



# Solutions to the problem of monotone likelihood in Cox and logistic regression

## Theory and applications

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# Outline

- Monotone likelihood: examples
- Options of analysis
- Penalized likelihood (PL) estimates
- Profile PL confidence intervals
- Application to Cox, unconditional and conditional logistic regression
- General comments about bias reduction and PL estimates

# Example 1: Preterm infants

From: Berger et al, J Perinat Med, 2003

<b>Group</b>	<b>No CLD</b>	<b>CLD</b>
No amniotic cavity culture found	40	0
Ureaplasma urealyticum found	17	4

OR estimate:  $4/0 / 40/17 = \text{infinite}$

# Example 2: Urinary tract infection

From: Foxman et al, Epidemiology, 1997

<b>Group</b>	<b>No infection</b>	<b>Infection</b>
No diaphragm use	109	123
Diaphragm use	0	7

Other variables in the model:

Age, use of condoms, use of lubricated condoms, use of spermicides, oral contraceptives

# Example 2: Urinary tract infection

- OR estimates of diaphragm use obtained by glm of SPLUS:
- Convergence criterion is change in deviance

<b>Criterion</b>	<b>Estimate</b>	<b>Lower</b>	<b>Upper</b>	<b>P-value</b>
0.001	220	0.8	6.5e5	0.06
0.0001	1726	4e-4	6.8e9	0.34
0.00001	34774	<1e-4	1.1e34	0.76

# Example 3: Breast cancer

- From Lösch et al, Brit J Cancer, 1998
- Survival of 100 patients, 74 censored
- 4 risk factors (pT, N, G, CD)
- Analysis via Cox regression:

<b>Faktor</b>	<b>RR (95% c.i.)</b>	<b>P-value</b>
pT	3.6 (1.3 – 9.6)	0.01
N	2.6 (1.1 – 5.9)	0.03
G	248054 (0 – 2 x 10 <sup>188</sup> )	0.95
CD	1.5 (0.6 – 3.6)	0.37

# Common to examples 1-3:

- Degenerate variation of outcome in one subgroup
  - Ex 1: no CLD+ for no ureaplasms found
  - Ex 2: only „infections“ for diaphragm users
  - Ex 3: no deaths for  $G=0$
- Parameter estimates  $\hat{\beta}$  are infinite
- Standard errors infinite
- Confidence interval  $[-\infty, +\infty]$  uninformative
- $\hat{\beta}/se \rightarrow 0$



# First occurrence in literature

- In Cox regression:

*„Monotone likelihood“*

(Bryson and Johnson, Technometrics, 1981)

- In logistic regression:

