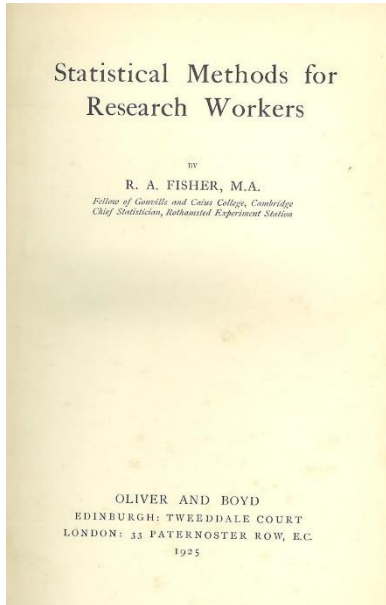


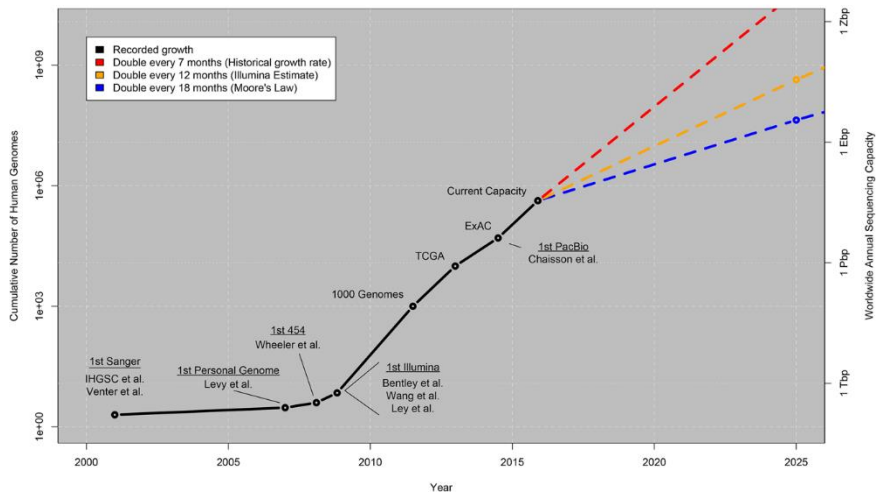
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# The data science revolution in scientific research

Jonathan Rowe  
Programme Director, Data Science for Science



Growth of DNA Sequencing



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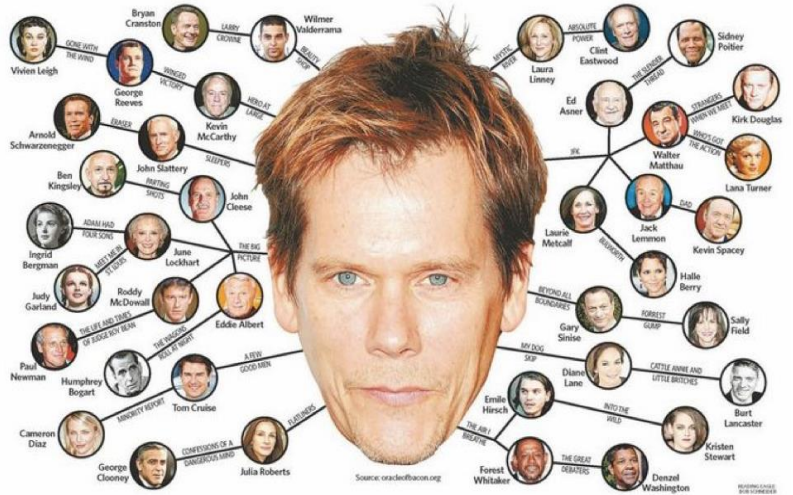
Published: July 7, 2015 • <https://doi.org/10.1371/journal.pbio.1002195>

