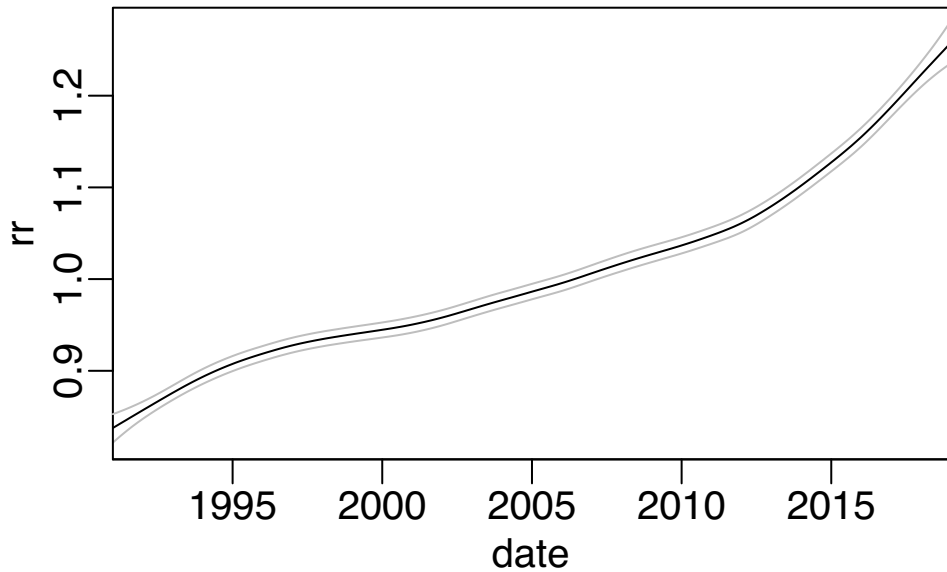


## Parameters

```
> Squant = paste0(c(0.5, 0.025, 0.975), "quant")  
> knitr::kable(deathsBayes$summary.fixed[, Squant],  
+   digits = 4)
```

	0.5quant	0.025quant	0.975quant
(Intercept)	5.5644	5.5530	5.5757
monthFebruary	-0.0299	-0.0461	-0.0138
monthMarch	-0.0528	-0.0689	-0.0367
monthApril	-0.1024	-0.1186	-0.0863
monthMay	-0.1350	-0.1511	-0.1189
monthJune	-0.1697	-0.1859	-0.1535
monthJuly	-0.1833	-0.1994	-0.1671
monthAugust	-0.1902	-0.2064	-0.1740
monthSeptember	-0.1719	-0.1881	-0.1557
monthOctober	-0.1159	-0.1320	-0.0997
monthNovember	-0.1002	-0.1164	-0.0841

