

Co-data course: GRridge

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Our group: www.bigstatistics.nl

Setting

- **Prediction or Diagnosis**
- **Primary data**
 - ▶ Variables $i = 1, \dots, p$; Individuals $j = 1, \dots, n$; $p > n$
 - ▶ Focus on binary response Y_j (e.g. case vs control)
 - ▶ Measurements $\mathbf{X}_j = (X_{1j}, \dots, X_{pj})$
 - ▶ Goal: find f such that $Y_j \approx f(\mathbf{X}_j)$
 - ▶ Here, f : *logistic regression*
 - ▶ Some form of regularization required
- **Focus**
 - ▶ Differential regularization based on prior information:
co-data

Co-data

Definition Co-data: any information on the *variables* that does not use the response labels of the primary data

Co-data

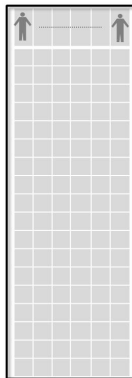
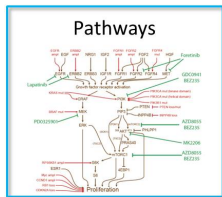
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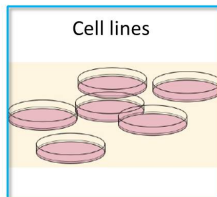
Databases



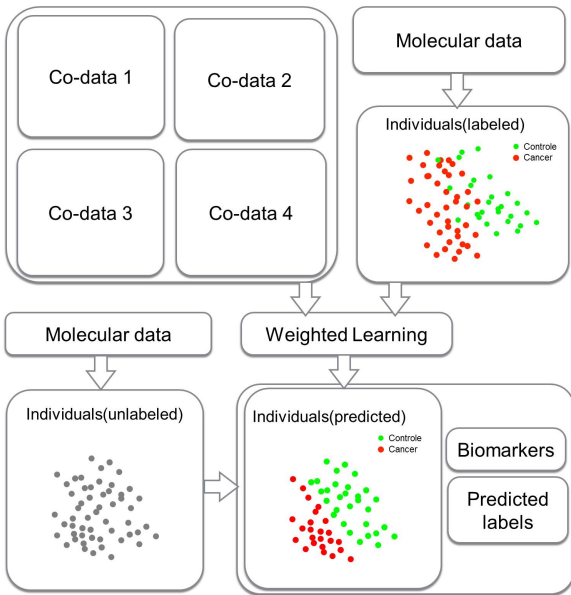
Related bio-molecules



Primary Data



Cell lines



Use of co-data

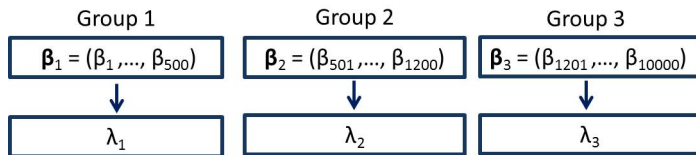
Groups: Co-data determine G prior groups of variables

Idea: Use different penalty weights $\boldsymbol{\lambda} = \lambda_1, \dots, \lambda_G$ across G co-data-based groups.

Use of co-data

Groups: Co-data determine G prior groups of variables

Idea: Use different penalty weights $\lambda = \lambda_1, \dots, \lambda_G$ across G co-data-based groups. $G = 3$:



E.g. Ridge: $\operatorname{argmax}_{\beta} \{ \mathcal{L}(\mathbf{Y}; \beta) - \sum_{g=1}^G \lambda_g \|\beta_g\|_2 \}$

→ **CV** not attractive

→ GRridge estimates λ by Empirical Bayes (EB)

Fitting

Likelihood contains term $\mathbf{X}'\beta$. Write $\lambda_g = \lambda'_g \lambda$

$$\operatorname{argmax}_{\beta} \{ \mathcal{L}(\mathbf{Y}; \mathbf{X}'\beta) - \lambda \sum_{g=1}^G \lambda'_g \|\beta_g\|_2 \}$$

