



Reply

Reply to the Letter of Charre et al. “Mis-Genotyping of Some Hepatitis D Virus Genotype 2 and 5 Sequences Using HDVdb”

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We thank Charre and colleagues for spotting the mis-annotation of sequences in our database, which was caused by human error. The problem has been corrected. In the future, we recommend submitting bug reports directly to one of the e-mail addresses indicated on the HDVdb Web page.

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