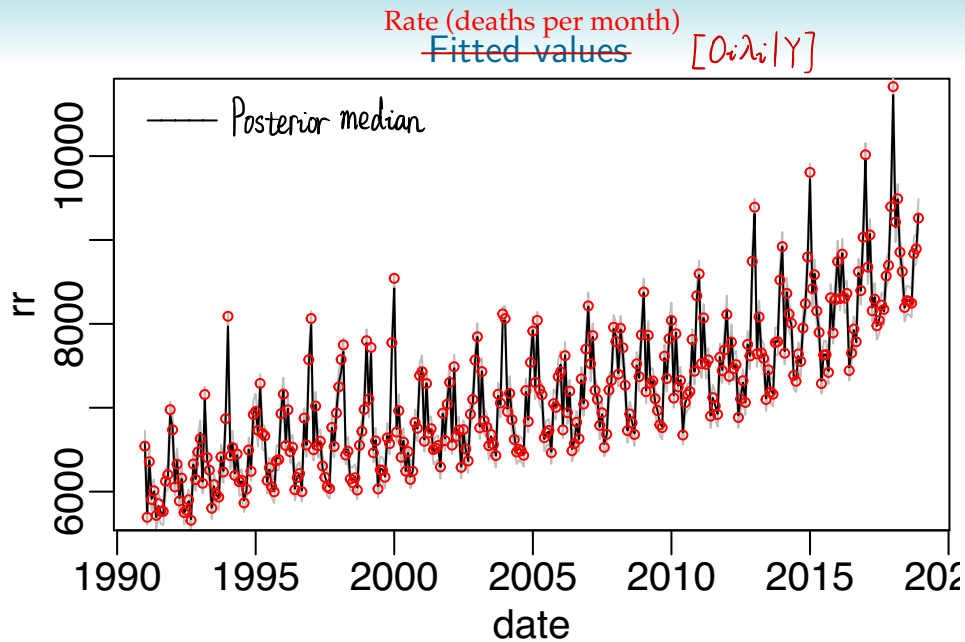
Time effect  $[u_i|Y]$



## Time effect

*X, in years since 1970*

```
> matplot(deathsBayes$summary.random$timeRw$ID,  
+   exp(deathsBayes$summary.random$timeRw[, c("0.025quant",  
+   "0.975quant", "0.5quant")])), xaxt = "n",  
+   xlab = "date", lty = 1, col = c("grey", "grey",  
+   "black"), type = "l", ylab = "rr", xaxs = "i")  
> Saxis = seq(as.Date("1990/1/1"), as.Date("2025/1/1"),  
+   by = "5 years")  
> axis(1, at = as.numeric(Saxis)/365.25, labels = format(Saxis,  
+   "%Y"))
```



## Fitted values

$$E(Y_i) = O_i \lambda_i$$

```
> matplot(oDeaths$date, deathsBayes$summary.fitted[,  
+   c("0.025quant", "0.975quant", "mean")], xaxt = "n",  
+   xlab = "date", lty = 1, col = c("grey", "grey",  
+   "black"), type = "l", ylab = "rr", yaxs = "i")  
> points(oDeaths$date, oDeaths$Value, cex = 0.5,  
+   col = "red")  
> axis(1, at = Saxis, labels = format(Saxis, "%Y"))
```

## Parameters

```
> deathsBayes$priorPost = Pmisc::priorPostSd(deathsBayes)
> knitr::kable(deathsBayes$priorPost$summary[, Squant],
+   digits = 4)
```

	0.5quant	0.025quant	0.975quant
SD for timeRw	0.0050	0.0026	0.0103
SD for timelid	0.0283	0.0258	0.0311

- How do we interpret Sd for timeRw?
- very hard, most people don't
- change in slope over one year, not surprising it's small

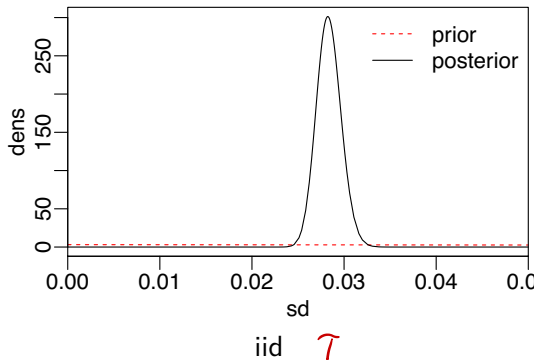
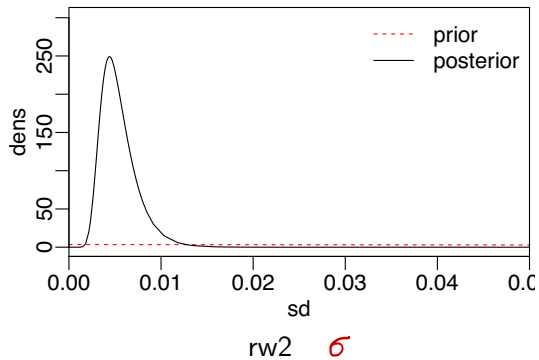
## Re-parametrized

```
> toPrint = diag(c(sqrt(25),1)) %*%  
+   as.matrix(deathsBayes$priorPost$summary[,  
+     c('mean','0.025quant','0.975quant')])  
> rownames(toPrint) = c('SD RW2, 25 yr', 'SD independent')  
> knitr::kable(toPrint, digits=3)
```

