

# Algorithmes efficaces pour la détection de ruptures

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Joint work with Charles Truong, Guillem Rigai, Vincent Runge



## Brief history of optimal change-point algorithms

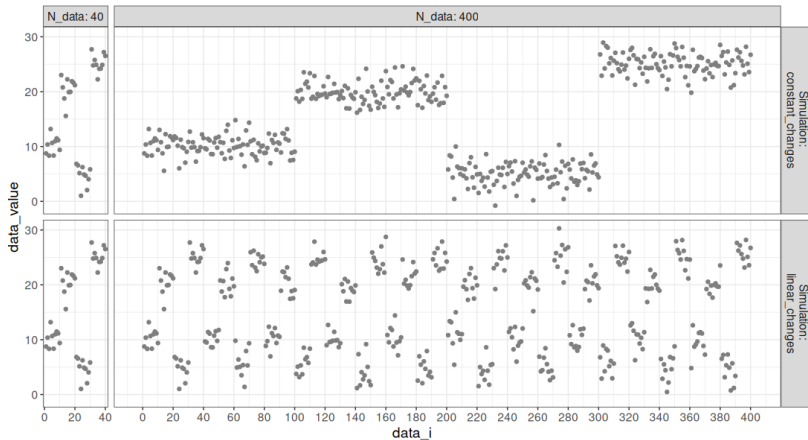
New algorithms with constraints between adjacent segments

Computing optimal changepoints subject to label constraints

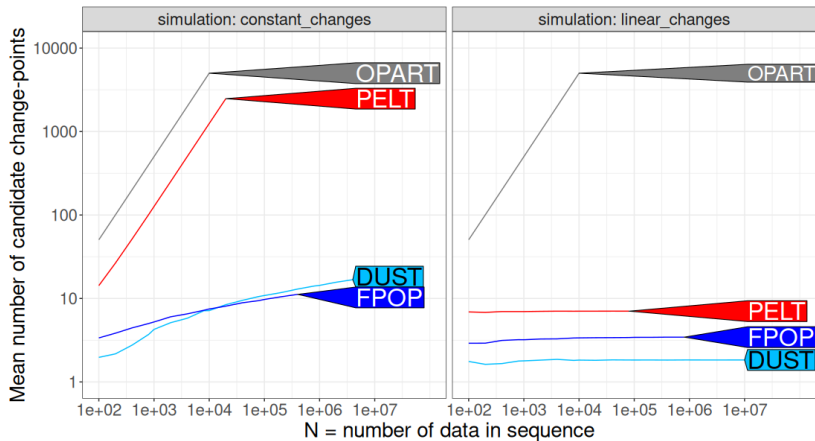
Learning to predict the number of changepoints

Summary and Discussion

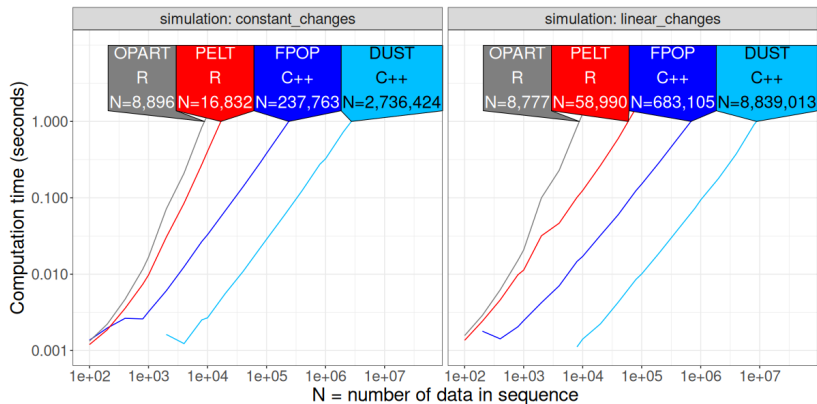
# Simulated data with constant and linear changes



# Number of candidates considered



# Overall computation time



Brief history of optimal change-point algorithms

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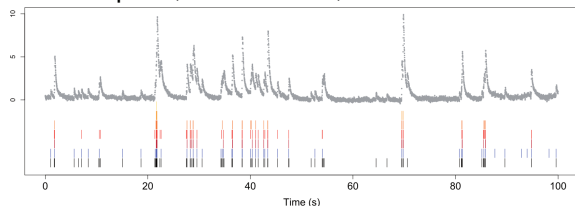
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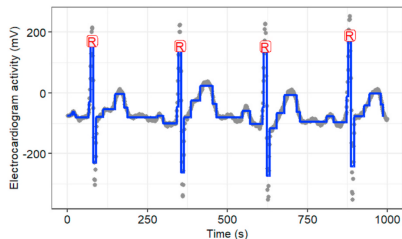
Summary and Discussion

# Changepoint detection algorithms for data over time

Neuron spikes, Jewell *et al.*, Biostatistics 2019.