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Methods and Applications

Stanley Wasserman and Katherine Faust





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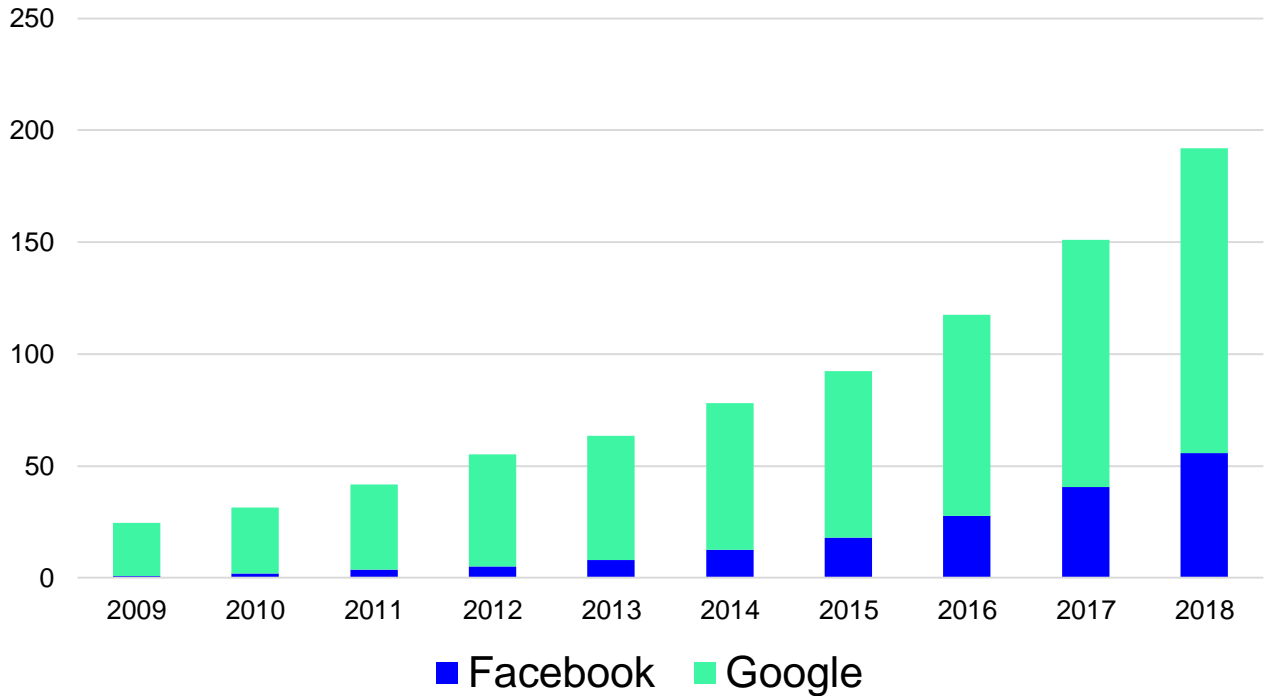
RESEARCH ARTICLE

# Testing for Questionable Research Practices in a Meta-Analysis: An Example from Experimental Parapsychology

**Dick J. Bierman<sup>1,2\*</sup>, James P. Spottiswoode<sup>3</sup>, Aron Bijl<sup>4</sup>**

**1** Experimental Psychology, University of Groningen, Groningen, The Netherlands, **2** Brain & Cognition, University of Amsterdam, Amsterdam, The Netherlands, **3** LFR, Palo Alto, California, United States of America, **4** Brain & Cognition, University of Amsterdam, Amsterdam, The Netherlands

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# ImageNet Classification with Deep Convolutional Neural Networks

---

**Alex Krizhevsky**  
University of Toronto  
kriz@cs.utoronto.ca

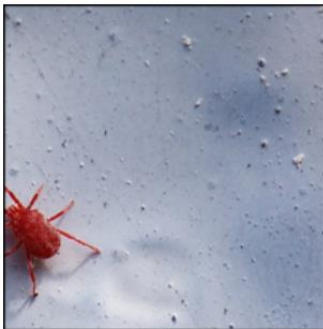
**Ilya Sutskever**  
University of Toronto  
ilya@cs.utoronto.ca

**Geoffrey E. Hinton**  
University of Toronto  
hinton@cs.utoronto.ca

## Abstract

We trained a large, deep convolutional neural network to classify the 1.2 million high-resolution images in the ImageNet ILSVRC-2010 contest into the 1000 different classes. On the test data, we achieved top-1 and top-5 error rates of 37.5% and 17.0% which is considerably better than the previous state-of-the-art. The neural network, which has 60 million parameters and 650,000 neurons, consists of five convolutional layers, some of which are followed by max-pooling layers, and three fully-connected layers with a final 1000-way softmax. To make training faster, we used non-saturating neurons and a very efficient GPU implementation of the convolution operation. To reduce overfitting in the fully-connected layers we employed a recently-developed regularization method called “dropout” that proved to be very effective. We also entered a variant of this model in the ILSVRC-2012 competition and achieved a winning top-5 test error rate of 15.3%, compared to 26.2% achieved by the second-best entry.