$\sqrt{250^2}$   
T

	mean	0.025quant	0.975quant
SD RW2, 25 yr	0.027	0.013	0.052
SD independent	0.028	0.026	0.031

## sd prior, post





## prior, post

```
> deathsBayes$priorPost[[1]]$matplot$ylim =  
+ deathsBayes$priorPost[[2]]$matplot$ylim =  
+   range(deathsBayes$priorPost[[2]]$matplot$y)  
> for(Dparam in 1:2){  
+   do.call(matplot,deathsBayes$priorPost[[Dparam]]$matplot)  
+   do.call(legend,deathsBayes$priorPost$legend)  
+ }
```

## Telling INLA we want forecasts

```
> newX = data.frame(  
+   date = seq(max(oDeaths$date),  
+     as.Date('2025/1/1'), by='month')[-1],  
+   Value = NA)  
> newX$timeId = newX$timeRw =  
+   as.numeric(newX$date)/365.25  
> newX$daysInMonth = Hmisc::monthDays(newX$date)  
> newX$nDays = log(newX$daysInMonth)  
> newX$month = months(newX$date)  
>  
> newData = rbind(oDeaths[,names(newX)], newX)
```

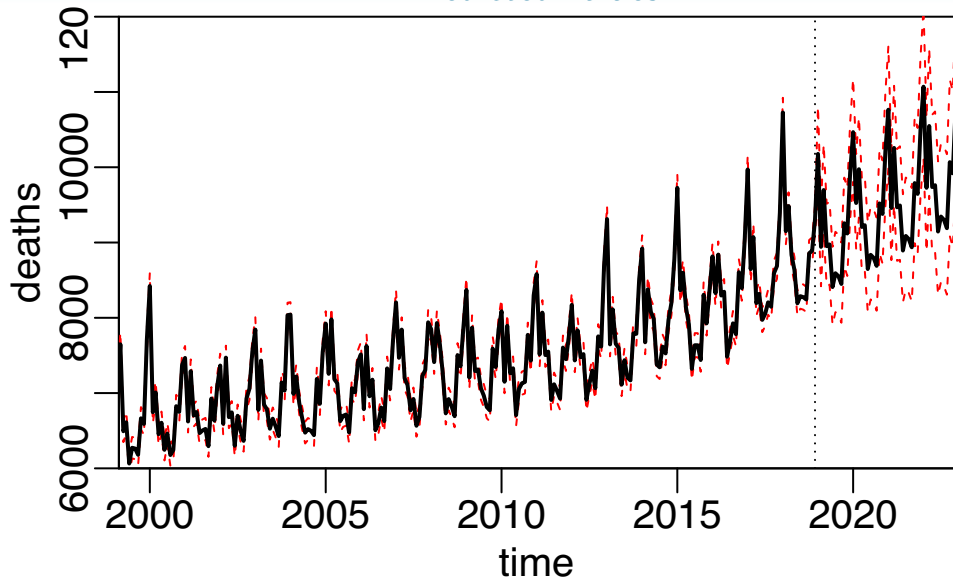
- Create a sequence of dates in a new data frame
- missing values for observed deaths
- convert dates to the values inla wants
- days in each month and name of month
- combine with the original data
- could have used `lincombs` instead

## Re-running with these new data

```
> deathsBayesRefit = inla(  
+   deathsBayes$.args$formula,  
+   data=newData, family='poisson',  
+   control.predictor=list(compute=TRUE, link=1))
```

- link=1 is a bizarre inla requirement
- it means use the link function from the first family
- ... note there's only one family

Predicted values



## Predicted values

```
> deathsPred = deathsBayesRefit$summary.fitted.values
> theXlim = as.Date(c("2000/1/1", "2022/1/1"))
> matplot(newData$date, deathsPred[, c("0.025quant",
+   "0.975quant", "0.5quant")], xlab = "time",
+   ylab = "deaths", xlim = theXlim, lwd = c(1,
+     1, 2), ylim = c(6000, 12000), type = "l",
+   lty = c(2, 2, 1), col = c("red", "red", "black"),
+   xaxt = "n", yaxs = "i")
> abline(v = as.numeric(max(oDeaths$date)), lty = 3)
> axis(1, at = Saxis, labels = format(Saxis, "%Y"))
```

