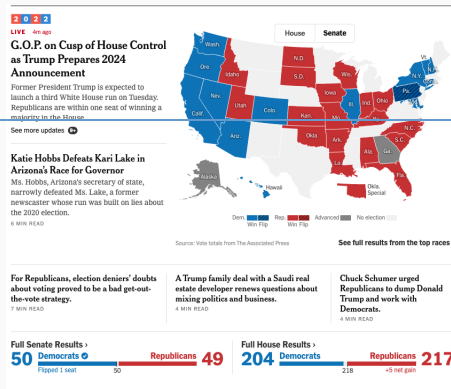


Methods of Applied Statistics I

STA2101H F LEC9101

Week 9

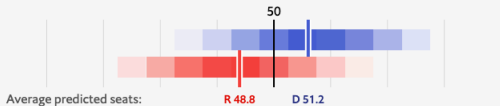
November 16 2022



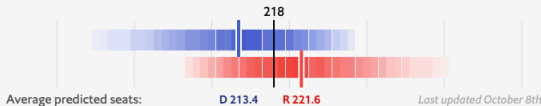
Predictions

The Economist's 2022 midterms model

→ **Senate** The Democrats are likely to keep their majority in the Senate



→ **House** The Republicans are slightly favoured to gain a majority in the House



[Click to view our full forecast for the midterm elections](#)

2 0 2 2

LIVE 4m ago

G.O.P. on Cusp of House Control as Trump Prepares 2024 Announcement

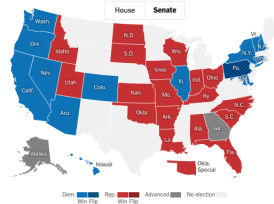
Former President Trump is expected to launch a third White House run on Tuesday. Republicans are within one seat of winning a majority in the House.

[See more updates](#)

Katie Hobbs Defeats Kari Lake in Arizona's Race for Governor

Ms. Hobbs, Arizona's secretary of state, narrowly defeated Ms. Lake, a former newscaster whose run was built on lies about the 2020 election.

6 MIN READ



[See full results from the top races](#)

For Republicans, election deniers' doubts about voting proved to be a bad get-out-the-vote strategy.

7 MIN READ

A Trump family deal with a Saudi real estate developer renews questions about mixing politics and business.

4 MIN READ

Chuck Schumer urged Republicans to dump Donald Trump and work with Democrats.

4 MIN READ

Full Senate Results >

50 Democrats

49 Republicans

Flipped 1 seat

Full House Results >

204 Democrats

217 Republicans

+5 net gain

And see "Land doesn't vote"

twitter.com/simongerman600/status/1591175192834965515?s=12

1. Upcoming events
2. Reminder re Project
3. Recap
4. Finish generalized linear models
5. Survival data

Project Guidelines

STA 2101F: Methods of Applied Statistics I 2022

Outline

- Part I 3-5 pages, non-technical 12 point type, 1.5 vertical spacing, thank you
 1. a description of the scientific problem of interest
 2. how (and why) the data being analyzed was collected
 3. preliminary description of the data (plots and tables)
 4. non-technical summary for a non-statistician of the analysis and conclusions
- Part II 3-5 pages, technical LaTeX or R markdown; submit .Rmd and .pdf files
 1. models and analysis
 2. summary for a statistician of the analysis and conclusions
- Part III Appendix submit .Rmd and .pdf or .html files

R script or .Rmd file; additional plots; additional analysis; References

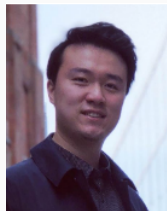
Project Marking

- 40 points total
- Part I:
description of data and scientific problem 5
suitability of plots and tables 5 clear, non-technical, concise but thorough
quality of the presentation 5
- Part II:
summary of the modelling and methods 5 justification for choices
suitability and thoroughness of the analysis 10 model checks, data checks
- Part III:
relevance of additional material 5
complete and reproducible submission 5

- November 17 3.30-4.30 Statistical Sciences Seminar
Room 9014, Hydro Building
and [online](#)

Qiongshi Li, U Wisconsin

“New Advances in Genetic Risk Prediction”



- November 18 12.00-1.00 Toronto Data Workshop
UY 9195 and [zoom](#)
Lindsay Katz, U Toronto
“... a new comprehensive database of all proceedings of Australian Parliamentary debates”

[link](#)

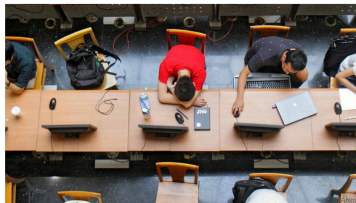
We are currently hiring a Statistical Programmer for our client, a global pharmaceutical company and one of the largest pharmaceutical companies in the world. The company is known for its success in researching developing and marketing innovative drugs.

The Statistical Programmer will be responsible for the development of SAS programs and statistical output for the management and reporting of clinical trial data managed by the Unit. Ensure quality of statistical output produced by external provider, programs tools to support data review activities and data visualization and collaborate on the interpretation and communication of trial results. ...

