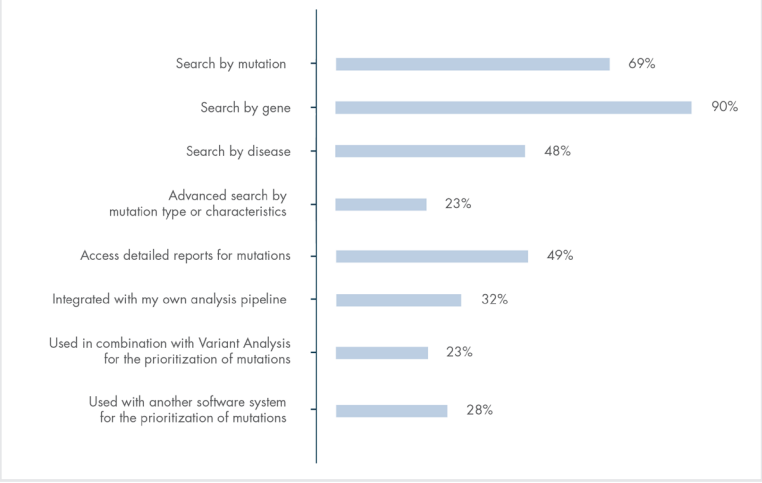
HGMD Survey Results

How is HGMD used by current customers in their work?

We conducted a survey of more than 200 users of the online and download offerings. The results provide a summary of viewpoints from customers spanning diverse backgrounds and institutions, giving insight into the many ways that HGMD can be applied and the challenges that it helps to resolve.

Primary Application Focus



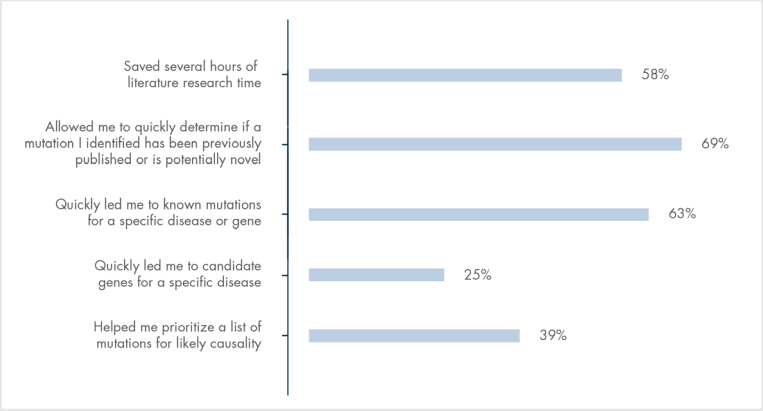
Listed above are examples of how HGMD can be applied. Please check all that you have used HGMD for in the past 12 months.

QIAGEN HGMD Online Customer Statistics

93% of surveyed research organizations rely on links to the mutation source (for example, the PubMed abstract) from HGMD, which contributes to their trust in the resource.

75% of surveyed research organizations have reduced the amount of time needed for identifying published mutations by 50% or more with HGMD compared with previous methods.

HGMD Contribution

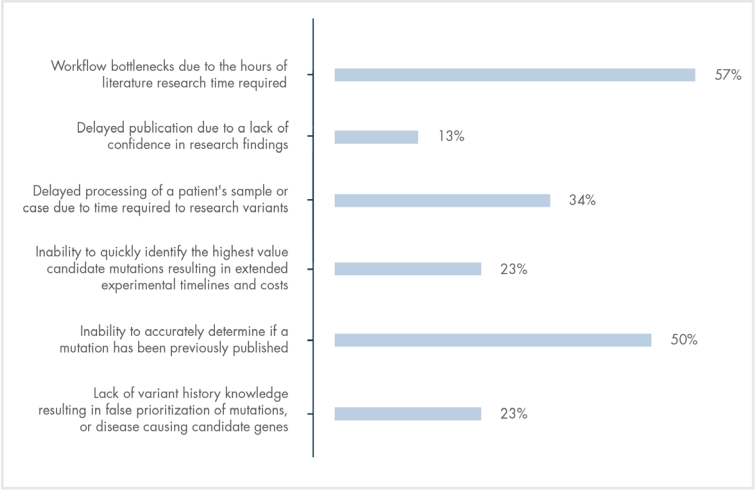


How has the use of HGMD contributed to your work, or the work of others that you support?

> “Fast reliable searching that we can use to integrate either into our own pipeline or being able to use in conjunction with other tools for variant analysis.”
>
> **Research Analyst, Federal Government**

> “It has provided up-to-date information about genes and mutations that help facilitate the interpretation of patient results. It is an excellent tool and saves me a lot of time.”
>
> **Lab Director, Health Care Company**

Challenges Solved



What bottlenecks has HGMD solved for your laboratory or organization?

> “The product has provided a fast, convenient way to prioritize previously described variants in relation to an exome or genome’s worth of variant data.”
>
> **Research Analyst, Medical College**

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