

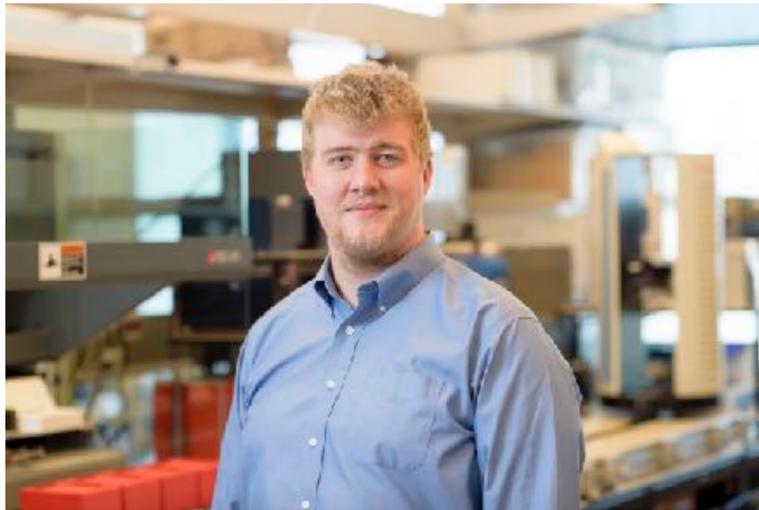
# BioExcel Webinar Series #47

## COVID-19 Molecular Structure and Therapeutics Hub

**Presenter:** Levi Naden (Molecular Sciences Software Institute (MolSSI))

**Host:** Rossen Apostolov (KTH Royal Institute of Technology)

# Today's Presenter



***Levi Naden***

*Software Scientist at MolSSI*  
*lnaden@vt.edu*

Levi Naden is a Software Scientist at MolSSI and lead for COVID-19 Molecular Structure and Therapeutics Hub project. Levi has a background in computational free energy algorithms, scientific software development and deployment, and software interoperability. He has a Bachelors in Chemical Engineering from Kansas State University and a PhD in Chemical Engineering from University of Virginia.

**The Molecular Sciences Software Institute:** <https://www.molssi.org/>

# What is MolSSI?

Designed to **serve** and **enhance** the software development efforts of the broad field of computational molecular science (CMS) – a domain that includes quantum chemistry, computational materials science, and biomolecular simulation.

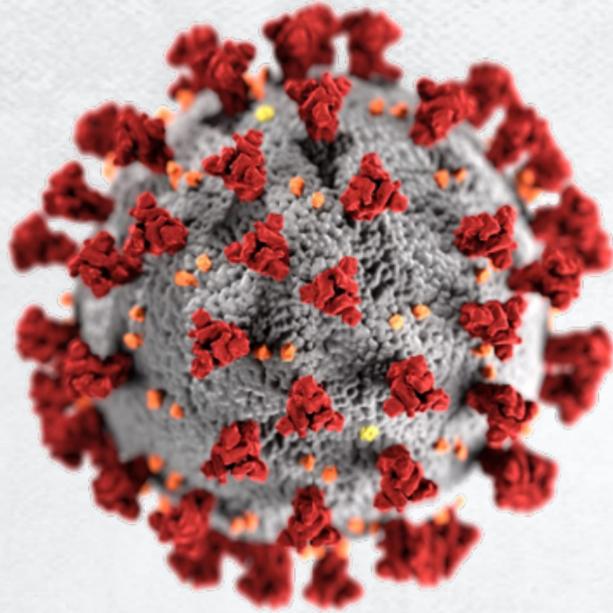
Goals:

1. Provide software expertise and infrastructure
2. Provide education and training
3. Provide community engagement and leadership



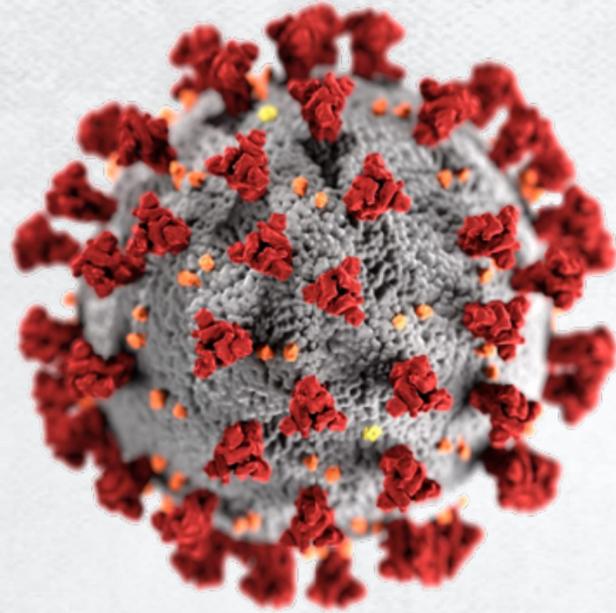
# There are ups and downs to the current state of research

The Enemy



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The Good

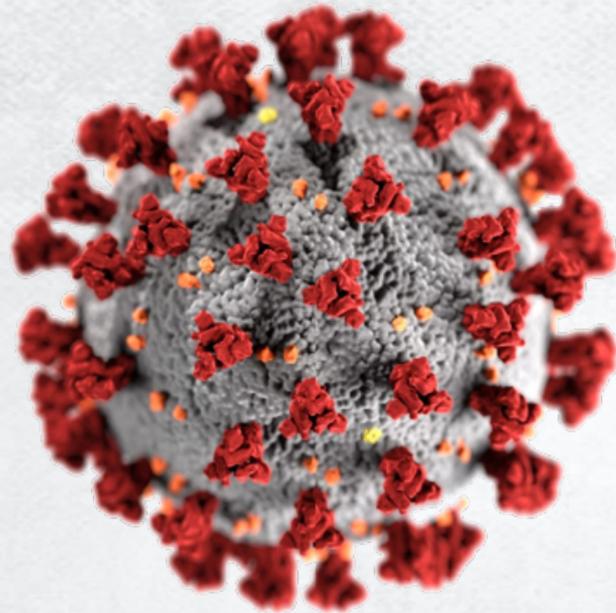
## ***Covid-19 Changed How the World Does Science, Together***

Never before, scientists say, have so many of the world's researchers focused so urgently on a single topic. Nearly all other research has ground to a halt.

Source: New York Times

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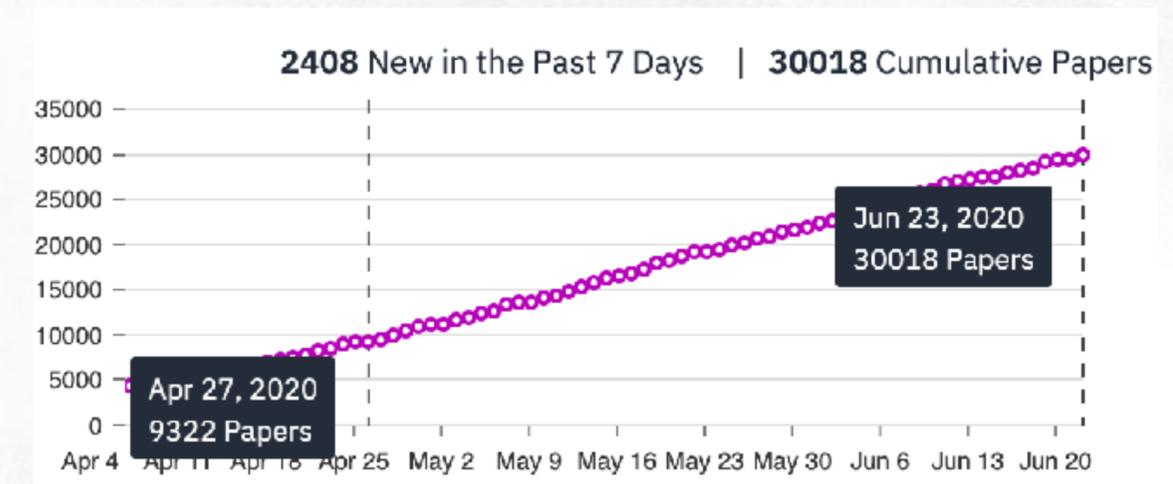
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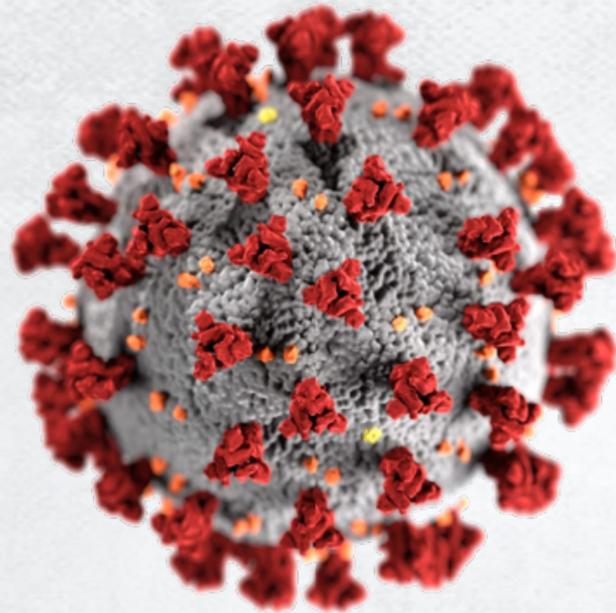
## The Problems



Total Volume of COVID-19 Research Papers  
Source: covid19primer by Primer.ai.

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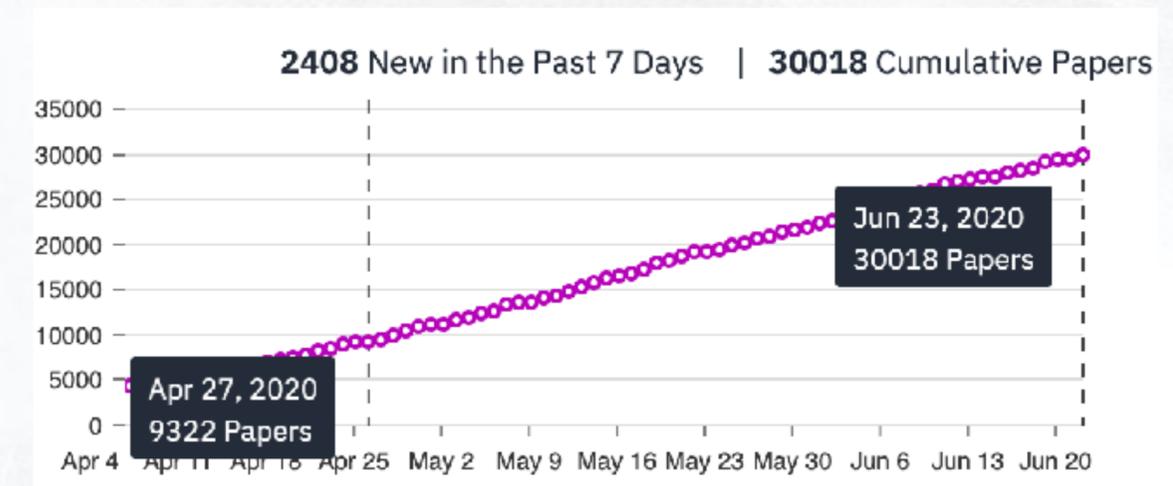
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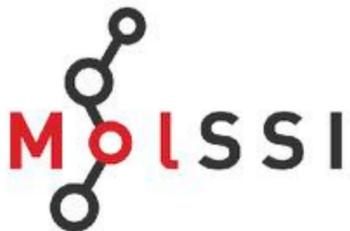


Total Volume of COVID-19 Research Papers  
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Lots of scientists stuck at home right now looking to help

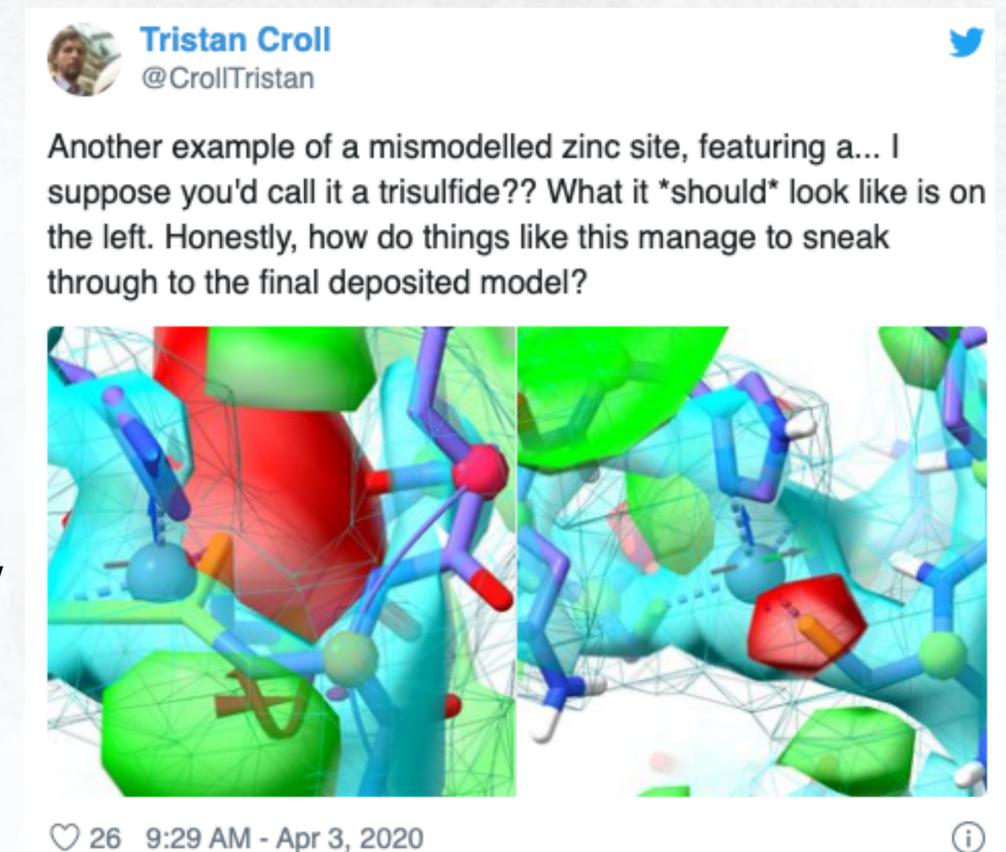


Source: First google search for "antsy person"