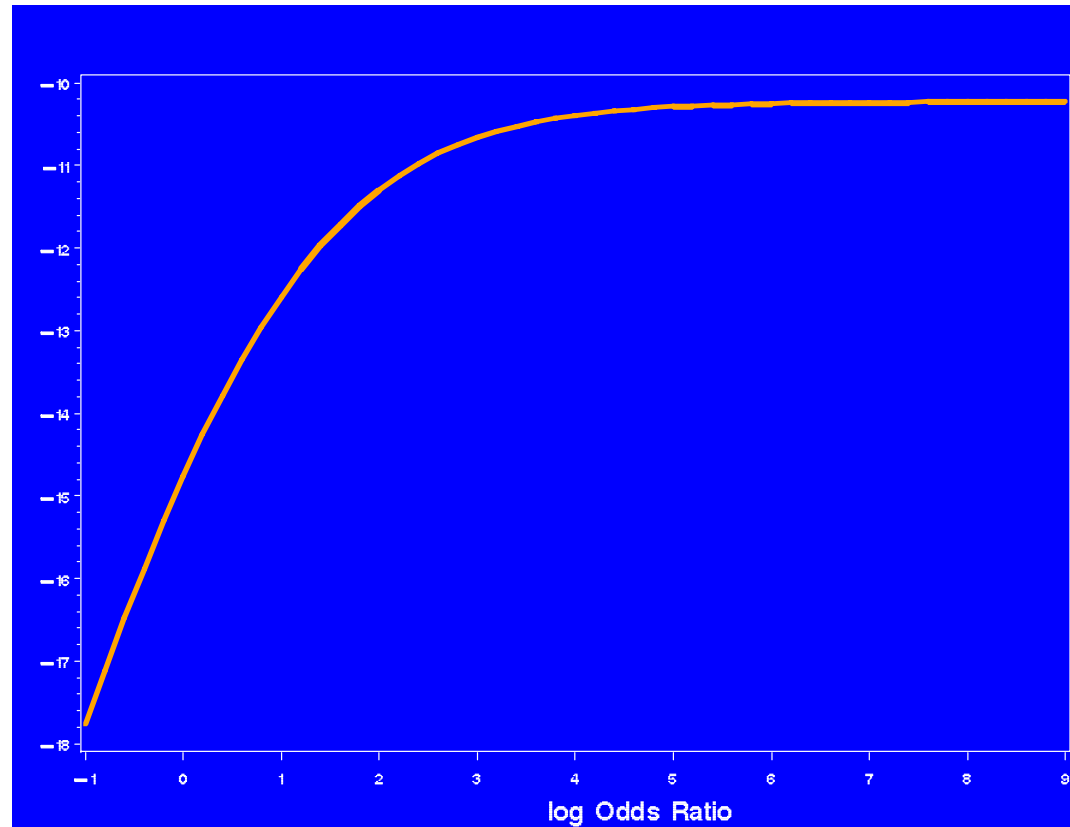
*„(Quasi-)complete Separation“*

(Day and Kerridge, Biometrics 1967)



# Monotone likelihood

- Likelihood is monotone
- no finite maximizer
  
- Likelihood is flat
- second derivative is 0
- variance is infinite





# Incidence of monotone likelihood

- High incidence if:
- Small N / heavy censoring
- Unbalanced covariates
- Large underlying effects
- Strong correlation among covariates

# Options of analysis



- Omit covariate X that is causing monotone likelihood
- Stratify analysis by covariate X
- Choose different type of model
  - Transform X (e.g. use quasi-metric scaling instead of dummies)
  - Use additive risk model instead of multiplicative risk model
- Ad hoc data adjustment
  - Haldane: add  $\frac{1}{2}$  to each cell (for rx2 tables)
  - Laplace: add 1 to each cell (for rx2 tables)
  - Clogg et al, JASA 1991: generalized data adjustment
- Exact logistic regression (LogXact, Cytel Software Corp.)

# Why exact logistic regression?

- Should read  
„*Exact conditional logistic regression*“
- Implementations:
  - LogXact
  - SAS/PROC LOGISTIC (from V8.1 on)
- *Exact*: inference based on exact distribution of sufficient statistic under null hypothesis
- *Conditional*: eliminate nuisance parameters by conditioning on sufficient statistics
- Point estimate is not „exact“, rather conditional

# Exact conditional log regression

- Maximum conditional likelihood estimate (MCLE):
  - $\Pr(T=T_{\text{obs}}|\beta)=\max!$
  - Infinite if  $T_{\text{obs}}$  is largest (smallest) possible value of  $T$
- Median unbiased estimate (MUE):
  - $\Pr(T \geq T_{\text{obs}} | \beta) \geq 0.5, \Pr(T \leq T_{\text{obs}} | \beta) \geq 0.5$
  - MUE is defined even if MCLE is infinite
- LogXact: MUE replaces MCLE if MCLE is infinite

# Exact conditional log regression

- Can even be (ab)used for estimating a Cox model:
  - Each risk set contributes a nuisance parameter that is eliminated by conditioning
  - Conditioning on risk sets improves on asymptotic Cox model, but still violates nominal significance level because of interdependence of risk sets
  - shown for exact logrank test in Heinze et al (2003)
- Problems if exact null distribution is (nearly) degenerate:
  - Conditioning on continuous covariates
  - Too many covariates, too many different levels of covariates
- If exact null distribution is degenerate:  
no estimation/inference possible



