France Mentré

List of Publications by Year in descending order

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91 papers 6,618 citations

94433 37 h-index 69250 77 g-index

98 all docs 98 docs citations 98 times ranked 9963 citing authors

#	Article	IF	CITATIONS
1	Clinical and virological data of the first cases of COVID-19 in Europe: a case series. Lancet Infectious Diseases, The, 2020, 20, 697-706.	9.1	953
2	Type 1 interferons as a potential treatment against COVID-19. Antiviral Research, 2020, 178, 104791.	4.1	425
3	Experimental Treatment with Favipiravir for Ebola Virus Disease (the JIKI Trial): A Historically Controlled, Single-Arm Proof-of-Concept Trial in Guinea. PLoS Medicine, 2016, 13, e1001967.	8.4	382
4	Computing normalised prediction distribution errors to evaluate nonlinear mixed-effect models: The npde add-on package for R. Computer Methods and Programs in Biomedicine, 2008, 90, 154-166.	4.7	370
5	Metrics for External Model Evaluation with an Application to the Population Pharmacokinetics of Gliclazide. Pharmaceutical Research, 2006, 23, 2036-2049.	3.5	268
6	Model Evaluation of Continuous Data Pharmacometric Models: Metrics and Graphics. CPT: Pharmacometrics and Systems Pharmacology, 2017, 6, 87-109.	2.5	261
7	Remdesivir plus standard of care versus standard of care alone for the treatment of patients admitted to hospital with COVID-19 (DisCoVeRy): a phase 3, randomised, controlled, open-label trial. Lancet Infectious Diseases, The, 2022, 22, 209-221.	9.1	233
8	Modeling SARS-CoV-2 viral kinetics and association with mortality in hospitalized patients from the French COVID cohort. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	181
9	Timing of Antiviral Treatment Initiation is Critical to Reduce SARS oVâ€2 Viral Load. CPT: Pharmacometrics and Systems Pharmacology, 2020, 9, 509-514.	2.5	170
10	Favipiravir pharmacokinetics in Ebola-Infected patients of the JIKI trial reveals concentrations lower than targeted. PLoS Neglected Tropical Diseases, 2017, 11, e0005389.	3.0	153
11	Are Population Pharmacokinetic and/or Pharmacodynamic Models Adequately Evaluated?. Clinical Pharmacokinetics, 2007, 46, 221-234.	3.5	149
12	Intracellular Pharmacokinetics of Antiretroviral Drugs in HIV-Infected Patients, and their Correlation with Drug Action. Clinical Pharmacokinetics, 2010, 49, 17-45.	3.5	140
13	Estimation of Population Pharmacokinetic Parameters of Saquinavir in HIV Patients with the MONOLIX Software. Journal of Pharmacokinetics and Pharmacodynamics, 2007, 34, 229-249.	1.8	135
14	Ebola Virus Infection: Review of the Pharmacokinetic and Pharmacodynamic Properties of Drugs Considered for Testing in Human Efficacy Trials. Clinical Pharmacokinetics, 2016, 55, 907-923.	3.5	135
15	Extension of the SAEM algorithm to left-censored data in nonlinear mixed-effects model: Application to HIV dynamics model. Computational Statistics and Data Analysis, 2006, 51, 1562-1574.	1.2	126
16	Protection of the Human Gut Microbiome From Antibiotics. Journal of Infectious Diseases, 2018, 217, 628-636.	4.0	124
17	Development and implementation of the population Fisher information matrix for the evaluation of population pharmacokinetic designs. Computer Methods and Programs in Biomedicine, 2001, 65, 141-151.	4.7	122
18	Antiviral efficacy of favipiravir against Ebola virus: A translational study in cynomolgus macaques. PLoS Medicine, 2018, 15, e1002535.	8.4	108

