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## **Standard Error of Area Under One Parameter Bi-exponential ROC Curve**

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#### **Abstract**

In this paper, properties of Bi-Exponential Receiver Operating Characteristic (ROC) curve and Area Under the ROC curve (AUC) of Bi-Exponential ROC model are discussed. The evaluation of AUC and its inference is a crucial part of ROC curve analysis. We proposed the estimation of asymptotic and exact variances for AUC of Bi-Exponential ROC curve. The Confidence Intervals for AUC using asymptotic and exact variance methods are also proposed. The proposed methods are validated by extensive simulation study as well as real life example.

**Keywords**: Area Under the Bi-Exponential ROC Curve, Bi-Exponential ROC model, Kullback-Leibler divergence, and TPR asymmetry. (10pt Times New Roman, Justified).

#### **Introduction**

In Medical diagnosis, a subject is categorized into either *healthy* or *diseased* based on some clinical measurement based on the selected cutoff *t*. If the clinical measurement is 'greater than or equal to' *t*, then the subject is labeled as diseased and if the measurement is 'less than' *t*, then the subject is labeled as healthy subjects. The clinical measurements are often called as test results or test scores or Biomarker.

Let us denote the test results of diseased subject by the random variable *Y* with Probability Density Function (PDF),  $g_Y(y)$  and Cumulative Distribution Function (CDF),  $G_Y(y)$ . Similarly, let us denote the test results of healthy subject by the random variable X with PDF,  $f_X(x)$  and CDF,  $F_X(x)$ . Assume that *X* and *Y* are independent and continuous.

*Sensitivity* of the diagnostic test is defined as  $G_Y(t) = P(Y > t)$ , which is the probability of correctly categorizing a diseased subject when a cutoff is given. Similarly, the *Specificity* of the diagnostic test is defined as  $F_X(t) = P(X \le t)$ , which is the probability of correctly categorizing a healthy subject.

*Receiver Operating Characteristic* (ROC) curve is defined as a plot of "*True Positive Rate*" (TPR),  $\overline{G}_Y(t)$  on the vertical axis versus the "*False* 

*Positive Rate*" (FPR),  $F_X(t)$  on the horizontal axis for different values of t, where  $-\infty < t < \infty$ . The mathematical model representing the ROC curve takes the form [8]

$$
\text{ROC}[t] = \overline{G}_{Y}^{\circ} \overline{F}_{X}^{-1}(t); 0 \le \overline{F}_{X}(t) \le 1 \tag{1}
$$

For an appropriate diagnostic test, the ROC curve should lie very close to upper left corner of the unit square. A typical ROC curve must satisfy the following properties.

- 1. ROC curve is invariant with respect to monotone increasing transformation of the test scores.
- 2. The test values of *X* are smaller than *Y*.
- 3. ROC curve is monotonically increasing function  $dROC(t)$

i.e. 
$$
\frac{\text{d} \overline{\text{d} \overline{\text{f}}_X(t)}}{\text{d} \overline{\text{f}}_X(t)} > 0
$$

- 4. ROC curve is said to be concave, if 2 2 X  $\frac{d^2ROC(t)}{d}$  < 0  $\frac{d^2 \text{ROC}(t)}{d\overline{F}_x^2(t)}$  < 0 and convex, if  $\frac{d^2}{dt^2}$ 2 X  $\frac{d^2 \text{ROC}(t)}{d} > 0.$  $d\overline{F}_X^2(t)$  $\mathbf{I}$
- 5. The slope of ROC curve at any operating point is equal to the ratio of PDF of diseased to PDF of healthy at a particular cut-off point [8] which is given by

slope = 
$$
\frac{g(t)}{f(t)}
$$
 (2)

6. Let  $KL(f, g)$  denote the Kullback – Leibler (K-L) divergence [6] between the distributions of healthy and diseased group with f(x) as the

comparison distribution and g(y) as the reference distribution. Then

$$
KL(f, g) = \int_{D} f(x) \ln \left( \frac{f(x)}{g(y)} \right) dx
$$
 (3)

where  $D \in x \cap y$ ;  $0 < x < \infty$ ;

 $0 < y < \infty$  since the range of x and y are same, we can represent x and y by x itself.

