

# CS 6474/CS 4803 Social Computing: Prediction & Forecasting II

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Examples of many  
successes...

# Incomplete history of cascade prediction

Who	Predicting	Features	Metric	Conclusion
HongD 10	Is item retweeted?	Topic Models	F1=0.47	Better than baseline
JendersKN 13	Will item reach some size $T$ ?	Content	F1>0.9	High accuracy
TanLP 14	Which of two does better?	Wording	Accu=65.6%	Computers are OK
ChengADKL 14	Will cascade double?	Temporal	AUC=0.88	Predictable
Lerman, Yang, Petrovic, Romero, Kupavskii, Ma, Weng, Zhao, Yu, etc				

# Progress?

All of this work examines a different **question** with a different **measure of success**, evaluated on a different subset of **data**, making it difficult to assess **overall progress**<sup>1</sup>

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<sup>1</sup><http://hunch.net/?p=22>

# Predicting success on Twitter?

Bakshy, Hofman,  
Mason, Watts (2011):  
How viral will my  
tweet be?

“Cascades are  
unpredictable!”



Mason Porter @masonporter · Jan 19

I took a brief break from work. :)



2



Reasons behind the  
inconsistencies



# Meaningless comparisons lead to false optimism in medical machine learning

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## Abstract

A new trend in medicine is the use of algorithms to analyze big datasets, e.g. using everything your phone measures about you for diagnostics or monitoring. However, these algorithms are commonly compared against weak baselines, which may contribute to excessive optimism. To assess how well an algorithm works, scientists typically ask how well its output correlates with medically assigned scores. Here we perform a meta-analysis to quantify how the literature evaluates their algorithms for monitoring mental wellbeing. We find that the bulk of the literature ( $\sim 77\%$ ) uses meaningless comparisons that ignore patient baseline state. For example, having an algorithm that uses phone data to diagnose mood disorders would be useful. However, it is possible to over 80% of the variance of some mood measures in the population by simply guessing that each patient has their own average mood - the patient-specific baseline. Thus, an algorithm that just predicts that our mood is like it usually is can explain the majority of variance, but is, obviously, entirely useless. Comparing to the wrong (population) baseline has a massive effect on the perceived quality of algorithms and produces baseless optimism in the field. To solve this problem we propose “user lift” that reduces these systematic errors in the evaluation of personalized medical monitoring.

Exploring limits to  
prediction in complex  
social systems



# A unified framework: Luck vs. skill<sup>2</sup>

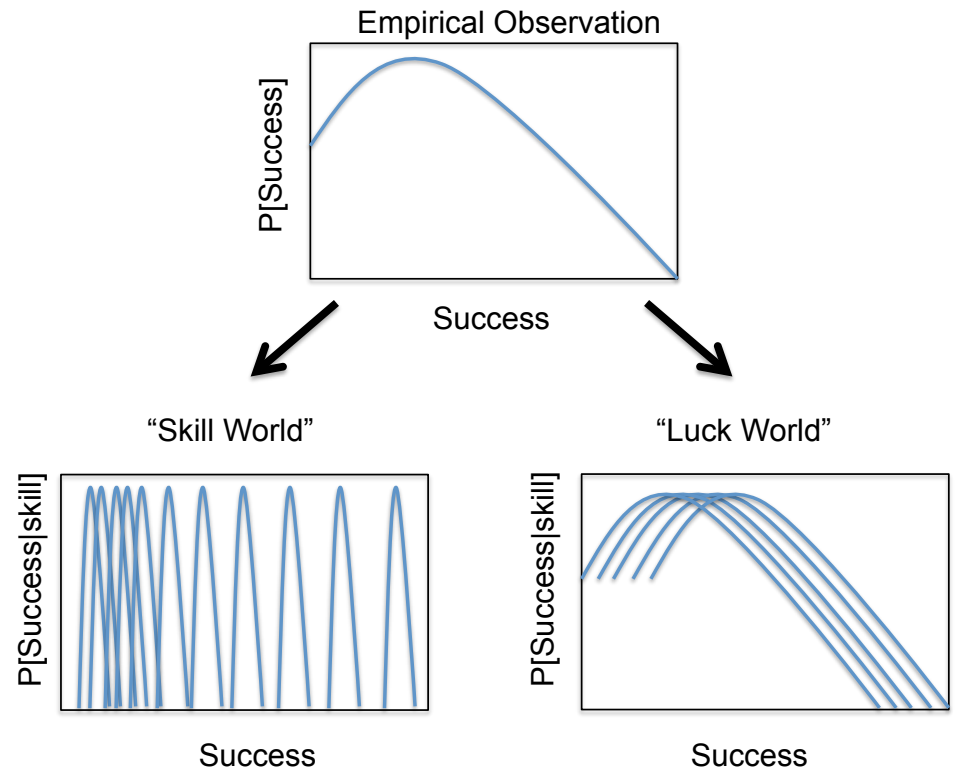
- Model success  $S$  as a mix of skill  $Q$  and luck  $\epsilon$ :

