# 國立交通大學

## 統計學研究所

## 碩 士 論 文

由混合變異建構基因表現分析之無母數檢定 Nonparametric Test based on Combined Variation for Gene Expression Analysis

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中華民國 一百 年 六 月

## 由混合變異建構基因表現分析之無母數檢定

### Nonparametric Test based on Combined Variation for Gene

## Expression Analysis

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Submitted to Institute of Statistics College of Science

National Chiao Tung University

in partial Fulfillment of the Requirements

for the Degree of Master

in

**Statistics** 

June 2011

Hsinchu, Taiwan, Republic of China

中華民國 一百 年 六 月

由混合變異建構基因表現分析之無母數檢定

研究生:連紫汝 2000 年 - 指導教授:陳鄰安博士

#### 國立交通大學統計學研究所



在致病基因的檢測問題上,因為 Tomlins et al.(2005)的發現使得 探討離群分配變成一個重要主題。不同於離群平均只能檢測中心位置 之改變,我們提出一個統計量它同時可以檢測中心與離心兩種變異。 這個統計量還有一個好處。由它所建立的檢定統計量不用估計未知分 配的密度函數值。我們利用模擬分析比較了幾種檢定方法的檢力並且 做了比較。我們也進一步做了一個簡單的實際資料分析。

## **Nonparametric Test based on Combined Variation for Gene Expression Analysis**

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 Observed by Tomlins et al. (2005), detection of the shift for outlierdistribution is a new topic useful in gene expression analysis. Alternative to the outlier mean test, we introduce a nonparametric statistic that can simultaneously detect the location shift and variation shift in the outlier distribution. There is an advantage, comparing with the outlier mean, that the test based on this statistic requires no prediction of distributional densities. Comparisons of this test statistic with some other methods in terms of mean square errors for estimation of their population parameters and powers for their abilities in detection of disease genes are simulated and displayed. Finally, a simple real data analysis is also performed and presented.

### 誌 謝

 畢業將至,回想起兩年的研究所生活,除了在專業上更精進之外,更 獲得許多學業以外的知識。

首先,最要感謝陳鄰安老師,謝謝您這些日子來的指導與教誨,從您 身上我學到做學問的態度,啟發我對研究的熱忱,都是支持我繼續努力的 動力。除此之外,老師也時常教導我許多人生哲理,使我獲益良多。一句 謝謝您,代表我最深的敬意。感謝口試委員許文郁老師、謝文萍老師以及 黃冠華老師對論文的指導與建議,使論文能更趨完善。

感謝所有的同學,彼此間的互相打氣,與論文奮戰的每一天,你們的 陪伴及鼓勵,都支持著我繼續努力下去,也因為有你們,讓我的碩士生涯 更多采多姿。謝謝親愛的室友們,不論在研究上、或在生活上妳們都是我 最佳的傾聽者,謝謝妳們無私的關心。感謝所辦的郭姐,協助我們處理大 小事務,讓我們可以專心做研究。感謝家人的支持,讓我能無後顧之憂的 繼續唸書,當我最強力的後盾?

謝謝每個曾經幫助過我的人,在此致上最深的謝意。

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#### 中華民國 一百 年 六 月





### Nonparametric Test based on Combined Variation for Gene Expression Analysis

#### SUMMARY

Observed by Tommins et al. (The signal-time al-al-time shift for the shift for the shift for  $\alpha$ is a new topic useful in gene expression analysis- Alternative to the outlier mean test, we introduce a nonparametric statistic that can simultaneously is an advantage, comparing with the outlier mean, that the test based on this statistic requires no prediction of distributional densities- Comparisons of this test statistic with some other methods in terms of mean square errors for estimation of their population parameters and powers for their abilities in detection of disease genes are simulated and displayed- Finally a simple real data analysis is also performed and presented-

key words over expression analysis Outlier and Outlier sum to a control to analysis of the sum to

#### - Introduction of the control of the

DNA microarray technology, which simultaneously probes thousands of gene expression profiles, has been successfully used in medical research for disease classication and all the contraction and all the contraction and all the contraction of the contract al- Sorlie et al- - Among the existed techniques in dieren tial genes detection, common statistical methods for two-group comparisons such as  $t$ -test, are not appropriate due to a large number of genes expressions and a limited number of subjects available-control statistical approaches approaches approaches have been proposed to identify those genes where only a subset of the sam  $\rho$  . And the matrix expression-them them the  $\lambda$  is the following them the  $\rho$  is the set of the that there is small number of outliers in samples of differential genes and then introduced a method called cancer outlier profile analysis that identifies outlier profiles by a statistic based on the median and the median absolute deviation of a generation product of the second contract of the sequence of  $\alpha$ approaches then concentrated on detecting differential genes based on outlier samples while Tibshirani and Hastie  $(2007)$  and Wu  $(2007)$  suggested to

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use an outlier sum the sum of all the gene expression values in the disease  $\alpha$  that are greater than a specific cuto point-disadvant-disadvant-disadvant-disadvant-disadvant-disadvant-disadvant-disadvant-disadvant-disadvant-disadvant-disadvant-disadvant-disadvant-disadvant-disadvant-disadvant-di vantage of these techniques is that the distribution theory of the proposed methods has not been discovered so that the distribution based  $p$  value can not been applied- Recently Chen Chen and Chan considered the outlier mean (average of outlier sum) and developed its large sample theory that allows us to formulate the  $p$  value based on its asymptotic distribution. For evaluation, they performed simulation studies in a parametric study by specifying the normal distribution- Although the outlier sum or outlier mean is shown interesting in detection of influential genes through statistical analysis and some real data analysis, however, these techniques can detect only the location shift in the outlier distribution, not the distributional variation.

We propose a statistic that can detect simultaneously the location shift and variation shift of the outlier distribution that is generalized from the combined control chart applied in quality control (see Cheng and Thaga , in a review- in Section , the reasons for the the compiler outlier quantity-  $\sim$  introduced and  $\sim$ tribution for the combined outlier quantity and use this theory to introduce a new test for gene expression analysis where a discussion of power based on the section that  $\pi$  are compared to the section between the section between the section and and and a test combined from the outlier mean and outlier variance is given- Finally the proofs of theorems are provided in Section 5.

