

Digital Discovery

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See Ryan C. Fortenberry *et al.*, pp. 1269–1288.

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See Jean-Louis Reymond *et al.*, pp. 1289–1296. Image reproduced by permission of Markus Orsi from *Digital Discovery*, 2023, 2, 1289.

Background: David Teniers the Younger, "The Alchemist". Mauritshuis, The Hague.

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14 examples of how LLMs can transform materials science and chemistry: a reflection on a large language model hackathon

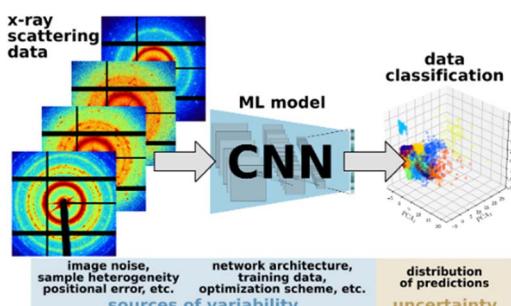
K. M. Jablonka,* Q. Ai, A. Al-Feghali, S. Badhwar, J. D. Bocarsly, A. M. Bran, S. Bringuier, L. C. Brinson, K. Choudhary, D. Circi, S. Cox, W. A. de Jong, M. L. Evans, N. Gastelli, J. Genzling, M. V. Gil, A. K. Gupta, Z. Hong, A. Imran, S. Kruschwitz, A. Labarre, J. Lála, T. Liu, S. Ma, S. Majumdar, G. W. Merz, N. Moitessier, E. Moubarak, B. Mouriño, B. Pelkie, M. Pieler, M. Ramos, B. Ranković, S. G. Rodrigues, J. N. Sanders, P. Schwaller, M. Schwarting, J. Shi, B. Smit, B. E. Smith, J. Van Herck, C. Völker, L. Ward, S. Warren, B. Weiser, S. Zhang, X. Zhang, G. A. Zia, A. Scourtas, K. J. Schmidt, I. Foster, A. D. White and B. Blaiszik*



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A rigorous uncertainty-aware quantification framework is essential for reproducible and replicable machine learning workflows

Line Pouchard, Kristofer G. Reyes, Francis J. Alexander and Byung-Jun Yoon*



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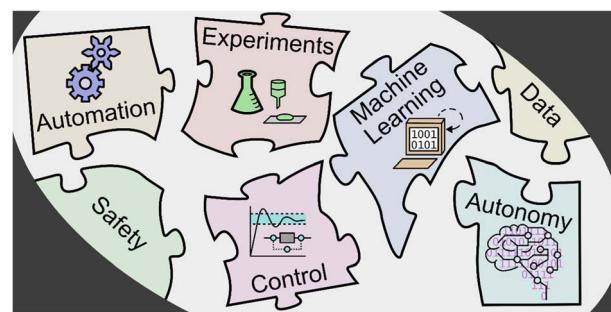


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Integrating autonomy into automated research platforms

Richard B. Carty, Brent A. Koscher, Matthew A. McDonald and Klavs F. Jensen*

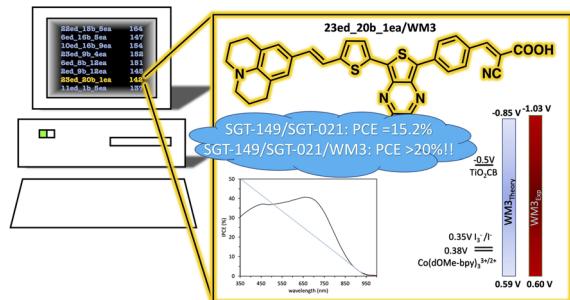


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An automated quantum chemistry-driven, experimental characterization for high PCE donor–π–acceptor NIR molecular dyes

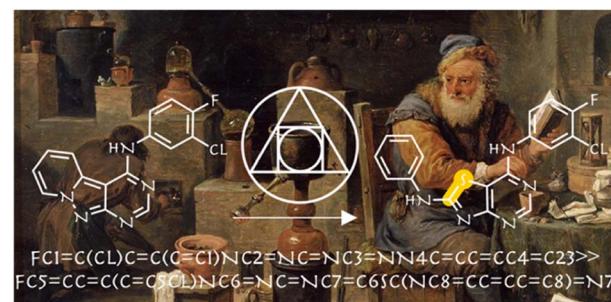
Taylor J. Santaloci, William E. Meador, Austin M. Wallace, E. Michael Valencia, Blake N. Rogers, Jared H. Delcamp and Ryan C. Fortenberry*