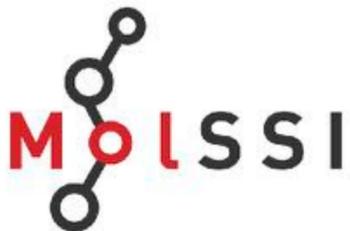
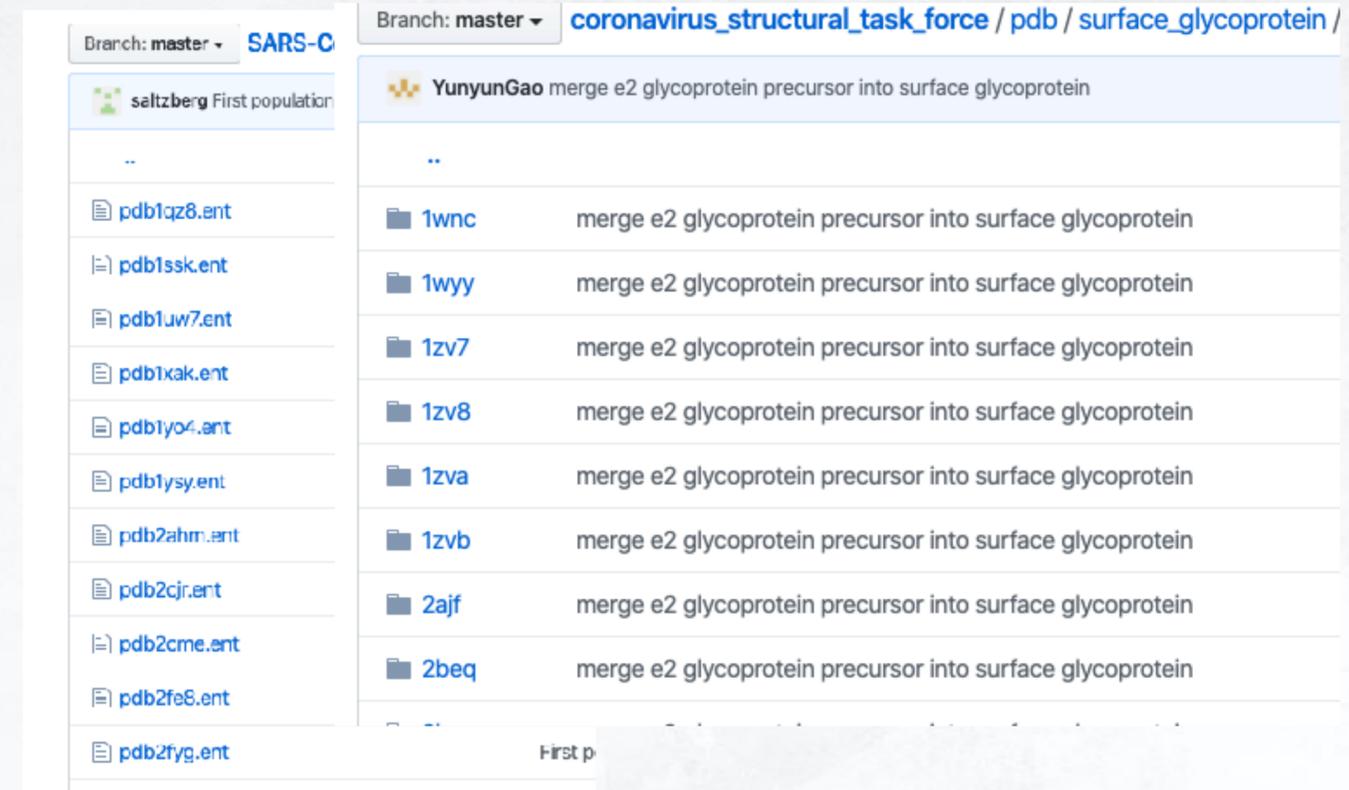
# Every institute and lab under the sun is doing something with SARS-CoV-2

- Overall a good thing, but has downsides
  - Editors: “We are being overwhelmed with papers”
  - Quality: Mixed bag
  - Collaborators: We have data, people want data, how do I get them data?



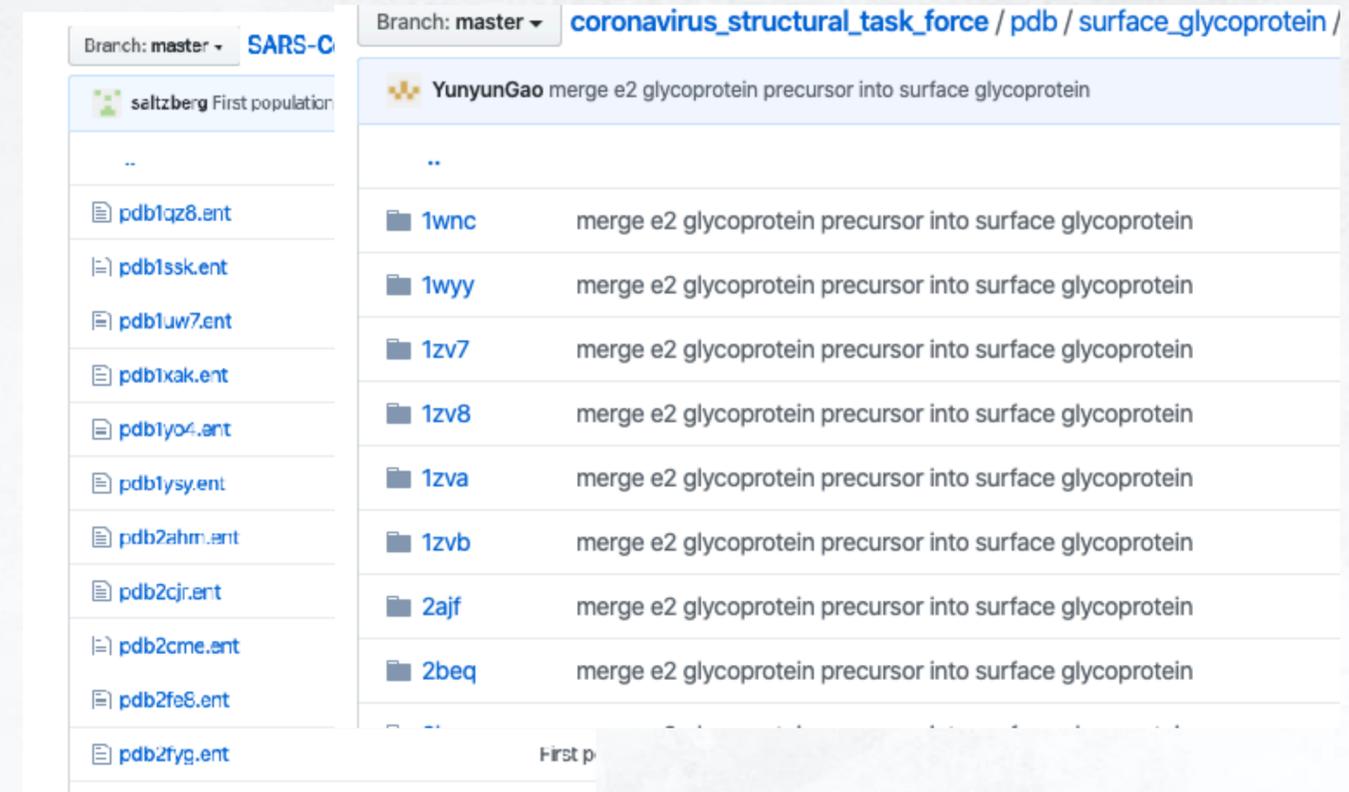
# Even good data are nebulous

- Everyone does their own work
- And is great in their own rights!



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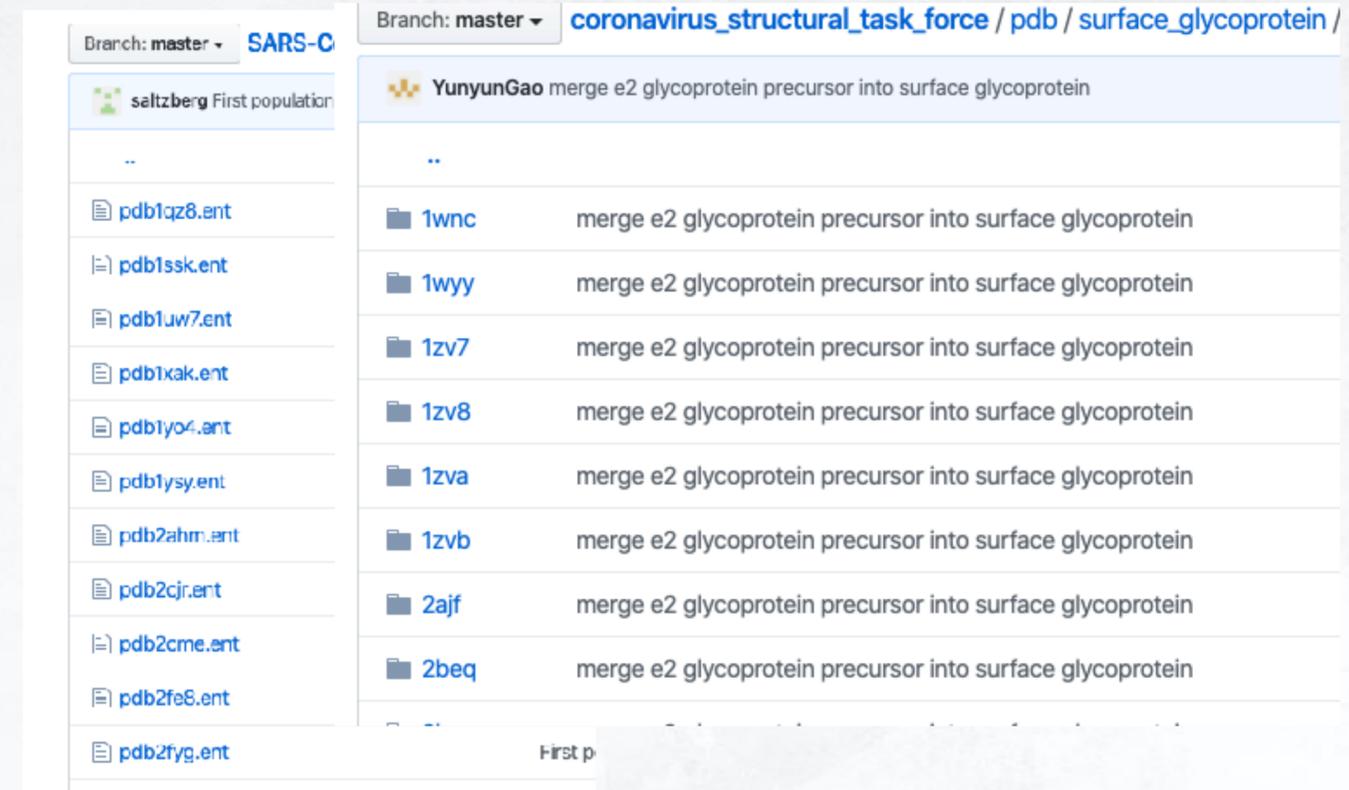
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- The data are all organized in domain specific ways



The screenshot shows a GitHub repository interface. The left sidebar displays a directory structure for the 'SARS-C' branch, listing files such as 'pdb1qz8.ent', 'pdb1ssk.ent', 'pdb1uw7.ent', 'pdb1xak.ent', 'pdb1yo4.ent', 'pdb1ysy.ent', 'pdb2ahm.ent', 'pdb2cjr.ent', 'pdb2cme.ent', 'pdb2fe8.ent', and 'pdb2fyg.ent'. The main content area shows the 'coronavirus\_structural\_task\_force / pdb / surface\_glycoprotein /' branch, displaying a list of merge commits by YunnyunGao, all with the message 'merge e2 glycoprotein precursor into surface glycoprotein'. The commits are identified by IDs: 1wnc, 1wy, 1zv7, 1zv8, 1zva, 1zvb, 2ajf, and 2beq.

