Understanding parallel analysis methods for rank selection in PCA



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1000G genetics data: n = 2318 individuals, p = 115019 SNPs





Rounak Dey



Xihong Lin

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*PC's can reveal population (and sub-population) structure, but how many are meaningful?* 

Parallel analysis for rank selection in PCA

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Question: how can we make principled selections and reason about them?

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Question: how can we make principled selections and reason about them?

The spectrum looks like a spiked covariance model...

Parallel analysis for rank selection in PCA

# Rank selection for PCA

Rank selection is important - it affects every downstream step!

- too many: add noise to downstream analyses
- too few: lose signals that were in the data

Many excellent and practical methods:

- Likelihood ratio test (Bartlett 1950)
- Fixed threshold (Kaiser 1960)
- Scree plot (Cattell 1966)

- $4/\sqrt{3}$  (Gavish & Donoho 2014)
- bi-cross-validation (Owen & Wang 2016)

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Today's talk: parallel analysis (Horn, 1965; Buja & Eyuboglu 1992)

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Today's talk: parallel analysis (Horn, 1965; Buja & Eyuboglu 1992)

PA is a popular method with extensive empirical evidence, but limited theoretical understanding – exciting area for work!

Parallel analysis is suggested in many reviews:

- Brown (2014): PA "is accurate in the vast majority of cases"
- Hayton et al. (2004): PA is "one of the most accurate factor retention methods" used in social science and management
- Costello and Osborne (2005): PA is "accurate and easy to use"
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Also gaining popularity in applied statistics (esp. biological sciences):

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But there remains limited theoretical understanding: PA is "at best a heuristic approach rather than a mathematically rigorous one" – Green et al. (2012)

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Idea: recover "null" by destroying correlations between features.

Parallel analysis for rank selection in PCA

# A quick sneak peak...

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Permutation provides a good estimate of the noise spectrum. ...let's begin characterizing this a bit!

Model: data is a linear combination of factors  $\lambda_{jk}$  with noise  $\varepsilon_{ij}$ 

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i.e., low-rank signal + noise

$$X = \underline{\eta} \Lambda^{\top} + \mathcal{E} = S + \mathcal{E}.$$





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Consequence: PA estimates noise spectrum (i.e., noise floor)

$$\sigma_k(X_{\pi}) = \sigma_k(S_{\pi} + \mathcal{E}_{\pi}) \approx \sigma_k(\mathcal{E}_{\pi}) =_d \sigma_k(\mathcal{E}_{\pi}).$$

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When does permutation successfully do this?

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Perceptible factor: singular value  $\sigma_k > b + \delta$  a.s. for some  $\delta > 0$ Imperceptible factors: singular value  $\sigma_k < b - \delta$  a.s. for some  $\delta > 0$
#### Important aside: small factors can fall below the noise

Example: Three factors, but only two rise above the phase transition.



Perceptible factor: singular value  $\sigma_k > b + \delta$  a.s. for some  $\delta > 0$ Imperceptible factors: singular value  $\sigma_k < b - \delta$  a.s. for some  $\delta > 0$ 

Question: when does parallel analysis identify perceptible factors?

#### Formalizing the intuition

**Theorem.** Suppose  $X = S + \mathcal{E}$  with signal  $S = \eta \Lambda^{\top}$  where

- ▶  $\eta = U\Psi^{1/2}$  for some  $\Psi$  where  $U \in \mathbb{R}^{n \times r}$  has ind. stand. entries;
- $\Lambda \Psi^{1/2} = (f_1, \dots, f_r)$  has bounded and delocalized columns, i.e.,  $\|f_k\|_2 \leq C n^{1/4-\delta/2}$  and  $\|f_k\|_4/\|f_k\|_2 \to 0$ ;

and with noise  $\mathcal{E} = Z \Phi^{1/2}$  where  $\Phi = \text{diag}(\phi)$  is diagonal,

- ▶  $Z \in \mathbb{R}^{n \times p}$  has ind. stand. entries with bounded fourth moment;
- entries of Z have bounded  $(6 + \Delta)$ th moments;

▶  $p^{-1}\sum_{j} \delta_{\phi_j} \Rightarrow H$  and  $\max_j \phi_j \to U(H)$  as  $n, p \to \infty$  with  $p/n \to \gamma > 0$ .

Then PA selects all perceptible and no imperceptible factors with prob  $\rightarrow$  1.

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# $\begin{array}{ll} \mbox{Key: Provide conditions so that} \\ \mbox{a) } \| {\it N} \| \rightarrow b > 0, \quad \mbox{b) } {\it N}_{\pi} =_d {\it N}, \quad \mbox{c) } \| {\it S}_{\pi} \| \rightarrow 0. \end{array}$

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Key: Provide conditions so that  
a) 
$$||N|| \rightarrow b > 0$$
, b)  $N_{\pi} =_d N$ , c)  $||S_{\pi}|| \rightarrow 0$ .

Involved deriving new moment bounds

#### Numerical experiment

Setup: n = 500 samples with p = 300 features, r = 1 latent factor.

$$\boldsymbol{X} = \boldsymbol{\theta} \sqrt{\gamma} \boldsymbol{\eta} \boldsymbol{\Lambda}^{\top} + \boldsymbol{\mathcal{E}},$$

where  $\eta \sim \text{Unif}(\mathbb{S}^{n-1})$ ,  $\Lambda \sim \text{Unif}(\mathbb{S}^{p-1})$ , and  $\varepsilon_{ij} \stackrel{iid}{\sim} \mathcal{N}(0, 1/n)$ .



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Example:  $\varepsilon_{ij} \stackrel{ind}{\sim} \mathcal{N}(0, \omega_i^2/n)$ , 90% have  $\omega_i^2 = 0.4$ , 10% have  $\omega_i^2 = 1$ .



This heterogeneous data is less noisy, should be easier!







But it performs much worse ...



But it performs <u>much</u> worse...what is happening?



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Permutation shrinks the noise spectrum, leading to overselection.

Given: data matrix  $X \in \mathbb{R}^{n imes p}$  and percentile  $\alpha \in [0, 1]$ 

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Sign-flipping also recovers the "null" by destroying correlations.

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For a larger version of the same problem, i.e., bigger n, p:



Signflip PA also provides a good estimate of the noise spectrum.

Recall:  $\varepsilon_{ij} \stackrel{ind}{\sim} \mathcal{N}(0, \omega_i^2/n)$ , 90% have  $\omega_i^2 = 0.4$ , 10% have  $\omega_i^2 = 1$ .



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Signflips preserve the noise spectrum (in distribution).



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Preserving the noise distribution with signflips addresses the overselection of permutation.

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$$X = S + (X - S) = S + N,$$

where N = X - S is centered (since  $\mathbb{E}X = S$ ), but has dep. entries.

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Ongoing work: how do our insights about PA apply here?

Prelim experiment: rank-10 S matrix, diverse total count rates, ...



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Permutations seem to shrink the noise spectrum sometimes and signflips seem to preserve them...

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Ongoing: theoretical analysis/characterization - how to deal with the dependence among noise entries?

# Conclusions

Today:

- explaination for how parallel analysis works using insights/tools from random matrix theory
- some theoretical guarantees/characterization for parallel analysis
- signflip variant to handle alternative noise models
- preliminary work on applications to scRNAseq

Ongoing:

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