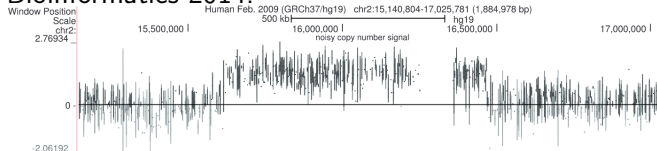


Electrocardiograms (heart monitoring), Fotoohinasab *et al.*, Asilomar conference 2020.

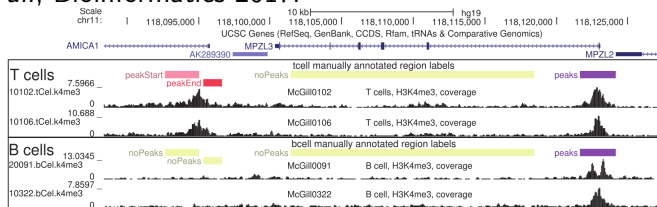


# Changepoint detection algorithms for data over space

DNA copy number data for cancer diagnosis, Hocking *et al.*, Bioinformatics 2014.

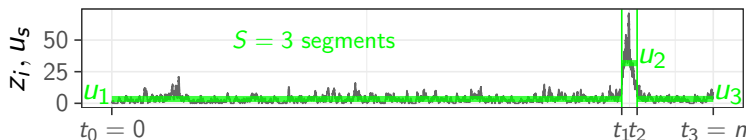


ChIP-seq data for understanding the human genome, Hocking *et al.*, Bioinformatics 2017.





# Optimal changepoint detection problem and algorithms



$$\min_{\substack{\mathbf{u} \in \mathbb{R}^S \\ 0=t_0 < t_1 < \dots < t_{S-1} < t_S=n}} \sum_{s=1}^S \sum_{i=t_{s-1}+1}^{t_s} \ell(u_s, z_i)$$

- ▶ Algorithm inputs  $n$  data  $z_1, \dots, z_n$  and  $\#$  of segments  $S$ .
- ▶ Goal is to compute best  $S - 1$  changepoints  $t_1 < \dots < t_{S-1}$  and  $S$  segment parameters  $u_1, \dots, u_S$ .
- ▶ Hard non-convex optimization problem, naïvely  $O(n^S)$  time.
- ▶ Auger and Lawrence 1989:  $O(Sn^2)$  time algorithm.
- ▶ Rigaiil 2015:  $O(n \log n)$  time, change in any direction.
- ▶ Hocking *et al.*, 2020:  $O(n \log n)$ , directional constraints.

# Constrained optimization algorithm speed

Hocking *et al.*, Journal of Machine Learning Research 2020.

$$\begin{aligned} & \min_{\substack{u \in \mathbb{R}^S \\ 0=t_0 < t_1 < \dots < t_{S-1} < t_S=n}} \sum_{s=1}^S \sum_{i=t_{s-1}+1}^{t_s} \ell(u_s, z_i) \\ & \text{subject to} \quad u_{s-1} \leq u_s \quad \forall s \in \{2, 4, \dots\}, \\ & \quad \quad \quad u_{s-1} \geq u_s \quad \forall s \in \{3, 5, \dots\}. \end{aligned}$$

Constraints used to force change up to peak state, then change down to background noise state.