Science & technology | Exam nerves' real effect

Anxiety does not cause bad results in exams

The problem is in the run-up, not the main event



Nov 2nd 2022

Save

Share

Give

“Anxiety on the day of the test did not predict exam performance at all. ... What actually hampered students, it turned out, were high levels of anxiety during the weeks before the exam took place.”

Economist



@emitanaka@fosstodon.org @statsgen · Nov 12

...

I highly recommend watching this talk by [@monjalexander](#) if you are interested in modern demography research or interesting use of facebook ad data. As someone who knows very little about demography, this was a great intro and an interesting application!



youtube.com

Toronto Data Workshop - Monica Alexander - Usin...
Talk on 3 December 2020 by Professor Monica
Alexander, Statistical Sciences and Sociology, ...

[YouTube Link](#)

Recap

- brief review of likelihood theory
- overdispersion; troutegg example; Pearson's χ^2 ; beta-binomial mgcv::betar
- measures of risk: odds ratio, risk ratio, risk difference; prospective and retrospective sampling
- binary responses: non-convergence; latent variables
- generalized linear models: families, density, linear predictor, link function, variance function

Generalized linear models

glm has several options for family

```
binomial(link = "logit")  
gaussian(link = "identity")  
Gamma(link = "inverse")  
inverse.gaussian(link = "1/mu^2")  
poisson(link = "log")  
quasi(link = "identity", variance = "constant")  
quasibinomial(link = "logit")  
quasipoisson(link = "log")
```