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Alchemical analysis of FDA approved drugs

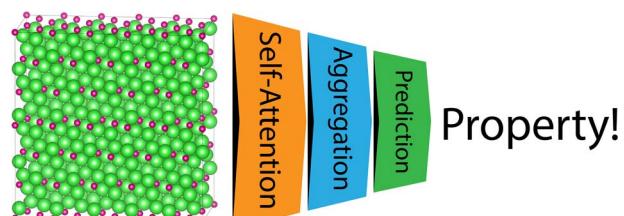
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Michael Moran, Michael W. Gaultois,* Vladimir V. Gusev and Matthew J. Rosseinsky



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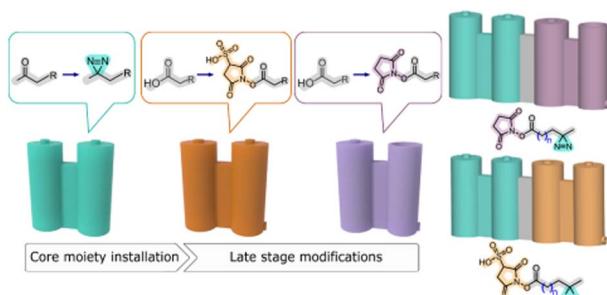
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Generating structural alerts from toxicology datasets using the local interpretable model-agnostic explanations method

Cayque Monteiro Castro Nascimento, Paloma Guimarães Moura and Andre Silva Pimentel*

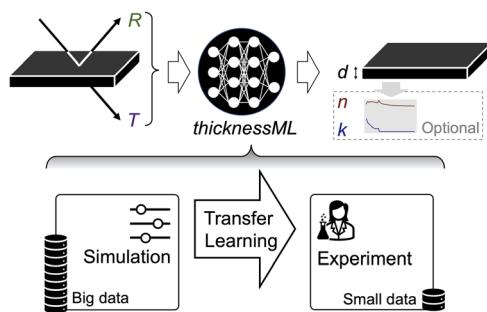
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Przemyslaw Frei, Philip J. Kitson, Alexander X. Jones and Leroy Cronin*

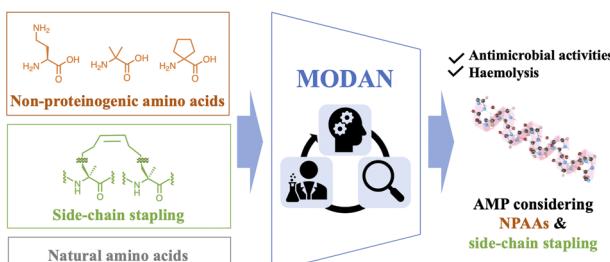
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Siyu Isaac Parker Tian, Zekun Ren, Selvaraj Venkataraj, Yuanhang Cheng, Daniil Bash, Felipe Oviedo, J. Senthilnath, Vijila Chellappan, Yee-Fun Lim, Armin G. Aberle, Benjamin P. MacLeod, Fraser G. L. Parlane, Curtis P. Berlinguette, Qianxiao Li, Tonio Buonassisi* and Zhe Liu

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Design of antimicrobial peptides containing non-proteinogenic amino acids using multi-objective Bayesian optimisation

Yuki Murakami, Shoichi Ishida, Yosuke Demizu and Kei Terayama*

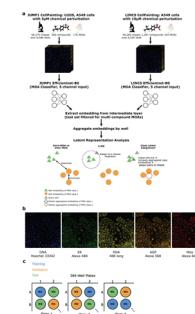


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Deep representation learning determines drug mechanism of action from cell painting images

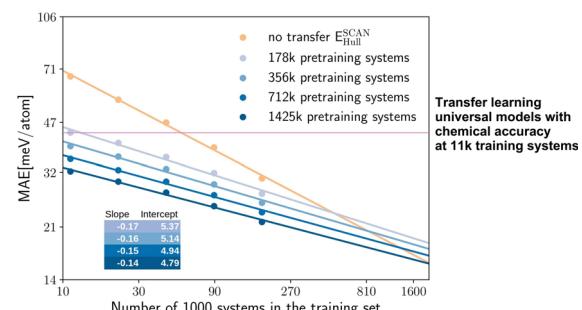
Daniel R. Wong,* David J. Logan, Santosh Hariharan, Robert Stanton, Djork-Arné Clevert and Andrew Kiruluta



