Simultaneous Inference for Latent Variables in Factor Analytic Models

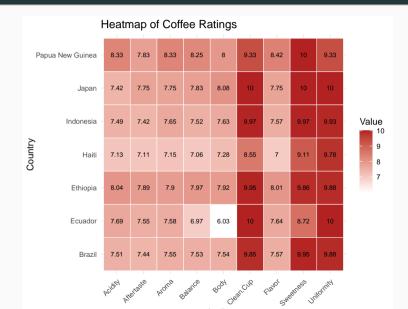
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Motivating Data: Coffee Bean Exploration

- The coffee bean dataset records consumer ratings (0–10) on nine sensory attributes (e.g., sweetness, aroma, acidity) and aggregates them by country. The dataset consists of n=32 countries in total.
 - Data are publicly available from Kaggle https://www.kaggle.com/datasets/ adampq/coffee-quality-with-locations-of-origin?resource=download, which itself is sourced from the Coffee Quality Institute (CQI) database https://database.coffeeinstitute.org/

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- We hypothesise a few latent factors can be used to represent these observed ratings e.g., latent axis representing aspects of coffee taste or richness. That is, we wish to perform dimension reduction.
- Our interest is to <u>perform statistical inference</u> in this latent space e.g., do two countries differ statistically in terms of coffee taste/richness? Is the "best" country statistically different from the rest?

The Factor Analytic Model

Suppose we have n independent observations $\mathbf{y}_i; i=1,2,\ldots,n$, each of dimension p. Assume the j-th response y_{ij} can be written as (West et al., 2003)

$$y_{ij} = \mathbf{x}_i^{\top} \boldsymbol{\beta}_j + \mathbf{f}_i^{\top} \boldsymbol{\lambda}_j + \epsilon_{ij},$$

where

- \mathbf{x}_i is a q-dimensional covariate and $\boldsymbol{\beta}_i$ the associated coefficients;
- \mathbf{f}_i is an m-dimensional latent factor assumed to be standard normal in distribution, $f_{ik} \sim \mathcal{N}(0,1)$;
- λ_j is an m-dimensional loading vector. Furthermore, we define the loading matrix $\mathbf{\Lambda} = [\lambda_1 \ \lambda_2 \ \cdots \ \lambda_p]^{\mathsf{T}};$
- ϵ_{ij} is an error term where we assume $\epsilon_{ij} \sim \mathcal{N}(0, \psi_{jj})$.

The Factor Analytic Model

· We write the fully vectorised form of the factor analytic model as,

$$y = (\mathbf{I}_p \otimes \mathbf{X}) \boldsymbol{\beta} + (\boldsymbol{\Lambda} \otimes \mathbf{I}_n) \boldsymbol{f} + \boldsymbol{\epsilon},$$

where $\boldsymbol{y} = (y_{11}, \cdots, y_{n1}, y_{12}, \cdots, y_{np})^{\top}$, $\mathbf{X}^{\top} = \begin{bmatrix} \mathbf{x}_1^{\top} \ \mathbf{x}_2^{\top} \ \dots \ \mathbf{x}_n^{\top} \end{bmatrix}$ is an $q \times n$ matrix, $\boldsymbol{\beta} = \begin{pmatrix} \boldsymbol{\beta}_1^{\top}, \boldsymbol{\beta}_2^{\top}, \boldsymbol{\beta}_p^{\top} \end{pmatrix}^{\top}$, $\boldsymbol{f} = (f_{11}, \cdots, f_{n1}, f_{12}, \cdots, f_{np})^{\top}$ and $\boldsymbol{\epsilon} = (\epsilon_{11}, \cdots, \epsilon_{n1}, \epsilon_{12}, \cdots, \epsilon_{np})^{\top}$.

Marginally, we have

$$y \sim \mathcal{N}_{np}\left(\left(\mathbf{I}_p \otimes \mathbf{X}\right) \boldsymbol{\beta}, \mathbf{V} \otimes \mathbf{I}_n\right),$$

where $\mathbf{V} = \mathbf{\Lambda} \mathbf{\Lambda}^{\top} + \mathbf{\Psi}$ is the marginal variance of each observation \mathbf{y}_i , and we call $\boldsymbol{\theta} := \left[\operatorname{vec}(\mathbf{\Lambda})^{\top}, \operatorname{diag}(\mathbf{\Psi})^{\top} \right]^{\top}$ the variance components.

Model Identifiability

- Without any constraints on the loading matrix Λ , the factor model is not identifiable, since for any orthogonal matrix M, the transformed loading matrix ΛM yields the same covariance structure and the same likelihood (Mardia et al., 1979).
- We put a **corner constraint** on the loading matrix by setting the upper-triangular elements of Λ to zero, and restrict the diagonal elements of Λ to be positive.

