

ClinVar Public Site Survey 2023 (NDB-428)

Start of Block: Default Question Block

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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.

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.

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?

End of Block: Default Question Block
