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Introduction

- Ensuring drug safety begins with extensive pre-approval clinical trials
- This process continues *after* approval when drugs are in widespread use: **post-marketing surveillance**
- Drugs taken by more people, for longer periods of time, and in different ways than in pre-approval trials
- May identify adverse health outcomes associated with drug exposure that were not previously detected



1997



2004

Statistical Objectives

- Identify drug-condition pairs that may be linked
- Find drug interactions linked with conditions
- Estimate the strength of these associations
- Fundamental Difficulties**
 - Large size: Millions of people, 10000's of conditions
 - High dimension: 10000's of drugs, millions of interactions

Current System: AERS

- Current approach to surveillance is based on the FDA's **Adverse Event Reporting System (AERS)**
- Anyone can voluntarily submit a report on adverse events (AEs) that may be related to drug exposures
- Difficulties with AERS**
 - Spelling – spelling errors, etc.
 - Bias – underreporting, duplicate reports, media
 - Unreliable temporal information
- Multiple drugs and AEs may be listed on one report

		AE j	
		Yes	No
Drug	Yes	a	b
	No	c	d
		Total: N	

- 15000 drugs x 16000 AEs = 240 million tables
- Most AEs do not occur with most drugs; small counts in a

- FDA uses 2 x 2 summaries, applies Bayesian shrinkage methods to deal with variability due to small counts
- Limitations**
 - No adjustment for confounding drugs
 - Ignores interactions
 - May not utilize temporal information

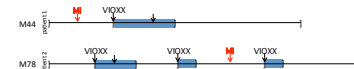
Longitudinal Health Databases

- Sentinel Initiative** - FDA plans to establish an active surveillance system using data from healthcare information holders
- Observational Medical Outcomes Partnership (OMOP)** - public/private partnership to research methods for statistical analysis of health databases
- Medical claims databases** - Time-stamped records of actions that generate insurance claims - filling a prescription, visiting a physician, etc.

- Advantages**
 - Automated
 - Better temporal data
- Disadvantages**
 - Little baseline data
 - No OTC information
- Many potential analysis techniques: maxSPRT, cohort methods, case control, case-crossover ...

Self Controlled Case Series

- Method developed to estimate relative incidence of AEs to assess vaccine safety [Farrington, 1995]
- One drug, one adverse event (AE)**



- Person i observed for τ_i days; (i,d) is their dth day
- y_{id} = # of events observed on (i,d)
- $x_{id} = 1$ if exposed to drug on (i,d), 0 otherwise

- Events arise according to a non-homogeneous Poisson process, exposure modulates the event rate

- Intensity on (i,d) = $e^{\phi_i + \beta x_{id}}$

$$y_{id} | x_{id} \sim \text{Poisson}(e^{\phi_i + \beta x_{id}})$$

$$L_i = P(y_{i1}, \dots, y_{i\tau_i} | x_{i1}, \dots, x_{i\tau_i}) = P(y_i | x_i) = \prod_{d=1}^{\tau_i} P(y_{id} | x_{id})$$

Condition to remove ϕ_i

- Could use ML to get estimates, but drug effect β is of interest and the ϕ_i 's are *nuisance* parameters

- Condition on sufficient statistic $n_i = \sum y_{id}$

$$n_i | x_i \sim \text{Poisson}\left(\sum_d e^{\phi_i + \beta x_{id}}\right)$$

- Conditional likelihood for i

$$L_i^c = P(y_i | x_i, n_i) = \frac{P(y_i | x_i)}{P(n_i | x_i)} \propto \prod_{d=1}^{\tau_i} \left(\frac{e^{\beta x_{id}}}{\sum_{d'} e^{\beta x_{id'}}} \right)^{y_{id}}$$

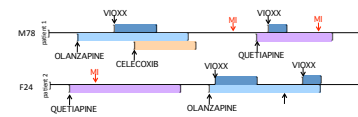
- Maximize $l^c = \sum \log L_i^c$ to get $\hat{\beta}_{EMLE}$ → consistent, asymptotically Normal [Cameron and Trivedi, 1998]

Data Reduction to Cases Only

- If i has no events ($y_i = 0$) then $L_i^c = 1$, so we **only need cases** (i.e. $n_i \geq 1$) in the analysis
- Computational advantage – incidence rate of most AEs is low, so may only have ~100,000 cases involved rather than many millions
- SCCS does within-person comparison of event rate during exposure to event rate while unexposed ('self-controlled')