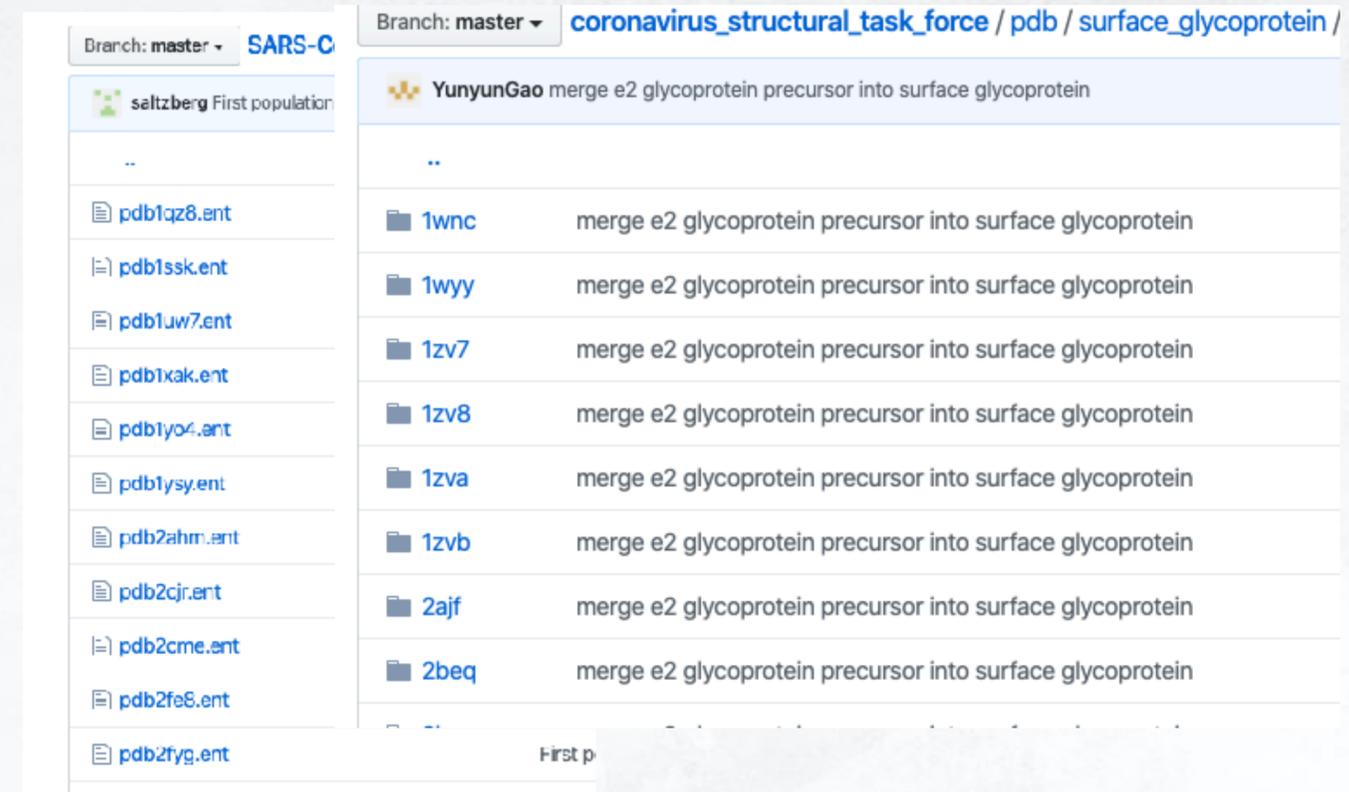
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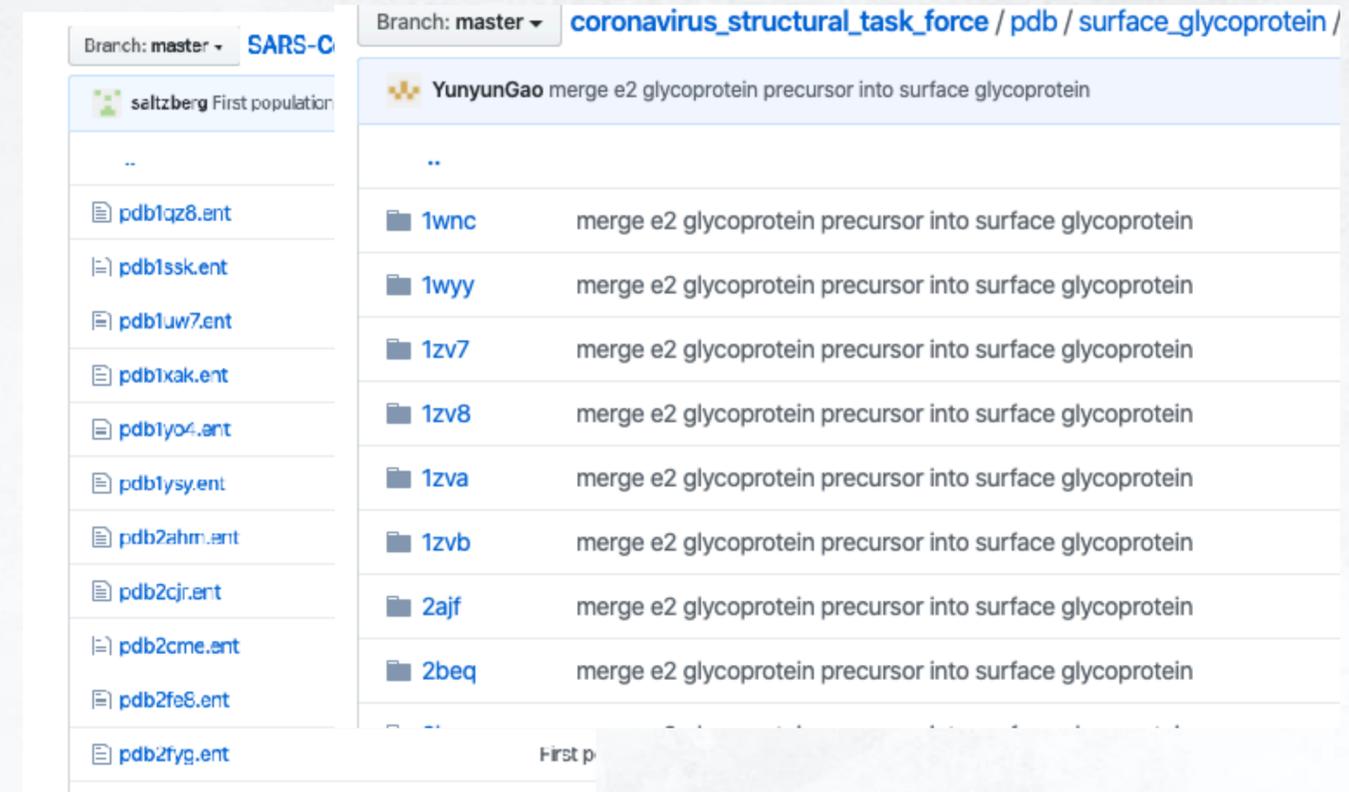
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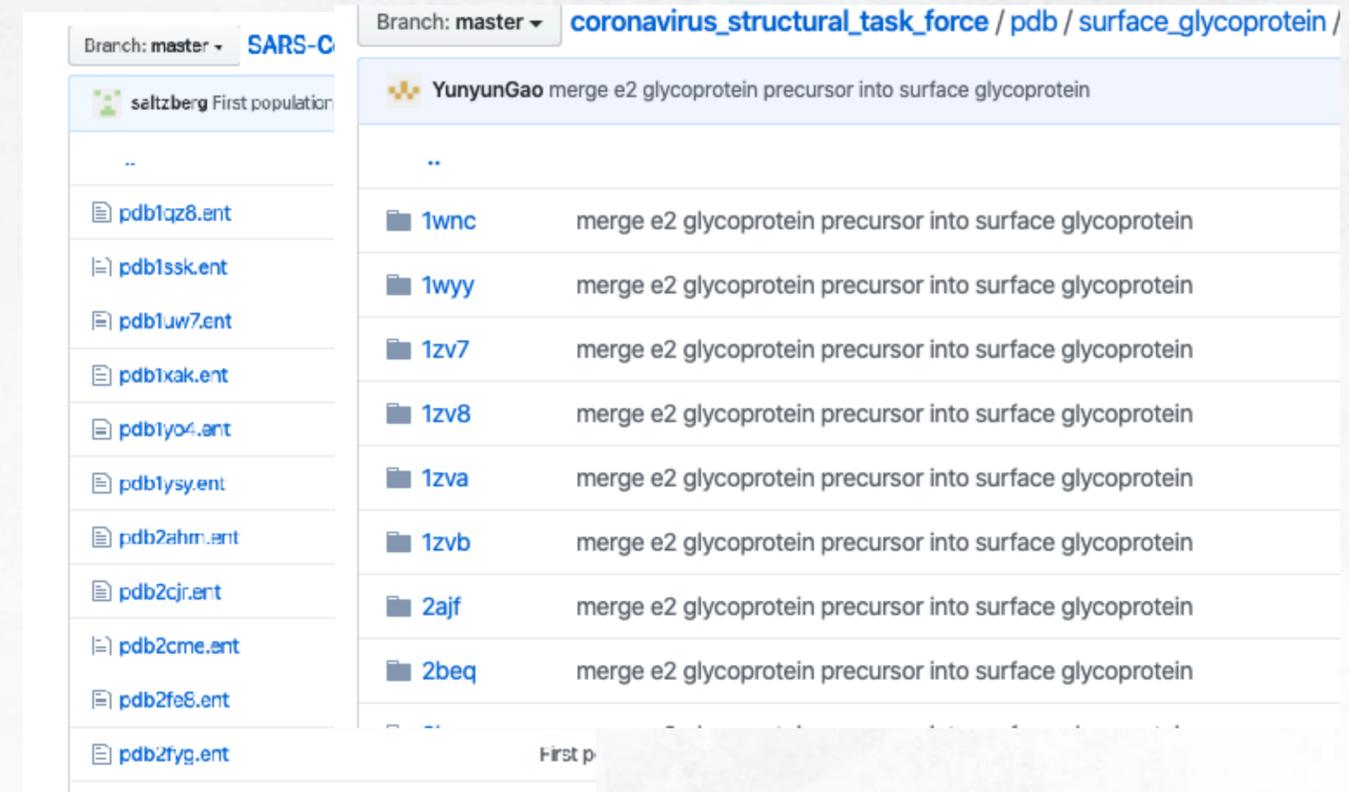
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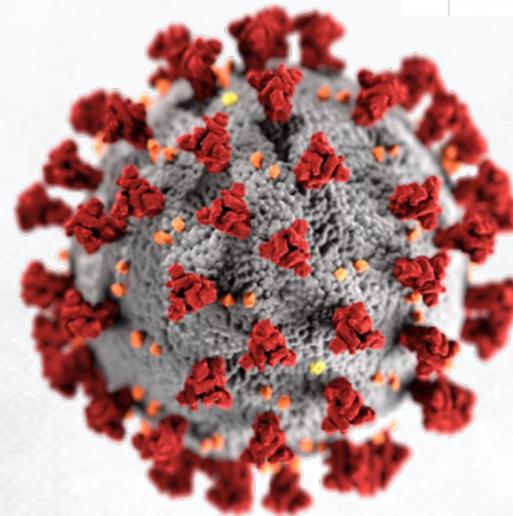
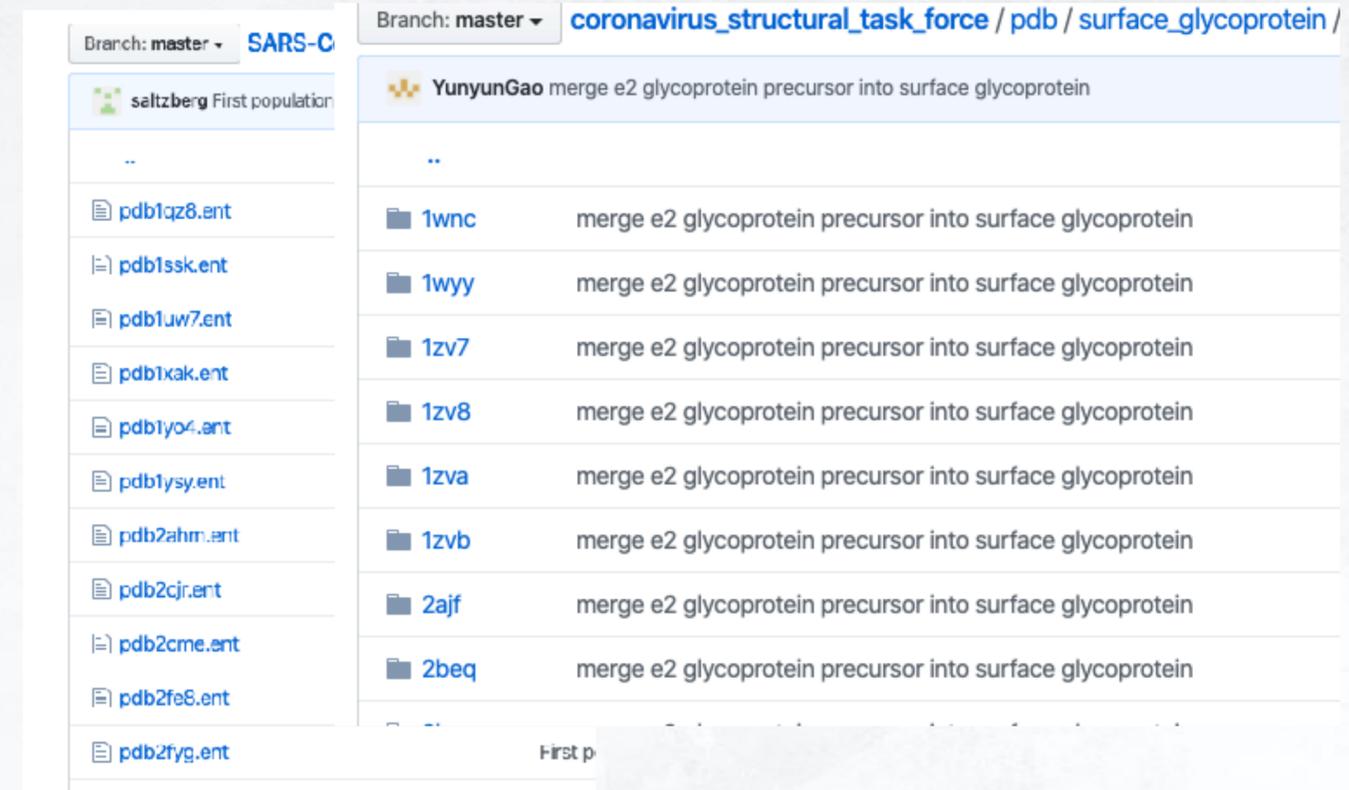
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- Whats good? Whats not?
  