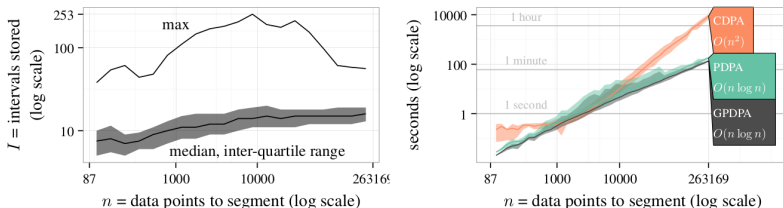
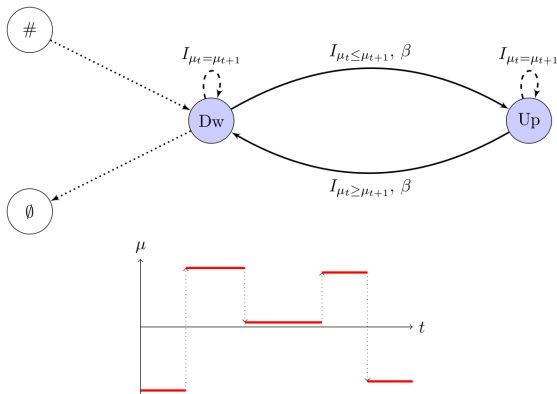


Figure 3: Empirical speed analysis on 2752 count data vectors from the histone mark ChIP-seq benchmark. For each vector we ran the GPDPA with the up-down constraint and a max of  $K = 19$  segments. The expected time complexity is  $O(KnI)$  where  $I$  is the average number of intervals (function pieces; candidate changepoints) stored in the  $C_{k,t}$  cost functions. **Left:** number of intervals stored is  $I = O(\log n)$  (median, inter-quartile range, and maximum over all data points  $t$  and segments  $k$ ). **Right:** time complexity of the GPDPA is  $O(n \log n)$  (median line and min/max band).

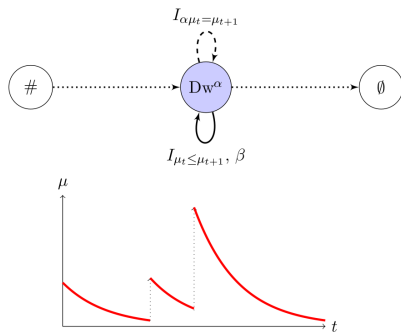
# Optimization constraints defined using a graph

Runge *et al.*, Pre-print arXiv:2002.03646 (graph figures).

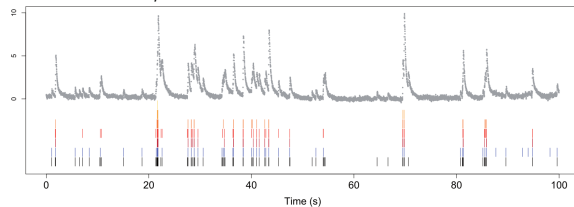


- ▶ Purple Dw/Up nodes represent hidden states.
- ▶  $\#/\emptyset$  nodes constrain start/end state.
- ▶ Edges represent possible state transitions.
- ▶ gfpop R package with C++ code computes optimal changepoints for user-defined constraint graphs.

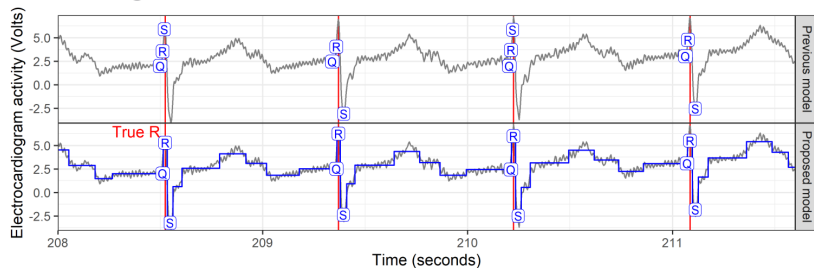
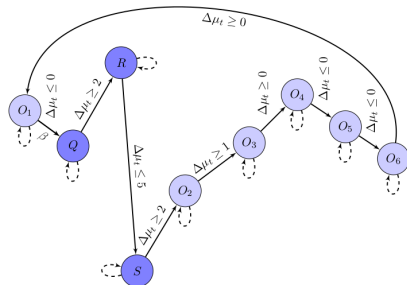
# All up changes, exponentially decaying segments



Jewell *et al.*, Biostatistics 2019.



# Complex graph for electrocardiogram data



Fotoohinasab *et al.*, Asilomar conference 2020.

