Applied Mathematics and Informatics In Drug Discovery (2021)

We are following **24 coronavirus treatments** for effectiveness and safety:



Coronavirus Drug and Treatment Tracker, New York Times, Visited on 23.09.2021

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Cases of SARS-COV-2 infection, colored per capita, New York Times, visited on 23.09.2021







Data as of: 13.09.2021 16:00:04

EUA = Emergency Use Authorization Source: <u>Biotechnology Innovation Organization (BIO)</u>



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Contact the author

Today's goals



- Introduction to the course
- Why mathematics and informatics matter for drug discovery?
- Two views of the drug discovery and development process
 - The linear view
 - The multiscale modelling view



Purity https://xkcd.com/435/



This course aims to bring people together and to promote interdisciplinary research

Our strength: we have a diverse classroom!



U N I B A S E L

Course information

- Lecturer: Jitao David Zhang
 - jitao-david.zhang@unibas.ch
- Website: <u>AMIDD.ch</u>
- Thirteen lectures this semester
 - Introduction to drug discovery (1 session)
 - Molecular level modelling (2 sessions)
 - Omics- and cellular modelling (2 sessions)
 - Organ- and system modelling (2 sessions)
 - Population modelling and reverse translation (2 sessions)
 - Dies Academicus Ask Me Anything (optional)
 - Invited guest speakers (1 sessions)
 - Near-end-term presentations (2 sessions)

- Fridays 12:15-14:00
- Slides, exercises, pre-reading and post-reading articles, as well as videos, are shared on the course's website <u>http://www.amidd.ch</u>.
- No exercise hours. One-to-one virtual sessions are possible upon request and reservation.
- The final note is given by participation (40%), presentation (30%), and project work (30%).
- The project work will be about concepts that we learned together and their applications in practice. Details will follow.
- Questions?

I am glad to share my expertise in drug discovery, and to learn from you!





Teaching is my personal engagement. My opinions and views do not necessarily reflect those by F. Hoffmann-La Roche, my employer.

Please be aware of my biases and limitations.

- I am neither a mathematician nor a computer scientist by training. I am a computational biologist working in drug discovery.
- I see my task is to share with you the mathematical concepts and computational approaches used in drug discovery that I find beautiful and useful.
- I look forward to learning from you mathematics and other expertise that I did not know.



Why applied mathematics and informatics in drug discovery, why now?

- Now is the best time in human history to fight diseases
- Applied mathematics and informatics approaches are indispensable to modern drug discovery
- Applied mathematics and informatics will join interdisciplinary efforts to transform drug discovery in the coming decades

The history of *Homo sapiens* is a history of living with, understanding, and fighting diseases



Trypanosomes

Plasmodium

Tropical diseases

~500,000 years ago



A young patient of smallpox, the first eradicated infectious disease

Hygiene, vaccination, and antibiotics ~250 years ago





Chloral hydrate, the first synthesized drug

Pharmaceutical drugs

~150 years ago

Nobel prize laureates 2018, immune checkpoints, and drugs targeting the pathways

Personalized precise healthcare

~20 years ago



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Now is the best time in human history to fight diseases UNI BASEL cultured stem cells muscle cells intestinal cells U1-C More biological, dsDNA **CRISPR-CAS9** gene chemical, and Cleavage editing system ipid-polymer hybrid nanoparticle medicinal knowledge (Lipid bilayer) iver cell blood ce cardiac cells nerve cell New New therapeutic Risdiplam, a **Stem cells** disease-modelling SMN2 splicing modalities modifier systems mRNA delivery with nano lipid particles Digitalization of Better algorithms, molecular models, and more mechanisms in living **Comprehensive Sensing** computing resources Single-cell biology, organisms

Single-cell biology, multi-modal omics profiling, and imaging

Novel drugs approved by the FDA's Center for Drug Evaluation and Research (CDER) in 2020

Top three modalities by approval in 2020

- Small molecules (molecular weight under 900 daltons)
- Antibodies
- Oligonucleotides



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A zoo of modalities









Small molecule Monoclonal antibody Oligonucleotides

Bispecific antibody



Chimeric Antigen Receptor (CAR) T-cells



mRNA vaccines

Comparison with historical data: most new drugs are small molecules, while antibodies and oligonucleotides are rising



2019 FDA drug approvals, Asher Mullard, Nature Reviews Drug Discovery, https://www.nature.com/articles/d41573-020-00001-7 mAb: monoclonal antibodies; ADC: antibody-drug conjugate. 2020 FDA drug approvals, Asher Mullard, Nature Reviews Drug Discovery, https://www.nature.com/articles/d41573-020-00001-7 mAb: monoclonal antibodies; ADC: antibody-drug conjugate. 2020 FDA drug approvals, Asher Mullard, Nature Reviews Drug Discovery, https://www.nature.com/articles/d41573-020-00001-7 U N I B A S E L

