## C Anthony Hunt

List of Publications by Year in descending order

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|          |                | 279798       | 345221         |
|----------|----------------|--------------|----------------|
| 78       | 1,553          | 23           | 36             |
| papers   | citations      | h-index      | g-index        |
|          |                |              |                |
|          |                |              |                |
| 93       | 93             | 02           | 1216           |
| 93       | 93             | 93           | 1216           |
| all docs | docs citations | times ranked | citing authors |
|          |                |              |                |

| #  | Article   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | Formation of Cysts by Alveolar Type II Cells in Three-dimensional Culture Reveals a Novel Mechanism for Epithelial Morphogenesis. Molecular Biology of the Cell, 2007, 18, 1693-1700.   | 2.1  | 91        |
| 2  | Engineering targeted in vivo drug delivery. I. The physiological and physicochemical principles governing opportunities and limitations. Pharmaceutical Research, 1986, 03, 333-344.  | 3.5  | 83        |
| 3  | At the Biological Modeling and Simulation Frontier. Pharmaceutical Research, 2009, 26, 2369-2400.   | 3.5  | 73        |
| 4  | Antisense c-myc oligodeoxyribonucleotide cellular uptake. Pharmaceutical Research, 1992, 09, 1010-1017.   | 3.5  | 59        |
| 5  | Physiologically Based Synthetic Models of Hepatic Disposition. Journal of Pharmacokinetics and Pharmacodynamics, 2006, 33, 737-772.   | 1.8  | 58        |
| 6  | Simulating Properties of In Vitro Epithelial Cell Morphogenesis. PLoS Computational Biology, 2006, 2, e129.   | 3.2  | 58        |
| 7  | Tuneable resolution as a systems biology approach for multiâ€scale, multiâ€compartment computational models. Wiley Interdisciplinary Reviews: Systems Biology and Medicine, 2014, 6, 289-309.                                     | 6.6  | 53        |
| 8  | LIGHT SENSITIVE LIPOSOMES. Photochemistry and Photobiology, 1983, 37, 491-494.  | 2.5  | 49        |
| 9  | Modeling and Simulation of Hepatic Drug Disposition Using a Physiologically Based, Multi-agent In Silico Liver. Pharmaceutical Research, 2008, 25, 1023-1036.   | 3.5  | 46        |
| 10 | Dynamics of in silico leukocyte rolling, activation, and adhesion. BMC Systems Biology, 2007, 1, 14.  | 3.0  | 45        |
| 11 | Challenges and rewards on the road to translational systems biology in acute illness: four case reports from interdisciplinary teams. Journal of Critical Care, 2007, 22, 169-175.  | 2.2  | 44        |
| 12 | Evidence that cannabidiol does not significantly alter the pharmacokinetics of tetrahydrocannabinol in man. Journal of Pharmacokinetics and Pharmacodynamics, 1981, 9, 245-260.   | 0.6  | 40        |
| 13 | Lymphatic transport of liposome-encapsulated drugs following intraperitoneal administration - effect of lipid composition. Pharmaceutical Research, 1985, 02, 271-278.  | 3.5  | 40        |
| 14 | Buffer effects on swelling kinetics in polybasic gels. Pharmaceutical Research, 1992, 09, 76-81.  | 3.5  | 40        |
| 15 | Predictions of Hepatic Disposition Properties Using a Mechanistically Realistic, Physiologically Based<br>Model. Drug Metabolism and Disposition, 2008, 36, 759-768.  | 3.3  | 40        |
| 16 | Essential operating principles for tumor spheroid growth. BMC Systems Biology, 2008, 2, 110.  | 3.0  | 39        |
| 17 | Dichotomies between computational and mathematical models. Nature Biotechnology, 2008, 26, 737-738.   | 17.5 | 36        |
| 18 | Agentâ€based modeling: a systematic assessment of use cases and requirements for enhancing pharmaceutical research and development productivity. Wiley Interdisciplinary Reviews: Systems Biology and Medicine, 2013, 5, 461-480. | 6.6  | 33        |

