Technical Specifications Report for Pharmacogenomics Of HLA Variants in 4 Asian populations

Computer facilities
- Acer E5-511, Windows 10, 64bits
- HackBio's remote server
- Linux terminal on https://shell.cloud.google.com/

Softwares
- R
- RStudio
- ggplot
- PLINK
- Winscp

Datasets
- Complete_1000_genomes_sample_list_.tsv for all populations
- Asia.bed - binary file contains data on individual identification and genotype
- Asia.bim -text file, contains data on SNPs position
- Asia.fam - text file, contains data on family relationship and gender

Metadata from complete 1000 genomes list