ClinVar Public Site Survey 2023 (NDB-428)

Start of Block: Default Question Block

OMB Control Number: **0925-0648**
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* Indicates required question

Please select one professional category that describes you best.

- ○ Life Sciences Researcher
- ○ Geneticist
- ○ Genetic Counselor
- ○ Laboratory Staff
- ○ Physician
- ○ Other Healthcare Professional
- ○ Bioinformatics Professional
- ○ Computer Scientist / Software Developer
- ○ Educator
- ○ Student
- ○ Librarian / Information Specialist
- ○ Patient and Family
- ○ Other (please specify) _____

Please select one type that describes your organization best.

- ○ College or University
- ○ Commercial or Industry
- ○ Hospital / Clinical / Medical Practice
- ○ Non-Profit Organization
- ○ Government
- ○ Other (please specify) _____

How likely are you to recommend this resource to a friend or colleague?

- ○ 0
- ○ 1
- ○ 2
- ○ 3
- ○ 4
- ○ 5
- ○ 6
- ○ 7
- ○ 8
- ○ 9
- ○ 10

Were you able to complete your work to your satisfaction today?

- ○ Yes
- ○ No
- ○ Unsure

Please order these tasks below from easiest to hardest.
_____ Finding my variant of interest
_____ Understand the impact of a variant on disease
_____ Determining the validity / credibility of a submission
_____ Evaluating discrepancies between ClinVar and other databases
_____ Providing feedback about data errors to ClinVar or ClinVar submitters
_____ Using ClinVar information in lab / patient reports
_____ Getting answers to my questions

If ClinVar created a mechanism to exclude submitted records (SCVs) with incorrect classifications from the calculation of the aggregate classification, do you foresee any issues? (Note: submissions would not be changed or removed from the database. The exclusion would only impact the summary data, e.g., the summary section at the top of a variant page.)

- ○ Yes, always
- ○ Yes, in certain cases
- ○ No, never
- ○ Not sure

Please explain your response in the previous question.

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Do you want the ability to provide corrections for data in ClinVar?

- ○ Yes, I would like to provide corrections for all types of data
- ○ Yes, I would like to provide corrections only for some types of data, e.g., citations
- ○ No, never
- ○ Not sure

If ClinVar annotated submitted records (SCVs) as “requires review” based on a community-raised error, under what circumstances would this annotation be appropriate?

- ○ Never
- ○ Sometimes – when the community-raised error is from an expert panel
- ○ Sometimes – when the community-raised error is from organizations / people that are known to ClinVar (e.g., has an NCBI account)
- ○ Always

If ClinVar annotated submitted records (SCVs) as “requires review” based on a community-raised error, which information would you want flagged?

- [ ] None, this feature is not important to me
- [ ] Classification
- [ ] Publications
- [ ] Assertion criteria
- [ ] Evidence
- [ ] Condition for the classification
- [ ] Other (please specify) _____

Please explain your response to the previous question.

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Please rank these potential improvements that you would like to see in ClinVar.

_____ Better search
_____ Better display of variant pages
_____ Better download options for variant pages
_____ Access to older versions of submitted classifications
_____ Better representation of CNVs
_____ Labeling of conflicting classifications with the type of classification, e.g., P/LP vs VUS
_____ Ability to investigate other variants at the same location
_____ More gene-centric views
_____ More graphical displays

Do you have any other suggestions for improving ClinVar?

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End of Block: Default Question Block