Litmaps

Mapping the citation network – augmenting human intelligence with AI

VALA 2024



Axton Pitt Co-founder and CEO

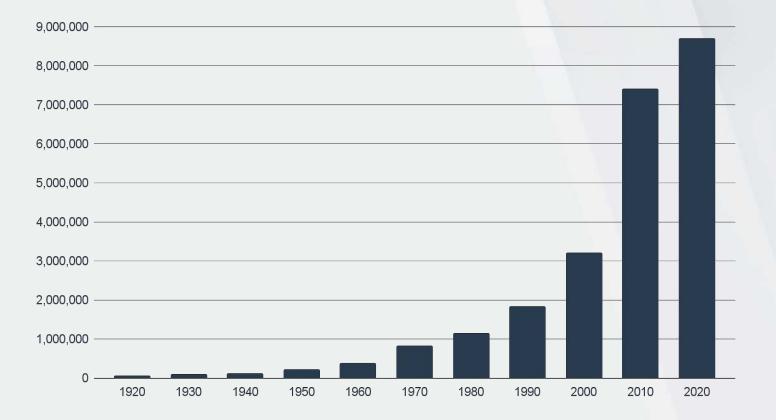
In this talk

Why Design challenges Data Human + Al Future



Why Litmaps?

SCIENTIFIC PAPERS PUBLISHED



Priem, J., Piwowar, H., & Orr, R. (2022). OpenAlex: A fully-open index of scholarly works, authors, venues, institutions, and concepts. ArXiv. https://arxiv.org/abs/2205.01833

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[HTML] Overview on peroxiredoxin

SG Rhee - Molecules and cells, 2016 - Elsevier

Peroxiredoxins (Prxs) are a very large and highly conserved family of peroxidases that reduce peroxides, with a conserved cysteine residue, designated the "peroxidatic" Cys (C P) ... \therefore Save $\overline{\mathfrak{M}}$ Cite Cited by 420 Related articles All 16 versions Web of Science: 260

Peroxiredoxin 1 and its role in cell signaling

CA Neumann, <u>J Cao</u>, Y Manevich - Cell cycle, 2009 - Taylor & Francis Peroxiredoxins (Prdxs) are a family of small (22-27kDa) non-seleno peroxidases currently known to possess six mammalian isoforms. Although their individual roles in cellular redox ... ☆ Save 切 Cite Cited by 276 Related articles All 10 versions Web of Science: 182

Advances in our understanding of **peroxiredoxin**, a multifunctional, mammalian redox protein

J Fujii, Y Ikeda - Redox Report, 2002 - Taylor & Francis

... A new family of proteins with an antioxidative function, now designated as **peroxiredoxin** (Prx), ... Genetic mapping of six mouse **peroxiredoxin** genes and fourteen **peroxiredoxin** related ...

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Peroxiredoxin evolution and the regulation of hydrogen peroxide signaling

ZA Wood, LB Poole, PA Karplus - Science, 2003 - science.org

Eukaryotic 2-Cys peroxiredoxins (2-Cys Prxs) not only act as antioxidants, but also appear to regulate hydrogen peroxide–mediated signal transduction. We show that bacterial 2-Cys ...

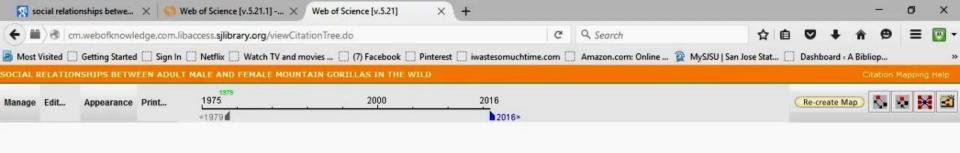
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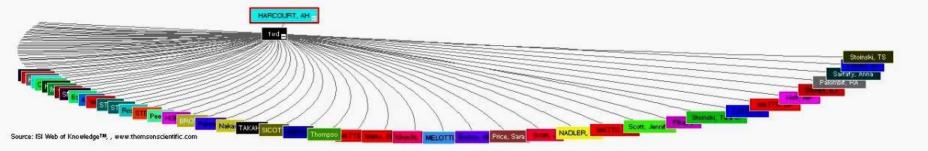
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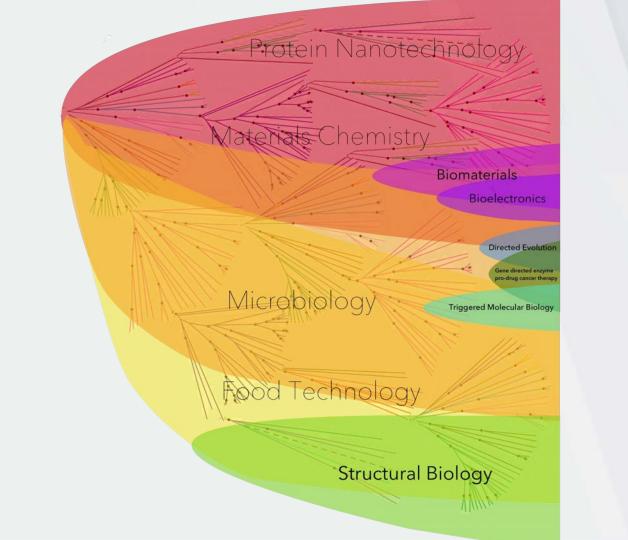
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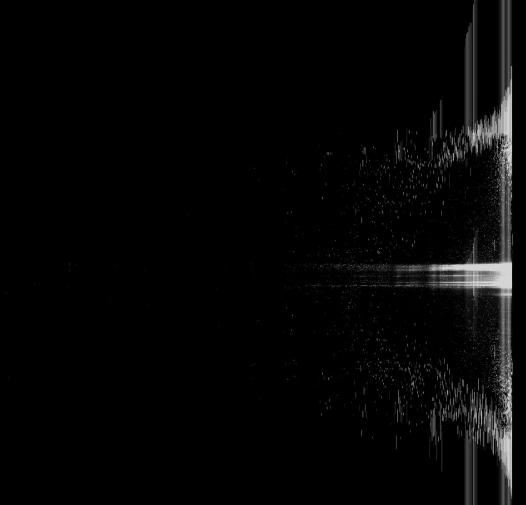


