Variable Selection In Additive Gene Environment Interactions with the Group Lasso

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1 Introduction

We consider a regression model for an outcome variable $\mathbf{Y} = (Y_1, \dots, Y_n)$ where n is the number of subjects. Let $E = (E_1, \dots, E_n)$ be a binary or continuous environment vector and $\mathbf{X} = (X_1, \dots, X_n)^T$ be the $n \times p$ matrix of high-dimensional data where $X_i = (X_{i1}, \dots, X_{ij}, \dots, X_{ip}) \in [0, 1]^p$. Consider the regression model with main effects and their interactions with E:

$$Y_{i} = \beta_{0}^{*} + \sum_{j=1}^{p} \beta_{j}^{*} X_{ij} + \beta_{E}^{*} E_{i} + \sum_{j=1}^{p} \alpha_{j}^{*} E_{i} X_{j} + \varepsilon_{i}, \qquad i = 1, \dots, n,$$

$$(1)$$

were $\beta_0^*, \beta_j^*, \beta_E^*, \alpha_j^*$ are the true unknown model parameters for $j = 1, \dots, p$. This can be extended to the more general additive model:

$$Y_i = \beta_0^* + \sum_{j=1}^p f_j^*(X_{ij}) + f_E^*(E_i) + \sum_{j=1}^p f_{jE}^*(X_{ij}, E_i) + \varepsilon_i \qquad i = 1, \dots, n$$
 (2)

As in (Radchenko and James, 2010), we can express (2) as

$$\mathbf{Y} = \sum_{j=1}^{p} \mathbf{f}_{j}^{*} + \mathbf{f}_{E}^{*}(E_{i}) + \sum_{j=1}^{p} \mathbf{f}_{jE}^{*} + \varepsilon_{i}$$

$$(3)$$

where
$$\mathbf{f}_{j}^{*} = (f_{j}^{*}(X_{1j}), \dots, f_{j}^{*}(X_{nj}))^{T}, \mathbf{f}_{jE}^{*} = (f_{jE}^{*}(X_{1j}, X_{1E}), \dots, f_{j}^{*}(X_{nj}, X_{nE}))^{T}$$

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