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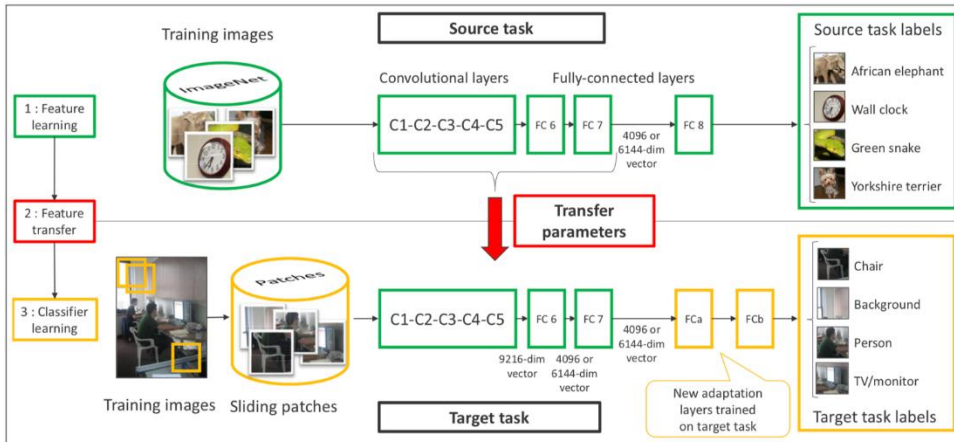


## A general reinforcement learning algorithm that masters chess, shogi and Go through self-play

David Silver,<sup>1,2\*</sup> Thomas Hubert,<sup>1\*</sup> Julian Schrittwieser,<sup>1\*</sup>  
Ioannis Antonoglou,<sup>1,2</sup> Matthew Lai,<sup>1</sup> Arthur Guez,<sup>1</sup> Marc Lanctot,<sup>1</sup>  
Laurent Sifre,<sup>1</sup> Dharshan Kumaran,<sup>1,2</sup> Thore Graepel,<sup>1,2</sup>  
Timothy Lillicrap,<sup>1</sup> Karen Simonyan,<sup>1</sup> Demis Hassabis<sup>1</sup>

<sup>1</sup>DeepMind, 6 Pancras Square, London N1C 4AG.

<sup>2</sup>University College London, Gower Street, London WC1E 6BT.



## Learning and Transferring Mid-Level Image Representations using Convolutional Neural Networks

Maxime Oquab<sup>1,\*</sup> Leon Bottou<sup>2</sup> Ivan Laptev<sup>1,\*</sup> Josef Sivic<sup>1,\*</sup>

<sup>1</sup>INRIA, Paris, France <sup>2</sup>MSR, New York, USA

Article | OPEN | Published: 16 October 2017

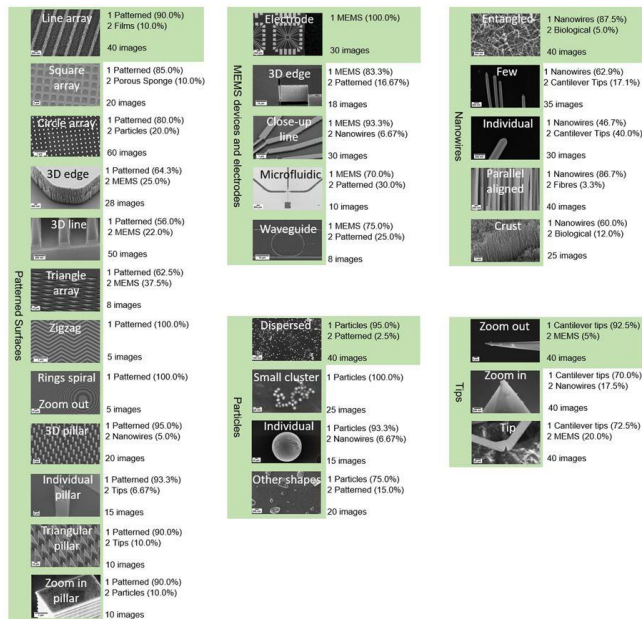
# Neural Network for Nanoscience Scanning Electron Microscope Image Recognition

Mohammad Hadi Modarres, Rossella Aversa , Stefano Cozzini, Regina Ciancio, Angelo Leto & Giuseppe Piero Brandino

Scientific Reports 7, Article number: 13282 (2017) | [Download Citation](#)

## Abstract

In this paper we applied [transfer learning](#) techniques for image recognition, automatic categorization, and labeling of nanoscience images obtained by scanning electron microscope (SEM). Roughly 20,000 SEM images were manually classified into 10 categories to form a labeled training set, which can be used as a reference set for future applications of deep learning enhanced algorithms in the nanoscience domain. The categories chosen spanned the range of 0-Dimensional (0D) objects such as particles, 1D nanowires and fibres, 2D films and coated surfaces, and 3D patterned surfaces such as pillars. The training set was used to retrain on the SEM dataset and to compare many





