

# Methods of Applied Statistics I

STA2101H F LEC9101

Week 11


November 26 2020



The Alan Turing Institute  @turinginst · Nov 20

Don't miss: Prof Neil Lawrence, Senior #AI Fellow at @turinginst & DeepMind Professor of #MachineLearning at @Cambridge\_Uni - our final #TuringCovidConf keynote speaker.

@lawrennd will discuss #policy, science and the convening power of #data

 [turing.ac.uk/CovidConference](https://turing.ac.uk/CovidConference)

#COVID19

AI and data science in the age of COVID-19

The Alan Turing Institute



**Professor Neil Lawrence**  
DeepMind Professor of Machine Learning, University of Cambridge and Senior AI Fellow, The Alan Turing Institute

Policy, science and the convening power of data

[turing.ac.uk/CovidConference](https://turing.ac.uk/CovidConference) **15:00 – 15:45** **24 Nov 2020**

Cambridge University and 6 others

  20  31 



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↻ 20

♥ 31



“statisticians have a feel for the data that is crucial”

1. HW3 due December 3
  2. Nonparametric regression continued
  3. Prediction and Explanation
  4. Strategies for Modelling
  5. In the News
- November 30 15.00 – 16.00 Tyler McCormick
  - <https://canssiontario.utoronto.ca/?mec-events>
  - “Identifying the latent space geometry of network models through analysis of curvature”
  - Dec. 4 noon – Kathryn Roeder
  - Dec. 7 – Margaret Roberts
  - Dec. 14 – Kosuke Imai



# Recap

- Generalized linear models theory
- flexible modelling via expected value and variance function
- inference using theory of likelihood functions
  
- nonparametric regression
- local averaging
- local polynomial regression
- kernel function (for averaging)
- bandwidth (for local)

local least squares

$$\mathbb{E}(y_i) = \mu_i; \quad g(\mu_i) = \mathbf{x}_i^T \boldsymbol{\beta}; \quad \text{Var}(y_i) = \phi_i \mathbf{V}(\mu_i) \quad \phi_i = \mathbf{a}_i \boldsymbol{\phi}$$

$$\hat{\beta} = (X^T W^{-1} X)^{-1} X^T W z$$

iteration

$$\hat{\beta}^{(t+1)} = (X^T W^{(t)} X)^{-1} X^T W^{(t)} z^{(t)} \quad z^{(t)} = X \hat{\beta}^{(t)} + W^{-1(t)} u^{(t)};$$

$$W^{(t)} = W(\hat{\beta}^{(t)}), \quad u^{(t)} = u(\hat{\beta}^{(t)})$$

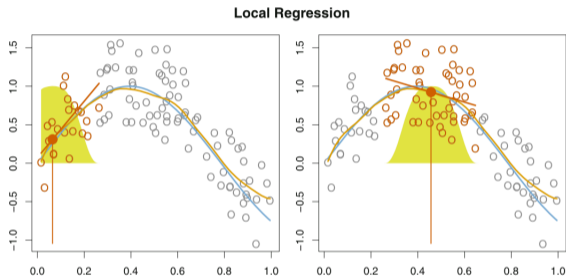
At convergence,

$$\hat{\beta} = (X^T \hat{W} X)^{-1} X^T \hat{z}$$

$$\widehat{\text{Var}}(\hat{\beta}) \doteq (X^T \hat{W} X)^{-1}$$

$W$  is diagonal

$$W_{ii} = \frac{1}{\phi a_i \{g'(\mu_i)\}^2 V(\mu_i)}, \quad u_i = \frac{y_i - \mu_i}{\phi a_i g'(\mu_i) V(\mu_i)}$$



**FIGURE 7.9.** Local regression illustrated on some simulated data, where the blue curve represents  $f(x)$  from which the data were generated, and the light orange curve corresponds to the local regression estimate  $\hat{f}(x)$ . The orange colored points are local to the target point  $x_0$ , represented by the orange vertical line. The yellow bell-shape superimposed on the plot indicates weights assigned to each point, decreasing to zero with distance from the target point. The fit  $\hat{f}(x_0)$  at  $x_0$  is obtained by fitting a weighted linear regression (orange line segment), and using the fitted value at  $x_0$  (orange solid dot) as the estimate  $\hat{f}(x_0)$ .

