This github contains the raw data and demonstration used in the original publication of OATH.

**1 How to get the data**

Please either "git clone <https://github.com/gc5k/OATH.git>" to have the whole data cloned to your local computer, or download it (notice the "clone or download”, a green button?)

**2 System requirement**

OATH has already been realized in GEAR [GEnetic Analysis Repository; gear.jar]. More details about GEAR can be found <https://github.com/gc5k/GEAR/wiki>.

gear.jar requires java installed. To check the details, type "java -version". If it shows "java 1.7" or "java 1.8", then your computer is ready; otherwise please download java online and install it first (from oracle website).

**3 Three demonstrations included**

**a)** Run Figure2Demo.R.  
DemonFig2 has an R script to generate Fig 2 as in the paper; and also it shows the kernel of OATH.

**b)** Run ArabAnalysis\_Fig6.R.  
ArabData has the cleaned genotype data for generating results for Fig 6. Although genotype data are not really needed by OATH, to provide a more comprehensive experience for users we still include the data for 295 Arabidopsis lines.

**c)** Run GWAMA\_Fig7.R.  
GWAMA analysis. Naive summary statistics provided two Chinese GWAS cohorts: NAcohort and SLEcohort.