Similarly, let KL(g, f) denote the K-L divergence between the distribution of diseased and healthy population with  $g(x)$  as the comparison distribution and f(x) as the reference distribution, then

$$
KL(g, f) = \int_{D} g(x) \ln \left( \frac{g(x)}{f(x)} \right) dx
$$
 (4)

where D is the common range of f and g. It is to be noted that  $KL(f, g)$  and  $KL(g, f)$  are positive and  $KL(f, g) = KL(g, f) = 0$ , if and only if  $f(x) = g(x)$ . These two measures tell us about the asymmetry of ROC curve about the negative diagonal. If  $KL(f, g) < KL(g, f)$ , then the ROC curve is said to be TPR asymmetric and if  $KL(f, g)$  >  $KL(g, f)$ , then the curve is said to be True Negative Rate (TNR) asymmetric.

Area under the ROC curve is the frequently used measure for quantifying the performance of the diagnostic test. It is defined as the probability that in a randomly selected pair of healthy and diseased subject, the test result of diseased subject is higher than the healthy subject. Mathematically, it is defined as

$$
AUC = \int_0^1 \overline{G}_Y(t) d\overline{F}_X(t).
$$
 (5)

The evaluation of AUC and its inference is a crucial part of ROC curve analysis. The estimation of ROC curve is done in three ways like Non-Parametric, Parametric and Semi-Parametric. In this paper, we have studied the parametric approach of plotting ROC curve.

In Parametric approach a specific distribution is assumed to X and Y with different parametric values. The conventional Bi-Normal ROC model assumes that X and Y or any monotone transformation of X and Y follow normal distribution with different parameters where the parameters with subscript '0' represents healthy parameters and with subscript '1' represents diseased parameter.

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The other distributional models that are considered in literature are Bi-Exponential ROC model [2], Bi-Gamma ROC model [4], Bi-Lomax ROC model [3], Generalized Bi-Exponential ROC model [7], Bi-Lognormal ROC model [1], Bi-Rayleigh ROC model [9], [10], Bi-Weibull ROC model [11] and a review of all parametric ROC models in case of continuous data [12].

This paper is organized as follows: In Section 2, Bi-Exponential ROC model, its properties and ML estimation of parameters are discussed. Section 3, provides estimation of AUC, asymptotic distribution of estimated AUC and confidence interval for  $\angle$ AUC. In Section 4, exact distribution of  $\widehat{AUC}$  and confidence interval for  $\widehat{AUC}$  are discussed. The proposed theory is validated by simulation studies and real life example in Section 5. Section 6 contains the concluding remarks.

### **Materials and methods Bi-Exponential ROC Model**

Let Z be a random variable that follows the one parametric Exponential distribution with inverse scale parameter  $\lambda$  is denoted by Exp ( $\lambda$ ). It possess the PDF as

$$
f_Z(z,\lambda) = \lambda e^{-\lambda z}; z > 0, \lambda > 0.
$$
 (6)

The CDF of random variable Z is given by

$$
F_Z(z) = P(Z \le z) = 1 - e^{-\lambda z}, \lambda > 0
$$
\n(7)

Bi-Exponential ROC model which is given by [2] λ1

$$
eROC(t) = \overline{F}_X(t)^{\overline{\lambda_0}}; 0 \le \overline{F}_X(t) \le 1, \lambda_0 > \lambda_1
$$
 (8)

where  $\lambda_0$  and  $\lambda_1$  are the parameter of healthy and diseased group respectively, "eROC (t) " represent the Bi-Exponential ROC model.

Now, we will discuss some of the properties of eROC curve

### **Properties**

1. eROC curve is monotonically increasing function.

Proof: A function is said to be a monotone increasing function, if the first derivative of the function is positive. Since, the first derivative of eROC curve with respect to  $F_X(t)$  is positive.

i.e. 
$$
\frac{\text{deROC}(t)}{\text{d}\overline{F}_X(t)} = \frac{\lambda_1}{\lambda_0} \left[ \overline{F}_X(t) \right] \left( \frac{\lambda_1}{\lambda_0} - 1 \right) > 0
$$
(9)

Hence, eROC curve is monotonically

increasing function.

