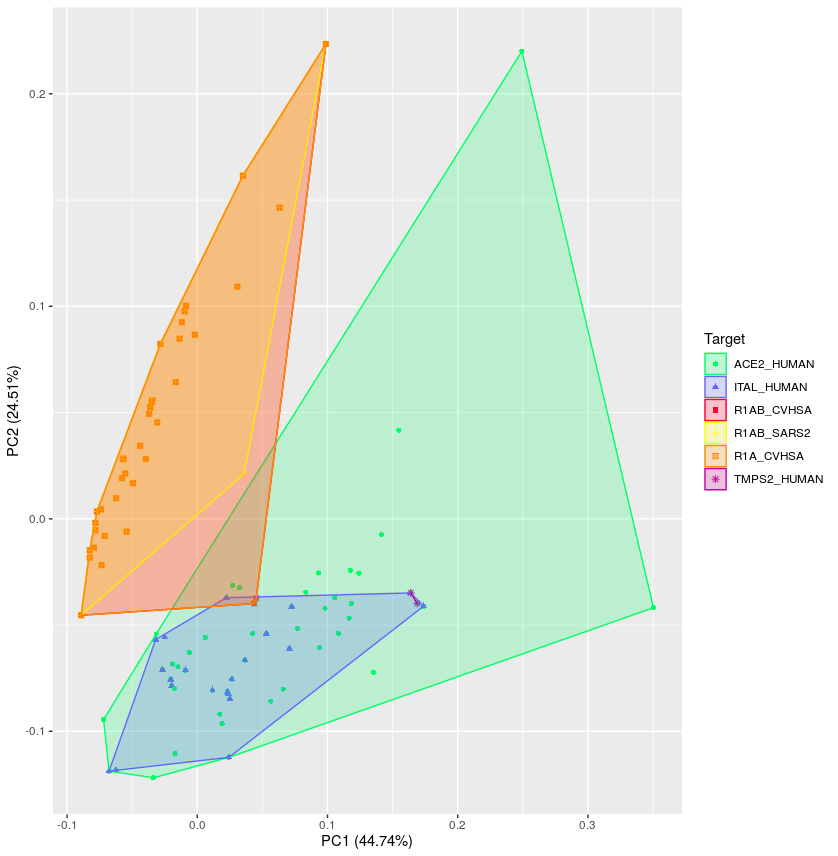
**Supporting Information**

**Drug Repurposing Pipeline in KNIME: A case study on COVID-19 Data**

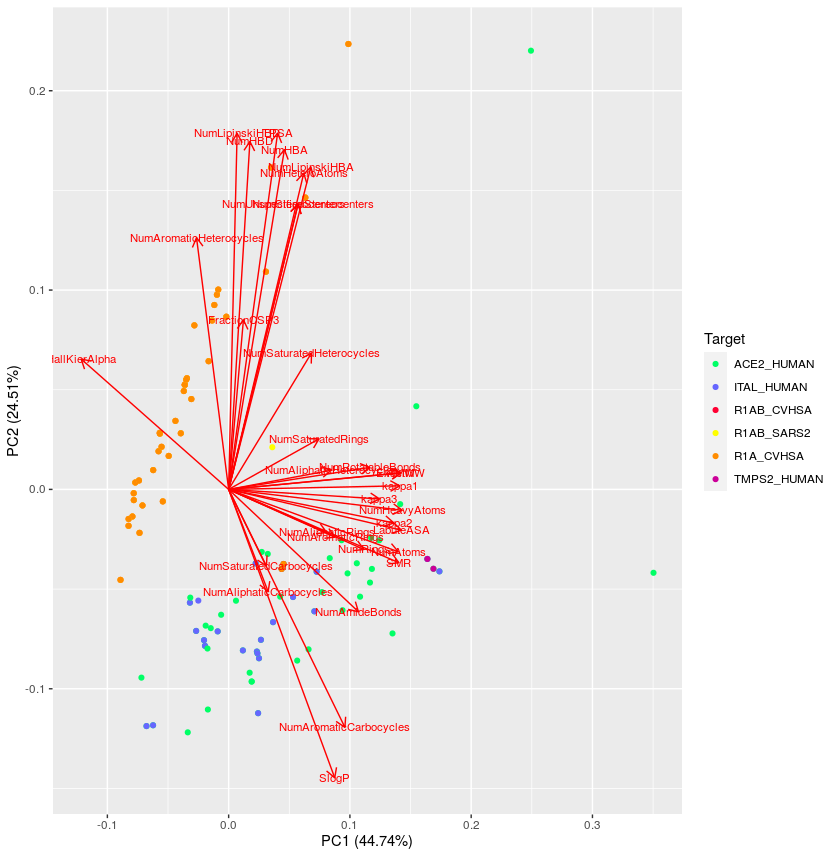
*Alzbeta Tuerkova and Barbara Zdrazil*

University of Vienna, Department of Pharmaceutical Chemistry, Division of Drug Design and Medicinal Chemistry, Althanstraße 14, A-1090 Vienna, Austria.