- Most data are only small parts of the large picture



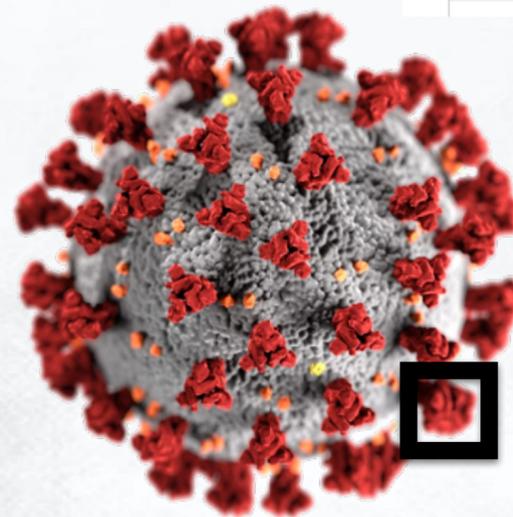
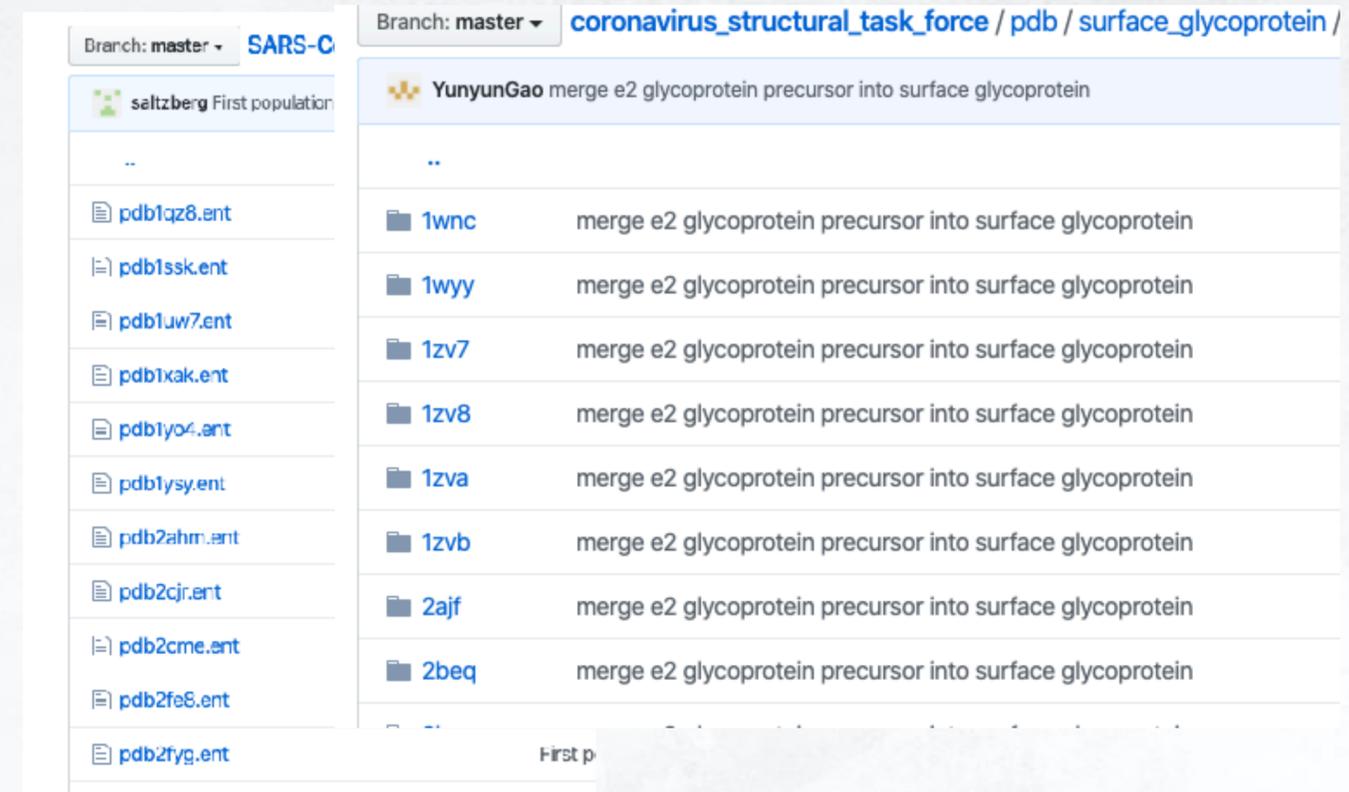
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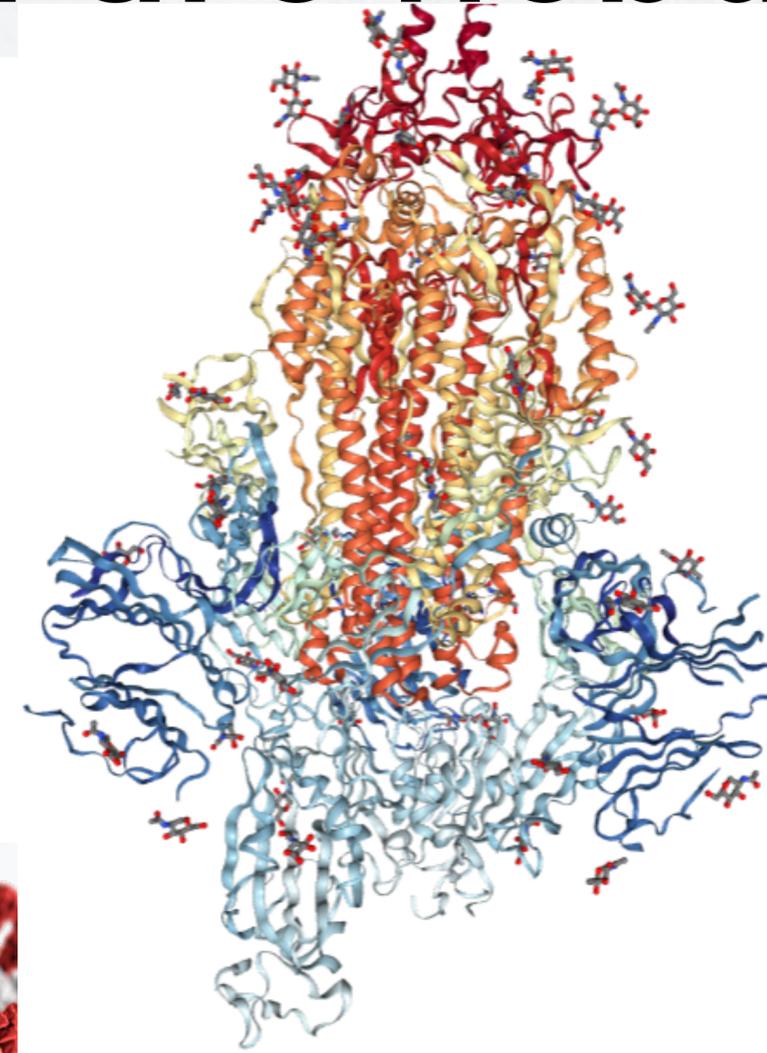
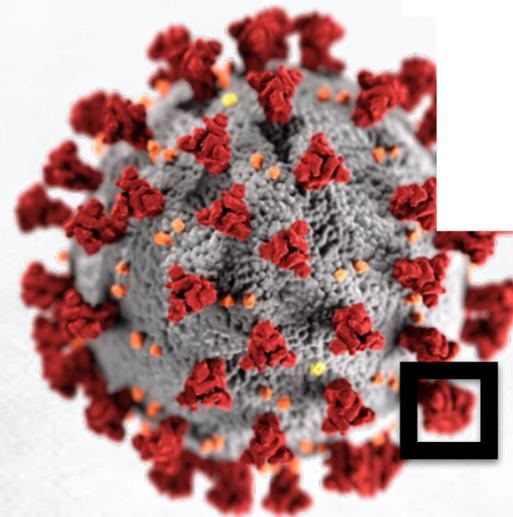
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Spike

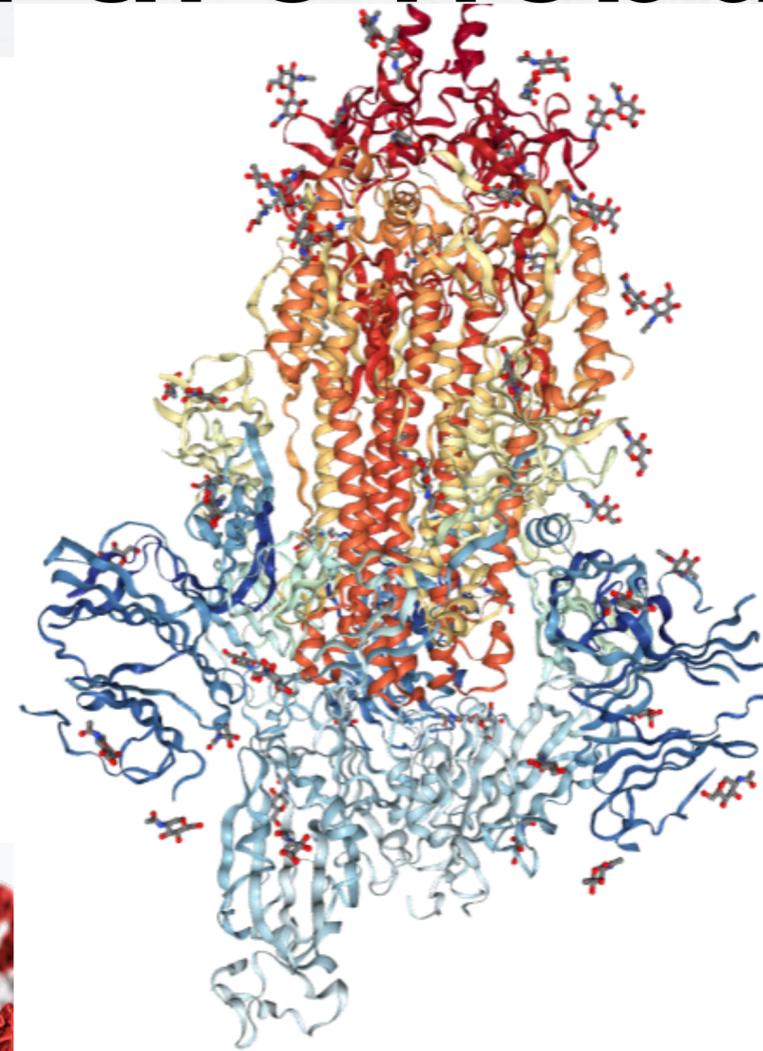
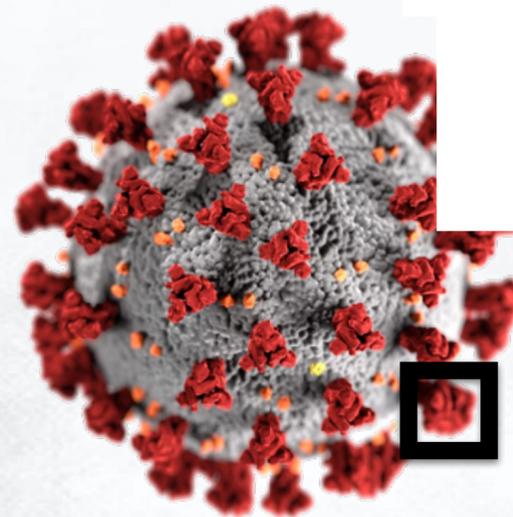
[al\\_task\\_force / pdb / surface\\_glycoprotein /](#)