$$S = f(Q) + \epsilon$$

- Measure the fraction of variance remaining after conditioning on skill:

$$F = \frac{\mathbb{E}[\text{Var}(S|Q)]}{\text{Var}(S)} = 1 - R^2$$

- $R^2 = 1$  in a pure skill world,  
 $R^2 = 0$  in pure luck world



<sup>2</sup>Formalizes Maboussin (2012)

# Data

- Examined all 1.4B tweets containing URLs posted in February 2015
- Eliminated spam using internal Microsoft classifier
- Restricted attention to tweets containing URLs from the top 100 English-speaking domains with the most unique adopters
- Resulted in 850M tweets from 50M distinct users covering news, entertainment, videos, images, and products
- Measured the total cascade size for each seed tweet

# Predictive features

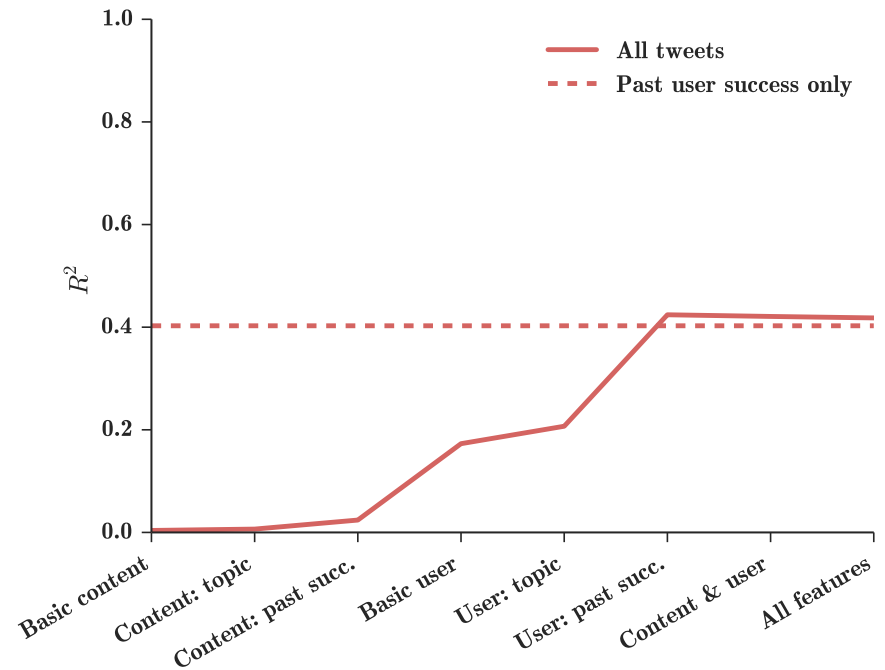
Used a random forest to estimate success (cascade size)  
given skill (available features)

- **Basic content features:** URL domain, time of tweet, spam score, ODP category
- **Basic user features:** number of followers, number of friends, number of posts, account creation time
- **Topic features:** the most probable Latent Dirichlet Allocation topic for each user and tweet, along with an interaction term
- **Past success:** the average number of retweets received by each URL and user in the past

