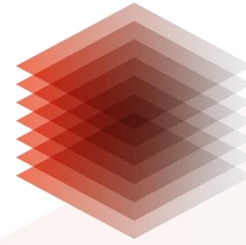
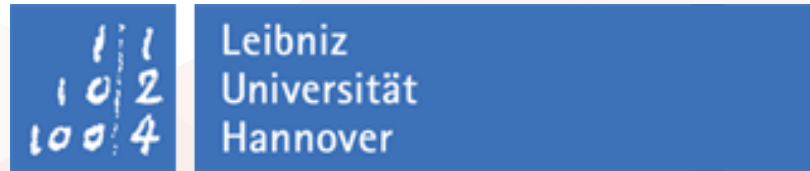


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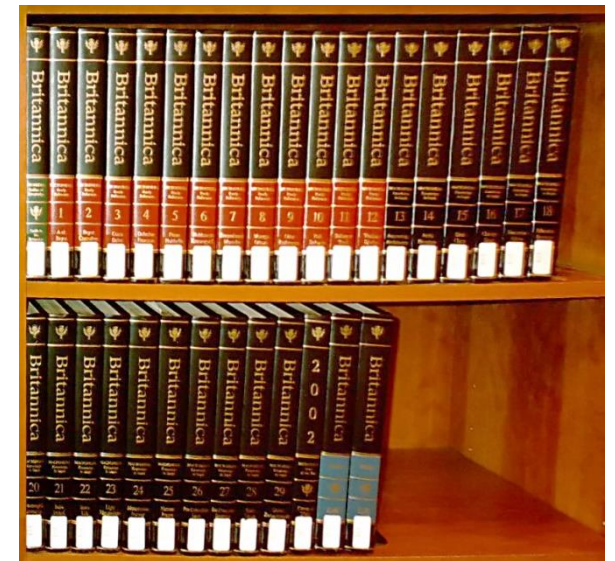
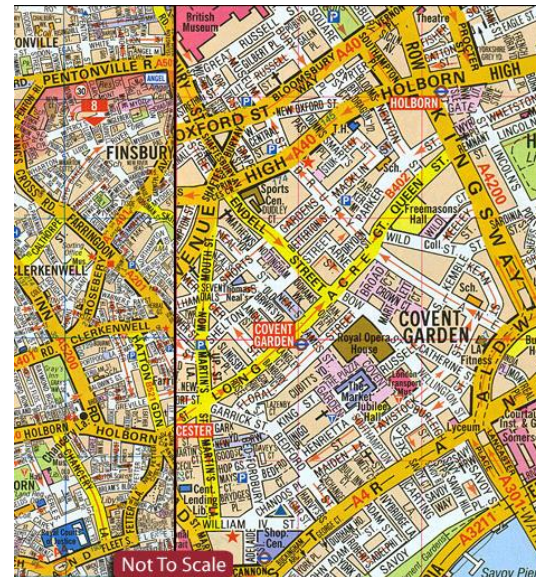
Organizing Scientific Contributions with the Open Research Knowledge Graph

Prof. Dr. Sören Auer

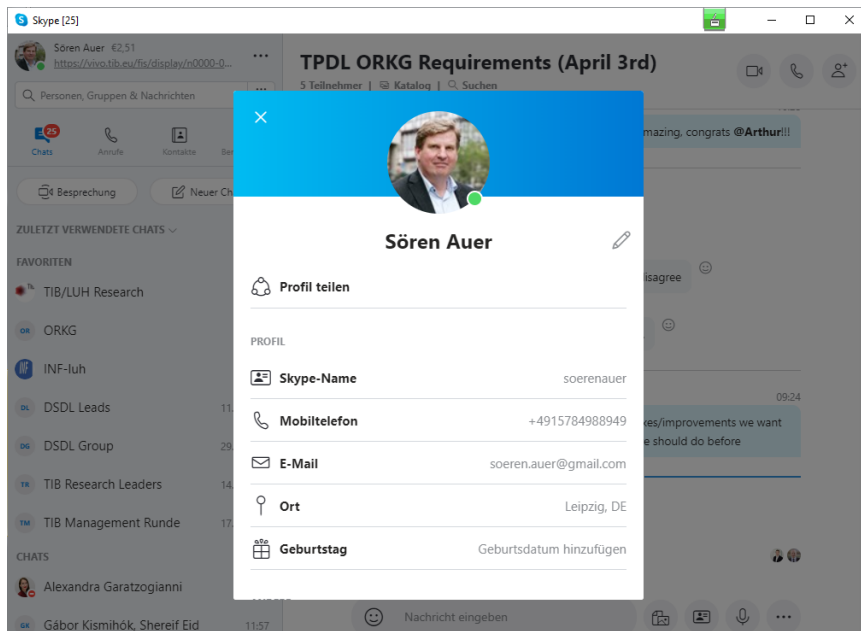
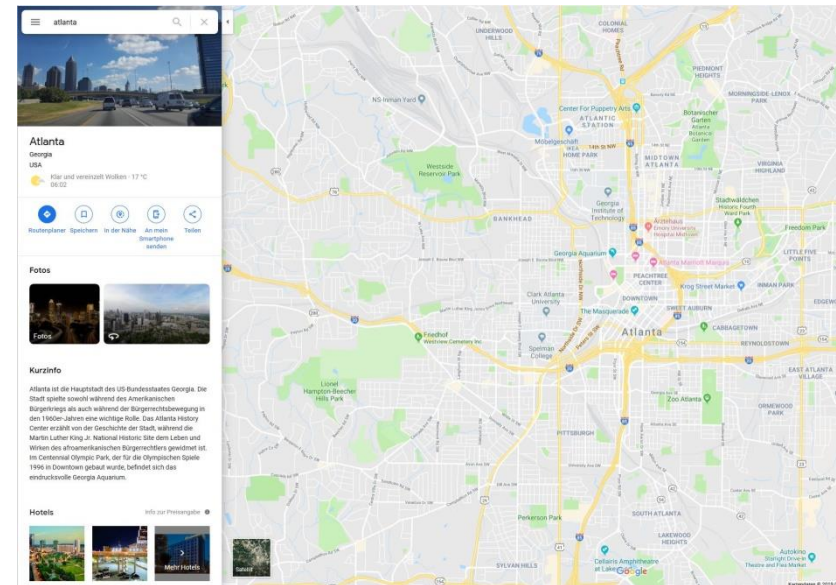
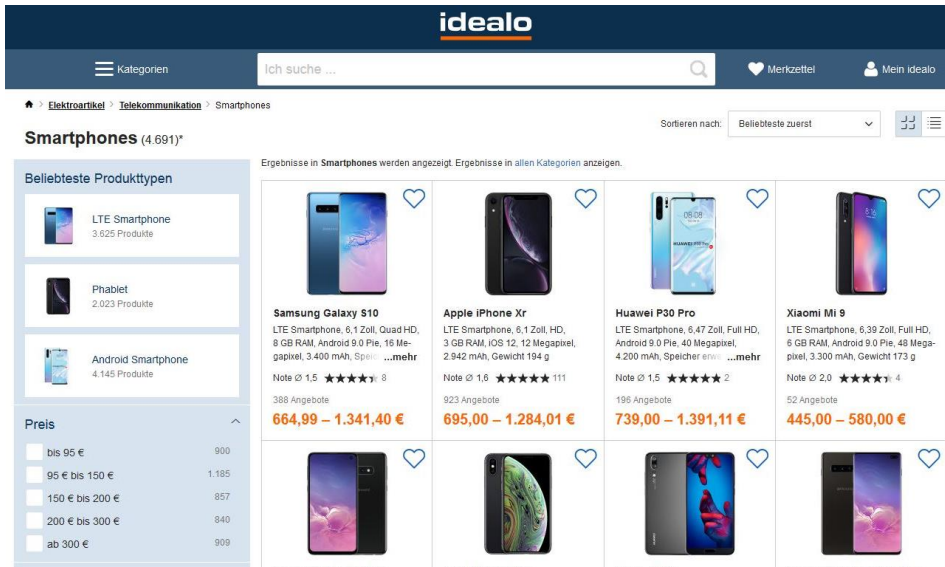
Leibniz University of Hannover