#### 2. Combined Outlier Quantity

In a general study that consists of  $n_1$  subjects in the normal control group and n-2 sub jects in the disease group suppose that the m genes are m genes are m genes  $\sim$  be investigated. Their gene expression can be represented as  $\mathcal{I}_{ij}$ ,  $-$ ;  $-$ ;  $\cdots$ ;  $\cdots$ ;  $\cdots$ ,  $\cdots$   $\cdots$ - - - <sup>m</sup> for the disease group- However in our study we restrict on one gene with expression variable  $X$  for group of normal subject and expression variable <sup>Y</sup> for group of disease subject where the distribution functions for the first and  $\Lambda$  and  $\Gamma$  respectively. The summation is that we have  $\Lambda$  and  $\Lambda$  $\mathbf{1}, \mathbf{1}, \mathbf{1}, \mathbf{1}, \mathbf{1}, \mathbf{1}$  and  $\mathbf{1}, \mathbf{1}, \mathbf{1}, \mathbf{1}, \mathbf{1}, \mathbf{1}, \mathbf{1}, \mathbf{1}, \mathbf{1}$ 

An important observation by Tomlins et alprostate cancer, outlier genes are over-expressed only in a small number of disease samples- with defining a cuto point  $\mu$  according to the control of  $\mu$ data of the variable X, Tibshirani and Hastie  $(2007)$  and Wu  $(2007)$  considered the sum of variables  $Y_i$ 's that are over higher cutoff point  $\hat{\eta}$  given by  $\sum_{i=1}^{n_2} Y_i I(Y_i \geq \hat{\eta})$  as a test statistic for detection if the disease group distribution is dierent from the normal group distribution- Latter Chen Chen and Chan  $(2010)$  developed the asymptotic distribution for its average, called the outlier mean,  $Y_{out} = (\sum_{i=1}^{n_2} I(Y_i \geq \hat{\eta}))^{-1} \sum_{i=1}^{n_2} Y_i I(Y_i \geq \hat{\eta})$ for constructing a distribution based <sup>p</sup> value- In this paper we choose  $\eta = F_X^-(\gamma)$ , the population  $\gamma$ th quantile, and  $\eta = F_X^-(\gamma)$ , the  $\gamma$ th empiri- $\alpha$  -  $\alpha$ means for distributions of  $X$  and  $Y$  are

$$
\mu_{X,out} = E(X|X \ge F_X^{-1}(\gamma)) \text{ and } \mu_{Y,out} = E(Y|Y \ge F_X^{-1}(\gamma)) \tag{2.1}
$$

and the population type outlier variances are

$$
\sigma_{X,out}^2 = Var(X|X \ge F_X^{-1}(\gamma)) \text{ and } \sigma_{Y,out}^2 = Var(Y|Y \ge F_X^{-1}(\gamma)). \quad (2.2)
$$

The outlier means of the outlier means is to the out is to the out of your contract of the statistically different from  $\mu_{X,out}$  and the outlier variance based analysis is to test if  $\sigma_{Y,out}^{\perp}$  is statistically different from  $\sigma_{\bar{X}, out}$  . Then

For the following two distribution settings

Normal : 
$$
X \sim N(0, 1), Y \sim N(\theta, \sigma^2), \sigma = 0.5
$$
  
Mixed normal :  $X \sim N(0, 1), Y \sim 0.9N(0, 1) + 0.1N(\theta, \sigma^2), \sigma = 0.5$ 

we choose parameter values of  $\theta$  such that either outlier means are equal, i.e.,  $\mu_{X,out} = \mu_{Y,out}$ , or outlier variances are equal, i.e.,  $\sigma_{X,out} = \sigma_{Y,out}$ . In Table 1, we display, for each distribution setting, two outlier means, two outlier variances-

Table -- Equal outlier means and equal outlier variances



We have several comments for the results in Table 1:

which is the output of the output  $\mathcal{M} \setminus \mathcal{O}(u)$  and  $\mathcal{M} \setminus \mathcal{O}(u)$  is the set of  $\mathcal{M} \setminus \{0\}$ and the outlier variances  $o_{\mathbf{X},out}$  and  $o_{\mathbf{Y},out}$  for two three  $\gamma$  s in (11) are all identical-dentical-dentical-dentical-density distribution that for any underlying distribution there is  $\mathcal{L}$ chance that using outlier mean or outlier variance to test equality of two distributions may not be a propriate-

We then consider a test that can simultaneously interpret the combined change in both outlier mean  $\mu_{Y,out}$  and outlier variance  $\sigma_{Y,out}$ . The combined outlier quantity is defined as

$$
\sigma_{Y,X}^2 = E\{(Y - \mu_{X,out})^2 | Y \geq F_X^{-1}(\gamma)\}.
$$

This combined outlier quantity when  $Y$  and  $X$  have the same distribution is

$$
\sigma_{X,out}^{2} = E\{(X - \mu_{X,out})^{2} | X \geq F_{X}^{-1}(\gamma)\}.
$$

The aim of combined outlier quantity is to verify if  $\sigma_{Y,X}^{\perp}$  and  $\sigma_{X,out}^{\perp}$  are identical. In Table 1, the values of combined outlier quantity  $\sigma_{Y,X}^-$  in all two distributions and different  $\gamma$  s are displayed. With a comparison of  $\sigma_{Y,X}^$ and  $\sigma_{\bar{X},out}^{-}$  in all situations, these two quantities are basically not identical.

This allows us to propose a combined outlier quantity based test for gene expression analysis.

We further consider the following three types of distribution setting,

Type 1: 
$$
X \sim N(0, 1), Y \sim (\chi^2(10) + \theta),
$$
  
Type 2:  $X \sim t(10), Y \sim 0.9t(10) + 0.1N(\theta, \sigma^2), \sigma = 1,$   
Type 3:  $X \sim t(10), Y \sim 0.9t(10) + 0.1(\chi^2(10) + \theta).$ 



in Table  $2$ .

