Accelerating Science
A Grand Challenge for AI?

CCC Task Force on Convergence of Data and Computing
Vasant Honavar, Mark Hill, Kathy Yelick
Presentation to DARPA Defense Science Office
March 16, 2017
The mission of Computing Research Association's Computing Community Consortium (CCC) is to catalyze the computing research community and enable the pursuit of innovative, high-impact research.

Promote Audacious Thinking:
- Community Initiated Visioning Workshops
- Blue Sky Ideas tracks at conferences

Inform Science Policy
- Outputs of visioning activities
- Task Forces – e.g., Artificial Intelligence, Data and Computing, Health, Internet of Things, Privacy

Engage the Community:
- CCC Blog - [http://cccblog.org/](http://cccblog.org/)
- Computing Research in Action Videos
- Research “Highlight of the Week”

Promote Leadership and Service:
- Computing Innovation Fellows Project
- Leadership in Science Policy Institute
Accelerating Science: A Grand Challenge for AI?

• Discussion based in part on:
  – AAAI Fall Symposium on Accelerating Science: A Grand Challenge for AI

• Other related events:
  – NSF Workshop on Discovery Informatics, February 2012
  – AAAI Fall Symposium on Discovery Informatics, November 2012
  – CMUSV Symposium on Cognitive Systems and Discovery Informatics, 2013
  – AAAI Fall Symposium on Discovery Informatics, November 2013
  – AAAI Workshop on Discovery Informatics, July 2014
  – ACM SIGKDD Workshop on Discovery Informatics, August 2014
  – PSB Workshop on Discovery Informatics, January 2015
Accelerating Science: A Grand Challenge for AI?

- All science is either stamp collecting or “physics”
- Big data = spectacular stamp collections!
- Big data ≠ Demise of the scientific method
- Accelerating science presents a grand challenge for AI:
  - Analysis and synthesis of computational abstractions of both
    - Universes of scientific discourse
    - Scientific artifacts and scientific process
  - Cognitive tools that augment and extend human intellect
  - Collaborative human-machine infrastructure for science
- Accelerating science calls for
  - **Foundational** advances within and across virtually all subfields of Artificial Intelligence
  - Concomitant advances in collaborative data and computing infrastructure
Big Data: Challenges and Opportunities

Omics

Human Sensors

Digital Media

Health Care

Source: Keith Marzullo
Big Data

Opportunities offered by big data are real

- Understanding the structure and dynamics of complex systems – cells, brains, individuals, organizations, societies
- Improving population health
- Anticipating and responding to crises
- Personalizing teaching and learning
- Defending critical infrastructure and services
- Making better decisions, e.g., public policy
- Making cities and communities smarter
- Improving food, energy, and water security
- ….
Big data = the end of the scientific method?


- “Petabytes allow us to say: “Correlation is enough.” We can stop looking for models. We can analyze the data without hypotheses about what it might show. We can throw the numbers into the biggest computing clusters the world has ever seen and let statistical algorithms find patterns where science cannot.”
- “Correlation supersedes causation, and science can advance even without coherent models, unified theories, or really any mechanistic explanation at all.”
- Most machine learning and data mining algorithms are essentially sophisticated ways of finding correlations from data
Perils of fishing for wisdom in oceans of data
Does cancer cause cell phone use?

Another huge study found no evidence that cell phones cause cancer. What was the WHO thinking?

Huh?

Well, take a look.

I think they just got it backward.

United States:

Total cancer incidence

Cell phone users

Per 100,000


100

75

50

25

You're not... There are so many problems with that.

Just to be safe, until I see more data I'm going to assume cancer causes cell phones.
Perils of fishing for wisdom in oceans of data
Fight global warming! Become a pirate!

- Big data ≠ End of theory!
- Correlation ≠ Causation!
Perils of fishing for wisdom in oceans of data
Eliminate science funding to save lives!

US spending on science, space, and technology
Millions of today's dollars (US OMB)

<table>
<thead>
<tr>
<th>Year</th>
<th>1999</th>
<th>2000</th>
<th>2001</th>
<th>2002</th>
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<td>19,753</td>
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<td>23,584</td>
<td>25,525</td>
<td>27,731</td>
<td>29,449</td>
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</table>

Suicides by hanging, strangulation and suffocation
Deaths (US) (CDC)

<table>
<thead>
<tr>
<th>Year</th>
<th>1999</th>
<th>2000</th>
<th>2001</th>
<th>2002</th>
<th>2003</th>
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<td>Value</td>
<td>5,427</td>
<td>5,688</td>
<td>6,198</td>
<td>6,462</td>
<td>6,635</td>
<td>7,336</td>
<td>7,248</td>
<td>7,491</td>
<td>8,161</td>
<td>8,578</td>
<td>9,000</td>
</tr>
</tbody>
</table>

