dbGaP Study Release Notes

Release Notes for NIMH Schizophrenia Swedish, phs000473.v2.p2
"Sweden-Schizophrenia Population-Based Case-Control Exome Sequencing"

For any questions or comments, please contact: dbgap-help@ncbi.nlm.nih.gov.

August 13, 2013 Version 1 Data set release date
October 26, 2016 Version 2 Data set release date

2016-10-26
Version 2 Data set release for NIMH Schizophrenia Swedish now available

This release includes phenotype tables, sequencing data including whole exome sequences, and genotype obtained from vcf files. Please refer to the latest study configuration report for a detailed description of each download component.

Consent group 1 (c1): General Research Use (GRU)

	Phenotype	SRA	Genotype
Subjects	12380	12380	12380
Samples	12380	12380	12380

1. **New Study Accession**
NIMH Schizophrenia Swedish version 1 phs000473.v1.p1 has been updated to version 2. The dbGaP accession for this study is phs000473.v2.p2. The participant number (p#) has not been changed in NIMH Schizophrenia Swedish, phs000473.v2.p2.

2. **Updated Datasets** (n=4; all variables have been updated)
a. phs000473.v2.pht002597.v2.p2.Schizophrenia_Swedish_Subject.MULTI;
b. phs000473.v2.pht002599.v2.p2.Schizophrenia_Swedish_Sample.MULTI;
c. phs000473.v2.pht002600.v2.p2.c1.Schizophrenia_Swedish_Subject_Phenotypes.GRU;
d. phs000473.v2.pht002601.v2.p2.c1.Schizophrenia_Swedish_Sample_Attributes.GRU

Additional notes for genotypes

1. Genotype data are accessioned under phg000773.v1. Please see "sample-info" component for genotyped samples, sample consent status and mapping of sample-vcf file.
2. Genotypes from 12380 samples are available in originally submitted variant-call-format (VCFv4.1), which are in the folder marked as "genotype-calls-vcf".
3. QC results, including consistency-checking results on sample gender and relatedness, are in the folder "phg000773.v1.SchizophreniaSwedish_Sklar_v2.genotype-qc.WES_markerset_grc37.MULTI.tar.gz".

Authorized Access (Individual Level Data)

Individual level data are available for download through the dbGaP Authorized Access System upon approval of the Data Access Request (DAR):

- https://dbgap.ncbi.nlm.nih.gov/aa/wga.cgi?page=login

FTP site
All data tables, data dictionaries, and documents will be housed under one directory for ease of downloading. The data_dict filenames have an added study version number (phs#.v#) and deleted participant set number (p#) from the table accession (pht#.v#). The var_report filenames have an added study version number (phs#.v#). In the var_report files, variables contain version numbers (phv#.v#) and summaries were created for each consent group (c#). These FTP files are available at:

- ftp://ftp.ncbi.nlm.nih.gov/dbgap/studies/phs000473/phs000473.v2.p1

2012-08-13
Version 1 Data set release for NIMH Schizophrenia Swedish now available

This release includes phenotype tables, SRA sequencing data, and genotype obtained from vcf files. Please refer to the latest study configuration report for a detailed description of each download component.

Consent group 1 (c1): General Research Use (GRU)

	Phenotype	SRA	Genotype
Subjects	5090	5090	5090
Samples	5090	5090	5090

Additional notes for genotypes

1. Genotype files are marked as “phg000347.v1” in the download components.
2. See components phg000347.v1.SchizophreniaSwedish_Sklar.genotype-calls-vcf.exome_vcf.c1. GRU.tar.gz and phg000347.v1.SchizophreniaSwedish_Sklar.genotype-calls-matrixfmt.exome_vcf.c1.GRU.tar.gz for original submissions, and converted matrix format files which are generated by dbGaP curator using the open-source library PLINK/SEQ.
3. The component phg000347.v1.SchizophreniaSwedish_Sklar.genotype-qc.exome_vcf.MULTI.tar.gz contains QC results from the toolset, PLINK.

FTP site

All data tables, data dictionaries, and documents will be housed under one directory for ease of downloading. The data_dict filenames have an added study version number (phs#.v#) and deleted participant set number (p#) from the table accession (pht#.v#). The var_report filenames have an added study version number (phs#.v#). In the var_report files, variables contain version numbers (phv#.v#) and summaries were created for each consent group (c#). These FTP files are available at:

- ftp://ftp.ncbi.nlm.nih.gov/dbgap/studies/phs000473/phs000473.v1.p1