# A Bayesian Approach for Envelope Models

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# Envelope Approach for Multivariate Regression

Standard multivariate linear regression model:  $Y_i = \mu + \beta X_i + \varepsilon_i$ ,  $i = 1, 2, \dots, n$ 

#### $\mathbf{Y} \in \mathbb{R}^{r}, \mathbf{X} \in \mathbb{R}^{p}, \boldsymbol{\mu} \in \mathbb{R}^{r}, \boldsymbol{\beta} \in \mathbb{R}^{r \times p}, \boldsymbol{\varepsilon} \in \mathbb{R}^{r}$

Envelopes arise by re-parametrization of the SLM in terms of the smallest subspace  $\mathcal{E} \subseteq \mathbb{R}^r$  such that ( $P_{\mathcal{E}}$  is projection onto the space  $\mathcal{E}$  and  $Q_{\mathcal{E}} = I - P_{\mathcal{E}}$ )

 $\mathbf{Q}_{\mathcal{E}}\mathbf{Y} \mid \mathbf{X} \sim \mathbf{Q}_{\mathcal{E}}\mathbf{Y}$  and  $\mathbf{P}_{\mathcal{E}}\mathbf{Y} \perp \mathbf{Q}_{\mathcal{E}}\mathbf{Y} \mid \mathbf{X}$ 

Impact of X on Y is concentrated in  $P_{\mathcal{E}}Y$ . Informally we refer

 $Q_{\mathcal{E}}Y$ : immaterial part of Y and  $P_{\mathcal{E}}Y$ : material part of Y

# **Posterior Distribution**

 $\pi((\boldsymbol{\mu},\boldsymbol{\eta},(\boldsymbol{\Gamma},\boldsymbol{\Gamma}_{0}),\boldsymbol{\omega},\boldsymbol{\omega}_{0}) \mid \boldsymbol{\mathbb{Y}}) \propto (2\pi)^{-(nr)/2} |\boldsymbol{\Omega}|^{-n/2} |\boldsymbol{\Omega}_{0}|^{-n/2} e^{-\frac{1}{2}\operatorname{tr}\left\{(\boldsymbol{\mathbb{Y}}-\mathbf{1}_{n}\boldsymbol{\mu}^{T}-\boldsymbol{\mathbb{X}}\boldsymbol{\eta}^{T}\boldsymbol{\Gamma}^{T})(\boldsymbol{\Gamma}\boldsymbol{\Omega}\boldsymbol{\Gamma}^{T}+\boldsymbol{\Gamma}_{0}\boldsymbol{\Omega}_{0}\boldsymbol{\Gamma}_{0}^{T})^{-1}(\boldsymbol{\mathbb{Y}}-\mathbf{1}_{n}\boldsymbol{\mu}^{T}-\boldsymbol{\mathbb{X}}\boldsymbol{\eta}^{T}\boldsymbol{\Gamma}^{T})^{T}\right\}}$  $\times |\Omega|^{-p/2} e^{-\frac{1}{2} \operatorname{tr} \left( \Omega^{-1} (\eta - \Gamma^{T} \mathbf{e})^{T} \right)} e^{-\frac{1}{2} \operatorname{tr} \left( \mathbf{D}^{-1} \mathbf{O}^{T} \mathbf{G} \mathbf{O} \right)} \prod^{u} \omega_{i}^{-\alpha - 1} e^{-\frac{\lambda}{\omega_{i}}} \prod^{r-u} \omega_{0,i}^{-\alpha_{0} - 1} e^{-\frac{\lambda_{0}}{\omega_{0,i}}}$ 

#### Theorem

- The posterior density in (3) is proper under either of the following conditions.
- $n > \max(r, p + 3)$
- $n + 2\alpha > 1$ ,  $\lambda$ ,  $\lambda_0 > 0$  and **C** is positive definite