2. eROC curve is concave and never lies below the chance line.

Proof: From equation (9), the second derivative of  $eROC(t)$  is given by

$$
\frac{d^2eROC(t)}{d\overline{F}_X^2(t)} = \frac{\lambda_1}{\lambda_0} \left(\frac{\lambda_1}{\lambda_0} - 1\right) \left[\overline{F}_X(t)\right] \left(\frac{\lambda_1}{\lambda_0} - 2\right) < 0
$$
\n(10)

The ratio  $\frac{\lambda_1}{\lambda_2}$ *0*  $\lambda_{\rm i}$ λ will always be less than zero since

we assumed that  $\lambda_0 > \lambda_1$  and hence the term  $\left( \begin{array}{ccc} & & & \end{array} \right)$ 

$$
\left(\frac{\lambda_1}{\lambda_0} - 1\right) < 0 \text{ and } \left[\overline{F}_X(t)\right] \left(\frac{\lambda_1}{\lambda_0} - 2\right) > 0 \text{ since }
$$

 $0 \le F_X(t) \le 1$ . On the whole, we will get  $\frac{d^2eROC(t)}{d} < 0$ . Hence, eROC curve is concave  $d\overline{F}_X^2(t)$ 

in nature. Now, let us prove that it never lies below the chance line.

ROC curve is said to be proper ROC curve if it never crosses the chance line or if the decision variable is a strictly increasing function of the likelihood ratio. Consider any two points *a* and *b* (say) where  $0 \le a$ ,  $b \le 1$  on eROC curve. Since we have proved that the eROC curve is concave, the line segment connecting the point *a* and *b* never lies above the curve. If we take the extreme point i.e.  $a = 0$  and  $b = 1$ , it becomes the chance line, so the chance line also never lies above the curve. Hence we have proved that the eROC curve never crosses the chance line.

3. The slope of the eROC curve at the threshold t is given by

slope = 
$$
\frac{\lambda_1}{\lambda_0} e^{\left\{t(\lambda_0 - \lambda_1)\right\}}
$$
 (11)

- 4. It is invariant with respect to monotone increasing transformation of the test scores.
- 5. eROC curve is TPR asymmetric [6]. Proof:

The K-L divergence between two exponential distributions has been studied by [6]. The K-L divergence between the distribution of diseased and healthy group with  $f(x)$  as the comparison distribution and  $g(x)$  as the reference distribution has been given as

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$$
KL(f, g) = \frac{\lambda_1}{\lambda_0} - 1 - \ln\left(\frac{\lambda_1}{\lambda_0}\right)
$$
 (12)

Similarly, the K-L divergence between the distribution of healthy and diseased group with  $g(x)$  as the comparison distribution and  $f(x)$  as the reference distribution has been given as

$$
KL(g, f) = \frac{\lambda_0}{\lambda_1} - 1 - \ln\left(\frac{\lambda_0}{\lambda_1}\right)
$$
 (13)

It was found that  $KL(g, f)$  >  $KL(f, g)$ . These two divergence measures would be zero, if the healthy and diseased group is identical. Hence, we have proved that, the eROC curve is TPR asymmetric.

# **Asymptotic Distribution of Bi-Exponential AÛC** and Confidence Interval for  $\widehat{\rm {AUC}}$

The area under the eROC curve is computed as

$$
AUC = P(Y > X) = \frac{\lambda_0}{\lambda_0 + \lambda_1}.
$$
\n(14)

where  $\lambda_0$  and  $\lambda_1$  are the parameters of healthy and diseased group respectively. In order to compute the estimated AUC, we need to find the MLE's of  $\lambda_0$  and  $\lambda_1$ . In the following section, we will find the MLEs of  $\lambda_0$  and  $\lambda_1$ .