Generalized linear models

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poisson(link = "log")
quasi(link = "identity", variance = "constant")
quasibinomial(link = "logit")
quasipoisson(link = "log")
```

Each of these is a member of the class of generalized linear models

Generalized: distribution of response is not assumed to be normal

Linear: some transformation of $E(y_i)$ is of the form $x_i^T \beta$

link function

`link`

a specification for the model link function. This can be a name/expression, a literal character string, a length-one character vector, or an object of class "link-glm" (such as generated by `make.link`) provided it is not specified via one of the standard names given next.

The gaussian family accepts the links (as names) identity, log and inverse;
the binomial family the links logit, probit, cauchit, (corresponding to logistic, normal and Cauchy CDFs respectively) log and cloglog (complementary log-log);
the Gamma family the links inverse, identity and log;
the poisson family the links log, identity, and sqrt;
and the inverse.gaussian family the links $1/\mu^2$, inverse, identity and log.

The quasi family accepts the links logit, probit, cloglog, identity, inverse, log

- density: $f(y_i; \mu_i, \phi_i) = \exp\left\{\frac{y_i\theta_i - b(\theta_i)}{\phi_i} + c(y_i; \phi_i)\right\}$

- moments: $E(y_i | x_i) = b'(\theta_i) = \mu_i \quad \text{Var}(y_i | x_i) = \phi_i b''(\theta_i) = \phi_i V(\mu_i)$

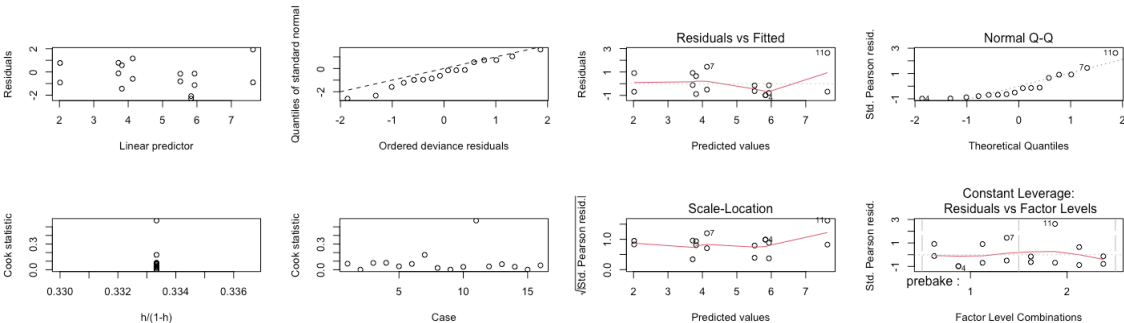
μ_i is a function of θ_i

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 μ_i is a function of θ_i
- link function: $g(\mu_i) = x_i^T \beta =$ links the n observations together via covariates
- linear predictor: $\eta_i = x_i^T \beta$ residual plots

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- linear predictor: $\eta_i = x_i^T \beta$ residual plots
- $\text{Var}(y_i | x_i) = \phi_i b''(\theta_i) = \phi_i V(\mu_i)$
- $V(\cdot)$ is the variance function

ELM has $a_i(\phi)$ instead of ϕ_i , later $a_i(\phi) = \phi/w_i$

SM has ϕ_i , later $\phi_i = a_i \phi$



[illegible]

Examples

- Normal: $f(y_i; \mu_i, \sigma^2) = \frac{1}{\sqrt{(2\pi)\sigma}} \exp\left\{-\frac{1}{2\sigma^2}(y_i - \mu_i^2)\right\}$
 $= \exp\left\{\frac{y_i\mu_i - (1/2)\mu_i^2}{\sigma^2} - (1/2)\log \sigma^2 - y_i^2/2\sigma^2 - (1/2)\log \sqrt{(2\pi)}\right\}$

$$\phi_i = \sigma^2, \quad \theta_i = \mu_i, \quad b(\mu_i) = \mu_i^2/2, \quad b'(\mu_i) = \mu_i, \quad b''(\mu_i) = 1$$

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- Binomial: $f(r_i; p_i) = \binom{m_i}{r_i} p_i^{r_i} (1 - p_i)^{m_i - r_i}; \quad y_i = r_i/m_i$
 $= \exp[m_i y_i \log\{p_i/(1 - p_i)\} + m_i \log(1 - p_i) + \log \binom{m_i}{m_i y_i}]$

$$\phi_i = 1/m_i, \quad \theta_i = \log\{p_i/(1 - p_i)\}, \quad b(p_i) = -\log(1 - p_i), \quad p_i = E(y_i)$$

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- ELM (§8.1/6.1) uses $a_i(\phi)$ in place of ϕ_i , later $a_i(\phi) = \phi/w_i$;
SM uses ϕ_i , later (p. 483) $\phi_i = \phi a_i$

Family	Canonical link	Variance function	ϕ_i
Normal	$\eta = \mu$	1	σ^2
Binomial	$\eta = \log\{\mu/(1 - \mu)\}$	$\mu(1 - \mu)$	$1/m_i$
Poisson	$\eta = \log(\mu)$	μ	1
Gamma	$\eta = 1/\mu$	μ^2	$1/\nu$
Inverse Gaussian	$\eta = 1/\mu^2$	μ^3	ξ

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Inverse Gaussian	$\eta = 1/\mu^2$	μ^3	ξ

$$\text{Gamma: } f(y_i; \mu_i, \nu) = \frac{1}{\Gamma(\nu)} \left(\frac{\nu}{\mu_i}\right)^\nu y_i^{\nu-1} \exp\left(-\frac{\nu}{\mu_i}\right) y_i$$

$$= \exp\left[-\frac{\nu}{\mu_i} y_i - \nu \log\left(\frac{1}{\mu_i}\right) + (\nu - 1) \log(y_i) + \nu \log(\nu) - \log\{\Gamma(\nu)\}\right]$$

$$= \exp\left[\nu \left\{ \frac{y_i}{-\mu_i} - \log\left(\frac{1}{\mu_i}\right) + \log(y_i) - \log \Gamma(\nu) + \log(\nu) - \frac{1}{\nu} \log(y_i) \right\}\right]$$

canonical link conventionally ignores the $-$ sign

- $\ell(\beta; \mathbf{y}) = \sum_{i=1}^n \left\{ \frac{y_i \theta_i - b(\theta_i)}{\phi_i} + c(y_i, \phi_i) \right\} \quad b'(\theta_i) = \mu_i; \quad b''(\theta_i) = V(\mu_i)$
- $g(\mu_i) = g\{b'(\theta_i)\} = \eta_i = \mathbf{x}_i^T \beta$