\equiv

$$\operatorname{argmax}_{\tilde{\beta}} \{ \mathcal{L}(\mathbf{Y}; \tilde{\mathbf{X}}'\tilde{\beta}) - \lambda \|\tilde{\beta}\|_2 \},$$

where

$$\tilde{\mathbf{X}} = \mathbf{X}(\operatorname{diag}(\lambda'))^{-1/2} \text{ and } \tilde{\beta}_g = \beta_g(\lambda'_g)^{1/2}$$

→ Existing software used for fitting (glmnet, penalized)

Iteration

One may iterate the hyper-parameter estimation:

$$\operatorname{argmax}_{\tilde{\beta}} \{ \mathcal{L}(\mathbf{Y}; \tilde{\mathbf{X}}' \tilde{\beta}) - \lambda \|\tilde{\beta}\|_2 \}, \quad \tilde{\mathbf{X}} = \mathbf{X}(\operatorname{diag}(\lambda'))^{-1/2}$$

→ Same group-structure on $\tilde{\beta}$:

$$\operatorname{argmax}_{\beta} \{ \mathcal{L}(\mathbf{Y}; \tilde{\mathbf{X}}' \tilde{\beta}) - \lambda \sum_{g=1}^G \lambda_g'' \|\tilde{\beta}_g\|_2 \}$$

Effective penalty multiplier: $\lambda_g' \lambda_g''$

Convergence is monitored by cross-validated likelihood

Multiple partitions I

Current solution: Iterative. Suppose first partition \mathcal{G} has been used.

Different group-structure \mathcal{H} on $\tilde{\beta}$:

$$\operatorname{argmax}_{\beta} \{ \mathcal{L}(\mathbf{Y}; \tilde{\mathbf{X}}' \tilde{\beta}) - \lambda \sum_{h=1}^H \lambda''_h \| \tilde{\beta}_h \|_2 \}$$

Effective penalty multiplier for variable k : $\lambda'_{g(k)} \lambda''_{h(k)}$

Disadvantage: order may matter, although partitions are interwoven with iterations

Multiple partitions II

A nicer solution is implemented in `gren`:

$$\begin{aligned}\hat{\beta} := \operatorname{argmax}_{\beta} & \ell(\mathbf{Y}; \beta) - \frac{\lambda_1}{2} \sum_{g=1}^G \sum_{h=1}^H \sqrt{\lambda'_g \lambda''_h} \sum_{\substack{j \in \mathcal{G}_g \\ \cap \mathcal{H}_h}} |\beta_j| \\ & - \frac{\lambda_2}{2} \sum_{g=1}^G \sum_{h=1}^H \lambda'_g \lambda''_h \sum_{\substack{j \in \mathcal{G}_g \\ \cap \mathcal{H}_h}} \beta_j^2\end{aligned}$$

Overlapping groups^{*}

Prior modeled as:

$$\beta_k \sim N(0, \sum_{g_k \in \mathcal{J}_k} \sigma_{g_k}^2 / |\mathcal{J}_k|)$$

→ Set of G estimation equations → $\hat{\sigma}_\ell^2$

Set $\lambda_\ell \propto \hat{\sigma}_\ell^{-2}$

^{*}See Novianti et al. (2017), *Bioinformatics*

Data

Data set 1

- Methylation data on cervical tissue, $p = 40,000$, $n = 37$
- Classification: Normal vs CIN3 (Precursor)
- Co-data: Location of the probes, 6 groups
- Reference: Farkas et al. (2013), *Epigenetics*
- Also analysed in: Van de Wiel et al. (2016), *Stat Med*

Data set 2

- RNAseq data from *blood* platelets, $p = 18,410$, $n = 81$
- Classification: Breast vs. Colon cancer
- Co-data: Several (see course text)
- Reference: Best et al. (2015), *Cancer Cell*
- Also analysed in: Novianti et al. (2017), *Bioinformatics*

Getting started

Instructions and Exercises:

See https://magnusmunch.github.io/co-data_learning/