

DORA and reinventing research assessment

EIFL, Istanbul, November, 2013 Mark Patterson, Executive Director, eLife

# **Outline**

- A few words about eLife
- Research assessment old and new approaches
- DORA recommendations



# Funders taking direct action





wellcometrust

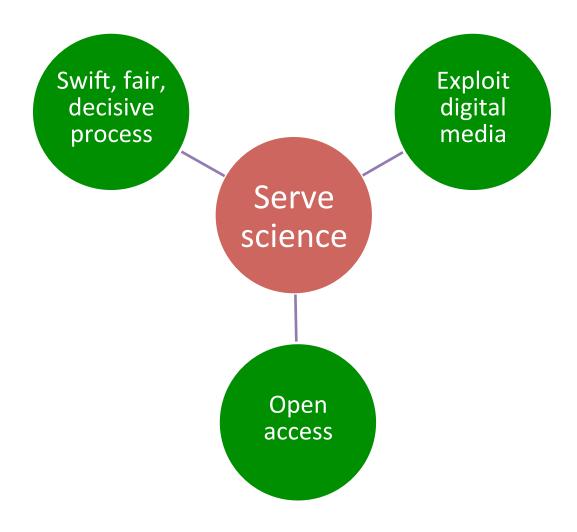






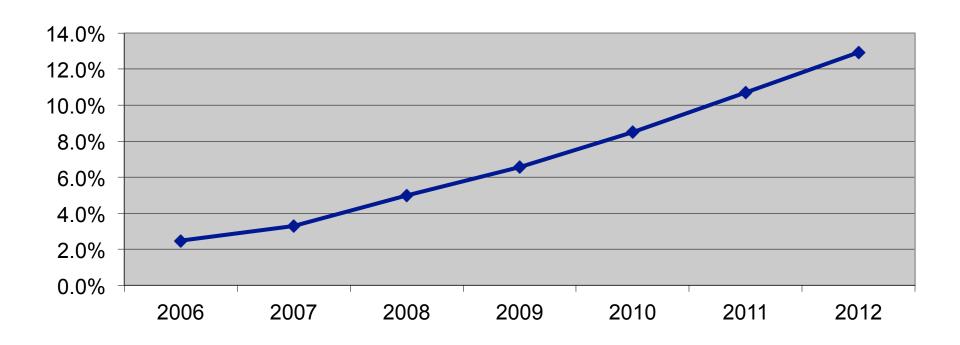


# **eLife:** motivations





# %PubMed available as open access in PMC



# Researchers (authors and readers)

#### **Institutions**





The public











**wellcome**trust

Policy makers



**Publishers** 







# What journal?



What impact factor?

The impact factor is...

- a journal-based metric
- proprietary
- incomplete

World politics

Business & finance Economics Science & technology Culture Blogs Debate Multimedia

Print edition

#### Unreliable research

# Trouble at the lab

Comment (196)

Print

E-mail

Reprints & permissions

Scientists like to think of science as self-correcting. To an alarming degree, it is not

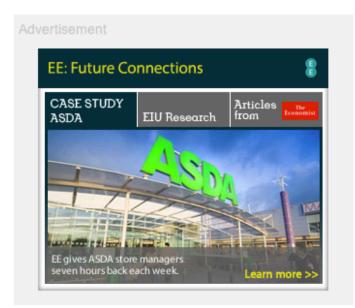
Oct 19th 2013 | From the print edition







"I SEE a train wreck looming," warned Daniel Kahneman, an eminent psychologist, in an open letter last year. The premonition concerned research on a phenomenon known as "priming". Priming studies suggest that decisions can be influenced by apparently irrelevant actions or events that took place just before the cusp of choice. They have been a boom area in psychology over the past decade, and some of their insights have already made it out of the lab and into the toolkits of policy wonks keen on "nudging" the populace.







s Zwarte Piet racism? ,230 people recommend this.

2,650 people recommend this.

About Submit Register Login

#### Search



### Statistical Genetics and Methodology

Home

About

Editorial Board

Archive

Research Topics

View Some Authors

Review Guidelines

Subscribe to Alerts

SEARCH



**AUTHOR INFO** 

Why Submit?