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## Generative Adversarial Nets

---

Ian J. Goodfellow<sup>\*</sup>, Jean Pouget-Abadie<sup>†</sup>, Mehdi Mirza, Bing Xu, David Warde-Farley,  
Sherjil Ozair<sup>‡</sup>, Aaron Courville, Yoshua Bengio<sup>§</sup>  
Département d'informatique et de recherche opérationnelle  
Université de Montréal  
Montréal, QC H3C 3J7

Published as a conference paper at ICLR 2018

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### PROGRESSIVE GROWING OF GANs FOR IMPROVED QUALITY, STABILITY, AND VARIATION

Tero Karras      Timo Aila      Samuli Laine      Jaakko Lehtinen  
NVIDIA          NVIDIA          NVIDIA          NVIDIA and Aalto University  
{tkarras, taila, slaine, jlehtinen}@nvidia.com

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# Accelerating Science with Generative Adversarial Networks: An Application to 3D Particle Showers in Multilayer Calorimeters

Michela Paganini, Luke de Oliveira, and Benjamin Nachman  
Phys. Rev. Lett. **120**, 042003 – Published 26 January 2018

Article

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## ABSTRACT

Physicists at the Large Hadron Collider (LHC) rely on detailed simulations of particle collisions to build expectations of what experimental data may look like under different theoretical modeling assumptions. Petabytes of simulated data are needed to develop analysis techniques, though they are expensive to generate using existing algorithms and computing resources. The modeling of detectors and the precise description of particle cascades as they interact with the material in the calorimeter are the most computationally demanding steps in the simulation pipeline. We therefore introduce a deep neural network-based generative model to enable high-fidelity, fast, electromagnetic calorimeter simulation. There are still challenges for achieving precision across the entire phase space, but our current solution can reproduce a variety of particle shower properties while achieving speedup factors of up to  $100\,000\times$ . This opens the door to a new era of fast simulation that could save significant computing time and disk space, while extending the reach of physics searches and precision measurements at the LHC and beyond.

# Active learning machine learns to create new quantum experiments



Alexey A. Melnikov, Hendrik Poulsen Nautrup, Mario Krenn, Vedran Dunjko, Markus Tiersch, Anton Zeilinger, and Hans J. Briegel

PNAS February 6, 2018 115 (6) 1221-1226; published ahead of print January 18, 2018 <https://doi.org/10.1073/pnas.1714936115>

Contributed by Anton Zeilinger, November 14, 2017 (sent for review August 24, 2017; reviewed by Jacob D. Biamonte and Jonathan P. Dowling)

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## Significance

Quantum experiments push the envelope of our understanding of fundamental concepts in quantum physics. Modern experiments have exhaustively probed the basic notions of quantum theory. Arguably, further breakthroughs require the tackling of complex quantum phenomena and consequently require complex experiments and involved techniques. The designing of such complex experiments is difficult and often clashes with human intuition. We present an autonomous learning model which learns to design such complex experiments, without relying on previous knowledge or often flawed intuition. Our system not only learns how to design desired experiments more efficiently than the best previous approaches, but in the process also discovers nontrivial experimental techniques. Our work demonstrates that learning machines can offer dramatic advances in how experiments are generated.



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**George Osborne, Chancellor of the Exchequer**  
Budget Speech, March 2014

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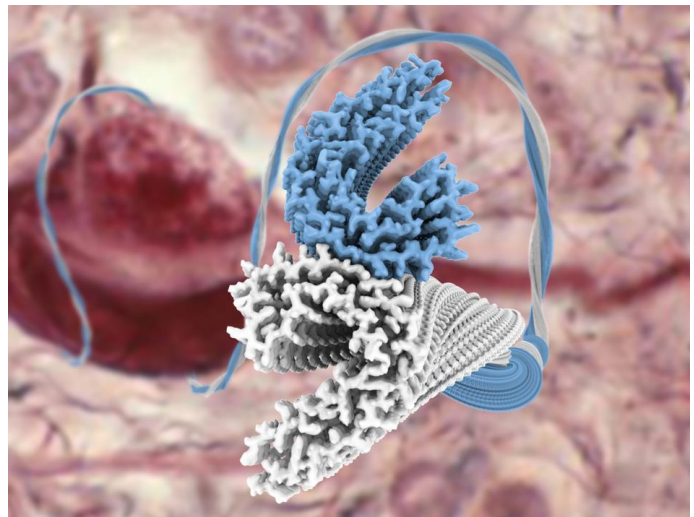