sor into surface glycoprotein

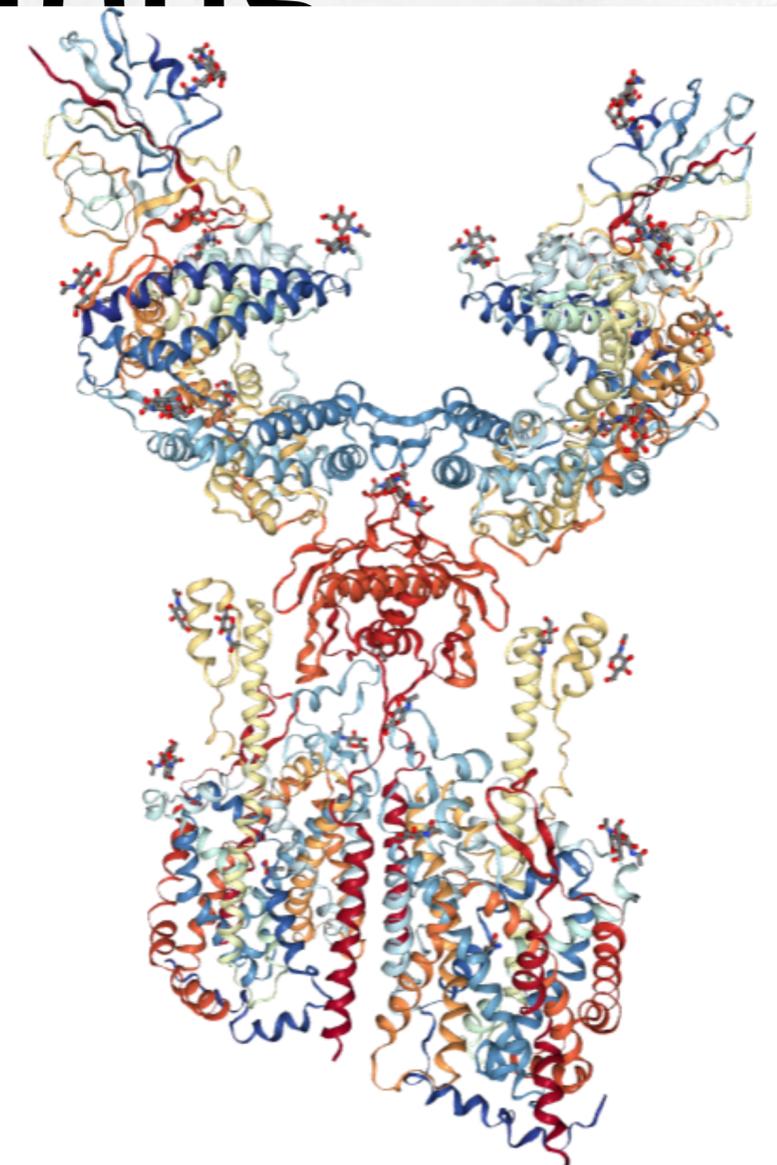
precursor into surface glycoprotein

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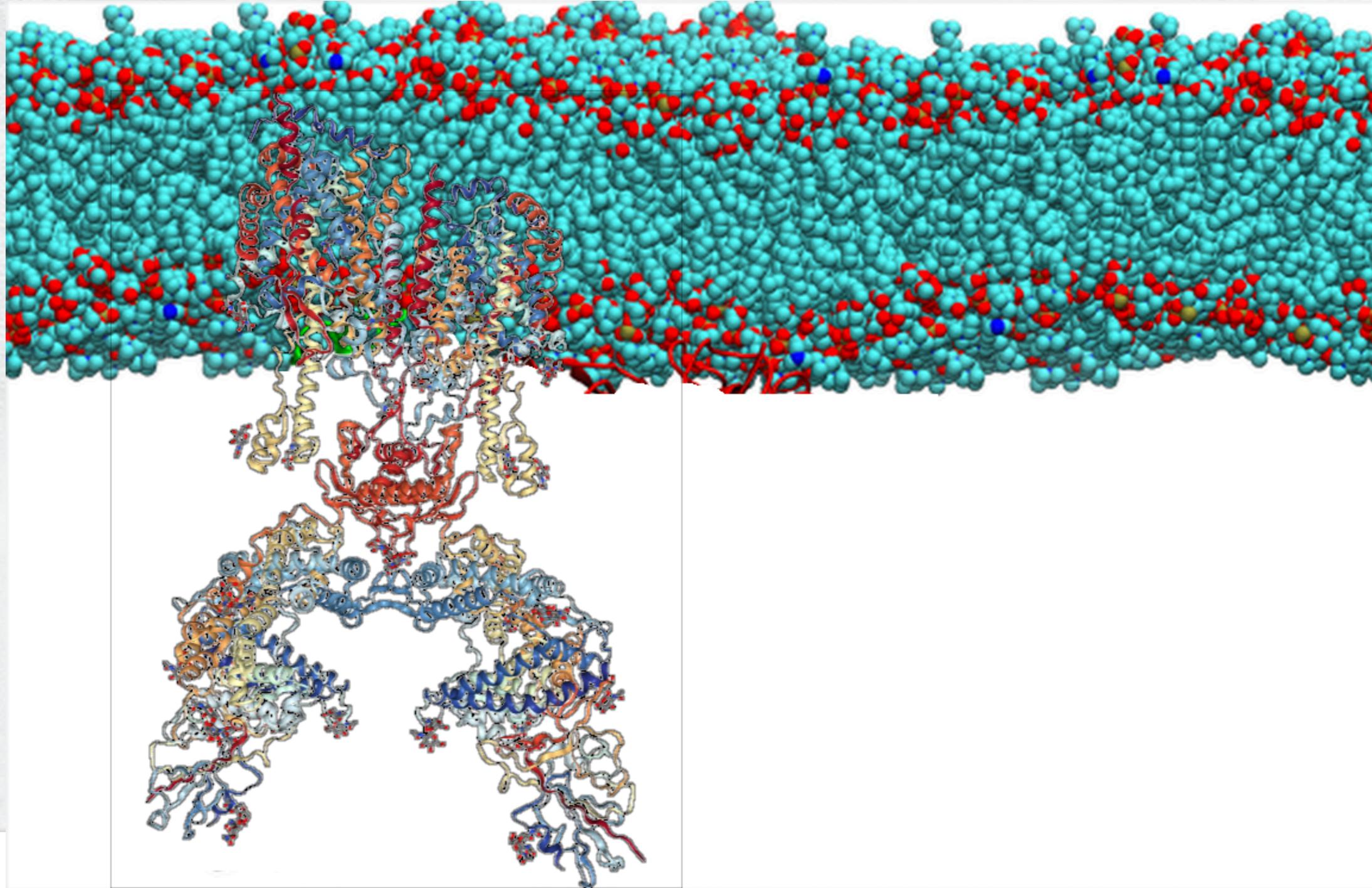
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ACE2

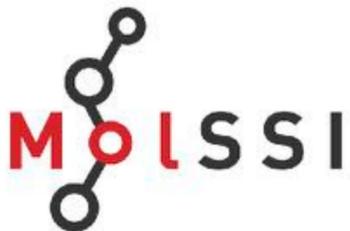


ACE2

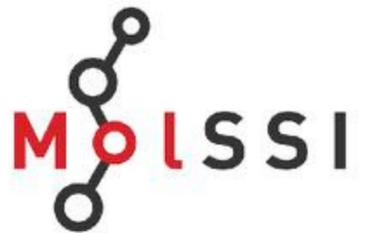
# We needed a hub for easily conveyed science

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Simple descriptors  
Community peer reviewed  
Centralized or pointers to data  
Accessible to experts in all domains



# Two PI's sparked what is now a global network of cooperation



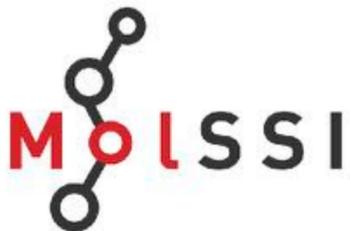
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Rommie Amaro



Adrian Mulholland



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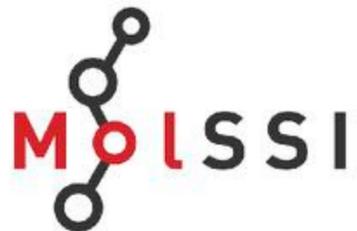
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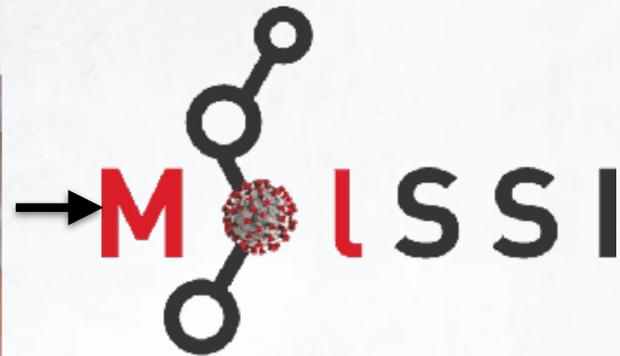
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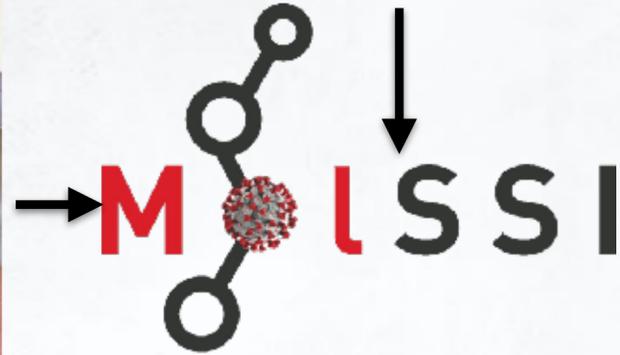
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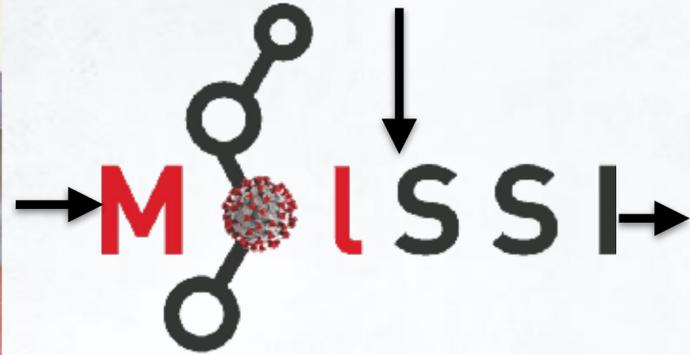
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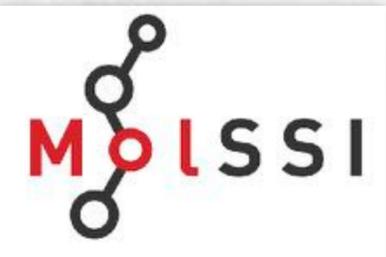
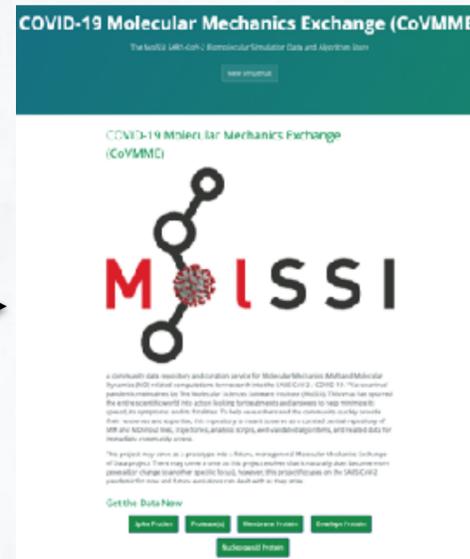
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Adrian Mulholland



Prototype (3 months ago)



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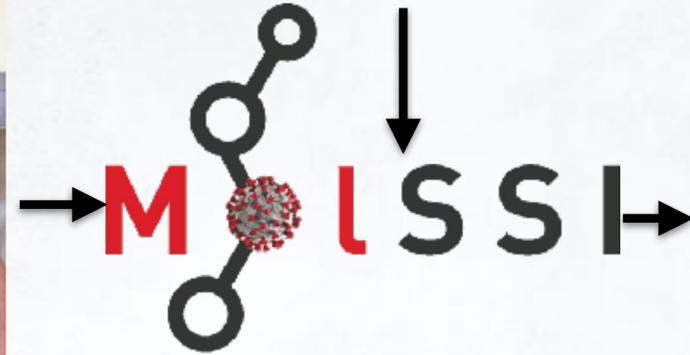
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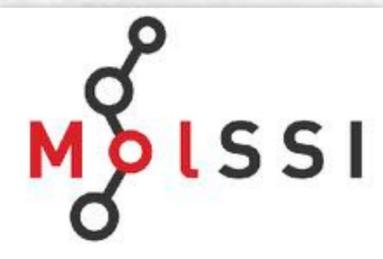
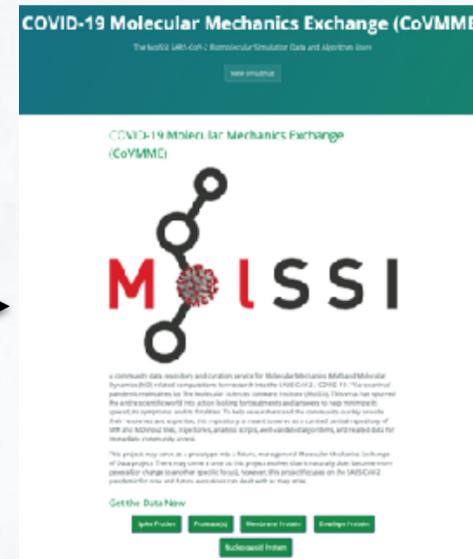
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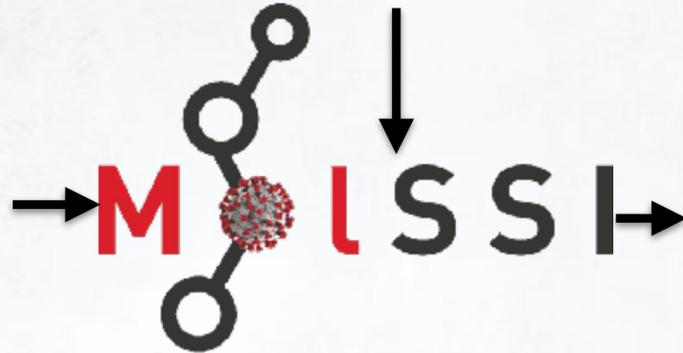
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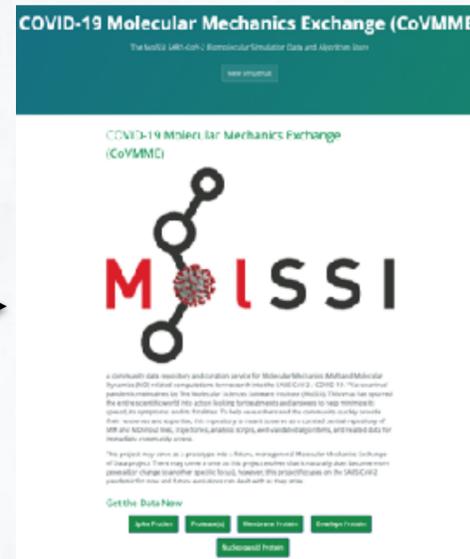
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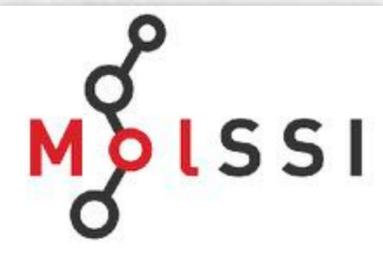
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D E Shaw Research