# References

- ▶ Auger IE and Lawrence CE. Algorithms for the optimal identification of segment neighborhoods. *Bull Math Biol* 51:39–54 (1989).
- ▶ G Rigaiil. A pruned dynamic programming algorithm to recover the best segmentations with 1 to kmax change-points. *Journal de la Société Française de la Statistique*, 156(4), 2015.
- ▶ **Hocking TD**, Boeva V, Rigaiil G, Schleiermacher G, Janoueix-Lerosey I, Delattre O, Richer W, Bourdeaut F, Suguro M, Seto M, Bach F, Vert J-P. SegAnnDB: interactive Web-based genomic segmentation. *Bioinformatics* (2014) 30 (11): 1539-1546.
- ▶ **Hocking TD**, Goerner-Potvin P, Morin A, Shao X, Pastinen T, Bourque G. Optimizing ChIP-seq peak detectors using visual labels and supervised machine learning. *Bioinformatics* (2017) 33 (4): 491-499.
- ▶ Jewell S, **Hocking TD**, Fearnhead P, Witten D. Fast Nonconvex Deconvolution of Calcium Imaging Data. *Biostatistics* (2019).
- ▶ Fotoohinasab A, **Hocking TD**, Afghah F. A Graph-Constrained Changepoint Learning Approach for Automatic QRS-Complex Detection. *Asilomar Conference on Signals, Systems, and Computers* (2020).
- ▶ **Hocking TD**, Rigaiil G, Fearnhead P, Bourque G. Constrained Dynamic Programming and Supervised Penalty Learning Algorithms for Peak Detection in Genomic Data. *Journal of Machine Learning Research* 21(87):1–40, 2020.
- ▶ Runge V, **Hocking TD**, Romano G, Afghah F, Fearnhead P, Rigaiil G. gfpop: an R Package for Univariate Graph-Constrained Change-point Detection. Under review at *Journal of Statistical Software*. Pre-print arXiv:2002.03646.

Brief history of optimal change-point algorithms

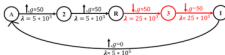
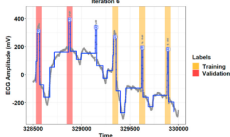
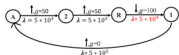
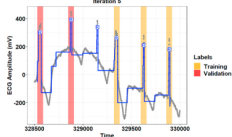
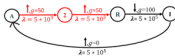
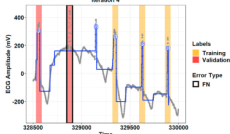
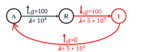
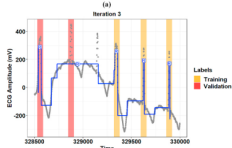
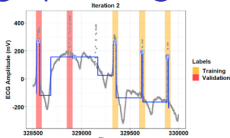
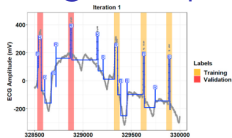
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Summary and Discussion

# Learning a complex graph using labels

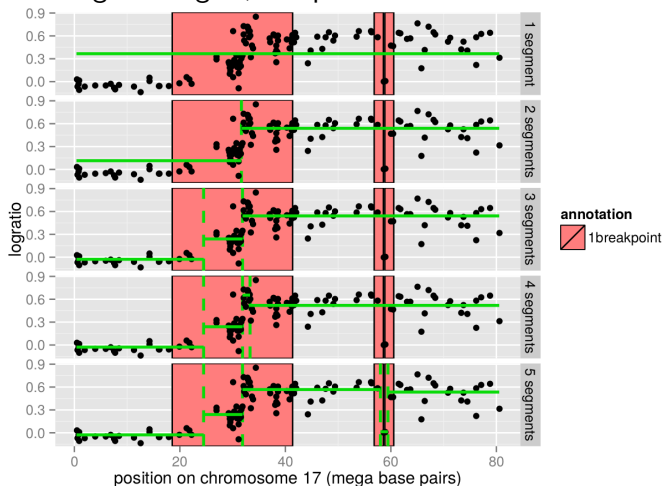


- Fotoohinasab *et al.*, 2021.
- Simple initial graph is iteratively edited (red) to agree with expert labeled regions (orange rectangles).
- Easier for expert to provide labels than graph.