Parameter Estimation and Factor Prediction

Given variance component heta,

• the MLE of $oldsymbol{eta}$ is

$$\hat{\boldsymbol{\beta}} = \left(\mathbf{I}_p \otimes (\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top\right) \boldsymbol{y},$$

 \cdot the best linear unbiased prediction (BLUP) (Searle et al., 2009) of f is

$$ilde{m{f}} = \mathbb{E}\left[m{f} \mid m{y}
ight] = \left(m{\Lambda}^ op \mathbf{V}^{-1} \otimes \mathbf{I}_n
ight) \left(m{y} - (\mathbf{I}_p \otimes \mathbf{X})\hat{m{eta}}
ight).$$

In practice, we can plug in estimated values of $\hat{\theta}$ (e.g. obtained via REML (Corbeil and Searle, 1976)) to obtain \hat{f} .

General Mixed Parameters

We are interested in conduct simultaneous inference for a set of L general mixed parameters. For known designed vectors c_l and κ_l , the general mixed parameter (Reluga et al., 2023) for level $l=1,\ldots,L$ is given by

$$\mu_l = \boldsymbol{c}_l^{\top} \boldsymbol{\beta} + \boldsymbol{\kappa}_l^{\top} \boldsymbol{f}$$

The BLUP for the general mixed parameter is then given by plugging $\hat{m{\beta}}$ and $\tilde{m{f}}$,

$$\tilde{\mu}_l = \boldsymbol{c}_l^{\top} \hat{\boldsymbol{\beta}} + \kappa_l^{\top} \tilde{\boldsymbol{f}},$$

and the empirical BLUP or EBLUP follows us

$$\hat{\mu}_l = \boldsymbol{c}_l^{\top} \hat{\boldsymbol{\beta}} + \kappa_l^{\top} \hat{\boldsymbol{f}}.$$

Prediction Mean Square Error

• For level $l=1,\ldots,L$, the prediction mean square error (PMSE) $\sigma^2(\hat{\mu}_l)$ can be decomposed as

$$\mathbb{E}[(\hat{\mu}_l - \mu_l)^2] = \mathbb{E}[(\hat{\mu}_l - \tilde{\mu}_l)^2] + \mathbb{E}[(\tilde{\mu}_l - \mu_l)^2] + 2\mathbb{E}[(\hat{\mu}_l - \tilde{\mu}_l)(\tilde{\mu}_l - \mu_l)]$$

= $\mathbb{E}[(\hat{\mu}_l - \tilde{\mu}_l)^2] + \sigma^2(\tilde{\mu}_l).$

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$$= \mathbb{E}[(\hat{\mu}_l - \tilde{\mu}_l)^2] + \sigma^2(\tilde{\mu}_l).$$

· As $n \to \infty$ the second term dominates the PMSE of $\hat{\mu}_l$, where

$$\sigma^{2}(\tilde{\mu}_{l}) = \boldsymbol{\kappa}_{l}^{\top} \boldsymbol{\kappa}_{l} - \boldsymbol{\kappa}_{l}^{\top} ((\boldsymbol{\Lambda}^{\top} \mathbf{V}^{-1} \boldsymbol{\Lambda}) \otimes \mathbf{I}_{n}) \boldsymbol{\kappa}_{l} + (\boldsymbol{c}_{l} - (\mathbf{I}_{p} \otimes \mathbf{X}^{\top}) \boldsymbol{o}_{l})^{\top} \mathbf{Q} (\boldsymbol{c}_{l} - (\mathbf{I}_{p} \otimes \mathbf{X}^{\top}) \boldsymbol{o}_{l})$$

where $o_l = (\mathbf{V}^{-1} \mathbf{\Lambda} \otimes \mathbf{I}_n) \kappa_l$ and $\mathbf{Q} = \mathbf{V} \otimes (\mathbf{X}^{\top} \mathbf{X})^{-1}$.

• We approximate $\hat{\sigma}^2(\hat{\mu}_l) \approx \hat{\sigma}^2(\tilde{\mu}_l)$, based on by plugging the estimated variance components.

