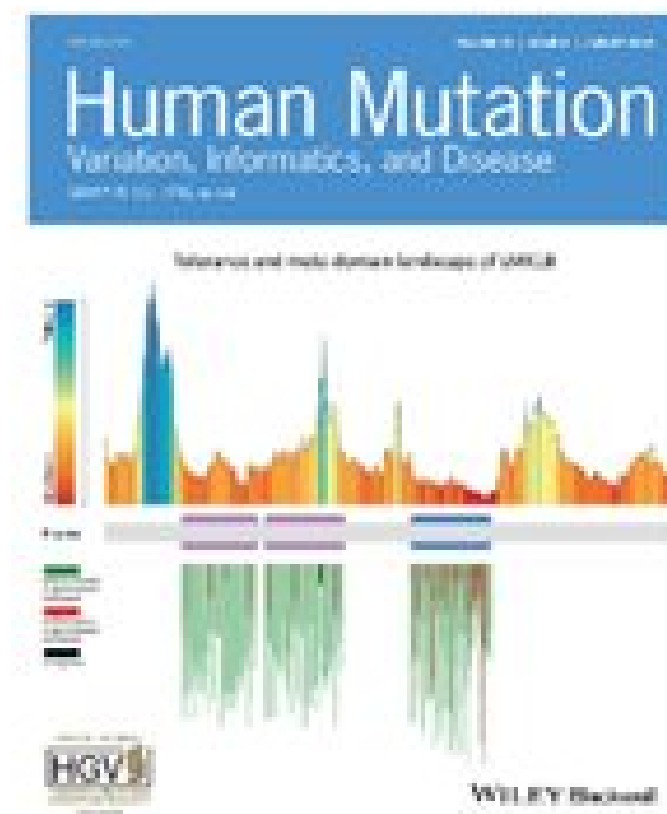


## Front cover

### Human Mutation

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21 August 2019



The MetaDome web server build to interpret genetic variants based on genetic tolerance and homologous protein domains is featured on the Cover of [Human Mutation](#)'s August edition. Additionally it is selected as an Editor's choice article.

MetaDome was developed by Laurens van de Wiel, Coos Baakman, Daan Gilissen, Joris Veltman, [Gert Vriend](#) and [Christian Gilissen](#) (Departments: CMBI & Human Genetics).

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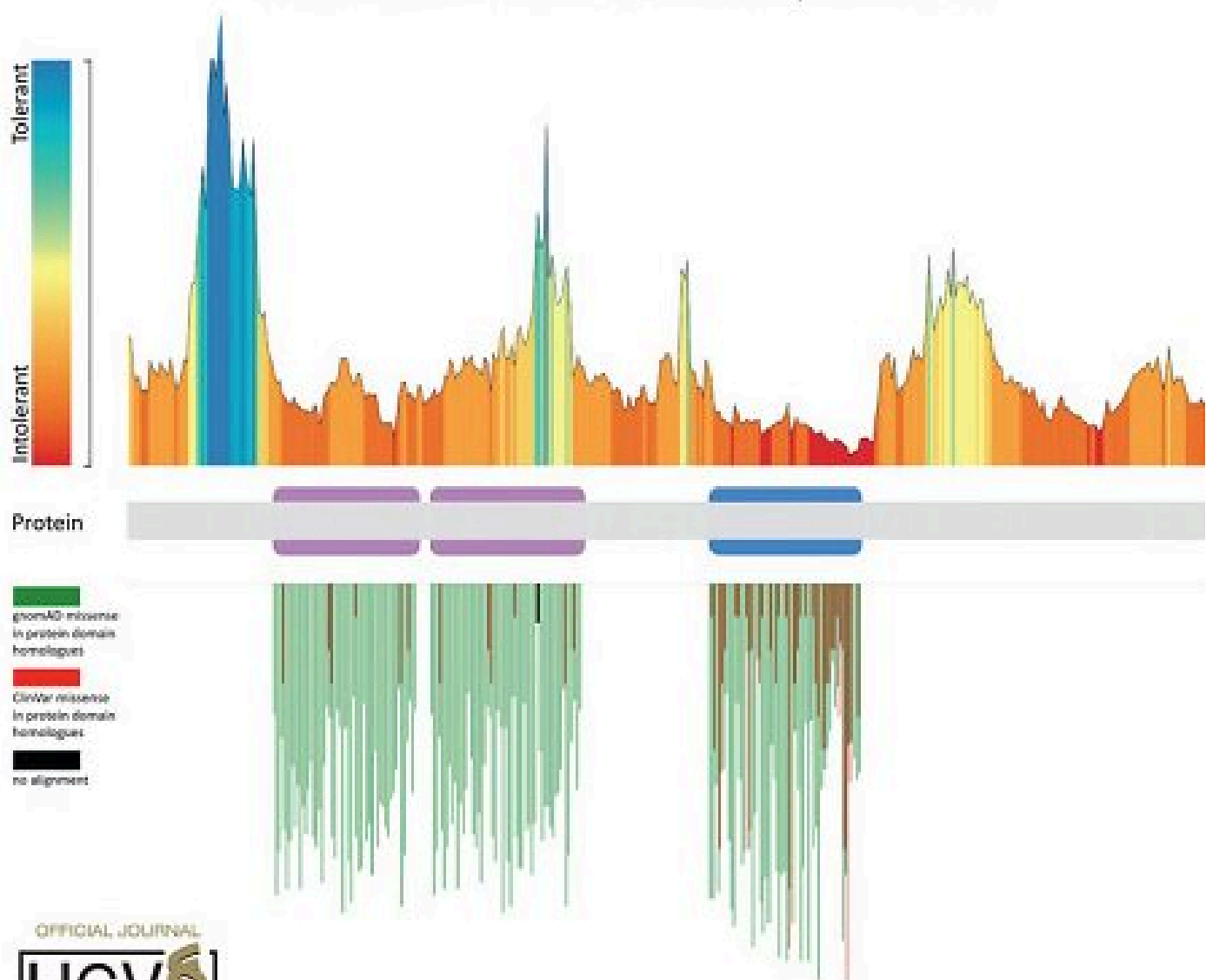
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# Human Mutation

## Variation, Informatics, and Disease

GARRY R. CUTTING, EDITOR

### Tolerance and meta-domain landscape of LMX1B



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