



NCBI Variation Resources Survey

Welcome

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NCBI is conducting this short survey in order to gain insight for future improvements to our variation resources.

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NCBI Variation Resources Survey

Data Access

***1. How do you access genetic/genomic data from NCBI?**

- Download batch files (.xml, .vcf, flat files) from our FTP website for command line processing
- Individual Websites
- Both

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NCBI Variation Resources Survey

Command Line User

2. Do you use the NCBI FTP site?

- Yes
- No

3. Do you find the FTP site easy to use for NCBI Variation Resources?

- Yes
- No
- N/A

Please specify

4. What format do you use/would you like to use for batch SNPs?

- .xml
- .txt
- .vcf

Other (please specify)

5. How would you like to see SNPs organized? Would you like to see them presented in a transcript-specific manner?

6. Do you use ClinVar?

- Yes
- No

7. Do you use 1,000 Genomes data?

Yes

No

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Variation resources survey

NCBI Variation Resources Survey

Website User

2. Have you ever used GRCh38?

- Yes
 No

3. Do you use BLAST for Variant Analysis?

- Yes
 No

4. Which other NCBI resources do you use?

- Variation Viewer
 Variation Reporter
 BLAST
 ClinVar
 GenBank
 1,000 Genomes Browser
 dbSNP RefSnp page
 dbGaP
 RefSeq
 Pubmed
 GTR
 SRA
 GEO
 OSIRIS
 PheGenI

Other (please specify)

5. Do you get all the information you want from those resources?

Yes

No

If no, why not?

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NCBI Variation Resources Survey

Command Line and Website User

2. Do you use the NCBI FTP site?

- Yes
- No

3. Do you find the FTP site easy to use for NCBI Variation Resources?

- Yes
- No
- N/A

Please specify

4. What format do you use/would you like to use for batch SNPs?

- .xml
- .txt
- .vcf

Other (please specify)

5. How would you like to see SNPs organized? Would you like to see them presented in a transcript-specific manner?

6. Have you ever used GRCh38?

- Yes
- No

7. Do you use BLAST for Variant Analysis?

Yes

No

8. Which other NCBI resources do you use?

Variation Viewer

Variation Reporter

BLAST

ClinVar

GenBank

1,000 Genomes Browser

dbSNP RefSnp page

dbGaP

RefSeq

PubMed

GTR

SRA

GEO

OSIRIS

PheGenI

Other (please specify)

9. Do you get all the information you want from those resources?

Yes

No

If no, why not?

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NCBI Variation Resources Survey

Thank you!

10. Additional comments:

11. How likely is it that you would recommend NCBI variation resources to a friend or colleague?

Not at all likely - 0	1	2	3	4	5	6	7	8	9	Extremely likely - 10
<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>

Thank you for helping NCBI provide better resources to the scientific community!

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Done