By substituting the ML estimates  $\hat{\lambda}_0$  and  $\hat{\lambda}_1$  we get the ML estimate of AUC i.e.

$$
\widehat{AUC} = \frac{m \sum_{j=1}^{n} y_j}{n \sum_{i=1}^{m} x_i + m \sum_{j=1}^{n} y_j}
$$
(15)

Now, we will derive the asymptotic distribution and confidence interval of AUC. To evaluate the significance of the statistic  $\angle AUC$ , we need to compute its variance and standard error. The following theorem evaluates the variance of the estimate, AÛC.

**Theorem 3.1**: The area under the eROC curve will converge in distribution to a Normal random variable

with mean zero and variance  ${}^{2}_{0}\lambda^{2}_{1}$  $(1 + \lambda_0)^4$  $(m+n)$ mn  $(\lambda_1 + \lambda_0)$  $+n)$   $\lambda_0^2 \lambda_1^2$  $\lambda_1 + \lambda_0$ for

 $\text{large } N(=m+n)$ .

#### **Proof:**

Let  $L(\theta \mid x, y)$ ;  $\theta = (\lambda_0, \lambda_1)$ be the likelihood function of the sample observations from X and Y which is given by

$$
\ln L = m \ln \lambda_0 - \lambda_0 \sum_{i=1}^{m} x_i + n \ln \lambda_1 - \lambda_1 \sum_{j=1}^{n} y_j
$$
 (16)

Asymptotic normality of MLE's, states that a consistent solution of the likelihood equation is asymptotically normally distributed about the true value  $\theta$  i.e.  $\hat{\theta} \sim N(\theta, I^{-1}(\theta)).$ 

$$
\Rightarrow \sqrt{N}(\hat{\theta} - \theta) \rightarrow N(0, I^{-1}(\theta)). \tag{17}
$$

where  $I(\theta)$  is the Fisher Information matrix which is given by

$$
I(\theta) = -\begin{bmatrix} E\left(\frac{\partial^2 \ln L}{\partial \lambda_0^2}\right) & E\left(\frac{\partial^2 \ln L}{\partial \lambda_0 \partial \lambda_1}\right) \\ E\left(\frac{\partial^2 \ln L}{\partial \lambda_1 \partial \lambda_0}\right) & E\left(\frac{\partial^2 \ln L}{\partial \lambda_1^2}\right) \end{bmatrix}
$$
(18)

where

$$
a_{11} = \frac{m}{\lambda_0^2}, \ a_{22} = \frac{n}{\lambda_1^2}, \qquad a_{12} = a_{21} = 0.
$$

The  $I^{-1}(\theta)$  is calculated as

$$
I^{-1}(\theta) = \begin{bmatrix} V(\hat{\lambda}_0) & Cov(\hat{\lambda}_0, \hat{\lambda}_1) \\ Cov(\hat{\lambda}_1, \hat{\lambda}_0) & V(\hat{\lambda}_1) \end{bmatrix}
$$

$$
= \begin{bmatrix} \frac{I}{a_{11}} & 0 \\ 0 & \frac{I}{a_{22}} \end{bmatrix}.
$$
(19)

where

$$
V(\hat{\lambda}_0) = \frac{\lambda_0^2}{m}, \text{Cov}(\hat{\lambda}_0, \hat{\lambda}_1) = 0
$$
  
\n
$$
\text{Cov}(\hat{\lambda}_1, \hat{\lambda}_0) = 0, \text{V}(\hat{\lambda}_1) = \frac{\lambda_1^2}{n}.
$$
\n(20)

Since area under the ROC curve is a function of parameters  $\theta = (\lambda_0, \lambda_1)$ , we will adopt the Delta method for finding the approximate variance.  $V(A\hat{U}C)$  can be defined as follows:

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$$
V(A\hat{U}C) = \left(\frac{\partial AUC}{\partial \lambda_0}\right)^2 V(\hat{\lambda}_0) + \left(\frac{\partial AUC}{\partial \lambda_1}\right)^2 V(\hat{\lambda}_1) +
$$
  

$$
2\left(\frac{\partial AUC}{\partial \lambda_0}\right) \left(\frac{\partial AUC}{\partial \lambda_1}\right) Cov(\hat{\lambda}_0, \hat{\lambda}_1).
$$
  

$$
= \frac{\lambda_0^2 \lambda_1^2}{(\lambda_0 + \lambda_1)^4} \left(\frac{m+n}{mn}\right)
$$
 (21)

where  $V(\hat{\lambda}_0)$ ,  $V(\hat{\lambda}_1)$  and  $Cov(\hat{\lambda}_0, \hat{\lambda}_1)$  and  $Cov(\hat{\lambda}_1, \hat{\lambda}_0)$  are taken from the equation (20). The estimate of variance is obtained by substituting the estimates of the parameters  $\lambda_0, \lambda_1$ .

