

Adjustment

Reading
Config
file

Ploidy
Geno/Pheno files
GWAS model
Significance
Correction
N associations
QC filters

Data preprocessing

Read

Clean

Filter

Format

Multi Analysis

Polyploids

GWASpoly

SHEsis

Diploids

Plink

Tassel

Integration

Scan

Summarize

Report

SNPs tables

Shared SNPs

SNPs profiles

Associations

Shared chromosomes