# What if no models agree with expert labels?

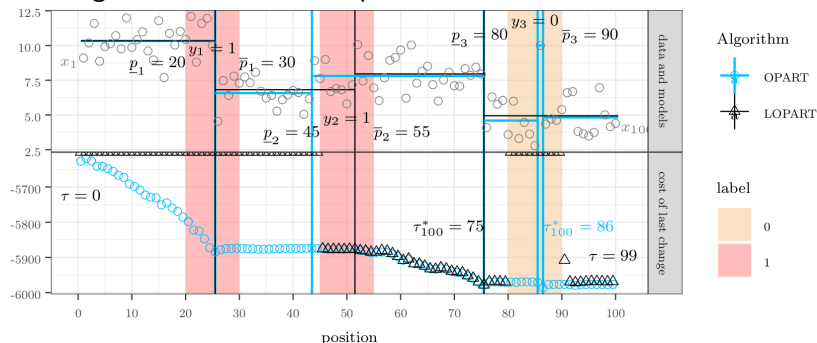
Hocking and Rigaiil, Pre-print hal-00759129.



- ▶ Expert wants: one changepoint in each label (red rectangle).
- ▶ No model is consistent with all three labels.

# Using expert labels as optimization constraints

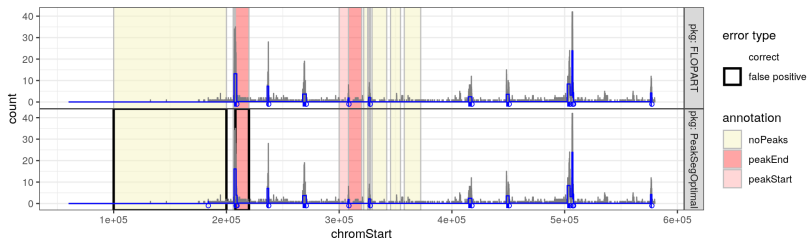
Hocking and Srivastava, Pre-print arXiv:2006.13967.



- ▶ Previous OPART model (blue) ignores labels (two errors).
- ▶ Main idea: add optimization constraints to ensure that there is the right number of changepoints predicted in each label.
- ▶ Proposed LOPART model (black) consistent with labels.

# Label constraints and directional constraints

Stenberg and Hocking, in progress.



- ▶ Previous PeakSegOptimal algorithm (bottom) ignores labels (two errors).
- ▶ Proposed FLOPART model (top) consistent with labels, and interpretable in terms of up changes to peaks and down changes to background noise.

# References

- ▶ **Hocking TD**, Rigai G. SegAnnot: an R package for fast segmentation of annotated piecewise constant signals, Pre-print hal-00759129.
- ▶ **Hocking TD**, Srivastava A. Labeled Optimal Partitioning. Under review at Computational Statistics. Pre-print arXiv:2006.13967.
- ▶ Fotoohinasab A, **Hocking TD**, Afghah F. A Greedy Graph Search Algorithm Based on Change-point Analysis for Automatic QRS-Complex Detection. Computers in Biology and Medicine 130 (2021).

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# How to predict the number of changes?

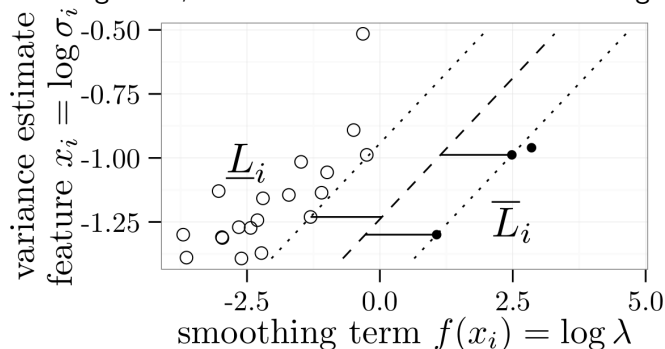
We assumed that the number of segments  $S$  is provided as an input parameter to our optimization algorithm.

$$\min_{\substack{\mathbf{u} \in \mathbb{R}^S \\ 0=t_0 < t_1 < \dots < t_{S-1} < t_S=n}} \sum_{s=1}^S \sum_{i=t_{s-1}+1}^{t_s} \ell(u_s, z_i)$$

In practice  $S$  is often unknown — what value should we use?