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19	Ciprofloxacin Dosage and Emergence of Resistance in Human Commensal Bacteria. Journal of Infectious Diseases, 2009, 200, 390-398.	4.0	105
20	Prediction Discrepancies for the Evaluation of Nonlinear Mixed-Effects Models. Journal of Pharmacokinetics and Pharmacodynamics, 2006, 33, 345-367.	1.8	94
21	Persistent COVID-19 symptoms are highly prevalent 6Âmonths after hospitalization: results from a large prospective cohort. Clinical Microbiology and Infection, 2021, 27, 1041.e1-1041.e4.	6.0	88
22	Further Developments of the Fisher Information Matrix in Nonlinear Mixed Effects Models with Evaluation in Population Pharmacokinetics. Journal of Biopharmaceutical Statistics, 2003, 13, 209-227.	0.8	87
23	Non-Linear Mixed Effects Modeling – From Methodology and Software Development to Driving Implementation in Drug Development Science. Journal of Pharmacokinetics and Pharmacodynamics, 2005, 32, 161-183.	1.8	87
24	Dose regimen of favipiravir for Ebola virus disease. Lancet Infectious Diseases, The, 2015, 15, 150-151.	9.1	86
25	Evaluation of different tests based on observations for external model evaluation of population analyses. Journal of Pharmacokinetics and Pharmacodynamics, 2010, 37, 49-65.	1.8	72
26	Design evaluation and optimisation in multiple response nonlinear mixed effect models: PFIM 3.0. Computer Methods and Programs in Biomedicine, 2010, 98, 55-65.	4.7	68
27	Fisher information matrix for non-linear mixed-effects models: evaluation and application for optimal design of enoxaparin population pharmacokinetics. Statistics in Medicine, 2002, 21, 2623-2639.	1.6	65
28	Methods and software tools for design evaluation in population pharmacokinetics–pharmacodynamics studies. British Journal of Clinical Pharmacology, 2015, 79, 6-17.	2.4	65
29	Favipiravir Pharmacokinetics in Nonhuman Primates and Insights for Future Efficacy Studies of Hemorrhagic Fever Viruses. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	59
30	Ebola virus dynamics in mice treated with favipiravir. Antiviral Research, 2015, 123, 70-77.	4.1	57
31	Design in nonlinear mixed effects models: Optimization using the Fedorov–Wynn algorithm and power of the Wald test for binary covariates. Statistics in Medicine, 2007, 26, 5162-5179.	1.6	55
32	Ebola viral dynamics in nonhuman primates provides insights into virus immuno-pathogenesis and antiviral strategies. Nature Communications, 2018, 9, 4013.	12.8	54
33	Evaluation of bootstrap methods for estimating uncertainty of parameters in nonlinear mixed-effects models: a simulation study in population pharmacokinetics. Journal of Pharmacokinetics and Pharmacodynamics, 2014, 41, 15-33.	1.8	53
34	The use of simulated annealing for finding optimal population designs. Computer Methods and Programs in Biomedicine, 2002, 69, 25-35.	4.7	49
35	Favipiravir for children with Ebola. Lancet, The, 2015, 385, 603-604.	13.7	43
36	Performance Comparison of Various Maximum Likelihood Nonlinear Mixed-Effects Estimation Methods for Dose–Response Models. AAPS Journal, 2012, 14, 420-432.	4.4	42

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37	Ceftriaxone and Cefotaxime Have Similar Effects on the Intestinal Microbiota in Human Volunteers Treated by Standard-Dose Regimens. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	41
38	SARS-CoV-2 viral dynamics in non-human primates. PLoS Computational Biology, 2021, 17, e1008785.	3.2	41
39	Global outbreak research: harmony not hegemony. Lancet Infectious Diseases, The, 2020, 20, 770-772.	9.1	40
40	Maximum Likelihood Estimation of Long-Term HIV Dynamic Models and Antiviral Response. Biometrics, 2011, 67, 250-259.	1.4	39
41	Nonlinear Mixed-Effect Models for Prostate-Specific Antigen Kinetics and Link with Survival in the Context of Metastatic Prostate Cancer: a Comparison by Simulation of Two-Stage and Joint Approaches. AAPS Journal, 2015, 17, 691-699.	4.4	38
42	Impact of Antibiotic Gut Exposure on the Temporal Changes in Microbiome Diversity. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	35
43	Impact on disease mortality of clinical, biological, and virological characteristics at hospital admission and overtime in COVIDâ€19 patients. Journal of Medical Virology, 2021, 93, 2149-2159.	5.0	35
44	Population pharmacokinetics of imipenem in critically ill patients with suspected ventilatorâ€associated pneumonia and evaluation of dosage regimens. British Journal of Clinical Pharmacology, 2014, 78, 1022-1034.	2.4	34
45	Dose Rationale for Favipiravir Use in Patients Infected With SARSâ€CoVâ€2. Clinical Pharmacology and Therapeutics, 2020, 108, 188-188.	4.7	34
46	Mathematical Modeling of Bacterial Kinetics to Predict the Impact of Antibiotic Colonic Exposure and Treatment Duration on the Amount of Resistant Enterobacteria Excreted. PLoS Computational Biology, 2014, 10, e1003840.	3.2	32
47	Population pharmacokinetic analysis of mizolastine and validation from sparse data on patients using the nonparametric maximum likelihood method. Journal of Pharmacokinetics and Pharmacodynamics, 1998, 26, 133-161.	0.6	28
48	Fisher information matrix for nonlinear mixed effects multiple response models: Evaluation of the appropriateness of the first order linearization using a pharmacokinetic/pharmacodynamic model. Statistics in Medicine, 2009, 28, 1940-1956.	1.6	28
49	Nonlinear joint models for individual dynamic prediction of risk of death using Hamiltonian Monte Carlo: application to metastatic prostate cancer. BMC Medical Research Methodology, 2017, 17, 105.	3.1	28
50	Association Between Tumor Size Kinetics and Survival in Patients With Urothelial Carcinoma Treated With Atezolizumab: Implication for Patient Followâ€Up. Clinical Pharmacology and Therapeutics, 2019, 106, 810-820.	4.7	27
51	PFIM 4.0, an extended R program for design evaluation and optimization in nonlinear mixed-effect models. Computer Methods and Programs in Biomedicine, 2018, 156, 217-229.	4.7	25
52	Effect of remdesivir on viral dynamics in COVID-19 hospitalized patients: a modelling analysis of the randomized, controlled, open-label DisCoVeRy trial. Journal of Antimicrobial Chemotherapy, 2022, 77, 1404-1412.	3.0	25
53	Implementation and Evaluation of the SAEM Algorithm for Longitudinal Ordered Categorical Data with an Illustration in Pharmacokinetics–Pharmacodynamics. AAPS Journal, 2011, 13, 44-53.	4.4	24
54	Extension of NPDE for evaluation of nonlinear mixed effect models in presence of data below the quantification limit with applications to HIV dynamic model. Journal of Pharmacokinetics and Pharmacodynamics, 2012, 39, 499-518.	1.8	23