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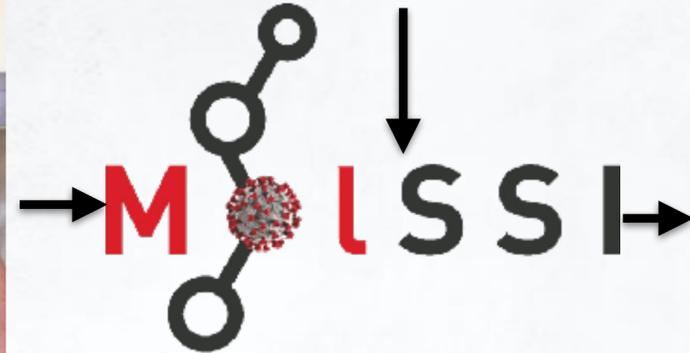
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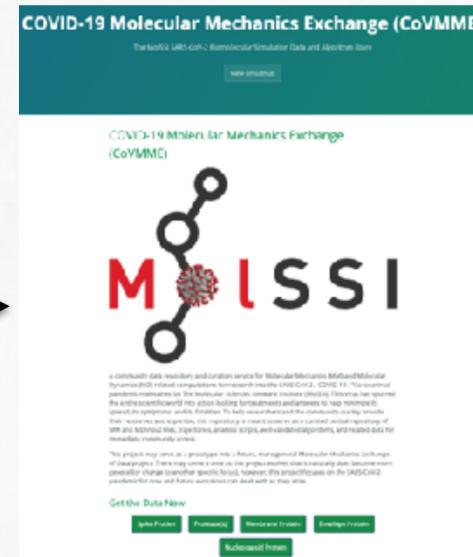
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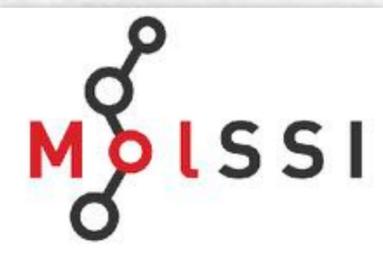
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D E Shaw Research



One month a whole bunch of this later...



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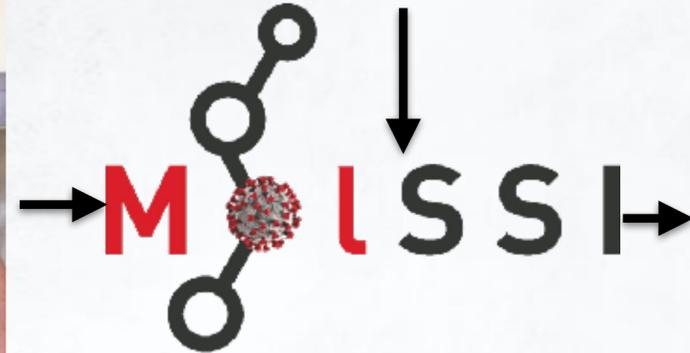
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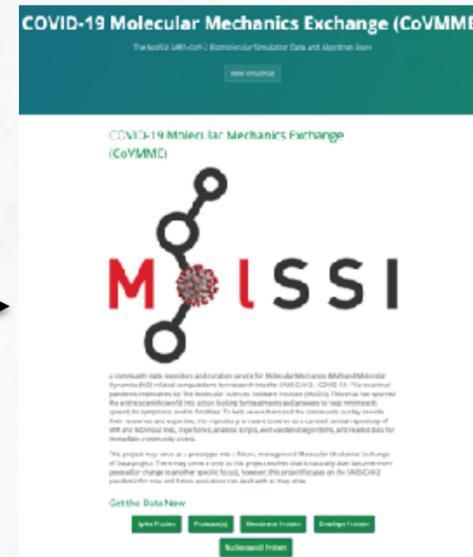
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Prototype (3 months ago)



D E Shaw Research



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Dozens of individual contributors  
Community in US, EU, Switzerland, Korea, Japan  
Upwards of a dozen Orgs./Institutes aiding or in talks



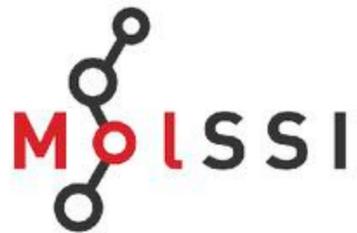
[covid.molssi.org](https://covid.molssi.org)



# [covid.molssi.org](https://covid.molssi.org)

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Let's take a look at what is there!



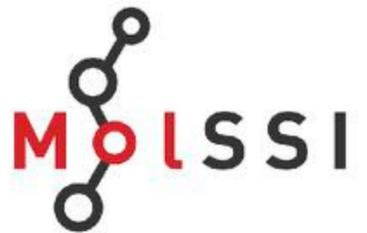
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[github.com/MolSSI/covid](https://github.com/MolSSI/covid)



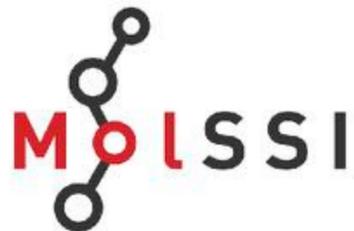
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- DataCite for issuing DOIs on demand
  - Make data citable in papers!



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- AWS Open Data Program for storage
  - 100+ TB of storage
  - Explore options of storing the data we generate correctly



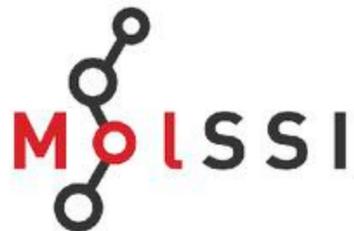
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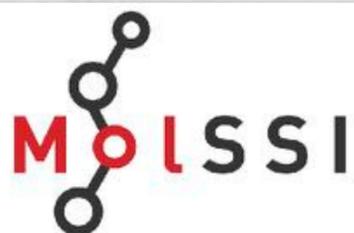
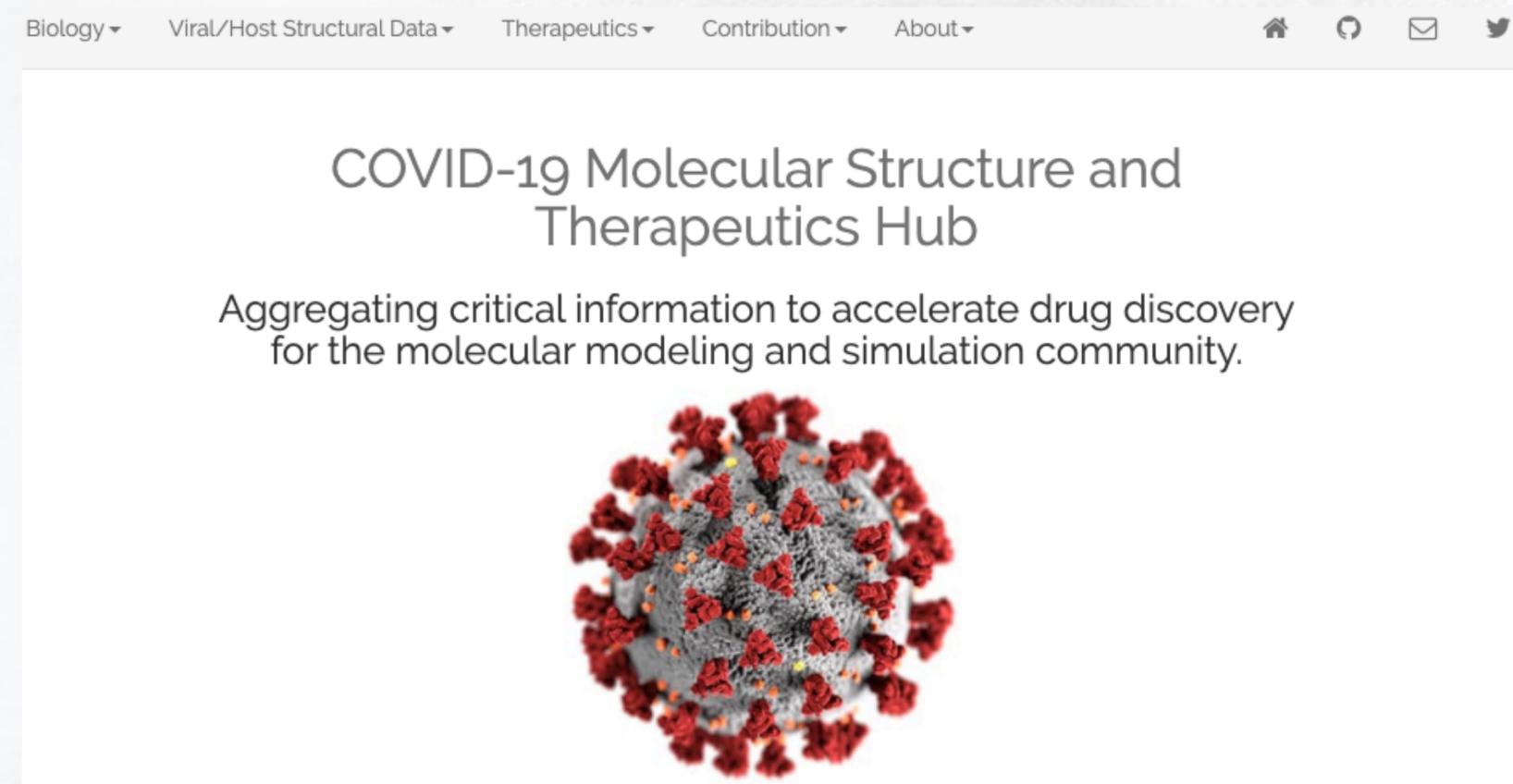
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- Potential Improvements:
  - Knowledge Graphs for data
  - Data licensing schemes