- $\hat{\beta} = (X^T W X)^{-1} X^T W y$        $W = \text{diag}(w_1, \dots, w_n)$   
weights from kernel function Nov 19 Slide 9
- $\hat{f}_\lambda(x_0) = \hat{\beta}_0 = \sum_{i=1}^n S(x_0; x_i, \lambda) y_i$       fit a poly, use only the intercept
- $S(x_0; x_1, \lambda), \dots, S(x_0; x_n, \lambda)$  first row of “hat” matrix  $(X^T W X)^{-1} X^T W$
- this makes it relatively easy to analyse the behaviour of local polynomial smoothers  
SM §10.7
- and to simplify the expression for the cross-validation criterion  $CV(\lambda)$
- fitting at each sample value gives

$$\hat{f}_\lambda(x_i) = \sum_{j=1}^n S(x_i; x_j, \lambda) y_j$$

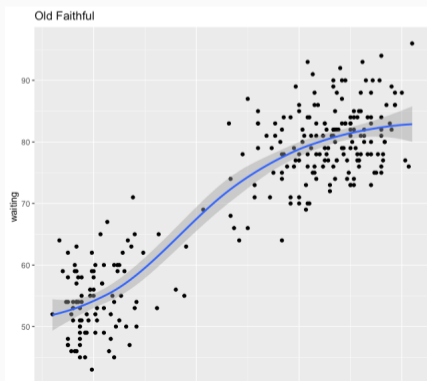
- model:  $y_i = f(x_i) + \epsilon_i$ ,  $i = 1, \dots, n$ ;  $E(\epsilon_i) = 0$ ;  $\text{var}(\epsilon_i) = \sigma^2$
- $\hat{f}_\lambda(x_0) = \hat{\beta}_0 = \sum_{i=1}^n S(x_0; x_i, \lambda) y_i$
- $E\{\hat{f}_\lambda(x_0)\} =$
- $\text{var}\{\hat{f}_\lambda(x_0)\} =$
- how many parameters did we fit?
- by analogy with least squares, estimates of 'degrees of freedom' are  $\nu_1 = \text{tr}(S_\lambda)$ , or  $\nu_2 = \text{tr}(S_\lambda^T S_\lambda)$

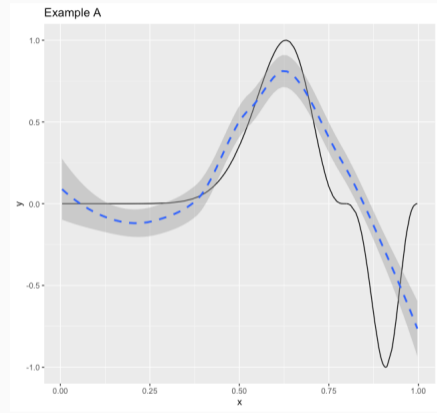
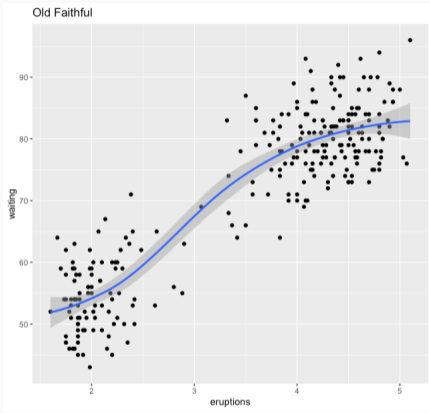
$$\tilde{\sigma}^2 = \frac{1}{n - 2\nu_1 + \nu_2} \sum \{y_i - \hat{f}_\lambda(x_i)\}^2$$



- $$E\{\hat{f}_\lambda(x_0)\} = \sum_{i=1}^n S(x_0; x_i, \lambda) f(x_i), \quad \text{var}\{\hat{f}_\lambda(x_0)\} = \sigma^2 \sum_{i=1}^n S^2(x_0; x_i, \lambda)$$

$$\frac{\hat{f}_\lambda(x_0) - E\{\hat{f}_\lambda(x_0)\}}{\widehat{\text{var}}\{\hat{f}_\lambda(x_0)\}^{1/2}} \sim N(0, 1)$$





*J. R. Statist. Soc. A* (2006)  
**169**, Part 2, pp. 179–203

## Model choice in time series studies of air pollution and mortality

Roger D. Peng, Francesca Dominici and Thomas A. Louis

*Johns Hopkins Bloomberg School of Public Health, Baltimore, USA*

[Received September 2004. Final revision July 2005]

### 4. National Morbidity, Morbidity, and Air Pollution Study data analysis

We apply our methods to the NMMAPS database which comprises daily time series of air pollution levels, weather variables and mortality counts. The original study examined data from 90 cities for the years 1987–1994 (Samet *et al.*, 2000a, b). The data have since been updated to include 10 more cities and six more years of data, extending the coverage until the year 2000. The entire database is available via the NMMAPSdata R package (Peng and Welty, 2004) which can be downloaded from the Internet-based health and air pollution surveillance system Web site at <http://www.ihapss.jhsph.edu/>.