## Posterior samples

$$Y_i \sim \text{Poisson}(O_i \lambda_i)$$

$$\log(\lambda_i) = X_i \beta + U(t_i) + V_i$$

$$[U_1 \dots U_T]^T \sim \text{RW2}(0, \sigma_U^2)$$

$$V_i \sim \text{N}(0, \sigma_V^2)$$

- we've been plotting the posterior medians and quantiles of  $\pi(U|Y)$
- we can take random samples of from the posterior of  $U$
- ... and plot those
- the method for doing this in `inla` isn't published, and is based on a Normal approximation.
- MCMC is an alternative with some advantages for this task

## Recompute

```
> deathsBayesC = inla(deathsBayes$.args$formula,  
+   data=newData, family='poisson',  
+   control.compute = list(config=TRUE),  
+   control.predictor=list(compute=TRUE, link=1))
```

```
> print(object.size(deathsBayesRefit), units='MB')
```

3.4 Mb

```
> print(object.size(deathsBayesC), units='MB')
```

16.4 Mb

- config means save a load of variance matrices
- which results in the new fitted object being much larger

## Posterior samples

- I want 64 samples from the posterior  $\pi(U(t)|Y)$
- run ~~16~~<sup>8</sup> threads in parallel, ~~4~~<sup>8</sup> samples per thread
- give me only the timeRw part, at all time points

```
> mySampleList = inla.posterior.sample(n = 64, result = deathsBayesC,  
+   num.threads = 8, selection = list(timeRw = seq(1,  
+   nrow(deathsBayesC$summary.random$timeRw))))
```

```
> length(mySampleList)
```

```
[1] 64
```

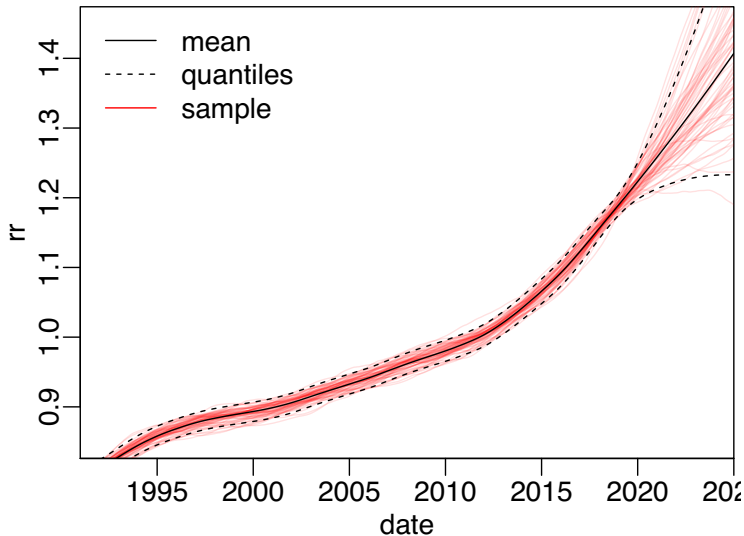
```
> mySample = do.call(cbind, Biobase::subListExtract(mySampleList,  
+   "latent"))
```

```
> dim(mySample)
```

```
[1] 409 64
```



## Plot of posterior sample $[\omega|\gamma]$



- samples are semi-transparent
- intervals are *pointwise*
- at each time point, interval contains 95% of samples
- can we get an envelope which contains 95% of samples?

