

Two-stage optimal designs in nonlinear mixed effect models: application to pharmacokinetics in children

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Abstract

Nonlinear mixed effect models (NLMEM) are used in pharmacometrics to analyse longitudinal data through models. Approaches based on the Fisher information matrix (M_F) can be used to optimise the design of these studies. A first-order linearization of the model was proposed to evaluate MF for these models [7] and is implemented in the R function PFIM [1]. Local optimal design needs some *a priori* values of the parameters which might be difficult to guess. Adaptive designs are useful to provide some exhibity and were applied in pharmacometrics [6, 9]. However, two articles in other contexts [2, 5] discussed that two-stage designs could be more efficient than fully adaptive designs. Moreover, two-stage designs are easier to implement in clinical practice. We implemented in a working version of PFIM the optimisation of the determinant of M_F for two-stage designs in NLMEM. We evaluated the approach by simulation. The example concerns a drug in development for which a pharmacokinetic study in children is needed and will be analysed through NLMEM as recommended [4, 8]. For the first stage, parameters were estimated using predictions from pharmaco-chemical properties of the drug [3]. We evaluated one and two-stage designs assuming that some parameter(s) is (are) different than the initial one(s). We evaluated the impact of the size of each cohort on the precision of population parameters estimation.

Keywords

Adaptive design, Design optimisation, Fisher information matrix, Nonlinear mixed effect models, PFIM, Population pharmacokinetics.

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