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DEPARTAMENTO DE GENÉTICA
LGN5825 Genética e Melhoramento de Espécies Alógamas



Bayesian methods of GS

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Piracicaba, November 13th, 2019

Estimation

$$\text{🍇} + \text{🍇} + \text{🍇} = 30$$

$$\text{🍇} + \text{🍌} + \text{🍌} = 18$$

$$\text{🍌} - \text{🍒} = 2$$

$$3A + 0B + 0C = 30$$

$$1A + 2B + 0C = 18$$

$$0A + 1B - 1C = 2$$

$$X\beta = Y$$

$$\begin{pmatrix} 3 & 0 & 0 \\ 1 & 2 & 0 \\ 0 & 1 & -1 \end{pmatrix} \begin{pmatrix} A \\ B \\ C \end{pmatrix} = \begin{pmatrix} 30 \\ 18 \\ 2 \end{pmatrix}$$

$$\begin{pmatrix} A \\ B \\ C \end{pmatrix} = \begin{pmatrix} 3 & 0 & 0 \\ 1 & 2 & 0 \\ 0 & 1 & -1 \end{pmatrix}^{-1} \begin{pmatrix} 30 \\ 18 \\ 2 \end{pmatrix}$$

$$\beta = X^{-1}Y$$

$$\begin{pmatrix} A \\ B \\ C \end{pmatrix} = \begin{pmatrix} 1/3 & 0 & 0 \\ -1/6 & 1/2 & 0 \\ -1/6 & 1/2 & -1 \end{pmatrix} \begin{pmatrix} 30 \\ 18 \\ 2 \end{pmatrix} = \begin{pmatrix} 10 \\ 4 \\ 2 \end{pmatrix}$$

Estimation and restrictions

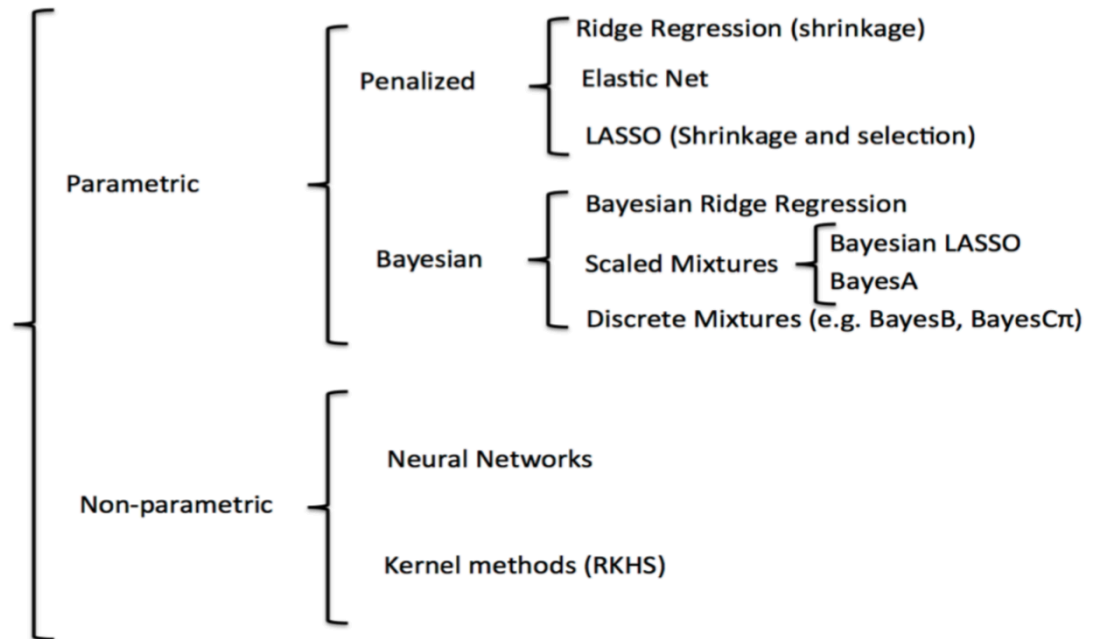
$$\begin{array}{l} \text{🍇} + \text{🍌} + \text{🍌} = 18 \\ \text{🍌} - \text{🍒} = 2 \end{array} \left. \vphantom{\begin{array}{l} \text{🍇} + \text{🍌} + \text{🍌} = 18 \\ \text{🍌} - \text{🍒} = 2 \end{array}} \right\} \text{Infinite solutions}$$

Markers > Observations

How do we solve it?



Restrictions



Bayes theory

$$P(\theta|y) = \frac{P(y|\theta)P(\theta)}{P(y)} \begin{cases} P(y|\theta): \text{pdf conditioned to } \theta \text{ given } y \\ P(\theta): \text{pdf of } \theta \text{ and } P(y): \text{pdf of } y \end{cases}$$

y : vector of observations
 θ : vector of parameters

$$P(\theta|y) \propto P(y|\theta)P(\theta)$$

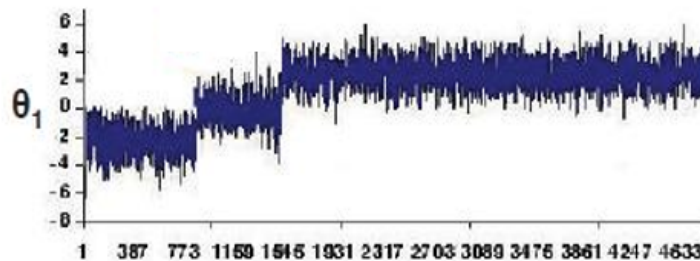
Posteriori distribution of θ

A priori distribution of θ

Genetic architecture (t, normal, exp, ...)