#### Table - Comparison of outlier means and outlier variances

	$Df_m$	$Df_v$	$Df_{com}$
Type 1			
$\theta = 0, \gamma = 0.85$	3.594	25.86	38.78
$\gamma=0.9$	4.340	27.38	46.22
$\gamma=0.95$	5.480	27.16	57.20
$\theta = 2$			
$\gamma=0.85$	4.444	35.10	54.85
$\gamma=0.9$	5.392	36.60	65.67
$\gamma=0.95$	6.853	34.83	81.80
$\theta = 4$			
$\gamma=0.85$	5.296	46.29	74.33
$\gamma=0.9$	6.444	47.81	89.35
$\gamma=0.95$	8.232	44.19	111.9
Type 2			
$\theta = 2, \gamma = 0.85$	0.222	0.167	0.216
$\gamma=0.9$	0.205	0.122	0.164
$\gamma=0.95$	0.153	0.044	0.067
$\theta = 4$			
$\gamma=0.85$	0.965	1.519	2.451
$\gamma=0.9$	$1.062$ and $\blacksquare$	1.337	2.467
$\gamma=0.95$	1.118	0.953	2.203
Type 3			
$\theta = 0, \gamma = 0.85$	3.517	25.05	37.42
$\gamma=0.9$	4.218	26.33	44.12
$\gamma=0.95$	5.245	25.85	53.37
$\theta=2$			
$\gamma=0.85$	4.368	189 534.11	53.19
$\gamma=0.9$	5.268	35.31	63.08
$\gamma=0.95$	6.614	33.23	76.99
$\theta = 4$			
$\gamma=0.85$	5.219	45.12	72.36
$\gamma=0.9$	6.321	46.29	86.26
$\gamma=0.95$	7.994	42.30	106.2

It is seen that the differences of combined outlier quantities are much more larger than the other two differences. This probably indicates that the combined outlier quantity may be more efficient in detecting the influential genes.

The sample estimator of combined outlier quantity is defined as

$$
S_{Y,X}^{2} = \left[\sum_{i=1}^{n_2} I(Y_i \geq \hat{F}_X^{-1}(\gamma))\right]^{-1} \sum_{i=1}^{n_2} (Y_i - \hat{\mu}_{X,out})^2 I(Y_i \geq \hat{F}_X^{-1}(\gamma)),
$$

where the sample outlier mean is  $\hat{\mu}_{X,out} = [\sum_{i=1}^{n_1} I(X_i \geq F_X^{-1}(\gamma))]^{-1} \sum_{i=1}^{n_1} X_i I(X_i \geq$  $F_{|X|}(\gamma)$ ). It is also interesting to evaluate the efficiencies in estimating the parameters of outlier mean, outlier variance and combined outlier quantity. We denote the mean square errors for  $\mu_{X,out}, \mu_{Y,out}, \sigma_{X,out}$ ,  $\sigma_{Y,out}$  and  $\sigma_{Y,X}$  are, respectively, as *MSE<sub>px*, out</sub>, *MSE<sub>py*, out</sub>, *MSE<sub>* $\sigma_{X,out}^2$ , *MSE<sub>* $\sigma_{X,out}^2$ ,</sub></sub>  $\frac{1}{\sigma}$   $\frac{1}{\sigma}$   $\frac{1}{\sigma}$   $\frac{1}{\sigma}$   $\frac{1}{\sigma}$   $\frac{1}{\sigma}$  and  $\frac{1}{\sigma}$  and  $\frac{1}{\sigma}$   $\frac{1}{\sigma}$   $\frac{1}{\sigma}$   $\frac{1}{\sigma}$ 

$$
X_1, ..., X_n
$$
 iid  $N(0, 1), Y_1, ..., Y_n$  iid  $0.9N(0, 1) + 0.1N(\mu, 1)$ 

we display these results in Table 3.

 $\blacksquare$  and  $\blacksquare$  . The comparison for parameters is the comparison for  $\setminus \setminus \cdot \bot$  ,  $\blacksquare$  ,  $\setminus \cdot \bot$  $30)$ 

	$\bar{M}SE_{\mu_{X,out}}$	$\overline{M}SE_{\mu_{Y,out}}$	. $\widehat{MSE}_{\sigma^2_{X,out}}$	, $\overline{MSE}_{\sigma^2_{Y,\underline{out}}}$	$\widetilde{MSE}_{\sigma^2_{Y,X}}$
$\mu = 1$					
$\gamma=0.85$	0.0977	0.0996	0.0288	0.0426	0.1007
$\gamma=0.9$	0.1235	0.1283	0.0283	0.0382	0.1269
$\gamma=0.95$	0.2191	0.1788	0.0263	0.0335	0.1580
$\mu=3$					
$\gamma=0.85$	0.0981	0.241996	0.0276	0.3354	1.2345
$\gamma=0.9$	0.1240	0.2895	0.0306	0.3415	1.3698
$\gamma=0.95$	0.2137	0.3587	0.0265	0.3270	1.8171

It is seen that the MSE's for combined outlier quantity are relatively larger than the other outlier mean and outlier variance quantity- This is due to that a quantity that can simultaneously predict the difference in outlier mean and outlier variance should be more dicult- The appropriateness of the test based on combined outlier quantity needs to be justied through the power comparisons-

#### 3. The Test based on Combined Outlier Quantity

We here introduce some asymptotic properties of the combined outlier quantity and then provide a test based on its asymptotic distribution $\mathbf{I}$  and  $\mathbf{I}$  and  $\mathbf{I}$  are proposed to the contract of  $\mathbf{I}$ 

$$
n_2^{1/2}(S_{Y,X}^2 - \sigma_{Y,X}^2) = n_1^{-1/2} \sum_{i=1}^{n_1} [\lambda_1(\gamma - I(X_i \le F_X^{-1}(\gamma))) + \lambda_2(X_i - \mu_{X,out})
$$
  

$$
I(X_i \ge F_X^{-1}(\gamma))] + \beta_Y^{-1} n_2^{-1/2} \sum_{i=1}^{n_2} [(Y_i - \mu_{X,out})^2 - \sigma_{Y,X}^2] I(Y_i \ge F_X^{-1}(\gamma)) + o_p(1)
$$

where we let

$$
\lambda_1 = -\left[\beta_Y^{-1}(F_X^{-1}(\gamma) - \mu_{X,out})^2 f_Y(F_X^{-1}(\gamma)) f_X^{-1}(F_X^{-1}(\gamma)) + 2\beta_X^{-1} F_X^{-1}(\gamma) (\mu_{Y,out} - \mu_{X,out})\right]
$$
  
\n
$$
\lambda_2 = -2\beta_X^{-1}(\mu_{Y,out} - \mu_{X,out})
$$
  
\nwith  $\beta_Y = P(Y \ge F_X^{-1}(\gamma)), \beta_X = 1 - \gamma.$ 

(b) We have  $n_2^{\gamma}$  (S<sub>V Y</sub> –  $_2$   $\sigma_{\bar{Y},X} - \sigma_{\bar{Y},out}$  converges in distribution to *I*V(0,  $v_y$ ) where

$$
v_y = \gamma (1 - \gamma) \lambda_1^2 + \lambda_2^2 E[(X - \mu_{X,out})^2 I(X \ge F_X^{-1}(\gamma))] - 2\lambda_1 \lambda_2 (1 - \gamma)
$$
  
