

## Adjustment

Genotype file  
Phenotype file  
GWAS type  
Filter values

Input config file

Cleaning

Filtering

Formating

## Multi Analysis

*GWASpoly*

*SHESis*

*Plink*

*Tassel*

## Integration

Summarizing

Reporting

Output files

Score tables  
Markers chart  
SNPs profiles  
Manhattan/QQ plots