The standard error of  $\angle AUC$  can be obtained by taking square root of  $V(A\hat{U}C)$  in equation (21). The  $100(1-\alpha)$ % confidence interval is obtained by 2  $[A\hat{U}C \pm SE(A\hat{U}C)Z_{\alpha}]$  (22)

where  $\alpha$  is the level of significance and  $Z_{\alpha/2}$  is the critical value.

It is observed that the asymptotic confidence interval do not perform well for small sample sizes. Hence, we propose the exact distribution of estimated AUC and confidence interval for small sample size in the following section.

### **Exact Confidence interval for AÛC**

From Section 3, the ML estimate of AUC was

found to be

$$
\hat{\text{AUC}} = \frac{\hat{\lambda}_0}{\hat{\lambda}_0 + \hat{\lambda}_0} = \frac{(\text{m/U})}{(\text{m/U}) + (\text{n/V})}
$$
(23)

where

$$
\hat{\lambda}_0 = \frac{m}{\sum_{i=1}^{m} x_i} = \frac{m}{U} \text{ and } \hat{\lambda}_0 = \frac{n}{\sum_{j=1}^{n} y_j} = \frac{n}{V}.
$$
\n(24)

It is well known that, if  $X \sim Exp(\lambda_0)$  then U follows gamma distribution with shape parameter *m* and scale parameter  $\lambda_0$  and similarly *V* follows gamma distribution with shape parameter *n* and scale parameter  $\lambda_1$ . Hence,  $2\lambda_0 V$  and  $2\lambda_1 U$  are independently distributed as  $\chi^2$  with 2m and 2n degrees of freedom.

**Lemma 1**: Let  $\delta_1$  and  $\delta_2$  be independent  $\chi^2$  random variables with m and n degrees of freedom respectively, the random variable [13]

$$
F = \frac{\delta_1 / m}{\delta_2 / n}
$$
 (25)

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[715]

is said to have an F- distribution with (m, n) degrees of freedom. From the Lemma, it follows that,

$$
F = \frac{(2\lambda_1 U/2m)}{(2\lambda_0 V/2n)} = \frac{(\lambda_1 U/m)}{(\lambda_0 V/n)} \sim F(2m, 2n)
$$
 (26)

The 100(1- $\alpha$ ) % confidence interval for AUC can be

obtained from F distribution through  $P(F_1 < F < F_u)$  where  $F_1$  and  $F_u$  satisfies the condition

where  $F_1$  and  $F_u$  satisfies the condition  $P(F_1 < F < F_n) = 1 - \alpha$ .

After a simple manipulation, the  $100 (1-\alpha)\%$ confidence interval is found to be

$$
\left[\frac{\text{A}\hat{\text{UC}} F_{(2\,\text{m},\,2\text{n},\,\alpha/2)}}{\text{A}\hat{\text{UC}} F_{(2\,\text{m},\,2\text{n},\,\alpha/2)} + (1 - \text{A}\hat{\text{UC}})} ,\, \frac{\text{A}\hat{\text{UC}} F_{(2\,\text{m},\,2\text{n},\,1-\alpha/2)}}{\text{A}\hat{\text{UC}} F_{(2\,\text{m},\,2\text{n},\,1-\alpha/2)} + (1 - \text{A}\hat{\text{UC}})}\right]
$$
\n(27)

where F2m,  $_{2n,\alpha/2}$  and  $F_{2m}$ ,  $_{2n,1-\alpha/2}$  are the lower and upper  $\alpha/2$ <sup>th</sup> percentile points of F distribution with 2m and 2n degrees of freedom.

