# Introduction to Applied Mathematics and Informatics in Drug Discovery (Autumn Semester 2019-2020)

Jitao David Zhang<sup>1,2</sup>

## Time and place

The lecture takes place on Fridays between 12:15 and 14:00 in Seminarraum 5.002, Department of Mathematics and Informatics, Spiegelgasse 1, 4051 Basel.

All course materials, unless otherwise specified, are shared via course's website, <a href="http://amidd.ch">http://amidd.ch</a>.

## **Syllabus**

- 1. Drug discovery: an overview (20.09.2019)
- 2. Biological sequence analysis (27.09)
- 3. Protein structure and function (4.10.)
- 4. Chemical structure representation and search (11.10)
- 5. Molecular interaction and modelling (18.10.)
- 6. Omics: genomics, transcriptomics, and proteomics (25.10.)
- 7. PK/PD and PBPK modelling (1.11.)
- 8. Bayesian modelling, machine learning, and causal inference (8.11.)
- 9. Multiscale modelling of drug mechanism and safety (15.11)
- 10. Guest speakers: Dr. Nicolas Frey and Dr. Lucy Hutchinson (titles to be announced) (22.11.)
- 11. \*Dies academicus no lecture\* (29.11.)
- 12. Guest speakers: Dr. Kaspar Rufibach and Dr. Benjamin Ribba (titles to be announced) (6.12.)
- 13. Student presentation (I) (13.12.)
- 14. Student presentation (II) (20.12.)

Changes, if necessary, be communicated during the course.

#### Assessment

The final note is given by participation (20%), presentation (30%), and an oral examination (50%). The oral examination will be about concepts that we learned together, and about

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explaining mathematical concepts (or concepts in your domain of experts) to a layman - that is, your lecturer.

### Further information

We focus on interdisciplinary research with mathematics as the language and informatics as the tool.

I do not offer exercise hour yet. Pre-reading and post-reading articles, as well as videos, are shared and recommended.

Both slides and board are used for the course. Slides and notes are shared.

## Further questions?

Please contact David, the lecturer, by following ways:

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