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55	Lessons learned from IDeAl $\hat{a}\in$ " 33 recommendations from the IDeAl-net about design and analysis of small population clinical trials. Orphanet Journal of Rare Diseases, 2018, 13, 77.	2.7	22
56	Modeling Favipiravir Antiviral Efficacy Against Emerging Viruses: From Animal Studies to Clinical Trials. CPT: Pharmacometrics and Systems Pharmacology, 2020, 9, 258-271.	2.5	20
57	Impact of imiglucerase on the serum glycosylated-ferritin level in Gaucher disease. Blood Cells, Molecules, and Diseases, 2011, 46, 34-38.	1.4	19
58	Protection of Hamsters from Mortality by Reducing Fecal Moxifloxacin Concentration with DAV131A in a Model of Moxifloxacin-Induced Clostridium difficile Colitis. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	19
59	Design evaluation and optimisation in crossover pharmacokinetic studies analysed by nonlinear mixed effects models. Statistics in Medicine, 2012, 31, 1043-1058.	1.6	18
60	Antibiotic-Induced Dysbiosis Predicts Mortality in an Animal Model of Clostridium difficile Infection. Antimicrobial Agents and Chemotherapy, 2018, 62, .	3.2	18
61	Pharmacometrics and Systems Pharmacology 2030. Clinical Pharmacology and Therapeutics, 2020, 107, 76-78.	4.7	18
62	Once-Daily Dosing of Saquinavir Soft-Gel Capsules and Ritonavir Combination in HIV-1-Infected Patients (Imea015 Study). Antiviral Therapy, 2004, 9, 247-256.	1.0	15
63	Design evaluation and optimization for models of hepatitis C viral dynamics. Statistics in Medicine, 2011, 30, 1045-1056.	1.6	14
64	An MCMC method for the evaluation of the Fisher information matrix for non-linear mixed effect models. Biostatistics, 2016, 17, 737-750.	1.5	13
65	Implementation of a non-human primate model of Ebola disease: Infection of Mauritian cynomolgus macaques and analysis of virus populations. Antiviral Research, 2017, 140, 95-105.	4.1	13
66	The safety profile of favipiravir should not be the first argument to suspend its evaluation in viral hemorrhagic fevers. PLoS Neglected Tropical Diseases, 2020, 14, e0008259.	3.0	13
67	A new method for evaluation of the Fisher information matrix for discrete mixed effect models using Monte Carlo sampling and adaptive Gaussian quadrature. Computational Statistics and Data Analysis, 2017, 111, 203-219.	1.2	12
68	Model Averaging in Viral Dynamic Models. AAPS Journal, 2020, 22, 48.	4.4	12
69	Evaluation of the Fisher information matrix in nonlinear mixed effect models using adaptive Gaussian quadrature. Computational Statistics and Data Analysis, 2014, 80, 57-69.	1.2	11
70	Modeling the Effect of DAV132, a Novel Colon†Targeted Adsorbent, on Fecal Concentrations of Moxifloxacin and Gut Microbiota Diversity in Healthy Volunteers. Clinical Pharmacology and Therapeutics, 2021, 109, 1045-1054.	4.7	11
71	Influence of the Size of Cohorts in Adaptive Design for Nonlinear Mixed Effects Models: An Evaluation by Simulation for a Pharmacokinetic and Pharmacodynamic Model for a Biomarker in Oncology. Pharmaceutical Research, 2015, 32, 3159-3169.	3 . 5	10
72	Ribavirin does not potentiate favipiravir antiviral activity against Ebola virus in non-human primates. Antiviral Research, 2020, 177, 104758.	4.1	10