\n
$$
E[(X - \mu_{X,out})I(X \ge F_X^{-1}(\gamma))] + \beta_Y^{-2} E\{ (Y - \mu_{X,out})^4 I(Y \ge F_X^{-1}(\gamma)) \} - \sigma_{Y,X}^4.
$$
  
\nwhere  $\beta_Y^{-2} E\{ (Y - \mu_{X,out})^4 I(Y \ge F_X^{-1}(\gamma)) \} - \sigma_{Y,X}^4 = Var[(Y - \mu_{X,out})^2 | Y \ge F_X^{-1}(\gamma)].$   
\n1896

From the above the theorem then  $\tau$  we have the  $\mu$  we have the following the following  $\alpha$ 

$$
P_{H_0}\left\{\sqrt{n_2}(\frac{S_{Y,X}^2-\sigma_{X,X}^2}{\sqrt{v_Y}})\leq z\right\}\to\int_{-\infty}^z\phi(z)dz
$$

for  $z \in R$  where  $\phi$  represents the probability density function of  $N(0,1)$ . If we further have  $\sigma_{X,X}^{\mathbf{x}}$  and  $v_Y$ , respectively, estimates of  $\sigma_{X,X}^{\mathbf{x}}$  and  $v_Y$ , we may define an outlier combined test as

rejecting 
$$
H_0
$$
 if  $n_2^{1/2} \left( \frac{S_{Y,X}^2 - \hat{\sigma}_{X,X}^2}{\sqrt{\hat{v}_Y}} \right) \ge z_\alpha$ . (3.1)

Having this outlier combined test, it is desired to verify the power performance of this test when there exists distributional shift for the disease group

distribution- An approximate power with signicant level may be derived as bellows

$$
\pi_{Y} = P_{F_{Y}} \{ \sqrt{n_{2}} (\frac{S_{Y,X}^{2} - \hat{\sigma}_{X,X}^{2}}{\sqrt{\hat{v}_{Y}}}) \ge z_{\alpha} \}
$$
\n
$$
= P_{F_{Y}} \{ \sqrt{n_{2}} (\frac{S_{Y,X}^{2} - \sigma_{Y,X}^{2}}{\sqrt{v_{Y}}}) \ge \frac{z_{\alpha} \sqrt{\hat{v}_{Y}} + \sqrt{n_{2}} (\hat{\sigma}_{X,X}^{2} - \sigma_{Y,X}^{2})}{\sqrt{v_{Y}}} \}
$$
\n
$$
\approx P \{ Z \ge z_{\alpha} + \sqrt{n_{2}} (\frac{\sigma_{X,X}^{2} - \sigma_{Y,X}^{2}}{\sqrt{v_{Y}}}) \}
$$
\n(3.2)

- test decreased in the form  $\alpha$  is consistent for parameters that we consider the state of  $\alpha$ rameter van die there is die there is die providing economic there is a monotory and the computational computa volved in the interest of the state in this dict that the complete and  $\eta$  there is no control to the control of test is restricted on size  $\alpha$  when two distributions  $F_Y$  and  $F_X$  are identical.

**Corollary 3.2.** When Y and X have the same distribution, we have, by the fact that  $\sigma_{X,X}^-=\sigma_{X,out}^-,$ 

$$
n_2^{1/2}(S_{Y,X}^2 - \sigma_{X,out}^2)
$$
  
=  $-\beta_X^{-1}(F_X^{-1}(\gamma) - \mu_{X,out})^2 n_1^{-1/2} \sum_{i=1}^{n_1} (\gamma - I(X_i \le F_X^{-1}(\gamma)))$   
+  $\beta_X^{-1} n_2^{-1/2} \sum_{i=1}^{n_2} [(X_i - \mu_{X,out})^2 - \sigma_{X,out}^2] I(X_i \ge F_X^{-1}(\gamma)) + o_p(1).$ 

We have  $n_2$  (S<sub>V v</sub> –  $_2$  ( $S_{Y,X} - o_{X,out}$ ) converges in distribution to  $N(0, v_X)$  where  $v_X = \beta_X^-\gamma(1-\gamma)(F_X^-(\gamma)\top\mu_{X,out})^+$ 

$$
+ \beta_X^{-2} E[(X - \mu_{X,out})^4 I(X \ge F_X^{-1}(\gamma))] - \sigma_{X,out}^4.
$$

Suppose that we have estimators  $\sigma_{X,X}^+$  and  $v_X$ , respectively, for estimation of  $\sigma_{X,X}$  and  $v_X$ . We then can define the following test

Combined test : rejecting 
$$
H_0
$$
 if  $n_2^{1/2} \frac{S_{Y,X}^2 - \hat{\sigma}_{X,X}^2}{\sqrt{\hat{v}_X}} > z_\alpha$ . (3.3)

The interest by applying the issue  $\{ \circ \circ \circ \}$  in the  $\Lambda$  itself involves no density, point so that estimation of it is much easier- We can similarly derive the approximate power for the above test as

$$
\pi_X = P_X \{ \sqrt{n_2} (\frac{S_{Y,X}^2 - \hat{\sigma}_{X,X}^2}{\sqrt{\hat{v}_X}}) \ge z_\alpha \}.
$$
 (3.4)

Power representations - and - provide approximate powers based on tests in the powers of t when the underlying distributions for control group and disease group as

$$
X \sim N(0, 1)
$$
 and  $Y \sim (1 - \delta)N(0, 1) + \delta N(\theta, 1)$ .

Table - Table -  $\frac{1}{2}$  for mixed normal distribution norma

	$\theta=1$	$\theta=3$	$\theta=5$	$\theta = 10$
$\delta = 0.1$				
$\gamma=0.8$	0.078	0.281	0.281	0.541
$\gamma=0.85$	0.074	0.247	0.414	0.534
$\delta = 0.2$				
$\gamma=0.8$	0.098	0.422	0.682	0.825
$\gamma=0.85$	0.090	0.345	0.618	0.810

with simulation study in the study in th propriate powers for the theory in the theory in the actually in the propriate  $\mathbf{r}_1$ the critical points  $\alpha$  require and will any mighting  $\alpha$  . This in this induction can add the model section.

