

General information on the code

Code in folder “general” is used in several projects; code in folder “gabriela” only for the data analysis of the GABRIELA study.

Documentation on the structure of the code can be found in the README-files in the several subfolders.

Data cannot be provided for legal reasons.

Code contains paths that are specific to our server and folder structure; these parts would have to be adjusted accordingly.

All paths containing “storage/cmbstore” have been and should be run on a server;

- analyses involving literature SNPs were run on a server with about 50 GB RAM and 24 cores
- multivariate analyses (glmnet, ranger, ipflasso) were run on a server with 512 GB RAM and 64 cores

In general, the numbering of the code file names shows in which order the code should be run.