

MSc in Bioinformatics & Systems Biology

Our international MSc programs are designed for students who desire focused training in the elements of biology, computer science, and information technology needed for a successful career in the exciting new discipline of Bioinformatics & Systems Biology. Students in our programs will receive comprehensive training in omics analysis, database design and management, software engineering and programming (including web-based development), simulation techniques and modeling, and data integration. Each student will apply their skills to a practical research project, where they will design and implement a solution to a real-world problem under the guidance of an experienced mentor.

PROGRAM LEARNING OUTCOMES

CURRICULUM

1) Solves biological problems using appropriate bioinformatics and systems biology approaches

1A: Explains foundations of biology, computing, statistics, and mathematics relating to bioinformatics and systems biology. 1B: Analyzes high-throughput biological data by integration of knowledge from different disciplines (biology, computing, statistics, and mathematics)

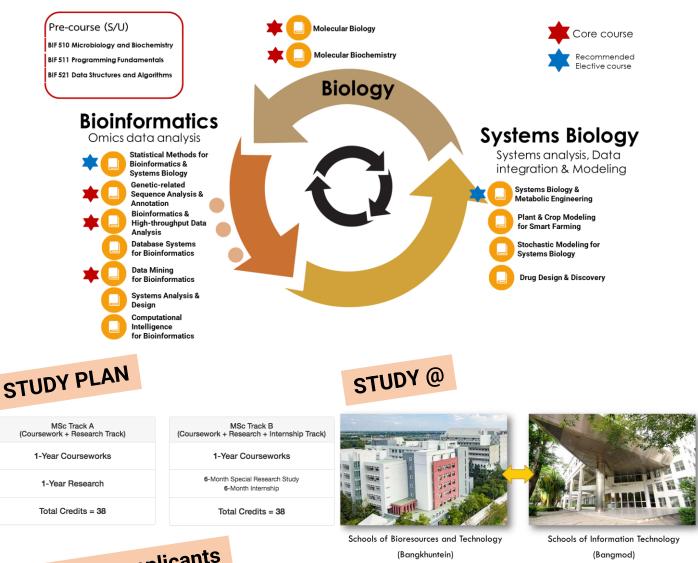
1C: Formulates research questions or hypotheses based on guidelines with an understanding of the role of bioinformatics and systems biology in it.

1D: Solves research problems/tests hypothesis by selecting effective bioinformatics and systems biology methods.

2) Communicates accurate information relating to bioinformatics and systems biology to diverse stakeholders

3) Takes part competently in diverse teams to accomplish a common goal by sharing their own ideas and accepting others' opinions

4) Values self and others with an understanding of ethical and social issues



Criteria for applicants

A bachelor's degree in biological sciences, computer science, computer engineering, medical sciences, chemistry, mathematics, statistics, or related disciplines.

🙂 (Optional) English test score (TOEFL iBT, IELTS or TETET)

Bioinformatics and Systems Biology Research

Plant

Modeling a (crop) plant to

predict the phenotype under the exposed

condition. Omics data

analysis, biological network

reconstruction and

mathematical modeling

have been applied to study

the dynamic regulation

inside plant cells, aiming to

precision science for tailor-

made yield and quality of

phyto-products.



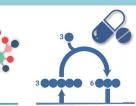


Metagenomic Analysis

Metagenomics is the study of genetic materials (of microbiome) derived directly from environmental samples using high-throughput sequencing technology. Microbiome is important for human, animal and plant health, including maintaining environmental balance.

Medical **Bioinformatics**

Bioinformatics plays an important role to analyze biomedical data for studying human health and diseases e.g. cancer, Alzheimer, Autism, etc. The research leads identifications of biomarkers for diagnosis, prognosis and prediction of drug response.



Drug Design Systems Biology and Discovery

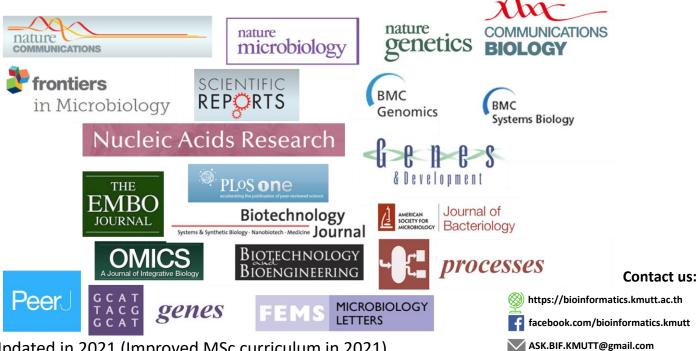
Bioinformatics & systems biology empower the research to gain insights into drug-receptor interactions, potential gene / protein targets, structures and functions of biomolecules, allowing the designing of new effective drugs and therapies while saving cost and time

Deep-tech Applications

- Personalized medicine
- Personalized nutrigenomics
- In silico drug discovery and design
- Microbiome for heath & well-being
- Smart farming & breeding
- Synthetic biology for future food



International journal publications from our academic staffs & students



Updated in 2021 (Improved MSc curriculum in 2021)