# Learning to predict number of changes similar to SVM

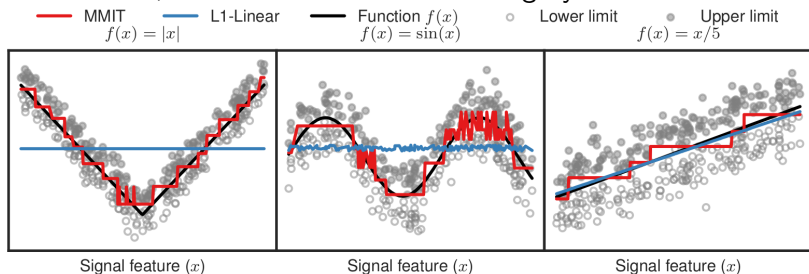
Hocking *et al.*, Int'l Conference on Machine Learning 2013.



- ▶ Train on several data sequences with labels (dots).
- ▶ Want to compute function between white and black dots.
- ▶ SVM margin is multi-dimensional (diagonal).
- ▶ Here margin to maximize is one-dimensional (horizontal).
- ▶ Learned function predicts number of changepoints/segments.

# Decision tree learns non-linear function of inputs

Drouin *et al.*, Neural Information Processing Systems 2017.

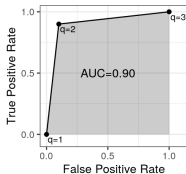


- ▶ Generalization of classical CART regression tree learning algorithm.
- ▶ Can learn non-linear functions of inputs.
- ▶ More recently we implemented a similar idea in xgboost, Barnwal *et al.*, Pre-print arXiv:2006.04920.

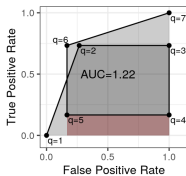


# Is maximizing Area Under the ROC Curve desirable?

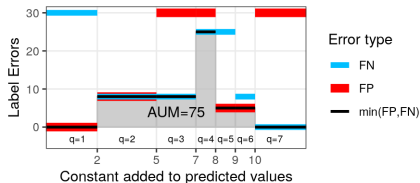
In binary classification the ROC curve is monotonic.



In changepoint detection it can have loops.

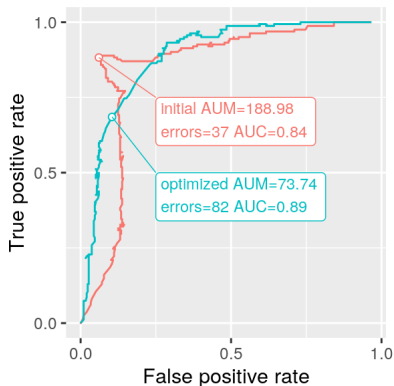
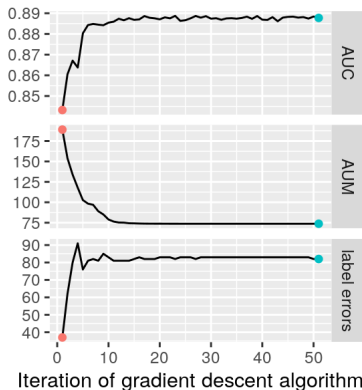


We propose instead to minimize the  $AUM = \text{Area Under the Minimum of false positives and false negatives}$ , as a function of prediction threshold.



# AUM gradient descent algorithm optimizes AUC

Hillman and Hocking, in progress.



- ▶ Initial predictions: minimum label errors.
- ▶ ROC curves become more regular/monotonic after optimization, but label error increases.
- ▶ Trade-off between AUC and label error optimization that does not exist in binary classification.

# References

- ▶ **Hocking TD**, Rigai G, Bach F, Vert J-P. Learning sparse penalties for change-point detection using max-margin interval regression. International Conference on Machine Learning 2013.
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- ▶ Hillman J, **Hocking TD**. Optimizing ROC Curves with a Sort-Based Surrogate Loss Function for Binary Classification and Change-point Detection. Preprint arXiv:2107.01285.

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# Summary and Discussion

- ▶ Optimal changepoint detection in  $n$  data is a non-convex problem, naively a  $O(n^S)$  computation for  $S$  segments.
- ▶ Recent algorithms can compute a globally optimal changepoint model much faster,  $O(n \log n)$ .
- ▶ Directional constraint graphs specified using domain prior knowledge, or learned using expert labels.
- ▶ Expert labels can also be used as optimization constraints, to ensure that predicted changepoints are consistent.
- ▶ Number of changes can be predicted with new learning algorithms, including ROC curve optimization.
- ▶ Let's collaborate! [toby.hocking@nau.edu](mailto:toby.hocking@nau.edu)