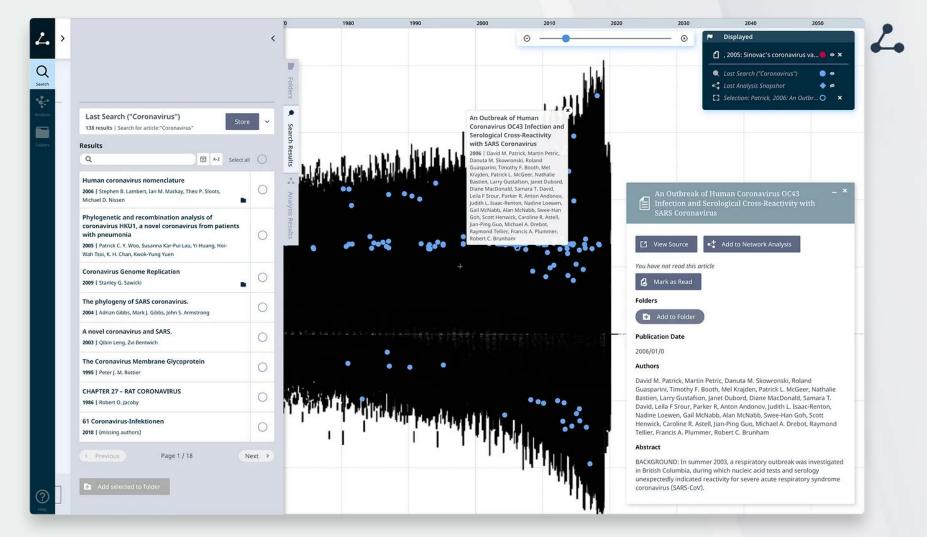
Accelerating Impactful Science



What didn't work?