- $\frac{\partial \ell(\beta; \mathbf{y})}{\partial \beta_j} = \sum \frac{\partial \ell_i}{\partial \theta_i} \frac{\partial \theta_i}{\partial \beta_j} = \sum \frac{y_i - b'(\theta_i)}{\phi_i} \frac{\partial \theta_i}{\partial \beta_j}$

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when $\phi_i = a_i \phi$

- $\ell(\beta; \mathbf{y}) = \sum_{i=1}^n \left\{ \frac{y_i \theta_i - b(\theta_i)}{\phi_i} + c(y_i, \phi_i) \right\} \quad b'(\theta_i) = \mu_i; \quad b''(\theta_i) = V(\mu_i)$

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when $\phi_i = a_i \phi$

- matrix notation:

$$\frac{\partial \ell(\beta)}{\partial \beta} = \mathbf{X}^T \mathbf{u}(\beta), \quad \mathbf{X} = \frac{\partial \boldsymbol{\eta}}{\partial \boldsymbol{\beta}^T}, \quad \mathbf{u} = (u_1, \dots, u_n), \quad u_i = \frac{y_i - \mu_i}{\phi_i g'(\mu_i) V(\mu_i)}$$

Scale parameter ϕ_i

- in most cases, either ϕ_i is known, or $\phi_i = \phi a_i$, where a_i is known
- Normal distribution, $\phi = \sigma^2$
- Binomial distribution $\phi_i = m_i^{-1}$
- Gamma distribution, $\phi = 1/\nu$

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$$\bullet \frac{\partial \ell(\beta; \mathbf{y})}{\partial \beta_j} = \sum \frac{y_i - \mu_i}{\phi_i g'(\mu_i) V(\mu_i)} \mathbf{x}_{ij} = \sum \frac{y_i - \mu_i}{a_i \phi g'(\mu_i) V(\mu_i)} \mathbf{x}_{ij}$$

when $\phi_i = a_i \phi$

- if $\theta_i = g(\mu_i)$ **canonical link**, then $g'(\mu_i) = 1/V(\mu_i)$, and

$$\sum \frac{y_i \mathbf{x}_{ij}}{a_i} = \sum \frac{y_i \hat{\mu}_i \mathbf{x}_{ij}}{a_i}$$

Solving maximum likelihood equation

- Newton-Raphson: $\ell'(\hat{\beta}) = 0 \approx \ell'(\beta) + \ell''(\beta)(\hat{\beta} - \beta)$

defines iterative scheme

- $\hat{\beta}^{(t+1)} = \hat{\beta}^{(t)} - \{\ell''(\hat{\beta}^{(t)})\}^{-1} \ell'(\hat{\beta}^{(t)})$

Solving maximum likelihood equation

- Newton-Raphson: $\ell'(\hat{\beta}) = 0 \approx \ell'(\beta) + \ell''(\beta)(\hat{\beta} - \beta)$

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- $\hat{\beta}^{(t+1)} = \hat{\beta}^{(t)} - \{\ell''(\hat{\beta}^{(t)})\}^{-1} \ell'(\hat{\beta}^{(t)})$

- Fisher scoring: $-\ell''(\beta) \leftarrow E\{-\ell''(\beta)\} = i(\beta)$

many books use $I(\beta)$

- $\hat{\beta}^{(t+1)} = \hat{\beta}^{(t)} + \{i(\hat{\beta}^{(t)})\}^{-1} \ell'(\hat{\beta}^{(t)})$

Solving maximum likelihood equation

- Newton-Raphson: $\ell'(\hat{\beta}) = \mathbf{0} \approx \ell'(\beta) + \ell''(\beta)(\hat{\beta} - \beta)$

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many books use $\mathbf{I}(\beta)$

- $\hat{\beta}^{(t+1)} = \hat{\beta}^{(t)} + \{\mathbf{i}(\hat{\beta}^{(t)})\}^{-1} \ell'(\hat{\beta}^{(t)})$

- applied to matrix version: $\mathbf{X}^T \mathbf{u}(\hat{\beta}) = \mathbf{0} \doteq \mathbf{X}^T \mathbf{u}(\beta) + (\hat{\beta} - \beta) \mathbf{X}^T \frac{\partial \mathbf{u}(\beta)}{\partial \beta^T}$ $u_i = \frac{y_i - \mu_i}{\phi_i g'(\mu_i) V(\mu_i)}$

- change to Fisher scoring: $\mathbf{X}^T \mathbf{u}(\hat{\beta}) = \mathbf{0} \doteq \mathbf{X}^T \mathbf{u}(\beta) + (\hat{\beta} - \beta) \mathbf{X}^T \mathbf{E} \left\{ \frac{\partial \mathbf{u}(\beta)}{\partial \beta^T} \right\}$

$$\hat{\beta} = \beta + \mathbf{i}(\beta)^{-1} \mathbf{X}^T \mathbf{u}(\beta)$$

... maximum likelihood equation

$$\hat{\beta} = \beta + i(\beta)^{-1} X^T u(\beta)$$

$$\bullet \frac{\partial^2 \ell(\beta; y)}{\partial \beta_j \partial \beta_k} =$$

... maximum likelihood equation

$$\hat{\beta} = \beta + i(\beta)^{-1} X^T u(\beta)$$

$$\bullet \frac{\partial^2 \ell(\beta; y)}{\partial \beta_j \partial \beta_k} = \sum \frac{-b''(\theta_i)}{\phi_i} \left(\frac{\partial \theta_i}{\partial \beta_j} \right) \left(\frac{\partial \theta_i}{\partial \beta_k} \right) + \sum \frac{y_i - b'(\theta_i)}{\phi_i} \frac{\partial^2 \theta_i}{\partial \beta_j \partial \beta_k}$$

... maximum likelihood equation

$$\hat{\beta} = \beta + i(\beta)^{-1} X^T u(\beta)$$

- $\frac{\partial^2 \ell(\beta; y)}{\partial \beta_j \partial \beta_k} = \sum \frac{-b''(\theta_i)}{\phi_i} \left(\frac{\partial \theta_i}{\partial \beta_j} \right) \left(\frac{\partial \theta_i}{\partial \beta_k} \right) + \sum \frac{y_i - b'(\theta_i)}{\phi_i} \frac{\partial^2 \theta_i}{\partial \beta_j \partial \beta_k}$