# **Results and discussion**

## **Simulation Studies**

In this section, we provide the results of estimation of asymptotic, exact variance and confidence interval for AUC using simulated datasets.

### **Asymptotic Variance Method**

In this section, we did simulation studies to observe how the asymptotic variance of AUC behaves using simulated data sets. We have considered four different samples of size  $(m, n)$  = (30, 30) with different parametric values for  $\lambda_0 =$ {0.0829, 0.1456, 0.2035, 0.4167} and  $\lambda_1$  = {0.0466, 0.0423, 0.0267, 0.0311}. The estimated parameters,  $\angle AUC$ ,  $SE(AUC)$ , Confidence

Interval for  $\angle AUC$  are shown in Table 1.

From the Table 1, it is observed that, as the accuracy increases and the standard error decreases simultaneously. The ROC curves for different parametric values are plotted in Figure 1. In table 2, we have simulated independent samples of *m* healthy and *n* diseased ( $m = n = 5, 10, 30, 60, 80$ ) to assess the behavior of asymptotic variance and confidence interval by fixing  $\lambda_0$  and by varying  $\lambda_1$ . Similarly, in Table 3, we have simulated independent samples of *m* healthy and *n* diseased of size ( $m = n = 5, 10, 30$ , 60, 80) by fixing  $\lambda_1$  and varying  $\lambda_0$ . In each row, the first element represents the test's  $\angle AUC$ , second element (bolded) represents the  $SE(A\hat{U}C)$ , third and

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fourth element represent the lower and upper confidence limit respectively. Figure 2 represents, the pattern of standard error with respect to the sample size by keeping  $\lambda_0$  constant as in Table 2 and Figure 3 represents the pattern of standard error with respect to the sample size by keeping  $\lambda_1$  constant as in Table 3 by using asymptotic variance method.

From Table 2 and 3, it is observed that  $SE(A\hat{U}C)$ , decreases with increase in sample size and increase in accuracy. The behavior is depicted in Figure 2 and 3. It is observed that the asymptotic MLE do not perform well for small sample size. If we observe the lower confidence limit, it is crossing the lowest accuracy 0.5 which is not being considered as a good AUC estimate. As we mentioned earlier, if the curve reaches upper left corner of unit square then AUC is close to one. The topmost curve corresponds to the estimates ( $\lambda_0$  $\hat{\lambda}_0$ ,  $\hat{\lambda}_1$  = (0.4001, 0.0305), the curve just below the topmost curve corresponds to (0.2134, 0.0238), the third curve from top corresponds to (0.1054, 0.0359) and the lowest curve corresponds to the estimates (0.0833, 0.0447).





#### **Exact Variance Method**

In this section, we used the same parameters and estimated value to observe how the exact variance method behaves. The estimated parameters, AÛC, SE(AÛC), Confidence Interval for  $A\hat{U}C$ , are shown in Table 4.

In Table 5 and 6, the AUC and SE are analyzed by fixing one parametric value and varying the other and vice versa for different sample sizes and results are presented.

By comparing Table 2 with Table 5 and Table 3 with Table 6, we observe that there is a considerable difference in the two method adopted. The exact variance method possesses a shorter confidence

interval when compared with asymptotic variance method.

## **Figure 2**:



**Figure 3**:



#### *Standard error versus sample size* **Real Life Example**

Biomarkers are extremely useful for detecting the disease at early stages. Prostate Specific Antigen (PSA) is a biomarker which is significant in detecting the prostate cancer. For the application of proposed model, we consider the biomarker called total serum PSA. The data has been obtained from a Phase 3 prostate cancer Case-Control study nested in the Beta Carotene [6]. The data used in this paper consisted randomly selected individuals who are affected by prostate cancer and 50 healthy individuals who were participated in a lung cancer prevention trial.

First of all, let us evaluate the goodness of fit test for one parameter exponential distribution using Kolmogorov-Smirnov, Anderson-Darling and Chi-Square statistics. The statistic and p-value for all the three procedures is shown in Table 7.