The full model that is used in the analysis for this section is larger than the simpler model that was described in Section 3. We use an overdispersed Poisson model where, for a single city,

$$\begin{aligned} \log\{E(Y_t)\} = & \text{age-specific intercepts} + \text{day of week} + \beta \text{PM}_t + f(\text{time, df}) \\ & + s(\text{temp}_t, 6) + s(\text{temp}_{1-3}, 6) + s(\text{dewpoint}_t, 3) + s(\text{dewpoint}_{1-3}, 3). \end{aligned}$$

- 90 largest cities in US by population (US Census)
- daily mortality counts from National Center for Health Statistics 1987–1994
- hourly temperature and dewpoint data from National Climatic data Center
- data on pollutants  $PM_{10}$ ,  $O_3$ ,  $CO$ ,  $SO_2$ ,  $NO_2$  from EPA
  
- **response:**  $Y_t$  number of deaths on day  $t$
- **explanatory variables:**  $X_t$  pollution on day  $t - 1$ , plus various confounders: age and size of population, weather, day of the week, time
- mortality rates change with season, weather, changes in health status, ...

NMMAPS: National Morbidity, Mortality and Air Pollution Study

- $Y_t \sim \text{Poisson}(\mu_t)$
- $\log(\mu_t) = \text{age specific intercepts} + \beta PM_t + \gamma DOW + g(t, df) + s(\text{temp}_t, 6) + s(\text{temp}_{t-1}, 6) + s(\text{dewpoint}_t, 3) + s(\text{dewpoint}_{t-1}, 3) + s_4(\text{dew}_0, 3) + s_5(\text{dew}_{1-3}, 3)$
- three ages categories; separate intercept for each ( $< 65, 65 - 74, \geq 75$ )
- dummy variables to record day of week
- $s(x, 7)$  a **smoothing spline** of variable  $x$  with 7 degrees of freedom
- estimate of  $\beta$  for each city; estimates pooled using Bayesian arguments for an overall estimate
- very difficult to separate out weather and pollution effects

see also: Crainiceanu, C., Dominici, F. and Parmigiani, G. (2008). *Biometrika* **95** 635–51

- $y_i = f(x_i) + \epsilon_i$
- there are **many** different methods for estimating  $f(\cdot)$
- local polynomial regression – `stats::loess`, `KernSmooth::locpoly` FELM 11.3; SM 10.7.1
- regression splines – `splines::bs` , `splines::ns` FELM 11.2b p 218ff
- **smoothing splines** – `stats::smooth.spline` FELM 11.2a; SM 10.7.2
- penalized splines – `pspline::smooth.Pspline` Peng et al. 2006
- wavelets – `wavethresh::wd` FELM 11.4
- and more... FELM 11.5; ISLR Ch.7

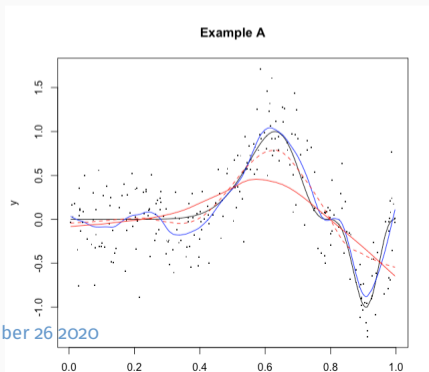
smoothing splines:

$$\min_f \sum_{i=1}^n \{y_i - f(x_i)\}^2 + \lambda \int \{f''(t)\}^2 dt$$

Elem Stat Learning, Hastie et al.

