Mid-term presentation

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- Introduction
- Variational inference
- ► Mean-field approximation
- ► Implementation
- ► Results
- ► Next steps

Introduction

- We introduce $X = (X_1, \dots, X_p)$, and $y = (y_1, \dots, y_q)$.
- \triangleright SNP X_s and a trait y_t .
- Estimate the association between SNP s and trait t.
- $y_{n \times q} = x_{n \times p} \beta_{p \times q} + \epsilon_{n \times q}, \ \epsilon_t \sim \mathcal{N}(0, \tau_t^{-1} I_n)$
- y is a response matrix, x are candidate predictors.
- ► Each response y_t is linearly related with the predictors and has a residual precision $\tau_t \sim \text{Gamma}(\eta_t, \kappa_t)$.
- \triangleright p and q large compared to n.

Introduction II

- $\beta_{st} \mid \gamma_{st}, \sigma^2, \tau_t \sim \gamma_{st} \mathcal{N}(0, \sigma^2 \tau_t^{-1}) + (1 \gamma_{st}) \delta_0,$ (spike and slab)
- $ightharpoonup \gamma_{st} \mid \omega_s \sim \text{Bernoulli}(\omega_s),$
- $\triangleright \omega_s \sim \text{Beta}(a_s, b_s),$
- ▶ a_s , b_s chosen to enforce sparsity. We choose p^* the expected number of predictors involved in the model. Then:

$$a_s \equiv 1$$
, $b_s \equiv q(p-p^*)/p^*$





Variational inference

- ▶ Observed data y, parameters θ , posterior distribution of parameters $p(\theta \mid y)$.
- Approximate the posterior density with a simpler density q, minimizing a "closeness" measure: the Kullback-Leibler divergence.
- $\blacktriangleright \text{ KL}(q \parallel p) := \int q(\theta) \log \left(\frac{q(\theta)}{p(\theta \mid \mathbf{y})} \right) d\theta.$
- Evidence lower bound (ELBO): $\mathcal{L}(q) = \mathbb{E}_q \left[\log p(\theta, \mathbf{y}) \right] \mathbb{E}_q \left[\log q(\theta) \right].$
- $\blacktriangleright \operatorname{KL}(q \parallel p) = \log(p) \mathcal{L}(q).$
- Minimizing KL is equivalent to maximizing ELBO.



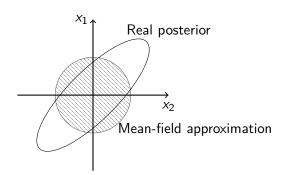


Mean-field approximation

▶ We assume independence for some of the parameters:

$$q(\boldsymbol{\theta}) = \left\{ \prod_{s=1}^p \prod_{t=1}^q q(\beta_{st}, \gamma_{st}) \right\} \left\{ \prod_{s=1}^p q(\omega_s) \right\} \left\{ \prod_{t=1}^q q(\tau_t) \right\} q(\sigma^{-2}).$$

► The mean-field approximation does not compute the correlations between parameters.





Parameters distributions

- $\blacktriangleright \ \beta_{\mathsf{st}} \mid \gamma_{\mathsf{st}} = 1, \mathbf{y} \sim \mathcal{N}\left(\mu_{\beta,\mathsf{st}}, \sigma_{\beta,\mathsf{st}}^2\right),$
- $ightharpoonup eta_{st} \mid \gamma_{st} = 0, m{y} \sim \delta_0,$
- $ightharpoonup \gamma_{st} \mid \mathbf{y} \sim \mathsf{Bernoulli}(\gamma_{st}^{(1)}),$
- $\blacktriangleright \mu_{\beta,st} = \sigma_{\beta,st}^2 \tau_t^{(1)} \boldsymbol{X}_s^T \left\{ \boldsymbol{y}_t \sum_{j=1,j\neq s}^p \gamma_{jt}^{(1)} \mu_{\beta,jt} \boldsymbol{X}_j \right\},\,$

Coordinate ascent variational inference - CAVI

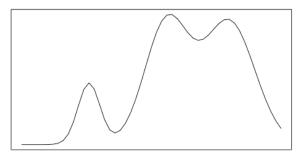
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▶ If we fix q_l(\theta_l), l \neq j, the optimal for q_i(\theta_i) verifies:
    q_i^*(\theta_i) \propto \exp\left\{\mathbb{E}_{-i}\left[\log p(\theta_i \mid \boldsymbol{\theta}_{-i}, \boldsymbol{y})\right]\right\}
▶ IN: p(x, z), data set x, tolerance tol,
    OUT: q(z) = \prod q_i(z_i).
    INIT: q_i(z_i),
    RFPFAT.
       FOR: i \in \{1, ..., m\},
           SET: q_i(z_i) \propto \exp \{\mathbb{E}_{-i} [\log p(z_i|z_{-i},x)]\}.
        COMPUTE:
           ELBO^{old}(q) \leftarrow ELBO(q).
           ELBO(q) = \mathbb{E} [\log p(z, x)] - \mathbb{E} [\log q(z)].
    UNTIL: |ELBO(q) - ELBO^{old}(q)| < tol.
    RETURN: q(z).
```





Coordinate ascent variational inference - CAVI II

- $ightharpoonup \mathcal{L}(q)$ is guaranteed to augment at every iteration.
- ► CAVI yields a local optimum, depending on the initialization of the parameters.



"Bayesian model averaging"

- ▶ Denote M_k , k = 1, ..., K the models yielded by the local optimums.

- ▶ $\mathcal{L}(q)$ serves as an approximation of $p(\mathbf{y} \mid M_k)$, as $\mathrm{KL}(q \parallel p) = \log p(\mathbf{y}) \mathcal{L}(q)$.
- ▶ $p(M_k)$ is the prior probability of the models, we consider them to be equiprobable: $p(M_k) = 1/K$, $\forall k = 1, ..., K$.

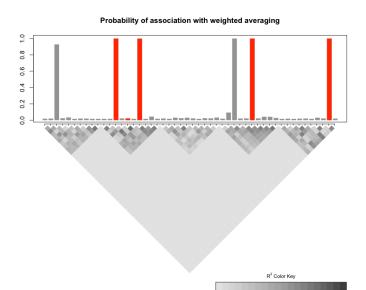




Implementation

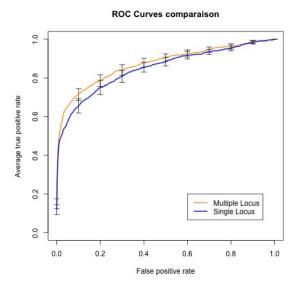
- ► Generate SNPs, traits, and dependences.
- ► Call function with different initial matrices, drawn at random.
- Generate ELBO and use it to calculate the weighted average ("BMA").

Probabilities of association with weighted averaging

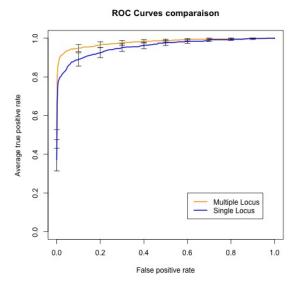




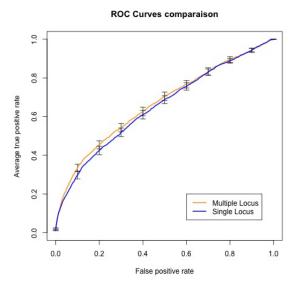
ROC curves comparison, $p_0 = 15$, max var. = 0.5



ROC curves comparison, $p_0 = 15$, max var. = 0.8



ROC curves comparison, $p_0 = 50$, max var. = 0.5





ROC curves comparison, $p_0 = 50$, max var. = 0.8