## Plot of posterior sample

```
> matplot(newData$date, exp(mySample), xaxt = "n",  
+   xlab = "date", lty = 1, col = "#FF000020",  
+   ylim = c(0.85, 1.45), type = "l", ylab = "rr",  
+   xaxs = "i")  
> matlines(newData$date, exp(deathsBayesRefit$summary.random$timeRw[,  
+   Squant]), lty = c(1, 2, 2), col = "black")  
> axis(1, at = Saxis, labels = format(Saxis, "%Y"))  
> legend("topleft", bty = "n", lty = c(1, 2, 1),  
+   col = c("black", "black", "red"), legend = c("mean",  
+   "quantiles", "sample"))
```

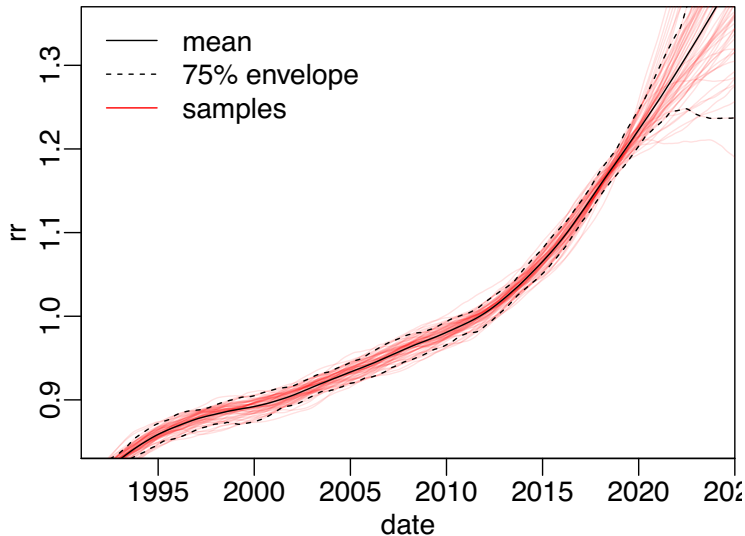
## Joint confidence regions

```
> cset <- GET::create_curve_set(list(  
+   r = as.numeric(newData$date),  
+   obs = mySample))  
> myEnv = GET::central_region(cset,  
+   coverage=0.75)
```

- global envelopes
- the GET package
- Myllymäki et al. (2017)
- this is *functional data analysis*, an active research topic

Mari Myllymäki et al. (2017). “Global envelope tests for spatial processes”. In: *Journal of the Royal Statistical Society: Series B (Statistical Methodology)* 79.2, pp. 381–404. DOI: 10.1111/rssb.12172

## Plot of posterior sample



- intervals are *global*
- 3/4 of samples are in the interval
- whatever the 'true' trend is, there's a 75% chance it's entirely contained within this envelope

## Plot of posterior sample

```
> matplot(newData$date, exp(mySample), xaxt = "n",  
+   xlab = "date", lty = 1, col = "#FF000020",  
+   ylim = c(0.85, 1.35), type = "l", ylab = "rr",  
+   xaxs = "i")  
> matlines(newData$date, exp(as.data.frame(myEnv)[,  
+   c("lo", "hi", "central")])), lty = c(2, 2,  
+   1), col = "black")  
> axis(1, at = Saxis, labels = format(Saxis, "%Y"))  
> legend("topleft", bty = "n", lty = c(1, 2, 1),  
+   col = c("black", "black", "red"), legend = c("mean",  
+   "75% envelope", "samples"))
```

## let's revisit the RW2 SD

```
> secondDiff = apply(mySample, 2,  
+   diff, differences=2)  
> dim(secondDiff)  
[1] 407  64  
> matplot(newData$date[-1],  
+   secondDiff, type='l',  
+   col='#FF000020', lty=1)
```

Error in matplot(newData\$date[-1], sec

## 1 year differences

```
> quantile(apply(secondDiff, 2, sd))
```

0%	25%	50%	75%	100%
9.065556e-05	1.297998e-04	1.396620e-04	1.708918e-04	2.742733e-04

```
> signif(sqrt(12) * quantile(apply(secondDiff, 2,  
+ sd)), 4)
```

0%	25%	50%	75%	100%
0.0003140	0.0004496	0.0004838	0.0005920	0.0009501

```
> Pmisc::priorPostSd(deathsBayes)$summary[1, Squant]
```

