2019 ML-MSM Posters

| Author Name | Affiliation | Poster Title | Group # | Poster # | Model Type |
|----------------------|---|--|------------|-------------|---------------|
| | University of California Los | Adapter-Layer RTK Signaling: Basic Understanding & Targeted Drug | | | |
| Aaron Meyer | Angeles | Resistance | 2 | 28 | data |
| Ahmet Erdemir | Cleveland Clinic | Software for Practical Annotation and Exchange of Virtual Anatomy (æva): Design Considerations | 1 | 5 | ODE |
| Amir Barati Farimani | Carnegie Mellon University | Discovery of Partial Differential Equations Using Deep Convolutional Networks | 1 | 10 | PDE |
| Andrzej Przekwas | CFD Research Corporation | Multiscale Simulation Framework for Personalized Pharmacology | 1 | 12 | PDE |
| Andy Somogyi | Indiana University | Multi-Cellular Model Specification and Simulation | 1 | 19 | theory |
| Ashlee Ford Versypt | Oklahoma State University | CAREER: Multiscale Modeling of a Virtual Kidney during the Onset and Progression of Diabetic Kidney Disease | 1 | 14 | theory |
| Assad Oberai | University of Southern California | Deep Generative Priors for Quantifying Uncertainty | 2 | 30 | data |
| Benjamin Gyori | Harvard Medical School | Ecosystem of Machine-maintained Models with Automated Analysis (EMMAA) | 1 | 2 | ODE |
| Chi-Hua Yu | Massachusetts Institute of Technology | Using Artificial Intelligence to Generate de novo Thermally Stable Collagen Sequences | 2 | 34 | data |
| Danh-Tai Hoang | NIDDK | Classification with multiscale hidden variables using Expectation Reflection | 2 | 27 | data |
| | University of | Precision medicine as a control problem: using simulation and deep reinforcement learning to discover adaptive, personalized multi-drug | | | |
| Daniel Faissol | Vermont | therapies. | 2 | 25 | data |
| David Nordsletten | University of Michigan | Towards Automated Biomechanical Analysis of Patients with Hypertrophic Cardiomyopathy | 2 | 29 | data |

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| | | A computational bridge between in vitro MIC and in vivo efficacy of | | | |
| Elsje Pienaar | Purdue University | antibiotics against MAC infection | 1 | 9 | ODE |
| Emily Greenspan | National Cancer Institute | NCI-DOE Collaborations: Extending Frontiers of Predictive Oncology and Exascale Computing | 2 | 26 | data |
| Eric D. | Galois Inc. | Automated Scientific Knowledge Extraction with AMIDOL | 1 | 3 | ODE |
| Farid Yaghouby | Center for Devices and Radiological Health | Sex-Specificity in a Safety of Vagus Nerve Stimulation: An Investigation on Cardiovascular and Immune Systems | 2 | 33 | data |
| Francisco Sahli Costabal | Pontificia Universidad Católica de Chile | Multi-fidelity classification using Gaussian processes: accelerating the prediction of large-scale computational models | 2 | 31 | data |
| Gary An | University of Vermont | Genetic Algorithms for model refinement and rule discovery in a high- dimensional agent-based model ofinflammation | 1 | 13 | theory |
| Herbert Sauro | University of Washington | Using perturbation data and ML to determine kinetic models | 1 | 4 | ODE |
| Jacob Barhak | None | Supervised Learning of Units of Measure | 2 | 23 | data |
| Jessica Zhang | Carnegie Mellon University | Artificial Intelligence Data-driven Model for Adolescent Idiopathic Scoliosis: Analysis, Prediction and Treatment | 2 | 35 | data |
| John Bachman | Harvard Medical School | An Automated Scientific Discovery Framework | 1 | 1 | ODE |
| Jorg Peters | University of Florida | Tri-variate C 1 elements for curved domains | 1 | 11 | PDE |
| Jungmin Han | NIDDK | Obesity and the Sustainability of Calcium Oscillations in Hepatocytes: Explicitly Modeling Mitochondria-associated ER Membranes | 1 | 8 | ODE |
| Kanaka Rajan | Icahcn School of Medicine at Mount Sinai | Multi-region 'Network of Networks' Recurrent Neural Network Models of Adaptive and Maladaptive Learning | 1 | 18 | theory |
| Krishna Garikipati | University of Michigan | Artificial Intelligence guided multi-scale multi-physics framework for discovering complex emergent materials phenomena | 1 | 6 | ODE |

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| | Rochester Institute | End-to-End Uncertainty Quantification in Multiscale Models via Bayesian | | | |
| Linwei Wang | of Technology | Active Learning | 2 | 32 | data |
| Manu Aggarwal | NIDDK | Topological data analysis applied to pancreatic-islet architecture | 2 | 20 | data |
| Misha Pavel | Northeastern University | Multiscale Digital Twin Models of Health Behaviors and Behavioral Change | 1 | 15 | theory |
| Paris Perdikaris | University of Pennsylvania | Machine learning in cardiovascular flows modeling: Predicting arterial blood pressure from non-invasive 4D flow MRI data using physics-informed neural networks | 1 | 16 | theory |
| Parya Aghasafari | University of California Davis | Predictive multiscale in silico cardio-pharmacology | 2 | 21 | data |
| Paul Aiyetan | Frederick National Laboratory for Cancer Research | Towards A Universal Multiscale Data Representation and Knowledge Integration | 2 | 22 | data |
| Peng Zhang | Stony Brook University | Machine Learning in Multiscale Modeling and Validation of In Vitro Experiments of Blood Flow and Platelet Mediated Thrombosis Initiation | 2 | 36 | data |
| Ravi Radhakrishnan | University of Pennsylvania | Computational Algorithms for In Silico Profiling of Activating Mutations in Cancer | 1 | 17 | theory |
| Samuel Britton | University of California Riverside | Learning Regulation and Optimal Control of Enzyme Activities to Preserve Solvent Capacity in the Cell | 2 | 24 | data |
| William Barnett | Georgia State University | Brainstem mechanisms of cardio-ventilatory coupling | 1 | 7 | ODE |