

## **BIM3007 Computational Genomics and Proteomics (2022-2023 Term 1)**

### **Assignment #3**

**Note:** Please write down your name and student ID when you submit the assignment to the blackboard e-learning system.

**Deadline:** Please submit your assignment through the system before **23:59:59 4<sup>th</sup> Dec**, 2022  
(Delayed submission is not allowed for any reasons)

**Percentage:** 10%

**Purpose:** to enhance the learning outcomes for the topics in “Association tests”, “Quality control” and “Genome-wide association study”.

Please download the data for this assignment through:

[https://connecthkuhk-my.sharepoint.com/:f/g/personal/yfwangbm\\_connect\\_hku\\_hk/EIBy4U1nQ11JpGHn0J9wvjoBtpjv8e45sxBJluOmKkY3Q](https://connecthkuhk-my.sharepoint.com/:f/g/personal/yfwangbm_connect_hku_hk/EIBy4U1nQ11JpGHn0J9wvjoBtpjv8e45sxBJluOmKkY3Q)

- (1) How many samples are included in the dataset? (10 points)
- (2) How many variants are included in the dataset? (10 points)
- (3) Please perform quality control for samples and record the details in each step. (20 points)
- (4) Please perform quality control for genetic variants and record the details in each step. (20 points)
- (5) Please perform association tests for the QCed data and illustrate the results using Manhattan plot. (30 points)
- (6) Please show the QQ plot for the association results and calculate the inflation factor. (10 points)