For healthy marker values, the probabilities of the modeled K-S (0.1342), A-D (0.9977) and Chi-Square (6.2572) was greater than or equal to the level of significance of 20% (0.2000). Hence, there is no

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evident that the null hypothesis "The healthy marker values fits the one parameter Exponential distribution" to be rejected. Similarly in diseased group, the p-values of the modeled statistics were greater than the level of significance of 10% (0.1000). Hence, we conclude that the diseased marker values fit the assumed Exponential distribution.

The estimated parameters are  $\lambda_0 = 2.3086$  and

 $\lambda_1$  = 0.6479. Hence the AUC, standard errors using asymptotic MLE and exact variance methods are estimated as 0.7799, 0.034498 and 0.034477 respectively. The 95% asymptotic and exact confidence intervals for estimated AUC are constructed as [0.7123, 0.8476] and [0.7133, 0.8484] respectively. From the result, it is obvious that the asymptotic MLE method and exact variance method both performs in the similar way. The sensitivity and specificity of the marker are found to be 71% and 71% respectively at the threshold value of 0.534. From the sensitivity and specificity rates, we infer that an individual whose is having the "ratio of free to total PSA" marker value greater than 0.534 are 71% likely to be detected with the prostate cancer. Similarly, an individual having the marker value less than 0.534 are 71% likely to be not having the prostate cancer. The ROC curve plotted is plotted for the PSA marker and the asymmetric property of the ROC curve is also studied which is shown in Figure 6.

The line segment connecting  $(0,1)$  and  $(1,0)$  is called the negative chace line and it is obtained by plotting FPR on X-axis and 1-FPR on Y-axis. The dashed vertical line segment let us call it as  $S_1$ corresponds to the co-ordinate  $[FPR=a (0.08, say), 0]$  $\leq$  TPR  $\leq$  1]. The dashed horizontal line (S<sub>2</sub>) segment corresponds to the co-ordinate  $[0 \leq FPR \leq 1, TPR=1$ a (0.95)]. Let A=[a,0.5], B=[0.5, 1-a] and C=[a\*>a, 1a \* ]. A ROC curve is said to be symmetric if it passes through the co-ordinate A, B and C. Any ROC curve is said be TPR asymmetric if it passess through S2 after the co-ordinate B and the one that passes though S2 before the co-ordinate B is called TNP asymmetric. Thus from Figure 6, it is proved that the Bi-Exponential ROC curve is TPR asymmetric graphically. From equation (3) and (4), the K-L divergence measures

#### $KL(g, f)(1.2927) > KL(f, g)(0.5514)$

Hence, the eROC curve is TPR asymmetry is proved numerically.

#### **Conclusion**

Some of the properties of the Bi-Exponential ROC model have been studied. It was found that Bi-Exponential ROC curve satisfies some common properties like monotonicity and invariance property. It also satisfies some specific properties like concavity and TPR asymmetric. We have developed two methods to estimate the variance of Bi-Exponential AUC namely asymptotic variance method and exact variance method. The  $100(1-\alpha)$  % confidence interval for AUC using both method have been computed. Through extensive simulation studies and by looking at the real life example, we found that exact variance methods perform slightly better than the asymptotic variance method. Each of the method has to be adopted according to the specific conditions i.e. asymptotic variance can be adopted if the sample size is large else one can go for exact variance.

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| on eROC through asymptotic variance method |        |        |  |                 |                                   |  |  |  |  |
|--|--------|--------|--|-----------------|-----------------------------------|--|--|--|--|
| <b>S.N.</b>                                | ^Ո     | /∿1    | $\mathbf{A}\mathbf{\hat{U}}\mathbf{C}$ | $SE(A\hat{U}C)$ | 95% Confidence<br><b>Interval</b> |  |  |  |  |
|  | 0.0833 | 0.0447 | 0.6508                                 | 0.0587          | [0.5358, 0.7658]                  |  |  |  |  |
|  | 0.1054 | 0.0359 | 0.7458                                 | 0.0489          | [0.6499, 0.8418]                  |  |  |  |  |
|  | 0.2134 | 0.0238 | 0.8996                                 | 0.0233          | [0.8540, 0.9453]                  |  |  |  |  |
|  | 0.4001 | 0.0305 | 0.9292                                 | 0.0170          | [0.8959, 0.9645]                  |  |  |  |  |