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How did information flows change in the digital era?



How does it work today?



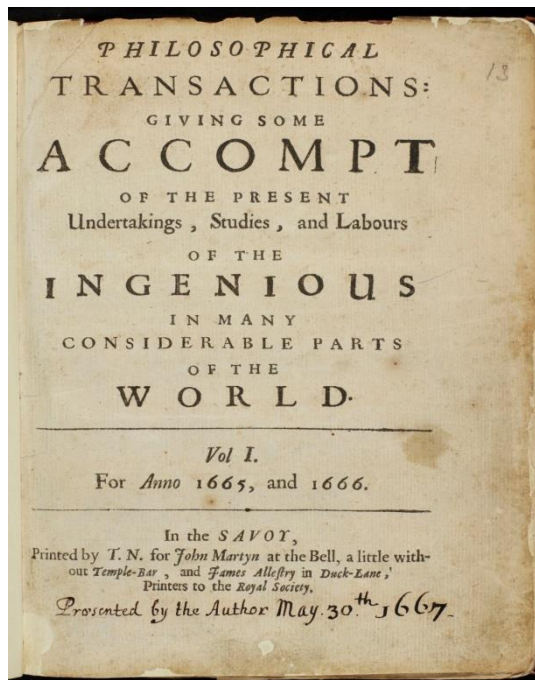
The World of Publishing & Communication has profoundly changed

- New means adapted to the new possibilities were developed, e.g. „zooming“, dynamics
- **Business models** changed completely
- More focus on data, interlinking of **data / services and search** in the data
- Integration, **crowdsourcing**, **data curation** play an important role

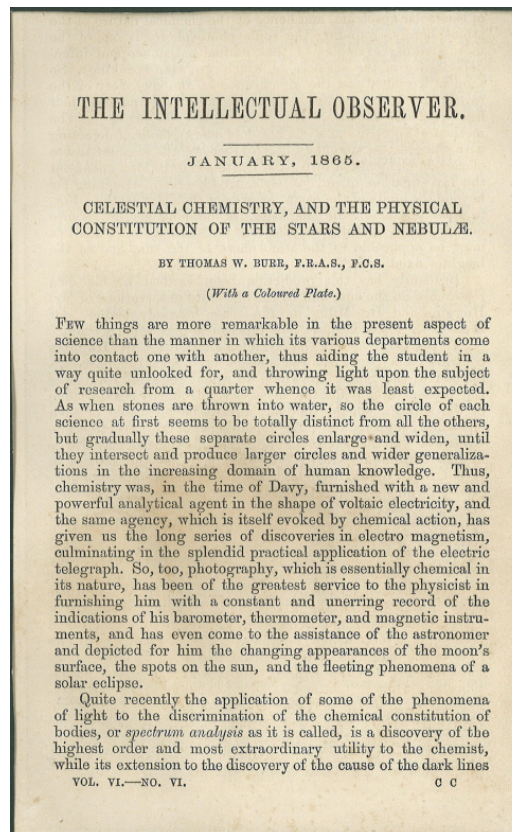
What about Scholarly Communication?

Scholarly Communication has not changed (much)

17th century



19th century



20th century



21th century



Meanwhile other information intense domains were completely disrupted: mail order catalogs, street maps, phone books, ...

We need to rethink the way how research is represented and communicated

Challenges we are facing:

Digitalisation of Science

- Data integration and analysis
- Digital collaboration

Monopolisation by commercial actors

- Publisher look-in effects
- Maximization of profits ^[1]

Reproducibility Crisis

- Majority of experiments are hard or not reproducible ^[2]

Proliferation of publications

- Publication output doubled within a decade
- continues to rise ^[3]

Deficiency of Peer Review

- Deteriorating quality ^[4]
- Predatory publishing

[1] <http://thecostofknowledge.com>, <https://www.projekt-deal.de>

[2] M. Baker: *1,500 scientists lift the lid on reproducibility*, *Nature*, 2016.

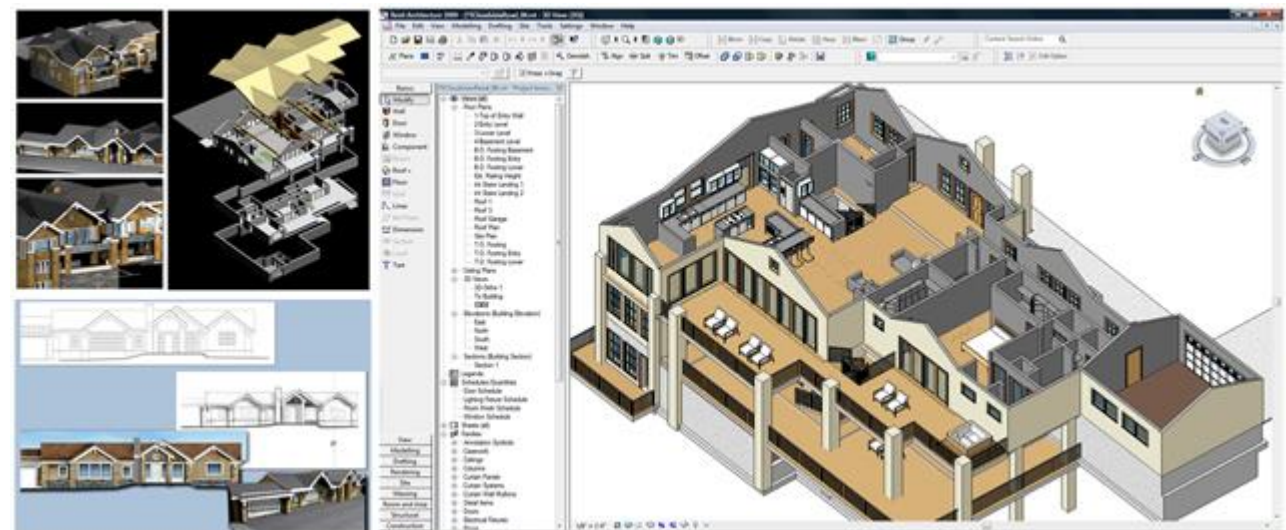
[3] *Science and Engineering Publication Output Trends*, National Science Foundation, 2018.