- $E \left(-\frac{\partial^2 \ell(\beta; y)}{\partial \beta_j \partial \beta_k} \right) = \sum \frac{V(\mu_i)}{\phi_i} \frac{x_{ij}}{g'(\mu_i)V(\mu_i)} \frac{x_{ik}}{g'(\mu_i)V(\mu_i)} = \sum \frac{x_{ij}x_{ik}}{\phi_i \{g'(\mu_i)\}^2 V(\mu_i)}$

... maximum likelihood equation

$$\hat{\beta} = \beta + i(\beta)^{-1} X^T u(\beta)$$

- $\frac{\partial^2 \ell(\beta; y)}{\partial \beta_j \partial \beta_k} = \sum \frac{-b''(\theta_i)}{\phi_i} \left(\frac{\partial \theta_i}{\partial \beta_j} \right) \left(\frac{\partial \theta_i}{\partial \beta_k} \right) + \sum \frac{y_i - b'(\theta_i)}{\phi_i} \frac{\partial^2 \theta_i}{\partial \beta_j \partial \beta_k}$
- $E \left(-\frac{\partial^2 \ell(\beta; y)}{\partial \beta_j \partial \beta_k} \right) = \sum \frac{V(\mu_i)}{\phi_i} \frac{x_{ij}}{g'(\mu_i)V(\mu_i)} \frac{x_{ik}}{g'(\mu_i)V(\mu_i)} = \sum \frac{x_{ij}x_{ik}}{\phi_i \{g'(\mu_i)\}^2 V(\mu_i)}$
-

$$\begin{aligned}\hat{\beta} &= \beta + (X^T W X)^{-1} X^T u(\beta) = (X^T W X)^{-1} \{X^T W X \beta + X^T u(\beta)\} \\ &= (X^T W X)^{-1} \{X^T W (X \beta + W^{-1} u(\beta))\} \\ &= (X^T W X)^{-1} X^T W z\end{aligned}$$

... maximum likelihood equation

$$\hat{\beta} = \beta + i(\beta)^{-1} X^T u(\beta)$$

- $\frac{\partial^2 \ell(\beta; y)}{\partial \beta_j \partial \beta_k} = \sum \frac{-b''(\theta_i)}{\phi_i} \left(\frac{\partial \theta_i}{\partial \beta_j} \right) \left(\frac{\partial \theta_i}{\partial \beta_k} \right) + \sum \frac{y_i - b'(\theta_i)}{\phi_i} \frac{\partial^2 \theta_i}{\partial \beta_j \partial \beta_k}$
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- does not involve ϕ_i iteratively re-weighted least squares W, z both depend on β
- **derived response** $z = X\beta + W^{-1}u$ linearized version of y

Summary

Model:

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

Estimation:

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

Variance:

$$\text{Var}(\hat{\boldsymbol{\beta}}) \doteq (\mathbf{X}^T \mathbf{W} \mathbf{X})^{-1} \quad \mathbf{W} \text{ is diagonal}$$

On pp. 118-119 of ELM, this iteration is carried out in R on the `bliss` data

Summary 2

$$\begin{aligned}\hat{\beta} &= (X^T W X)^{-1} X^T W z; & z &= X\beta + W^{-1}u; & z(\beta) &= X\beta + W^{-1}(\beta)u(\beta) \\ \text{Var}(\hat{\beta}) &\doteq (X^T W X)^{-1} & & & W &\text{ is diagonal}\end{aligned}$$

$$W_{ii} = \frac{1}{\phi a_i \{g'(\mu_i)\}^2 V(\mu_i)}$$

$$u_i = \frac{y_i - \mu_i}{\phi a_i g'(\mu_i) V(\mu_i)}$$

Note $\hat{\beta}$ is free of ϕ because of W and W^{-1} , but $\text{Var}(\hat{\beta})$ depends on ϕ

Warnings

1. in ELM W is defined slightly differently (no ϕ), so he writes $\widehat{\text{Var}}(\hat{\beta}) = (X^T W X)^{-1} \hat{\phi}$
2. ELM uses w_i where SM uses $1/a_i$

Analysis of data using GLMs: overview

- choose a model, often based on type of response or on mean/variance relationship
- fit a model, using maximum likelihood estimation convergence (almost) guaranteed
- inference for individual coefficients $\hat{\beta}_j$ from summary
- inference for groups of coefficients by analysis of deviance

Analysis of data using GLMs: overview

- choose a model, often based on type of response or on mean/variance relationship
- fit a model, using maximum likelihood estimation convergence (almost) guaranteed
- inference for individual coefficients $\hat{\beta}_j$ from summary
- inference for groups of coefficients by analysis of deviance
- estimation of ϕ based on Pearson's Chi-square

typo in ELM p.121: cross out = $\text{var}(\hat{\mu})$

$$\hat{\phi} = \frac{1}{n-p} \sum_{i=1}^n \frac{(y_i - \hat{\mu}_i)^2}{V(\hat{\mu}_i)}$$

- analysis of deviance: see p. 121 (near bottom) likelihood ratio tests
- diagnostics: same as for `lm` ELM p.124; SM p.477
 - residuals: deviance or Pearson; can be standardized ELM likes 1/2 normal plots
 - influential observations: uses hat matrix `SMPracticals` has very good GLM diagnostics

`glm.diag`, `plot.glm.diag`

- response is **time to event** – failure of a unit, death of a patient, length of unemployment
