NCBI Health Resources Survey

Welcome

OMB Control Number: 0925-0648

Expiration Date: 03/31/2018

Public reporting burden for this collection of information is estimated to average 4 minutes per response, including the time for reviewing instructions, searching existing data sources, gathering and maintaining the data needed, and completing and reviewing the collection of information. An agency may not conduct or sponsor, and a person is not required to respond to, a collection of information unless it displays a current valid OMB control number. Send comments regarding this burden estimate or any other aspect of this collection of information, including suggestions for reducing this burden, to NIH, Project clearance Branch, 6705 Rockledge Drive, MSC 7974, Bethesda, MD 20892-7974, ATTN: PRA (0925-0648). Do not return the completed form to this address.

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

Next

NCBI Health Resources Survey

Data Access

*** 1. In what primary role do you use NCBI Health Resources?**

- ○ Clinician
- ○ Scientific/Clinical Researcher
- ○ Student
- ○ Health consumer
- ○ Other (please specify)

*** 2. What type of content/information are you primarily interested in?**

- ○ Conditions & Tests
- ○ Genetic Variants

Prev Next

NCBI Health Resources Survey

ClinVar

3. Do you find it useful to have variant interpretations from OMIM included in ClinVar?

- ◯ Yes
- ◯ No
- ◯ I don't know what this is

4. When searching in Clinvar, do you use the links to other resources (MedGen, GTR, Variation Viewer, PubMed, etc.)

- ◯ Yes
- ◯ No
- ◯ I didn't know these existed

5. How do you define co-occurring variants?

- ◯ Variants in cis or trans occurring in the same gene, where one variant is interpreted to impact the clinical significance (pathogenicity) of the other variant.
- ◯ Variants in cis or trans occurring in the same gene whether or not they impact each other's clinical significance.
- ◯ Variants occurring in the same patient.

Other (please specify)

6. Do you obtain data via the command-line or website?

- [ ] Website
- [ ] Command-line
- [ ] Both

7. What format do you use/would you like to use for batch SNPs from ClinVar?

- [ ] .xml
- [ ] .txt
- [ ] .vcf

Other (please specify)

8. How would you like to see these batch SNPs from ClinVar organized? Would you like to see them presented in a transcript-specific manner?

9. Have you ever used GRCh38 (hg38)?

- ( ) Yes
- ( ) No
- ( ) I don't know what this is

10. How do you usually search in ClinVar?

- ◯ Gene Symbol
- ◯ Variant
- ◯ Disease
- ◯ Accession Number

Other (please specify)

11. Which other NCBI resources do you use?

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

*** 12. Do you use MedGen and/or GTR?**

- ◯ Yes
- ◯ No

Prev Next

NCBI Health Resources Survey

MedGen & GTR

13. Were you aware that GTR provides information on Genetic Tests?

- ◯ Yes
- ◯ No

14. Were you aware that MedGen provides information on medical conditions?

- ◯ Yes
- ◯ No

Prev Next

NCBI Health Resources Survey

MedGen & GTR

13. Would you find it helpful to search by medical specialty (oncology, cardiology, neurology, etc.)?

- Yes
- No

14. Do you search by feature/symptom in MedGen?

- Yes
- No
- I didn't know you could do that

15. Do you use the advanced search feature in GTR?

- Yes
- No
- I didn't know this existed

16. When using MedGen and/or GTR, do you utilize links to other resources (ClinVar, MedGen, GTR, Pubmed, etc.)?

- Yes
- No
- I didn't know these existed

Prev Next

NCBI Health Resources Survey

MedGen & GTR

3. Would you find it helpful to search by medical specialty (oncology, cardiology, neurology, etc.)?

- ◯ Yes
- ◯ No

4. Do you search by feature/symptom in MedGen?

- ◯ Yes
- ◯ No
- ◯ I didn't know you could do that

5. Do you use the advanced search feature in GTR?

- ◯ Yes
- ◯ No
- ◯ I didn't know this existed

6. When using MedGen and/or GTR, do you utilize links to other resources (ClinVar, MedGen, GTR, Pubmed, etc.)?

- ◯ Yes
- ◯ No
- ◯ I didn't know these existed

*** 7. Do you use ClinVar?**

- ◯ Yes
- ◯ No

Prev Next

NCBI Health Resources Survey

ClinVar

8. Were you aware that ClinVar provides information on genetic variants?

- ◯ Yes
- ◯ No

Prev Next

NCBI Health Resources Survey

ClinVar

8. Do you find it useful to have variant interpretations from OMIM included in ClinVar?

- ◯ Yes
- ◯ No
- ◯ I don't know what this is

9. When searching in Clinvar, do you utilize the links to other resources (MedGen, GTR, Variation Viewer, PubMed, etc.)

- ◯ Yes
- ◯ No
- ◯ I didn't know these existed

10. How do you define co-occurring variants?

- ◯ Variants in cis or trans occurring in the same gene, where one variant is interpreted to impact the clinical significance (pathogenicity) of the other variant.
- ◯ Variants in cis or trans occurring in the same gene whether or not they impact each other's clinical significance.
- ◯ Variants occurring in the same patient.

Other (please specify)

11. Do you obtain data via the command-line or website?

- [ ] Website
- [ ] Command-line
- [ ] Both

12. What format do you use/would you like to use for batch SNPs from ClinVar?

- [ ] .xml
- [ ] .txt
- [ ] .vcf

Other (please specify)

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

14. Have you ever used GRCh38 (hg38)?

- ( ) Yes
- ( ) No

15. How do you usually search in ClinVar?

- ( ) Gene Symbol
- ( ) Variant
- ( ) Disease
- ( ) Accession Number

Other (please specify)

16. Which other NCBI resources do you use?

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

Prev Next

NCBI Health Resources Survey

Thank you!

17. What would you like this resource(s) to do that it currently doesn't? Or what would you like it to do better?

18. How likely is it that you would recommend ClinVar to a friend or colleague?

Not at all likely — Extremely likely

0	1	2	3	4	5	6	7	8	9	10

19. How likely is it that you would recommend MedGen to a friend or colleague?

Not at all likely — Extremely likely

0	1	2	3	4	5	6	7	8	9	10

20. How likely is it that you would recommend GTR to a friend or colleague?

Not at all likely — Extremely likely

0	1	2	3	4	5	6	7	8	9	10

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

Prev Done