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### Nonparametric computation of survival functions in the presence of interval censoring

Stephen M. Taylor

MSc Candidate Supervisor: Dr Yong Wang Department of Statistics The University of Auckland stay020@aucklanduni.ac.nz

1 September 2008

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Research Aims

- Create a robust algorithm for solving the NPMLE problem
- One that is fastest in all circumstances

Adaptive Constrained Newton Method (ACNM)

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### Survival Analysis

Time to event data

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#### Survival Analysis

- Time to event data
- Want to model the distribution of times to 'failure'

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### Survival Analysis

- Time to event data
- Want to model the distribution of times to 'failure'
- Interested in the survival function, S(t) = P(T > t)

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#### Survival Analysis

- Time to event data
- Want to model the distribution of times to 'failure'
- Interested in the survival function, S(t) = P(T > t)
- Example: Time to healing

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#### Censoring

#### • Time of event may not be directly measurable

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- Time of event may not be directly measurable
- Check periodically to see if it has occurred

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- Time of event may not be directly measurable
- Check periodically to see if it has occurred
- Example: healing occurred some time between doctor visits

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- Time of event may not be directly measurable
- Check periodically to see if it has occurred
- Example: healing occurred some time between doctor visits
- The event may never occur for some subjects

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- Time of event may not be directly measurable
- Check periodically to see if it has occurred
- Example: healing occurred some time between doctor visits
- The event may never occur for some subjects
- Example: end of study or "lost to followup"

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#### Interval Censoring

• Event times are not known exactly, only within intervals

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- Event times are not known exactly, only within intervals
- · Perhaps no event time is observed exactly

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- Event times are not known exactly, only within intervals
- Perhaps no event time is observed exactly
- Interval censored: event occurred somewhere in  $(t_L, t_R]$

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- Event times are not known exactly, only within intervals
- Perhaps no event time is observed exactly
- Interval censored: event occurred somewhere in  $(t_L, t_R]$
- Right censored:  $(t_L,\infty)$

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- Event times are not known exactly, only within intervals
- Perhaps no event time is observed exactly
- Interval censored: event occurred somewhere in  $(t_L, t_R]$
- Right censored:  $(t_L,\infty)$
- Left censored:  $(0, t_R]$

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- Event times are not known exactly, only within intervals
- Perhaps no event time is observed exactly
- Interval censored: event occurred somewhere in  $(t_L, t_R]$
- Right censored:  $(t_L,\infty)$
- Left censored:  $(0, t_R]$
- Exact observation: event occurred at time t

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- Event times are not known exactly, only within intervals
- Perhaps no event time is observed exactly
- Interval censored: event occurred somewhere in  $(t_L, t_R]$
- Right censored:  $(t_L,\infty)$
- Left censored:  $(0, t_R]$
- Exact observation: event occurred at time t
- Call these intervals  $O_i$  for  $i = 1, \ldots, n$

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Why Nonparametric?

• Let the data speak for itself

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Why Nonparametric?

- Let the data speak for itself
- Don't make assumptions about the distribution

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Why Nonparametric?

- Let the data speak for itself
- Don't make assumptions about the distribution
- Maximise the likelihood

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Why Nonparametric?

- Let the data speak for itself
- Don't make assumptions about the distribution
- Maximise the likelihood
- Explore the data before choosing a parametric model

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Summing Up 00

# The NPMLE Survival Function with Interval Censored Data

• Partition the positive real line





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# The NPMLE Survival Function with Interval Censored Data

- Partition the positive real line
- All unique values of  $t_L$  and  $t_R$





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# The NPMLE Survival Function with Interval Censored Data

- Partition the positive real line
- All unique values of  $t_L$  and  $t_R$
- Potential support intervals



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# The NPMLE Survival Function with Interval Censored Data

- Partition the positive real line
- All unique values of  $t_L$  and  $t_R$
- Potential support intervals
- Only use maximal cliques



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Summing Up

The NPMLE Survival Function with Interval Censored Data

- Partition the positive real line
- All unique values of  $t_L$  and  $t_R$
- Potential support intervals
- Only use maximal cliques
- Support set:  $I_j$  for  $j = 1, \ldots, m$



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Summing Up

The NPMLE Survival Function with Interval Censored Data

- Partition the positive real line
- All unique values of  $t_L$  and  $t_R$
- Potential support intervals
- Only use maximal cliques
- Support set:  $I_j$  for  $j = 1, \ldots, m$
- The clique matrix A<sub>n×m</sub> gives δ<sub>ij</sub> membership of each O<sub>i</sub> in each I<sub>j</sub>