	0.5quant	0.025quant	0.975quant
SD for timeRw	0.005047952	0.00264209	0.01030015

- sd of secondDiff is monthly sd
- yearly var is 12 times monthly var
- agrees with parameter estimates

## Notes on INLA/Bayes

- INLA fit an RW2 model,
- not quite the same as the  $ARIMA(0,2,1)$
- and not a cubic spline
- so INLA is rougher, second derivatives are independent, not continuous



## Math

$$Y_{ijk} \sim \mathcal{N}(\mu_{ijk}, \tau^2)$$

$$\mu_{ijk} = X_{ijk}\beta + U_i + f_j(W_{ijk})$$

$$U_i \sim \mathcal{N}(0, \sigma_U^2)$$

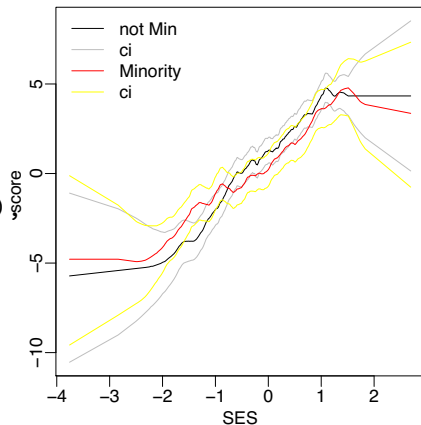
$$f_j(s+1) - f_j(s) \sim \mathcal{N}(0, \sigma_f^2)$$

- school  $i$ , ethnicity  $j$ , individual  $k$
- $W_{ijk}$  is the SES variable
- non-parameteric SES effect
  - `f(SES, model='rw1')`
- SES/ethnicity interaction
  - `f(SES, model='rw1', replicate = ethnicity)`

## Math

```
> MathAchieve$minorityNum = as.numeric(MathAchieve$Minority)
> mathInla = inla(MathAch~f(SES,
+   replicate=minorityNum,
+   model='rw1', prior='pc.prec',
+   param=c(1, 0.5), scale.model=FALSE
+ ) + Minority*Sex,
+ control.family = list(hyper = list(
+ prec=list(prior='pc.prec',param=c(1,0.5))))
+ data=MathAchieve)
```

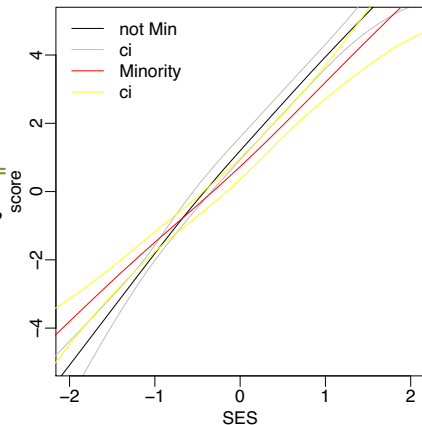
	mean	2.5	97.5
(Intercept)	11.7	11.3	12.0
MinorityYes	-2.5	-3.0	-1.9
SexMale	1.4	1.1	1.7
MinorityYes:SexMale	-0.1	-0.8	0.5



## Math, rw2

```
> mathInla2 = inla(  
+   MathAch ~ f(SES, replicate=minorityNum,  
+               model = 'rw2', prior='pc.prec',  
+               param=c(0.2, 0.5), scale.model=FALSE  
+   ) + Minority*Sex,  
+   control.inla = list(h=0.00001),  
+   control.family = list(hyper = list(prec=  
+ list(prior='pc.prec', param = c(1, 0.5)))),  
+   data=MathAchieve)
```

	mean	2.5	97.5
(Intercept)	11.7	11.4	12.0
MinorityYes	-2.5	-3.1	-2.0
SexMale	1.4	1.1	1.7
MinorityYes:SexMale	-0.1	-0.8	0.5