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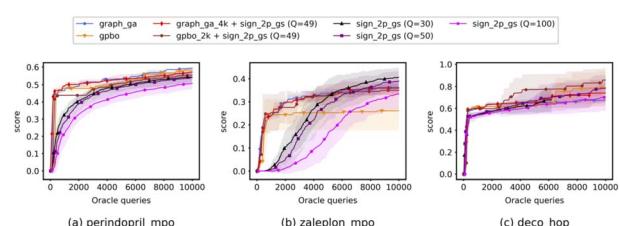
Noah Hoffmann, Jonathan Schmidt, Silvana Botti and Miguel A. L. Marques*



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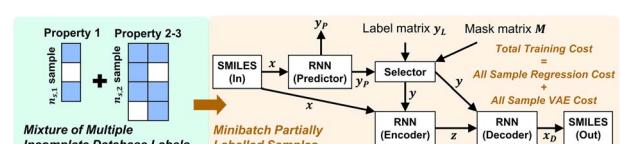
Elvin Lo and Pin-Yu Chen*



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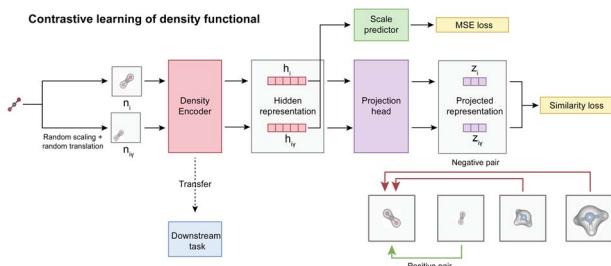
Multi-constraint molecular generation using sparsely labelled training data for localized high-concentration electrolyte diluent screening

Jonathan P. Mailoa,* Xin Li, Jiezhong Qiu and Shengyu Zhang*



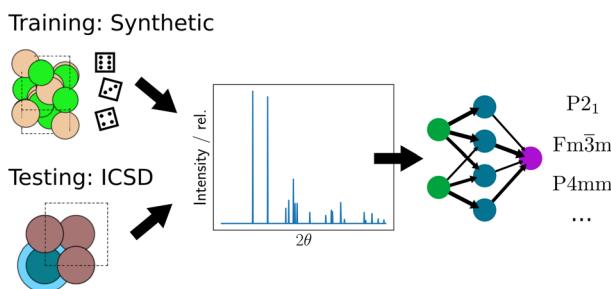
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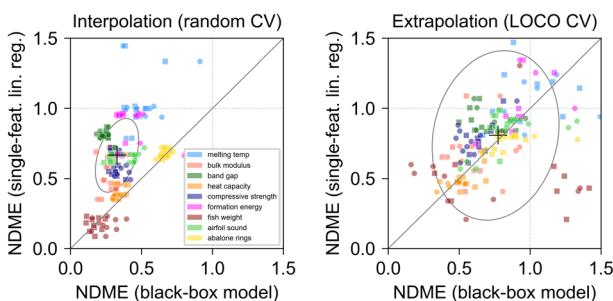
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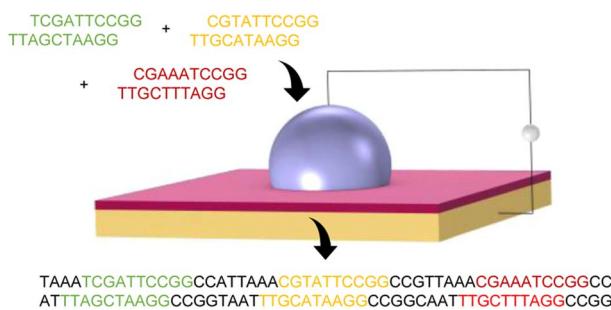
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Ajay Manicka, Andrew Stephan, Sriram Chari, Gemma Mendonsa, Peyton Okubo, John Stolzberg-Schray, Anil Reddy and Marc Riedel

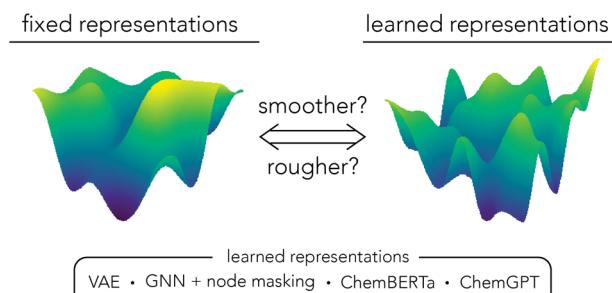


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Evaluating the roughness of structure–property relationships using pretrained molecular representations