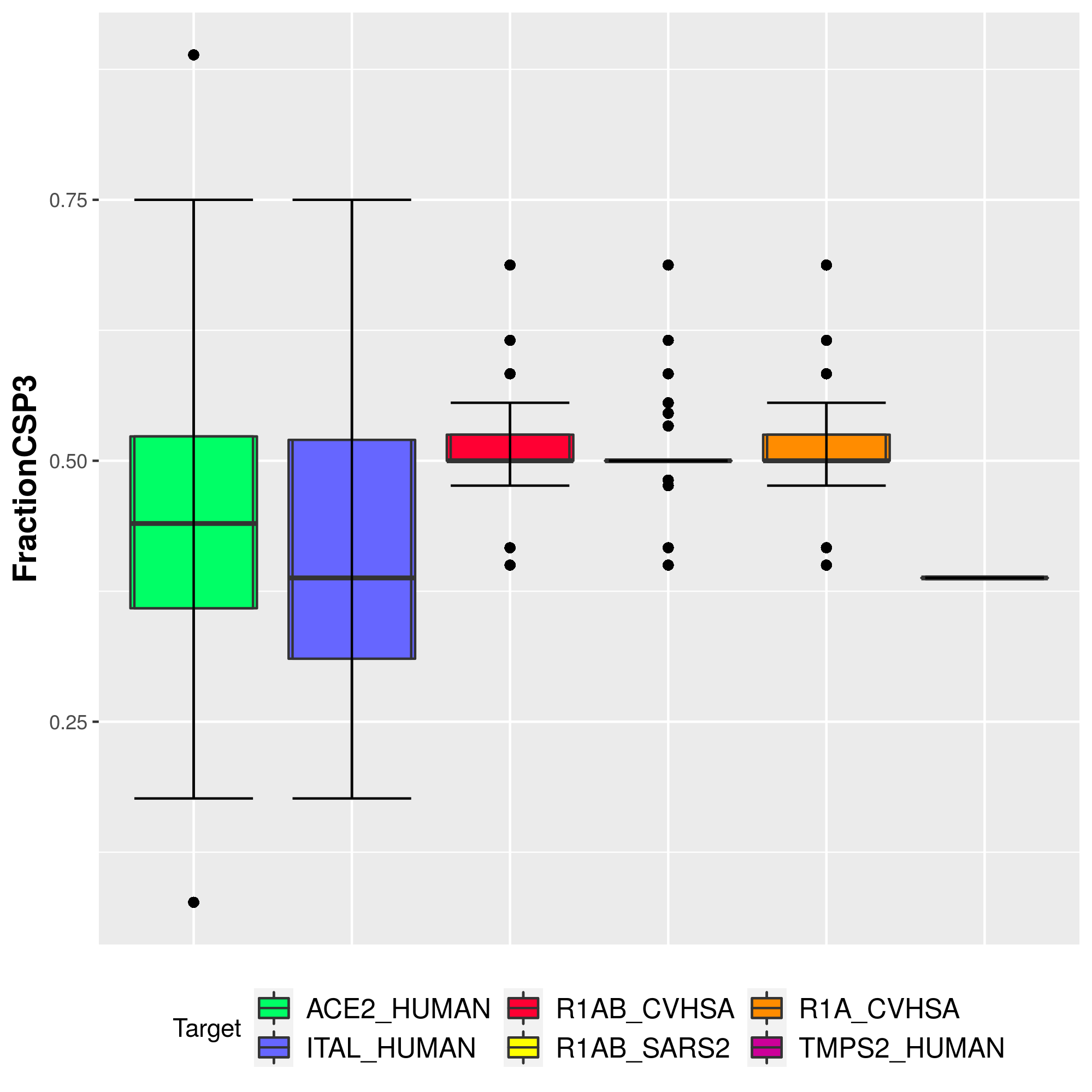
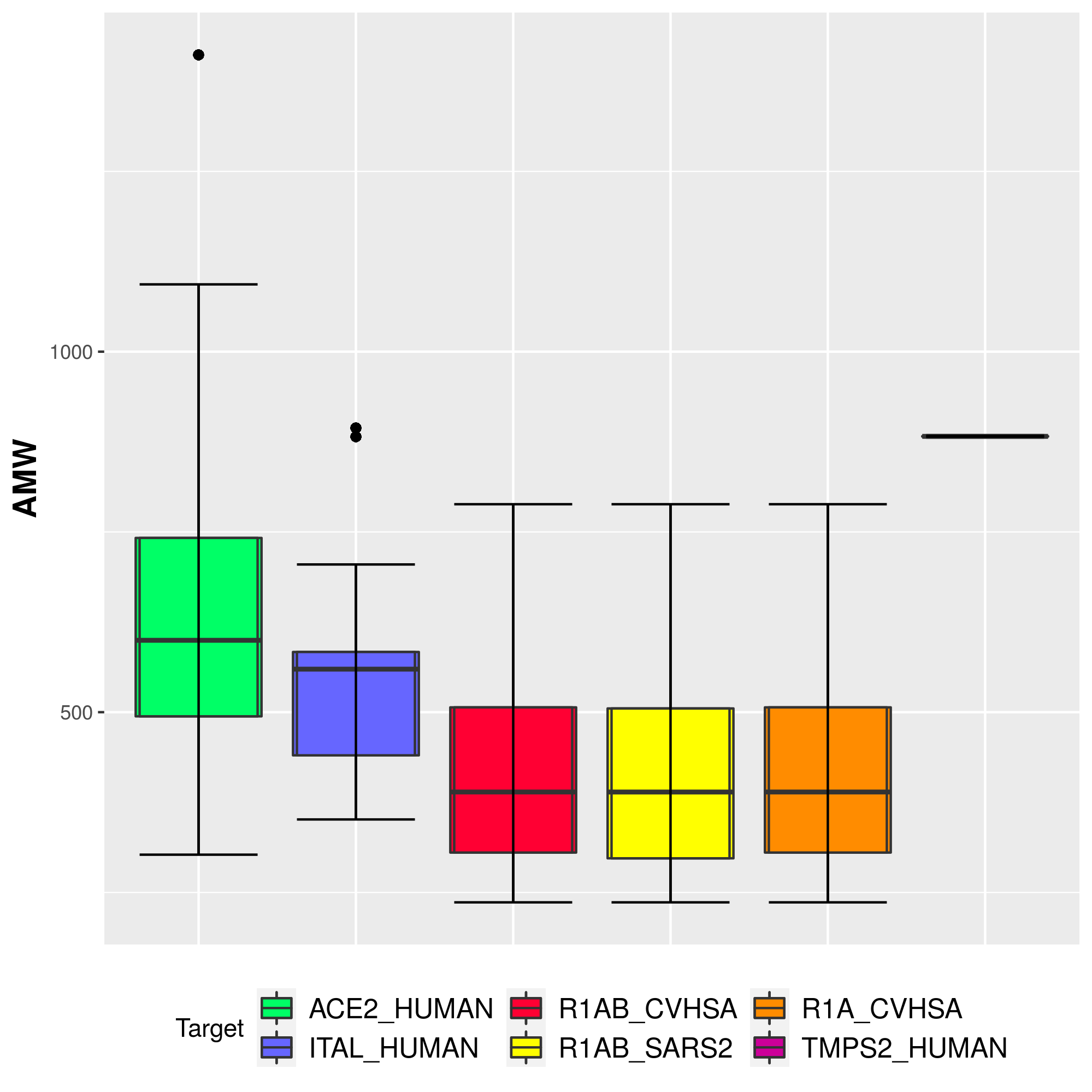
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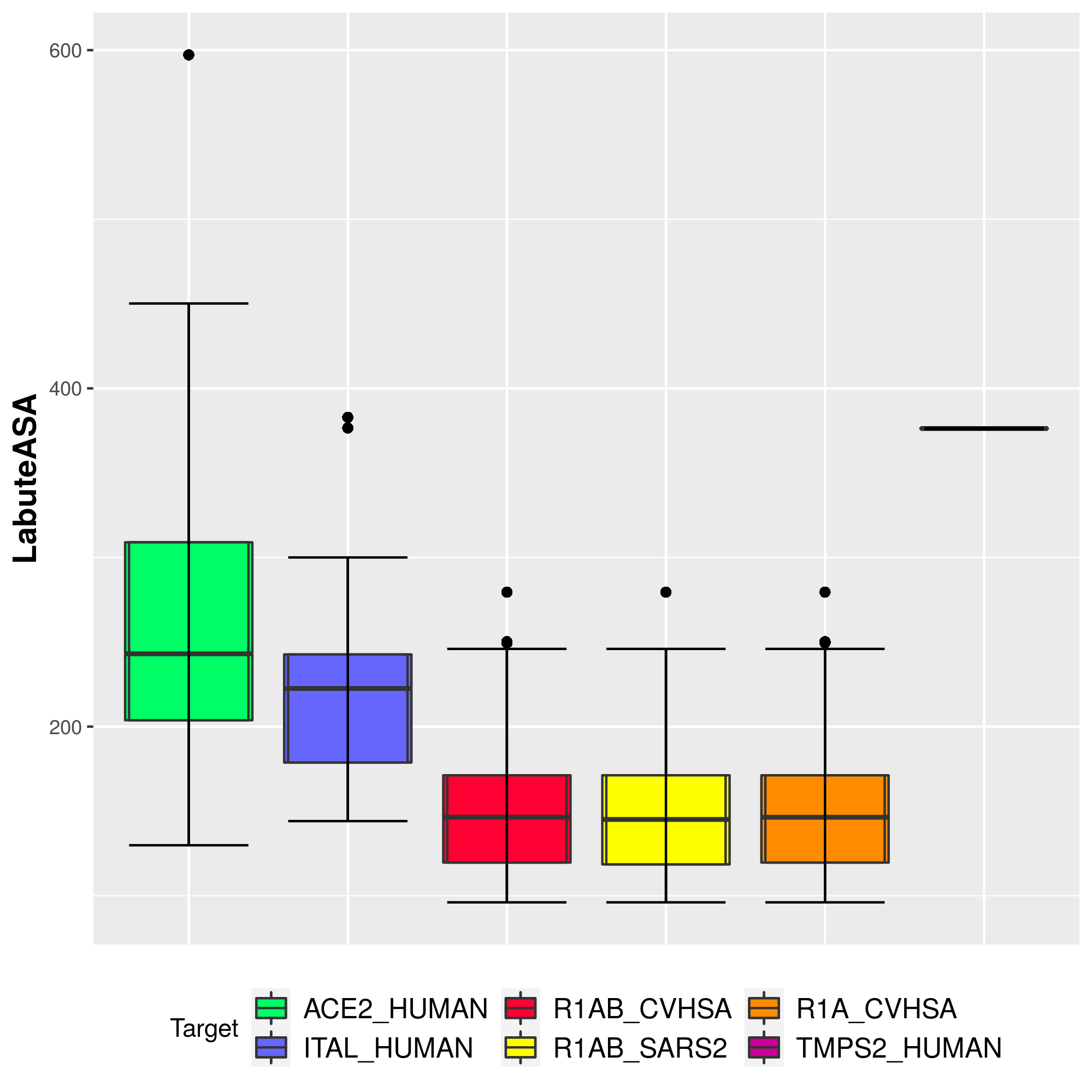
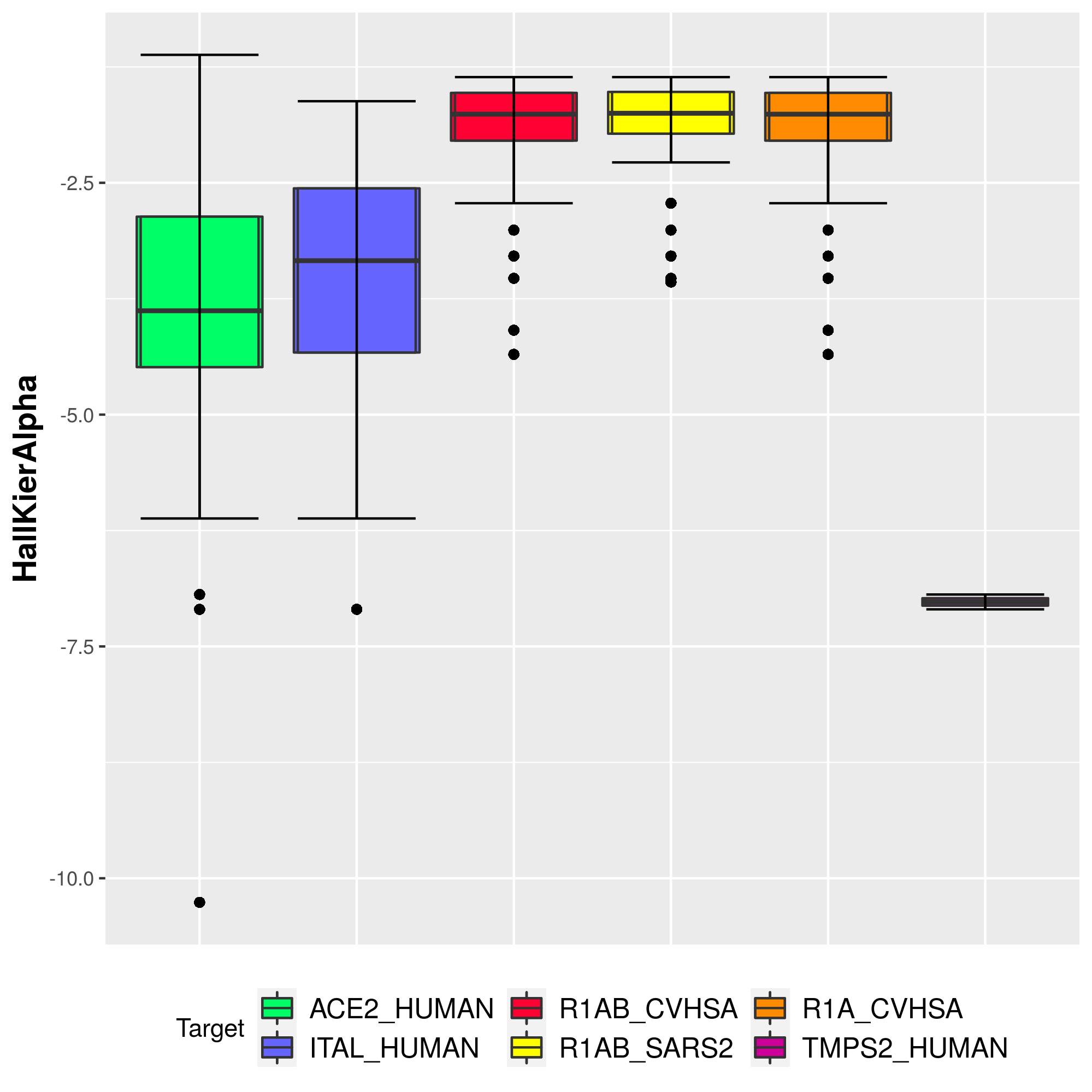
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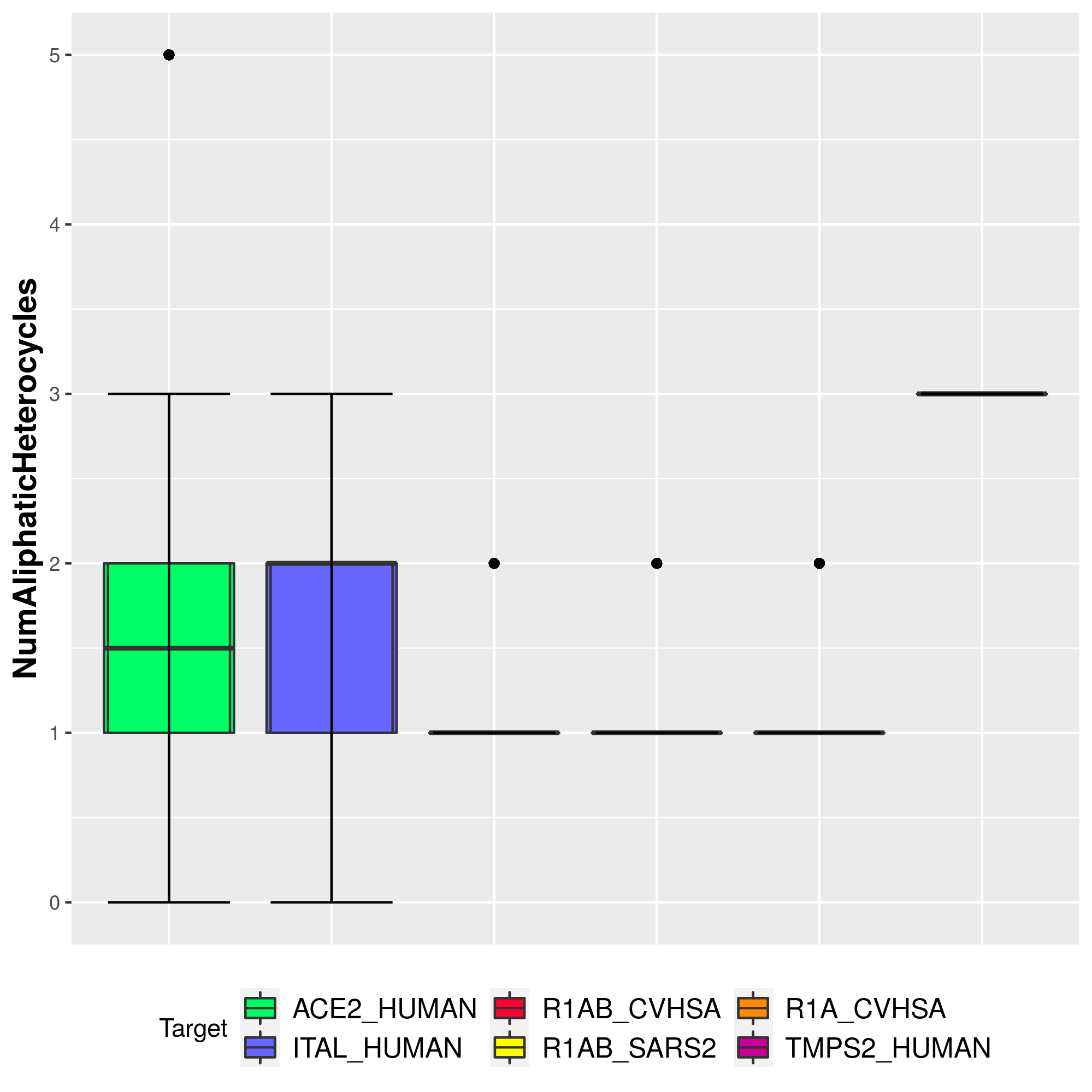
