Machine Learning Network-Constrained Regression of Epigenetic Data

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Outline

Epigenetic background

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Regression method similarities

Breast cancer dataset

Epigenetic background

Penalized regression methods

Lasso
$$\lambda \sum_{i=1}^{p} |\beta_i|$$

Elastic Net $\lambda_1 \sum_{i=1}^{p} |\beta_i| + \lambda_2 \sqrt{\sum_{i=1}^{p} \beta_i^2}$

Grace $\lambda_1 \sum_{i=1}^{p} |\beta_i| + \lambda_2 \sum_{u \sim v} \left(\frac{\beta_u}{\sqrt{d_u}} - \frac{\beta_v}{\sqrt{d_v}} \right)^2 w(u, v)$

aGrace $\lambda_1 \sum_{i=1}^{p} |\beta_i| + \lambda_2 \sum_{u \sim v} \left(\frac{sign(\tilde{\beta}_u)\beta_u}{\sqrt{d_u}} - \frac{sign(\tilde{\beta}_v)\beta_v}{\sqrt{d_v}} \right)^2 w(u, v)$

GBLasso $\lambda \sum_{u \sim v} \left[\left(\frac{|\beta_u|}{\sqrt{d_u}} \right)^{\gamma} + \left(\frac{|\beta_v|}{\sqrt{d_v}} \right)^{\gamma} \right]^{1/\gamma}$

Linf $\lambda \sum_{u \sim v} \max \left(\frac{|\beta_u|}{\sqrt{d_u}}, \frac{|\beta_v|}{\sqrt{d_v}} \right)$

aLinf $\lambda \sum_{u \sim v} \left| \frac{sign(\tilde{\beta}_u)\beta_u}{\sqrt{d_u}} - \frac{sign(\tilde{\beta}_v)\beta_v}{\sqrt{d_v}} \right|$

TTLP $\lambda_1 \sum_{i=1}^{p} J_\tau |\beta_i| + \lambda_2 \sum_{u \sim v} \left| J_\tau \left(\frac{|\beta_u|}{w_u} \right) - J_\tau \left(\frac{|\beta_v|}{w_v} \right) \right|$

LTLP $\lambda_1 \sum_{i=1}^{p} |\beta_i| + \lambda_2 \sum_{u \sim v} \left| J_\tau \left(\frac{|\beta_u|}{w_u} \right) - J_\tau \left(\frac{|\beta_v|}{w_v} \right) \right|$

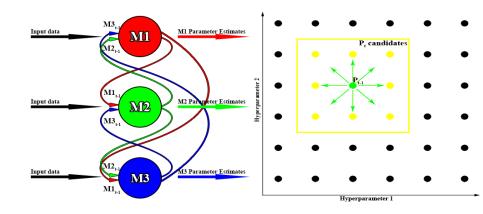
Composite voting regression

	X_1	X_2	 X_p
Method 1	$M_1(\beta_1)$	$M_1(\beta_2)$	 $M_1(\beta_p)$
Method 1 Method 2	$M_2(\beta_1)$	$M_2(\beta_2)$	 $M_2(\beta_p)$
Method k	$M_k(\beta_1)$	$M_k(\beta_2)$	 $M_k(\beta_p)$

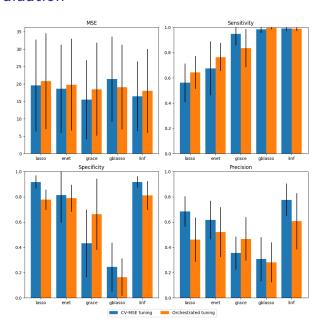
$$X_j = \begin{cases} important, & \text{if } \frac{\sum_{i=1}^k [M_i(eta_j) \neq 0]}{k} \geq \text{fraction of votes threshold} \\ unrelated, & \text{otherwise} \end{cases}$$

Final model obtained from OLSE on the set of important predictors

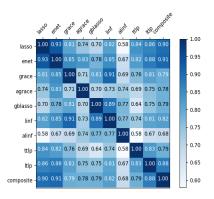
Orchestrated hyperparameter tuning

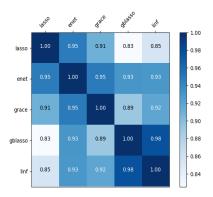


Model evaluation



Regression method similarities





Breast cancer dataset