Article | Published: 05 July 2017

# Cryo-EM structures of tau filaments from Alzheimer's disease

Anthony W. P. Fitzpatrick, Benjamin Falcon, Shaoda He, Alexey G. Murzin, Garib Murshudov, Holly J. Garringer, R. Anthony Crowther, Bernardino Ghetti, Michel Goedert  & Sjors H. W. Scheres 

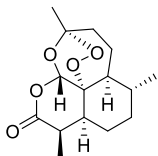
*Nature* **547**, 185–190 (13 July 2017) | [Download Citation](#) 





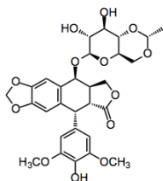
### Anti-malarial

Artemisinin  
Wormwood  
(*artemisia annua*)



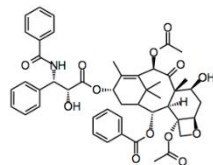
### Anti-cancer

Etoposide Mayapple  
(*podophyllum*)



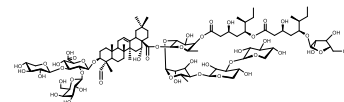
### Anti-cancer

Taxol Yew  
(*taxus baccata*)

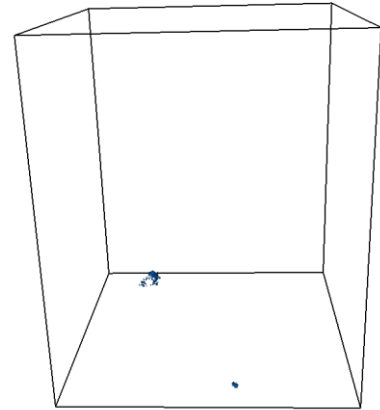
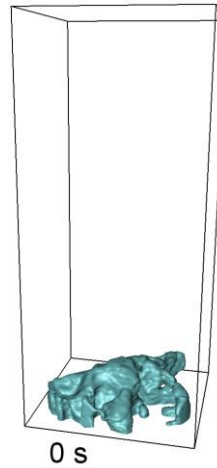
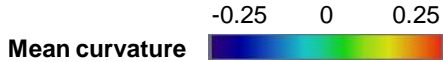
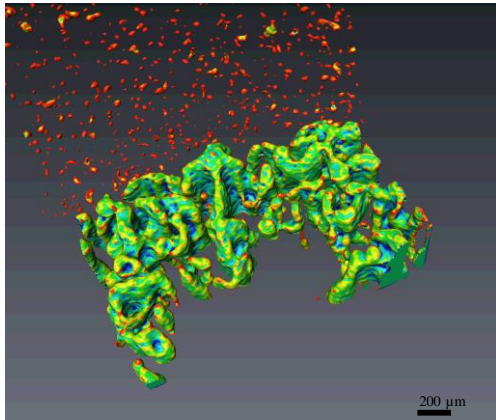


### Adjuvant

Qs-21 Soapbark  
(*quillaja saponaria*)



The majority of plant natural products cannot be accessed – and most of them have not yet been discovered



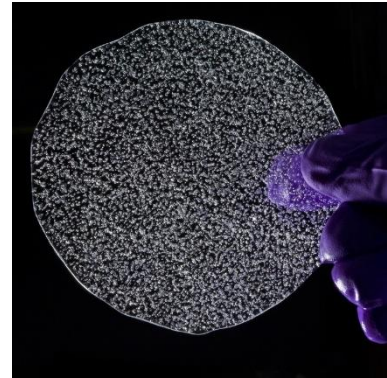
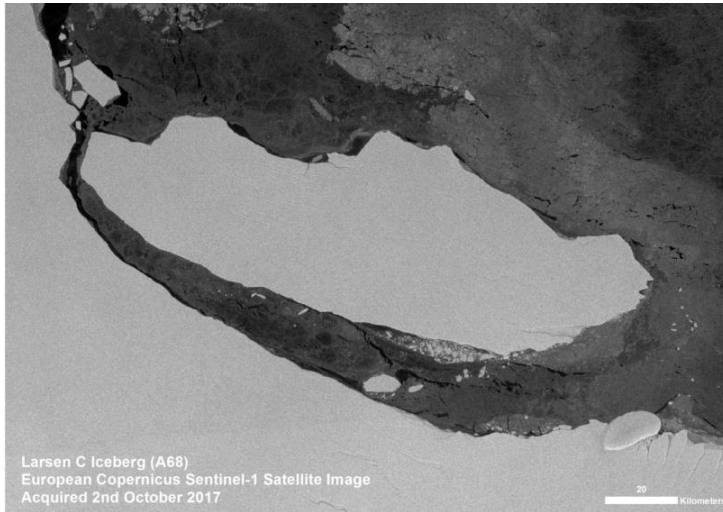
A positive mean curvature means  
concave towards the liquid

B. Cai et al. Acta Mater. 2014(76): 371-380; B. Cai et al. Acta Mater. 2017(117):160-169;  
B. Cai et al. Scripta Mater. submitted.