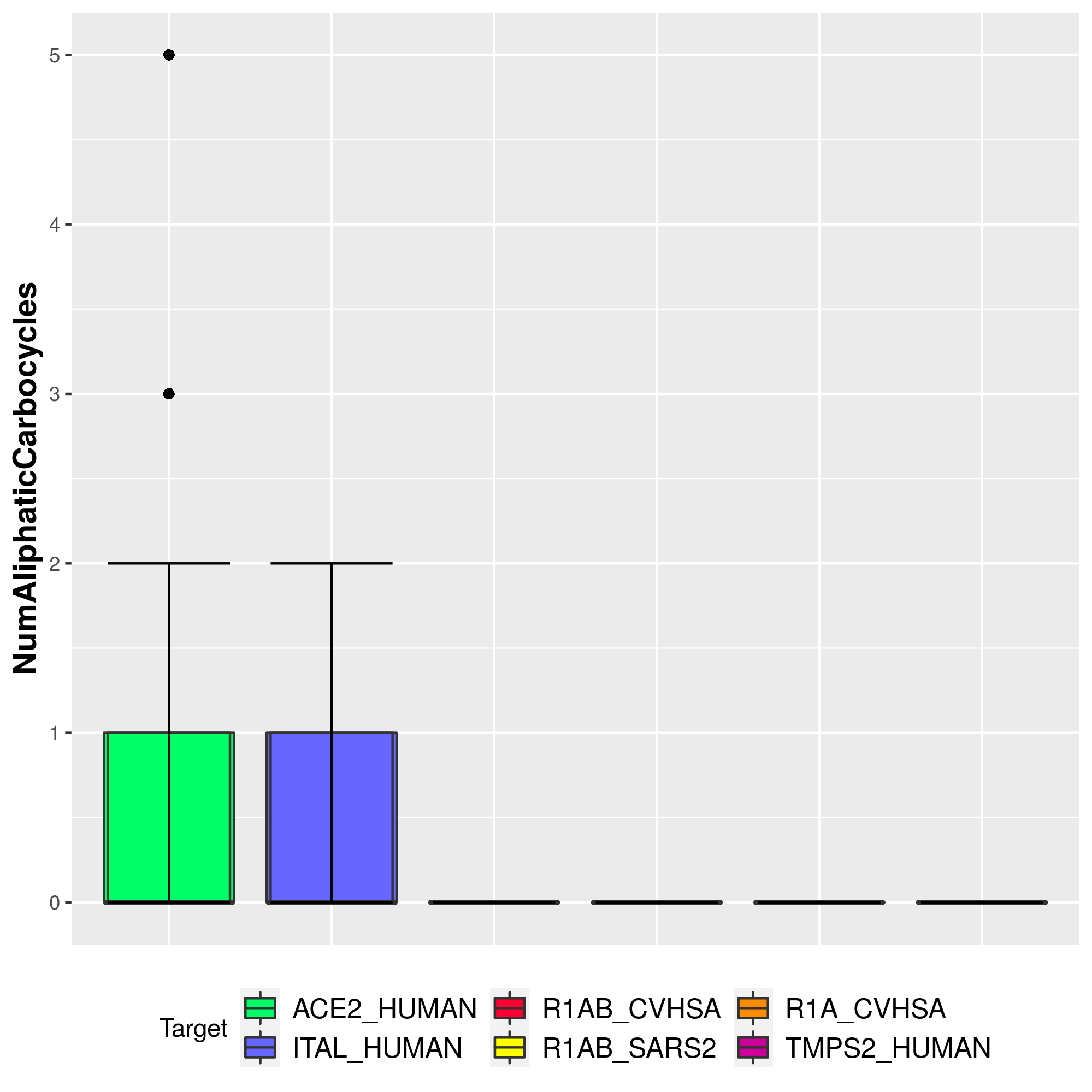
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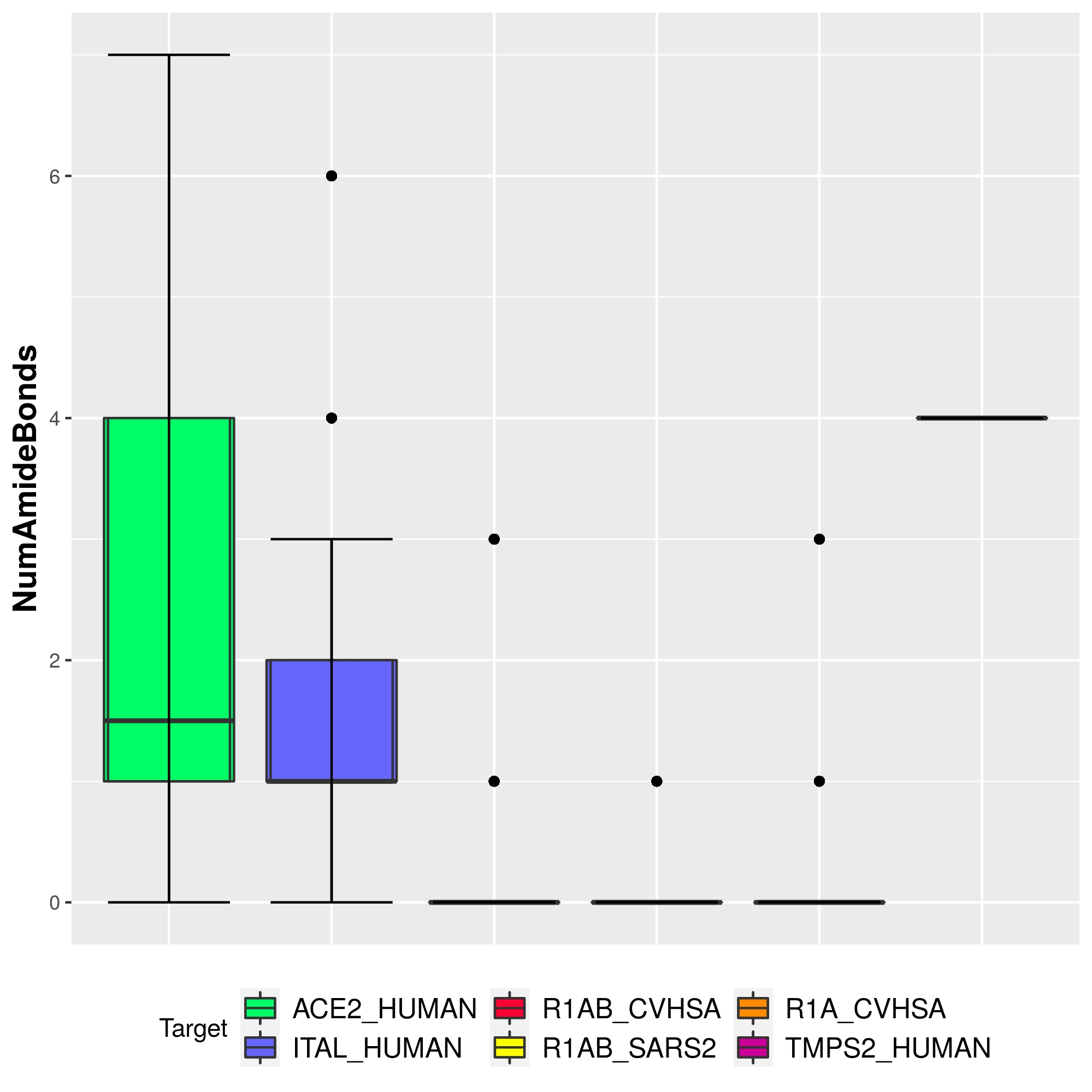
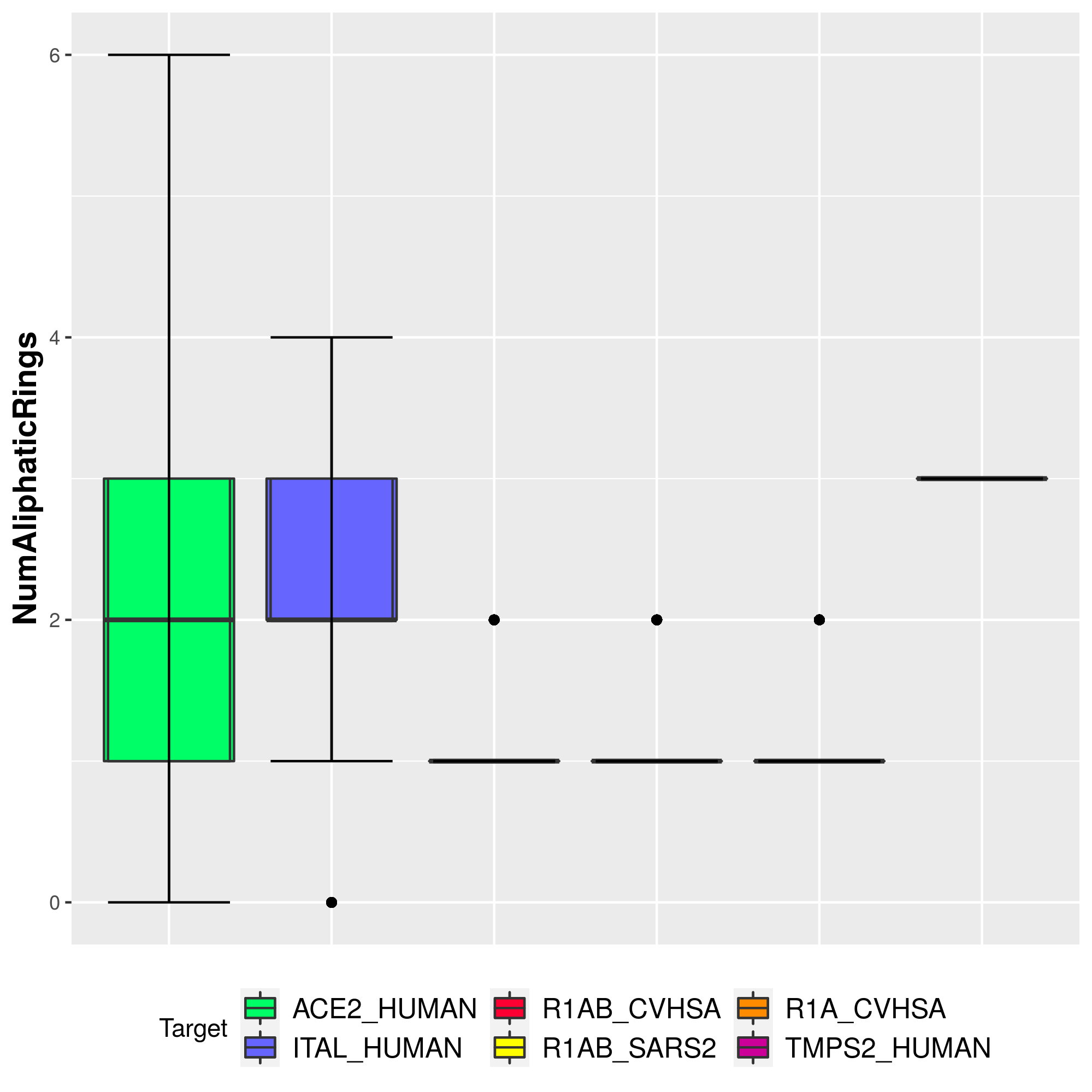
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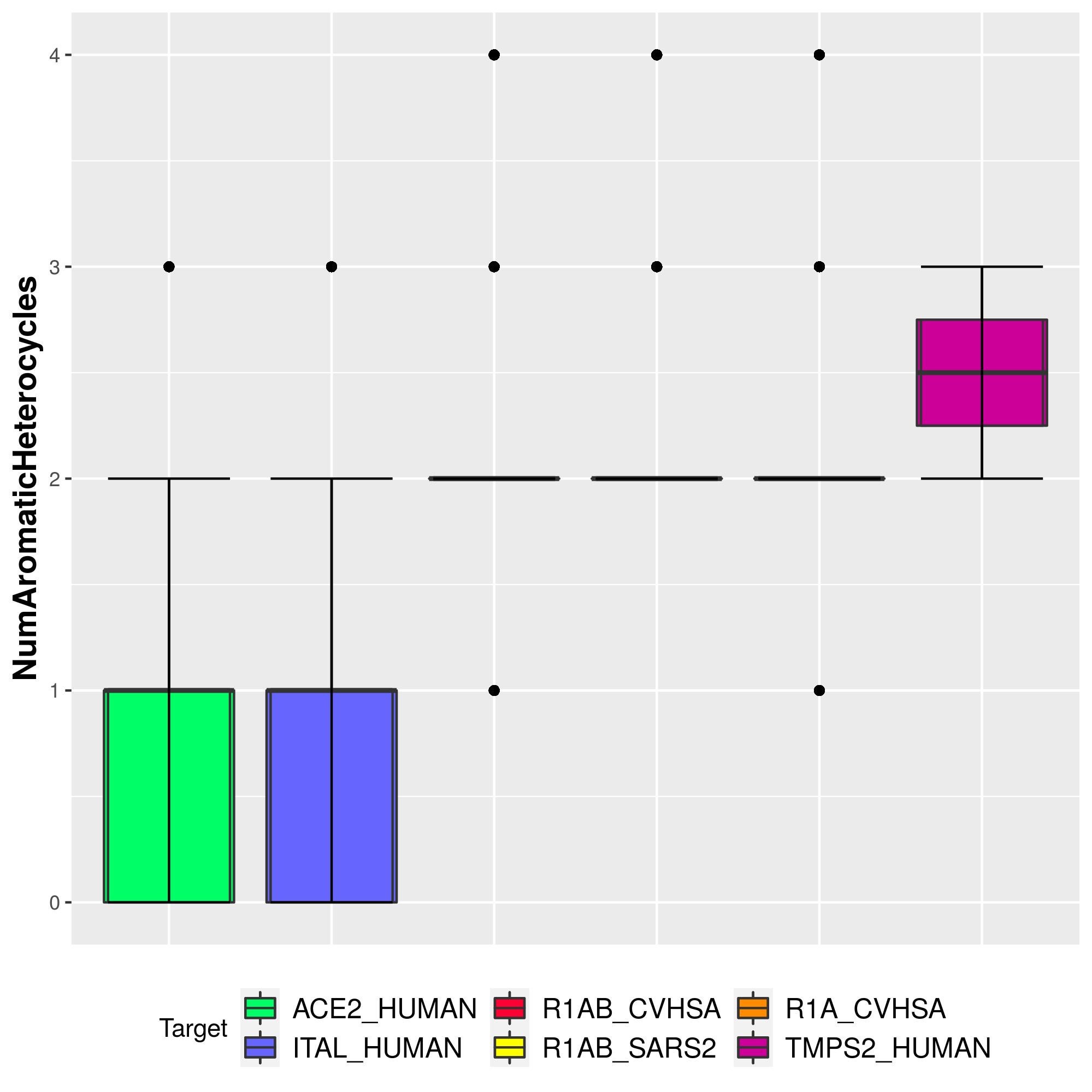
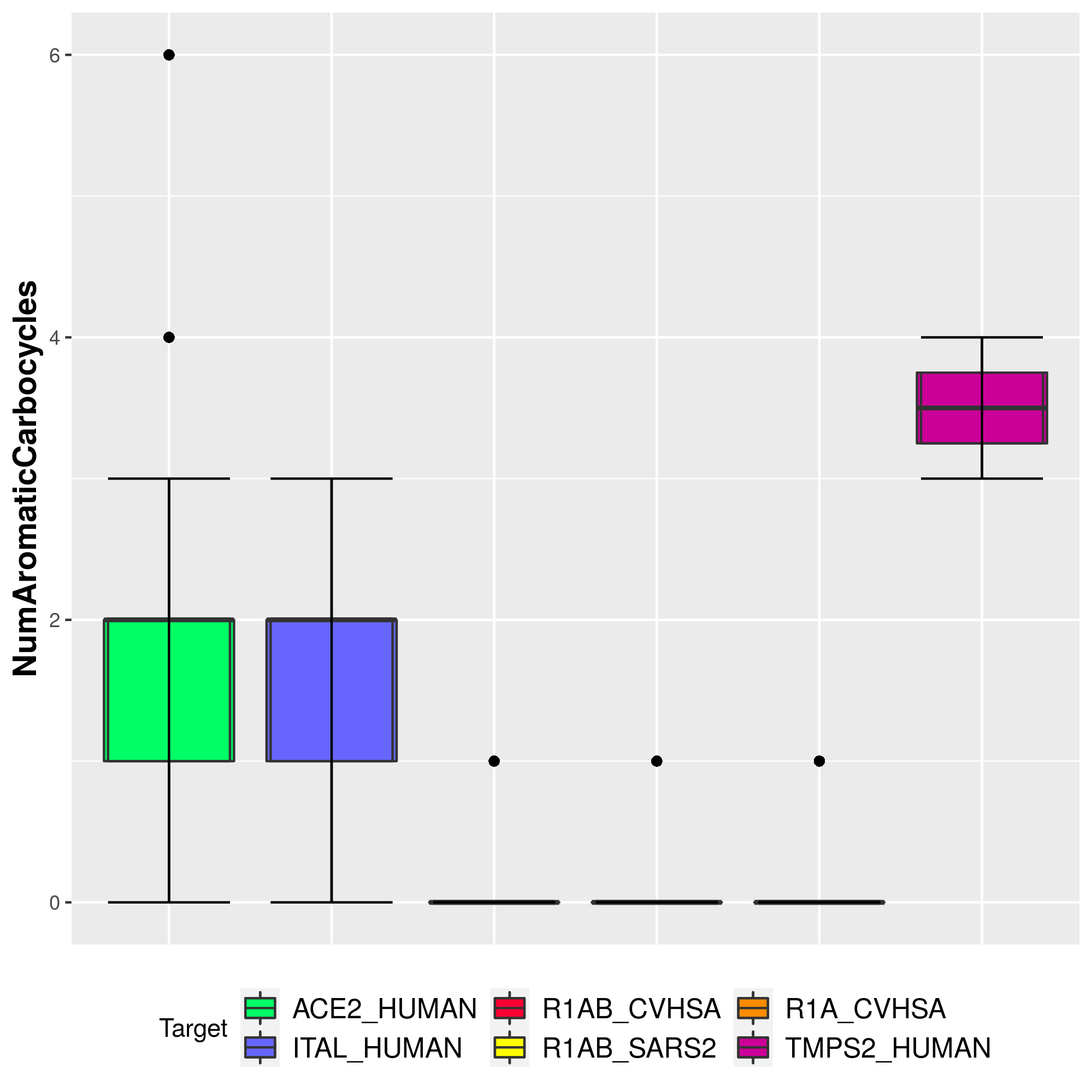