- response takes values in $[0, \infty)$
- responses may be **censored**

we know $y > c$ only

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- density function $f(y)$; cumulative distribution function $F(y)$
- **survivor function** $S(y) = 1 - F(y) = \text{pr}(Y \geq y)$ assume distribution of y is continuous
- **hazard function** $\lambda(y) = \text{pr}(\text{failure "at" } y \mid \text{survival to } y)$ force of mortality

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$$\lambda(y) = \lim_{h \rightarrow 0} \frac{1}{h} \text{pr}(y \leq Y < y + h \mid Y \geq y) = \frac{f(y)}{S(y)}$$

- cumulative hazard function

$$\Lambda(y) = \int_0^y \lambda(u) du$$

SM uses $h(\cdot)$ for $\lambda(\cdot)$

- exponential: density $f(y) = \lambda e^{-\lambda y}$
- Weibull: survivor fn $S(y) = \exp\{-(y/\theta)^\alpha\}$; hazard function $\lambda(y) = \alpha\theta^{-\alpha}y^{\alpha-1}$

- Gamma: hazard function

$$\lambda(y) = \frac{\lambda^\alpha y^{\alpha-1} e^{-\lambda y}}{\int_y^\infty \lambda^\alpha u^{\alpha-1} e^{-\lambda u} du}$$

- log-logistic: survivor function

$$S(y) = \frac{1}{1 + (\lambda y)^\alpha}$$

- sometimes it's more convenient to specify the density, sometimes the hazard, sometimes the survivor function

- sample of size n , we observe $(y_1, d_1), \dots, (y_n, d_n)$
- $y_i = \min(y_i^o, c_i)$ where $Y_i^o \sim f(\cdot)$ and $C_i \sim g(\cdot)$
- C_i is an associated censoring time for unit i
- $d_i = 1$ if y_i is uncensored; $d_i = 0$ if y_i is censored

independent

192

5 · Models

0+	1+	1+	3+	3+	7	10+	11+	12+	12+	15+	18+
20+	22+	22+	24+	25+	26+	31+	36+	36+	36	38	40
47+	47+	49+	53+	53+	55+	56+	57+	61+	67+	67+	70
73	75+	77+	83+	84+	88+	89+	99	121+	122+	123+	141+
0+	0+	2+	2+	2+	2+	3	3+	4+	5+	9+	10+
11	12+	13	13+	18+	22+	22+	24+	24+	24+	25+	26+
27	28	32+	35+	36	40+	43+	50+	54			

Table 5.3

Blalock–Taussig shunt data (Oakes, 1991). The table gives survival time of shunt (months after operation) for 48 infants aged over one month at time of operation, followed by times for 33 infants aged 30 or fewer days at operation. Infants whose shunt has not yet failed are marked +.

- data $(y_1, d_1), \dots, (y_n, d_n)$
- likelihood function

$$\begin{aligned} L(\theta; y, d) &= \prod_{i=1}^n f(y_i; \theta)^{d_i} \{1 - F(y_i; \theta)\}^{1-d_i} \\ &= \prod_{i=1}^n \left[\frac{f(y_i; \theta)}{1 - F(y_i; \theta)} \right]^{d_i} \{1 - F(y_i; \theta)\} \\ &= \prod_{i=1}^n \lambda(y_i; \theta)^{d_i} \{1 - F(y_i; \theta)\} \end{aligned}$$

- log-likelihood function

cumulative hazard function

$$\ell(\theta; y, d) = \sum_{i=1}^n d_i \log\{\lambda(y_i; \theta)\} - \Lambda(y_i; \theta)$$

- any of the models above could be used for inference

- order the observed times $y_1 < y_2 < \dots < y_n$
- Kaplan-Meier estimator of survivor function:

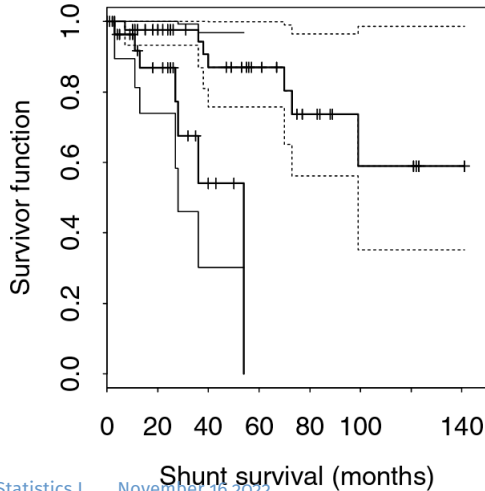
life-table

$$\hat{S}(y) = \prod_{y_i \leq y} \left(1 - \frac{1}{r_i}\right)^{d_i}$$

•

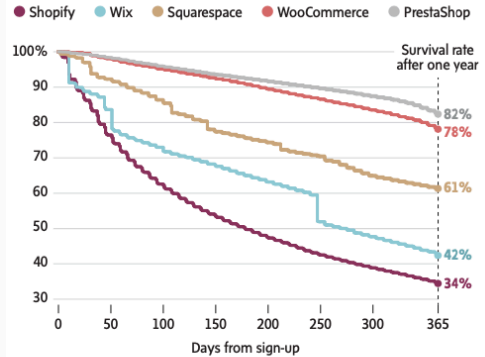
$$\text{var}\{\hat{S}(y)\} \doteq \{\hat{S}(y)\}^2 \sum_{y_i \leq y} \frac{d_i}{r_i(r_i - d_i)}$$

- censoring indicator $d_i = 1$ if y_i is a failure time
- r_i = number of items available to fail at time y_i
= $\#\{j; y_j \geq y_i\}$
= $n - i + 1$ if there are no ties in the data



E-commerce domain survival rates, by platform, 2019–2021

Percentage of domains that survive by number of days after sign-up



MURAT YÜKSELİR AND MAHIMA SINGH / THE GLOBE AND MAIL,
SOURCE: GLOBE AND MAIL ANALYSIS

- data $(y_i, d_i, x_i), i = 1, \dots, n$ independent still ordered
- parametric inference: model hazard function or density function in terms of $y_i \mid x_i$
- log-likelihood function $\ell(\beta; y, d, x) = \sum_{i=1}^n d_i \log\{\lambda(y_i; x_i, \beta)\} - \Lambda(y_i; x_i, \beta)$

SM Example 10.36

- **proportional hazards model**

$$\lambda(y_i; x_i, \beta) = \lambda_o(y_i) \exp\{x_i^T \beta\}$$

- **partial likelihood**

$$L_{\text{part}}(\beta; t, x) = \prod_{\text{failures}} \frac{\exp(x_i^T \beta)}{\sum_{j \in \mathcal{R}_i} \exp(x_j^T \beta)}$$

- **risk set** \mathcal{R}_i set of individuals still alive at the time the i th item fails

- partial log likelihood