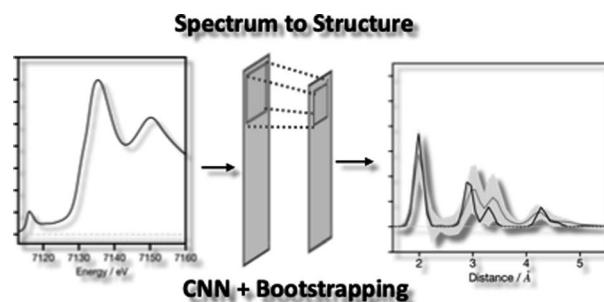
David E. Graff, Edward O. Pyzer-Knapp, Kirk E. Jordan, Eugene I. Shakhnovich and Connor W. Coley



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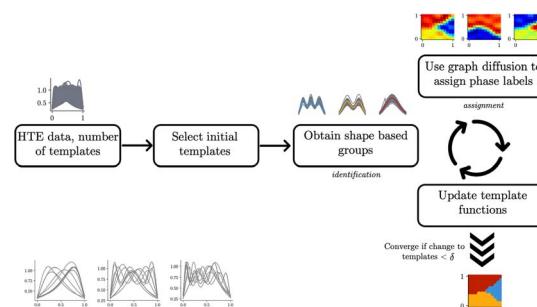
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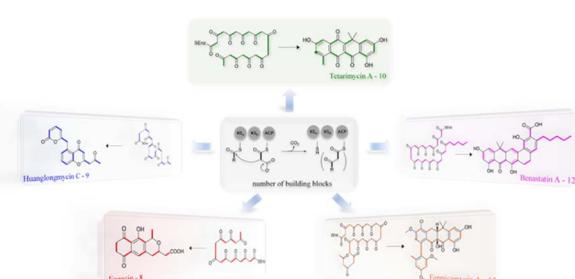
Kiran Vaddi,* Karen Li and Lilo D. Pozzo



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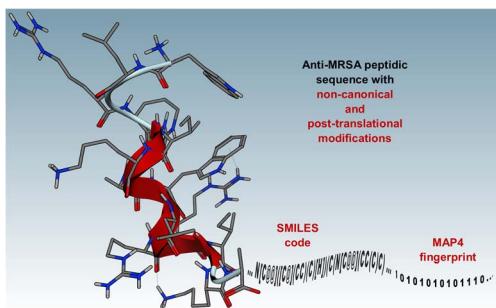
A deep learning model for type II polyketide natural product prediction without sequence alignment

Jiaquan Huang, Qiandi Gao, Ying Tang, Yixin Wu, Heqian Zhang* and Zhiwei Qin*



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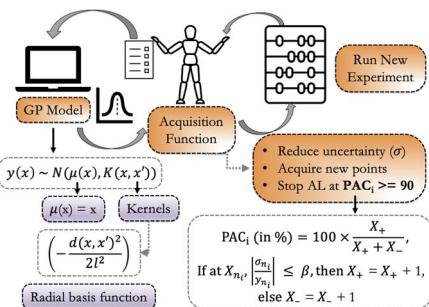
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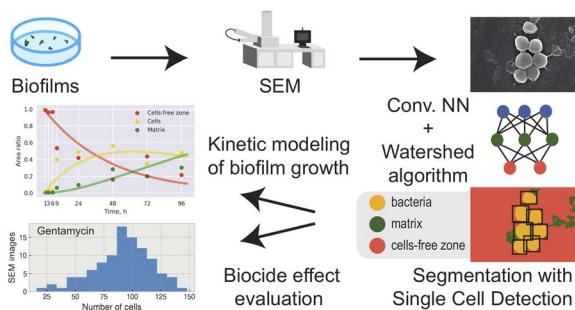
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Krishnendu Mukherjee, Etinosa Osaro and Yamil J. Colón*

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Go with the flow: deep learning methods for autonomous viscosity estimations

Michael Walker, Gabriella Pizzuto, Hatem Fakhruldeen and Andrew I. Cooper*

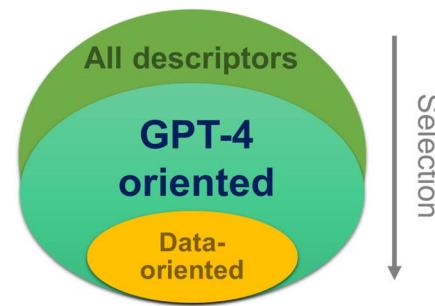


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Using GPT-4 in parameter selection of polymer informatics: improving predictive accuracy amidst data scarcity and 'Ugly Duckling' dilemma

Kan Hatakeyama-Sato,* Seigo Watanabe, Naoki Yamane, Yasuhiko Igarashi and Kenichi Oyaizu*