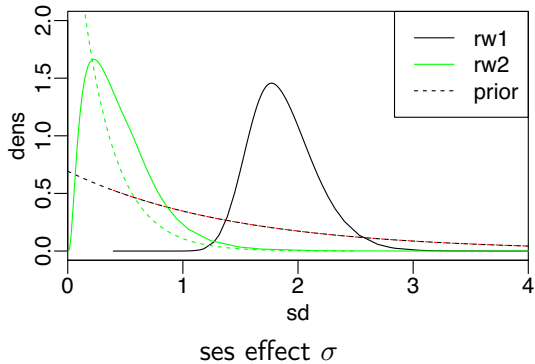
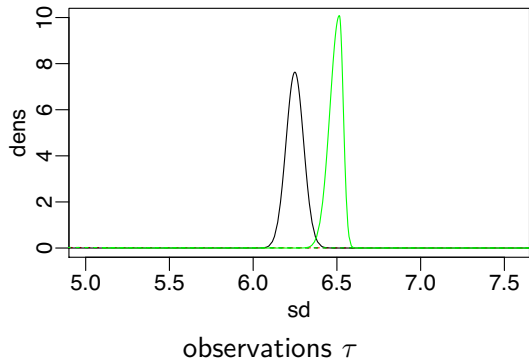


## Parameters

```
> mathInla$priorPost = Pmisc::priorPost(mathInla)
> mathInla2$priorPost = Pmisc::priorPost(mathInla2)
>
> knitr::kable(rbind(mathInla$priorPost[[1]]$summary[,
+   Squant], mathInla2$priorPost[[1]]$summary[,
+   Squant], mathInla$priorPost[[2]]$summary[,
+   Squant], mathInla2$priorPost[[2]]$summary[,
+   Squant]), digits = 5)
```

	0.5quant	0.025quant	0.975quant
SD for the Gaussian observations	6.24889	6.14644	6.35186
SD for the Gaussian observations1	6.49417	6.39819	6.55571
SD for SES	1.84044	1.38494	2.53273
SD for SES1	0.39571	0.07779	1.24123

## sd prior, post



## RW1 v RW2

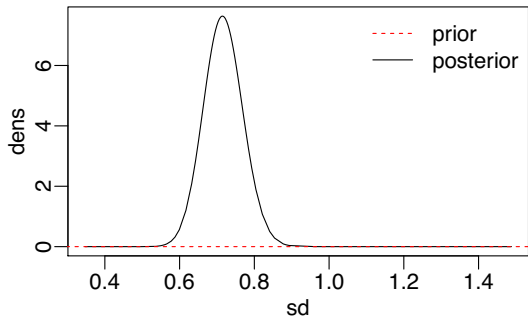
- The difference is what happens when  $\sigma = 0$
- RW1,  $U_i = U_0$
- RW2,  $U_i = U_0 + i \cdot (U_1 - U_0)$
- RW2 doesn't need to work to make straight lines
- RW1 needs large  $\sigma$  to capture SES effect
- Use RW2 if you regard linear regression as the 'default' or 'null'

## An inappropriate model for deaths

```
> deathsBayesUnwise = inla(
+   Value ~ month +
+   f(timeRw, model='rw2',
+     hyper = list(theta=list(
+       prior='pc.prec', scale.model=FALSE,
+       param=c(0.02, 0.05)))) +
+   offset(nDays),
+   data=newData,
+   family='poisson',
+   control.predictor = list(compute = TRUE, link=1)
+ )
```

- Eliminating the iid random effect
- Any variation in death counts about the mean is due to the Poisson
- Not consistent with flu outbreaks