# Example

```
with(exa, plot(x,y, cex = 2, main = "Example A", pch = "."))  
lines(m ~ x, exa)  
lines(exa.sm1$x, exa.sm1$y, col="blue")  
lines(loess.smooth(exa$x, exa$y), col="red")  
lines(loess.smooth(exa$x, exa$y, span = 1/3), col="red", lty = 2)
```



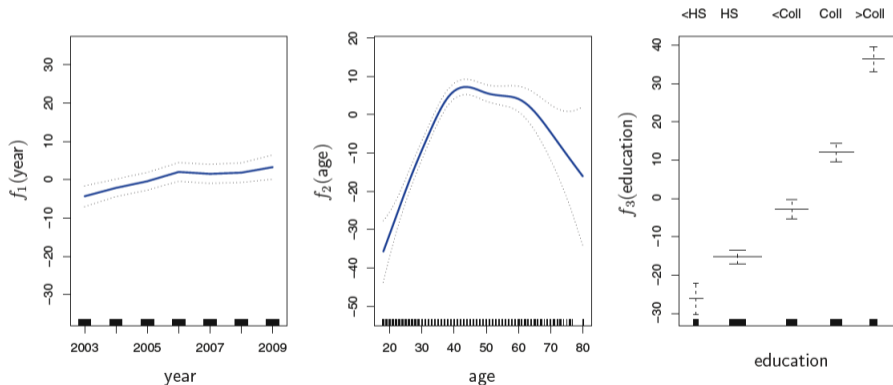
⇒ smooth.R

- more than 1 x
- local polynomials in two (or three) dimensions difficult to view/fit
- thin-plate splines  $d = 2, 3$
- **additive models:**  $y_i = \beta_0 + f_1(x_{1i}) + f_2(x_{2i}) + \dots + f_p(x_{pi}) + \epsilon_i$  FELM Ch.12  
`gam::gam` and `mgcv::gam`
- beyond least squares
- **GAM:** e.g.  $\log\{p_i/(1 - p_i)\} = \beta_0 + f_1(x_{1i}) + f_2(x_{2i}) + \dots + f_p(x_{pi})$  ISLR 7.7
- penalized generalized linear models
- e.g. linear predictor  $\eta_i = \mathbf{x}_i^T \beta + f(t_i)$  SM 10.7.3



smooth.html

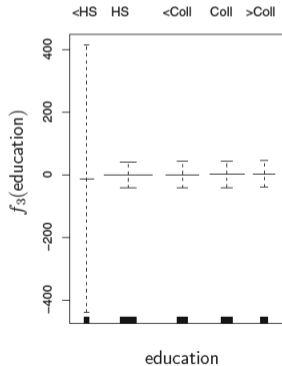
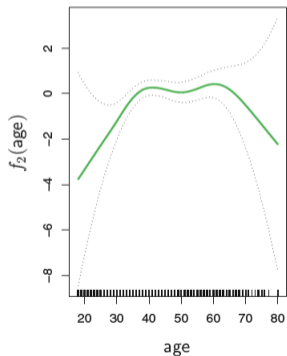
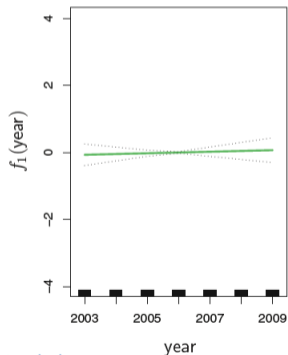
## 7.7 Generalized Additive Models 285



**FIGURE 7.12** Details are as in Figure 7.11, but now  $f_1$  and  $f_2$  are smoothing splines with four and five degrees of freedom, respectively.

`smooth.html`

## 7.8 Lab: Non-linear Modeling 287



## Aside: Explanation vs Prediction

- regression (and other) models may be fit in order to uncover some structural relationship between the response and one or more predictors
  - How do wages depend on education?
  - How does numeracy score affect probability of saying yes to vaccine?
- statistical analysis will focus on **estimation** and/or **testing**
- it is a remarkable fact that the data provides both an **estimate** of a model parameter **and** an estimate of uncertainty
  
- the focus might instead be on predicting responses for new values of  $x$
- or classifying new observations on the basis of their  $x$  values
- the statistical analysis will focus on the **accuracy and precision** of the prediction/classification
- the data used to fit the model **does not** provide a good assessment of the prediction or classification error — motivates the division of data into training and test sets

- in many fields of study the models used as a basis for interpretation do not have a special subject-matter base
- rather represent broad patterns of haphazard variation quite widely seen
- this is typically combined with a specification of the systematic part of the variation
- which is often the primary focus
- modelling then often reduces to a choice of distributional form
- and of the independence structure of the random components