Basic Idea: CPI vs. SPI

• A cluster-level prediction interval (CPI) is an region $C_{1-\alpha}$ such that it has probability $100(1-\alpha)\%$ of covering the general mixed parameter for a single level.

$$\mathbb{P}(\mu_l \in \mathcal{C}_{1-\alpha}) = 1 - \alpha, \ \forall l \in [L]$$

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• A simultaneous prediction interval (SPI) is a region $\mathcal{I}_{1-\alpha}$ such that it has probability $100(1-\alpha)\%$ of covering the general mixed parameters across all levels,

$$\mathbb{P}(\mu_l \in \mathcal{I}_{1-\alpha}, \forall l \in [L]) = 1 - \alpha.$$

Constructing SPIs

• From the equation $\mathbb{P}(\mu_l \in \mathcal{I}_{1-\alpha}, \forall l \in [L]) = 1 - \alpha$ suggests a maximum t-type statistic:

$$\begin{split} &\alpha = \mathbb{P}\big(\mu_l \notin \mathcal{I}_{1-\alpha}, \exists l \in [L]\big) \\ &= \mathbb{P}\left(\left|\frac{\hat{\mu}_l - \mu_l}{\hat{\sigma}(\hat{\mu}_l)}\right| \geqslant c_{1-\alpha}, \exists l \in [L]\right)\right) \\ &= \mathbb{P}\left(\max_{l=1,2,...,L} \left|\frac{\hat{\mu}_l - \mu_l}{\hat{\sigma}(\hat{\mu}_l)}\right| \geqslant c_{1-\alpha}\right). \end{split}$$

• Theoretically, the critical value $c_{1-\alpha}$ is then the $(1-\alpha)^{\rm th}$ -quantile of the maximum t-type statistic i.e.,

$$c_{1-\alpha} = \inf_{t} \{ t \in \mathbb{R} : \mathbb{P}(\tau \leqslant t) \geqslant 1 - \alpha \},$$

where $\tau = \max_{l=1,2,...,L} |\tau_l|$ and $\tau_l = \frac{\hat{\mu}_l - \mu_l}{\hat{\sigma}(\hat{\mu}_l)}$.

· If c_{1-lpha} is known, the SPI can be constructed as

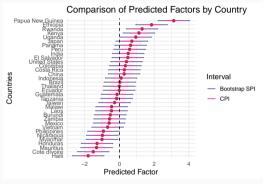
$$\mathcal{I}_{1-\alpha} = \underset{l=1}{\overset{L}{\times}} \left[\hat{\mu_l} \mp c_{1-\alpha} \hat{\sigma}(\hat{\mu}_l) \right].$$

- 1. Fit the model and obtain the EBLUP $\hat{\mu}_l$ and the estimated PMSE $\hat{\sigma}(\hat{\mu}_l)$ for levels $l=1,2,\ldots,L$.
- 2. Create B bootstrap datasets using the estimated parameters.

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- 2. **Create** *B* **bootstrap datasets** using the estimated parameters.
- For each bootstrap dataset, fit the model again to obtain the bootstrap version of the maximum t-type statistics.
- 4. Use the $1-\alpha$ quantile of the bootstrap empirical distribution of the maximum t-type statistics $c_{1-\alpha}^{BS}$.

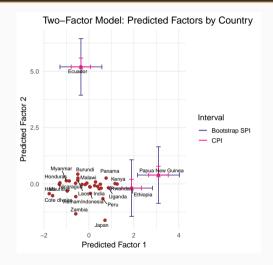
- 1. Fit the model and obtain the EBLUP $\hat{\mu}_l$ and the estimated PMSE $\hat{\sigma}(\hat{\mu}_l)$ for levels $l=1,2,\ldots,L$.
- 2. **Create** *B* **bootstrap datasets** using the estimated parameters.
- 3. For each bootstrap dataset, fit the model again to **obtain the bootstrap version of the maximum t-type statistics**.
- 4. Use the $1-\alpha$ quantile of the bootstrap empirical distribution of the maximum t-type statistics $c_{1-\alpha}^{BS}$.
- 5. Construct SPI using $c_{1-\alpha}^{BS}$ and the results from Step 1, $\mathcal{I}_{1-\alpha}^{BS} = \times_{l=1}^L \left[\hat{\mu}_l \mp c_{1-\alpha}^{BS} \hat{\sigma}(\hat{\mu}_l) \right]$.

Application to Coffee Bean Data: 1-factor Model





Application to Coffee Bean Data: 2-factor Model





Discussion

- Simultaneous inference can be useful in the context of factor analytic models, if we want to perform (say) formal statistical inference on the latent variables.
- We proposed constructing SPI using a bootstrap approach. In simulations (not shown), we show this generically performs better than simpler Monte Carlo- and Bonferroni-based SPIs.
- In future work, we will run more simulations under more complicated models and general mixed parameters, as well as investigate simultaneous inference the case of non-Gaussian response factor analytic models.

Thank You!

Let's drink a cup of coffee and see whether my research aligns with your tastes!

Questions?