Table  $1\,\mathrm{A}\hat{\text{U}}\text{C}$ , SE( $\mathrm{A}\hat{\text{U}}\text{C}$ ) and 95% confidence interval for  $\mathrm{A}\hat{\text{U}}\text{C}$  based





| <b>Sample Size</b><br>$\mathbf{v_0}$ | (5,5)      | (10,10)  | (30, 30) | (60, 60)   | (80, 80)   |
|--------------------------------------|------------|----------|----------|------------|------------|
| 0.0829                               | 0.6402     | 0.6402   | 0.6402   | 0.6402     | 0.6402     |
|                                      | 0.1457     | 0.1030   | 0.0595   | 0.0421     | 0.0364     |
|                                      | [0.3546]   | [0.4382] | [0.5236] | [0.5577]   | [0.5688]   |
|                                      | $0.9257$ ] | 0.8421]  | 0.7567   | $0.7226$ ] | $0.7115$ ] |
| 0.1278                               | 0.7328     | 0.7328   | 0.7328   | 0.7328     | 0.7328     |
|                                      | 0.1238     | 0.0876   | 0.0506   | 0.0358     | 0.0310     |
|                                      | [0.4901]   | [0.5612] | [0.6337] | [0.6627]   | [0.6721]   |
|                                      | $0.9755$ ] | 0.9044]  | 0.83191  | 0.8029]    | 0.79351    |
| 0.2056                               | 0.8152     | 0.8152   | 0.8152   | 0.8152     | 0.8152     |
|                                      | 0.0953     | 0.0674   | 0.0389   | 0.0275     | 0.0238     |
|                                      | [0.6285]   | [0.6832] | [0.7390] | [0.7613]   | [0.7685]   |
|                                      | 1.00001    | 0.94731  | 0.89151  | 0.86911    | 0.86191    |
| 0.5562                               | 0.9227     | 0.9227   | 0.9227   | 0.9227     | 0.9227     |
|                                      | 0.0451     | 0.0319   | 0.0184   | 0.0130     | 0.0113     |
|                                      | [0.8343]   | [0.8602] | [0.8866] | [0.8972]   | [0.9006]   |
|                                      | 1.0000]    | 0.98521  | 0.95881  | 0.94821    | $0.9448$ ] |

 $\bf{Table 3 \, A \hat{U}C}$  ,  $\bf{SE}(\bf{A \hat{U}C})$  , 95% confidence interval for  $\bf{A \hat{U}C}$  by fixing  $\lambda_1$  = 0.0466 and varying  $\lambda_0$ 

 $\hat{\textbf{a}}$   $\textbf{b}$  and  $\hat{\textbf{b}}$  and  $\hat{\textbf{c}}$  and  $\hat{\textbf{c}}$  and  $\hat{\textbf{c}}$  and  $\hat{\textbf{c}}$  and  $\hat{\textbf{c}}$  and  $\hat{\textbf{c}}$ 

| based on Bi-Exponential ROC through exact variance method |         |                |                          |                  |  |  |  |  |
|---|---------|----------------|--------------------------|------------------|--|--|--|--|
| <b>S.N.</b>   |         |                | $\boldsymbol{\hat{AUC}}$ | 95% Confidence   |  |  |  |  |
|   | $n_{0}$ | n <sub>1</sub> |                          | <b>Interval</b>  |  |  |  |  |
|   | 0.0833  | 0.0447         | 0.6508                   | [0.5279, 0.7545] |  |  |  |  |
|   | 0.1054  | 0.0359         | 0.7458                   | [0.6377, 0.8302] |  |  |  |  |
| 3   | 0.2134  | 0.0238         | 0.8996                   | [0.8432, 0.9373] |  |  |  |  |
|   | 0.4001  | 0.0305         | 0.9292                   | [0.8873, 0.9563] |  |  |  |  |





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### **Table 7 Results of Goodness of Fit test**



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