$$\ell_{\text{part}}(\beta; \mathbf{t}, \mathbf{x}) = \sum_{\text{failures}} [\mathbf{x}_i^T \beta - \log\{\sum_{j \in \mathcal{R}_i} \exp(\mathbf{x}_j^T \beta)\}]$$

- inference

$$\ell'_{\text{part}}(\hat{\beta}) = \mathbf{0}; \quad -\ell''_{\text{part}}(\hat{\beta}) \doteq \{\widehat{\text{var}}(\hat{\beta})\}^{-1}$$

- partial log likelihood

$$\ell_{\text{part}}(\beta; \mathbf{t}, \mathbf{x}) = \sum_{\text{failures}} [\mathbf{x}_i^T \beta - \log \{ \sum_{j \in \mathcal{R}_i} \exp(\mathbf{x}_j^T \beta) \}]$$

- inference

$$\ell'_{\text{part}}(\hat{\beta}) = \mathbf{0}; \quad -\ell''_{\text{part}}(\hat{\beta}) \doteq \{\widehat{\text{var}}(\hat{\beta})\}^{-1}$$

$$\hat{\beta} - \beta \sim \mathbf{N}(\mathbf{0}, \widehat{\text{var}}(\hat{\beta}))$$

$$2\{\ell_{\text{part}}(\hat{\beta}) - \ell_{\text{part}}(\beta_0)\} \sim \chi_p^2$$

- can be proved (but it's hard) that the usual likelihood theory applies to ℓ_{part}

- estimation of survivor function $S(y; x) = \text{pr}(Y \geq y \mid x)$
- under PH model $S(y; x) = \{S_0(y)\}^{\exp(x^T \beta)}$
- estimate baseline survivor function

$$\hat{S}_0(y) = \prod_{i: y_i \leq y} \left(1 - \frac{d_i}{\sum_{j \in \mathcal{R}_y} \exp(x_j^T \hat{\beta})} \right)$$

- estimate survivor function for individual with covariates x_+ :

$$\hat{S}(y; x_+) = \{\hat{S}_0(y)\}^{\exp(x_+^T \hat{\beta})}$$

Research

JAMA | Original Investigation | CARING FOR THE CRITICALLY ILL PATIENT

Effect of a Resuscitation Strategy Targeting Peripheral Perfusion Status vs Serum Lactate Levels on 28-Day Mortality Among Patients With Septic Shock

The ANDROMEDA-SHOCK Randomized Clinical Trial

Glenn Hernández, MD, PhD; Gustavo A. Ospina-Tascón, MD, PhD; Lucas Petri Damiani, MSc; Elisa Estenssoro, MD; Arnaldo Dubin, MD, PhD; Javier Hurtado, MD; Gilberto Friedman, MD, PhD; Ricardo Castro, MD, MPH; Leyla Alegría, RN, MSc; Jean-Louis Teboul, MD, PhD; Maurizio Cecconi, MD, FFICM; Giorgio Ferri, MD; Manuel Jibaja, MD; Ronald Pairumani, MD; Paula Fernández, MD; Diego Barahona, MD; Vladimir Granda-Luna, MD, PhD; Alexandre Biasi Cavalcanti, MD, PhD; Jan Bakker, MD, PhD; for the ANDROMEDA-SHOCK Investigators and the Latin America Intensive Care Network (LIVEN)

[link](#)

“the treatment effect on the primary outcome was calculated with Cox proportional hazards, with adjustment for 5 pre-specified baseline covariates”

“results are reported as hazard ratio with 95% confidence interval
and Kaplan-Meier curves”

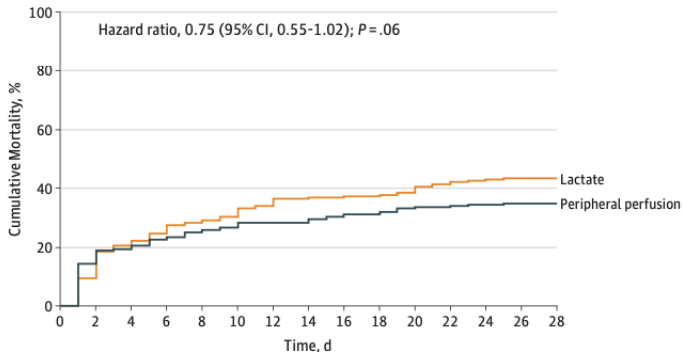
Research Original Investigation

Effect on Septic Shock Mortality of Resuscitation Targeting Peripheral Perfusion vs Serum Lactate Levels

Table 2. Main Outcomes of the Study of Resuscitation Strategies in Septic Shock

Outcome	Peripheral Perfusion-Targeted Resuscitation (n = 212)	Lactate Level-Targeted Resuscitation (n = 212)	Unadjusted Absolute Difference (95% CI)	Adjusted Relative Measure (95% CI)	P Value
Primary Outcome					
Death within 28 d, No. (%)	74 (34.9)	92 (43.4)	-8.5 (-18.2 to 1.2) ^b	HR, 0.75 (0.55 to 1.02) ^a	.06 ^a
Secondary Outcomes					
Death within 90 d, No. (%)	87 (41.0)	99 (46.7)	-5.7 (-15.6 to 4.2) ^b	HR, 0.82 (0.61 to 1.09) ^a	.17 ^a

Figure 2. Kaplan-Meier Estimates of Cumulative Mortality Within 28 Days Among Patients Treated With Peripheral Perfusion-Targeted Resuscitation vs Lactate Level-Targeted Resuscitation



Hazard ratio, 95% confidence interval, and *P* value were calculated with a Cox proportional hazards model that included as covariates baseline Acute Physiology and Chronic Health Evaluation (APACHE) II score,²³ Sequential Organ Failure Assessment (SOFA) score,²⁴ lactate level, capillary refill time, and source of infection. Median follow-up for peripheral perfusion-targeted resuscitation was 28 days (interquartile range, 8-28 days) and for lactate level-targeted resuscitation was 28 days (interquartile range, 6-28 days).

No. at risk																	
Lactate	212	192	168	160	152	148	140	135	134	133	130	124	122	120	120		
Peripheral perfusion	212	182	171	164	159	155	152	152	148	146	142	141	139	138	138		

See *Appendix to An R Companion to Applied Regression*

