



LAVIELLE
Marc

Domaine(s) de compétence :

Statistique

Etablissement /organisme de rattachement ou d'origine :

Inria Saclay

Fonction et position :

Directeur de Recherche

Marc Lavielle is a statistician specialized in computational statistics and healthcare applications. He created and directed the Monolix team at Inria. He has worked on the development of new statistical methods, applied to a broad area of applications (geophysics, signal processing, neuro-imaging, agronomy, genetics, PKPD,...). Marc Lavielle is the author of the book Mixed Effects Models for the Population Approach: Models, Tasks, Methods and Tools (Chapman & Hall/CRC Biostatistics Series, 2014). He has also co-authored several papers on various aspects of statistical estimation and on SAEM algorithm. Several of the methods that he developed are implemented in the Monolix software. He recently developed Simulx and the mlxR package for clinical trial simulations.

Marc Lavielle holds a Ph.D. in applied mathematics from Université Paris-XI, Orsay, France (1991). He was named Assistant Professor in 1991 and Professor in 1998 at Paris Descartes University, and joined Inria in 2007 as Research Director.

Publications marquantes

Lavielle M. "Mixed Effects Models for the Population Approach. Models, Tasks, Methods & Tools. ", Chapman & Hall/CRC Biostatistics Series, 2014

Mbogning C., Bleakley K., Lavielle M. "Joint modeling of longitudinal and repeated time-to-event data with maximum likelihood estimation via the SAEM algorithm", Journal of Statistical Computation and Simulation, vol. 85, n. 8, 1512--1528, 2015

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Lavielle M. "Rôle et limites de la statistique dans l'évaluation des risques sanitaires liés aux OGM", Statistique et Société, vol. 1, 2013

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Dubois A., Lavielle M., Gsteiger S., Pigeolet E., Mentré F. "Model-Based Analyses of Bioequivalence Crossover Trials Using the SAEM Algorithm", Statistics in Medicine, vol. 30, pp. 2582-600, 2011

Lavielle M., Samson A., Fermin A.K., Mentré F. "Maximum likelihood estimation of long term HIV dynamic models and antiviral response", *Biometrics*, vol. 67, pp. 250-259, 2011.

Chan P., Jacqmin P., Lavielle M., McFadyen L., Weatherley B. "The Use of the SAEM Algorithm in MONOLIX Software for Estimation of Population Pharmacokinetic-Pharmacodynamic-Viral Dynamics Parameters of Maraviroc in Asymptomatic HIV Subjects" *Journal of Pharmacokinetics and Pharmacodynamics*, vol. 38, pp. 41-61, 2011

Snoeck E., Chanu P., Lavielle M., Jacqmin P., Jonsson N., Jorga K., Goggin T., Jumbe S. , Frey N. "Hepatitis C Viral Dynamics Explaining Breakthrough, Relapse or Response after Chronic Treatment", *Clinical Pharmacology and Therapeutics*, Vol 87 (6), pp 706-713, 2010

Perry JN, Ter Braak CJ, Dixon PM, Duan JJ, Hails RS, Huesken A, Lavielle M, Marvier M, Scardi M, Schmidt K, Tothmeresz B, Schaarschmidt F, van der Voet H. "Statistical aspects of environmental risk assessment of GM plants for effects on non-target organisms" *Environ Biosafety Res.*, vol. 8 pp 65-78, 2009