Correlation: 0.992082
Perils of fishing for wisdom in oceans of data

• Sally Clark’s 1st son died in 1996, due to SIDS
• Her 2nd son died in 1999, also as a result of SIDS
• Prosecutors charged Sally Clark for murder on the grounds that both deaths were too unlikely to be due to SIDS
• Rationale:
  – One in 8,543 infant deaths is due to SIDS.
  – So chance of 2 deaths = $1/ (8,543^2)$ (or 1 in 73 million)
• – What’s wrong with this?
Data acquisition no longer the rate limiting step in science

- “We are close to having a $1,000 genome sequence, but this may be accompanied by a $1,000,000 interpretation”
- >1300 NAR gene databases
- 1M new biomedical journal articles published per year (2700/day)

1Bruce Korf, Former President, American College of Medical Genetics

Slide courtesy Larry Hunter
Many aspects of data management and analytics no longer the rate limiting steps in science

Most of the recent advances and efforts are focused on:

• **Data Management**
  – Organizing
  – Indexing
  – Integrating
  – Storing
  – Querying

• **Data Analytics**
  – Machine learning
  – Scaling up
  – High dimensionality
  – Heterogeneity
Big Data = the end of the scientific method? A lesson from Physics

Transformation of physics from a descriptive science (pre Newton) into a predictive science (post Newton)

• Tycho Brahe gathered 20 years of extremely accurate astronomical measurements: positions of the stars and planets: **big data**

• Johannes Kepler, working for Tycho Brahe, fit the data in every way imaginable to discover laws of planetary motion: **big data analytics**

• Isaac Newton’s invention of calculus provided the language to express, analyze, and communicate the unified laws of motion: **knowledge representation for physics**

• Big data did not make obsolete the scientific method then, and it does not do so now!
Big Data ≠ The end of the scientific method!

- Automation of “Big data” acquisition, management and analytics accelerates
  - Brahe’s part of the scientific endeavor (data acquisition, management)
  - And thanks to advances in machine learning increasingly, Kepler’s part (data analytics and model building)
  - But for the most part, leaves untouched, the other aspects of science, which become the rate limiting steps in science

➢ Accelerating science in the era of big data requires accelerating the rate-limiting steps of the scientific method!
How close are we to fully automating science?


Vasant Honavar, AAAI Fall Symposium on Accelerating Science: A Grand Challenge for AI, 2016
How close are we to fully automating science?

Science 3 April 2009: Vol. 324 no. 5923 pp. 85–89

The Automation of Science

Ross D. King, Jem Rowland, Stephen G. Oliver, Michael Young, Wayne Aubrey, Emma Byrne, Maria Liakata, Magdalena Markham, Pinar Pir, Larisa N. Soldatova, Andrew Sparkes, Kenneth E. Whelan and Amanda Clare

ABSTRACT

The basis of science is the hypothetico-deductive method and the recording of experiments in sufficient detail to enable reproducibility. We report the development of Robot Scientist “Adam,” which advances the automation of both. Adam has autonomously generated functional genomics hypotheses about the yeast Saccharomyces cerevisiae and experimentally tested these hypotheses by using laboratory automation. We have confirmed Adam's conclusions through manual experiments. ...
How close are we to fully automating science?

- Not very, except perhaps in very carefully constrained settings
- Collaborative human-machine systems might offer a more realistic approach to accelerating science
- Accelerating science presents a grand challenge for AI:
  - Analysis and synthesis of computational abstractions of both
    - Universes of scientific discourse
    - Scientific artifacts and the scientific process
  - Cognitive tools that augment and extend human intellect
  - Collaborative human-machine infrastructure for science
Computational abstractions of the universes of scientific discourse

• Church-Turing Thesis: Anything that can be described can be described by a computer program
• In any domain of scientific discourse, we need computational abstractions that describe objects, their properties, inter-relationships
Example: Computational abstractions of bio-molecular networks

- Genes a, b, c, d code for proteins A, B, C, D
- Proteins A and B form a hetero-dimer that activates the expression of gene c
- Protein C inhibits the expression of (and co-regulates) genes b and d
- Protein D is necessary for the transcription of protein B
Example: Undirected Graphs as abstractions of biomolecular interaction networks

- Protein-protein interaction networks
  - Nodes represent proteins
  - Edges represent interactions e.g., protein \( a \) binds to protein \( b \)
  - Topological analysis reveals functional roles
  - Connected components suggest complexes or pathways
  - Comparative analyses (across species, tissues, etc.) reveal shared sub-networks
Directed labeled graphs as abstractions of biomolecular interaction networks

