The “GWASnetwork\_discoverableGenesOnly\_2021\_10\_14.RData” file is too large to upload in GitHub. Please find this file in the following link of the Harvard Dataverse: <https://nam12.safelinks.protection.outlook.com/?url=https%3A%2F%2Fdataverse.harvard.edu%2Fdataset.xhtml%3FpersistentId%3Ddoi%3A10.7910%2FDVN%2FB32FRP&data=05%7C01%7Ckhanmsr%40pitt.edu%7Cc22d973dc2464e13bdd608db83e60a5f%7C9ef9f489e0a04eeb87cc3a526112fd0d%7C1%7C0%7C638248796268228403%7CUnknown%7CTWFpbGZsb3d8eyJWIjoiMC4wLjAwMDAiLCJQIjoiV2luMzIiLCJBTiI6Ik1haWwiLCJXVCI6Mn0%3D%7C3000%7C%7C%7C&sdata=Kr13ycc24z5BVkTQzUc7BcFSw9z1adMV1jB8gU0FVPE%3D&reserved=0> )

# The title is: objects files of the supplementary material-1 of metGWAS 1.0 (The objects folder of supplementary material 1 of metGWAS 1.0 platform ( https://github.com/saifurbd28/metGWAS-1.0)