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|----|---|--------------|-----------|
| 19 | MDCK Cystogenesis Driven by Cell Stabilization within Computational Analogues. PLoS Computational Biology, 2011, 7, e1002030.   | 3.2          | 32        |
| 20 | Tracing Multiscale Mechanisms of Drug Disposition in Normal and Diseased Livers. Journal of Pharmacology and Experimental Therapeutics, 2010, 334, 124-136.   | 2.5          | 29        |
| 21 | Mechanistic Study of the Cellular Interplay of Transport and Metabolism Using the Synthetic Modeling Method. Pharmaceutical Research, 2006, 23, 493-505.  | 3 <b>.</b> 5 | 28        |
| 22 | Identifying the Rules of Engagement Enabling Leukocyte Rolling, Activation, and Adhesion. PLoS Computational Biology, 2010, 6, e1000681.  | 3.2          | 27        |
| 23 | A computational approach to resolve cell level contributions to early glandular epithelial cancer progression. BMC Systems Biology, 2009, 3, 122.   | 3.0          | 25        |
| 24 | Computational Strategies Unravel and Trace How Liver Disease Changes Hepatic Drug Disposition. Journal of Pharmacology and Experimental Therapeutics, 2009, 328, 294-305.                                   | 2.5          | 24        |
| 25 | Relational grounding facilitates development of scientifically useful multiscale models. Theoretical Biology and Medical Modelling, 2011, 8, 35.  | 2.1          | 24        |
| 26 | Mechanistic Insight from In Silico Pharmacokinetic Experiments: Roles of P-glycoprotein, Cyp3A4 Enzymes, and Microenvironments. Journal of Pharmacology and Experimental Therapeutics, 2010, 332, 398-412.  | 2.5          | 22        |
| 27 | Toward modular biological models: defining analog modules based on referent physiological mechanisms. BMC Systems Biology, 2014, 8, 95.   | 3.0          | 22        |
| 28 | Competing Mechanistic Hypotheses of Acetaminophen-Induced Hepatotoxicity Challenged by Virtual Experiments. PLoS Computational Biology, 2016, 12, e1005253.   | 3.2          | 22        |
| 29 | Studies of intestinal drug transport using an in silico epithelio-mimetic device. BioSystems, 2005, 82, 154-167.  | 2.0          | 20        |
| 30 | Discovering Plausible Mechanistic Details of Hepatic Drug Interactions. Drug Metabolism and Disposition, 2009, 37, 237-246.   | 3.3          | 20        |
| 31 | Cloud computing and validation of expandable in silico livers. BMC Systems Biology, 2010, 4, 168.   | 3.0          | 19        |
| 32 | The Spectrum of Mechanism-Oriented Models and Methods for Explanations of Biological Phenomena. Processes, 2018, 6, 56.   | 2.8          | 19        |
| 33 | An In Silico Transwell Device for the Study of Drug Transport and Drug–Drug Interactions.<br>Pharmaceutical Research, 2007, 24, 2171-2186.  | 3.5          | 18        |
| 34 | Computational investigation of epithelial cell dynamic phenotype in vitro. Theoretical Biology and Medical Modelling, 2009, 6, 8.   | 2.1          | 17        |
| 35 | Computational experiments reveal plausible mechanisms for changing patterns of hepatic zonation of xenobiotic clearance and hepatotoxicity. Journal of Theoretical Biology, 2010, 265, 718-733.             | 1.7          | 17        |
| 36 | Contrasting model mechanisms of alanine aminotransferase (ALT) release from damaged and necrotic hepatocytes as an example of general biomarker mechanisms. PLoS Computational Biology, 2020, 16, e1007622. | 3.2          | 17        |