New drug approvals vary between disease areas







Top pharmaceuticals by retail sails in 2020

Poster compiled by the Jon Njardarson group/ University of Arizona. Source: <u>https://njardarson.lab.arizona.edu</u>, accessed on 24.09.2021. Citation: J. Chem. Ed. 2010, 87, 1348

Questions:

- 1. How many are small molecules, proteins, and oligonucleotides each? What other modalities are there?
- 2. What patterns do you observe? Do you have explanations for these patterns? 18



How Do You Make A Drug?

It sounds simple, but...



Source and copyright: roche.com, assessed on 1.2.2019



Even if we understood everything, the search space of drug hunting is huge

Prerequisites to make a good drug that works



- Potency
- Safety
- Efficacy
- **Diagnosis**: doctors' judgement + *biomarkers*
 - Biomarkers are informative features derived from measurements of patient or patient material, *e.g.* blood chemistry, genetic make-up, imaging, *etc.*
- Other criteria: commercial rationale, development ability, intellectual property, *etc.*

Success in drug discovery is determined by potent, safe, efficacious drugs and accurate diagnosis

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The essence & THE challenge of Drug Discovery



Constrained optimization and decision making based on incomplete, noisy and heterogeneous data, and limited prior knowledge.

The linear view of drug discovery





Adapted from Paul *et al.* "How to Improve R&D Productivity: The Pharmaceutical Industry's Grand Challenge." Nature Reviews Drug Discovery, 2010



Risks and costs associated with each stage of the linear view of drug discovery



pTS: probability of technical success. **WIP**: work in progress; **Capitalized cost**: Out-of-pocket cost corrected for cost of capital, standard for long-term investments; **Out-of-pocket cost**: total cost required to expect one drug launch, taking into account attrition, but not the cost of capital; **Cost of capital**: annual rate of return expected by investors based on the level of risk of the investment. Paul *et al.*, Nature Reviews Drug Discovery, 2010.

Increasing cost and decreasing return of investment in drug discovery



Modified from Smietana *et al.* "Improving R&D Productivity." Nature Reviews Drug Discovery, 2015

Finding new drugs has become more challenging and expensive

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R&D of leading pharmaceutical companies neccesates both high spending and huge organizations



Schuhmacher, Alexander, Lucas Wilisch, Michael Kuss, Andreas Kandelbauer, Markus Hinder, and Oliver Gassmann. "R&D Efficiency of Leading Pharmaceutical Companies – A 20-Year Analysis." *Drug Discovery Today* 26, no. 8 (August 1, 2021): 1784–89. <u>https://doi.org/10.1016/j.drudis.2021.05.005</u>.







Danger + Opportunity

Applied mathematics empowers drug discovery in many ways

Applied mathematics *in drug discovery* is not a definable scientific field but a human attitude.

Quantitative critical thinking in communication and decision making Statistics, Data Mining and **Applied Combinatorics** Dynamical Systems Causal inference Machine Learning and Graph Theory Ordinary / Partial/ Stochastic Molecular, Quantum, and Multiscale modelling Network Analysis **Continuum Mechanics Differential Equations**

Richard Courant (1888-1972)





The alternative, multiscale-modelling view of drug discovery



Zhang, Jitao David, Lisa Sach-Peltason, Christian Kramer, Ken Wang, and Martin Ebeling. 2020. "Multiscale Modelling of Drug Mechanism and Safety." Drug Discovery Today 25 (3): 519–34. <u>https://doi.org/10.1016/j.drudis.2019.12.009</u>.





Introduction to Applied Mathematics and Informatics in Drug Discovery (*AMIDD*)

A course series at DMI, University of Basel

- Introduction to drug discovery
- Molecular modelling
 - Biological sequence analysis
 - Protein sequence and structure
 - Molecular modelling and dynamics
- Omics and cellular modelling
 - From drug-target interactions to networks
 - Gene expression profiling
 - Cell-based phenotypic drug discovery

- Mathematical modelling
 - Principles and applications of modelling in pharmacology
 - Pharmacokinetics (PK) and pharmacodynamics (PD) modelling
 - Clinical pharmacology and pharmacometrics
- Population modelling
 - Non-linear mixed-effect models (NLMEs)
 - Essentials of clinical trials
- Guest lectures
- Your presentations

It is hoped that AMIDD builds a bridge between students and quantitative aspects of drug discovery