Multiple Drugs and Interactions

- We extend the model to one AE and **multiple** drugs



- Intensity on (i,d) = $e^{\phi_i + \beta' x_{id}} = e^{\phi_i + \beta_1 x_{id1} + \dots + \beta_p x_{idp}}$

$$x_{id} = (x_{id1}, \dots, x_{idp})^T \quad \beta = (\beta_1, \dots, \beta_p)^T$$

- $x_{idj} = 1$ if exposed to drug j; 0 otherwise
- Intensity with drug interactions, time-varying covariates:

$$e^{\{\phi_i + \beta' x_{id} + \sum_{r \neq s} \gamma_{rs} x_{idr} x_{ids} + \alpha' z_{id}\}}$$

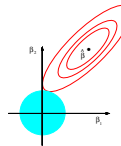
Bayesian Extension of SCCS

- Longitudinal databases have 10000's of potential drugs
- Intensity model: $e^{(\text{main effects}) + (2\text{-way interactions})}$ → high dimensionality with *millions of predictors*
- Standard ML leads to overfitting; need to regularize
- Our approach** – put a prior on β parameters to shrink the estimates toward zero, smooth out estimation, and reduce overfitting

1. Normal prior (ridge regression)

$$\beta_j \sim N(0, \sigma_j^2)$$

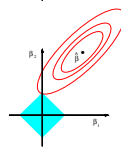
$$\text{max lik subject to } \sum_{j=1}^p \beta_j^2 \leq s$$



2. Laplacian prior (lasso)

$$\beta_j \sim \text{Laplace}(0, 1/\lambda)$$

$$\text{max lik subject to } \sum_{j=1}^p |\beta_j| \leq s$$



- Convex optimization: Posterior modes via cyclic coordinate descent [Genkin et al, 2007]
- Handles millions of predictors in logistic case (BBR)

Results: OMOP Evaluation

- Methods evaluation:
 - Chose 10 drugs, 10 conditions of interest
 - 9 drug-condition pairs with a *true association*
 - Pairs determined to have no link serve as *negative controls*
- Evaluation is based on **mean average precision (mAP) score**: measures how much a method maximizes 'true positives' while minimizing 'false positives'

MSLR database (1.5M people)

Method	mAP score
22 PRR	0.2251486
22 OR	0.228857
23 BCPMA	0.209257
22 EBOP	0.2173618
23 CHI-SQ	0.2144375
22 PRRS	0.204662
23 ROBES	0.2044231
12 BCPMA05	0.1823317
12 EBOS	0.180992
SCCS (1 AE, 1 drug)	0.2126972
Bayesian SCCS, Normal prior, precision 1 (1 AE, 1 drug)	0.2686558
Bayesian Logistic Regression, Normal prior, precision 1 (1 AE, multiple drugs)	0.2655139
Case-Control	0.186743

Allowing Event Dependence

- SCCS assumes conditional independence of events

$$y_{id} \perp\!\!\!\perp y_{id'} | x_i \text{ for } d \neq d'$$
- In practice, occurrence of an event may increase future risk of that event (e.g. MI)
- We generalize the model by allowing the occurrence of events to additively increase the baseline event rate

$$\lambda_i(t | H_i(t)) = (e^{\phi_i} + \delta N_i(t-)) e^{x_i(t)^T \beta}$$

- If person i has n_i events at times t_{i1}, \dots, t_{in_i} , their likelihood contribution is (Cook and Lawless, 2007):

$$L_i = \prod_{j=1}^{n_i} (e^{\phi_i} + \delta N_i(t_{ij}-)) e^{x_i(t_{ij})^T \beta} \times \exp\left\{-\int_0^{t_{i1}} (e^{\phi_i} + \delta N_i(u-)) e^{x_i(u)^T \beta} du\right\}$$

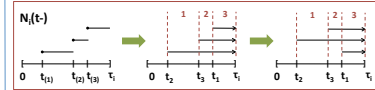
$$= \prod_{j=1}^{n_i} e^{x_i(t_{ij})^T \beta} \times \frac{(e^{\phi_i} + \delta(n_i - 1))}{(e^{\phi_i} - 1)^{n_i}} \exp\left\{-e^{\phi_i} \int_0^{t_{i1}} e^{x_i(u)^T \beta} du\right\} \exp\left\{-\delta \int_0^{t_{i1}} (n_i - u) e^{x_i(u)^T \beta} du\right\}$$

