# Generalized Group Lasso for Patient Subgroup Identification

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## Introduction

Prognostic biomarkers and predictive biomarkers.

Why decision trees is not workable? Because the sample size in real clinical datasets is too small, typically no more than 100 patients.

Group lasso <sup>1</sup>

Elastic net <sup>2</sup> adaptive weights for elastic net <sup>3</sup>

Hierarchical Group lasso for interactions <sup>4</sup>

Overlapping group lasso <sup>567</sup>

Sparse Group Lasso <sup>8</sup> <sup>9</sup>

Structured group lasso <sup>10</sup>

Group lasso for logistic regression <sup>11</sup>

Other variable selection methods:

GUIDE: a regression tree <sup>1213</sup>

SIS: screening <sup>1415</sup>

SIR: 1617

Stepwise selection: 18

## **Methods**

#### Model

#### **Ordinary Linear Model**

$$Y = X\beta + W\tau + G\alpha + W \otimes G\gamma + \epsilon$$

Where X is the baseline variables, W is the treatment variable, G is the high dimensional design matrix of genes, i.e. gene expression levels, SNP and mutations, and  $W \otimes G$  is the interaction between genes and treatment.  $\theta = (\beta, \tau, \alpha, \gamma)$  is the corresponding coefficients. e is random error.

#### **Loss Function**

We used group lasso and elastic net for variables selection when  $n \ll p$ , and assumed the hierarchical relationship between prognostic biomarkers and predictive biomarkers, that is the predictive biomarkers should be a prognostic biomarkers. The loss function is

$$\begin{aligned} \min_{\theta} f(\theta|Y, X, W, G) + g(\theta) \\ g(\theta) &= \lambda_1 \sum_{i} \phi_i |\gamma_i| + \lambda_1 \sum_{i} \psi_i \sqrt{\alpha_i^2 + \gamma_i^2} + \lambda_2 (\parallel \alpha \parallel_2^2 + \parallel \gamma \parallel_2^2) \end{aligned}$$

Where  $\theta=(\alpha,\tau,\alpha,\gamma)$  is the parameter, and  $f(\theta|Y,X,W,G)$  is L-2 loss function. When the model is the ordinary linear model, the L-2 loss function is  $\|Y-(X\beta+W\tau+G\alpha+W\otimes G\gamma)\|^2$ . Penalty function  $g(\theta)$  can construct a complex hierarchical selection of  $\alpha$  and  $\gamma$ , that nonzero  $\gamma$  is a sufficient but not necessary condition for nonzero  $\alpha$ . The contour plot for a pair of  $\alpha$  and  $\gamma$  is shown in Figure 1.  $\lambda_1$  and  $\lambda_2$  are regularization parameters.

# Contour plot for alpha and gamma

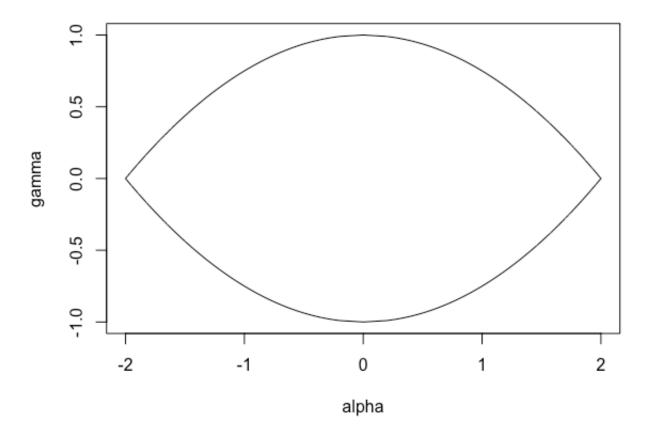


Figure 1

### **Criterion and Adaptive Weights**

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# **Algorithms**

Fast iterative shrinkage-thresholding algorithm with backtracking<sup>20</sup>

Proximal operator for group lasso 21

Adaptive restart for rippling behavior 22

Adaptive stepwise of cyclic Barzilai-Borwein spectral approach 23

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