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Luca Piali	Lue Dai
John Young	Ravi Jagasia
Lisa Sach-Peltason	Marco Prunotto
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Tony Kam-Thong	Detlef Wolf
Corinne Solier	Ken Wang
Thomas Singer	Nikolaos Berntenis



















Maria Anisimova

Lorenzo Gatti

- Erhard van der Vries
- Ab Osterhaus
- Nevan Krogan

Oliv Eidam













Conclusions and perspectives



- It is now probably the best time in human history to join the fight against diseases.
- We learned about modalities and the drug discovery and development process.
- Interdisciplinary research, especially applying mathematical approaches and tools to biological, chemical and medicinal questions, is imperative to fill the knowledge gaps and to make potent, safe, and efficacious drugs and to perform accurate diagnosis.
- Mathematics and informatics will continue transforming drug discovery
 - From correlation to causation
 - From qualitative description to quantitative prediction
 - From trial-and-error to systematic understanding
 - From population inference to individual prediction and continuous intervention
 - From descriptive studies of biology to predictive and counterfactual models
 - From observations to engineering and synthesis of the biological system
- In the AMIDD course, we will learn some basic concepts and tools we use to model interactions between biological systems and drugs at multiple levels (*multiscale-modeling of drug mechanism and safety*). The Mathematical and Computational Biology in Drug Discovery (MCBDD) course in spring semesters builds upon the basics in AMIDD and introduces advanced topics.



Offline Activities



OA1: Questions on the video on Herceptin by Susan Desmond-Hellmann

Link to the video

Questions for the video

- 1. What is the **indication** of *Herceptin*? What is its generic (USAN, or United States Adopted Name) name?
- 2. What is the gene target of Herceptin?
- 3. In which year was the **target** of Herceptin described? When was Herceptin **approved**?
- 4. What was the **improvement** of Herceptin compared with earlier antibodies?
- 5. Why does a **biomarker** matter besides developing drugs?
- 6. In the clinical trial of *Herceptin* for **metastatic breast cancer**, how much improvement in the **median survival** did Herceptin achieve? And how much improvement is in the **adjuvant setting** (Herceptin applied directly after operation)?

Questions for further thinking

- Susan Desmond-Hellmann summarizes successful drug development in four aspects: (1) having a deep understanding of the basic science and the characteristics of the drug, (2) targeting the right patients, (3) setting a high bar in the clinic, and (4) working effectively with key regulatory decision makers. Where do you think mathematics and computer science play a crucial role?
- She emphasized the importance of collaboration. What skill sets do we need for that?
- How do you like her presentation? Anything that you can learn from her about presentation and storytelling?

OA2: Required and recommended reading



[Required]

Principles and workflow of early drug discovery:

Hughes, JP, S Rees, SB Kalindjian, and KL Philpott. 2011. "Principles of Early Drug Discovery." British Journal of Pharmacology 162 (6): 1239–49. <u>https://doi.org/10.1111/j.1476-5381.2010.01127.x</u>.

[Recommended]

History of drug discovery and the rise of pharmaceutical company:

Jones, Alan Wayne. 2011. "Early Drug Discovery and the Rise of Pharmaceutical Chemistry." *Drug Testing and Analysis* 3 (6): 337–44. <u>https://doi.org/10.1002/dta.301</u>.

Mathematics and biology:

Cohen, Joel E. 2004. "Mathematics Is Biology's Next Microscope, Only Better; Biology Is Mathematics' Next Physics, Only Better." PLOS Biology 2 (12): e439. <u>https://doi.org/10.1371/journal.pbio.0020439</u>.

Extensive reading about history of medicine:

• <u>Taking the Medicine: A Short History of Medicine's Beautiful Idea, and our Difficulty Swallowing It</u> by Druin Burch (ISBN: 1845951506, ISBN13: 9781845951504)

OA3: Preparation for Lecture 2 and 3



If you want to learn more about the **Central Dogma of Molecular Biology**:

- 1. If you are a film person, watch <u>this video on Youtube about the Central Dogma of Life</u>, and <u>this</u> <u>animated film</u>.
- If you are a reading person, read <u>Biology Briefs</u>, a six-part introductory article series run by *The Economist*, or if you prefer textbooks, read <u>DNA & The Central Dogma of Biology</u> by Prof. Henry Jakubowski in the Biology LibreTexts.

If you are already familiar with the Central Dogma and wishes to refresh relevant **mathematical knowledge**:

- 1. Find on Youtube or other websites introductory courses to probability (like this)
- 2. Read the first chapter of *Biological Sequence Analysis* by Richard Durbin *et al.*, <u>available here</u>