***Supplementary Figure S1.*** *Principal Component Analysis (PCA) along principal component 1 (PC1) and principal component 2 (PC2). (A) PCA depicting clusters of compounds which are linked to a certain target. (B) Eigenvectors of molecular features along PC1 and PC2.*

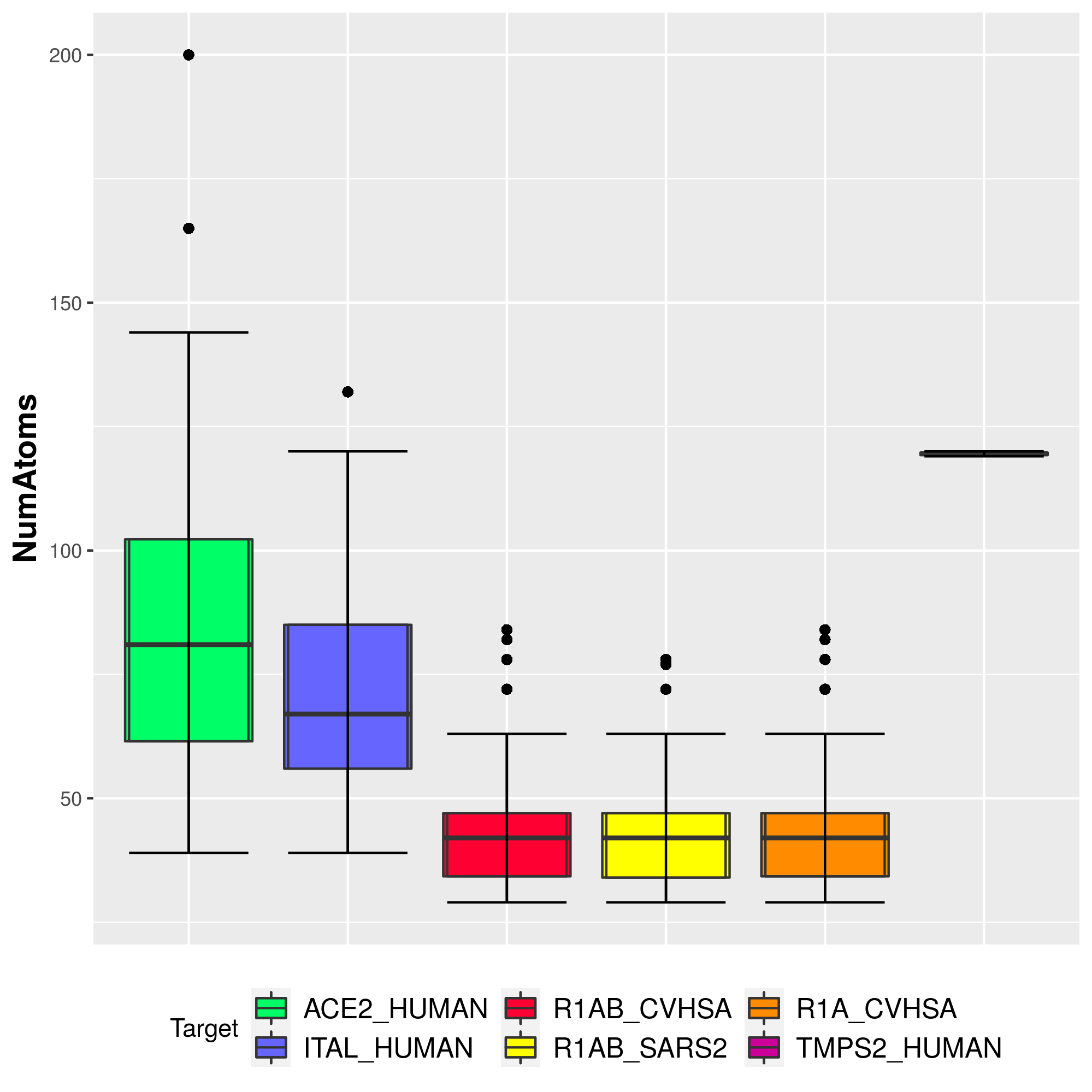