# Example 4: Lung cancer

- Case-control study, 18 matched sets, 1:m matching
- Factors smoking (S), radiation (R), RxS
- Options of analysis:
  - CML (conditional maximum likelihood)
    - conditions on matched sets
    - but estimates effects S, R, RxS simultaneously
  - CXL (conditional exact maximum likelihood),
    - conditions on matched sets
    - and eliminates other effects by conditioning

## Example 4: Lung cancer, OR (95% ci)

Method	Radiation	Smoking	RxS
CML	1.2 (0.17, 8.5)	21 (2.6, 167)	$\infty$ (0.0, $\infty$ )
CXL	1.2 (0.14, 20)	20 (2.6, 859)	2.5 (0.06, $\infty$ )

- CXL estimate for RxS is a MUE,  
based on a distribution consisting of 2 possible values only  
(overconditioning)
- Had the other of these two values been observed:  
CML estimate = -  $\infty$ ,  
but CXL estimate = still 2.5
- Therefore, CXL estimate and c.i. are very conservative!



# A solution through bias reduction

- Firth, Biometrika 1993:
- Eliminate  $O(1/n)$ -bias from maximum likelihood parameter estimates
- Bias reduction is applied while estimating the parameter estimates:

*bias preventing, not bias correcting*

- Maximize a penalized likelihood:

$$\log L^* = \log L + \frac{1}{2} \log \det (I)$$

with  $I$  denoting Fisher information matrix

# A solution through bias reduction (2)

- Firth's paper remained undiscovered for 8 years
- Application to log reg and Cox reg possible (Heinze and Schemper, 2001, 2002)
- We showed that parameter estimates are always finite
- Small-sample bias is greatly reduced

# Penalized maximum likelihood estimation

- Penalisation by  $\frac{1}{2} \log \det(I)$  is like adding pseudo-observations with total weight  $k$  to the data
- Log Reg: each observation  $(x_i, y_i)$  is splitted into two new observations:
  - A: Outcome  $y_i$ , weight  $1+h_i/2$
  - B: Outcome  $1-y_i$ , weight  $h_i/2$
  - $h_i$  are diagonal elements of hat matrix  $H$  (leverages)
  - $\sum h_i = k$ ;  $0 \leq h_i \leq 1$
  - Balance of pseudo-observations guarantees finite estimates
- Weighting of pseudo-observations is done iteratively during estimation

# Inference

A decorative graphic consisting of six circles arranged in two rows. The top row has two circles: a solid light purple one on the left and an outlined light purple one on the right. The bottom row has three circles: a solid light purple one on the left, an outlined light purple one in the middle, and a solid light purple one on the right.

- Standard errors deduced from second derivative of  $\log L$  (rather than  $\log L^*$ )
- Although penalized likelihood has maximum, its shape is very asymmetric in case of monotone likelihood
- Normal approximation unsuitable
- Better: Profile penalized likelihood confidence intervals, penalized likelihood ratio tests

# Profile penalized likelihood

- Likelihood ratio statistic

$$LR(\hat{\gamma}, \gamma_0) = 2 \left\{ \log L(\hat{\gamma}, \hat{\delta})^* - \log L(\gamma_0, \hat{\delta}_{\gamma_0})^* \right\}$$

- Under  $H_0: \gamma = \gamma_0$ ,  $LR \sim \chi^2$

- 95% confidence interval:

set of values  $\gamma_0$  for which  $LR < \chi^2_{1;0.95}$

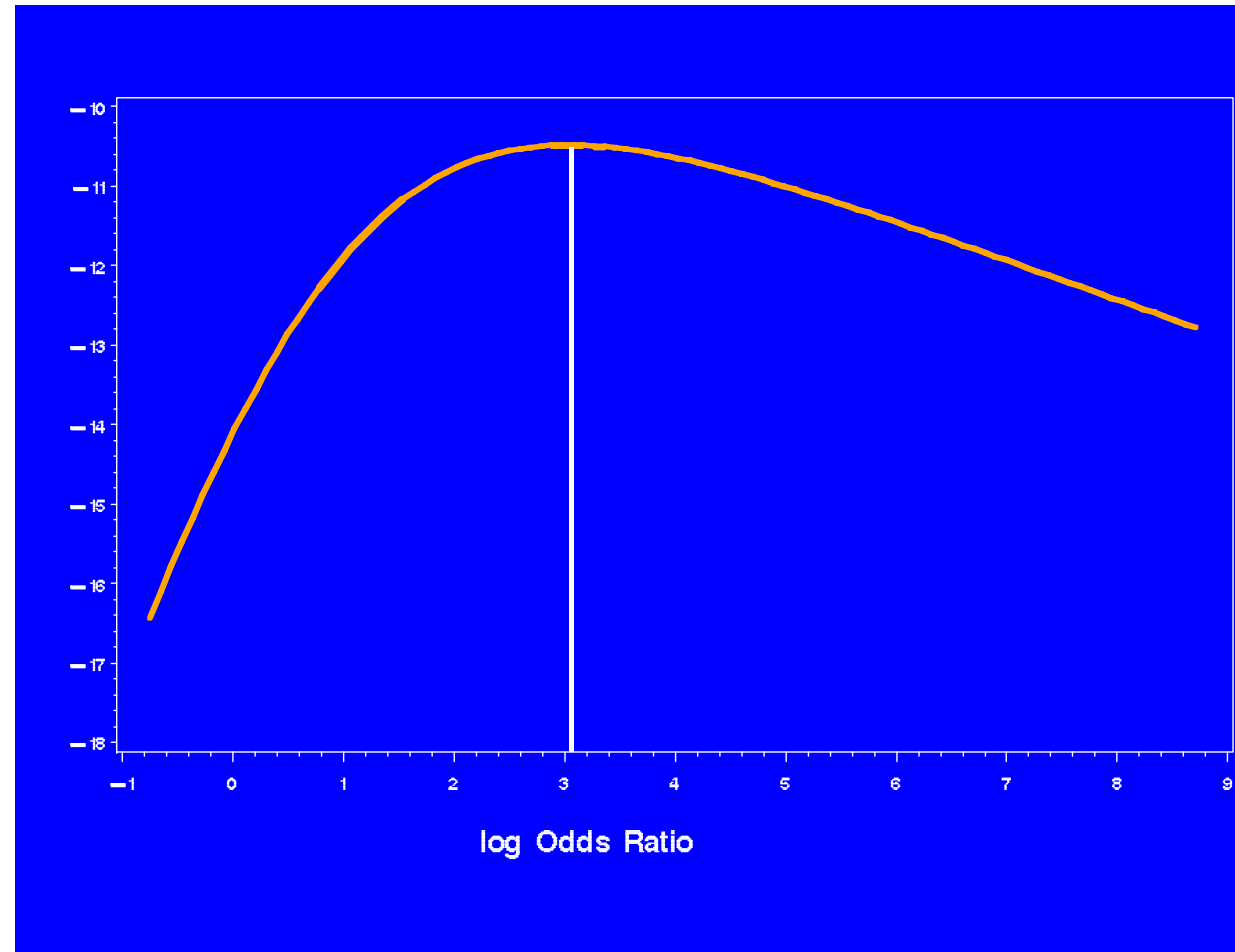
# Example 1: Preterm infants

OR (95% CI):

**20.8**

(2.1 – 2017)

$p=0.007$



## Example 2: Urinary tract infection

Factor	OR estimate	
	ML	PML
Age	3.2	3.0
Oral contraceptives	0.9	0.9
Condom use	11.1	9.7
Lubricated condom	0.1	0.1
Spermicide use	0.4	0.5
Diaphragm use	INF	22.1

# Example 3: Breast cancer

	<b>ML RR (95% CI)</b>	<b>P</b>	<b>PML RR (95% CI)</b>	<b>P</b>
pT	3.6 (1.3 – 9.6)	0.01	3.4 (1.4 – 9.5)	0.01
N	2.6 (1.1 – 5.9)	0.03	2.5 (1.1 – 5.8)	0.03
G	248054 (0 – 2 x 10 <sup>188</sup> )	0.95	<i>11.3 (1.5 – 1452)</i>	<i>0.01</i>
CD	1.5 (0.6 – 3.6)	0.37	1.5 (0.6 – 3.5)	0.36