- It is clear from the expression that n_i and τ_i are sufficient for Φ_i , so we will condition on $\{N_i(\tau_i) = n_i\}$

$$P(N_i(\tau_i) = n_i) = \int \dots \int P(n_i \text{ events at } t_{i1}, \dots, t_{in_i} \text{ in } [0, \tau_i]) dt_{i1} \dots dt_{in_i}$$

$$\propto \int \dots \int \prod_{j=1}^{n_i} e^{x_i(t_{ij})^T \beta} \times \exp\left\{-\delta \int_0^{t_{i1}} \sum_{j=1}^{n_i} e^{x_i(t_{ij})^T \beta} du\right\} dt_{i1} \dots dt_{in_i}$$

$$= \int \dots \int \prod_{j=1}^{n_i} e^{x_i(t_{ij})^T \beta} \times \exp\left\{-\delta \int_0^{t_{in_i}} e^{x_i(u)^T \beta} du\right\} dt_{i1} \dots dt_{in_i}$$



$$P(N_i(\tau_i) = n_i) \propto \frac{1}{n_i!} \prod_{j=1}^{n_i} \int_0^{\tau_i} e^{x_i(t_{ij})^T \beta} \exp\left\{-\delta \int_{t_{ij}}^{\tau_i} e^{x_i(u)^T \beta} du\right\} dt_{ij}$$

$$= \frac{1}{n_i!} \left(\frac{1}{\delta}\right) \left[\int_0^{\tau_i} \frac{d}{du} \exp\left\{-\delta \int_0^u e^{x_i(t)^T \beta} dt\right\} du\right]^{n_i}$$

$$= \frac{1}{n_i!} \left(\frac{1}{\delta}\right) \left[1 - \exp\left\{-\delta \int_0^{\tau_i} e^{x_i(t)^T \beta} dt\right\}\right]^{n_i}$$

- The conditional likelihood no longer depends on ϕ_i

$$L_i^c = \frac{P(n_i \text{ events at } t_{i1}, \dots, t_{in_i} \text{ in } [0, \tau_i] | n_i)}{P(N_i(\tau_i) = n_i)}$$

$$= n_i! \exp\left\{-\delta \int_0^{\tau_i} N_i(u-) e^{x_i(u)^T \beta} du\right\} \prod_{j=1}^{n_i} \left(\frac{\delta e^{x_i(t_{ij})^T \beta}}{1 - \exp\left\{-\delta \int_0^{t_{ij}} e^{x_i(t)^T \beta} dt\right\}}\right)$$

Further Work

- Hierarchical modeling of drugs into drug classes

$$e^{\beta_1 x_{i1} + \dots + \beta_j x_{ij} + \dots + \beta_k x_{ik} + \dots + \beta_p x_{ip}}$$

$$N(\mu_{(1)}, \sigma_{(1)}^2) \dots N(\mu_{(p)}, \sigma_{(p)}^2)$$

- Hierarchical modeling of conditions into classes

$$\begin{matrix} \text{condition class } C \\ \vdots \\ y_m \end{matrix} \sim \begin{matrix} \text{drug class } [1] & \dots & \text{drug class } [D] \\ \beta_1 x_{i1} + \dots + \beta_j x_{ij} + \dots + \beta_k x_{ik} + \dots + \beta_p x_{ip} \\ \vdots \\ \beta_1 x_{i1} + \dots + \beta_j x_{ij} + \dots + \beta_k x_{ik} + \dots + \beta_p x_{ip} \end{matrix}$$

$$N(\mu_{(1)}, \sigma_{(1)}^2) \dots N(\mu_{(p)}, \sigma_{(p)}^2)$$

References

- Cameron and Trivedi (1998) Regression Analysis of Count Data. Cambridge University Press.
- Farrington (1995) "Relative incidence estimation from case series for vaccine safety evaluation," *Biometrics*, Vol. 51, No. 1, pg. 228-235.
- Genkin et al. (2007) "Large-scale Bayesian logistic regression for text categorization," *Technometrics*, Vol. 49, No. 3, pg. 291-304.
- Cook and Lawless (2007) The Statistical Analysis of Recurrent Events.