Fees

Article Types

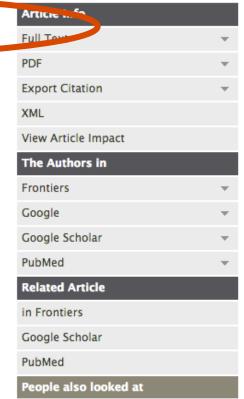


# The hidden factors in impact factors: a perspective from Brazilian science

Renata C. Ferreira, Fernando Antoneli and Marcelo R. S. Briones\*

Laboratório de Genômica Evolutiva e Biocomplexidade, Departamento de Microbiologia, Imunologia e Parasitologia, Universidade Federal de São Paulo, São Paulo, Brazil

Contrary to all current international recommendations on evaluation of academic achievement the evaluation of graduate programs in Brazil relies heavily on journal impact factors (Garfield, 2006; San Francisco Declaration on Research Assessment, 2013). The governmental agency CAPES from the Education Ministry monopolize this evaluation and pressure programs by the distribution of funding resources and departmental fellowships conditioned to adherence to a journal classification system called "Qualis" which is a discretization of the continuous distribution of journals ranking by their impact factors (Greenwood, 2007). In several institutions the graduate committee authorizes professors to act as thesis advisors only if in a certain period (e.g., 4 years) they publish at least one paper in a journal classified as "Qualis A2." This classification has seven categories with decreasing impact factor ranges (A1, A2, B1, B2, B3, B4, B5, and C.



Q,

#### Probabilistic inference of short-term synaptic plasticity in neocortical microcircuits

Rui P. Costa, P. Jesper Sjostrom and Mark C. W. van Rossum

MicroRNA discovery by similarity

#### **Citations**



## **Policy and practice**



#### Media

# theguardian

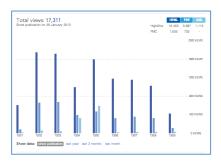
The New York Times







## Usage



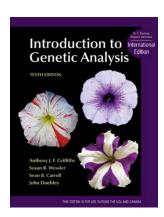


# 





## **Textbooks**













**Reference managers** 



Search

a

About Us

RESEARCH ARTICLE | FEATURED IN PLOS COLLECTIONS

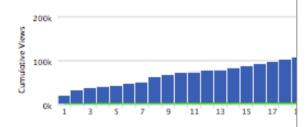
## How Many Species Are There on

Camilo Mora . Derek P. Tittensor, Sina Adl, Alastair G. B. Simo

Article	About the Authors	Metrics

#### Viewed







\*Although we update our data on a daily basis, there may be a 48 recent numbers are available. PMC data is posted on a monthly be Cited 0

Browse

For Authors











Saved





#### Discussed @

















Recommended @





Software ~

Research ~

Partners ~

Support ~

Q

PKP is a multi-university initiative developing (free) open source software and conducting research to improve the quality and reach of scholarly publishing.

Public Knowledge Project > News > PKP launches Article Level Metrics for OJS journals

# PKP launches Article Level Metrics for OJS journals

Published on October 9th, 2013 by Karen Meijer-Kline in News | 4 Comments

The Public Knowledge Project, developer of publishing software used by thousands of scholarly journals continues to be among those leading innovation in scholarly publishing with its introduction of article level metrics (ALMs).

The purpose of ALMs is to provide a more accurate means for measuring the impact of research at the article level rather than the journal level, as is tradition, said Juan Pablo Alperin of PKP. "ALMs have the potential to be a truly valuable tool for both publishers and readers. We urge all journals using our Open Journal Systems (OJS) to sign-up for the service." PKP's ALM offering is built on an open source application developed by the Public Library of Science (PLOS) and has been implemented with the guidance and support from PLOS. (See PLOS blog for their announcement)



Co-Action Publishing, a close partner of PKP and one of its first hosted clients in 2007, is the first publisher to implement the service. "Co-Action's wishes, requirements and feedback have been instrumental in the development and evolution of our publishing platform, and most recently our implementation of ALMs," Alperin said.

Co-Action Publishing Co-founder Caroline Sutton, who has been a leading voice in the OA community for many years, is gratified by the role Co-Action is playing in making a new generation of publishing tools available to OA publishers everywhere. "Together, we're working to develop many tools that can be used and leveraged across the thousands of journals that use OJS. We're excited about what's in the pipeline, and how these new developments will help further academic publishing."