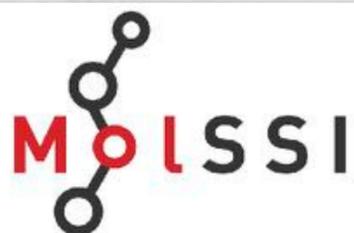
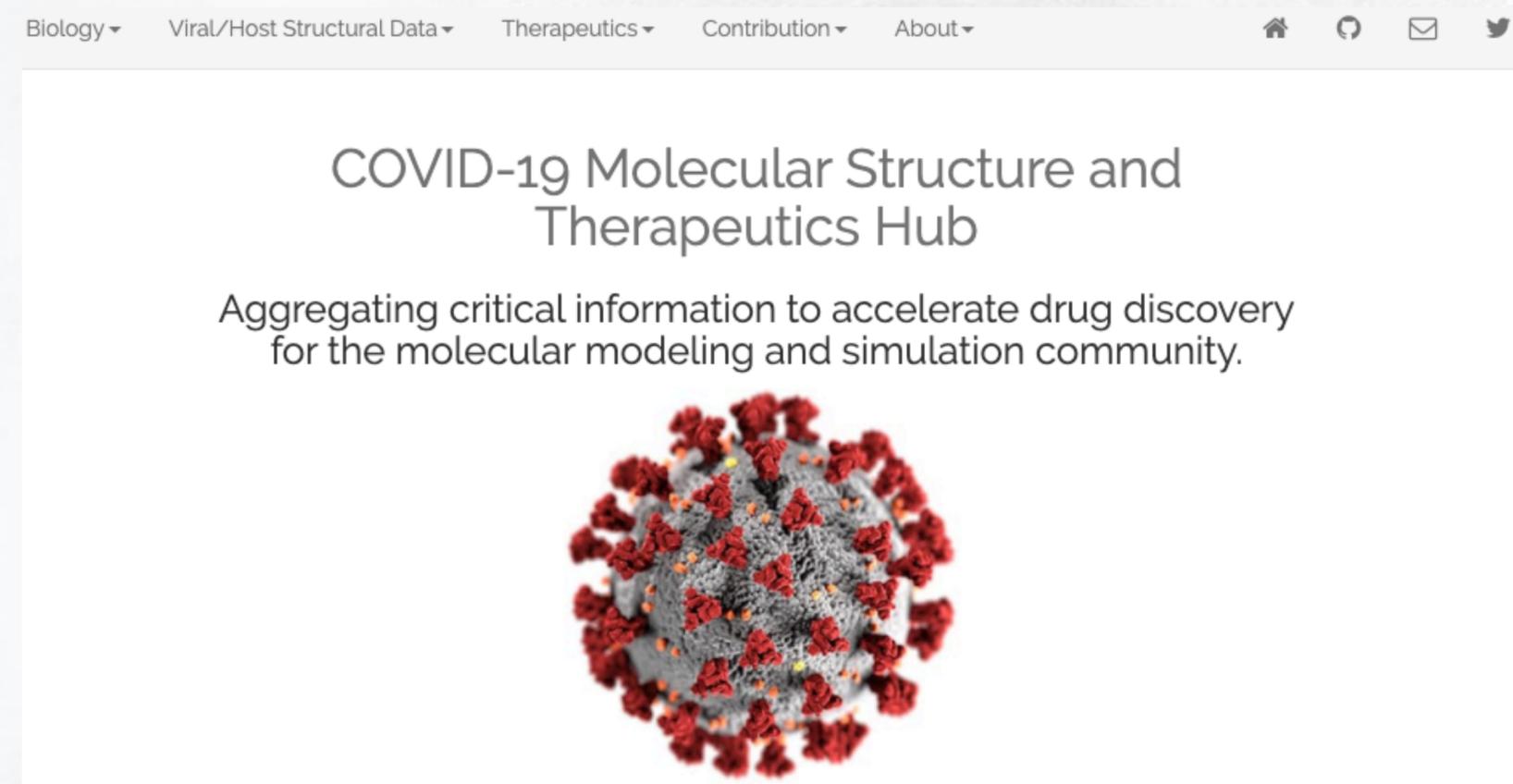
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- Simple descriptors:
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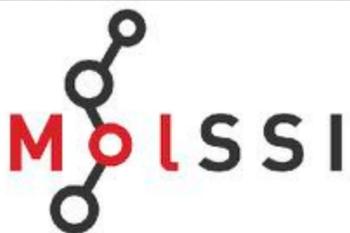
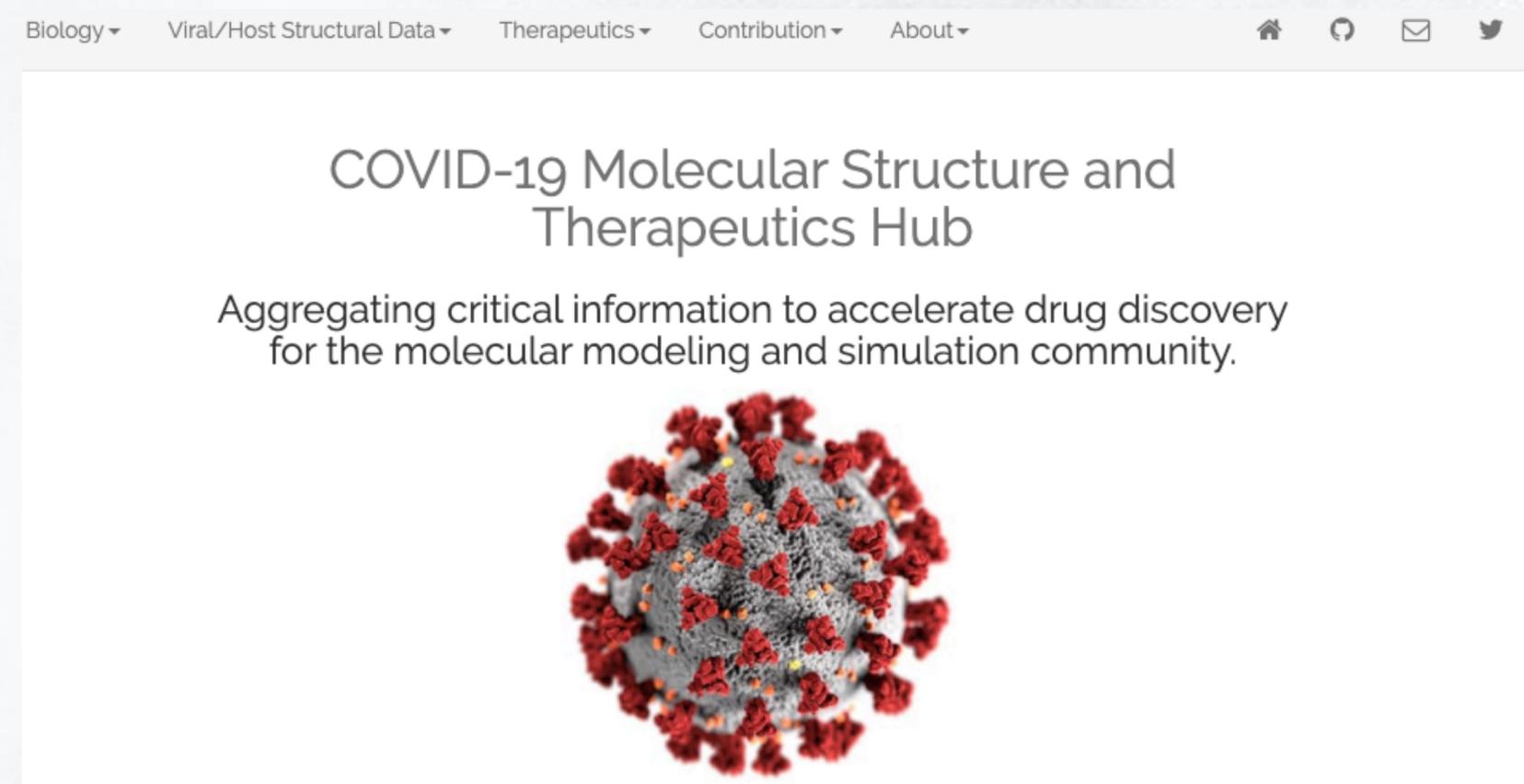
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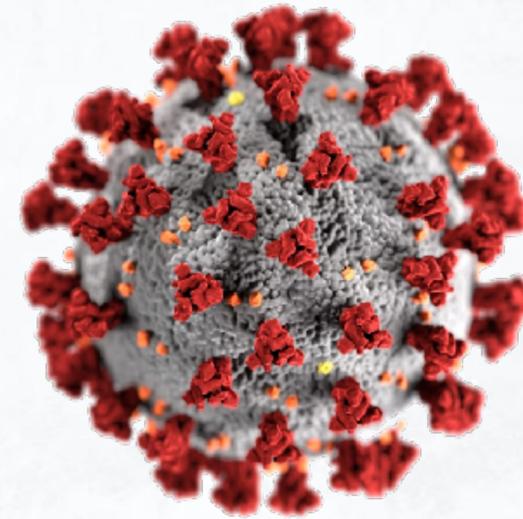


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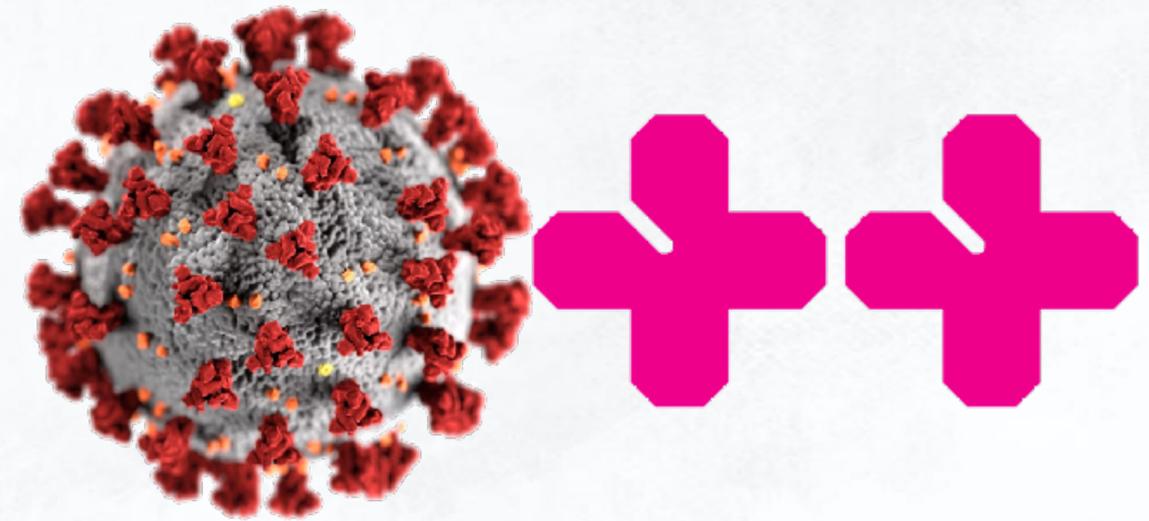
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  - Actual: Reluctant\* data additions
- Centralized or pointers to data
  - Can we link to it? We'll share it.
  - Can we quantify access? Even better.



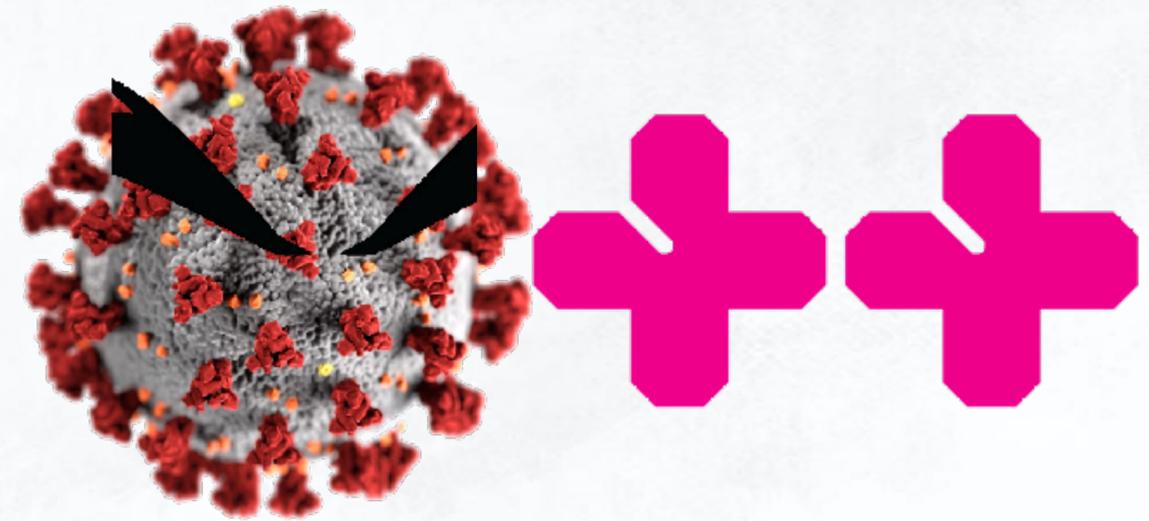
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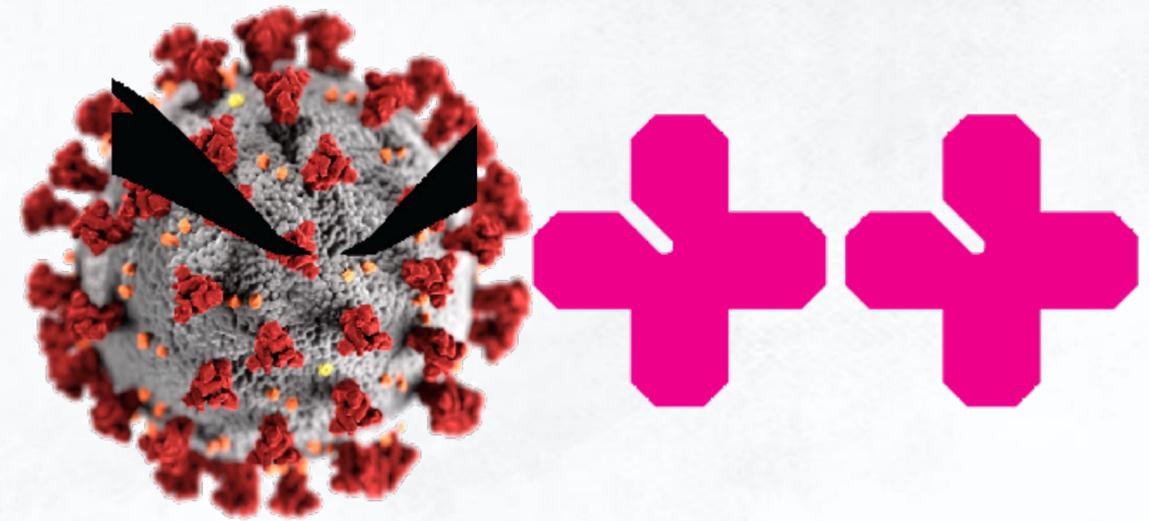