## Example 4: Lung cancer, OR (95% ci)

Method	Radiation (main effect)	Smoking (main effect)	RxS
CML	1.2 (0.17, 8.5)	21 (2.6, 167)	$\infty$ (0.0, $\infty$ )
CXL	1.2 (0.14, 20)	20 (2.6, 859)	2.5 (0.06, $\infty$ )
<i>CPML</i>	<i>0.99 (0.2, 3.1)</i>	<i>14 (3.1, 128)</i>	<i>11 (0.4, 1800)</i>

- Our approach can easily be adopted to CML log reg

## Example 5: Childhood leukemia matched-pairs study

- Ebi et al, Epidemiology 1999; Greenland, 2000
- Case: house with an index case of leukemia
- Control: reflection of case house across the street
- Exposure: backyard power line (3-phase, secondary, none)
- Standard analysis via conditional maximum likelihood (CML) log reg

# Example 5: Childhood leukemia matched-pairs study

Case exposure	Control exposure		
	Three-phase	Secondary	None
Three-phase	15	24	11
Secondary	11	107	9
None	0	1	81

# Example 5: Childhood leukemia matched-pairs study

- No separation, but sparse data: CML expected to be biased away from 0

Method	OR (95% CI)	
	Three-phase vs none	Secondary vs none
CML	32 (4.0, 0.253)	14 (1.8, 107)
CXL	30 (4.5, 1328)	14 (2.1, 507)
CPML	21 (3.7, 124)	9.6 (2.4, 87)
Haldane (add ½ to cells)	16 (3.5, 78)	7.4 (1.6, 34)
Laplace (add 1 to cells)	11 (2.9, 43)	5.2 (1.4, 19)

CML: conditional ML; CXL: conditional exact ML;  
 CPML: conditional penalized ML

# Penalized likelihood/bias reduction

- $\log L^* = \log L + c \log \det (I)$
- Jeffreys prior:  $c = \frac{1}{2}$ , removes  $O(1/n)$  bias
  - shrinks parameter estimates toward the point of minimum variance
  - shrinkage not equal for each parameter
- $c > \frac{1}{2}$ : reduces bias on  $\exp(\beta)$  scale, but introduces negative bias on estimate of  $\beta$  (Greenland, 2000)
  - $c=1$ : generalization of Laplace ad-hoc estimator



# Comments on bias reduction

- Firth:  $O(n^{-1})$ -bias reduction
  - Optimal to reduce bias and MSE
- Should we try to obtain higher-order bias reduction?
  - $MSE = \text{bias}^2 + \text{variance}$
  - bias smaller, but variance greater  $\Rightarrow$  MSE worse



# Bias reduction if $n/k$ is small

- Since estimates exist in each and every situation, we are seduced to analyze samples with very small  $n$  and very large  $k$
- Watch out! Firth's bias reduction overcorrects bias if  $n/k$  is small
- This means, a negative bias (bias towards zero) is introduced
- Overcorrection becomes severe (effects are halvened) if  $y$  is unbalanced AND  $n/k$  is small



# Why the overcorrection?

- Bias reduction implicitly estimates bias  $b(\beta)$  replacing  $\beta$  by its estimate
- If estimate is inconsistent, approximation fails (Leung and Wang, 1998)
- Common to all bias correcting approaches that need to estimate the bias
- Pseudo-observations obtain too much weight ( $k$  compared to  $n$ )





## Bias reduction if $n/k$ is small (2)

- Smaller amount of correction necessary
- Maximize  $\text{Log } L^* = \log L + c \log \det (I)$  with  $c < \frac{1}{2}$
- Optimization of  $c$  via simulation (?)
- Or switch to ridge regression (leCessie and Van Houwelingen, 1992)

# Penalized likelihood: ridge regression

- leCessie and Van Houwelingen, JRSS C 1992:
- $\text{Log } L^* = \log L - \lambda \|\beta\|^2$ 
  - $\|\beta\|^2 = (\sum \beta_j^2)^{1/2}$
  - $\lambda$  optimized to yield small prediction error
  - Shrinks parameter estimates towards 0
  - Can be used to apply restrictions on estimation, e.g.
    - Smooth transitions of parameter estimates corresponding to neighbouring categories
    - Smooth transitions of hazards in piecewise exponential models
  - Purpose: primarily for prediction, not for estimation of parameter estimates
  - In case of separation, produces finite estimates