# Predictive performance

best model explains roughly half of the variance in outcomes

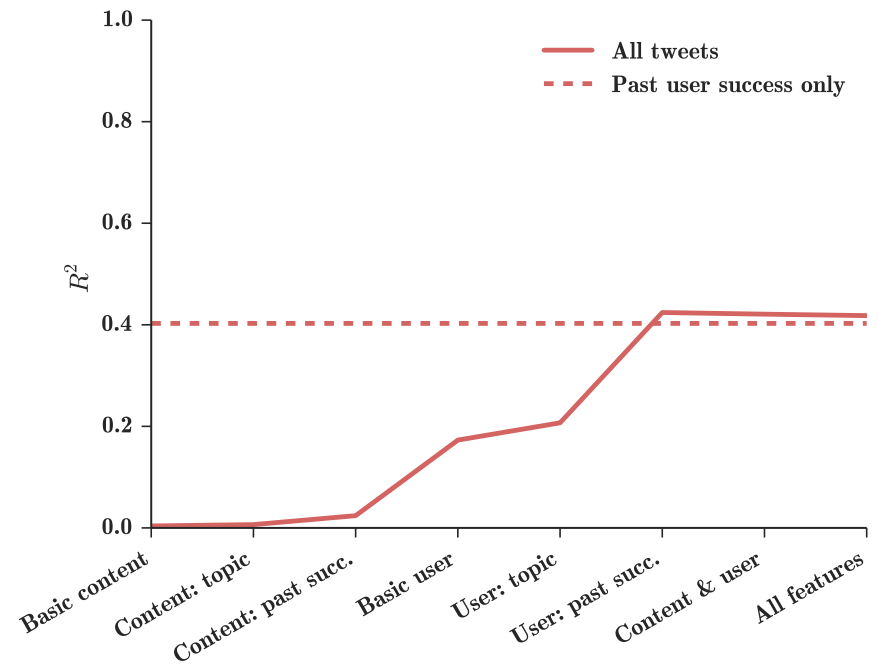
Model	<i>Tweet time</i>	<i>Domain</i>	<i>Spam score</i>	<i>Category</i>	<i>Tweet topic</i>	<i>Past url success</i>	<i>User time</i>	<i>Followers</i>	<i>Friends</i>	<i>Statuses</i>	<i>User topic</i>	<i>Past user success</i>	<i>Topic interaction</i>
1. Basic content	✓	✓	✓	✓									
2. Content, topic	✓	✓	✓	✓	✓								
3. Content, past succ.	✓	✓	✓	✓	✓	✓							
4. Basic user							✓	✓	✓	✓			
5. User, topic							✓	✓	✓	✓	✓		
6. User, past succ.							✓	✓	✓	✓	✓	✓	
7. Content, user	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓
8. All	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓