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Francisco de Goya, Doña Isabel de Porcel (NG1473), before 1805



Color Photograph



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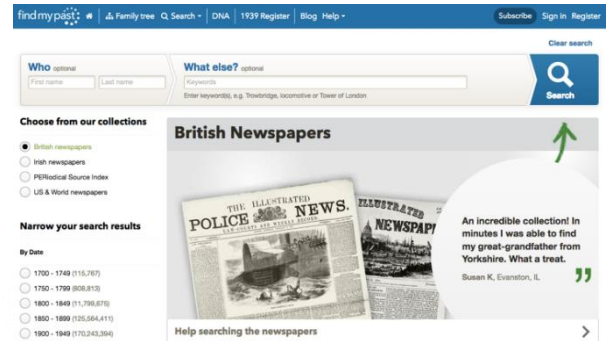
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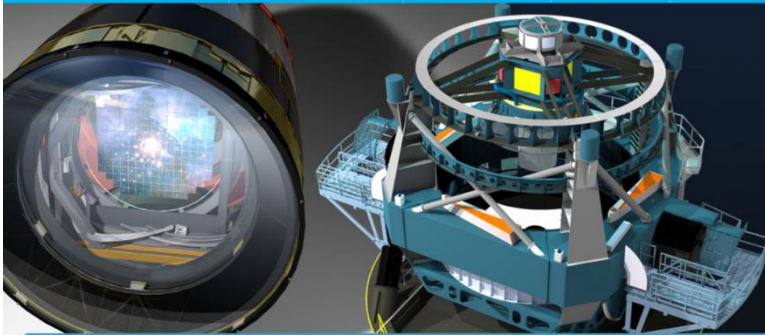
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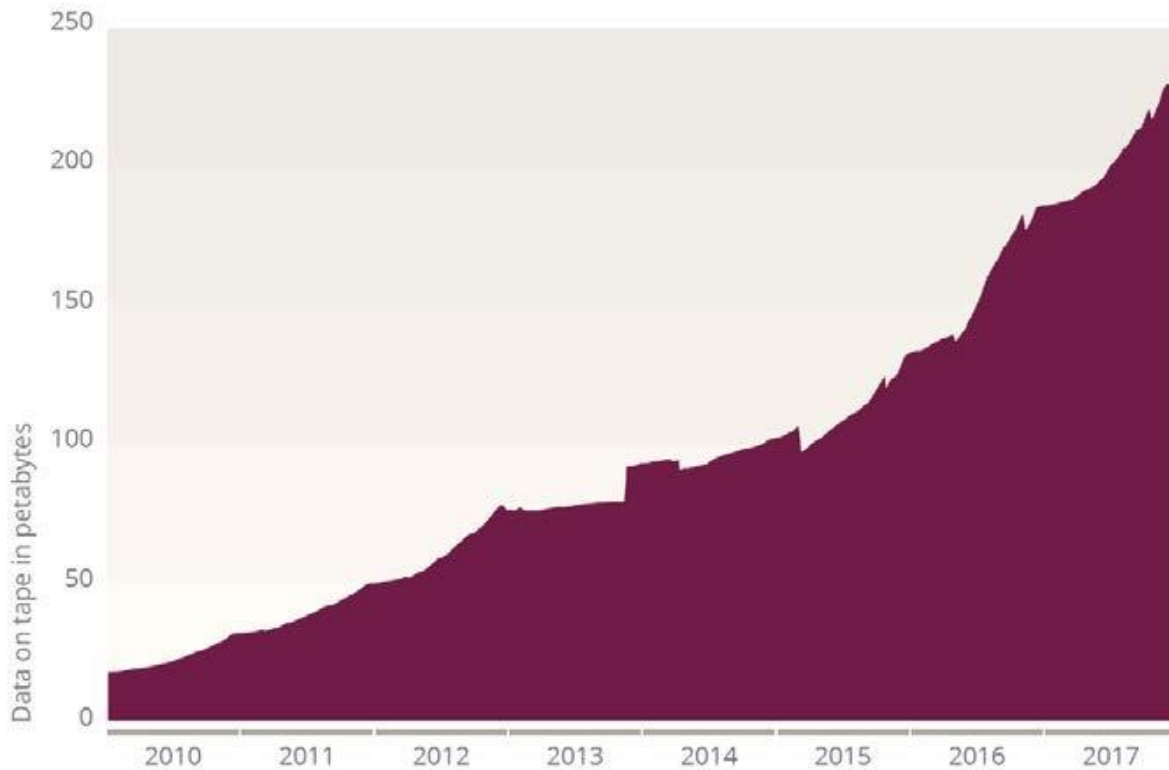
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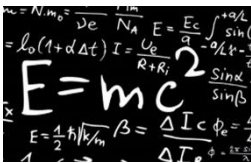
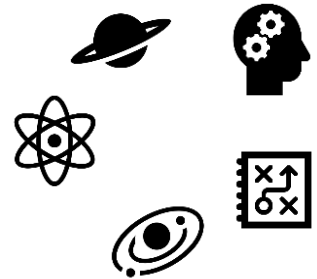
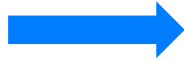