[4] J. Couzin-Frankel: *Secretive and Subjective, Peer Review Proves Resistant to Study*, *Science*, 2013.

Duplication and Inefficiency

How can we avoid duplication if the terminology, research problems, approaches, methods, characteristics, evaluations, ... are not properly defined and identified?

How would you build an engine / building without properly defining their parts, relationships, materials, characteristics ...?



Root Cause – Deficiency of Scholarly Communication?

Lack of...

Transparency

information is hidden
in text

Integratability

fitting different
research results
together

Machine assistance

unstructured content
is hard to process

Identifiability

of concepts beyond
metadata

Collaboration

one brain barrier

Overview

Scientists look for the
needle in the haystack

Ungefähr 238.000 Ergebnisse (0,04 Sek.)

Zeitraum wä

☆ 55 Zitiert von: 3650 Ähnliche Artikel Alle 19 Versionen Web of Science: 2423

**How can
we fix it?**

Concepts

Overarching Concepts

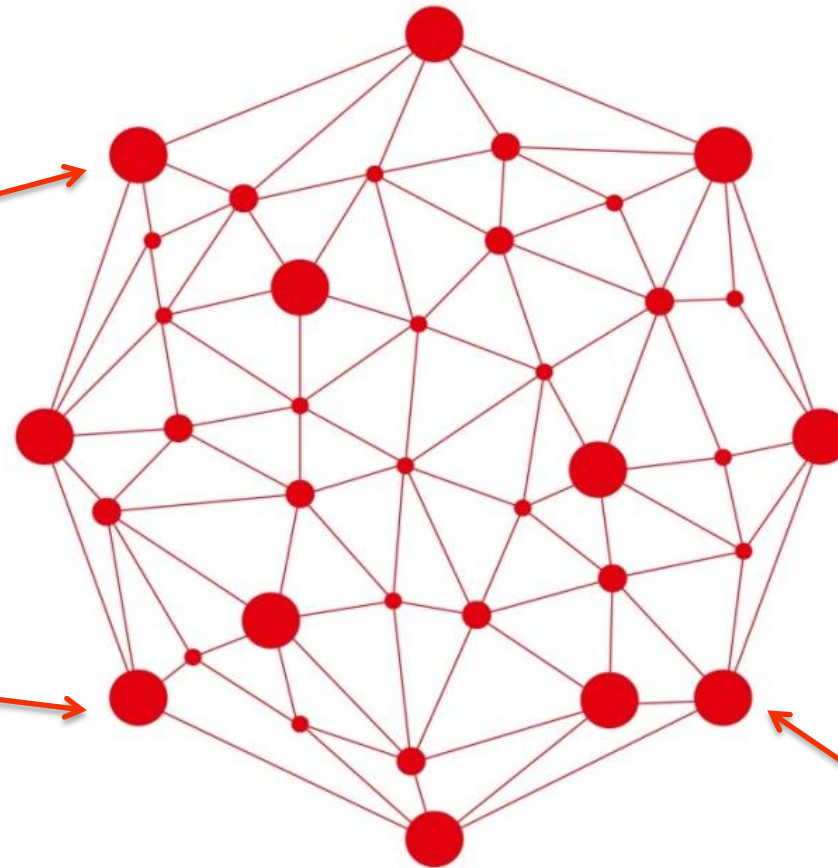
- Research problems
- Definitions
- Research approaches
- Methods

Artefacts

- Publications
- Data
- Software
- Image/Audio/Video
- Knowledge Graphs / Ontologies

Domain specific Concepts

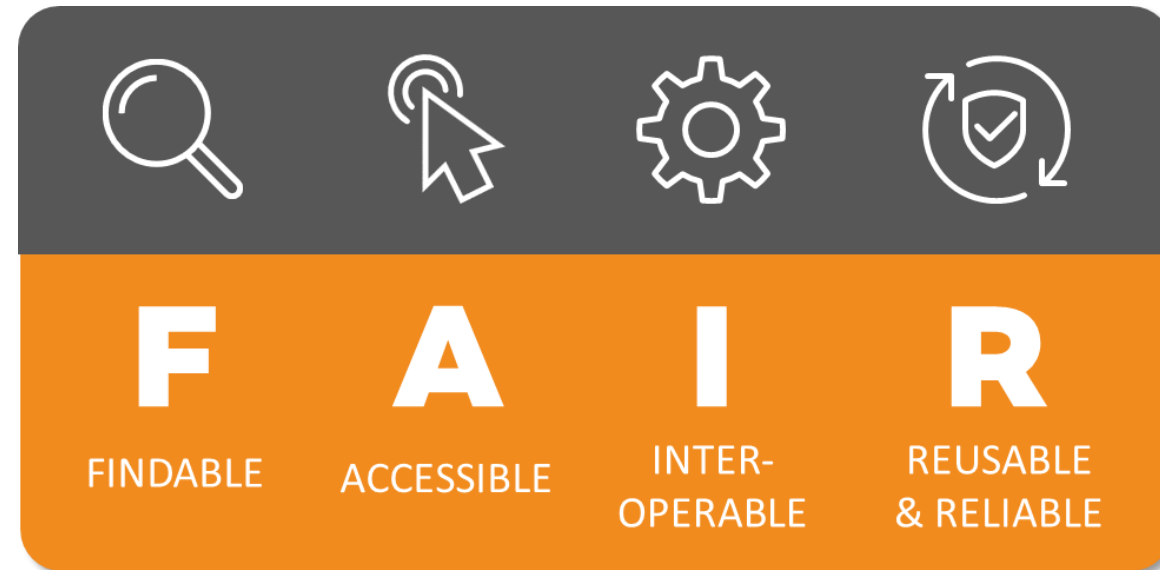
Mathematics	Physics	Chemistry	Computer Science	Technology	Architecture
• Definitions • Theorems • Proofs • Methods • ...	• Experiments • Data • Models • ...	• Substances • Structures • Reactions • ...	• Concepts • Implementations • Evaluations • ...	• Standards • Processes • Elements • Units, Sensor data	• Regulations • Elements • Models • ...