The conditions  $\mathbf{Q}_{\mathcal{E}}\mathbf{Y} \mid \mathbf{X} \sim \mathbf{Q}_{\mathcal{E}}\mathbf{Y}$  and  $\mathbf{P}_{\mathcal{E}}\mathbf{Y} \perp \mathbf{Q}_{\mathcal{E}}\mathbf{Y} \mid \mathbf{X}$  hold if and only if (Cook 2010) •  $\mathcal{E}$  envelopes  $\mathcal{B} = span\{\boldsymbol{\beta}\}$ , i.e.  $\mathcal{B} \subseteq \mathcal{E}$ •  $\mathcal{E}$  is reducing subspace of  $\Sigma$ , i.e.  $\Sigma = P_{\mathcal{E}}\Sigma P_{\mathcal{E}} + Q_{\mathcal{E}}\Sigma Q_{\mathcal{E}}$ 

The intersection of all subspace  $\mathcal{E}$  with the above properties is called  $\Sigma$ -Envelope of  $\mathcal{B}$  and denoted by  $\mathcal{E}_{\Sigma}(\mathcal{B})$  with  $u = dim(\mathcal{E}_{\Sigma}(\mathcal{B}))$ 

#### **Coordinate representation of Envelope model**

 $\mathbf{Y} = \boldsymbol{\mu} + \boldsymbol{\Gamma} \boldsymbol{\eta} \mathbf{X} + \boldsymbol{\varepsilon}$ , where  $\boldsymbol{\beta} = \boldsymbol{\Gamma} \boldsymbol{\eta}, \boldsymbol{\Sigma} = \boldsymbol{\Gamma} \boldsymbol{\Omega} \boldsymbol{\Gamma}^{T} + \boldsymbol{\Gamma}_{0} \boldsymbol{\Omega}_{0} \boldsymbol{\Gamma}_{0}^{T} \boldsymbol{\Omega}, \boldsymbol{\Omega}_{0} > 0$ 

 $\Gamma$  and  $\Gamma_0$  be a basis for the space  $\mathcal{E}_{\Sigma}(\mathcal{B})$  and  $\mathcal{E}_{\Sigma}^{\perp}(\mathcal{B})$ . Note that the choice of  $\Gamma$  and  $\Gamma_0$  are not unique.

# How Envelope Model Works? Toy Example

Consider the multivariate linear regression model with response  $Y_1$ ,  $Y_2$  and one predictor variable X with two label 0 and 1.

 $\mathbf{Y} = \begin{pmatrix} Y_1 \\ Y_2 \end{pmatrix} = \begin{pmatrix} \mu_1 \\ \mu_2 \end{pmatrix} + \begin{pmatrix} \beta_1 \\ \beta_2 \end{pmatrix} X + \begin{pmatrix} \epsilon_1 \\ \epsilon_2 \end{pmatrix}$ (1)

> $\mu_1 = E(Y_1 \mid X = 0), \ \beta_1 = E(Y_1 \mid X = 1) - E(Y_1 \mid X = 0)$  $\mu_2 = E(Y_2 | X = 0), \quad \beta_2 = E(Y_2 | X = 1) - E(Y_2 | X = 0)$

Schematic representation: standard model	Sc
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### Sampling Scheme

#### Generalized Bingham distribution

A random matrix  $\mathbf{Z} = [\mathbf{Z}_1 : \mathbf{Z}_2]$  is defined to have a generalized matrix Bingham distribution on  $S_{2,2}$  with parameters  $A_1$  and  $A_2$  ( $GB_{2,2}(A_1, A_2)$ ) if the probability density function of Z (w.r.t the Haar measure on  $S_{2,2}$ ) is proportional to  $e^{-Z_1^T A_1 Z_1 - Z_2^T A_2 Z_2}$ . An efficient algorithm has been de-

•  $\mu \mid ((\Gamma, \Gamma_0), \eta, \omega, \omega_0, \mathbb{Y})$  is Normal and  $\eta \mid ((\Gamma, \Gamma_0), \omega, \omega_0, \mathbb{Y})$  is Multivariate normal •  $\omega_i \mid (\Gamma, \Gamma_0), \omega^{-i}, \omega_0, \mathbb{Y}$  and  $\omega_{0,i} \mid (\Gamma, \Gamma_0), \omega, \omega_0^{-i}, \mathbb{Y}$  are Truncated-Inverse-Gamma • Sampling of O involves samling from  $GB_{2,2}$  distribution

We start at a given initial value of the parameters, and repeat the following steps.