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|----|---|--------------|-----------|
| 37 | Liposome Dialysis for Improved Size Distributions. Journal of Pharmaceutical Sciences, 1982, 71, 806-812.   | 3.3          | 16        |
| 38 | A Computational Approach to Understand In Vitro Alveolar Morphogenesis. PLoS ONE, 2009, 4, e4819.   | 2.5          | 15        |
| 39 | In Silico Methods for Unraveling the Mechanistic Complexities of Intestinal Absorption:<br>Metabolism-Efflux Transport Interactions. Drug Metabolism and Disposition, 2008, 36, 1414-1424.            | 3.3          | 14        |
| 40 | Virtual Experiments Enable Exploring and Challenging Explanatory Mechanisms of Immune-Mediated P450 Down-Regulation. PLoS ONE, 2016, 11, e0155855.  | 2.5          | 14        |
| 41 | Synthesis of DNA Dumbbells: Chemical vs. Enzymatic Ligation of Self-Complementary Oligonucleotides. Nucleosides & Nucleotides, 1997, 16, 41-51.   | 0.5          | 12        |
| 42 | Bootstrapping for pharmacokinetic models: visualization of predictive and parameter uncertainty. Pharmaceutical Research, 1998, 15, 690-697.  | 3 <b>.</b> 5 | 12        |
| 43 | Enabling Clearance Predictions to Emerge from In Silico Actions of Quasi-Autonomous Hepatocyte Components. Drug Metabolism and Disposition, 2011, 39, 1910-1920.                                      | 3.3          | 12        |
| 44 | Propagation of Pericentral Necrosis During Acetaminophen-Induced Liver Injury: Evidence for Early Interhepatocyte Communication and Information Exchange. Toxicological Sciences, 2019, 169, 151-166. | 3.1          | 12        |
| 45 | Murine plasma fibronectin depletion after intravenous injection of liposomes. International Journal of Pharmaceutics, 1987, 37, 233-238.  | <b>5.</b> 2  | 11        |
| 46 | Prediction of in Vitro Hepatic Biliary Excretion using Stochastic Agent-Based Modeling and Fuzzy Clustering., 2006,,.   |              | 10        |
| 47 | Simulation of lung alveolar epithelial wound healing in vitro. Journal of the Royal Society Interface, 2010, 7, 1157-1170.  | 3.4          | 8         |
| 48 | Moving beyond in silico tools to in silico science in support of drug development research. Drug Development Research, 2011, 72, 153-161.   | 2.9          | 8         |
| 49 | A Model Mechanism-Based Explanation of an In Vitro-In Vivo Disconnect for Improving Extrapolation and Translation. Journal of Pharmacology and Experimental Therapeutics, 2018, 365, 127-138.         | 2.5          | 8         |
| 50 | A cell-centered, agent-based framework that enables flexible environment granularities. Theoretical Biology and Medical Modelling, 2016, 13, 4.   | 2.1          | 7         |
| 51 | Individualized, discrete event, simulations provide insight into inter- and intra-subject variability of extended-release, drug products. Theoretical Biology and Medical Modelling, 2012, 9, 39.     | 2.1          | 6         |
| 52 | In Silico, Experimental, Mechanistic Model for Extended-Release Felodipine Disposition Exhibiting Complex Absorption and a Highly Variable Food Interaction. PLoS ONE, 2014, 9, e108392.              | 2.5          | 6         |
| 53 | Artificial Red Cells. A Link Between the Membrane Skeleton and Res Detectability?. Biomaterials, Artificial Cells, and Artificial Organs, 1990, 18, 329-343.  | 0.2          | 5         |
| 54 | Evaluating an hepatic enzyme induction mechanism through coarse―and fineâ€grained measurements of an in silico liver. Complexity, 2009, 14, 28-34.  | 1.6          | 5         |