# Predictive performance

Content features alone perform poorly

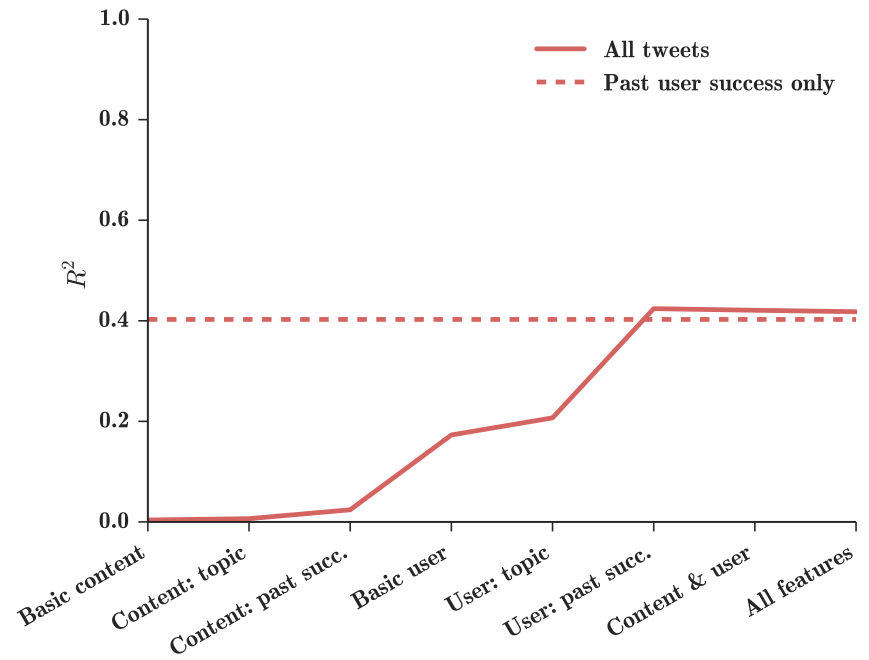
Model	<i>Tweet time</i>	<i>Domain</i>	<i>Spam score</i>	<i>Category</i>	<i>Tweet topic</i>	<i>Past url success</i>	<i>User time</i>	<i>Followers</i>	<i>Friends</i>	<i>Statuses</i>	<i>User topic</i>	<i>Past user success</i>	<i>Topic interaction</i>
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3. Content, past succ.	✓	✓	✓	✓	✓	✓							
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5. User, topic							✓	✓	✓	✓	✓		
6. User, past succ.							✓	✓	✓	✓	✓	✓	
7. Content, user	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	
8. All	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓



# Predictive performance

Basic user features provide a reasonable boost in performance

Model	<i>Tweet time</i>	<i>Domain</i>	<i>Spam score</i>	<i>Category</i>	<i>Tweet topic</i>	<i>Past url success</i>	<i>User time</i>	<i>Followers</i>	<i>Friends</i>	<i>Statuses</i>	<i>User topic</i>	<i>Past user success</i>	<i>Topic interaction</i>
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2. Content, topic	✓	✓	✓	✓	✓								
3. Content, past succ.	✓	✓	✓	✓	✓	✓							
4. Basic user							✓	✓	✓	✓			
5. User, topic							✓	✓	✓	✓	✓		
6. User, past succ.							✓	✓	✓	✓	✓	✓	
7. Content, user	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	
8. All	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓