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Element similarity in high-dimensional materials representations

Anthony Onwuli, Ashish V. Hegde, Kevin V. T. Nguyen, Keith T. Butler* and Aron Walsh*

Chemical Elements as Vectors

One hot encoding

[1 0 0 0 0 0 0 0 ...]
H

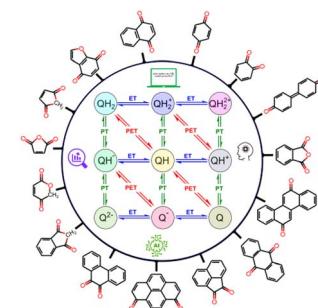
Distributed embedding

[1 1 0 1 0 0 1 0 0 ...]
From machine learning

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Density functional theory and machine learning for electrochemical square-scheme prediction: an application to quinone-type molecules relevant to redox flow batteries

Arsalan Hashemi,* Reza Khakpour, Amir Mahdian, Michael Busch, Pekka Peljo and Kari Laasonen

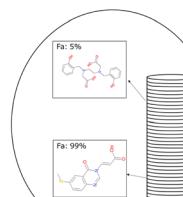


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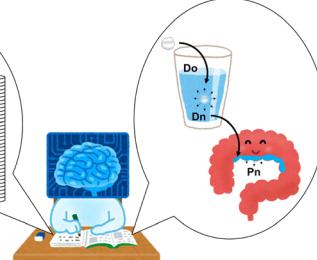
Combined data-driven and mechanism-based approaches for human-intestinal-absorption prediction in the early drug-discovery stage

Koichi Handa,* Sakae Sugiyama, Michiharu Kageyama and Takeshi Iijima

Data-driven Approach

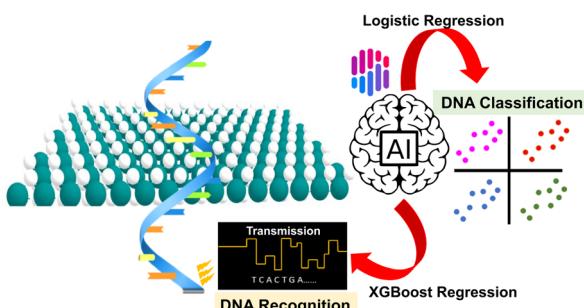


Mechanism-based Approach



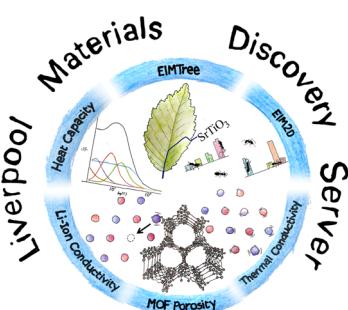
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Artificial intelligence aided recognition and classification of DNA nucleotides using MoS₂ nanochannels

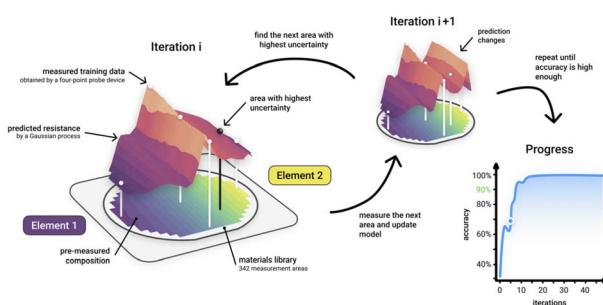
Sneha Mittal, Souvik Manna, Milan Kumar Jena and Biswarup Pathak*

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The Liverpool materials discovery server: a suite of computational tools for the collaborative discovery of materials

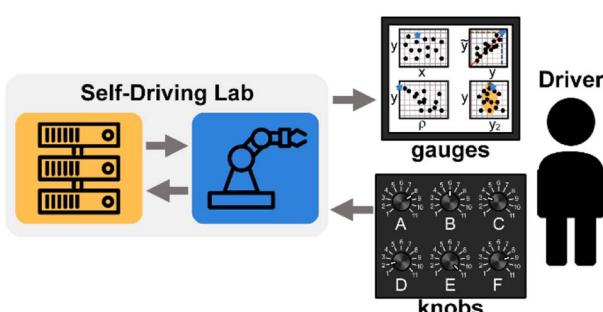
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Speeding up high-throughput characterization of materials libraries by active learning: autonomous electrical resistance measurements

Felix Thelen, Lars Banko, Rico Zehl, Sabrina Baha and Alfred Ludwig*

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Driving school for self-driving labs

Kelsey L. Snapp and Keith A. Brown*

