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Participants Enrolled

N=3561

Bio-samples Collected

N=3470

Genome wide genotyping completed

N=3010

Did not pass genotyping quality control, N=460

Incomplete Phenotype Data, N=89

Genotype-Phenotype dataset

N=2921

Final Data Set

N=2850

Exclude 4-hr D/P Cr > 1.15, N=5

Exclude 4-hr D/P Cr < 0.3, N=4

Data Set

N=2912

Exclude genetically related participants, N=62