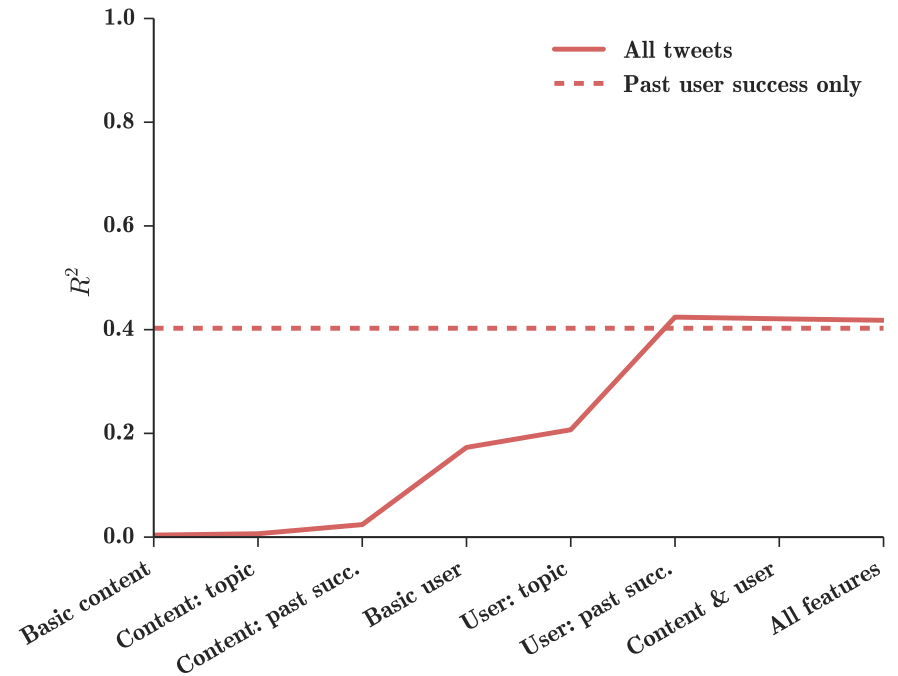




# Predictive performance

Past user success alone accounts for almost all of predictive power

Model	<i>Tweet time</i>	<i>Domain</i>	<i>Spam score</i>	<i>Category</i>	<i>Tweet topic</i>	<i>Past url success</i>	<i>User time</i>	<i>Followers</i>	<i>Friends</i>	<i>Statuses</i>	<i>User topic</i>	<i>Past user success</i>	<i>Topic interaction</i>
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3. Content, past succ.	✓	✓	✓	✓	✓	✓							
4. Basic user							✓	✓	✓	✓			
5. User, topic							✓	✓	✓	✓	✓		
6. User, past succ.							✓	✓	✓	✓	✓	✓	
7. Content, user	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	
8. All	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓



- Both models derive their predictive power from the same simple feature: a user's past success
- Content features are only weakly informative
- Performance plateaus as we add more features, suggesting a possible limit to the predictability of diffusion outcomes

# How can you *prove* a limit?

- Results robust to other ML models
  - Decision tree, linear regression
- Consistent with prior work
- Asymptote, dependency between features
- Can't rule everything out
  - Simulation

# Simulation

- SIR disease model
- Scale free network similar to Twitter
  - 7M users,  $\alpha = 2.05$
  - 8B simulated cascades
- *Quality*:  $R_0$  = average neighbors infected
  - $p(\text{infect over edge}) \times \text{mean-degree}$
- Prediction task
  - Given (possibly noisy) estimate of  $R_0$  and the seed node, predict cascade size

# Conclusions

Most things **don't spread**, but when they do, it's **difficult to predict success**

# Conclusions

Despite a great deal of research on the topic, it's difficult to assess long-term progress in predicting success



# Conclusions

State-of-the-art models explain roughly half of the variance in outcomes, based primarily on past success

# Conclusions

This is likely due to randomness in diffusion process itself, rather than our ability to estimate or model it

[nature](#) > [letters](#) > [article](#)


Published: 19 February 2009

# Detecting influenza epidemics using search engine query data

[Jeremy Ginsberg](#), [Matthew H. Mohebbi](#) , [Rajan S. Patel](#), [Lynnette Brammer](#), [Mark S. Smolinski](#) & [Larry Brilliant](#)