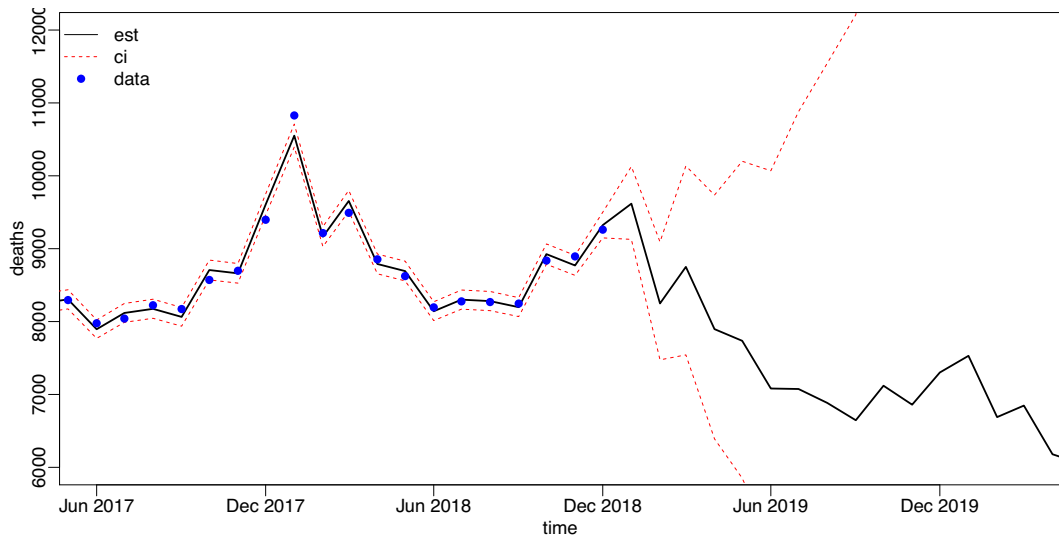
## prior, post



- The time effect will try to capture all extra-Poisson variation
- so the SD is much larger
- larger even than the iid SD was
- If we believe this model, deaths are very unstable over time



## Predicted values



## Constraints and identifiability

- Suppose  $U_{t+1} \sim N(U_t, \sigma^2)$ , an RW1
- and  $\eta_t = \mu + U_t$
- not identifiable
- $\mu_2 = \mu + 10000$  and  $V_t = U_t - 10000$
- $\gamma_t = \mu_2 + V_t$  has the same distribution as  $\eta_t$
- ... and  $V_t$  is also an RW1
- must *constrain*  $U_t$  for the model to be identifiable
- default in `inla` is  $\sum_t U_t = 0$
- interpret  $\mu$  as an average  $\eta_t$
- alternative is to set  $U_k = 0$  for some  $k$
- then  $\mu = \eta_k$

## Ontario deaths with constraint

a bit complicated, so not shown

## Generalized Additive Mixed Models

$$Y_{ij} \sim (\lambda_{ij}, \theta)$$

$$g(\lambda_{ij}) = X_{ij}\beta + f(W_{ij}) + U_i$$

$$[U_1 \dots U_M]^T \sim \text{MVN}(0, \Sigma)$$

- Random effects are possible in a semi-parametric model
- School level random effect should be in the math model
- with Bayes/INLA this is conceptually and computationally straightforward
- These models are possible, but far from rigorous, using penalized likelihood with cross validation.
- the `gamm` function alternates between `gam` and `glmmPQL`

# INLA v GAM

...or more accurately, joint prob + xv versus Bayes

## GAM's

- fast and seem less likely to break
- give the appearance of making few assumptions

## Bayes

- fast enough using INLA (or BayesX)
- explicit specification of assumptions
- rigorous inference methodology

# Bayes v GAM

## Anti-GAM

- Uncertainty and inference with GAM's much be approached with caution
- GAM's assumptions are implicit
- ...and are quite strong
- so GAM's probably don't always adequately reflect uncertainty in their estimates






## Pro-GAM

- Stability of GAM's make them a better sludgehammer than Bayes
- ...for prediction problems with Big Data
- Brieman would approve

## Arguments for semi-parametric Bayes

- Forces one to be specific about the model
- Cox would prefer Bayes to GAM, provided you approach it properly
  - Think carefully about the model before you start, because an overly complex model will break
  - Use scientific knowledge to decide where to go non-parametric, before looking at the data
  - If you use models which the data don't support, your results should be crazy!
- Bayes is more extensible
  - Spatial correlations
  - Censored survival times

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