

# The probability to select the correct model using likelihood-ratio based criteria in choosing between two nested models of which the more extended one is true

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## Abstract

The probability to select the correct model is calculated for likelihood-ratio-based criteria to compare two nested models. If the more extended of the two models is true, the difference between twice the maximised log-likelihoods is approximately noncentral chi-square distributed with d.f. the difference in the number of parameters. The noncentrality parameter of this noncentral chi-square distribution can be approximated by twice the minimum Kullback–Leibler divergence (MKLD) of the best-fitting simple model to the true version of the extended model.

The MKLD, and therefore the probability to select the correct model increases approximately proportionally to the number of observations if all observations are performed under the same conditions. If a new set of observations can only be performed under different conditions, the model parameters may depend on the conditions, and therefore have to be estimated for each set of observations separately. An increase in observations will then go together with an increase in the number of model parameters. In this case, the power of the likelihood-ratio test will increase with an increasing number of observations. However, the probability to choose the correct model with the AIC will only increase if for each set of observations the MKLD is more than 0.5. If the MKLD is less than 0.5, that probability will decrease. The probability to choose the correct model with the BIC will always decrease,

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sometimes after an initial increase for a small number of observation sets. The results are illustrated by a simulation study with a set of five nested nonlinear models for binary data.

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## 1. Introduction

Several methods have been developed to determine which of two nested models,  $f_1$  and  $f_2$  with  $f_1$  nested in  $f_2$ , conform better to a set of data. Many of the general applicable methods such as likelihood-ratio (LR) testing, Akaikean information criterion (AIC) and Bayesian information criterion (BIC) all use the same test statistic

$$T = 2 \times (\ln(L_2) - \ln(L_1)), \quad (1)$$

where  $L_1$  and  $L_2$  are the maximum of the likelihood functions of  $f_1$  and  $f_2$ , respectively. If the simpler nested model  $f_1$  is true, the distribution of  $T$  converges for a large number of observations,  $n$ , to the chi-square-distribution with  $\nu$  degrees of freedom (d.f.) (see for instance, Cox and Hinkley, 1974) with  $\nu = k_2 - k_1$ , where  $k_1$  and  $k_2$  are the number of parameters of functions  $f_1$  and  $f_2$ , respectively. This holds as long as a set of not very restrictive regularity conditions are met.

For power analysis, the distribution of  $T$  should be known if the extended model ( $f_2$ ) is true. In this paper, a general solution for the distribution of  $T$  will be given. For this general solution, it is assumed that the functions  $f_1$  and  $f_2$  are locally linear in their parameters and that the deviance between the true model ( $f_2$ ) and the simpler model ( $f_1$ ) is small.

In many biological problems, the number of observations which can be obtained under constant conditions is limited. Therefore, experiments are often repeated several times within a study, each experiment having its own conditions. If models are fitted to such sets of experiments, it will often be necessary to estimate the model parameters for each experiment separately. The consequence of the distribution of  $T$  for the LR test, the AIC and the BIC are explored. The statistical results are illustrated by a simulation study on some nonlinear models for binary data.

## 2. Three methods to compare two nested models

### 2.1. The LR test

The LR test is a general hypothesis testing method to evaluate a simpler model ( $f_1$ ) against a more complex alternative ( $f_2$ ). This test uses the ratio of the maximum likelihood (ML) for the simple model and the ML for the more complex model. If  $f_1$  is true the difference between twice the log of the ML's,  $T$ , is approximately  $\chi^2$  distributed. This means that for large numbers of observations the  $\alpha$ -critical value for  $T$  is approximately  $\chi^2_{\alpha, \nu}$ . The d.f.  $\nu$  is

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