

The following Genome in a Bottle files were used to design and analyze data for the NIST Genome Editing Consortium Mixture Study.

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## GENOME REFERENCE FILES:

### GRCh37 (hg19):

<https://ftp-trace.ncbi.nlm.nih.gov/ReferenceSamples/giab/release/references/GRCh37/>

- [hs37d5.fa.gz](#) 2021-04-12 09:56 851M
- [hs37d5.fa.gz.fai](#) 2021-04-12 10:31 2.7K
- [hs37d5.fa.gz.gzi](#) 2021-07-01 08:17 761K

### GRCh38 (hg38):

<https://ftp-trace.ncbi.nlm.nih.gov/ReferenceSamples/giab/release/references/GRCh38/>

- [GCA\\_000001405.15\\_GRCh38\\_no\\_alt\\_analysis\\_set.fasta.gz](#) 2021-04-11 22:58 845M
- [GCA\\_000001405.15\\_GRCh38\\_no\\_alt\\_analysis\\_set.fasta.gz.fai](#) 2021-04-12 10:31 7.6K
- [GCA\\_000001405.15\\_GRCh38\\_no\\_alt\\_analysis\\_set.fasta.gz.gzi](#) 2021-04-12 10:31 753K

## GIAB BENCHMARK VCF FILES:

### HG002 GRCh37 V4 Benchmark:

[https://ftp-trace.ncbi.nlm.nih.gov/ReferenceSamples/giab/release/AshkenazimTrio/HG002\\_NA24385\\_son/NISTv4.2.1/GRCh37/](https://ftp-trace.ncbi.nlm.nih.gov/ReferenceSamples/giab/release/AshkenazimTrio/HG002_NA24385_son/NISTv4.2.1/GRCh37/)

- [HG002\\_GRCh37\\_1\\_22\\_v4.2.1\\_benchmark.vcf.gz](#) 2020-12-07 11:37 168M
- [HG002\\_GRCh37\\_1\\_22\\_v4.2.1\\_benchmark.vcf.gz.tbi](#) 2020-12-07 11:37 1.6M
- [HG002\\_GRCh37\\_1\\_22\\_v4.2.1\\_benchmark\\_noinconsistent.bed](#) 2020-12-02 10:35 9.3M

### HG002 GRCh38 V4 Benchmark:

[https://ftp-trace.ncbi.nlm.nih.gov/ReferenceSamples/giab/release/AshkenazimTrio/HG002\\_NA24385\\_son/NISTv4.2.1/GRCh38/](https://ftp-trace.ncbi.nlm.nih.gov/ReferenceSamples/giab/release/AshkenazimTrio/HG002_NA24385_son/NISTv4.2.1/GRCh38/)

- [HG002\\_GRCh38\\_1\\_22\\_v4.2.1\\_benchmark.vcf.gz](#) 2020-12-07 11:41 149M
- [HG002\\_GRCh38\\_1\\_22\\_v4.2.1\\_benchmark.vcf.gz.tbi](#) 2020-12-07 11:41 1.6M
- [HG002\\_GRCh38\\_1\\_22\\_v4.2.1\\_benchmark\\_noinconsistent.bed](#) 2020-12-04 11:31 11M