• Sampling from  $((\Gamma, \Gamma_0), \omega, \omega_0) \mid \mathbb{Y}$  $\rightarrow$  For  $i = 1, 2, \cdots, u$ , update  $\omega_i$ 

 $\rightarrow$  For  $i = 1, 2, \cdots, r - u$ , update  $\omega_{0,i}$ 

The corresponding MC is Harris ergodic.

veloped to sample from  $GB_{2,2}$ .

(3)

 $\rightarrow$  For every pair or randomly chosen pair (i, j) such that  $1 \le i < j \le r$ , update  $\mathbf{O}_{i}$  and  $\mathbf{O}_{j}$ 

• Sample from full conditional distribution for  $\eta \mid ((\Gamma, \Gamma_0), \omega, \omega_0), \mathbb{Y}$  and  $\mu \mid \mathbb{Y}$ 

## Model Selection: DIC Criteria

• Need to select  $u \in 0, 1, \ldots, r$ •  $\theta := (\mu, \eta, (\Gamma, \Gamma_0), \omega, \omega_0)$  be the parameter vector • For each *u* construct the appropriate Markov chain

# Bayesian Envelope Model

Features that a Bayesian approach would offer are

- Comprehensive uncertainty characterization through the posterior distribution
- A framework to incorporate prior information
- Ability to deal with the case when n < r

#### A reparameterization of Envelope model

$$\mathbf{Y} = \boldsymbol{\mu} + \boldsymbol{\Gamma} \boldsymbol{\eta} \mathbf{X} + \boldsymbol{\varepsilon}, \qquad \mathbf{\Sigma} = \boldsymbol{\Gamma} \boldsymbol{\Omega} \boldsymbol{\Gamma}^{T} + \boldsymbol{\Gamma}_{0} \boldsymbol{\Omega}_{0} \boldsymbol{\Gamma}_{0}^{T},$$

(2)

•  $\boldsymbol{\beta} = \boldsymbol{\Gamma}\boldsymbol{\eta}$ ,  $\boldsymbol{\Gamma} \in S^+_{\boldsymbol{\Gamma},\boldsymbol{\mu}}$ ,  $\boldsymbol{\Gamma}_0 \in S^+_{\boldsymbol{\Gamma},\boldsymbol{\Gamma}-\boldsymbol{\mu}}$  with  $\boldsymbol{\Gamma}_0^T \boldsymbol{\Gamma} = 0$ ,  $\boldsymbol{\eta} \in M_{\boldsymbol{\mu},\boldsymbol{p}}$ .

•  $\Omega$  and  $\Omega_0$  are diagonal matrices with diagonal entries arranged in decreasing order

• To estimate  $(\mu, \eta, (\Gamma, \Gamma_0), \omega, \omega_0)$ 

If u = r, envelope model (2) is equivalent to the standard model by the one to one transformation  $(\Gamma, \eta, \Omega) \rightarrow (\beta = \Gamma \eta, \Sigma = \Gamma \Omega \Gamma')$ 

 $\to \{\theta^{(i)}\}_{i=1}^M$  the samples from the relevant posterior distribution (after an appropriate burn-in)  $\rightarrow \text{Calculate } D/C = \bar{D} + \frac{1}{2(M-1)} \sum_{i=1}^{M} \left( D\left(\theta^{(i)}\right) - \bar{D} \right)^2 \text{ where}$  $D(\theta) := -2 \log L(\theta) \text{ and } \bar{D} := \sum_{i=1}^{M} D\left(\theta^{(i)}\right) / M.$ 

• Select the value of *u* which corresponds to the minimum DIC.