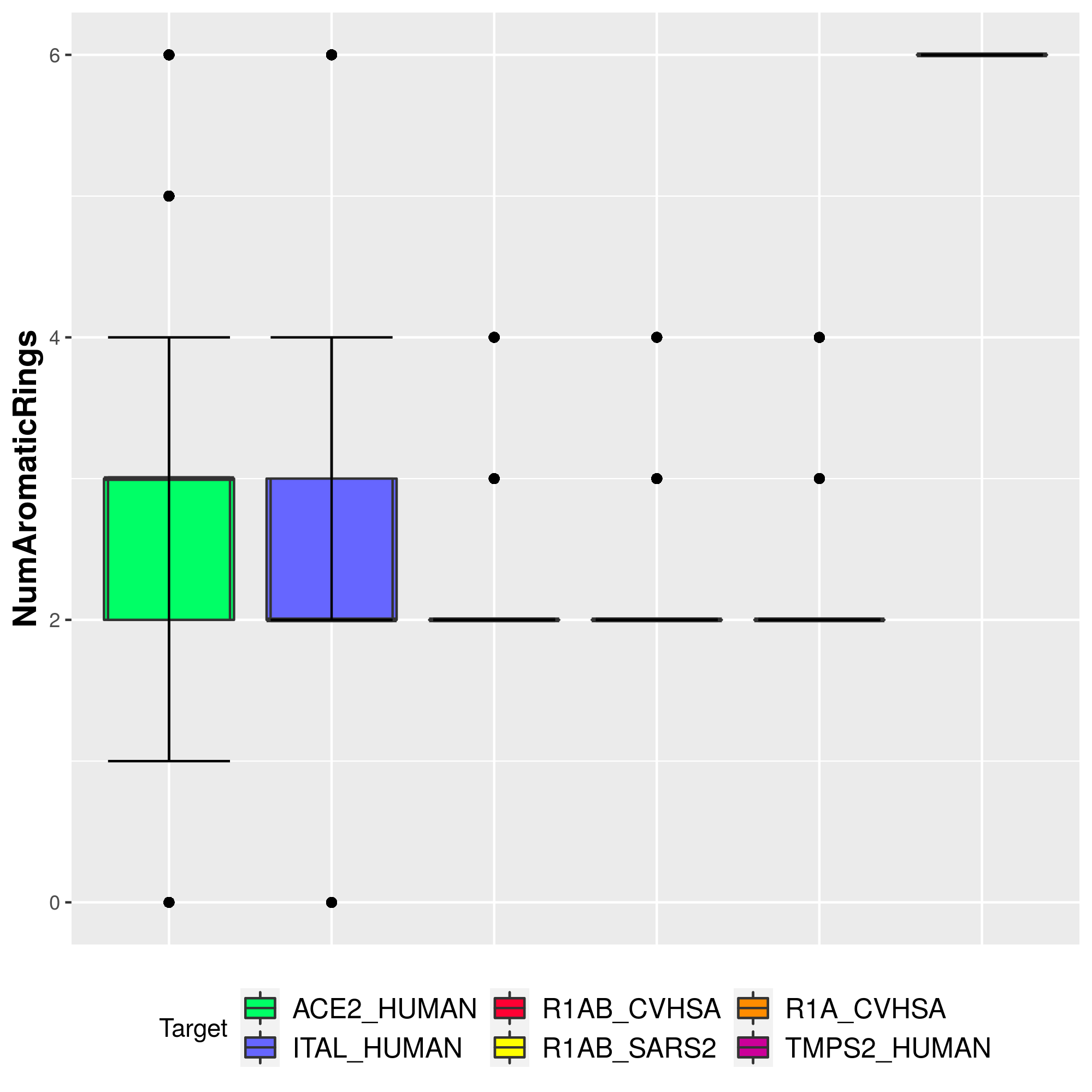
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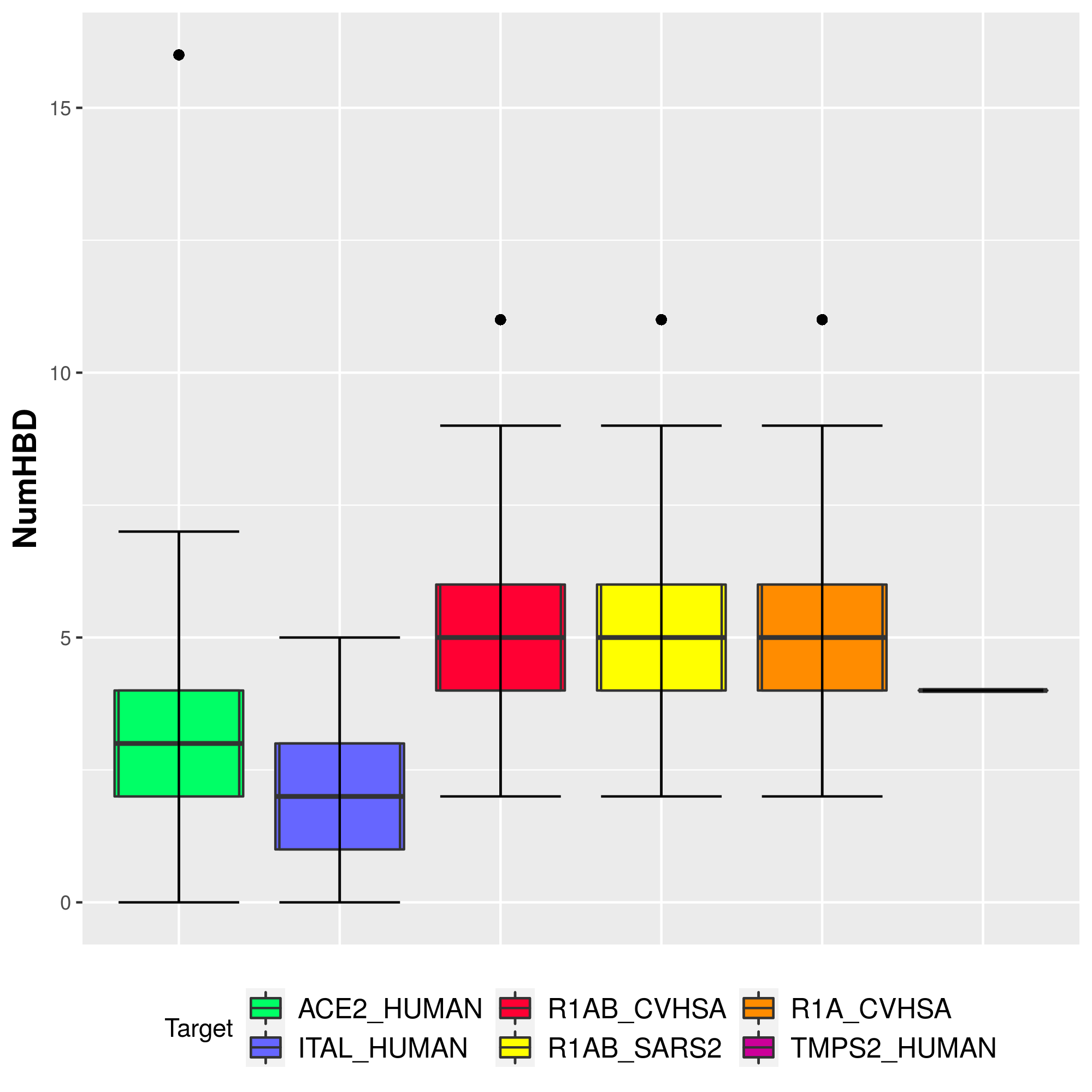
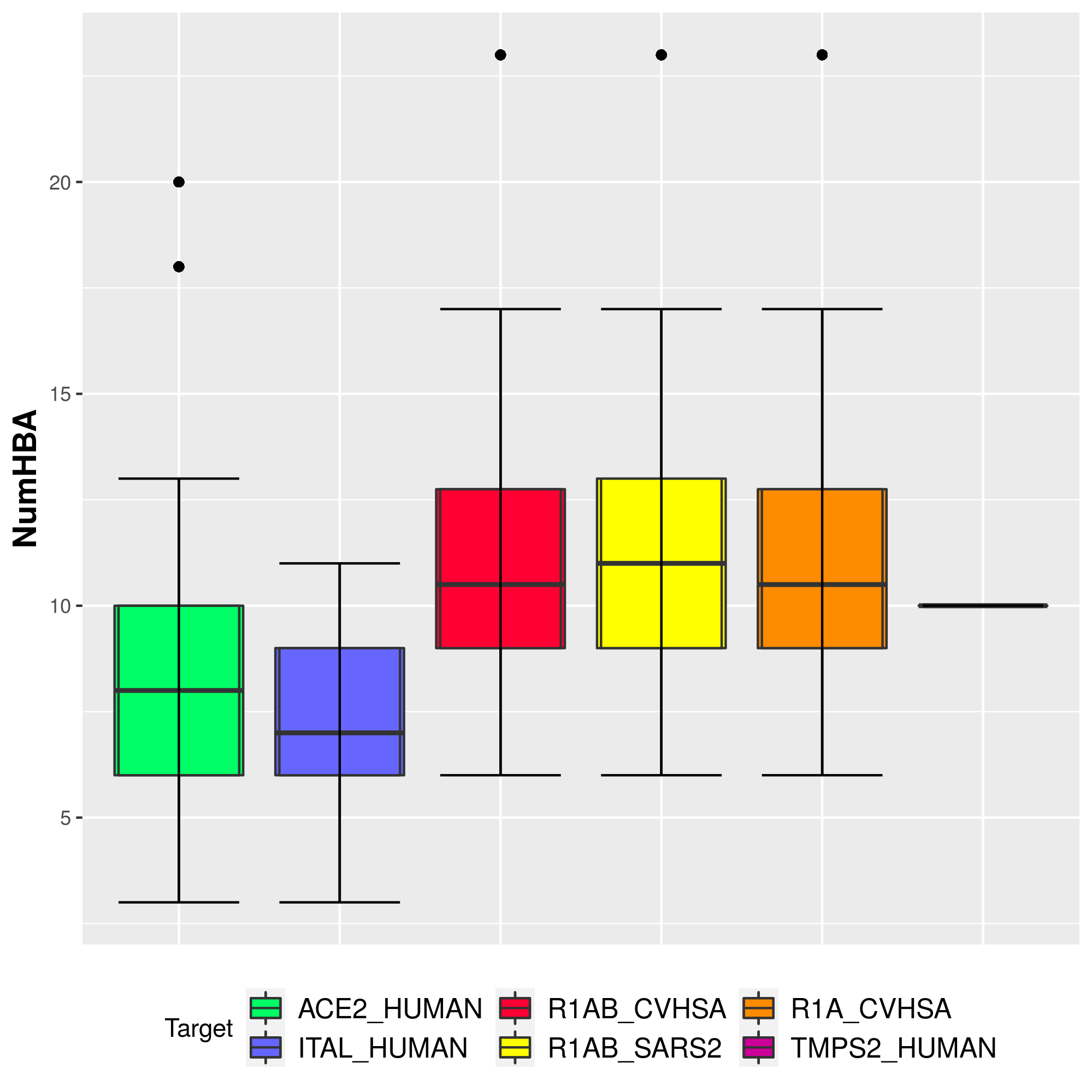
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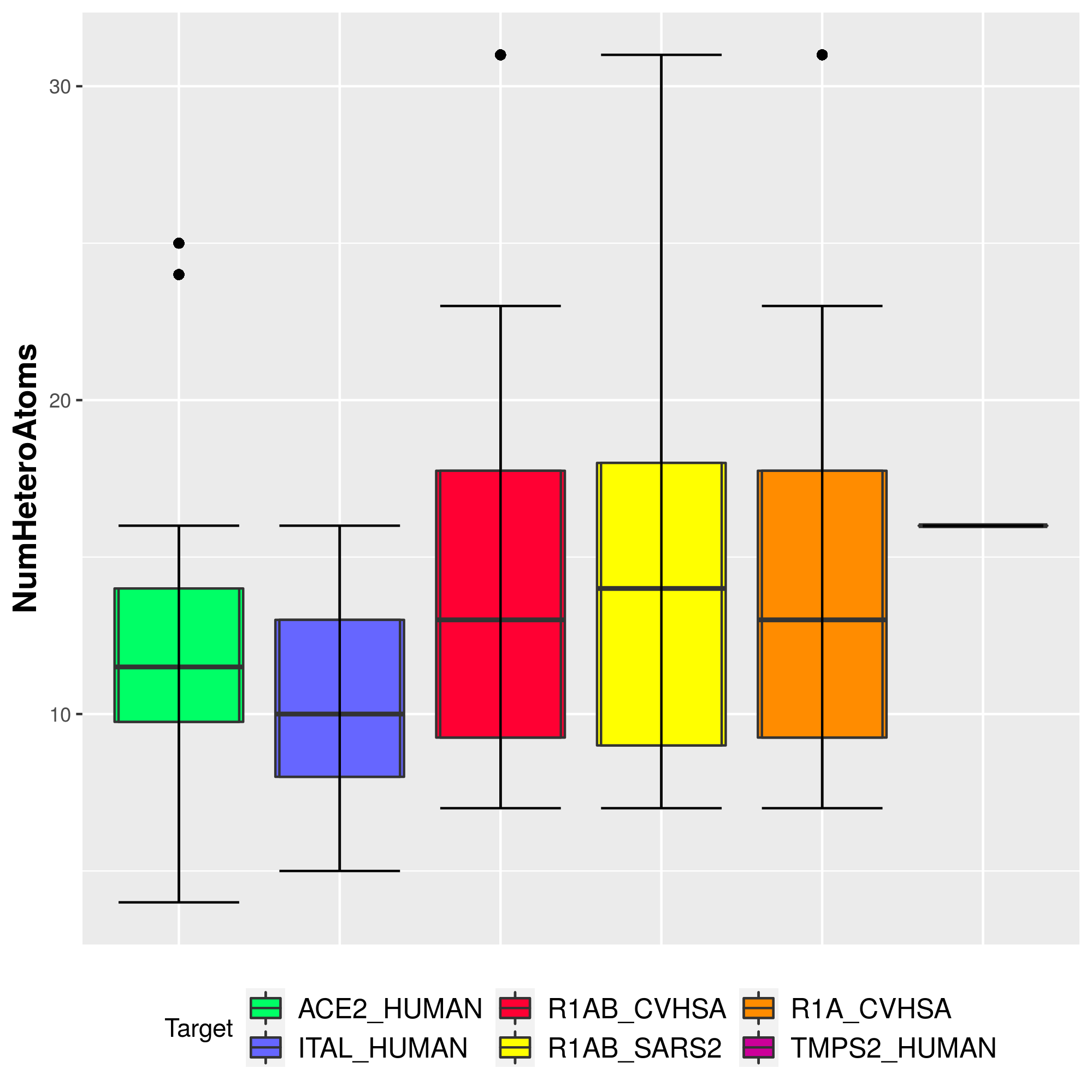
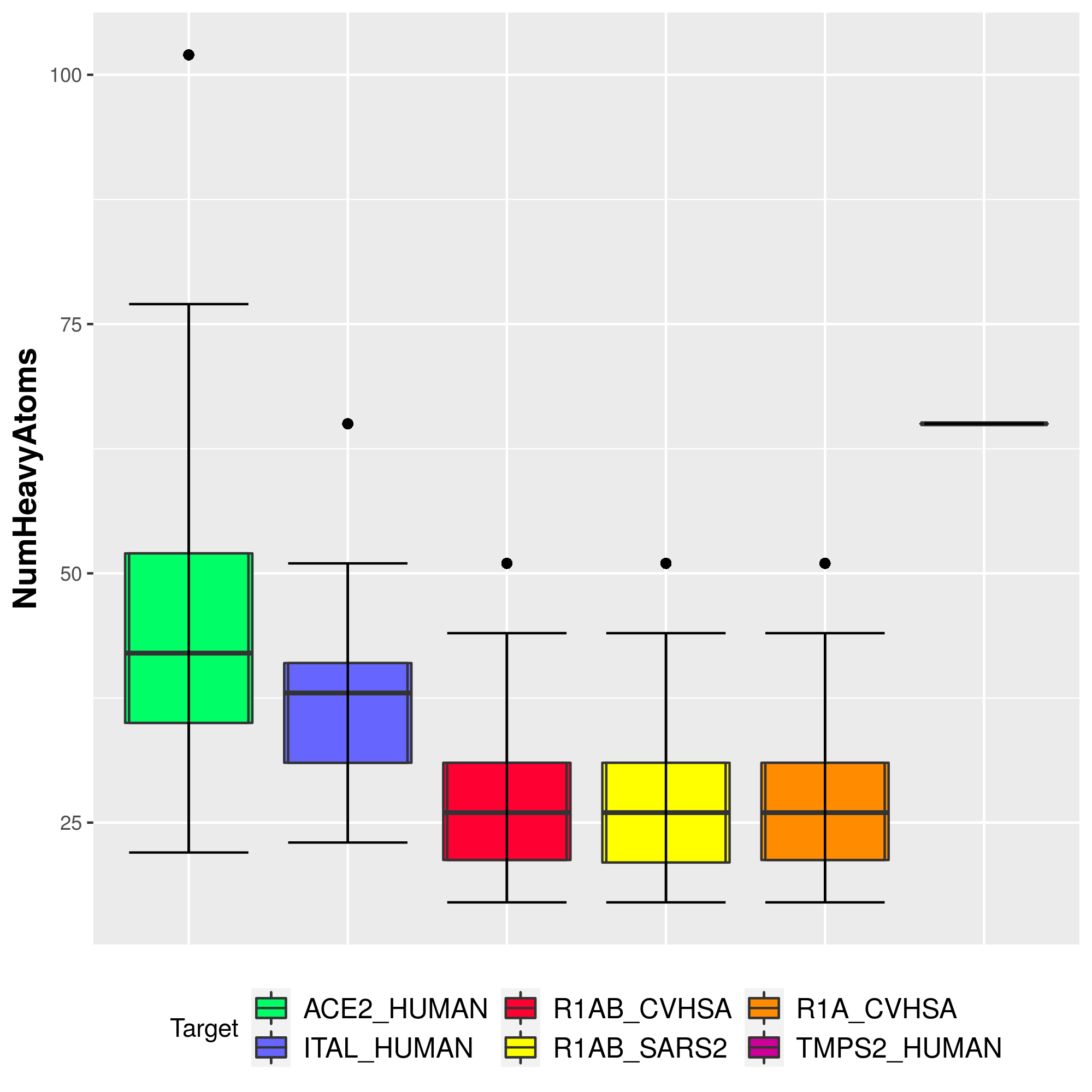