# Model comparison with penalized maximum likelihood

- Comparison of models is difficult:
  - Hierarchical models: use penalization term of larger model
    - Model 1:  $y=A+B+C$
    - Model 2:  $y=A+B$
    - $\text{Log } L1^* := \log L(A,B,C) + \frac{1}{2} \log \det I(A,B,C)$
    - $\text{Log } L2^* := \log L(A,B,C) + \frac{1}{2} \log \det I(A,B,C)$  with  $\beta_C=0$
    - Please note that in this comparison,  
 $\text{Log } L2^* \neq \log L(A,B) + \frac{1}{2} \log \det I(A,B)$  !!!
    - That's how inference is performed in our programs

# Model comparison (2)

- Non-hierarchical model comparison
  - Model 1:  $y=A+B+C$
  - Model 2:  $y=A+B+D$
- Penalized likelihoods cannot be compared, because the structure of penalization term  
 $\frac{1}{2} \log \det (I)$  is not comparable
- Comparison via some information criterion (DIC? BIC?)
- Still an open issue

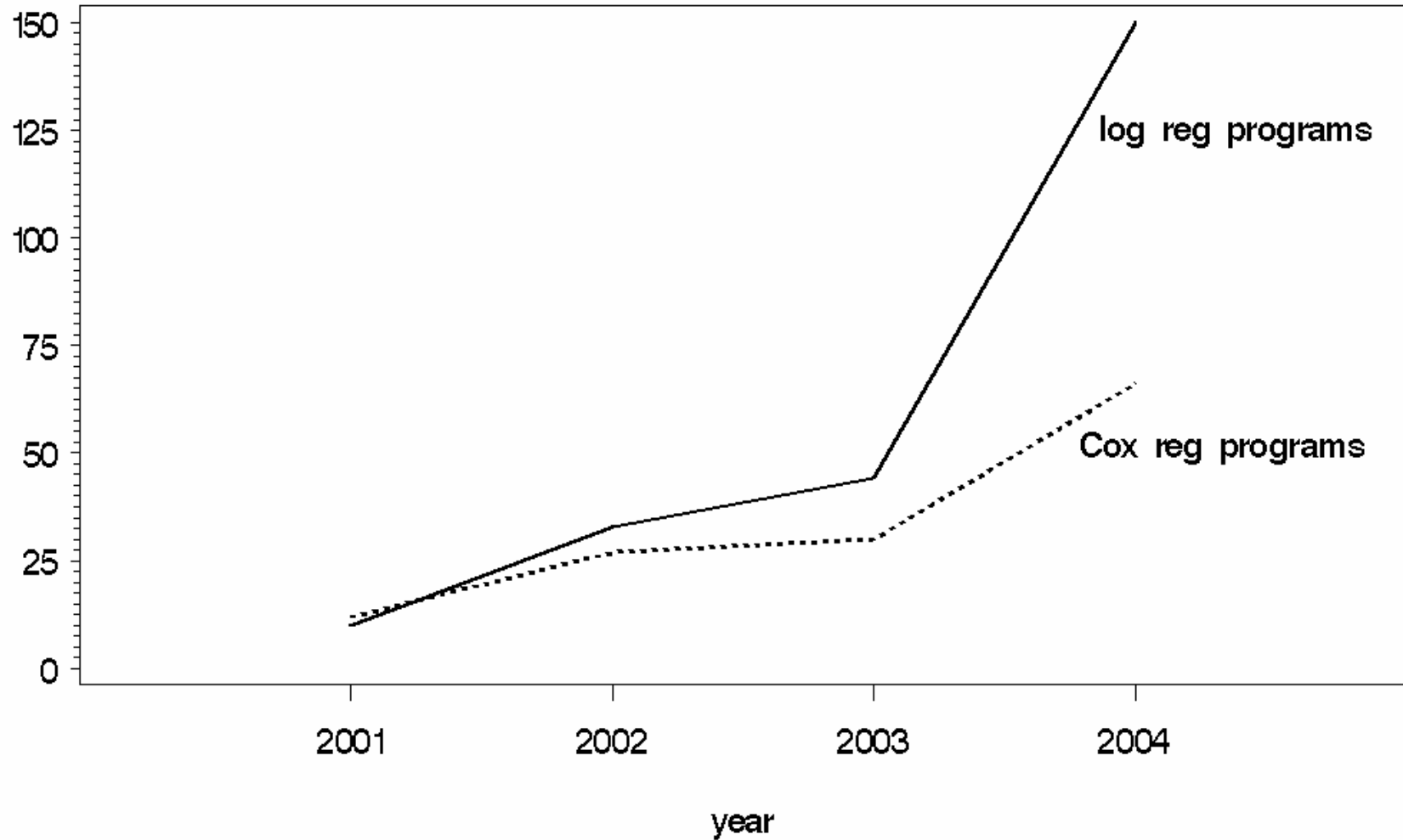


# Software

- Cox regression
  - SAS macro **FC** (Heinze and Ploner, 2002)
  - SPLUS function **coxphf** (Heinze and Ploner, 2002)
  - New SAS macro **FGCSS** (time-dependent variables/effects, counting process style, stratified analysis)
  - All based on FORTRAN
- Logistic regression
  - SAS macro **FL** (Heinze and Ploner, 2003)
  - SPLUS function **logistf** (Heinze and Ploner, 2003)
  - R package **logistf** (Heinze and Ploner, 2004)
  - R package **brlr** (by D. Firth)
- Conditional logistic regression:
  - 1:1 matching: FL/logistf, suppress estimation of intercept
  - 1:m matching: FGCSS

# Software: registered downloads

downloads



# Conclusions



- Penalized likelihood approach removes the problem of reporting infinite odds/risk ratios
- PPL confidence intervals account properly for asymmetry of likelihood
- PML estimates have smaller bias than ML estimates
- PPL confidence intervals have better coverage than PL or Wald ci
- Approach works, better than others, for all normal problems
