

# Binormal confidence intervals for AUC in R

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AUC curves are used to measure the accuracy of a classification of two groups  $X$  and  $Y$ :

$$\begin{aligned}X_1, \dots, X_{n_X} &\sim \mathcal{N}(\mu_X, \sigma_X^2) \\ Y_1, \dots, Y_{n_Y} &\sim \mathcal{N}(\mu_Y, \sigma_Y^2)\end{aligned}$$

$Y$  could be denoted as the healthy controls and  $X$  the cases with a particular disease. When having small sample size (and therefore small values in the contingency table) the confidence interval given with Wald (as in function `biostatUZH::confIntAUC`) will not perform well (fails). Hence, another way has to be found to compute the confidence interval. Pepe (2003) illustrates how AUC curves can be described using the normal distribution:

$$\begin{aligned}a &= \frac{\mu_Y - \mu_X}{\sigma_Y} \\ b &= \frac{\sigma_X}{\sigma_Y} \\ AUC &= \Phi\left(\frac{a}{\sqrt{1 + b^2}}\right)\end{aligned}$$

## Assumption: equal variances

Under the assumption that both variances  $\sigma_X$  and  $\sigma_Y$  are equal and known, the equations take a much simpler form:

$$\begin{aligned}\sigma &= \sigma_X = \sigma_Y \\ a &= \frac{\mu_Y - \mu_X}{\sigma} \\ b &= \frac{\sigma_X}{\sigma_Y} = 1 \\ AUC &= \Phi\left(\frac{a}{\sqrt{1 + b^2}}\right) = \Phi\left(\frac{a}{\sqrt{2}}\right)\end{aligned}$$

The expected value of  $X$  and  $Y$  is estimated using the average and variance:  $\hat{\mu}_X = \bar{x}$ ,  $\hat{\mu}_Y = \bar{y}$ .

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To calculate the confidence interval, the  $SE(\hat{a})$  is needed. Given that  $\hat{\mu}_X \sim \mathcal{N}(\mu_X, \sigma^2/n_X)$  and  $\hat{\mu}_Y \sim \mathcal{N}(\mu_Y, \sigma^2/n_Y)$

$$SE(\hat{a}) = SE\left(\frac{\hat{\mu}_X - \hat{\mu}_Y}{\sigma}\right) = \sqrt{\widehat{Var}\left(\frac{\hat{\mu}_X - \hat{\mu}_Y}{\sigma}\right)} = \sqrt{\widehat{Var}\left(\frac{\hat{\mu}_X}{\sigma}\right) + \widehat{Var}\left(\frac{\hat{\mu}_Y}{\sigma}\right)} = \sqrt{\frac{1}{n_X} + \frac{1}{n_Y}}.$$

The  $(1 - \alpha)$ -confidence interval for  $a$  has the following form:

$$\begin{aligned} \text{from } a_{lower} &= a - z \cdot SE(\hat{a}) \\ \text{to } a_{up} &= a + z \cdot SE(\hat{a}) \end{aligned}$$

where  $z$  refers to the  $(1-\alpha/2)$ -quantile of the standard normal distribution.

The confidence interval limits of  $AUC$  are derived by calculating the percentile of the confidence interval limits of  $\hat{a}$ :

$$\begin{aligned} \text{from } \Phi\left(\frac{a_{lower}}{\sqrt{2}}\right) \\ \text{to } \Phi\left(\frac{a_{up}}{\sqrt{2}}\right) \end{aligned}$$

### **Assumption: not equal variances**

If the assumption of equal variances would not hold, the standard error of  $AUC$  could be derived using the multivariate delta method.

## **References**

PEPE, M. S. (2003). *The statistical evaluation of medical tests for classification and prediction*, vol. 28 of *Oxford Statistical Science Series*. Oxford University Press, Oxford.