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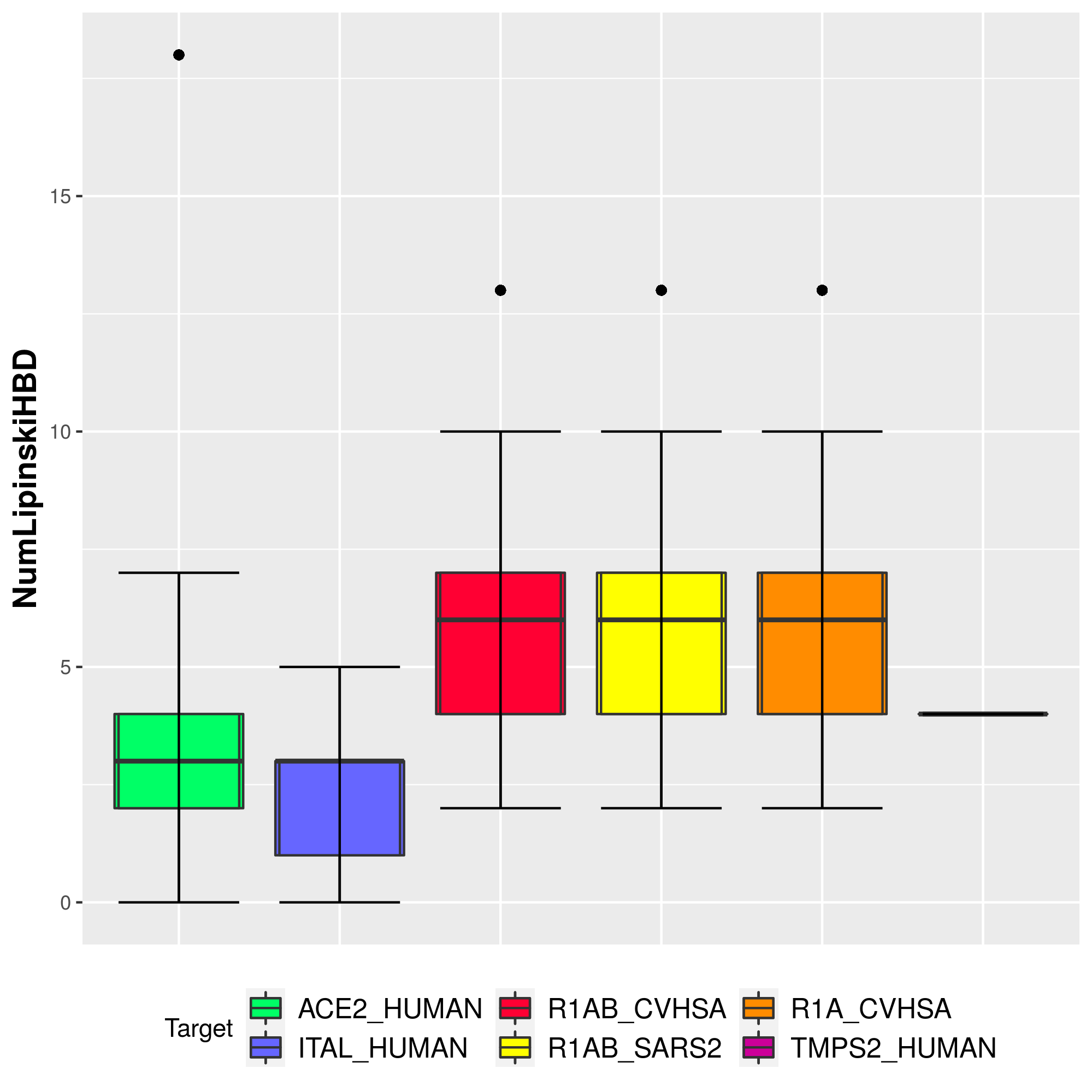
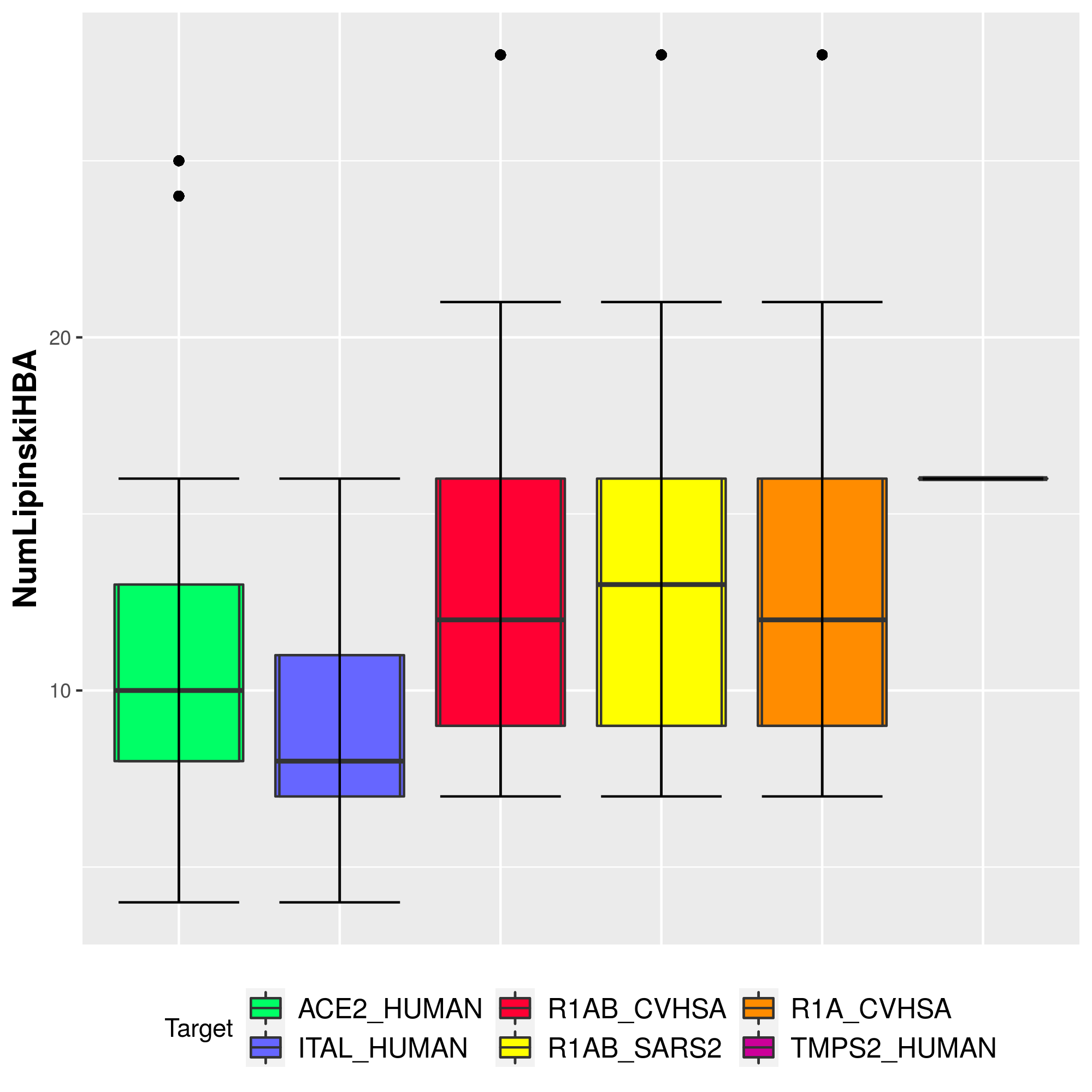
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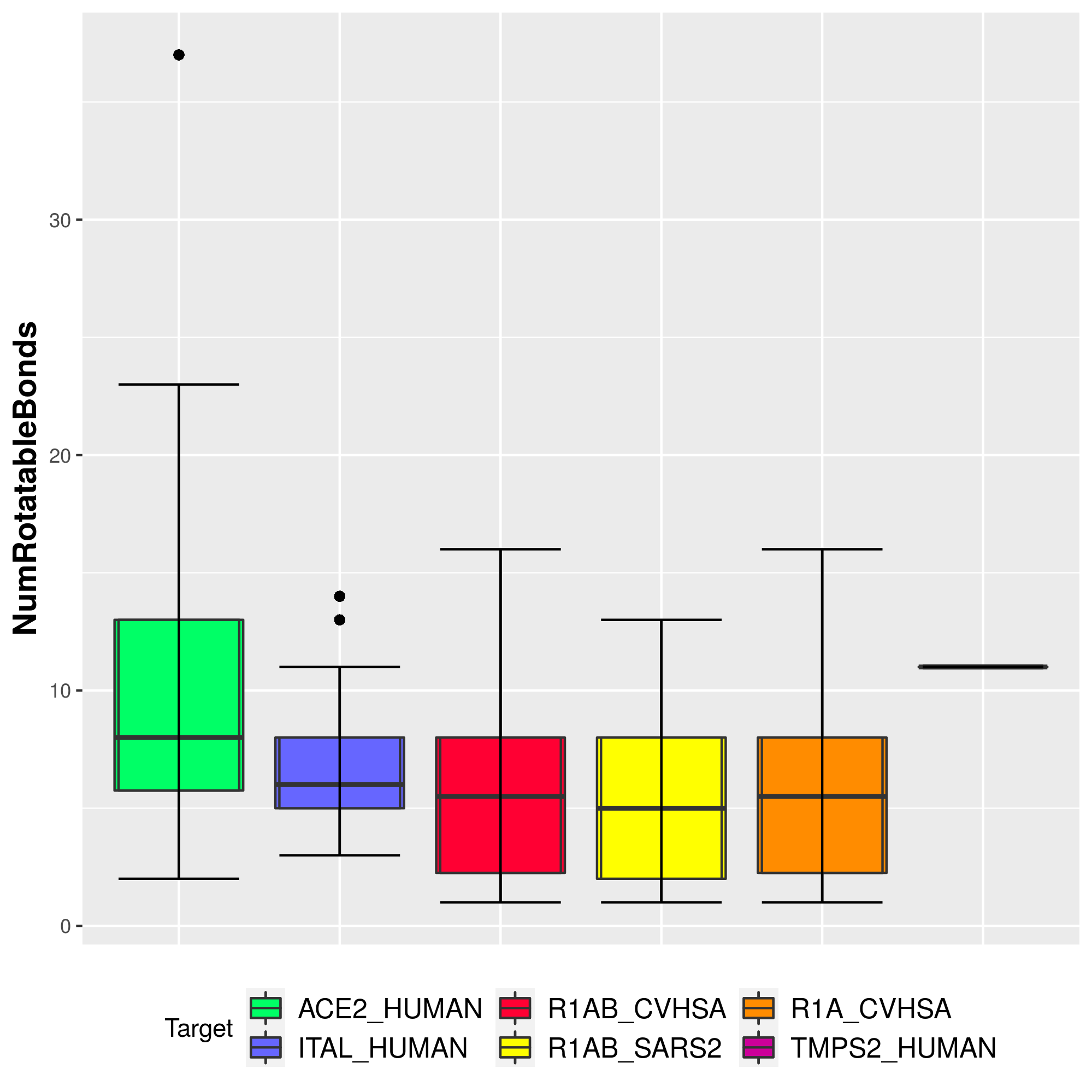
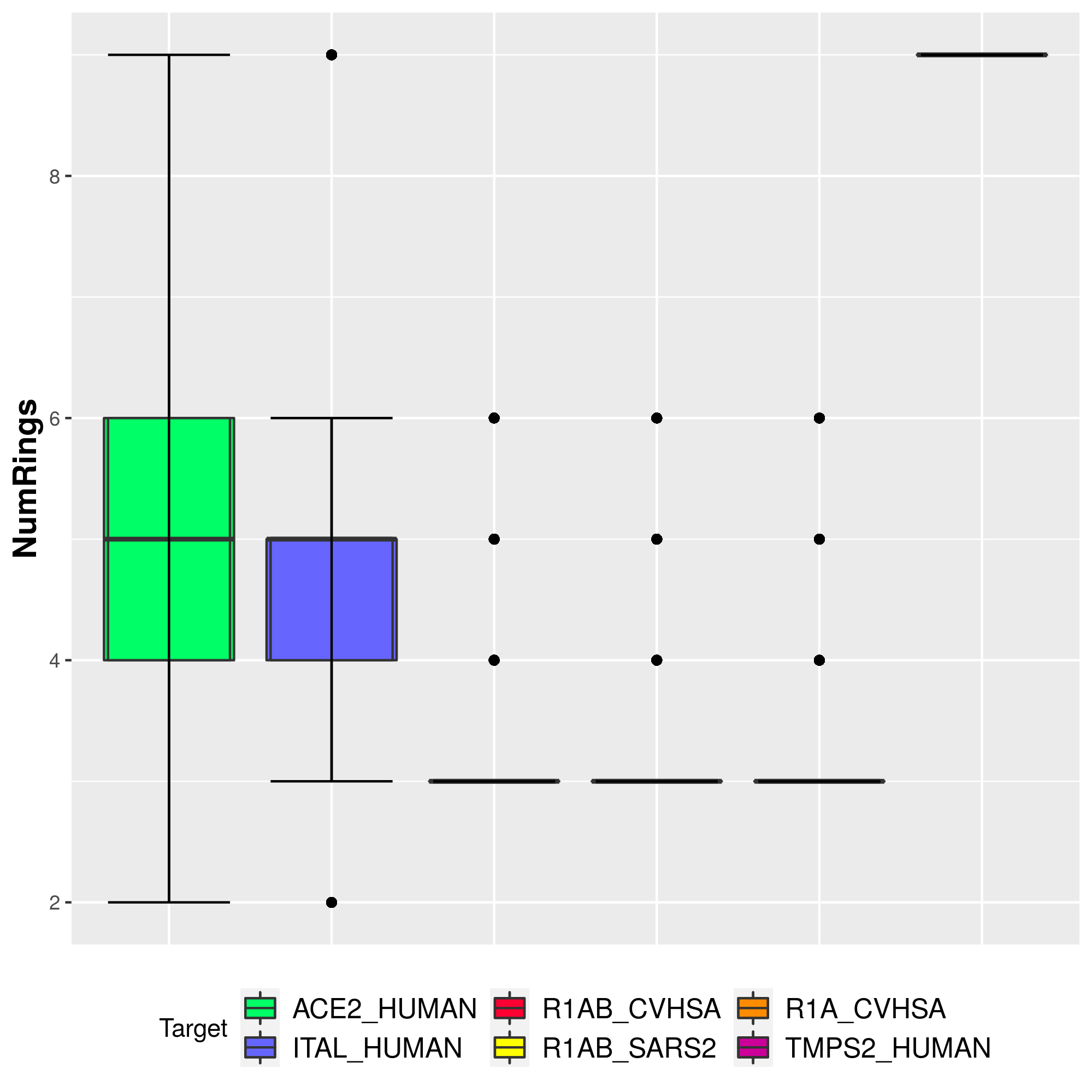
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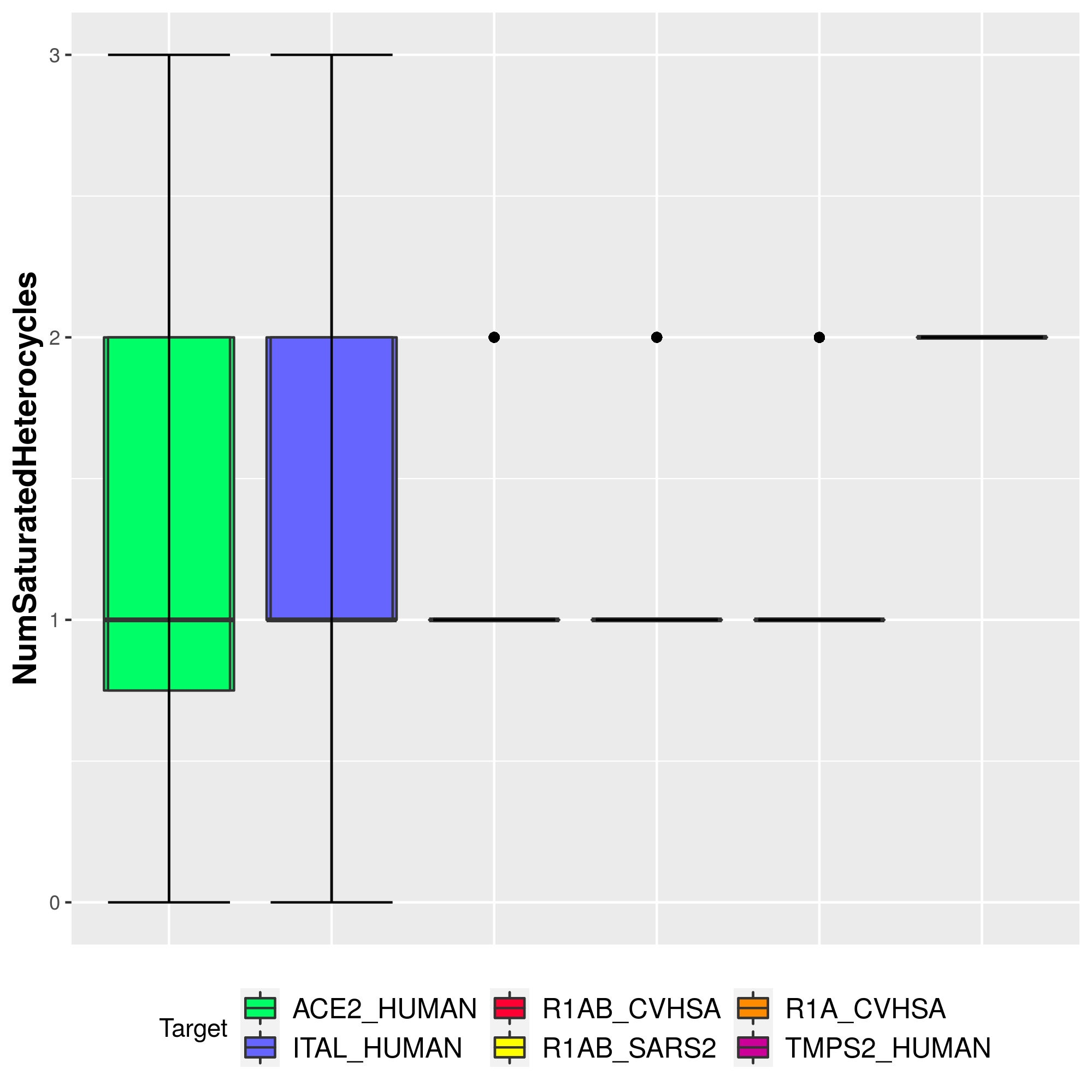