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Appendix

The Monte Carlo Procedure

Using the mixed model equation, we find

$$\begin{bmatrix} \tilde{\boldsymbol{\beta}} - \boldsymbol{\beta} \\ \tilde{\boldsymbol{f}} - \boldsymbol{f} \end{bmatrix} \sim \mathcal{N}_{qp+nm} \left(\mathbf{0}_{qp+nm}, \left(\hat{\mathbf{A}}^{\top} \hat{\mathbf{K}}^{-1} \hat{\mathbf{A}} + \hat{\mathbf{B}} \right)^{-1} \right),$$

where
$$\hat{\mathbf{A}} := \begin{bmatrix} \mathbf{I}_p \otimes \mathbf{X} & \hat{\mathbf{\Lambda}} \otimes \mathbf{I}_n \end{bmatrix}$$
, $\hat{\mathbf{B}} := \begin{bmatrix} \mathbf{0}_{qp \times qp} & \mathbf{0}_{qp \times nm} \\ \mathbf{0}_{nm \times qp} & \mathbf{I}_{nm} \end{bmatrix}$, and $\hat{\mathbf{K}} := \hat{\mathbf{\Psi}} \otimes \mathbf{I}_n$.

Therefore for a Monte Carlo Procedure, for $s=1,2,\ldots,S$:

- 1. Sample $\begin{bmatrix} \tilde{\beta} \beta \\ \tilde{f} f \end{bmatrix}^{(s)}$ from its asymptotic distribution.
- 2. Calculate the centred general mixed effect $(\mu_l \mu)^{(s)} = c_l^{\top} (\tilde{\boldsymbol{\beta}} \boldsymbol{\beta})^{(s)} + \boldsymbol{\kappa}_l^{\top} (\tilde{\boldsymbol{f}} \boldsymbol{f})^{(s)}$.
- 3. Calculate $\tau_{MC}^{(s)} := \max_{l} \frac{|(\mu_l \mu)^{(s)}|}{\hat{\sigma}(\hat{\mu}_l)}$.

Order $\tau_{MC}^{(s)}$ for all $s \in [S]$ and find the upper $1-\alpha$ quantile as the empirical critical value $c_{1-\alpha}^{\rm MC}$.

Bonferroni Procedure

In a Bonferroni procedure, we directly approximate the critical value as $c_{1-\alpha}^{\mathrm{BO}}=\Phi^{-1}\left(1-\frac{\alpha}{2L}\right)$, where $\Phi^{-1}(\cdot)$ is the quantile function of the standard normal distribution.

Simulation Settings

• It is of interest to see how SPIs constructed by different methods behave under different n,p,q,m, along with the true variance explained (VE) by the loadings

$$VE = \frac{\operatorname{tr}(\boldsymbol{\Lambda}\boldsymbol{\Lambda}^\top)}{\operatorname{tr}(\boldsymbol{\Lambda}\boldsymbol{\Lambda}^\top + \boldsymbol{\Psi})}.$$

- We ran 200 simulation rounds for different combinations of (n, p, q, m, VE).
- The **target** of the intervals was all the latent variables in a one-factor model i.e., L=n.
- \cdot We used B=500 bootstrap datasets and S=2000 Monte Carlo samples.

Simulation Results

Table 1: Summary of SPI methods across different simulation settings (p = 20, m = 1, q = 3). The nominal level of intervals is 0.95.

Setting	Method (Intervals)	ECP	Mean Len.	Avg Var-Width ($\times 10^{-2}$)
n = 500, VE= 0.7	Bootstrap SPI	0.949	1.30	0.13
	Monte Carlo SPI	0.913	1.26	0.11
	Bonferroni SPI	0.924	1.27	0.10
	CPI	0	0.57	0.03
n = 200, VE = 0.7	Bootstrap SPI	0.965	1.43	0.27
	Monte Carlo SPI	0.915	1.35	0.22
	Bonferroni SPI	0.930	1.37	0.19
	CPI	0	0.57	0.09
n = 500, VE= 0.5	Bootstrap SPI	0.940	1.82	0.31
	Monte Carlo SPI	0.935	1.78	0.25
	Bonferroni SPI	0.935	1.79	0.25
	CPI	0	0.85	0.07

Takeaways from Simulations

- Bootstrap SPI performs well, achieving empirical coverage closest to the nominal 0.95 level.
- As sample size increases, Monte Carlo and Bonferroni SPIs are still slightly undercovered.
- Lower VE increases uncertainty, resulting in longer interval lengths and larger average variance of interval widths.
- **CPI consistently fails to provide valid coverage**, highlighting the necessity of simultaneous inference.