Knowledge Graphs built on FAIR and Linked Data Principles

1. Use **URIs** to **identify** the “things” in your data
2. Use **http:// URIs** so people (and machines) can **look** them **up** on the web
3. When a URI is looked up, **return** a **description** of the thing **in** the W3C **Resource Description Format (RDF)**
4. Include **links to related things**

<http://www.w3.org/DesignIssues/LinkedData.html>



Chemistry Example: CRISPR Genome Editing





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Advanced Search

New Results

Previous

Next

A practical guide to CRISPR/Cas9 genome editing in Lepidoptera

Linlin Zhang,  Robert Reed

doi: <https://doi.org/10.1101/130344>

Now published in *Diversity and Evolution of Butterfly Wing Patterns* doi: [10.1007/978-981-10-4956-9_8](https://doi.org/10.1007/978-981-10-4956-9_8)

Abstract

Info/History

Metrics

 Preview PDF

Abstract

CRISPR/Cas9 genome editing has revolutionized functional genetic work in many organisms and is having an especially strong impact in emerging model systems. Here we summarize recent advances in applying CRISPR/Cas9 methods in Lepidoptera, with a focus on providing practical advice on the entire process of genome editing from experimental design through to genotyping. We also describe successful targeted GFP

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Subject Area

Genetics

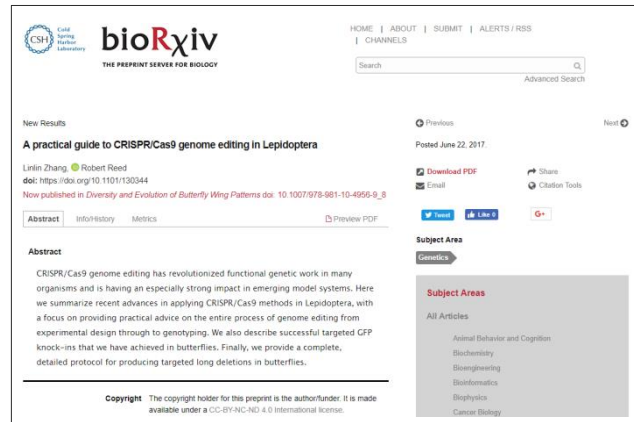
Subject Areas

All Articles

Animal Behavior and Cognition

Chemistry Example: Populating the Graph

2. Adaptive Graph Curation & Completion



Author Robert Reed

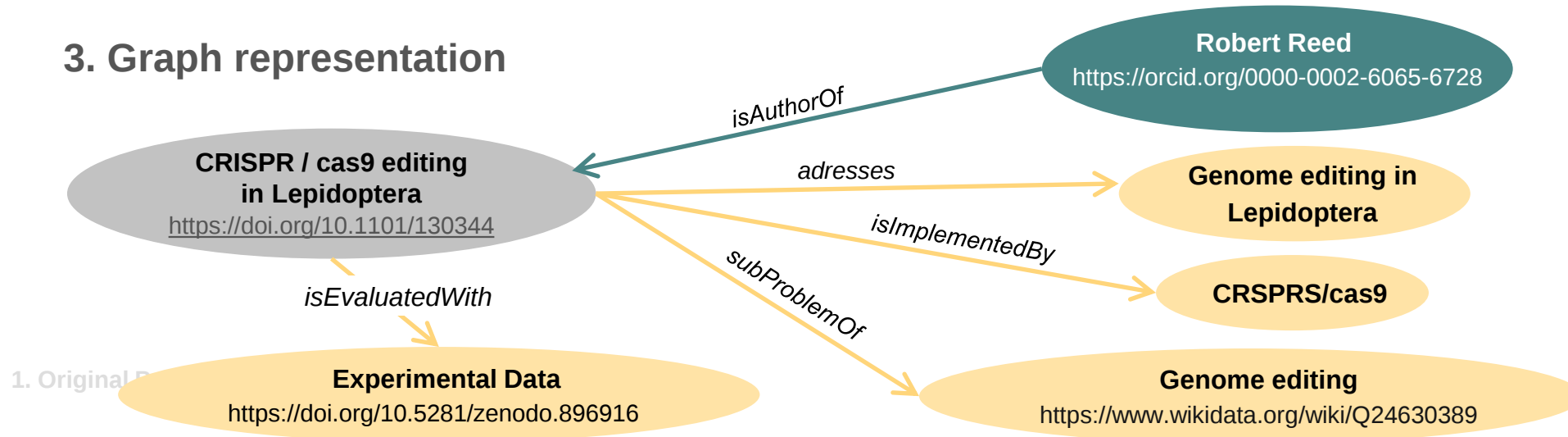
Research Problem Genome editing in Lepidoptera