## 4. Power Comparison by Simulation and a Simple Real Data Analysis

Two tasks will be done in this section- First we will show bysimulation that the setting of critical point  $\mathbf{p}$  and  $\mathbf{p}$  is  $\mathbf{p}$  of  $\mathbf{p}$  of  $\mathbf{p}$  and  $\mathbf{p}$  and  $\mathbf{p}$ conservative and we will study present the appropriate level  $\alpha$  critical point. Second, we will compare this outlier combined test with a combination of  $\mathbf{F}$ a change in distributional mean and F-test is to detect a change in distributional variation-test and Free and Fre shift in mean and variation simultaneously- It is then desired to compare powers of these two combined tests-

A t and F combined test is

rejecting 
$$
H_0
$$
 if  $\frac{\bar{Y} - \bar{X}}{S_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}} > t_{\alpha/2}(n_1 + n_2 - 2)$   
or  $\frac{S_X^2}{S_Y^2} > F_{\alpha/2}(n_1 - 1, n_2 - 1)$  or  $< \frac{1}{F_{\alpha/2}(n_2 - 1, n_1 - 1)}$ 

where  $S_p^2 = \frac{\sum_{i=1}^{n_1} (X_i - \bar{X})^2 + \sum_{i=1}^{n_2} (Y_i - \bar{Y})^2}{n_1 + n_2 - 2}, S_X^2 = \frac{1}{n_1 - 1} \sum_{i=1}^{n_1} (X_i - \bar{X})^2$  and  $\frac{X_1 + X_2}{n_1 + n_2 - 2}$ ,  $S_X^2 = \frac{1}{n_1 - 1} \sum_{i=1}^{n_1} (X_i - X)^2$  and  $S_Y^2 = \frac{1}{n_2-1}\sum_{i=1}^{n_2} (Y_i - Y)^2.$ 

We consider a simulation with sample size n  $\alpha$  and  $\beta$  and  $\alpha$  and  $\alpha$   $\alpha$   $\alpha$   $\alpha$  and  $\alpha$   $\alpha$   $\alpha$   $\alpha$   $\alpha$ <sup>m</sup> - to evaluate the power when <sup>X</sup> and<sup>Y</sup> are from the following setting of distribution

$$
X \sim N(0, 1)
$$
 and  $Y \sim 0.9N(0, 1) + 0.1N(\theta, 1)$ .

In Tables 5 and 6, we display the simulated results for  $n = 50$  and  $n = 100$ when level of significance is  $0.05$  and in Table 7, we display the simulated results for  $n = 50$  when  $\alpha = 0.1$ .

We have comments for the results in Tables  $5, 6$  and  $7$ :

a Although the contamination percentage of outlier in mixed normal dis tribution is small as  $0.1$  the combined outlier quantity of cutoff with small  $\gamma$ 's are more powerful than it with larger  $\gamma$ 's.

(b) The tests based on the combined outlier quantity of cutoff with small  $\gamma$ 's are relatively more powerful than the t and F combined than the t and F combined test-ful than the t and F combi that simultaneously detect the shift in outlier mean and outlier variance is appropriate when we choose  $\gamma$  appropriately for the cutoff.

 $(c)$  The power for the test based on the combined outlier quantity is increasing when the contaminated location shift  $\theta$  is increasing.

We next consider that alternative distribution has a constant shift as

77 T N

Setting 
$$
I: X \sim N(0, 1)
$$
 and  $Y \sim (1 - \delta)N(0, 1) + \delta\{\theta\}$   
Setting  $II: X \sim N(0, 1)$  and  $Y \sim (1 - \delta)t(10) + \delta\{\theta\}$ 

We list the simulated results in Tables 8-11.

We consider a real data of control group and disease group that includes  $22$ , 285 genes. Considering the significance level  $\alpha = 0.05$ , the constants  $z$  is in table are the critical points designed to ensure that the sizes of the tests and s are appropriately vivve control in a control percentages of general control of  $\mathcal{A}$ numbers to be regulated for all the respective tests in all the respective tests in a linear position of the c results are displayed in Table 12.

	Outlier mean	Outlier variance	Combined q	$t$ -test
$\gamma=0.6$	$0(z^* = 2.68)$	$0.1247(z^* = 2.85)$	$0.1195(z^* = 3.07)$	0.0325
$\gamma=0.65$	$0(z^* = 2.46)$	$0.1263(z^* = 2.97)$	$0.1168(z^* = 3.28)$	
$\gamma=0.7$	$0(z^* = 2.21)$	$0.1286(z^* = 3.11)$	$0.1165(z^* = 3.51)$	
$\gamma=0.75$	$0(z^* = 1.86)$	$0.1256(z^* = 3.42)$	$0.1157(z^* = 3.85)$	
$\gamma=0.8$	$0(z^* = 1.54)$	$0.1242(z^* = 3.74)$	$0.1092(z^* = 4.35)$	
$\gamma=0.85$	$0(z^* = 1.23)$	$0.1230(z^* = 4.45)$	$0.1102(z^* = 5.28)$	
$\gamma=0.9$	$0.00004(z^* = 1.01)$	$0.1247(z^* = 5.54)$	$0.1049(z^* = 7.25)$	
$\gamma=0.95$	$0.0004(z^* = 0.77)$	$0.1267(z^* = 9.15)$	$0.1154(z^* = 15.5)$	

Table -- Percentages of genes larger than critical values

We have several comments on the results in this table

(a) It is seen that the outlier mean test performed poorly with very low percentages of genes to be rejected. This shows that it can not detect any that  $\mathcal{C}$ gene as influentials. WWW.

b The tests based on outlier variance and outlier combined quantity are with relatively moderatively percentages of genes between constraints in an interesting  $\sim$ the genes are measured simultaneously from the same subjects, there is need a simultaneous test that would remarkedly reduce the percentages of genes to be considered in the case of the pure we pure we will not further the case of the pure we were asset that the see that only outlier variance and outlier combined quantity are with hope to be able to be able to general between in the second state of the second state o

#### Appendix

Three assumptions for the two sample outlier variance test are as follows.

ASSUMPTION 1: The limit  $\gamma = \lim_{n_1, n_2 \to \infty} n_1^{-n_2}$  exists.

assumption in the state  $\alpha$  assumption for distribution for distribution for  $\Lambda$  is a contract of  $\Lambda$ bounded away from zero in neighborhoods of  $F_X^{-1}(\alpha)$  for  $\alpha \in (0,1)$  and the population cutoff point  $\eta$ .

assumption in the contraction of the contraction for the contract and the contract of the cont zero in a neighborhood of the population cutoff point  $\eta$ .