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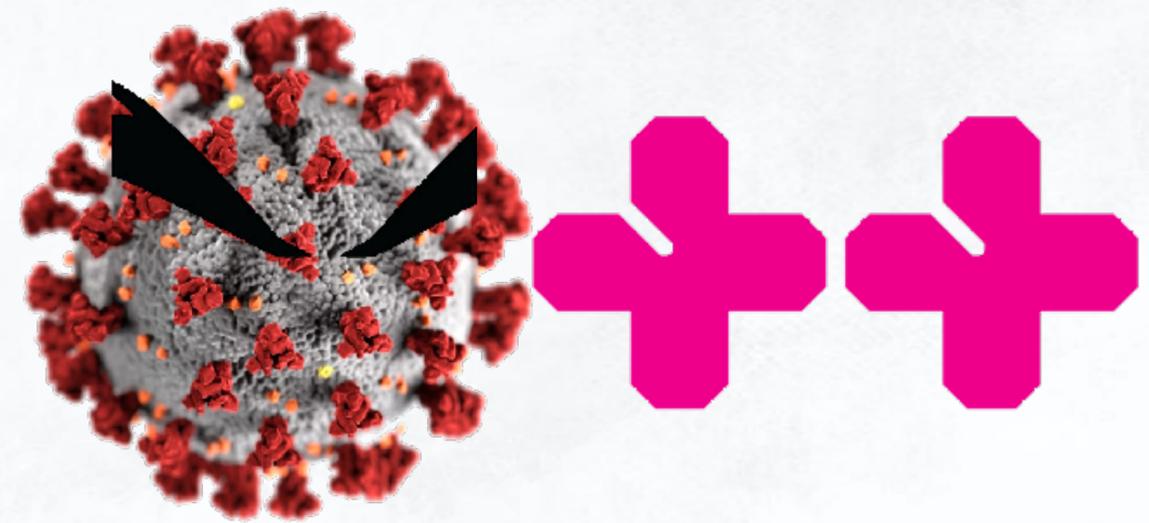
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- Encourage easy contribution



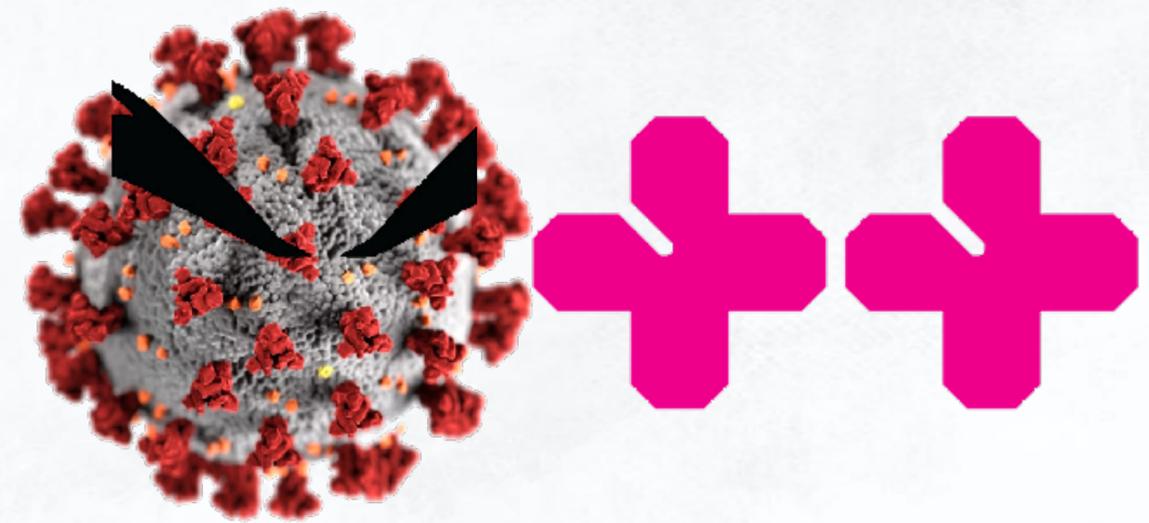
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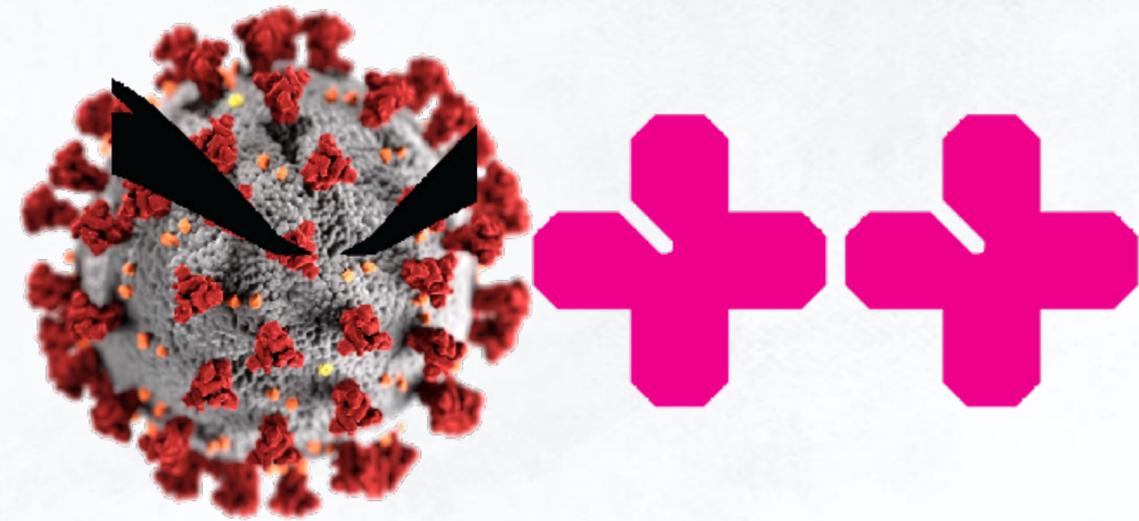
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- Encourage easy contribution
- Keep metadata simple with small iterations
- Highlight and promote research gaps



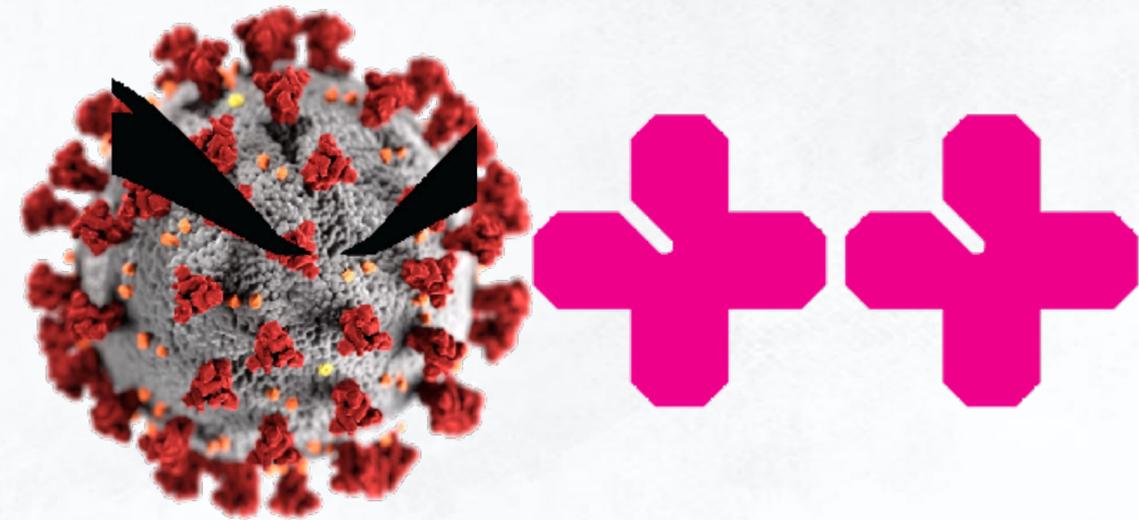
# What can we learn for the next research cohesion?

- Encourage easy contribution
- Keep metadata simple with small iterations
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- Network data then connect people



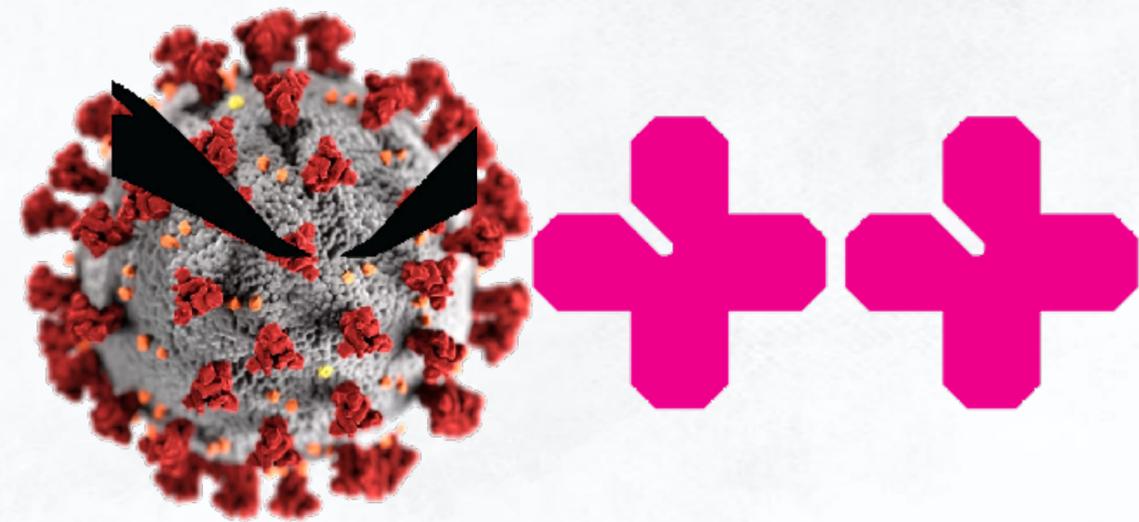
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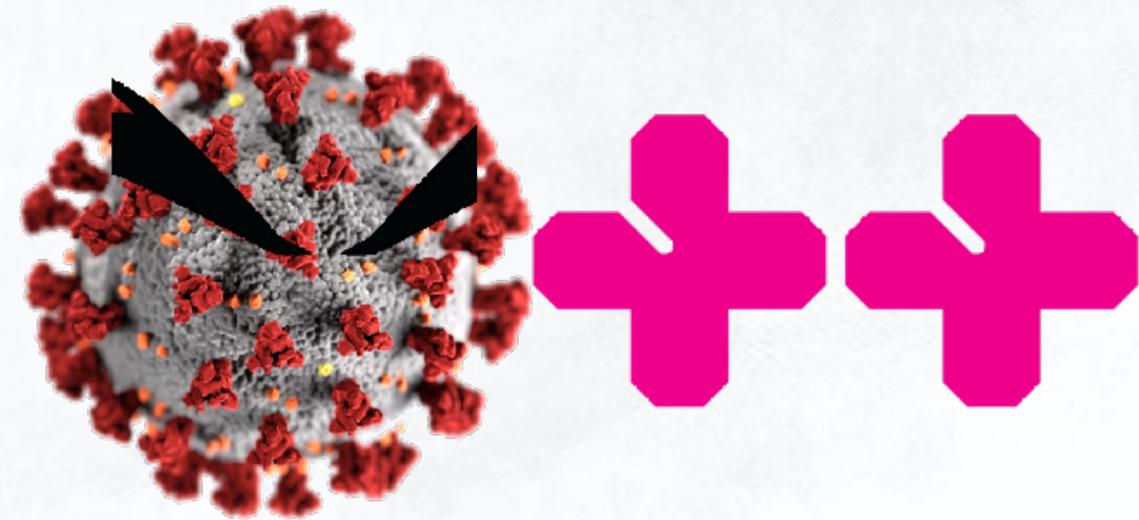
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- Replace review teams with outreach teams
- Secure large storage early with download metrics
- Bring in UX expertise



# Additional Thanks



Sam Ellis

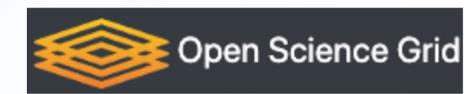


Andrew Abi-Mansour

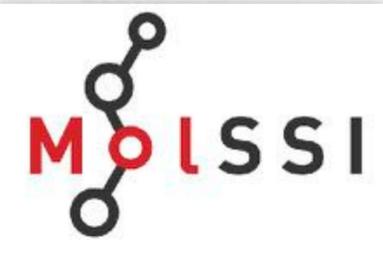
Levi Naden	MOLSSI	Andrew Abi-Mansour	MOLSSI	Sam Ellis	MOLSSI
Ben Richard	MOLSSI	Doss Alkaway	MOLSSI	Jessica Nash	MOLSSI
Daniel Crawford	MOLSSI	Teresa Heach-Gordon		Coelia Clementi	
Rumiko Amari		Achran Muthellane		Julia Chodas	
Eric Goldberger		Rafal Wlasiak		Alan Payne	
Elisita Karki		Hanna Bruce-Macdonald		Chris Mancinelli	
Melissa Roby		Julme Rodriguez-Cuama		Andrea Thom	
Yunyun Gao		Michael Fay		Lin Heo	
Richard Friesner		Peter Televant		Syma Khalid	
Shantana Jha		Dusan Apostolov	bioExcel	Bar de Groot	bioExcel
Erik Lindahl	bioExcel	Emily Morozov	bioExcel	Gerit Grothof	bioExcel
Medina Oezgen		Thomas M. Hermans		Ernst Wilmanns	
Heidi Elmer		Frank Wurstwein		Tim Smith	
Lukas Hennrich		Markus Dising		Alessandra Forti	
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Jean-Philip Flouquet		Callie Rodgers-Chappell		Thomas Subtil	
Charlie Brooks		William Jorgensen		D. E. Shaw Research	D E Shaw Research
Tristan Crell		Vincent Voez		Marek Tajik	
Yuki Sugita		Tomoko Komatsu		Natali Okamoto	
Stefan Solandt-Reyes		Alexandre Honvin		Yang Zhang	Zhang Lab
Changxin Zhang	Zhang Lab	Voth Group		Sonweda Group	
David Huggins	Amazon Web Services	Kurzban Group		Daniel McKay	VENTUS THERAPEUTICS



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# Thank you for listening!