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|----|---|-----|-----------|
| 55 | Falsifying an Enzyme Induction Mechanism within a Validated, Multiscale Liver Model. International Journal of Agent Technologies and Systems, 2012, 4, 1-14.                                  | 0.1 | 5         |
| 56 | Agent-Directed Tracing of Multi-Scale Drug Disposition Events within Normal and Diseased In Silico Livers. International Journal of Agent Technologies and Systems, 2010, 2, 1-19.            | 0.1 | 4         |
| 57 | Cost-based Partitioning for Distributed and Parallel Simulation of Decomposable Multiscale Constructive Models. Simulation, 2006, 82, 809-826.  | 1.8 | 3         |
| 58 | Mechanistic simulations explain paradoxical saquinavir metabolism during in vitro vectorial transport study., 2008, 2008, 5462-5.   |     | 3         |
| 59 | Simulation enabled search for explanatory mechanisms of the fracture healing process. PLoS Computational Biology, 2018, 14, e1005980.   | 3.2 | 3         |
| 60 | Advanced Concepts and Generative Simulation Formalisms for Creative Discovery Systems Engineering. Intelligent Systems Reference Library, 2011, , 233-258.                                    | 1.2 | 3         |
| 61 | Biomimetic in Silico Devices. Lecture Notes in Computer Science, 2005, , 34-42.   | 1.3 | 2         |
| 62 | In silico simulation of epithelial cell tubulogenesis. , 2008, 2008, 1036-9.  |     | 2         |
| 63 | Using an In Silico Liver to evaluate a hepatic enzyme induction mechanism., 2008, 2008, 2415-8.   |     | 2         |
| 64 | An In Silico Analogue of In Vitro Systems Used to Study Epithelial Cell Morphogenesis. Lecture Notes in Computer Science, 2006, , 285-297.  | 1.3 | 2         |
| 65 | DEVS Peer-to-Peer Protocol for Distributed and Parallel Simulation of Hierarchical and Decomposable DEVS Models., 2007,,.   |     | 1         |
| 66 | Some Unique Properties of a Bilayer and Liposome Forming System. Materials Research Society Symposia Proceedings, 1987, 110, 413.   | 0.1 | 0         |
| 67 | Synthesis of Artificial Models of Sickle Red Cells. Materials Research Society Symposia Proceedings, 1987, 110, 99.   | 0.1 | 0         |
| 68 | Characterization of apocytochrome C binding to human erythrocytes. American Journal of Hematology, 1994, 47, 132-134.   | 4.1 | 0         |
| 69 | In Silico Analogues of Epithelial Cell Growth and Morphogenesis. , 2006, , .  |     | 0         |
| 70 | Simulation modeling of in vitro epithelial morphogenesis and malignancy. Journal of Critical Care, 2007, 22, 347-348.   | 2.2 | 0         |
| 71 | New Simulation Methods to Facilitate Achieving a Mechanistic Understanding of Basic Pharmacology Principles in the Classroom. Journal of Science Education and Technology, 2008, 17, 366-372. | 3.9 | 0         |
| 72 | Agent-based simulation of drug disposition in cirrhotic liver. , 2010, , .  |     | 0         |

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|----|---|-----|-----------|
| 73 | Simulating plausible mechanisms for changing hepatic xenobiotic clearance patterns. , 2010, , .   |     | O         |
| 74 | Simulating Properties of In Vitro Epithelial Cell Morphogenesis. PLoS Computational Biology, 2005, preprint, e129.  | 3.2 | 0         |
| 75 | Synthetic Models and Methods. , 2013, , 2046-2050.  |     | O         |
| 76 | Establishing model mechanismâ€based causal linkages between APAPâ€induced hepatic necrosis and serum ALT. FASEB Journal, 2019, 33, 506.11.                            | 0.5 | 0         |
| 77 | In vitro–in vivo extrapolation of hepatic clearance: using virtual experiments to identify a plausibly influential source of inaccuracies. FASEB Journal, 2019, 33, . | 0.5 | 0         |
| 78 | A framework and mechanistically focused, in silico method for enabling rational translational research. Summit on Translational Bioinformatics, 2008, 2008, 46-50.    | 0.7 | 0         |