Proof of Theorem - First we consider the following expansion

$$
\sum_{i=1}^{n_2} (Y_i - \hat{\mu}_{X,out})^2 I(Y_i \ge \hat{F}_X^{-1}(\gamma)) = \sum_{i=1}^{n_2} (Y_i - \mu_{X,out})^2 I(Y_i \ge \hat{F}_X^{-1}(\gamma))
$$
  
+  $(\hat{\mu}_{X,out} - \mu_{X,out})^2 \sum_{i=1}^{n_2} I(Y_i \ge \hat{F}_X^{-1}(\gamma)) - 2(\hat{\mu}_{X,out} - \mu_{X,out}) \sum_{i=1}^{n_2} [(Y_i - \mu_{Y,out}) + n_2(\mu_{Y,out} - \mu_{X,out})] I(Y_i \ge \hat{F}_X^{-1}(\gamma)).$  (5.1)

From the theory for the outlier mean by Chen, Chen and Chan (2010), we may see that  $n_{2}^{-}$  ( $\mu_{Y,out}$ )  $\overline{Q}^{\prime\;\;\tau}(\mu_{Y,out}-\mu_{Y,out})=O_{p}(1),\ n_{1}^{\;\;\tau\;\;\tau}(\mu_{X,out}-\mu_{X,out})=O_{p}(1).$ and  $n_{\alpha}$   $\rightarrow$   $\alpha$ .  $\sum_{i=1}^{n/2} \sum_{i=1}^{n_2} (Y_i - \mu_{Y,out}) I(Y_i \geq F_X^{-1}(\gamma)) = O_p(1)$ . We then, from (5.1), may re-write the combined quantity as

$$
n_2^{1/2}(S_{Y,X}^2 - \sigma_{Y,X}^2)
$$
  
\n
$$
= n_2^{1/2}(\sum_{i=1}^{n_2} I(Y_i \ge \hat{F}_X^{-1}(\gamma)))^{-1}\left{\sum_{i=1}^{n_2} (Y_i - \mu_{X,out})^2 [I(Y_i \ge F_X^{-1}(\gamma) + n_2^{-1/2}T)\right}
$$
  
\n
$$
- I(Y_i \ge F_X^{-1}(\gamma))] + \sum_{i=1}^{n_2} [(Y_i - \mu_{X,out})^2 - \sigma_{Y,X}^2] I(Y_i \ge F_X^{-1}(\gamma))\}
$$
  
\n
$$
- 2(\mu_{Y,out} - \mu_{X,out}) n_2^{1/2} (\hat{\mu}_{X,out} - \mu_{X,out}) + o_p(1),
$$
  
\nwhere we let  $T = n_1^{1/2}(\hat{F}_X^{-1}(\gamma) - F_X^{-1}(\gamma)).$  (5.2)

with Assumptions and techniques from Ruppert and techniques from Ruppert and techniques from Ruppert and Rupper  $(1980)$  and Chen  $\&$  Chiang  $(1996)$ , we may see that

$$
n_2^{-1/2} \sum_{i=1}^{n_2} (Y_i - \mu_{X,out})^2 [I(Y_i \ge F_X^{-1}(\gamma) + n_2^{-1/2} T^*) - I(Y_i \ge F_X^{-1}(\gamma))]
$$
  
= -(F\_X^{-1}(\gamma) - \mu\_{X,out})^2 f\_Y(F\_X^{-1}(\gamma)) T^\* (5.3)

for any sequence  $T^* = O_p(1)$ .

We may also see from Chen, Chen and Chang  $(2010)$  that the outlier mean and following representation and the following representation of the following representa

$$
n_1^{1/2}(\hat{\mu}_{X,out} - \mu_{X,out}) = \beta_X^{-1} F_X^{-1}(\gamma) n_1^{-1/2} \sum_{i=1}^{n_1} (\gamma - I(X_i \le F_X^{-1}(\gamma)))
$$
  
+  $\beta_X^{-1} n_1^{-1/2} \sum_{i=1}^{n_1} (X_i - \mu_{X,out}) I(X_i \ge F_X^{-1}(\gamma)) + o_p(1).$  (5.4)

The result of this theorem is induced by plugging - and - into and applying a representation for empirical quantile  ${F}_X^-(\gamma)$  in Chen, Chen and Chang -



	$p_{tF}$	$p_{com}$
$\gamma=0.6, \theta=0$	$0.049(\alpha^* = 0.034)$	$0.049(z_{\alpha^*} = 4.02)$
$\gamma = 0.65, \theta = 0$		$0.049(z_{\alpha^*} = 4.47)$
$\gamma = 0.7, \theta = 0$		$0.049(z_{\alpha^*} = 4.89)$
$\gamma = 0.75, \theta = 0$		$0.049(z_{\alpha^*}=5.74)$
$\gamma = 0.8, \theta = 0$		$0.0473(z_{\alpha^*}=6.83)$
$\gamma = 0.85, \theta = 0$		$0.0493(z_{\alpha^*} = 9.55)$
$\gamma = 0.9\theta = 0$		$0.0492(z_{\alpha^*}=14.53)$
$\gamma=0.6$		
$\theta=1$	0.092	0.101
$\theta=3$	0.539	0.755
$\theta=5$	0.921	0.978
$\theta=10$	0.994	0.995
$\gamma = 0.65$		
$\theta=1$		0.097
$\theta=3$		0.733
$\theta=5$		0.978
$\theta=10$		0.995
$\gamma=0.7$		
$\theta=1$		0.095
$\theta=3$		0.717
$\theta=5$		0.975
$\theta=10$		0.995
$\gamma = 0.75$		
$\theta = 1$		0.090
$\theta=3$		0.683
$\theta=5$	1896	0.972
$\theta=10$		0.995
$\gamma=0.8$		
$\theta=1$		0.085
$\theta=3$		0.639
$\theta=5$		0.966
$\theta=10$		0.994
$\gamma=0.85$		
$\theta = 1$		0.078
$\theta = 3$		0.562
$\theta=5$		0.949
$\theta=10$		0.995
$\gamma=0.9$		
$\theta=1$		0.075
$\theta=3$		0.478
$\theta=5$		0.912
$\theta=10$		0.995

Table 5. Power comparison between t and F combined test and outlier combined test  $(n=50,\alpha=0.05)$ 