# New metrics and indicators of scholarship

- From one measure to many
- From journal to article
- From one output to many

# San Francisco

# DARA

**Declaration on Research Assessment** 

- Recommendations for publishers, funders, institutions, metrics suppliers, and researchers
- >9000 signatories

# Three themes

- eliminate the use of impact factors in decisions about funding and careers
- opportunities provided by online publication (such as new indicators of significance and impact)
- assess research on its own merits

# Funding agencies and institutions

- be transparent about processes
- base judgement on outputs rather than journal names and metrics
- consider multiple outputs/outcomes
- consider multiple metrics and indicators

# Taking steps to improve assessment (1)

# National Cancer Institute

at the National Institutes of Health

"Varmus wants to replace a section that now lists major publications with a narrative describing the investigator's five major accomplishments."

Science, 25 October 2013

# Taking steps to improve assessment (2)

**Sandra L. Schmid**, chair of the Department of Cell Biology at the University of Texas Southwestern Medical Center.

"We believe we can recognize excellence that has been missed by journal editors."

Applicants for faculty positions provide a cover letter describing:

- (1) most significant scientific accomplishment as a graduate student
- (2) most significant scientific accomplishment as a postdoc
- (3) overall goals/vision for their research program
- (4) experience and qualifications that make them able to achieve those goals

# Taking steps to improve assessment (3)

No sub-panel will make any use of journal impact factors, rankings, lists or the perceived standing of publishers in assessing the quality of research outputs.

An underpinning principle of the REF is that all types of research and all forms of research outputs across all disciplines shall be assessed on a fair and equal basis.



# **Publishers**

- don't promote the impact factor (ideally)
- provide new metrics and indicators
- make reference lists available as open data
- promote responsible authorship and author contributions

Article

## Evolutionary principles of modular gene regulation in 8 yeasts

Dawn A Thompson , Sushmita Roy, Michelle Chan, Mark P Styczynsky, Jenna Pfiffner, Courtney French, Amanda Socha, Anne Thielke, Sara Napolitano, Paul Muller, Manolis Kellis, Jay H Konieczka, Ilan Wapinski, Aviv Regev

Broad Institute of MIT and Harvard, United States; Massachusetts Institute of Technology, United States; Howard Hughes Medical Institute, Massachusetts Institute of Technology, United States

DOI: http://dx.doi.org/10.7554/eLife.00603

Published June 18, 2013 Cite as eLife 2013;2:e00603

#### Abstract

Divergence in gene regulation can play a major role in evolution. Here, we used a phylogenetic framework to measure mRNA profiles in 15 yeast species from the phylum Ascomycota and reconstruct the evolution of their modular regulatory programs along a time course of growth on glucose over 300 million years. We found that modules have diverged proportionally to phylogenetic distance, with prominent changes in gene regulation accompanying changes in lifestyle and ploidy, especially in carbon metabolism. Paralogs have significantly contributed to regulatory divergence, typically within a very short window from their duplication. Paralogs from a whole genome duplication (WGD) event have a uniquely substantial contribution that extends over a longer span. Similar patterns occur when considering the evolution of the heat shock regulatory program measured in eight of the species, suggesting that these are general evolutionary principles.

DOI: http://dx.doi.org/10.7554/eLife.00603.001

#### 1+1 eLife digest

#### Main text

#### Introduction

Divergence in the regulation of gene expression has been repeatedly postulated to play a major



POF	SHILL	Ġ







Reference tools:

DOWNLOAD OPEN SHARE

Related content

#### Yeast rises to the occasion

MA Ragan

Insight — Genomics and evolutionary biology

#### Correction: Evolutionary principles of modular gene regulation in yeasts

DA Thompson, S Roy, M Chan, MP Styczynski, J Pfiffner, C French, A Socha, A Thielke, S Napolitano, P Muller, M Kellis, JH Konieczka, I Wapinski, A Regev

Correction — Genomics and evolutionary biology

#### Jump to:

#### Article

Abstract

eLife digest

Main text

Introduction

Results

Discussion

Materials and methods

References

Decision letter

Author response Leave a comment

Figures & data

Article

titute of Technology, United States; Howard

ted States



Dawn A Thompson , Sushmita Roy, Michelle Chan, Mark P Styczynsky, Jenna Pfiffner, Courtney French Amend Soche Anne Thielke Sere Napolitano, Paul Muller, Manolis