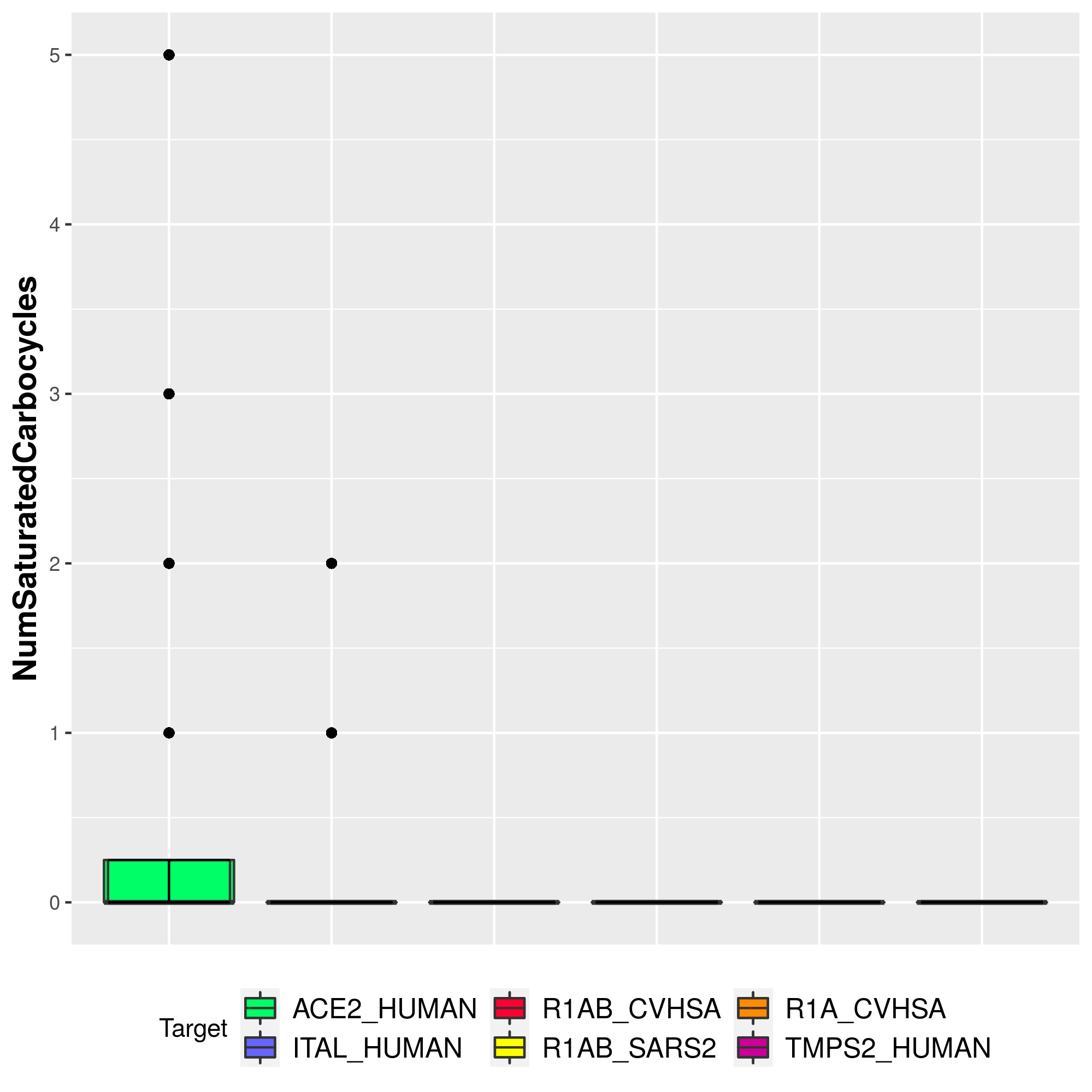
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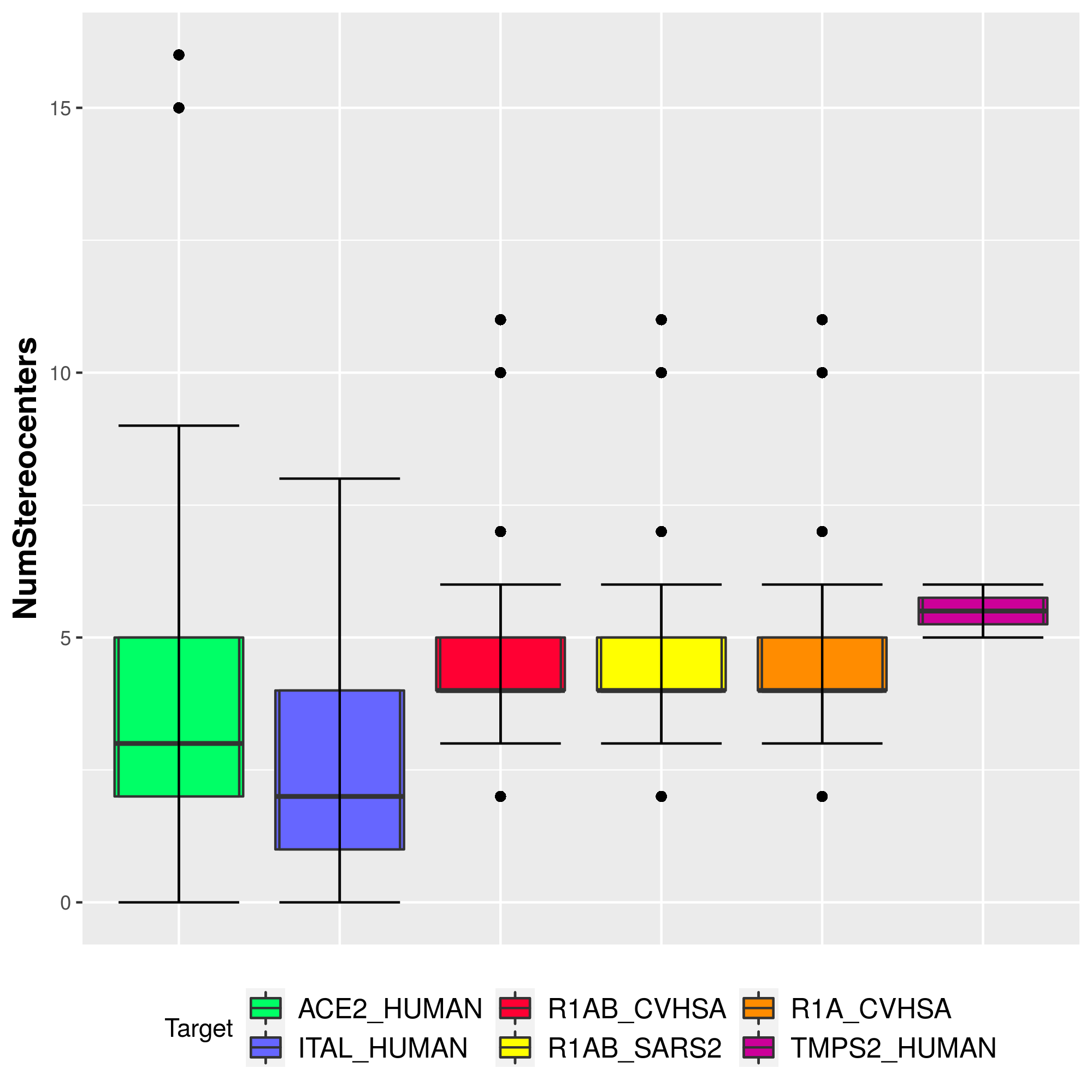
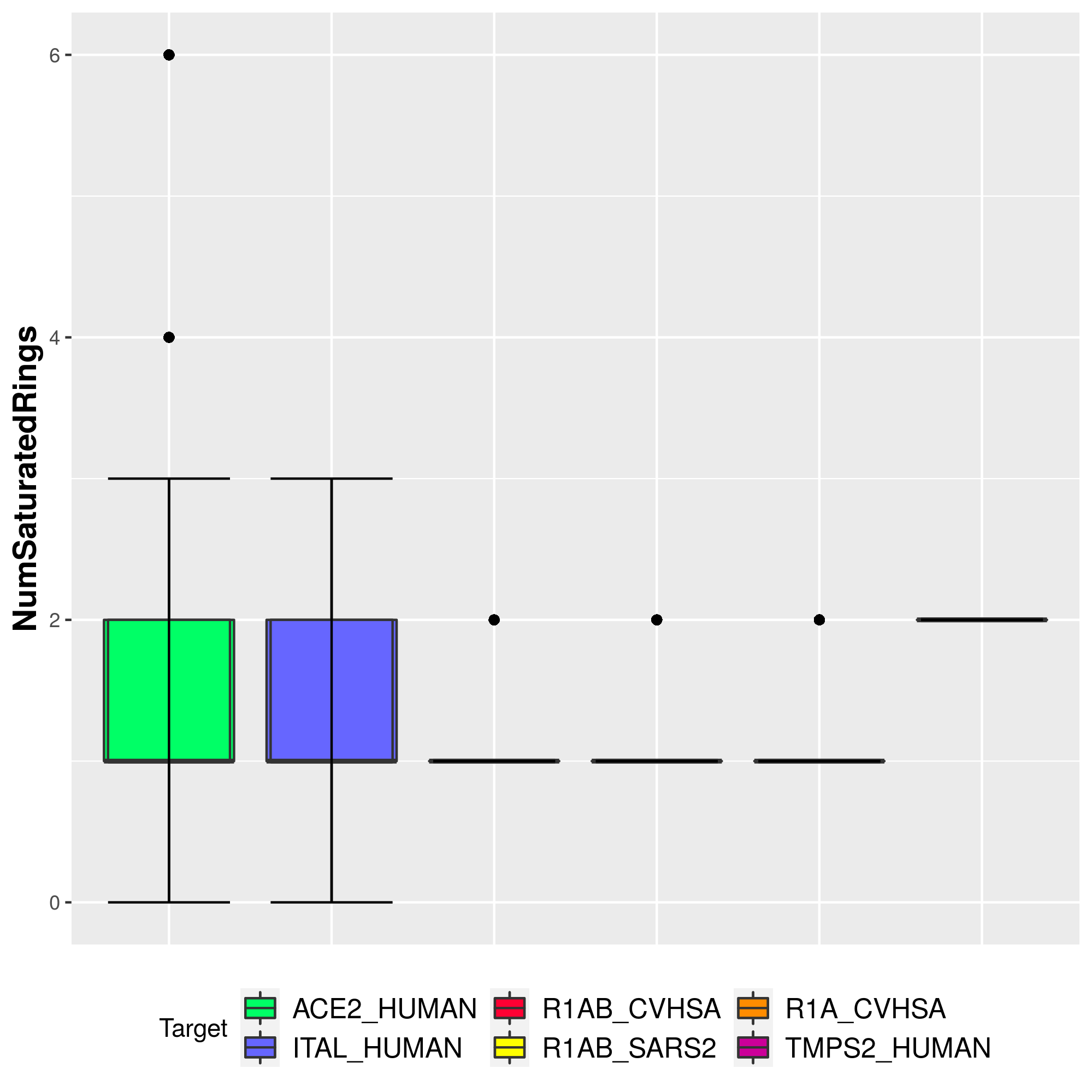
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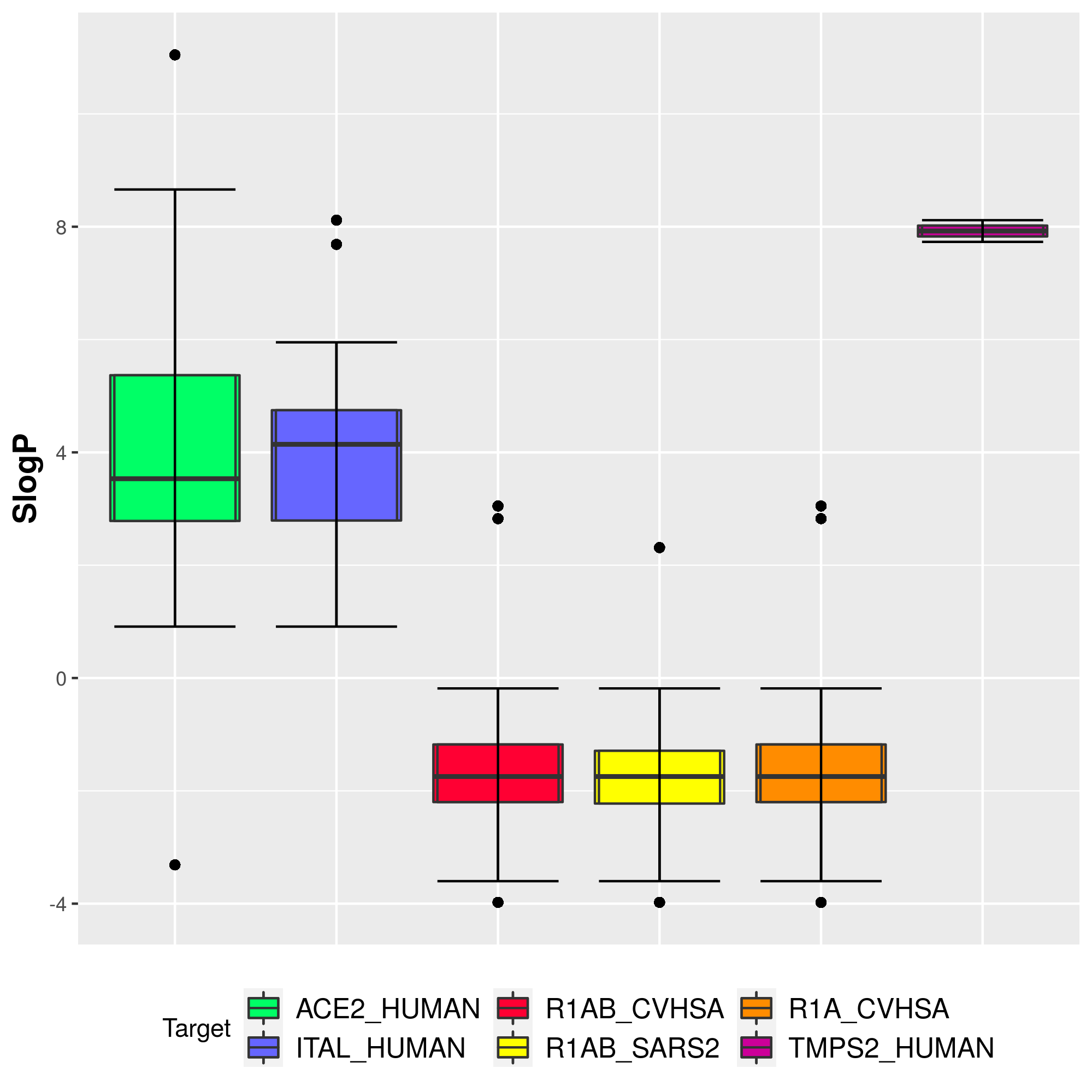
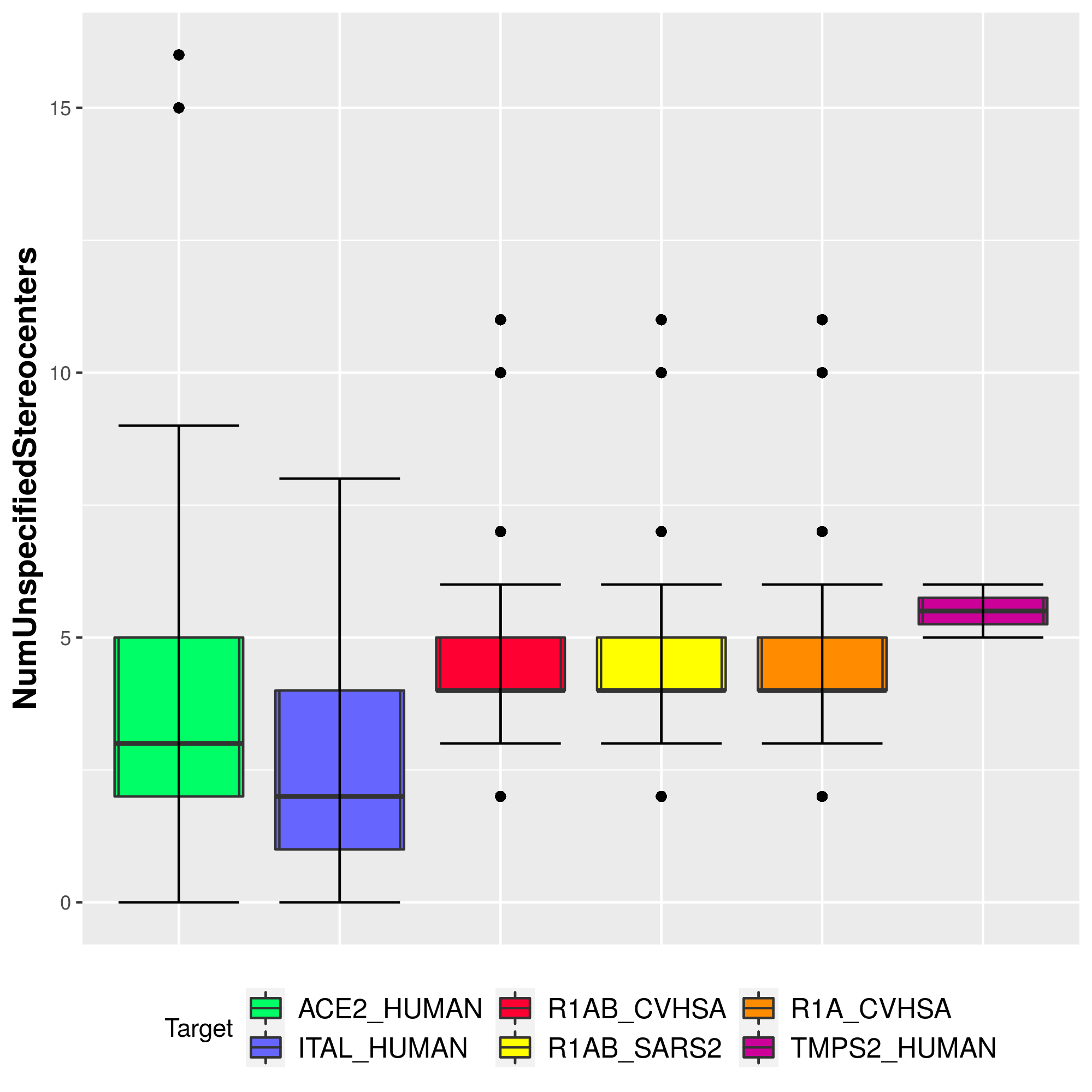