	$p_{tF}$	$p_{com}$
$\gamma = 0.6, \theta = 0$	$0.049(\alpha^* = 0.034)$	$0.049(z_{\alpha^*}=3.07)$
$\gamma = 0.65, \theta = 0$		$0.049(z_{\alpha^*}=3.28)$
$\gamma = 0.7, \theta = 0$		$0.049(z_{\alpha^*}=3.51)$
$\gamma=0.75, \theta=0$		$0.049(z_{\alpha^*}=3.85)$
$\gamma = 0.8, \theta = 0$		$0.0484(z_{\alpha^*}=4.35)$
$\gamma = 0.85, \theta = 0$		$0.0494(z_{\alpha^*} = 5.28)$
$\gamma = 0.9\theta = 0$		$0.0497(z_{\alpha^*}=7.25)$
$\gamma=0.6$		
$\theta=1$	0.123	0.136
$\theta=3$	0.821	0.949
$\theta=5$	0.994	0.999
$\theta = 10$	0.999	0.999
$\gamma = 0.65$		
$\theta=1$		0.128
$\theta=3$		0.943
$\theta=5$		0.999
$\theta = 10$		0.999
$\gamma = 0.7$		
$\theta=1$		0.121
$\theta=3$		0.933
$\theta=5$		0.999
$\theta = 10$		0.999
$\gamma = 0.75$		
$\theta=1$	<b>ETER</b>	0.116
$\theta=3$		0.920
$\theta=5$	1896	0.999
$\theta = 10$		0.999
$\gamma=0.8$		
$\theta=1$		0.108
$\theta=3$		0.898
$\theta=5$		0.999
$\theta = 10$		0.999
$\gamma=0.85$		
$\theta=1$		0.098
$\theta=3$		0.854
$\theta=5$		0.998
$\theta = 10$		0.999
$\gamma=0.9$		
$\theta=1$		0.089
$\theta=3$		0.763
$\theta=5$		0.997
$\theta = 10$		0.999

Table 6. Power comparison between t and F combined test and outlier combined test  $(n=100, \alpha=0.05)$ 

	$p_{tF}$	$p_{com}$
$\gamma = 0.6, \theta = 0$	$0.099(\alpha^* = 0.067)$	$0.099(z_{\alpha^*}=2.18)$
$\gamma = 0.65, \theta = 0$		$0.099(z_{\alpha^*}=2.30)$
$\gamma = 0.7, \theta = 0$		$0.099(z_{\alpha^*}=2.440)$
$\gamma=0.75, \theta=0$		$0.099(z_{\alpha^*}=2.620)$
$\gamma = 0.8, \theta = 0$		$0.0983(z_{\alpha^*}=2.93)$
$\gamma = 0.85, \theta = 0$		$0.0993(z_{\alpha^*} = 3.41)$
$\gamma = 0.9\theta = 0$		$0.1001(z_{\alpha^*} = 4.41)$
$\gamma=0.6$		
$\theta=1$	0.205	0.229
$\theta=3$	0.887	0.975
$\theta=5$	0.997	0.999
$\theta = 10$	0.999	0.999
$\gamma = 0.65$		
$\theta=1$		0.217
$\theta=3$		0.973
$\theta=5$		0.999
$\theta = 10$		0.999
$\gamma=0.7$		
$\theta=1$		0.212
$\theta=3$		0.968
$\theta=5$		0.999
$\theta = 10$		0.999
$\gamma = 0.75$		
$\theta=1$		0.202
$\theta=3$		0.962
$\theta=5$	1896	0.999
$\theta = 10$		0.999
$\gamma=0.8$		
$\theta=1$		0.189
$\theta=3$		0.950
$\theta=5$		0.999
$\theta = 10$		0.999
$\gamma=0.85$		
$\theta=1$		0.180
$\theta=3$		0.931
$\theta=5$		0.999
$\theta = 10$		0.999
$\gamma=0.9$		
$\theta=1$		0.166
$\theta=3$		0.883
$\theta=5$		0.999
$\theta = 10$		0.999

Table 7. Power comparison between t and F combined test and outlier combined test  $(n=50,\alpha=0.1)$ 

	$p_{tF}$	$p_{com}$
$\gamma = 0.6, \theta = 0$	$0.049(\alpha^* = 0.034)$	$0.049(z_{\alpha^*} = 4.02)$
$\gamma = 0.65, \theta = 0$		$0.049(z_{\alpha^*}=4.47)$
$\gamma=0.7, \theta=0$		$0.049(z_{\alpha^*} = 4.89)$
$\gamma = 0.75, \theta = 0$		$0.049(z_{\alpha^*} = 5.74)$
$\gamma = 0.8, \theta = 0$		$0.0473(z_{\alpha^*}=6.83)$
$\gamma = 0.85, \theta = 0$		$0.0493(z_{\alpha^*} = 9.55)$
$\gamma = 0.9\theta = 0$		$0.0492(z_{\alpha^*}=14.53)$
$\gamma=0.6$		
$\theta=3$	0.482	0.717
$\theta=5$	0.925	0.984
$\theta=10$	0.995	0.996
$\gamma = 0.65$		
$\theta=3$		0.685
$\theta=5$		0.984
$\theta=10$		0.995
$\gamma=0.7$		
$\theta=3$		0.658
$\theta=5$		0.982
$\theta=10$		0.995
$\gamma = 0.75$		
$\theta=3$		0.608
$\theta=5$		0.978
$\theta=10$	1896	0.995
$\gamma=0.8$		
$\theta=3$		0.554
$\theta=5$		0.974
$\theta=10$		0.995
$\gamma = 0.85$		
$\theta=3$		0.451
$\theta=5$		0.956
$\theta=10$		0.995
$\gamma=0.9$		
$\theta=3$		0.361
$\theta=5$		0.917
$\theta=10$		0.995

Table 8. Power comparison between t and F combined test and outlier combined test  $(n=50,\alpha=0.05,\delta=0.1)$ 

	$p_{tF}$	$p_{com}$
$\gamma = 0.6, \theta = 0$	$0.049(\alpha^* = 0.034)$	$0.049(z_{\alpha^*} = 4.02)$
$\gamma = 0.65, \theta = 0$		$0.049(z_{\alpha^*} = 4.47)$
$\gamma = 0.7, \theta = 0$		$0.049(z_{\alpha^*} = 4.89)$
$\gamma = 0.75, \theta = 0$		$0.049(z_{\alpha^*} = 5.74)$
$\gamma = 0.8, \theta = 0$		$0.0473(z_{\alpha^*}=6.83)$
$\gamma = 0.85, \theta = 0$		$0.0493(z_{\alpha^*} = 9.55)$
$\gamma = 0.9\theta = 0$		$0.0492(z_{\alpha^*} = 14.53)$
$\gamma=0.6$		
$\theta=3$	0.883	0.939
$\theta=5$	0.999	0.999
$\theta = 10$	0.999	0.999
$\gamma = 0.65$		
$\theta=3$		0.913
$\theta=5$		0.999
$\theta=10$		0.999
$\gamma=0.7$		
$\theta=3$		0.887
$\theta=5$		0.999
$\theta = 10$		$\mathbf{1}$
$\gamma=0.75$		
$\theta=3$		0.824
$\theta=5$		0.999
$\theta = 10$	1896	0.999
$\gamma=0.8$		
$\theta=3$		0.753
$\theta=5$		0.999
$\theta = 10$		$\mathbf{1}$
$\gamma = 0.85$		
$\theta=3$		0.605
$\theta=5$		0.995
$\theta = 10$		0.999
$\gamma=0.9$		
$\theta=3$		0.467
$\theta=5$		0.975
$\theta = 10$		0.999