Methods CRISPR / cas9

Applied on Lepidoptera

Experimental Data <https://doi.org/10.5281/zenodo.896916>

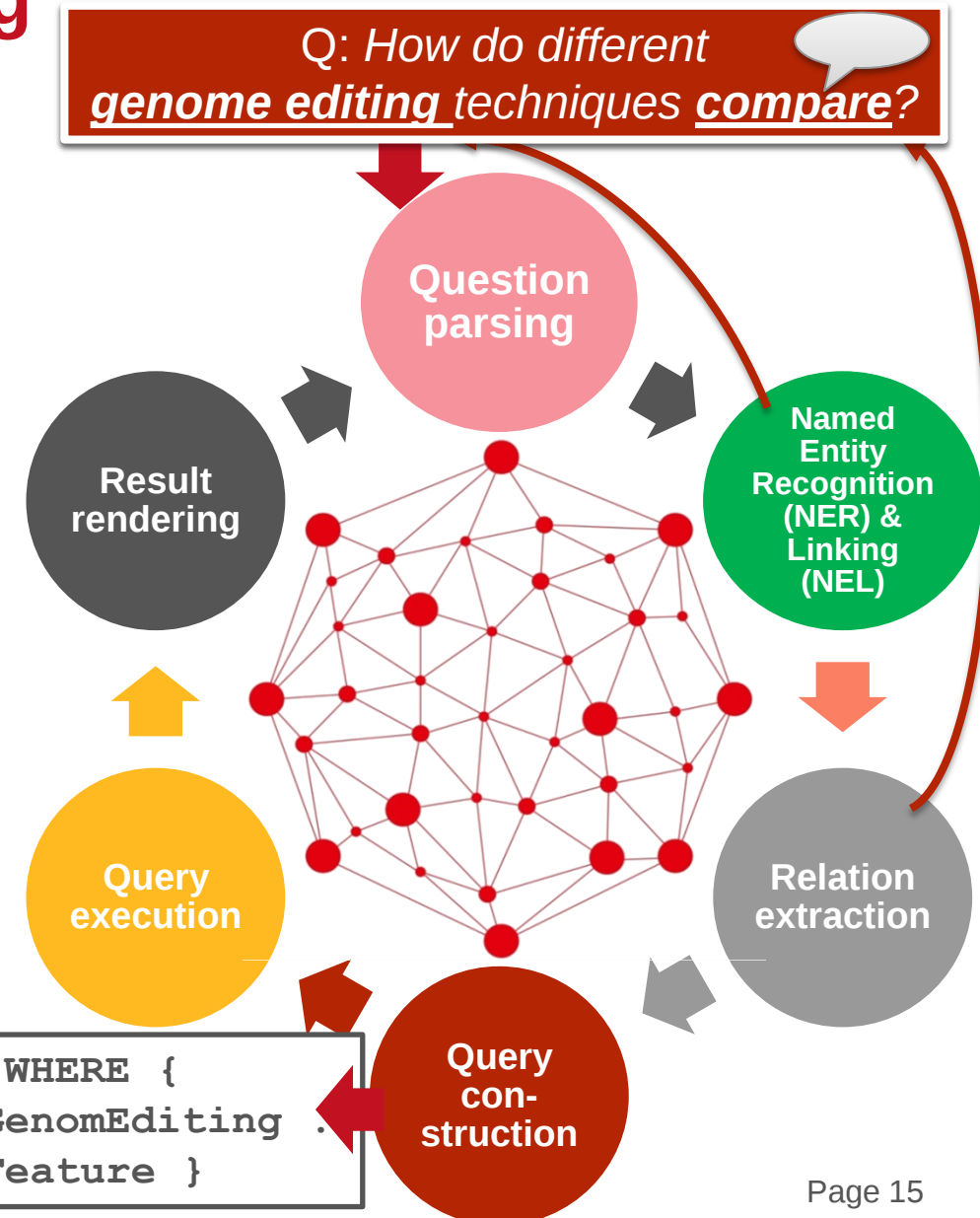
3. Graph representation



Exploration and Question Answering

Research Challenge:

- Intuitive exploration leveraging the rich semantic representations
- Answer natural language questions



[1] K. Singh, S. Auer et al: *Why Reinvent the Wheel? Let's Build Question Answering Systems Together*. The Web Conference (WWW 2018).

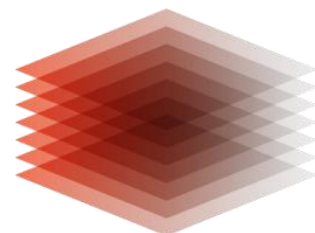
Result:

Automatic Generation of Comparisons / Surveys

Q: How do different **genome editing** techniques compare?

Engineered Nucleases	Site-specificity	Safety	Ease-of-use / costs/ speed
zinc finger nucleases (ZFN)	++ 9-18nt	+	-- \$\$\$: screening, testing to define efficiency
transcription activator-like effector nucleases (TALENs)	+++ 9-16nt	++	++ Easy to engineer 1 week / few hundred dollar
engineered meganucleases	+++ 12-40 nt	0	-- \$\$\$ Protein engineering, high-throughput screening
CRISPR system/cas9	++ 5-12 nt	-	+++ Easy to engineer few days / less 200 dollar

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Open Research Knowledge Graph
<https://orkg.org>

Properties	Estimation of the epidemic properties of the 2019 novel coronavirus: A mathematical modeling study Contribution 1 - 2020	Estimation of the epidemic properties of the 2019 novel coronavirus: A mathematical modeling study Contribution 2 - 2020	Estimation of the epidemic properties of the 2019 novel coronavirus: A mathematical modeling study Contribution 3 - 2020	Transmission potential of COVID-19 in Iran Contribution 1 - 2020	Transmission potential of COVID-19 in Iran Contribution 2 - 2020	Estimating the generation interval for COVID-19 based on symptom onset data Contribution 1 - 2020
Has research problem	COVID-19 reproductive number	COVID-19 reproductive number	COVID-19 reproductive number	COVID-19 reproductive number	COVID-19 reproductive number	COVID-19 reproductive number
Location	Wuhan City, China	Wuhan City, China	Wuhan City, China	Iran	Iran	Singapore
Study date	2020-01-10/2020-01-23	2020-01-23/2020-02-08	2020-01-10/2020-02-08	2020-02-19/2020-02-29	2020-02-19/2020-02-29	2020-01-21/2020-02-26
R0 estimates (average)	4.38	3.41	3.39	3.6	3.58	1.27
95% confidence interval	3.63-5.13	3.16-3.65	3.09-3.70	3.2-4.2	1.29-8.46	1.19-1.36
Method	a weighted average of Exponential growth, Maximum likelihood, Sequential Bayesian, Time-dependent reproduction numbers, and SEIR model basic reproduction numbers by calculating weights from a Poisson loss function	a weighted average of Exponential growth, Maximum likelihood, Sequential Bayesian, Time-dependent reproduction numbers, and SEIR model basic reproduction numbers by calculating weights from a Poisson loss function	a weighted average of Exponential growth, Maximum likelihood, Sequential Bayesian, Time-dependent reproduction numbers, and SEIR model basic reproduction numbers by calculating weights from a Poisson loss function	generalized growth model	based on the calculation of the epidemic's doubling times: estimated epidemic doubling time of 1.20 (95% CI, 1.05, 1.44) days	generation interval



View paper

View paper

Graph view

Edit

Transmission interval estimates suggest pre-symptomatic spread of COVID-19

March 2020

Virology

Lauren Tindale

Michelle Coombe

Jessica Stockdale

Emma Garlock

Wing Yin Venus Lau

Manu Saraswat

Yen-Hsiang Brian Lee

Louxin Zhang

Dongxuan Chen

Jacco Wallinga

Caroline Colijn

DOI: 10.1101/2020.03.03.20029983

Share this paper:  

Contribution 1

Contribution 2

Research problems

☐ Add to comparison

COVID-19 reproductive number

Contribution data

95% Confidence interval	1.45-2.48
Location	Singapore
Mean incubation period	7.1 (6.13, 8.25) days
Mean serial interval	4.56 (2.69, 6.42) days
R0 estimates (average)	1.97
Study date	2020-01-19/2020-02-26

Edit mode Every change you make is automatically saved

 View paper

 Graph view

 Stop editing

 March 2020

 Virology

 Lauren Tindale

 Michelle Coombe

 Jessica Stockdale

 Emma Garlock

 Wing Yin Venus Lau

 Manu Saraswat

 Yen-Hsiang Brian Lee

 Louxin Zhang

 Dongxuan Chen

 Jacco Wallinga

 Caroline Colijn

DOI: [10.1101/2020.03.03.20029983](https://doi.org/10.1101/2020.03.03.20029983)

 Edit data

Share this paper:  