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73	Optimal Design for Informative Protocols in Xenograft Tumor Growth Inhibition Experiments in Mice. AAPS Journal, 2016, 18, 1233-1243.	4.4	9
74	Influence of Covariance Between Random Effects in Design for Nonlinear Mixed-Effect Models with an Illustration in Pediatric Pharmacokinetics. Journal of Biopharmaceutical Statistics, 2014, 24, 471-492.	0.8	7
75	CPT: Pharmacometrics & Dystems Pharmacology â€" Inception, Maturation, and Future Vision. CPT: Pharmacometrics and Systems Pharmacology, 2021, 10, 649-657.	2.5	6
76	Bayesian Individual Dynamic Predictions with Uncertainty of Longitudinal Biomarkers and Risks of Survival Events in a Joint Modelling Framework: a Comparison Between Stan, Monolix, and NONMEM. AAPS Journal, 2020, 22, 50.	4.4	5
77	Once-daily dosing of saquinavir soft-gel capsules and ritonavir combination in HIV-1-infected patients (IMEA015 study). Antiviral Therapy, 2004, 9, 247-56.	1.0	5
78	Robust designs in longitudinal studies accounting for parameter and model uncertainties – application to count data. Journal of Biopharmaceutical Statistics, 2020, 30, 31-45.	0.8	4
79	Robust designs accounting for model uncertainty in longitudinal studies with binary outcomes. Statistical Methods in Medical Research, 2020, 29, 934-952.	1.5	4
80	Tutorial for \$DESIGN in NONMEM: Clinical Trial Evaluation and Optimization. CPT: Pharmacometrics and Systems Pharmacology, 2021, , .	2.5	4
81	Modeling the bacterial dynamics in the gut microbiota following an antibioticâ€induced perturbation. CPT: Pharmacometrics and Systems Pharmacology, 2022, 11, 906-918.	2.5	4
82	Individual Bayesian Information Matrix for Predicting Estimation Error and Shrinkage of Individual Parameters Accounting for Data Below the Limit of Quantification. Pharmaceutical Research, 2017, 34, 2119-2130.	3.5	3
83	Developing Tools to Evaluate Non-linear Mixed Effect Models: 20ÂYears on the npde Adventure. AAPS Journal, 2021, 23, 75.	4.4	3
84	Finding optimal design in nonlinear mixed effect models using multiplicative algorithms. Computer Methods and Programs in Biomedicine, 2021, 207, 106126.	4.7	3
85	A Colonâ€Targeted Adsorbent (DAV132) Does Not Affect the Pharmacokinetics of Warfarin or Clonazepam in Healthy Subjects. Clinical Pharmacology in Drug Development, 2021, 10, 908-917.	1.6	2
86	High doses of favipiravir in two men survivors of Ebola virus disease carrying Ebola virus in semen in Guinea. IDCases, 2022, 27, e01412.	0.9	2
87	Comment on Jaki et al., A proposal for a new PhD level curriculum on quantitative methods for drug development <i>>. Pharmaceutical Statistics</i> > 17 (5):593–606, Sep/Oct 2018, DOI: 10.1002/pst.1873. Pharmaceutical Statistics, 2019, 18, 278-281.	1.3	1
88	Comparison of Various Phase I Combination Therapy Designs in Oncology for Evaluation of Early Tumor Shrinkage Using Simulations. CPT: Pharmacometrics and Systems Pharmacology, 2020, 9, 686-694.	2.5	1
89	Welcome to the statistics and pharmacometrics themed issue. CPT: Pharmacometrics and Systems Pharmacology, 2021, 10, 273-274.	2.5	1
90	Impact of study design and statistical model in pharmacogenetic studies with geneâ€treatment interaction. CPT: Pharmacometrics and Systems Pharmacology, 2021, 10, 340-349.	2.5	1

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91	<i>CPT: Pharmacometrics & Systems Pharmacology (i) 2.0. CPT: Pharmacometrics and Systems Pharmacology, 2019, 8, 195-196.</i>	2.5	0