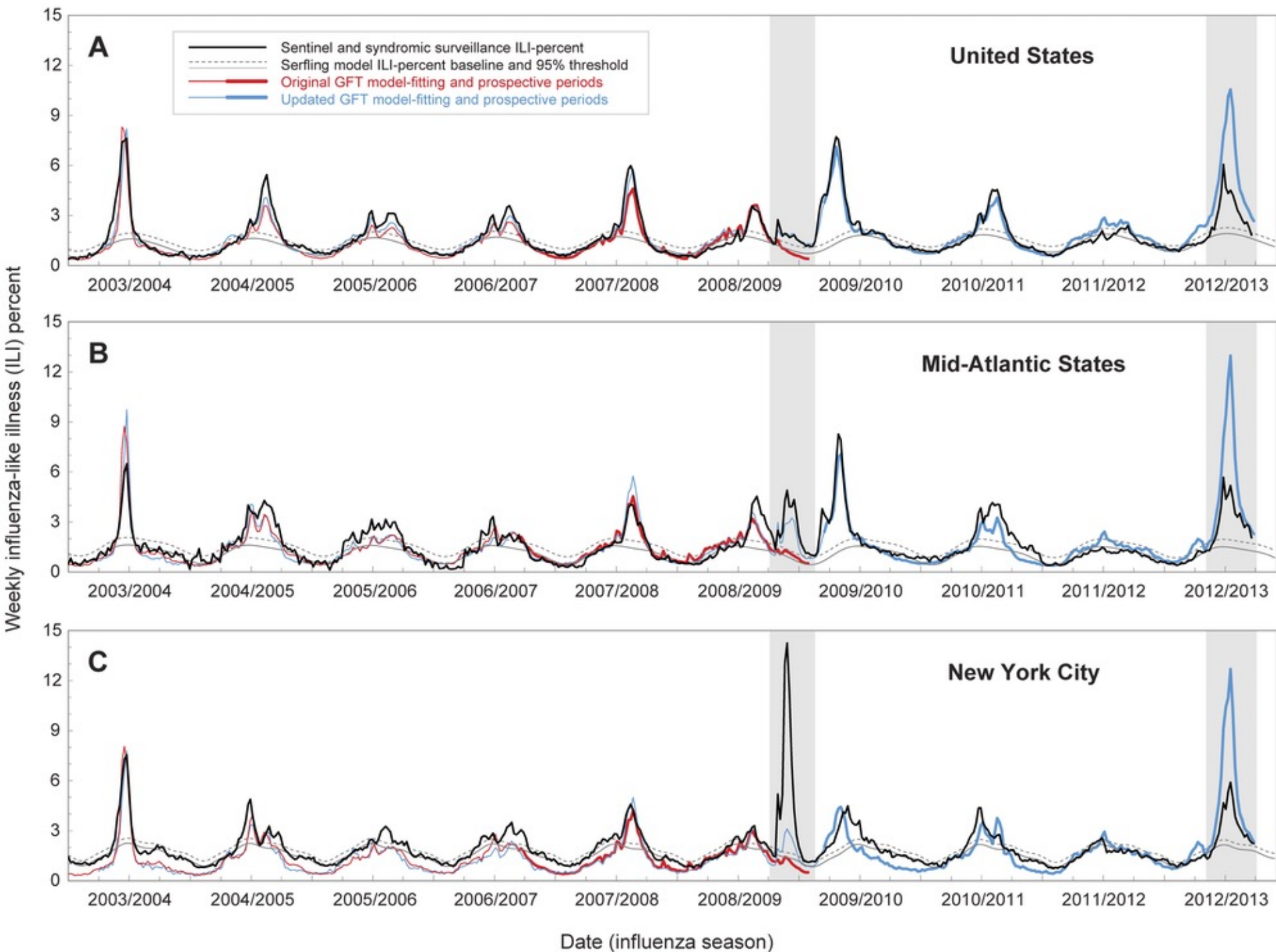
*Nature* **457**, 1012–1014(2009) | [Cite this article](#)

**16k** Accesses | **2217** Citations | **548** Altmetric | [Metrics](#)

 This article has been [updated](#)

## Abstract

Seasonal influenza epidemics are a major public health concern, causing tens of millions of respiratory illnesses and 250,000 to 500,000 deaths worldwide each year<sup>1</sup>. In addition to seasonal influenza, a new strain of influenza virus against which no previous immunity exists



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## RESEARCH ARTICLE



# Accurate estimation of influenza epidemics using Google search data via ARGO

Shihao Yang, Mauricio Santillana, and S. C. Kou

[+ See all authors and affiliations](#)

PNAS November 24, 2015 112 (47) 14473–14478; first published November 9, 2015;

<https://doi.org/10.1073/pnas.1515373112>

Edited by Wing Hung Wong, Stanford University, Stanford, CA, and approved September 30, 2015 (received for review August 6, 2015)

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## Significance

Big data generated from the Internet have great potential in tracking and predicting massive social activities. In this article, we focus on tracking influenza epidemics. We

## Reassessing Google Flu Trends Data for Detection of Seasonal and Pandemic Influenza: A Comparative Epidemiological Study at Three Geographic Scales