- Software is available, have a look at
  - [www.muw.ac.at/msi/biometrie/programme/fl](http://www.muw.ac.at/msi/biometrie/programme/fl)
  - [www.muw.ac.at/msi/biometrie/programme/fc](http://www.muw.ac.at/msi/biometrie/programme/fc)

# References

- Day and Kerridge (Biometrics 1967). A general maximum likelihood discriminant.
- Bryson and Johnson (Technometrics 1981). The incidence of monotone likelihood in Cox regression.
- Clogg, Rubin, Schenker, Schultz and Weidman (J Am Stat Assoc 1991). Multiple imputation of industry and occupation codes in census public-use samples using Bayesian logistic regression.
- Le Cessie and van Houwelingen (Applied Statistics 1992). Ridge estimators in logistic regression.
- Firth (Biometrika 1993). Bias reduction of maximum likelihood estimates.
- Leung and Wang (Austr New Zeal Journal of Statistics 1998). Bias reduction using stochastic approximation.
- Greenland (Biostatistics 2000). Small-sample bias and corrections for conditional maximum-likelihood odds-ratio estimators.
- **Heinze and Schemper (Biometrics 2001). A solution to the problem of monotone likelihood in Cox regression.**
- **Heinze and Schemper (Statistics in Medicine 2002). A solution to the problem of separation in logistic regression.**
- Heinze and Ploner (Comp Meth Prog Biomed 2002). SAS and SPLUS programs to perform Cox regression without convergence problems.
- Heinze, Gnant and Schemper (Biometrics 2003). Exact logrank tests for unequal follow-up.
- Heinze and Ploner (Comp Meth Prog Biomed 2003). Fixing the nonconvergence bug in logistic regression using SPLUS and SAS.
- Heinze and Ploner (TechRep, 2004). A SAS macro, SPLUS library and R package to perform logistic regression without convergence problems.  
[http://www.meduniwien.ac.at/msi/biometrie/programme/fl/tr2\\_2004.pdf](http://www.meduniwien.ac.at/msi/biometrie/programme/fl/tr2_2004.pdf)