- Nodes correspond to genes
- Edges correspond to regulatory interactions
- Edge labels can be used to denote the types of interactions, or lists of regulators and their influence on the specific edge, e.g., KEGG pathways
- Can help uncover sequences of regulatory events, cycles (feedback regulation), redundancy...
Boolean networks as abstractions of bio-molecular interaction networks

- Genes are modeled by binary variables - on, off (1, 0)
- States of genes are updated in discrete time steps
- State of a gene at time $t+1$ is a Boolean function of the states at time $t$ of the genes that influence it
- An $N$ gene Boolean network can in principle be in one of $2^N$ states
- Can help determine if the network can get from one state to another, the effect of gene knockout, etc.
Differential equations as abstractions of bio-molecular interaction networks

Considering only the mRNA abundances $a, b, c, d$

\[
\frac{da}{dt} = f_a(a) \quad \frac{db}{dt} = f_b(b, c, d) \\
\frac{dc}{dt} = f_c(a, b, c) \quad \frac{dd}{dt} = f_d(c, d)
\]

Differential equations can provide detailed information about kinetics.
Accelerating science requires computational abstractions of the universes of scientific discourse

- Church-Turing Thesis: Anything that can be described can be described by a computer program
- In any domain of scientific discourse, we need computational abstractions that describe objects, their properties, inter-relationships, in domains of scientific discourse
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Computational abstractions of scientific artifacts and scientific processes

Examples of scientific artifacts
- Experimental protocols
- Data, metadata, provenance
- Assumptions
- Conjectures
- Hypotheses
- Analysis tools
- Findings
- Arguments
- Models
- Explanations
- Theories
- Workflows

Examples of scientific processes
- Designing, prioritizing, planning, executing, documenting, replicating experiments
- Acquiring and organizing data
- Building, evaluating, linking models
- Generating and ranking conjectures
- Generating and testing hypotheses
- Testing, refining, comparing theories
- Producing and ranking explanations
- Sharing data and other artifacts
Computational abstractions of model construction

- Models can be built from knowledge (using inference, e.g., abstraction, specialization), observations (machine learning), experiments (e.g., causal inference).
- The use of off the shelf machine learning methods (SVM, DNN, Bayesian Networks, Stochastic grammars, etc.) introduces a language gap between model builders and model users.
- Need principled and generalizable approaches to
  - Construct and refine models that are
    - Accurate yet comprehensible
    - Explanatory
    - Communicable
    - Consistent with accepted background knowledge (e.g., laws of physics)
    - Lead to testable hypotheses
    - Models that span multiple levels of abstraction and scale
  - Assessing models with respect to not only predictive accuracy but also
    - Explanatory power
    - Coherence with models at higher and lower levels of abstraction
    - Simplicity …
How close are we to fully automating science?

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Cognitive tools for scientists that augment and extend the human intellect

- Acquiring, organizing and maintaining background “knowledge”
- Assessing and finding gaps in scientific knowledge
- Formulating and prioritizing questions
- Formulating, prioritizing, planning, and documenting studies
- Designing, prioritizing, planning, executing, monitoring experiments
- Drawing inferences, constructing explanations and hypotheses
- Synthesizing findings from disparate observational and experimental studies
- Building accurate, communicable, testable models from knowledge, observations and experiments
- Linking data, models, scientific arguments, hypotheses, experiments
- Linking and reasoning with models at different levels of abstraction or across different facets
- Sharing data, models, hypotheses, and other scientific artifacts
- Integrating results into the larger body of knowledge
A representative cognitive tool for scientists

A scientist’s associate that

• Learns what you and others in your field and related fields are working on
• Finds and reads relevant literature
• Locates and ingests available knowledge and data
• Offers assistance
  – Here are some data that contradict your hypothesis
  – Here are arguments for and against your hypothesis
  – Here is some data from lab X that explains your finding
  – Here is why you should prefer model A to model B
Collaborative human-machine infrastructure for science

- Distributed collaboratories that support:
  - Sharable and communicable representations of scientific artifacts
  - Data and computational resources
- Organizational structures and processes for collaboration
  - Assembling teams
  - Prioritizing, assigning and scheduling tasks
  - Decomposing tasks, combining results
  - Incentivizing and engaging participants
  - Organizing citizen science
Accelerating science: A grand challenge for AI?

- Accelerating science calls for synergistic advances across multiple areas of AI (and computing)
  - Knowledge representation and inference
    - How to represent and reason about computational abstractions of scientific domains, scientific artifacts, and scientific processes?
  - Planning and robotics
    - How to design, plan, execute, monitor, experiments?
  - Machine learning and causal inference
    - How to build accurate, comprehensible, predictive, explanatory or causal models from knowledge, observations, and experiments?
  - Computer supported collaborative work
    - How to optimally organize and incentivize scientific collaborative teams?
  - Data and computational infrastructure for science
    - How to share and reuse scientific artifacts at scale?
Discussion