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Topic 18: Eigenvalue and Spike Models

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Key points: .

Disclaimer: The note is built on Prof. Jinchi Lv's lectures of the course at USC, DSO 607, High-Dimensional Statistics and Big Data Problems.

18.1 Motivation

Consider n independent observations $X_i \in \mathbb{R}^p$ drawn from a $\mathcal{N}(\mathbf{0}, \Sigma)$, then the covariance can be decomposed into 2 parts, white noise and low rank

$$\Sigma = \text{Cov}(\mathbf{X}_i) = \mathbf{I} + \sum_{k=1}^{M} \theta_k \nu_k \nu_k' = \Sigma_0 + \mathbf{\Phi}$$

where M denotes the **number of spikes** in the distribution of eigenvalues. The idea is: spikes deviate from a reference model along a <u>small fixed number</u> of unknown directions. If $\Phi = 0$, then none of the sample eigenvalues is separated from the bulk.

Why a spike model is interesting? A spike model can help determine the latent dimension of the data, some examples being

- Principal component analysis (PCA): spikes are related to the directions of the most variations of the data, i.e., the principal components
- Clustering model: M spikes is equivalent to M+1 clusters
- Economic significance: *M* is related to the number of factor loadings

Then the question is threefold:

- How to determine *M*
- How to estimate v_k
- How to test θ_k

Under rank one alternative, we would like to test the hypothesis

$$the H_1: \mathbf{\Sigma} = \mathbf{I}_p + \theta \mathbf{v} \mathbf{v}', \theta > 0$$

against the null

$$H_0: \mathbf{\Sigma} = \mathbf{I}_p$$

References