Donald R. Olson , Kevin J. Konty, Marc Paladini, Cecile Viboud, Lone Simonsen

Published: October 17, 2013 • <https://doi.org/10.1371/journal.pcbi.1003256>

Article	Authors	Metrics	Comments	Media Coverage
				

### Abstract

Author Summary

Introduction

Methods

Results

Discussion

Supporting Information

Acknowledgments

Author Contributions

References

### Abstract

The goal of influenza-like illness (ILI) surveillance is to determine the timing, location and magnitude of outbreaks by monitoring the frequency and progression of clinical case incidence. Advances in computational and information technology have allowed for automated collection of higher volumes of electronic data and more timely analyses than previously possible. Novel surveillance systems, including those based on internet search query data like Google Flu Trends (GFT), are being used as surrogates for clinically-based reporting of influenza-like-illness (ILI). We investigated the reliability of GFT during the last decade (2003 to 2013), and compared weekly public health surveillance with search query data to characterize the timing and intensity of seasonal and pandemic influenza at the national (United States), regional (Mid-Atlantic) and local (New York City) levels. We identified substantial flaws in the original and updated GFT models at all three geographic scales, including completely missing the first wave of the 2009 influenza A/H1N1 pandemic, and greatly overestimating the intensity of the A/H3N2 epidemic during the 2012/2013 season. These results were obtained for both the original (2008) and the updated (2009) GFT algorithms. The performance of both models was



The parable of google flu

Big data hubris

# Algorithmic Dynamics

*It's Not Just About  
Size of the Data*

# danah boyd & Kate Crawford

## CRITICAL QUESTIONS FOR BIG DATA

Provocations for a cultural,  
technological, and scholarly  
phenomenon

*The era of Big Data has begun. Computer scientists, physicists, economists, mathematicians, political scientists, bio-informaticists, sociologists, and other scholars are clamoring for access to the massive quantities of information produced by and about people, things, and their interactions. Diverse groups argue about the potential benefits and costs of analyzing genetic sequences, social media interactions, health records, phone logs, government records, and other digital traces left by people. Significant questions emerge. Will large-scale search data help us create better tools, services, and public goods? Or will it usher in a new wave of privacy incursions and invasive marketing? Will data analytics help us understand online communities and political movements? Or will it be used to track protesters and suppress speech? Will it transform how we study human communi-*

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## Deeper data: a response to boyd and Crawford

[Andre Brock](#)

First Published August 24, 2015 | Research Article



<https://doi.org/10.1177/0163443715594105>

[Article information](#) ▾

5



## Abstract

Data analysis of any sort is most effective when researchers first take account of the complex ideological processes underlying data's originating impetus, selection bias, and semiotic affordances of the information and communication technologies (ICTs) under examination.

## Keywords

[Big Data](#), [critical cultural informatics](#), [critical information studies](#), [data and society](#), [digital sociology](#), [social media and society](#)

In 2013, Lois Scheidt and I organized a panel for the International Congress of Qualitative Inquiry titled 'Small data in a big data world' as a response to 'Six Provocations for Big Data'. Our panelists presented incredible work conceptualizing new approaches in an age of 'big data' to qualitative social media research,



# Class Exercise

Assess what “small” or “deep” data, in each of the following cases might be considered.

- i) Predict how people react on a new product release (e.g., the latest version of iPhone), as observed on social media
- ii) Predict whether greater anonymity leads to greater hate speech on social media
- iii) Predict whether following recommended videos on YouTube leads down a more politically extreme rabbit hole
- iv) Predict whether deplatforming reduces misinformation on social media

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# Prediction and explanation in social systems

[JAKE M. HOFMAN](#), [AMIT SHARMA](#), AND [DUNCAN J. WATTS](#) [Authors Info & Affiliations](#)[SCIENCE](#) • 3 Feb 2017 • Vol 355, Issue 6324 • pp. 486-488 • DOI: [10.1126/science.aal3856](#)

788 143

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## Abstract

Historically, social scientists have sought out explanations of human and social phenomena that provide interpretable causal mechanisms, while often ignoring their predictive accuracy. We argue that the increasingly computational nature of social science is beginning to reverse this traditional bias against prediction; however, it has also highlighted three important issues that require resolution. First, current practices for evaluating predictions must be better standardized. Second, theoretical limits to predictive accuracy in complex social systems must be better

