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Multivariate Theory for Analyzing High Dimensional Data

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Abstract: In this article, we develop a multivariate theory for analyzing multivariate datasets that have fewer observations than dimensions. More specifically, we consider the problem of testing the hypothesis that the mean vector $\boldsymbol{\mu}$ of a *p*-dimensional random vector \boldsymbol{x} is a zero vector where *N*, the number of independent observations on \boldsymbol{x} , is less than the dimension *p*. It is assumed that \boldsymbol{x} is normally distributed with mean vector $\boldsymbol{\mu}$ and unknown nonsingular covariance matrix \sum . We propose the test statistic $F^+ = n^{-2} (p - n + 1) N^- \boldsymbol{x}' S^{+-} \boldsymbol{x}$, where n = N - 1 < p, $^-\boldsymbol{x}$ and *S* are the sample mean vector and the sample covariance matrix respectively, and S^+ is the Moore-Penrose inverse of *S*. It is shown that a suitably normalized version of the F^+ statistic is asymptotically normally distributed under the hypothesis. The asymptotic non-null distribution in one sample case is given. The case when the covariance matrix \sum is singular of rank *r* but the sample size *N* is larger than *r* is also considered. The corresponding results for the case of two-samples and *k* samples, known as MANOVA, are given.

Key words: distribution of test statistics, DNA microarray data, fewer observations than dimension, multivariate analysis of variance, singular Wishart

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