- functional form of the probability distribution sometimes critical, for example where an implicit assumption is involved of a relationship between variance and mean: geometric, Poisson, binomial
- the simple situations that give rise to binomial, Poisson, geometric, exponential, normal and log normal are some guide to empirical model choice in more complex situations
- In some specific contexts there is a tradition establishing the form of model
- illustration: financial time series –  $Y(t) = \log\{P(t)/P(t-1)\}$  has a long-tailed distribution, small serial correlation, large serial correlation in  $Y^2(t)$  stock price
- often have a long tail of large values; exponential distribution is a natural starting point
- extensions may be needed, including Weibull, gamma or log-normal

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- often helpful to develop random and **systematic** parts of the model separately
- models should obey natural or known constraints, even if these lie outside the range of the data
- example  $P(Y = 1) = \alpha + \beta x \implies \log \frac{P(Y = 1)}{P(Y = 0)} = \alpha' + \beta' x$
- however,  $\beta$  measures the change in probability per unit change in  $x$   $\beta'$  does not
- when relationship between  $y$  and several variables  $x_1, \dots, x_p$  is of interest
  - unlikely that the system is wholly linear
  - impractical to study nonlinear systems of unknown form
  - therefore reasonable to begin with a linear model
  - and seek isolated nonlinearities

- often helpful to develop **random** and **systematic** parts of the model separately
- naive approach: one random variable per study individual
- values for different individuals independent
  
- more realistic: possibility of structure in the random variation
- dependence in time or space, or a hierarchical structure corresponding to levels of aggregation
- ignoring these complications may give misleading assessments of precision, or bias the conclusions



- example: standard error of mean  $\sigma/\sqrt{n}$
- but, under mutual correlation, becomes  $(\sigma/\sqrt{n})(1 + \sum \rho_{ij})^{1/2}$
- if each observation correlated with  $k$  others, at same level,  
 $(\sigma/\sqrt{n})(1 + k\rho)^{1/2}$

0.1 0.2 0.4 0.8

-----  
 1.14 1.26 1.48 1.84

1.18 1.34 1.61 2.05

1.22 1.41 1.73 2.24

1.26 1.48 1.84 2.41

1.30 1.55 1.95 2.57

1.34 1.61 2.05 2.72

$k = 3 : 8$

- important to be explicit about the unit of analysis
- has a bearing on independence assumptions involved in model formulation
- example: if all patients in the same clinic receive the same treatment
- then the clinic is the unit of analysis
  
- in some contexts there may be a clear hierarchy
- assessment of precision comes primarily from comparisons **between** units
- modelling of variation **within** units is necessary only if of intrinsic interest
  
- when relatively complex responses are collected on each individual, the simplest way of condensing these is through a number of **summary descriptive measures**
- in other situations it **may** be necessary to represent explicitly the different hierarchies of variation

- “Toronto researchers have found a new way to speed up the body’s ability to rid itself of alcohol” [Globe & Mail, Nov 12](#)
- “Williams prof disavows own finding of mishandled GOP ballots” [Twitter C. Bergstrom](#)
- “The association between early career informal mentorship in academic collaborations and junior author performance” [Gelman’s blog](#)
- “What We Know About AstraZeneca’s Head-Scratching Vaccine Results” [NY Times Nov 24](#)
- “A very, very bad look for remdesivir” [Science, Nov 6](#)

**scientific** reports[Explore our content](#) ▾[Journal information](#) ▾[nature](#) > [scientific reports](#) > [articles](#) > [article](#)[Article](#) | [Open Access](#) | [Published: 12 November 2020](#)**Accelerated ethanol elimination via the lungs**

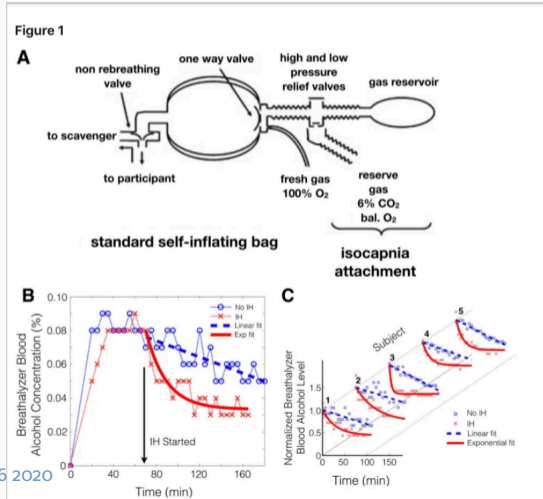
Jesse M. Klostranec, Diana Vucevic, Adrian P. Crawley, Lashmi Venkatraghavan, Olivia Sobczyk, James Duffin, Kevin Sam, Royce Holmes, Ludwik Fedorko, David J. Mikulis & Joseph A. Fisher [✉](#)