Contribution 1 

Contribution 2



Research problems

☐ Add to comparison

COVID-19 reproductive number 



Contribution data

95% Confidence interval

1.45-2.48



Location

Singapore



Mean incubation period

7.1 (6.13, 8.25) days



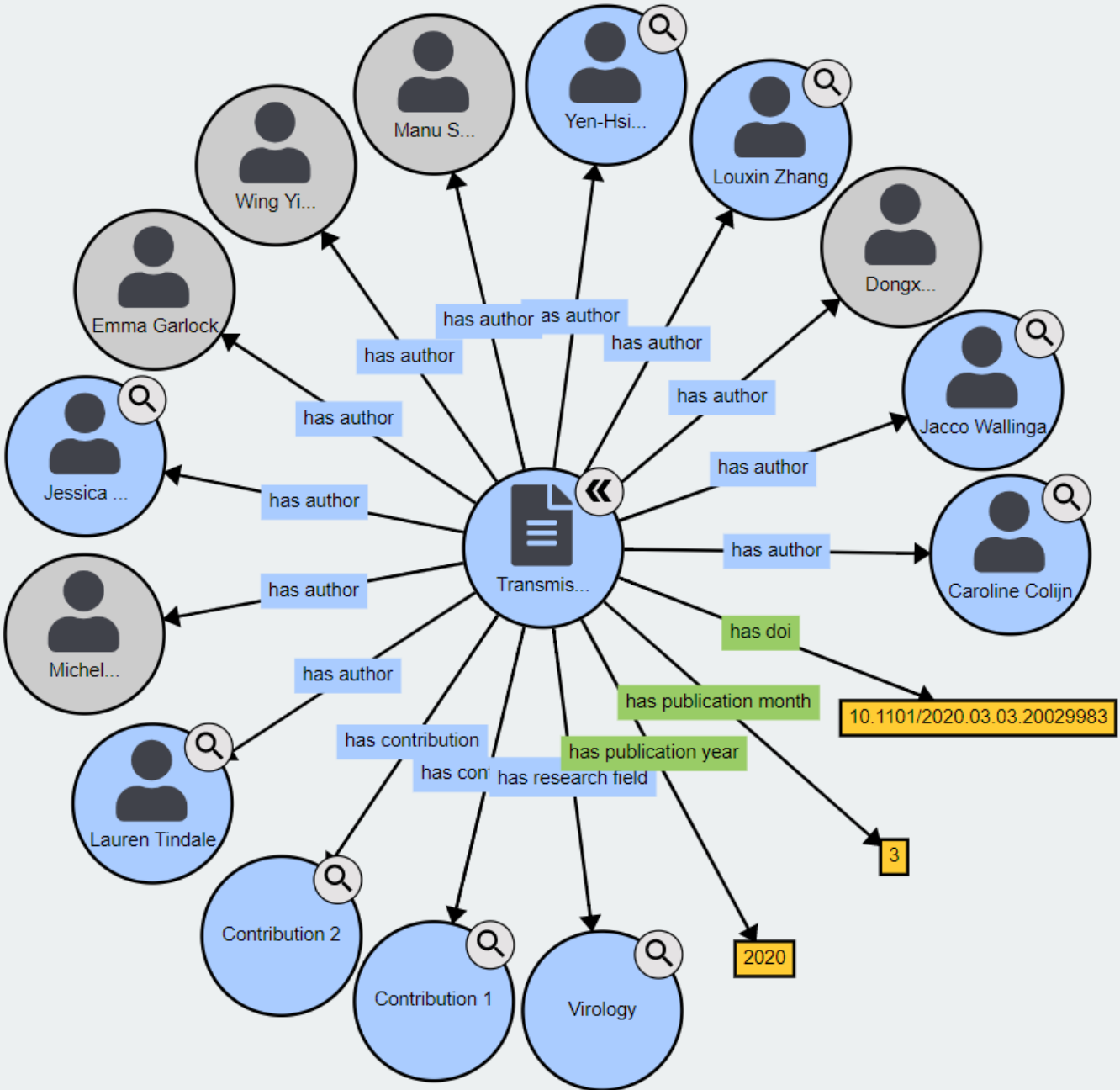
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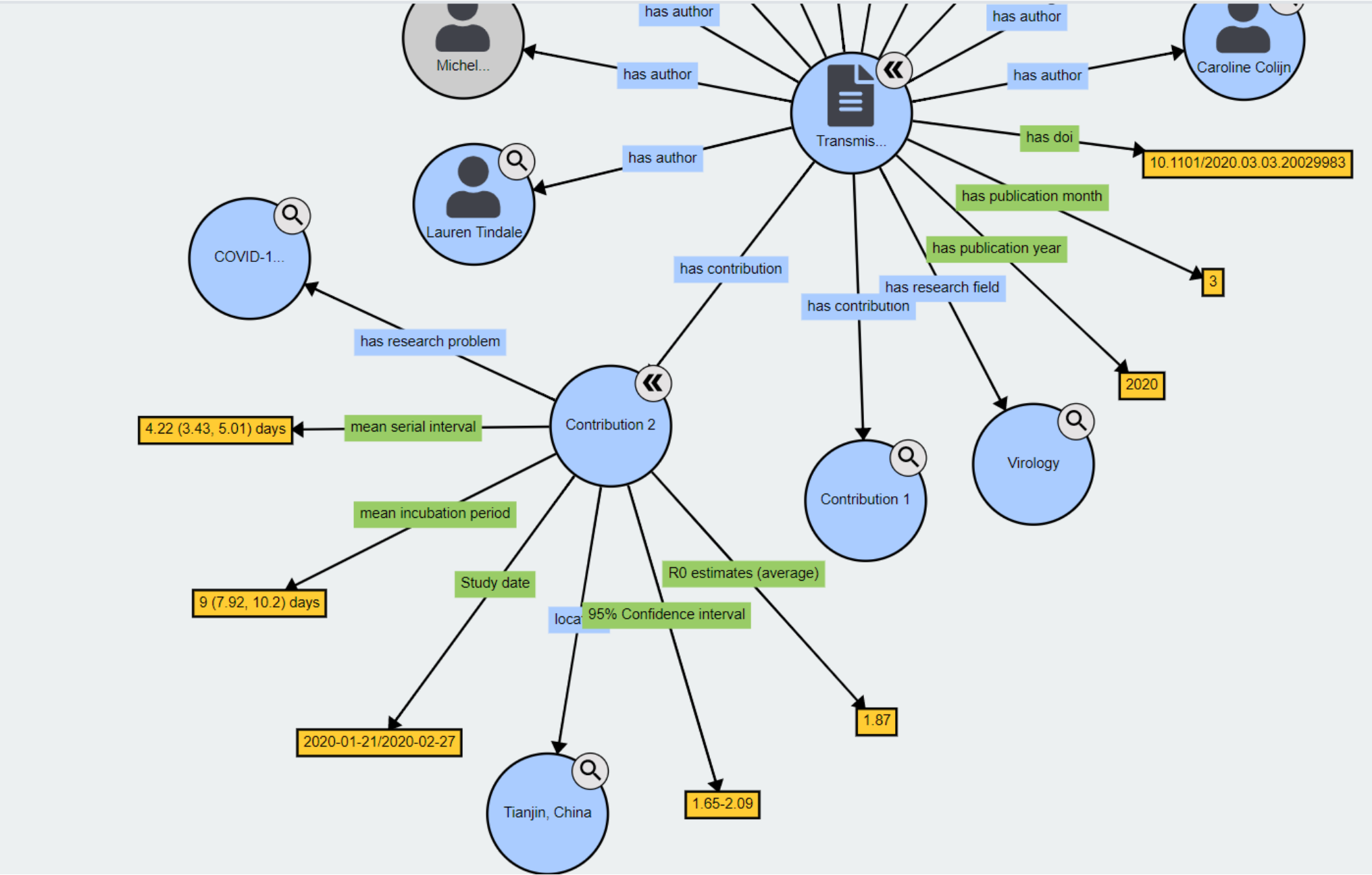
Mean serial interval

