

Plasma proteome analyses in individuals of European ancestry identify cis-pQTLs and models for proteome-wide association studies

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Citation Report

#	ARTICLE	IF	CITATIONS
3	Multi-ancestry Mendelian randomization of omics traits revealing drug targets of COVID-19 severity. <i>EBioMedicine</i> , 2022, 81, 104112.	6.1	7
5	Quantitative trait locus (xQTL) approaches identify risk genes and drug targets from non-coding genome findings. <i>Human Molecular Genetics</i> , 0, , .	2.9	2
6	Proteomic profiling platforms head to head: Leveraging genetics and clinical traits to compare aptamer- and antibody-based methods. <i>Science Advances</i> , 2022, 8, .	10.3	57
8	Genetics-informed precision treatment formulation in schizophrenia and bipolar disorder. <i>American Journal of Human Genetics</i> , 2022, 109, 1620-1637.	6.2	12
9	How dysregulation of the immune system promotes diabetes mellitus and cardiovascular risk complications. <i>Frontiers in Cardiovascular Medicine</i> , 0, 9, .	2.4	11
10	Differences and commonalities in the genetic architecture of protein quantitative trait loci in European and Arab populations. <i>Human Molecular Genetics</i> , 2023, 32, 907-916.	2.9	9
11	Leveraging omics data to boost the power of genome-wide association studies. <i>Human Genetics and Genomics Advances</i> , 2022, 3, 100144.	1.7	0
12	Cross-Ancestry Investigation of Venous Thromboembolism Genomic Predictors. <i>Circulation</i> , 2022, 146, 1225-1242.	1.6	25
13	Proteome-wide Mendelian randomization in global biobank meta-analysis reveals multi-ancestry drug targets for common diseases. <i>Cell Genomics</i> , 2022, 2, 100195.	6.5	8
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29	Proteome-wide Mendelian randomization implicates nephronectin as an actionable mediator of the effect of obesity on COVID-19 severity. <i>Nature Metabolism</i> , 2023, 5, 248-264.	11.9	24
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31	Alterations in the Circulating Proteome Associated with Albuminuria. <i>Journal of the American Society of Nephrology: JASN</i> , 2023, 34, 1078-1089.	6.1	2
34	Integrative Post-Genome-Wide Association Study Analyses Relevant to Psychiatric Disorders: Imputing Transcriptome and Proteome Signals. <i>Complex Psychiatry</i> , 2023, 9, 130-144.	0.9	0

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36	Gene expression in African Americans, Puerto Ricans and Mexican Americans reveals ancestry-specific patterns of genetic architecture. <i>Nature Genetics</i> , 2023, 55, 952-963.	21.4	9
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79	Longitudinal plasma proteomics reveals biomarkers of alveolar-capillary barrier disruption in critically ill COVID-19 patients. <i>Nature Communications</i> , 2024, 15, .	12.8	2
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