*Scientific Reports* **10**, Article number: 19249 (2020) | [Cite this article](#)

**17k** Accesses | **561** Altmetric | [Metrics](#)

**Abstract**

Link



## Link

 **Carl T. Bergstrom**  @CT\_Bergstrom · 20h

How it started . . . . . How it's going

### In sworn statement, prominent mathematician flags up to 100,000 Pennsylvania ballots

Federal Elections Commission Chairman Trey Trainor says new analysis by professor Steven Miller "adds to the conclusions that some level of voter fraud took place in this year's election."



TOP STORY

### Williams prof disavows own finding of mishandled GOP ballots

By Francesca Paris, The Berkshire Eagle 16 hrs ago



Steven Miller, a professor of mathematics at Williams College, issued a statement Monday in which he apologized for a "lack of clarity and due diligence" after his statistical analysis of Pennsylvania mail-in votes was used by Pennsylvania lawmakers to push a disputed claim of voter

 4  12  56 

 **Carl T. Bergstrom**  @CT\_Bergstrom · Nov 24

You know? Instead of doubling down, he corrected the record when he made the mistake. I give him credit for that.

 2   37 

- “In a signed affidavit, Miller claimed to show that more than 89,000 ballots requested by Pennsylvania Republicans either were not counted by the state or requested by someone other than the registered Republican [Link](#)
- “He used data provided by former Donald Trump campaign staffer Matt Braynard.
- “Election officials have called the voting process secure
- “Miller told The Eagle that he made a mistake separating his analysis of the data from questions about the reliability of the data itself.
- “Braynard collected the data by contracting call centers to get in touch with Republican voters across six swing state
- “In his analysis, Miller wrote that the group called 20,000 Republican voters in Pennsylvania who, according to state records, had requested but not returned ballots. In all, 2,684 agreed to answer questions
- “Of the respondents, 463 reported that they actually had mailed in a ballot and 556 reported that they had not requested a ballot ... Miller extrapolated from those numbers
- ‘To apply naive statistical formulas to biased data and publish this is both irresponsible and unethical,’ De Veaux wrote in a statement

## Link

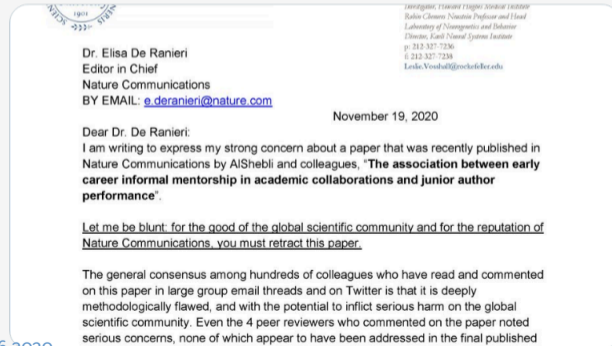
 Daniela Witten Retweeted



**Leslie Vosshall PhD** @polyp1 · Nov 19



Open letter to the Editor-in-Chief of @NatureComms about the AIShebli paper, which claims that training with #WomenInSTEM damages the careers of young scientists





[Link](#) to Gelman's blog post

Thanks to Emma, here's a link to a re-analysis of the data that points out several problems with it. [Link](#)



## Link



Carl T. Bergstrom  @CT\_Bergstrom · 19h ...

Bullshit.

[nytimes.com/2020/11/24/hea...](https://nytimes.com/2020/11/24/hea...)

**Why did the researchers test two different doses?**

It was a lucky mistake. Researchers in Britain had been meaning to give volunteers the initial dose at full strength, but they made a miscalculation and accidentally gave it at half strength, [Reuters reported](#). After discovering the error, the researchers gave each affected participant the full strength booster shot as planned about a month later.

 24  55  351 

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Carl T. Bergstrom  @CT\_Bergstrom · 19h ...

Seriously, why is the @NYTimes spinning this as a lucky mistake instead of a monumental fuck-up?

 21  23  427 

[Link to Science article](#) [Link to BMJ article](#)



President Donald Trump and FDA Commissioner Stephen Hahn (right) met with Gilead CEO Daniel O'Day (left) after remdesivir received an emergency use authorization.

COVID-19

## 'A very, very bad look' for remdesivir

FDA and Europe anointed it as a key therapy just after a major study found it has little value

Downloaded from