4.56 (2.69, 6.42) days

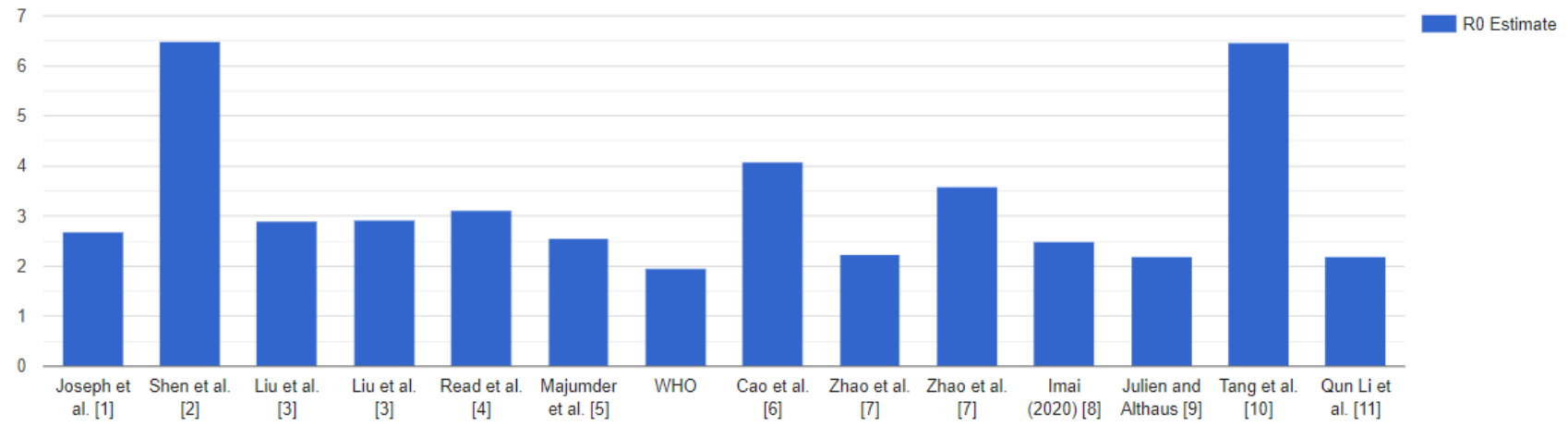
 



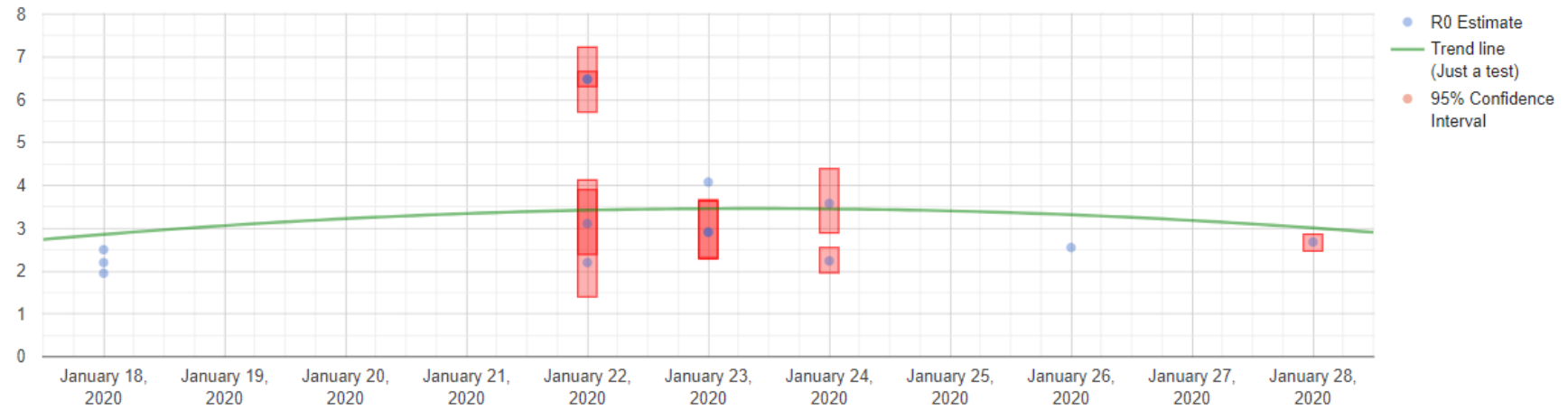




Bar Chart



Combo Chart



 ORKG

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» Life Sciences » Microbiology » Virology

Research field | Virology

Edit

Description

Virology is the study of viruses – submicroscopic, parasitic particles of genetic material contained in a protein coat – and virus-like agents. It focuses on the following aspects of viruses: their structure, classification and evolution, their ways to infect and exploit host cells for reproduction, their interaction with host organism physiology and immunity, the diseases they cause, the techniques to isolate and culture them, and their use in research and therapy.

Contributors

All time ▾



Research problems

- COVID-19 reproductive number
- identification of SARS-Cov-2
- Predicting the evolution of 2019-nCoV
- COVID-19 case fatality rate
- assess the impact of COVID-19 by SEIR model predictions

View more

Last updates

a month ago

Markus Stocker added visualization **Hazard ratio SARS-CoV-2 B.1.1.7**

a month ago

Markus Stocker added comparison **Case fatality risk of SARS-CoV-2 B.1.1.7**

a month ago

Anonymous user added contribution **Contribution 1**

View more

Observatories

COVID-19

Demonstration observatory with research problems and comparisons related to COVI...

