

NLM Citation: SNP FAQ Archive [Internet]. Bethesda (MD): National Center for Biotechnology Information (US); 2005-. Ancestral Allele Data. 2005 Jul 7 [Updated 2014 Feb 18].

Bookshelf URL: https://www.ncbi.nlm.nih.gov/books/



Ancestral Allele Data

Created: July 7, 2005; Updated: February 18, 2014.

Where do I find ancestral allele information in dbSNP?

You can get these data from two the tab-delimited table dump files on the dbSNP FTP site: SNPAllele.bcp and Allele.bcp. When the SNPAllele.ancestral_flag is 1, the allele_id is an ancestral allele. To get the actual allele, please use the allele table (Allele.allele_id and Allele.allele). The DDL for shared tables is located in the shared_schema subdirectory, and the organism table DDL is located in the organism sub-directory (for this example I'm using human as the example organism). (uda: 3/21/08)

How do I download a flat file that contains the ancestral state of each SNP(when this ancestral state is known)?

You can the get ancestral state of each SNP (when this ancestral state is known) by using the following two tables from the FTP dump files:

- 1. Go to the organism_shared_data file located in the dbSNP FTP site, and download the Allele.bcp.gz file.
- 2. Go to your organism's (human, in this case) "organism data" file located in the dbSNP ftp site and download the "SNPAncestralAllele.bcp.gz" file.
- 3. You can get the column definitions for the SNPAncestralAllele and Allele.bcp.gz tables from the dbSNP main table.sql.gz, which is located in the shared_schema file of the dbSNP ftp site. (11/16/07)

Determining Ancestral Alleles

Allele.bcp.gz and SNPAncestralAllele.bcp.gz show the ancestral allele for rs10465407 is "A", but my multigenome alignment shows "G" is ancestral.

The method used to derive the ancestral alleles that are stored in SNPAncestralAllele may be different from your method or possibly outdated as more sequence evidence becomes available suggesting an alternate allele.

Please see the FAQ "How was the SNPAncestralAllele table derived?" (08/29/08)

How does dbSNP determine the ancestral allele of a SNP?

dbSNP ancestral allele data was submitted by Dr. Jim Mullikin of the National Human Genome Research Institute (NHGRI). Dr. Mullikin derived his ancestral allele data by by the comparison of human DNA to chimpanzee DNA, and his methodology is available online at PLoS. The last time Dr. Mullikin provided us with ancestral data was in May 2004.(6/13/07)

Could you please define the ancestor upon which SNP ancestral allele information is based?

2 SNP FAQ Archive

The National Human Genome Research Institute (NHGRI) determines the ancestral alleles for human SNPs by the comparison of human DNA to chimpanzee DNA. The methodology used for finding these ancestral alleles is available online at PLoS. (6/13/07)

How do I determine if a SNP is an ancestral allele?

Human SNP ancestral alleles are determined by comparison with primate DNA, so in general, they're based on chimpanzee sequence.

You can get ancestral allele data from dbSNP in a couple of ways:

- 1. If the ancestral state of the SNP in which you are interested is known, and is contained in dbSNP, look in the "Allele" section of the RefSNP page, which is located at the top right hand side of the report.
- 2. You can also get the ancestral state of a SNP(when this ancestral state is known) by using the following 2 tables from the FTP dump files:
 - a. Go to the organism_shared_data file located in the dbSNP FTP site, and download the Allele.bcp.gz file.
 - b. Go to your organism's (human, in this case) "organism data" file located in the dbSNP ftp site and download the "SNPAncestralAllele.bcp.gz" file.
 - c. You can get the column definitions for the SNPAncestralAllele and Allele.bcp.gz tables from the dbSNP main table.sql.gz, which is located in the shared_schema file of the dbSNP ftp site. (2/7/05)

I want the ancestral allele for a number of rsIDs, but can't find rsID numbers in Allele.bcp.gz and SNPAncestralAllele.bcp.gz.

If you go to the dbSNP Data Dictionary and search for "SNPAncestralAllele", you'll find that the rsID is stored as an integer in the first column of the table.

Here's an example to help you — say you want to find the ancestral allele for rs3:

- 1. Look for the number "3" in first column of the SNPAncestralAllele table and you'll find that the corresponding ancestral allele ID (located in the second column) is 7.
- 2. Now, look in the Allele table for the allele letter that corresponds to Ancestral allele ID = 7. In the allele table, the allele ID number is again located in the first column:

7 C 2003-02-22 01:11:00.0 2 2003-10-06 17:42:00.0

So you can see from this example, that the ancestral allele for rs3 is "C". (08/29/08)

Annotating Ancestral Alleles

Is there a program that I can use which will allow me to annotate ancestral alleles to existing SNPs? Would dbSNP accept submissions of such data?

Ancestral allele data is provided to dbSNP as part of the SNP assay data that is provided in dbSNP submissions, so we don't have a program that allows us to annotate ancestral alleles, and we don't currently have a mechanism for third party annotation of SNPs. (3/29/05)

Interpreting Ancestral Allele Field in refSNP Report

What does the "?" in the ancestral allele field in a refSNP report mean?

"?" in the ancestral allele field indicates that we do not know the ancestral allele for the SNP. (3/3/05)