

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 8, 2026

Simulations Plus

RRID:SCR_003946

Type: Tool

Proper Citation

Simulations Plus (RRID:SCR_003946)

Resource Information

URL: <http://www.simulations-plus.com/>

Proper Citation: Simulations Plus (RRID:SCR_003946)

Description: Commercial developer of Absorption, Distribution, Metabolism, Excretion and Toxicity (ADMET) and Physiologically-Based Pharmacokinetic (PBPK) modeling and simulation software for the pharmaceutical and biotechnology, industrial chemical, cosmetic, herbicide, and food ingredient industries. Their software allows pharmaceutical scientists to predict certain key potential endpoints and dynamics, in silico, thereby reducing research & development costs and helping clients make better projects decisions sooner. Consulting services: Their team of highly-skilled scientists is available to assist with your discovery and development activities on a project-by-project basis. They now offer expanded pharmacodynamic modeling services, NONMEM analysis, and regulatory report writing assistance. Their ultimate goal is to develop tools and systems which bridge PBPK modeling with clinical trial data analysis. Training workshops & User Groups: they offer hands-on introductory and advanced training workshops in the United States, Europe, and Asia throughout the year.

Synonyms: Simulations Plus Inc.

Resource Type: commercial organization

Keywords: pharmaceutical, modelling, simulation, pharmacodynamic, biotechnology, chemical, cosmetic, food ingredient, herbicide, drug development, drug, pharmacokinetic

Availability: License required

Resource Name: Simulations Plus

Resource ID: SCR_003946

Alternate IDs: nlx_158339, ISNI: 0000 0004 0506 5380, grid.418738.1, Wikidata: Q7521341

Alternate URLs: <https://ror.org/02p0yhm49>

Record Creation Time: 20220129T080221+0000

Record Last Update: 20260905T132511+0000

Ratings and Alerts

No rating or validation information has been found for Simulations Plus.

No alerts have been found for Simulations Plus.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 37 mentions in open access literature.

Listed below are recent publications. The full list is available at [PRECISE-TBI](#) .

Gajewska M, et al. (2026) A comprehensive evaluation of the human Tg32 and Tg276 FcRn transgenic mouse and non-human primate models for predicting antibody pharmacokinetics in human. *mAbs*, 18(1), 2668888.

Park H, et al. (2026) Stratification of viral shedding patterns in saliva of COVID-19 patients. *eLife*, 13.

Najjar A, et al. (2025) Grouping of chemicals for safety assessment: the importance of toxicokinetic properties of salicylate esters. *Archives of toxicology*, 99(3), 995.

Pansari A, et al. (2024) A tutorial on physiologically based pharmacokinetic approaches in lactation research. *CPT: pharmacometrics & systems pharmacology*, 13(11), 1841.

Chaturvedi K, et al. (2023) Identification of the Putative Binding Site of a Benzimidazole Opioid (Etazene) and Its Metabolites at μ -Opioid Receptor: A Human Liver Microsomal Assay and Systematic Computational Study. *Molecules (Basel, Switzerland)*, 28(4).

Komura H, et al. (2023) The Trends and Future Prospective of In Silico Models from the Viewpoint of ADME Evaluation in Drug Discovery. *Pharmaceutics*, 15(11).

Yuan Y, et al. (2023) Pharmacokinetics of Novel Furoxan/Coumarin Hybrids in Rats Using LC-MS/MS Method and Physiologically Based Pharmacokinetic Model. *Molecules (Basel, Switzerland)*, 28(2).

Scior T, et al. (2023) Targeting the Human Influenza a Virus: The Methods, Limitations, and Pitfalls of Virtual Screening for Drug-like Candidates Including Scaffold Hopping and Compound Profiling. *Viruses*, 15(5).

Punt A, et al. (2022) Predictive Performance of Next Generation Physiologically Based Kinetic (PBK) Model Predictions in Rats Based on In Vitro and In Silico Input Data. *Toxicological sciences : an official journal of the Society of Toxicology*, 186(1), 18.

Lai Y, et al. (2022) Recent advances in the translation of drug metabolism and pharmacokinetics science for drug discovery and development. *Acta pharmaceutica Sinica. B*, 12(6), 2751.

Frechen S, et al. (2022) Quality Assurance of PBPK Modeling Platforms and Guidance on Building, Evaluating, Verifying and Applying PBPK Models Prudently under the Umbrella of Qualification: Why, When, What, How and By Whom? *Pharmaceutical research*, 39(8), 1733.

Koczurkiewicz-Adamczyk P, et al. (2022) Cinnamamide derivatives with 4-hydroxypiperidine moiety enhance effect of doxorubicin to cancer cells and protect cardiomyocytes against drug-induced toxicity through CBR1 inhibition mechanism. *Life sciences*, 305, 120777.

Khalidi H, et al. (2022) SimRFlow: An R-based workflow for automated high-throughput PBPK simulation with the Simcyp® simulator. *Frontiers in pharmacology*, 13, 929200.

Ooka M, et al. (2022) Identification of environmental chemicals that activate p53 signaling after in vitro metabolic activation. *Archives of toxicology*, 96(7), 1975.

Fatima K, et al. (2021) Neomenthol prevents the proliferation of skin cancer cells by restraining tubulin polymerization and hyaluronidase activity. *Journal of advanced research*, 34, 93.

Krzywik J, et al. (2021) An insight into the anticancer potential of carbamates and thiocarbamates of 10-demethoxy-10-methylaminocolchicine. *European journal of medicinal chemistry*, 215, 113282.

Alagumuthu M, et al. (2021) Structure-Based Design of Novel Peptidomimetics Targeting the SARS-CoV-2 Spike Protein. *Cellular and molecular bioengineering*, 14(2), 177.

Acquah FA, et al. (2021) Simulations of Promising Indolizidine- α -Nicotinic Acetylcholine Receptor Complexes. *International journal of molecular sciences*, 22(15).

Djokovic N, et al. (2021) An Integrative in silico Drug Repurposing Approach for Identification of Potential Inhibitors of SARS-CoV-2 Main Protease. *Molecular informatics*, 40(5), e2000187.

Rácz A, et al. (2021) Machine learning models for classification tasks related to drug safety. *Molecular diversity*, 25(3), 1409.