

TIGP Bio 2021 Spring Syllabus
Advanced Statistical Methods in Bioinformatics (S2)

Place: Room 107, New Building of the Institute of Information Science, Academia Sinica.

Time: Thursday 9:00-12:00

Chair: Dr. Grace S. Shieh (gshieh@stat.sinica.edu.tw)

Outline: Introduction to useful and advanced statistical methods in computational biology. The topics include: Analysis of next generation sequencing (NGS) Data (e.g., RNA-Seq and ChIP-Seq), maximum likelihood estimation, the EM algorithm, Bayesian inference, Monte Carlo methods, Resampling (Bootstrap & permutation test), Human Genetics, clustering and classification, dimension-reduction and missing data.

Textbook: N/A

Reference book: papers will be given by instructors

TA: N/A (Please refers to the lectures respectively shall you have any questions of each class)

E-mail: Contact the Lecturers Directly.

Office hours: N/A

Grades: Midterm exam 50%. Final exam 50%.

Note: For **Non-BP student** to register/sit-in any BP course, it is required to gain course chair's permission and follow the steps:

(1) Submit the hard copy or PDF file of the completed [TIGP Bioinformatics Course Registration Consent Form](#) to the TIGP BP office

(2) Provide the information via the google form at [BP Class Enrollment Information](#).

The deadline for above requirement is **the 4th week** of each semester. Signature of corresponding BP Course Chair should be collected and incomplete form will not be accepted.

※Course grade will **NOT** be given (even class enrollment is completed at school) if fail to follow the above procedures.

For the most up-to-date syllabus, please visit <https://tigppb.iis.sinica.edu.tw/tigpbio/index.html>

Week	Date	Topics/Brief Description	sub-topics/Detail Descriptions	Lecturers
1	2021/02/25	Analysis of NGS data I	N/A	Dr. Hao Ho
2	2021/03/04	Analysis of NGS data II	N/A	Dr. Hao Ho
3	2021/03/11	Maximum likelihood estimates and the EM algorithm I	Maximum likelihood estimates	Dr. Yen-Tsung Huang
4	2021/03/18	Maximum likelihood estimates and the EM algorithm II	The EM algorithm	Dr. Yen-Tsung Huang
5	2021/03/25	Bayesian methods	Bayesian methods	Dr. Grace S. Shieh
6	2021/04/01	Monte Carlo Markov Chains	Monte Carlo Markov Chains	Dr. Shin-Sheng Yuan
7	2021/04/08	Resampling methods I	Bootstrap	Dr. Grace S. Shieh

8	2021/04/15	Midterm Exam		
9	2021/04/22	Resampling methods II	Permutation test, jackknife	Dr. Grace S. Shieh
10	2021/04/29	Clustering, classification and data visualization (I)	Clustering: K-Means, Hierarchical Clustering, Cluster Validation	Dr. Han-Ming Wu
11	2021/05/06	Clustering, classification and data visualization (II)	KNN, Decision Tree, Support Vector Machine, Data visualization in R	Dr. Han-Ming Wu
12	2021/05/13	Statistics in Human Genetics I	Concept of Statistical Genetics, International Genomic Projects, Linkage Analysis	Dr. Hsin-Chou Yang
13	2021/05/20 @Webex	Statistics in Human Genetics II	Genetic Association, Gene-Gene interaction, Rare Variant Analysis	Dr. Hsin-Chou Yang
14	2021/05/27 @Webex	Advanced regression and dimension reduction I	N/A	Dr. Shin-Sheng Yuan
15	2021/06/03 @Webex	Advanced regression and dimension reduction II	N/A	Dr. Shin-Sheng Yuan
16	2021/06/10	Final Exam		