Table 9. Power comparison between t and F combined test and outlier combined test  $(n=50,\alpha=0.05,\delta=0.2)$ 

	$p_{tF}$	$p_{com}$
$\gamma = 0.6, \theta = 0$	$0.049(\alpha^* = 0.034)$	$0.049(z_{\alpha^*} = 4.02)$
$\gamma = 0.65, \theta = 0$		$0.049(z_{\alpha^*} = 4.47)$
$\gamma = 0.7, \theta = 0$		$0.049(z_{\alpha^*} = 4.89)$
$\gamma = 0.75, \theta = 0$		$0.049(z_{\alpha^*} = 5.74)$
$\gamma = 0.8, \theta = 0$		$0.0473(z_{\alpha^*}=6.83)$
$\gamma = 0.85, \theta = 0$		$0.0493(z_{\alpha^*} = 9.55)$
$\gamma = 0.9\theta = 0$		$0.0492(z_{\alpha^*}=14.53)$
$\gamma=0.6$		
$\theta=3$	0.620	0.772
$\theta=5$	0.951	0.987
$\theta = 10$	0.995	0.996
$\gamma = 0.65$		
$\theta=3$		0.739
$\theta=5$		0.986
$\theta = 10$		0.996
$\gamma=0.7$		
$\theta=3$		0.709
$\theta=5$		0.986
$\theta = 10$		0.996
$\gamma = 0.75$		
$\theta=3$		0.658
$\theta=5$	1896	0.982
$\theta=10$		0.996
$\gamma=0.8$		
$\theta=3$		0.596
$\theta=5$		0.976
$\theta = 10$		0.996
$\gamma = 0.85$		
$\theta=3$		0.484
$\theta=5$		0.956
$\theta=10$		0.996
$\gamma=0.9$		
$\theta=3$		0.384
$\theta=5$		0.912
$\theta = 10$		0.995

Table 10. Power comparison between t and F combined test and outlier combined test  $(n=50,\alpha=0.05,\delta=0.1)$ 

	$p_{tF}$	$p_{com}$
$\gamma = 0.6, \theta = 0$	$0.049(\alpha^* = 0.034)$	$0.049(z_{\alpha^*} = 4.02)$
$\gamma = 0.65, \theta = 0$		$0.049(z_{\alpha^*} = 4.47)$
$\gamma = 0.7, \theta = 0$		$0.049(z_{\alpha^*}=4.89)$
$\gamma = 0.75, \theta = 0$		$0.049(z_{\alpha^*} = 5.74)$
$\gamma = 0.8, \theta = 0$		$0.0473(z_{\alpha^*}=6.83)$
$\gamma = 0.85, \theta = 0$		$0.0493(z_{\alpha^*} = 9.55)$
$\gamma = 0.9\theta = 0$		$0.0492(z_{\alpha^*} = 14.53)$
$\gamma=0.6$		
$\theta=3$	0.923	0.950
$\theta=5$	0.999	0.999
$\theta=10$	0.999	0.999
$\gamma = 0.65$		
$\theta=3$		0.925
$\theta=5$		0.999
$\theta = 10$		$\mathbf{1}$
$\gamma=0.7$		
$\theta=3$		0.900
$\theta=5$		0.999
$\theta = 10$		0.999
$\gamma=0.75$		
$\theta=3$		0.842
$\theta=5$		0.999
$\theta=10$	1896	0.999
$\gamma=0.8$		
$\theta=3$		0.768
$\theta=5$		0.999
$\theta = 10$		0.999
$\gamma = 0.85$		
$\theta=3$		0.617
$\theta=5$		0.995
$\theta = 10$		0.999
$\gamma=0.9$		
$\theta=3$		0.473
$\theta=5$		0.973
$\theta = 10$		0.999

Table 11. Power comparison between t and F combined test and outlier combined test  $(n=50,\alpha=0.05,\delta=0.2)$ 

#### **REFERENCES**

- Agrawal, D., Chen, T., Irby, R., et al. (2002). Osteopontin identified as lead marker of colon cancer progression, using pooled sample expression profiling. J. Natl. Cancer Inst., 94, 513-521.
- Alizadeh, A. A., Eisen, M. B., Davis, R. E., et al. (2000). Distinct types of diffuse large B-cell lymphoma identified by gene expression profiling. Nature, 403, 503-511.
- Chen, L.-A., Chen, Dung-Tsa and Chan, Wenyaw. (2010). The p Value for the Outlier Sum in Differential Gene Expression Analysis. *Biometrika*, 97, 246-253.
- Chen, L.-A. and Chiang, Y. C. (1996). Symmetric type quantile and trimmed means for location and linear regression model. *Journal of Nonpara*metric Statistics.,  $7, 171-185$ .
- Cheng, S. W., and Thaga, K. (2006). On single variable control charts: an overview. Quality and Reliability Engineering International, 22, 811-820.
- Hoaglin, D. C., Mosteller, F. and Tukey, J. W. (1983). Understanding Robust and Exploratory Data Analysis, Wiley: New York.
- Ohki, R., Yamamoto, K., Ueno, S., et al. (2005). Gene expression profiling of human atrial myocardium with atrial fibrillation by DNA microarray analysis. *Int. J. Cardiol.* **102**, 233-238.
- Ruppert, D. and Carroll, R.J. (1980). Trimmed least squares estimation in the linear model. Journal of American Statistical Association 75, 828-838.
- Sorlie, T., Tibshirani, R., Parker, J., eta l. (2003). Repeated observation of breast tumor subtypes in independent gene expression data sets. *Proc.* Natl. Acad. Sci. U.S.A., 100, 8418-8423.
- Tibshirani, R. and Hastie, T. (2007). Outlier sums differential gene expression analysis. *Biostatistics*, 8, 2-8.

Tomlins, S. A., Rhodes, D. R., Perner, S., eta l. (2005). Recurrent

fusion of TMPRSS2 and ETS transcription factor genes in prostate cancer. Science, 310, 644-648.

Wu, B. (2007). Cancer outlier differential gene expression detection. Biostatistics, 8, 566-575.