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The NPMLE Survival Function with Interval Censored Data

- Partition the positive real line
- All unique values of  $t_L$  and  $t_R$
- Potential support intervals
- Only use maximal cliques
- Support set:  $I_j$  for  $j = 1, \ldots, m$
- The clique matrix A<sub>n×m</sub> gives δ<sub>ij</sub> membership of each O<sub>i</sub> in each I<sub>j</sub>
- NPMLE assigns probability mass to each support interval



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#### Likelihood Function for the NPMLE

• Likelihood of an interval  $(t_1, t_2]$  is  $S(t_1) - S(t_2)$ 



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- Likelihood of an interval  $(t_1, t_2]$  is  $S(t_1) S(t_2)$
- Assign probability p<sub>j</sub> to support interval I<sub>j</sub>

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- Likelihood of an interval  $(t_1, t_2]$  is  $S(t_1) S(t_2)$
- Assign probability  $p_j$  to support interval  $I_j$
- Probability of observation O<sub>i</sub> using A and **p**

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- Likelihood of an interval  $(t_1, t_2]$  is  $S(t_1) S(t_2)$
- Assign probability  $p_j$  to support interval  $I_j$
- Probability of observation O<sub>i</sub> using A and **p**
- Take logs and add them up

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- Likelihood of an interval  $(t_1, t_2]$  is  $S(t_1) S(t_2)$
- Assign probability  $p_j$  to support interval  $I_j$
- Probability of observation O<sub>i</sub> using A and **p**
- Take logs and add them up
- Goal: find  $\hat{\mathbf{p}} \in \mathbb{R}^m$  to maximise  $\ell(\hat{\mathbf{p}})$
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Likelihood Function for the NPMLE

- Likelihood of an interval  $(t_1, t_2]$  is  $S(t_1) S(t_2)$
- Assign probability p<sub>j</sub> to support interval I<sub>j</sub>
- Probability of observation O<sub>i</sub> using A and **p**
- Take logs and add them up
- Goal: find  $\hat{\mathbf{p}} \in \mathbb{R}^m$  to maximise  $\ell(\hat{\mathbf{p}})$
- Subject to:  $\mathbf{\hat{p}} \ge \mathbf{0}$  and  $\mathbf{\hat{p}}^{T}\mathbf{1} = 1$

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Honey as Adjuvant Leg Ulcer Therapy (HALT)

• Randomised Clinical Trial, 368 participants

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- Randomised Clinical Trial, 368 participants
- Clinical Trials Research Unit in Auckland

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- Randomised Clinical Trial, 368 participants
- Clinical Trials Research Unit in Auckland
- Effect of Manuka Honey dressings for treatment of leg ulcers

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- Randomised Clinical Trial, 368 participants
- Clinical Trials Research Unit in Auckland
- Effect of Manuka Honey dressings for treatment of leg ulcers
- Participants assessed weekly and also at a 12-week follow-up

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- Randomised Clinical Trial, 368 participants
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- Participants assessed weekly and also at a 12-week follow-up
- Nurse changes dressing and assesses healing status

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- Randomised Clinical Trial, 368 participants
- Clinical Trials Research Unit in Auckland
- Effect of Manuka Honey dressings for treatment of leg ulcers
- Participants assessed weekly and also at a 12-week follow-up
- Nurse changes dressing and assesses healing status
- Event times cannot be observed exactly

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- Randomised Clinical Trial, 368 participants
- Clinical Trials Research Unit in Auckland
- Effect of Manuka Honey dressings for treatment of leg ulcers
- Participants assessed weekly and also at a 12-week follow-up
- Nurse changes dressing and assesses healing status
- Event times cannot be observed exactly
- Thanks to Andrew Jull and Varsha Parag of CTRU for providing the data

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#### **Censor Intervals for each Participant**



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# Existing Algorithms for finding the NPMLE

• The Icens package in R provides five algorithms:

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- The Icens package in R provides five algorithms:
  - EM, ISDM, EMICM, VEM and PGM

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- The Icens package in R provides five algorithms:
  - EM, ISDM, EMICM, VEM and PGM
- Subspace-based Newton method (Dümbgen et al. 2006)

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- The Icens package in R provides five algorithms:
  - EM, ISDM, EMICM, VEM and PGM
- Subspace-based Newton method (Dümbgen et al. 2006)
- Wang (2008) introduced:

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- The Icens package in R provides five algorithms:
  - EM, ISDM, EMICM, VEM and PGM
- Subspace-based Newton method (Dümbgen et al. 2006)
- Wang (2008) introduced:
  - Constrained Newton Method

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- The Icens package in R provides five algorithms:
  - EM, ISDM, EMICM, VEM and PGM
- Subspace-based Newton method (Dümbgen et al. 2006)
- Wang (2008) introduced:
  - Constrained Newton Method
  - Dimension-reduced approach to improve any algorithm

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Times to compute the NPMLE survival function for 100 Bootstrap samples of the HALT data using:

•	EMICM, PGM and VEM from the	
	Icens package	E
•	Methods SBN(DR) and	
	EMICM(DR) from Wang (2008)	

• The new ACNM algorithm (and CNM)

	Time (s)
EMICM	113.03
PGM	791.00
VEM	610.42
SBN(DR)	14.34
EMICM(DR)	26.93
ACNM	9.41

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#### Problems with Existing Algorithms

• Some are very slow and may fail to converge

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Problems with Existing Algorithms

- Some are very slow and may fail to converge
- No algorithm outperforms the others in all situations

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Problems with Existing Algorithms

- Some are very slow and may fail to converge
- No algorithm outperforms the others in all situations
- Inefficent use of Hessian matrix or gradient

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Problems with Existing Algorithms

- Some are very slow and may fail to converge
- No algorithm outperforms the others in all situations
- Inefficent use of Hessian matrix or gradient
- Best choice depends on size of dataset and proportion of exact observations

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#### Constrained Newton Method

• Calculates gradient S of  $\ell(\mathbf{p})$  at current estimate  $\mathbf{p}$ 

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- Calculates gradient S of  $\ell(\mathbf{p})$  at current estimate  $\mathbf{p}$
- Makes use of mixture structure of solution

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- Calculates gradient S of  $\ell(\mathbf{p})$  at current estimate  $\mathbf{p}$
- Makes use of mixture structure of solution
- Uses NNLS to find new estimate of **p**

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- Calculates gradient S of  $\ell(\mathbf{p})$  at current estimate  $\mathbf{p}$
- Makes use of mixture structure of solution
- Uses NNLS to find new estimate of **p**
- Computation time of NNLS is of order  $O(nm^2)$

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- Calculates gradient S of  $\ell(\mathbf{p})$  at current estimate  $\mathbf{p}$
- Makes use of mixture structure of solution
- Uses NNLS to find new estimate of **p**
- Computation time of NNLS is of order  $O(nm^2)$
- Very fast for fully censored datasets

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- Calculates gradient S of  $\ell(\mathbf{p})$  at current estimate  $\mathbf{p}$
- Makes use of mixture structure of solution
- Uses NNLS to find new estimate of **p**
- Computation time of NNLS is of order  $O(nm^2)$
- Very fast for fully censored datasets
- · Can be slow in cases with many exact observations

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# Adaptive CNM

• Uses a divide and conquer approach

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- Uses a divide and conquer approach
- Breaks the support set up into blocks

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- Uses a divide and conquer approach
- Breaks the support set up into blocks
- Adapts to the data to make efficient use of Hessian

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- Uses a divide and conquer approach
- Breaks the support set up into blocks
- Adapts to the data to make efficient use of Hessian
- Examines data to choose number/size of blocks

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- Uses a divide and conquer approach
- Breaks the support set up into blocks
- Adapts to the data to make efficient use of Hessian
- Examines data to choose number/size of blocks
- Solves each block using NNLS

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- Uses a divide and conquer approach
- Breaks the support set up into blocks
- Adapts to the data to make efficient use of Hessian
- Examines data to choose number/size of blocks
- Solves each block using NNLS
- Globally reallocates probability among blocks, calling itself recursively

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- Uses a divide and conquer approach
- Breaks the support set up into blocks
- Adapts to the data to make efficient use of Hessian
- Examines data to choose number/size of blocks
- Solves each block using NNLS
- Globally reallocates probability among blocks, calling itself recursively
- Guaranteed convergence to the solution

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# Heatmap of HALT Hessian



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## Conclusions

- Where Interval Censoring is present in survival data, it can be allowed for in the analysis.
- The NPMLE Survival Function combined with Bootstrap methods can create an informative picture of survival progression in such cases.
- The ACNM algorithm provides a fast and robust solution to this problem.

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Thanks to:

- My supervisor, Dr Yong Wang
- Andrew Jull and Varsha Parag of CTRU for providing the HALT data