#### Sushmita Roy

Broad Institute of MIT and Harvard, Cambridge, United States; Computer Science and Artificial Intelligence Laboratory, Massachusetts Institute of Technology, Cambridge, United States

Present address: Biostatist University of Wisconsi Madison, United States

contribution: Conception and design, Analysis and interpretation of data

No competing interests declared

Contributed equally with: Dawn A Thompson

Here, we used a phylogenetic framework to measure mRNA profiles in 15 yeast species from the phylum Ascomycota and reconstruct the evolution of their modular regulatory programs along a time course of growth on glucose over 300 million years. We found that modules have diverged proportionally to phylogenetic distance, with prominent changes in gene regulation accompanying changes in lifestyle and ploidy, especially in carbon metabolism. Paralogs have significantly contributed to regulatory divergence, typically within a very short window from their duplication. Paralogs from a whole genome duplication (WGD) event have a uniquely substantial contribution that extends over a longer span. Similar patterns occur when considering the evolution of the heat shock regulatory program measured in eight of the species, suggesting that these are general evolutionary principles.

DOI: http://dx.doi.org/10.7554/eLife.00603.001

1+1 eLife digest

I-I Main text

Introduction

Divergence in the regulation of gene expression has been repeatedly postulated to play a major

View article with eLIFE Lens

POF	STREET, STREET,	<





Reference tools:

DOWNLOAD OPEN SHARE

Related content

#### Yeast rises to the occasion

MA Ragan

Insight — Genomics and evolutionary biology

#### Correction: Evolutionary principles of modular gene regulation in yeasts

DA Thompson, S Roy, M Chan, MP Styczynski, J Pfiffner, C French, A Socha, A Thielke, S Napolitano, P Muller, M Kellis, JH Konieczka, I Wapinski, A Regev

Correction — Genomics and evolutionary biology

Jump to:

#### Article

Abstract

eLife digest Main text

Introduction

Results

Discussion

Materials and methods

References

Decision letter Author response

Leave a comment

Figures & data

# Metrics service providers

- be transparent about methods
- make data open
- consider how to respond to gaming

# Metrics service providers









# Researchers

- challenge use impact factor in research assessment
- promote assessment on the basis of research content
- use new metrics/indicator to support your own career

# **Dr Stephen Robert Pettifer**

#### **Journal Articles**

- J1. A. J. Williams, L. Harland, P. Groth, S. Pettifer, C. Chichester, E. L. Willighagen, C. T. Evelo, N. Blomberg, G. Ecker, C. Goble, and B. Mons. Open PHACTS: Semantic interoperability for drug discovery. *Drug Discovery Today*, 2012.

  [CS:13,] Highly Viewed Highly Discussed
- J2. A. J. G. Gray, P. Groth, A. Loizon, S. Askjaer, C. Brenninkmeijer, K. Burger, C. Chichester, C. T. Evelo, C. Goble, L. Harland, S. Pettifer, M. Thompson, A. Waagmeester, and A. J. Williams. Applying linked data approaches to pharmacology: Architectural decisions and implementation. *Semantic Web Journal*, 2012. [0 citations]
- J3. B. Vroling, D. Thorne, P. McDermott, T. K. Attwood, G. Vriend, and S. Pettifer. Integrating GPCR-specific information with full text articles. *BMC Bioinformatics*, 12(362), Sep 2011. [GS:4] Highly Discussed .
- J4. B. Vroling, D. Thorne, P. McDermott, HJ. Joosten, T. K. Attwood, S. Pettifer, and G. Vriend. Nucleardb: information system for nuclear receptors. *Nucleic Acids Research*, Nov 2011. [GS:2 ] ❖
- J5. A. Brown, R. Stevens, and S. Pettifer. Making graph-based diagrams work in sound: the role of annotation. Human Computer Interaction, 2011. [GS:35]

# Summary

- Current methods are poor
- New approaches are possible
- And steadily, improvements are being made

# San Francisco DER RA

**Declaration on Research Assessment** 

- Signing DORA is a start, but what happens next?
- Further discussion amongst key stakeholders
- Identify specific actions
- Monitor and report progress

# Thank you

Mark Patterson m.patterson@elifesciences.org