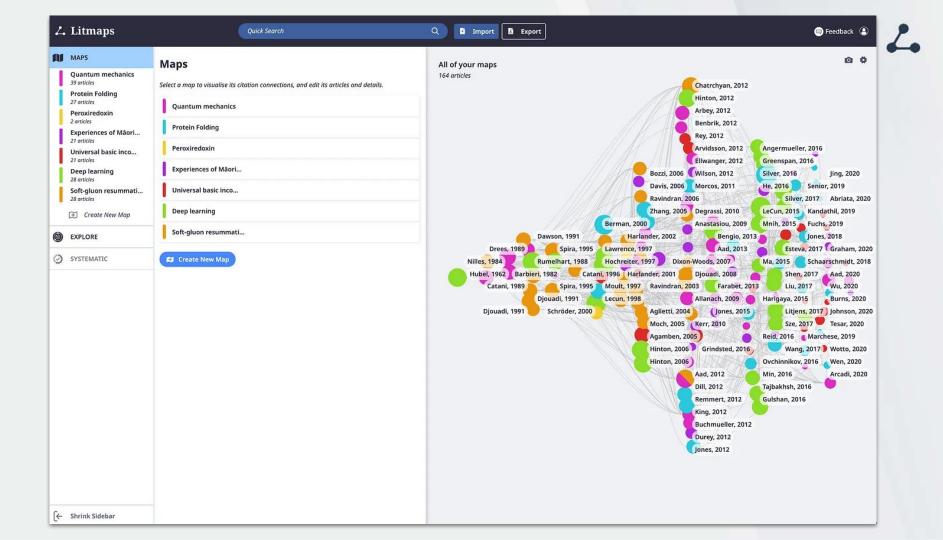


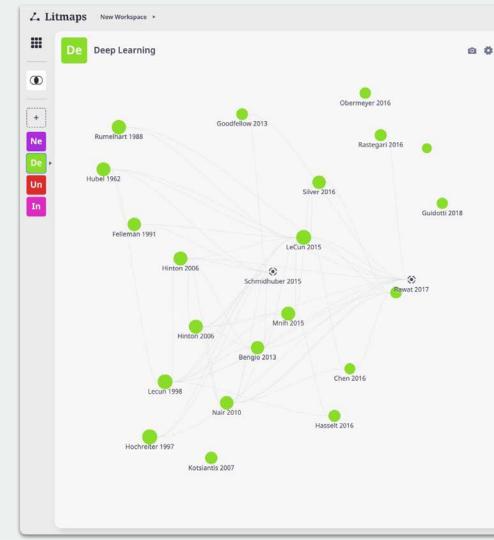




How did we get here?

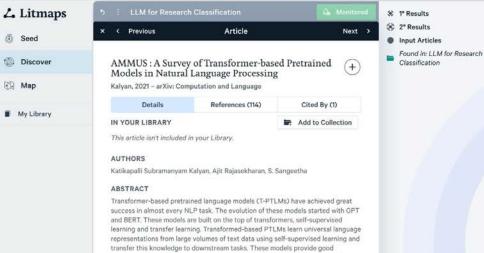
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he oligomeric conformation of peroxiredoxins links redox state to function.	A A A A A A A A A A A A A A A A A A A
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Discrimination of single-stranded DNA homopolymers by sieving out G-	
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019 Wei Si, Haojie Yang, Jingjie Sha, Yin Zhang, Yunfei Chen	Wood, 2003 Barranco-Medina, 2009 Phillips, 2014 Yewdall, 2016 Sharapov, 2018
Peroxiredoxins: a historical overview and speculative preview of novel	
nechanisms and emerging concepts in cell signaling, 005 Sue Goo Rhee, Ho Zoon Chae, Kanghwa Kim	
	Angelucci, 2015
Peroxiredoxin Evolution and the Regulation of Hydrogen Peroxide Signaling	
2003 Zachary A. Wood, Leslie B. Poole, P. Andrew Karplus	
A primer on peroxiredoxin biochemistry,	
2015 P. Andrew Karplus	
Formation, TEM study and 3D reconstruction of the human erythrocyte	
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	Zhong, 2019 From shallow feature learning to deep learning: Benefits from the width and depth of deep architectures					
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	Zhong. 2016 An overview on data representation learning: From traditional feature learning to recent deep learning					
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representations from large volumes of text data using self-supervised learning and transfer this knowledge to downstream tasks. These models provide good background knowledge to downstream tasks. These models provide good background knowledge to downstream tasks. These models provide good background knowledge to downstream tasks. These models provide good pretraining, pretraining methods, pretraining tasks, embeddings and downstream adaptation methods. Next, we present a new taxonomy of T-PTLMs and then give brief overview of various benchmarks including both intrinsic and extrinsic. We present a summary of various useful libraries to work with T-PTLMs. Finally, we highlight some of the future research directions which will further improve these models. We strongly believe that this comprehensive survey paper will serve as a good reference to learn the core concepts as well as to stay updated with the recent happenings in T-PTLMs.

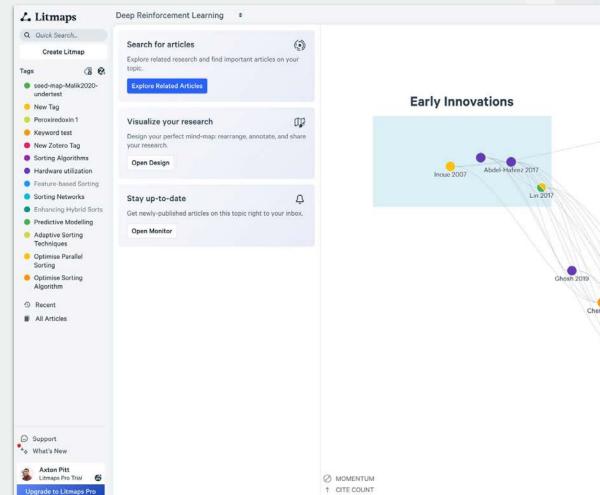
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Axton Pitt

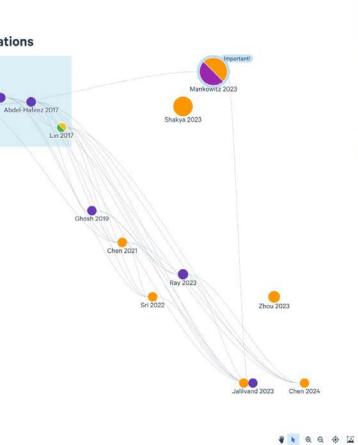


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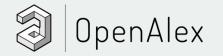
Dataset



Semantic **Scholar**

Import Process







+270M Scholarly Items