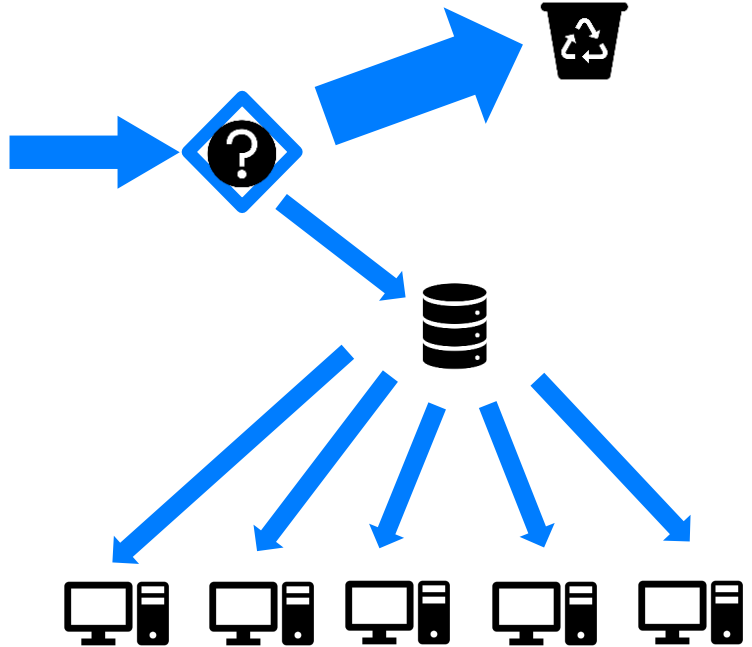
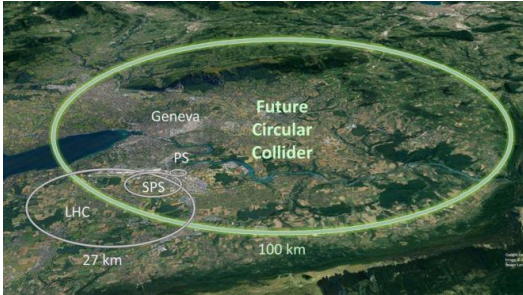
“A just machine to make big decisions  
Programmed by fellows with  
compassion and vision  
We'll be clean when their work is done  
We'll be eternally free, yes and eternally  
young.

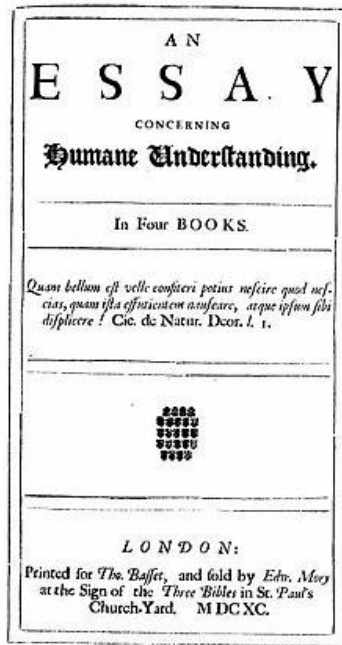
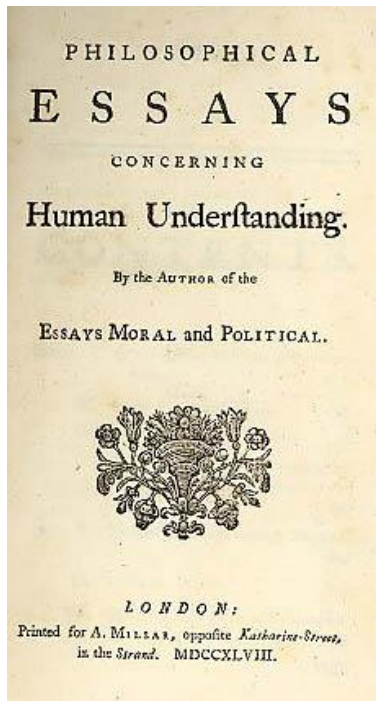
What a beautiful world this will be  
What a glorious time to be free.”

