The summary statistics can be downloaded are listed as below:

- MAGIC1000G_2hGlu_TA.tsv.gz: trans-ancestry meta-analysis summary statistics of 2 hour glucose adjusted for BMI
- MAGIC1000G_2hGlu_AA.tsv.gz: meta-analysis summary statistics of 2 hour glucose adjusted for BMI in African American ancestry
- MAGIC1000G_2hGlu_EAS.tsv.gz: meta-analysis summary statistics of 2 hour glucose adjusted for BMI in East Asian ancestry
- MAGIC1000G_2hGlu_EUR.tsv.gz: meta-analysis summary statistics of 2 hour glucose adjusted for BMI in European ancestry
- MAGIC1000G_2hGlu_HISP.tsv.gz: meta-analysis summary statistics of 2 hour glucose adjusted for BMI in Hispanic ancestry
- MAGIC1000G_FG_TA.tsv.gz: trans-ancestry meta-analysis summary statistics of fasting glucose adjusted for BMI
- MAGIC1000G_FG_AA.tsv.gz: meta-analysis summary statistics of fasting glucose adjusted for BMI in African American ancestry
- MAGIC1000G_FG_EAS.tsv.gz: meta-analysis summary statistics of fasting glucose adjusted for BMI in East Asian ancestry
- MAGIC1000G_FG_EUR.tsv.gz: meta-analysis summary statistics of fasting glucose adjusted for BMI in European ancestry
- MAGIC1000G_FG_HISP.tsv.gz: meta-analysis summary statistics of fasting glucose adjusted for BMI in Hispanic ancestry
- MAGIC1000G_FG_SAS.tsv.gz: meta-analysis summary statistics of fasting glucose adjusted for BMI in South Asian ancestry
- MAGIC1000G_FI_TA.tsv.gz: trans-ancestry meta-analysis summary statistics of fasting insulin adjusted for BMI
- MAGIC1000G_FI_AA.tsv.gz: meta-analysis summary statistics of fasting insulin adjusted for BMI in African American ancestry
- MAGIC1000G_FI_EAS.tsv.gz: meta-analysis summary statistics of fasting insulin adjusted for BMI in East Asian ancestry
- MAGIC1000G_FI_EUR.tsv.gz: meta-analysis summary statistics of fasting insulin adjusted for BMI in European ancestry
- MAGIC1000G_FI_HISP.tsv.gz: meta-analysis summary statistics of fasting insulin adjusted for BMI in Hispanic ancestry
- MAGIC1000G_FI_SAS.tsv.gz: meta-analysis summary statistics of fasting insulin adjusted for BMI in South Asian ancestry
- MAGIC1000G_HbA1c_TA.tsv.gz: trans-ancestry meta-analysis summary statistics of HbA1c
- MAGIC1000G_HbA1c_AA.tsv.gz: meta-analysis summary statistics of HbA1c in African American ancestry
- MAGIC1000G_HbA1c_EAS.tsv.gz: meta-analysis summary statistics of HbA1c in East Asian ancestry
- MAGIC1000G_HbA1c_EUR.tsv.gz: meta-analysis summary statistics of HbA1c in European ancestry
- MAGIC1000G_HbA1c_HISP.tsv.gz: meta-analysis summary statistics of HbA1c in Hispanic ancestry

- MAGIC1000G_HbA1c_SAS.tsv.gz: meta-analysis summary statistics of HbA1c in South Asian ancestry
- MAGIC1000G_HbA1c_Uganda.tsv.gz: meta-analysis summary statistics of HbA1c in Uganda

For each variant, we have provided the following information:

- chromosome: the chromosome that the variant is located on
- base_pair_location: the base pair location of the variant based in build 37
- effect_allele: the effect allele of the variant (+ strand)
- other_allele: the non-effect allele (+ strand)
- effect_allele_frequency: frequency of the effect allele
- beta: the effect size of the effect allele with respect to the trait being analysed
- standard_error: the standard error of the effect size
- p_value: the p-value of the variant/trait association
- sample_size: the number of individuals after QC being tested for association at a specific variant
- log10BF: the 10 based logarithm of Bayes factor of the variant/trait association
- het_p_value: the p-value of the heterogeneity test

^{*} Fasting insulin was natural log transformed.