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5 comparisons 31 papers



Comparisons | 11 comparisons

Newest first ▾

Case fatality risk of SARS-CoV-2 B.1.1.7

3 Contributions 01-04-2021

Hazard ratio estimates for SARS-CoV-2 variant of concern B.1.1.7 compared with non variant of concern cases

» Virology



SARS-CoV-2 BioAssays

38 Contributions 26-01-2021

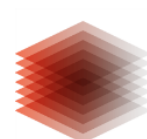
Comparison of over 35 Bio Assays regarding SARS-CoV-2

» Virology



Research field overviews
organize research problems,
comparisons, visualizations etc.

Show latest updates,
contributors, observatories



» Life Sciences » Microbiology » Virology

Research problem | COVID-19 reproductive number

Edit

Description

In epidemiology, the basic reproduction number, or basic reproductive number (sometimes called basic reproduction ratio or basic reproductive rate), denoted R_0 (pronounced R nought or R zero), of an infection can be thought of as the expected number of cases directly generated by one case in a population where all individuals are susceptible to infection.

Summary from Wikipedia

Failed loading summary from Wikipedia.



Contributors



Top Authors

Dongxuan Chen

6 papers

Jacco Wallinga

6 papers

Yanni Xiao

5 papers

View more

Research fields

Research fields of papers that are addressing this problem

- Virology 41

Super-problems

Research problems that has this problem as Sub Problem.

No super research problems.

Papers

Estimation of the Transmission Risk of the 2019-nCoV and Its Implication for Public Health Interventions -

Contribution 1

Tang, Biao, Wang, Xia, Li, Qian, Bragazzi, Nicola Luigi, Tang, Sanyi et al. 2020

Virology



Estimating the Unreported Number of Novel Coronavirus (2019-nCoV) Cases in China in the First Half of January

2020: A Data-Driven Modelling Analysis of the Early Outbreak - Contribution 1

Virology

Organize Research Contributions along research problems



View all observatories

Observatories organize research contributions in a particular research field and are curated by research

Further information about observatories can be also found in the [ORKG wiki](#).

All research fields

Oceanography

Virology

Computer Sciences

Life Sciences

Artificial Intelligence/Robotics

Information Science

**UNIKLINIK
RWTHAACHEN**



Occupants' perception and behaviour

Papers: 2

Comparisons: 0

**Leibniz
AILab**

Artificial Intelligence

Papers: 0

Comparisons: 0



Digital Libraries

Papers: 0

Comparisons: 0



Microbiodata

Papers: 0

Comparisons: 0

Domain Observatories

Make organizations supporting the curation of research in a certain domain more visible

 ORKG

PapersTools ▾About ▾

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Observatory

COVID-19

Demonstration observatory with research problems and comparisons related to COVID-19.

Research field: [Virology](#)

Research Problems

1. Inhibition of sars-cov 3-chymotrypsin-like protease
2. Identification of sars-cov-2 by metagenomic rna sequencing
3. Predicting the evolution of 2019-ncov
4. Sequencing of the sars-cov-2 genome in iceland
5. Sequencing of the sars-cov-2 genome
6. Sars-cov-2 infection of residents of a nursing facility in the usa
7. Determination of the covid-19 basic reproduction number

Organizations



Contributors

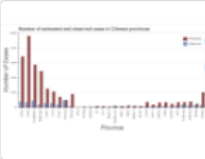
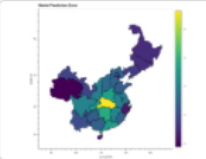
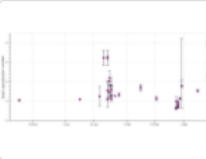
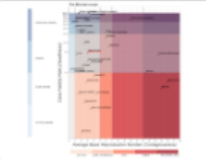
 **Maria-Esther Vidal**
TIB Leibniz Information Centre for Science and Technology

 **Jennifer D'Souza**
TIB Leibniz Information Centre for Science and Technology

 **Philipp Dillschneider**

Content

Figures



Comparisons

[COVID-19 Reproductive Number Estimates](#) 31 Contributions  24-06-2020

Observatories are curated by organizations and organize research problems. Contributions, comparisons and visualizations in a particular field



View all organizations



InfAI Institut für Angewandte Informatik



International Future Lab for Artificial Intelligence in Hannover



Leibniz University of Hannover



Uniklinik RWTH Aachen



TIB Leibniz Information Centre for Science and Technology



L3S Research Center



Northwestern University



ZB MED Informationszentrum Lebenswissenschaften



LifeGlimmer



Vrije Universiteit Amsterdam



EUROPEAN OPEN SCIENCE CLOUD



NFDI4Ing

Showcase contributing organizations

The Team



Group Leaders



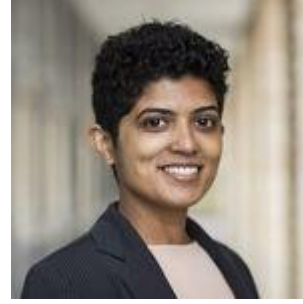
Dr. Markus Stocker



Dr. Gábor Kismihók



Dr. Javad Chamanara



Dr. Jennifer D'Souza



Dr. Lars Vogt



Dr. Oliver Karras

PostDocs

Doctoral Researchers



Allard Oelen



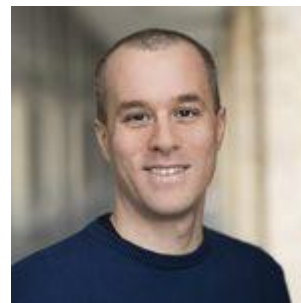
Yaser Jaradeh



Vitalis Wiens



Muhammad Haris



Manuel Prinz



Kheir Eddine Farfar



Suhas Murthy

Software Development



Golsa Heidary



Salmonon Kabongo



Dr. Michael Martin



Natanael Arndt

Collaborators InfAI Leipzig / AKSW



Katja Bartel



Simone Matern

Administration

Conclusions

- We need to **reinvent scholarly communication**
- **Knowledge Graphs** are perfectly suited to capture research contributions in a structured and semantic way making them human and machine interpretable
- With our **Open Research Knowledge Graph initiative** we aim to establish a registry for research contributions
- Curation and **synergistic combination of human, expert and machine intelligence** is a challenge

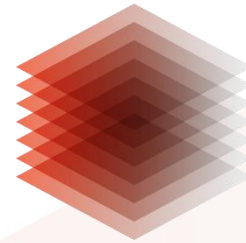
Stay tuned

- <https://tib.eu>
- Consider creating an ORKG observatory for your domain
- Mailinglist/group:
<https://groups.google.com/forum/#!forum/orkg>
- Open Research Knowledge Graph:
<https://orkg.org>
- ERC Consolidator Grant ScienceGRAPH on the topic



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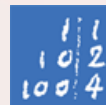
https://www.xing.com/profile/Soeren_Auer



http://www.researchgate.net/profile/Soeren_Auer

Contact

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TIB & Leibniz University of Hannover
auer@tib.eu



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