



5/5 Stars

Please share details on your experience with it.

“ I’m satisfied in using HGMD as it enables quick data acquisition and mutation assesment. In particular, identification of mutations incorrectly described in the literature is enhanced, making the HGMD unique and very helpful.

Would you recommend our software to others, and if so, why?

“ Yes, I would, as I have used it for several years and I’m very satisfied to have it as an external, curated database.

Source: Andrzej Tysarowski, Associate Professor, Maria Sklodowska-Curie Institute of Oncology, PhD



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