**Donald Fagen**

I.G.Y. (International Geophysical Year)







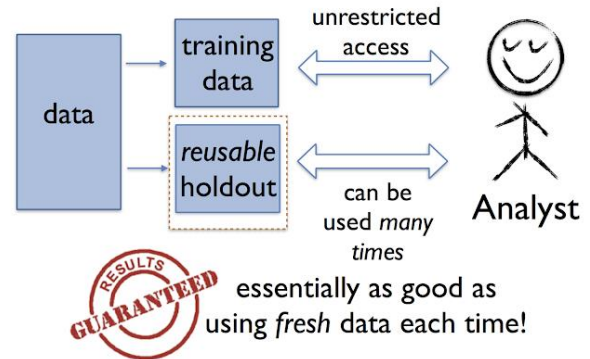
## STATISTICS

# The reusable holdout: Preserving validity in adaptive data analysis

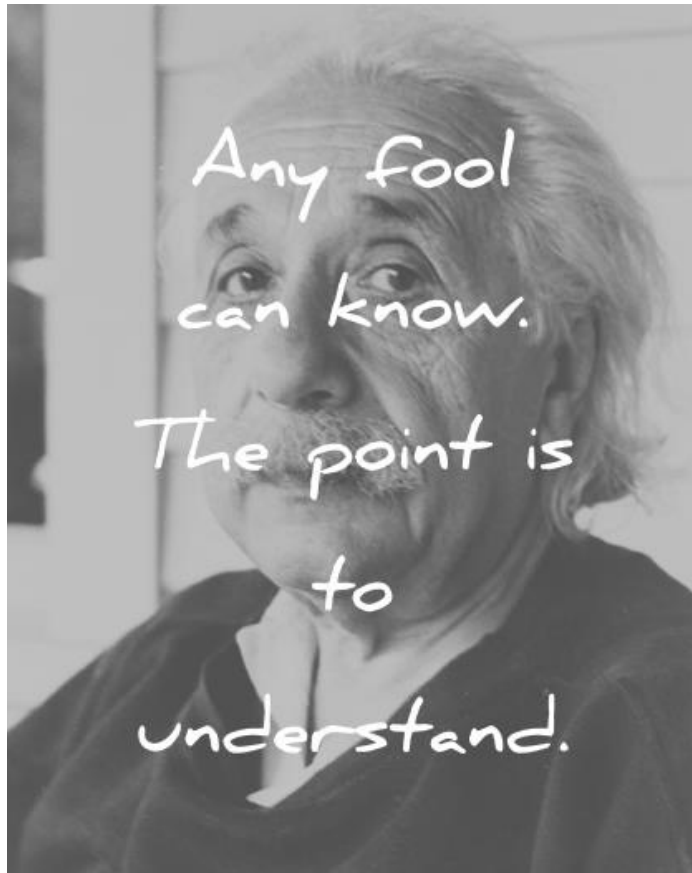
Cynthia Dwork,<sup>1\*</sup> Vitaly Feldman,<sup>2\*</sup> Moritz Hardt,<sup>3\*</sup> Toniann Pitassi,<sup>4\*</sup>  
Omer Reingold,<sup>5\*</sup> Aaron Roth<sup>6\*</sup>

Misapplication of statistical data analysis is a common cause of spurious discoveries in scientific research. Existing approaches to ensuring the validity of inferences drawn from data assume a fixed procedure to be performed, selected before the data are examined. In common practice, however, data analysis is an intrinsically adaptive process, with new analyses generated on the basis of data exploration, as well as the results of previous analyses on the same data. We demonstrate a new approach for addressing the challenges of adaptivity based on insights from privacy-preserving data analysis. As an application, we show how to safely reuse a holdout data set many times to validate the results of adaptively chosen analyses.

## Reusable holdout method







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