Jointed distribution of data given the parameters

Likelihood function (gaussian or ordinal)



Main difference among Bayesian methods

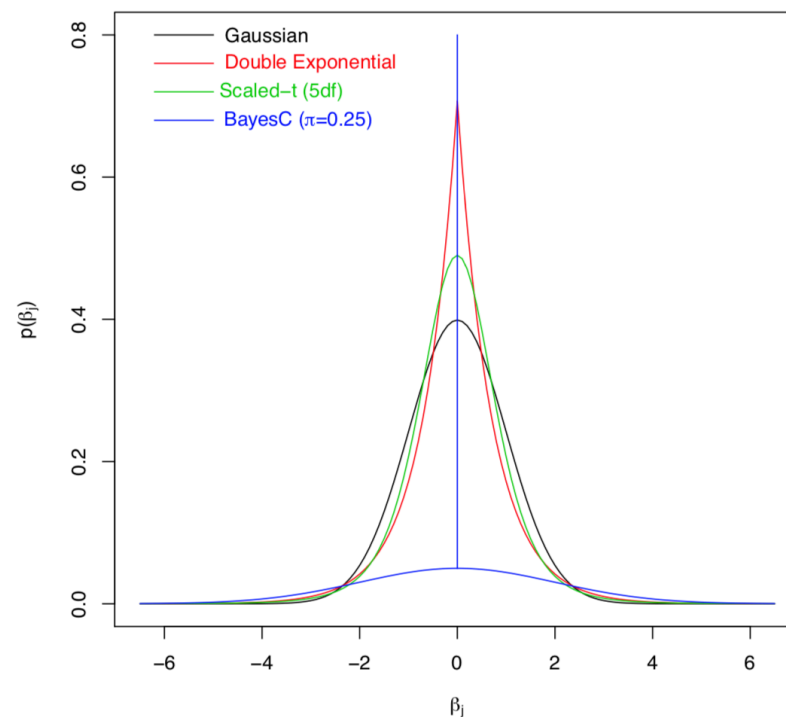
SNPs effect (a_i)

$$y = 1u + \sum_{i=1}^m M_i a_i + e$$

$i = 1, 2, \dots, m$ (SNPs)

Bayes alphabet

Model	Genetic variance (Vu)	Genetic control
Ridge Regression	Just one for all SNPs	Polygenic, small effects and explain equal amounts of variance
Bayes A	Specific for each SNP	Polygenic, small effects and explain different amounts of variance
Bayes B	Specific for each SNP	Polygenic, but some SNPs does not affect the trait (π), where π is fixed
Bayes C π	Just one for all SNPs	Polygenic, but some SNPs does not affect the trait (π), where π is random
Bayesian LASSO	Specific for each SNP	Oligogenic. Many SNPs does not affect the trait. Not necessary to define π



Bayes alphabet

Model	Conditioned fdp	Prior fdp	Vu
RR	$a_i \sigma^2 \sim N(0, \sigma^2)$	$\sigma^2 \sim \chi^{-2}(S, v)$	$\sigma_u^2 = 2\sigma^2 \sum_{i=1}^m p_i(1 - p_i)$
Bayes A	$\alpha_i \sigma_i^2 \sim N(0, \sigma_i^2)$	$\sigma_i^2 \sim \chi^{-2}(S, v)$	$\sigma_u^2 = 2 \sum_{i=1}^m p_i(1 - p_i) \sigma_i^2$
Bayes B	$a_i \boldsymbol{\pi}, \sigma_{i1}^2 \sim (1 - \gamma_i)N(0, \sigma_{i0}^2 = 0) + \gamma_i N(0, \sigma_{i1}^2)$ $P(y_i=0) = \boldsymbol{\pi}; P(y_i=1) = 1 - \boldsymbol{\pi}$	$\sigma_{i1}^2 \sim \chi^{-2}(S, v)$	$\sigma_u^2 = 2 \sum_{i=1}^m p_i(1 - p_i) \sigma_i^2$
Bayes C $\boldsymbol{\pi}$	$a_i \boldsymbol{\pi}, \sigma^2 \sim \boldsymbol{\pi}N(0, \sigma^2 = 0) + (1 - \boldsymbol{\pi})N(0, \sigma^2)$ $\boldsymbol{\pi} \sim U(0, 1)$	$\sigma^2 \sim \chi^{-2}(S^*, v)$ $S^* = \tilde{\sigma}^2 (v - 2) / v$ expected priori for SNP variance (fixed)	$\sigma_u^2 = 2(1 - \pi) \sigma^2 \sum_{i=1}^m p_i(1 - p_i)$
Lasso	$a_i \tau_i, \sigma_e^2 \sim N(0, \tau_i^2 \sigma_e^2)$ $a_i \tau_i, \sigma_e^2 \sim N(0, \sigma_i^2)$ $a \lambda, \sigma_e^2 \sim \prod_i \frac{\lambda}{2\sigma_e} \exp\left(\frac{-\lambda a_i }{\sigma_e}\right)$	$\tau_i^2 \lambda^2 \sim \text{Exp}(\lambda^2)$ $\lambda^2 \sim \text{Gamma}(\varphi_1, \varphi_2)$ $a_i \lambda, \sigma_e^2 \sim \text{Double-Exp}(0, \lambda)$	$\sigma_u^2 = \sum_{i=1}^m \underbrace{\tau_i^2 \sigma_e^2}_{\sigma_i^2} 2p_i(1 - p_i)$