# Application: Analysis of Wheat Protein Data

A brief summary of the Wheat protein data, (Cook 2010) can be summarized as follows

- Consists of r = 6 responses, which measure the log infrared reflectance at six different wavelengths for 50 ground wheat samples
- The predictor is a binary indicator, taking 0 or 1 if a sample has high or low protein content
- There are 26 samples with high protein content, and 24 samples with low protein content

Model selection: DIC scores								
	u	Uniform	Emperical					
	0	1257.7	1254.5					
	1	1201.9	1197.5					
	2	1206.3	1374.3					
	3	1208	1245.8					
	4	1209.3	1266.2					
	5	1211.1	1333					

1668.1

1216

6

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	Coefficient	Bayesian envelope model		Bayesian standard model		
		Post. mean	Post. SD	Post. mean	Post. SD	
-	$\beta_1$	-1.039	0.378	2.934	10.479	
	$\beta_2$	4.406	0.498	7.745	8.630	
	$\beta_3$	3.630	0.417	7.219	9.273	
	$eta_4$	-5.880	0.644	-2.395	10.157	
	$eta_5$	0.594	0.224	2.799	14.601	
	$oldsymbol{eta}_6$	-1.610	0.904	0.410	5.759	

Extensive simulation has been conducted for: model selection accuracy, comparison with non-Bayesian envelope model and different versions of standard Bayesian models

# **Prior Specification**

#### • $\pi(\mu) \propto 1$

#### • $\eta \mid (\Gamma, \Gamma_0), \omega, \omega_0 \sim \mathcal{MN}_{u,p}(\Gamma^T \mathbf{e}, \Omega, \mathbf{C}^{-1})$ $\Omega := diag(\omega)$

#### • $\mathbf{O} \sim B_{r,r}(\mathbf{G}, \mathbf{D}^{-1})$ $\mathbf{O} := [\Gamma, \Gamma_0]$

The **prior mode for O** is an appropriately permuted version of the eigenvectors of **G** 

- ( $\boldsymbol{\omega}, \boldsymbol{\omega}_0$ ) and **O** are apriori independent  $\rightarrow$  The entries of  $\omega$ : order statistics of u i.i.d. observations from the Inverse-Gamma( $\alpha, \lambda$ ) dis-
- tribution
- $\rightarrow$  The entries of  $\omega_0$ : order statistics of r u i.i.d observations from the Inverse-Gamma( $\alpha_0, \lambda_0$ ) distribution

#### Uniform improper prior The joint improper prior corresponding to uniform improper prior is given by $\pi(\mu, \eta, (\Gamma, \Gamma_0), \omega, \omega_0) \propto 1$

This corresponds to following hyperparameter choices

• e = 0, C = 0, G = 0•  $\alpha = -(1 + \frac{p}{2}), \lambda = 0, \ \alpha_0 = -1, \lambda_0 = 0$ 

#### Emperical prior

 $(\boldsymbol{\eta}^*, (\boldsymbol{\Gamma}^*, \boldsymbol{\Gamma}^*_0), \boldsymbol{\omega}^*, \boldsymbol{\omega}^*_0)$  obtained by a naive method • Set  $\mathbf{e} = \Gamma^* \eta^*$  and  $\mathbf{C} = \mathbf{0}$ 

•  $\alpha$  ,  $\lambda$  and  $\alpha_0$  ,  $\lambda_0$ : estimated by moment estimator using values  $\omega^*$  and  $\omega_0^*$ • **D** diagonal with diagonal elements ( $\omega^*$ ,  $\omega_0^*$ ) • we employ a procedure to ensured that  $O^*$  is the prior mode of **O** 

#### Summary

- We developed a comprehensive Bayesian framework for estimation and model selection is in the context of envelope model
- A parameterization available for Bayesian analysis has been introduced
- Class of priors introduced has desirable proprieties:
- $\rightarrow$  flexible
- $\rightarrow$  sensible interpretation as well as specification of hyperparameters
- $\rightarrow$  easy to sample from the corresponding posterior
- conditions for posterior propriety have been investigated
- A new distribution  $GB_{2,2}$  along with efficient sampling scheme is developed
- R package "BENV" is developed for data analysis
- The method is applied successfully on simulated and real datasets

# References

- Khare K., Pal S. and Su Z. A Bayesian Approach for Envelope Models, Submitted to (and tentatively Accepted in) Annals of Statistics.
- Cook, R.D., Li, B. and Chiaromonte, F. (2010). Envelope Models for Parsimonious and Efficient Multivariate Linear Regression, Statistica Sinica, 20, 927–960.