```
> library(car)
```

```
> data(Rossi)
```

```
> Rossi[1:5, 1:10]
```

	week	arrest	fin	age	race	wexp	mar	paro	prio	educ
1	20	1	no	27	black	no	not married	yes	3	3
2	17	1	no	18	black	no	not married	yes	8	4
3	25	1	no	19	other	yes	not married	yes	13	3
4	52	0	yes	23	black	yes	married	yes	1	5
5	52	0	no	19	other	yes	not married	yes	3	3

```
> mod.allison <- coxph(Surv(week,arrest) ~ fin + age + race + wexp + mar + paro  
+ prio, data = Rossi)
```



```
> summary(mod.allison)
```

```
Call:
```

```
coxph(formula = Surv(week, arrest) ~ fin + age + race + wexp +  
      mar + paro + prio, data = Rossi)
```

```
n= 432, number of events= 114
```

	coef	exp(coef)	se(coef)	z	Pr(> z)	
finyes	-0.37942	0.68426	0.19138	-1.983	0.04742	*
age	-0.05744	0.94418	0.02200	-2.611	0.00903	**
raceother	-0.31390	0.73059	0.30799	-1.019	0.30812	
wexpyes	-0.14980	0.86088	0.21222	-0.706	0.48029	
marnot married	0.43370	1.54296	0.38187	1.136	0.25606	
paroyes	-0.08487	0.91863	0.19576	-0.434	0.66461	
prio	0.09150	1.09581	0.02865	3.194	0.00140	**

```
> plot(survfit(mod.allison), ylim=c(.7,1), xlab="weeks",  
      ylab = "proportion not re-arrested")
```

