Controlled Discovery and Localization of Astronomical Point Sources via Bayesian Linear Programming (BLiP)

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Lucas Janson (Harvard Statistics)

BLiP: Signal Discovery and Localization

- Motivation
- Problem statement
- Methodological contribution: Bayesian Linear Programming (BLiP)
- Simulations
- Application to genetic fine-mapping
- Application to astronomical point-source detection

Astronomical point source detection



Figure: Cartoon of partial point source data

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Solution: adaptively selected regions

Don't want to narrow potential discovery regions until after seeing data

$$\begin{split} \max & \quad \mathbb{E}\left[\mathsf{Power}(G_1,\ldots,G_R)\right] \\ \text{s.t.} & \quad \mathsf{FDR} := \mathbb{E}\left[\frac{\#\{G_r \text{ containing no signal}\}}{\max(1,R)}\right] \leq q, \\ & \quad G_1,\ldots,G_R \subset \mathcal{L} \text{ are disjoint.} \end{split}$$

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Existing work: no formalization of what "power" means, so cannot optimize it

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Optimizing resolution-adjusted power

Sum weights of true rejections to get Power():

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Then the power of a Bayesian method that discovers G_1, \ldots, G_R is

$$\mathbb{E}[\mathsf{Power}(G_1,\ldots,G_R) \mid \mathsf{Data}] = \mathbb{E}\left[\sum_{r=1}^R I_{G_r} w(G_r) \mid \mathsf{Data}\right] = \sum_{G \subseteq \mathcal{L}} p_G w(G) x_G,$$

x_G ∈ {0,1} is indicator that G is one of the method's discoveries
p_G = E[I_G | Data] is posterior inclusion probability (PIP)

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Optimal Bayesian method would solve:

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Linearizing the FDR constraint

$$\mathsf{FDR} = \frac{\sum_G (1 - p_G) x_G}{\sum_G x_G} \le q \qquad \Leftrightarrow \qquad \sum_G (1 - p_G - q) x_G \le 0$$

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Mixed-integer linear program (MILP) non-convex; fast solvers for small problems

Convexifying the integer constraint

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Empirically, Very few $x_G^{\star} \notin \{0,1\}$, and $\{x_G^{\star\star}\}$ very nearly MILP-optimal

BLiP is verifiably nearly optimal

An exact upper-bound for the suboptimality of $\{x_G^{\star\star}\}$ is

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Figure: Expected power (objective function) of $\{x_G^{\star\star}\}$ (BLiP) vs. $\{x_G^{\star}\}$ (Upper bound). Optimization dimension $\geq 50,000$.

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Can also solve problem as if $|\mathcal{G}|$ were much bigger via adaptive pruning

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Figure: p denotes dimension of linear model being fit, with n=p/2

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Figure: Linear model w/ autocorrelated X, sparsity s, sample size n, and dimension p

BLiP idea works for other error rates: local FDR, PFER, FWER



Figure: BLiP's solution indistinguishable from upper-bound for optimal solution

BLiP applies out of the box to change point detection



Figure: Green bands denote LSS+BLiP's outputted regions; left is example SuSiE fails on due to variational approximation

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Figure: 20×20 pixel sub-image; green dots = ground truth, red regions = false discoveries, blue regions = true discoveries

Point-source detection (contd)

Inverse Radius Weight Fn.



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Fine-mapping

UK Biobank data: $n \approx 337,000$, $p \approx 19,000,000$; BLiP takes $\leq 1 \text{ min}$ per trait

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Cumulative Frequency of Discovered Group Sizes



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paper available at: https://arxiv.org/abs/2203.17208 all code posted at: https://github.com/amspector100
BLiP is a powerful, principled, efficient, and flexible method for resolution-adaptive signal discovery

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all code posted at: https://github.com/amspector100

Thank you! http://lucasjanson.fas.harvard.edu ljanson@fas.harvard.edu

- Katsevich, E., Sabatti, C., and Bogomolov, M. (2021). Filtering the rejection set while preserving false discovery rate control. *Journal of the American Statistical Association*, 0(0):1–12.
- Lee, Y., Luca, F., Pique-Regi, R., and Wen, X. (2018). Bayesian multi-snp genetic association analysis: Control of fdr and use of summary statistics. *bioRxiv*.
- Liu, R., McAuliffe, J. D., and Regier, J. (2021). Variational inference for deblending crowded starfields.
- Wang, G., Sarkar, A., Carbonetto, P., and Stephens, M. (2020). A simple new approach to variable selection in regression, with application to genetic fine mapping. *Journal of the Royal Statistical Society: Series B (Statistical Methodology)*, 82(5):1273–1300.
- Yekutieli, D. (2008). Hierarchical false discovery rate-controlling methodology. *Journal of the American Statistical Association*, 103(481):309–316.