# ACTGATCGTATGGGGCCAAGAGATATATCT <br> CAGGTACGGCTGTCATCACTTACACCTCAC <br> CAGCGCTGOGCATAAAAGTCAGGCCACAGC CCATGCTGCATCTCACTCCTCAGGAGAAGT GCAGGTTGGTATCAAGGTTACAAGACAGGT GGCACTGACTCTCTCTGCCTATTGGTCTAT <br> ClinVar Quarterly User Survey <br> ClinVar aggregates information about genomic variation and its relationship to human health. 

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## 1. How often do you use ClinVar?

Every daySeveral times a weekSeveral times a monthAbout once a monthLess than once a monthThis is my first visit
## 2. What were you looking for on your current visit to ClinVar?

A specific variantAll variants in a geneVariants associated with a disease or phenotypeVariants submitted by a specific groupOther (please specify)$\square$
3. How easy was it to find what you were looking for on ClinVar?Extremely easyVery easySomewhat easyNot so easyNot at all easy
4. How easy is it to understand the information on ClinVar?Extremely easyVery easySomewhat easyNot so easyNot at all easy

## 5. Overall, how satisfied are you with ClinVar?

Extremely satisfiedVery satisfiedModerately satisfiedSlightly satisfiedNot at all satisfied
## 6. Which category describes you best?

Lab directorGenetic counselor
$\square$ Medical geneticistPathologistOther type of physicianResearcher
Student

Other (please specify)
$\square$
7. What would make you more likely to use ClinVar?
$\square$

## 8. Any other comments?

9. 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 |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |

Done