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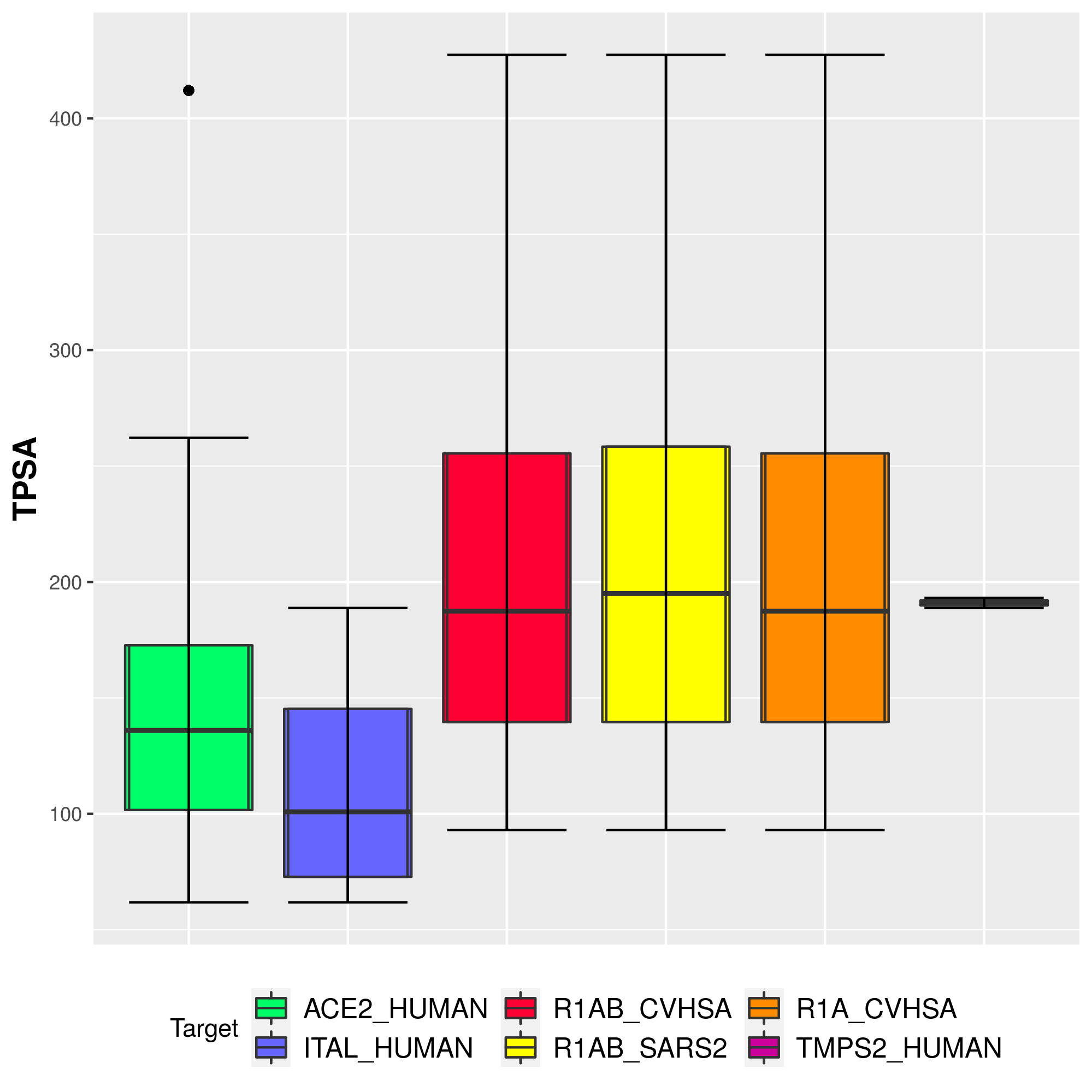
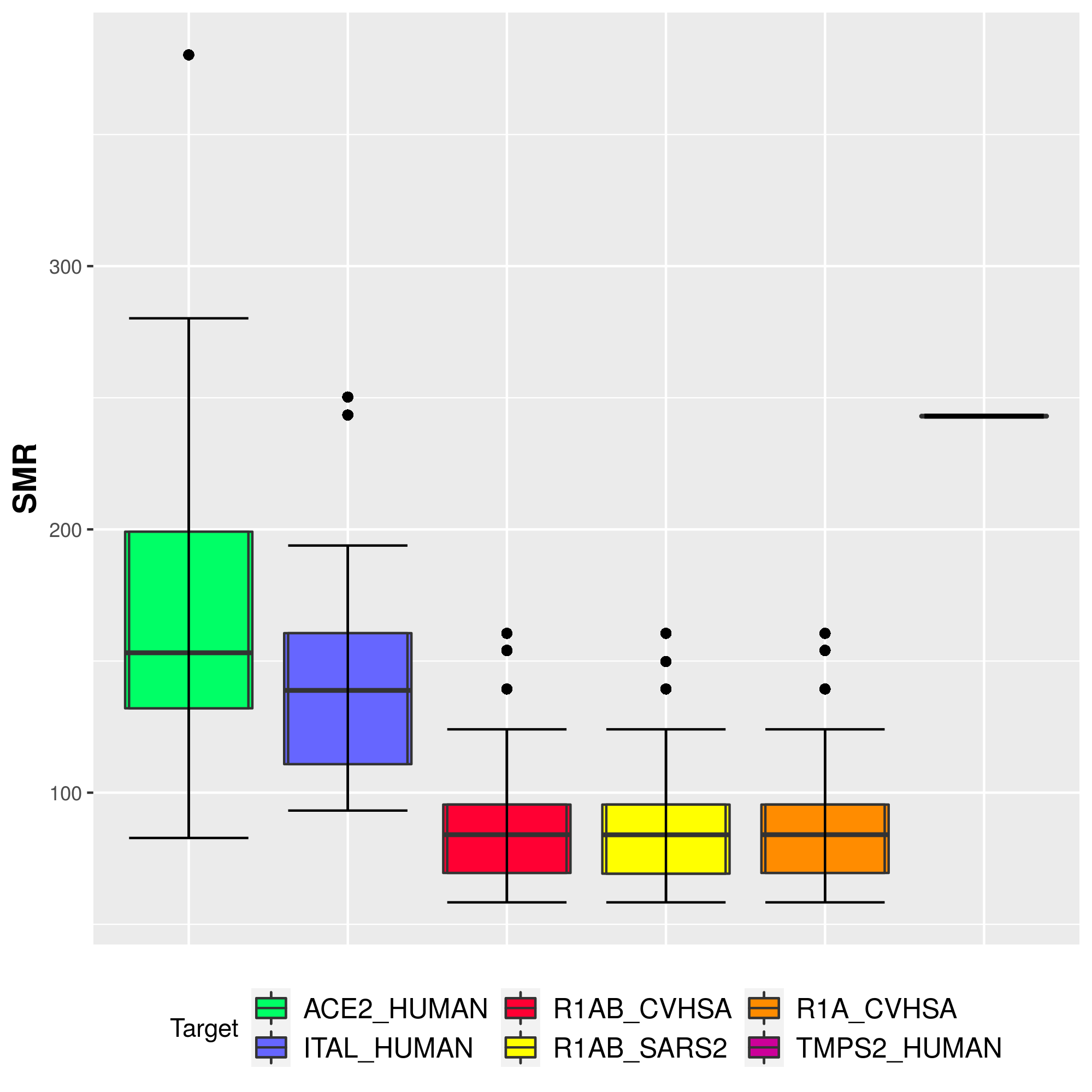
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***Supplementary Figure S2.*** *Distribution of distinct physicochemical descriptors of found hits across different targets.*