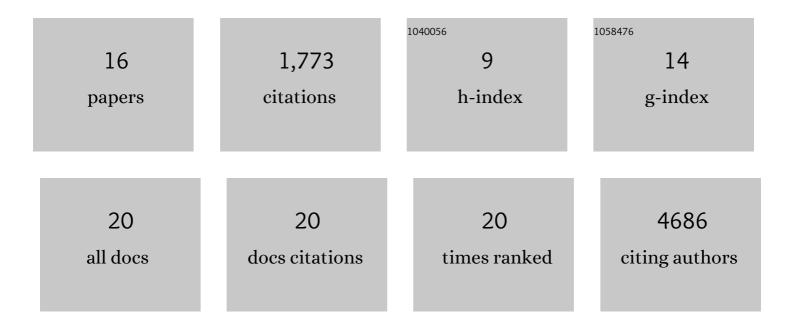
## Raili Ermel

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3700918/publications.pdf Version: 2024-02-01



**RAILLEDMEL** 

#	Article	IF	CITATIONS
1	Opportunities and challenges for transcriptome-wide association studies. Nature Genetics, 2019, 51, 592-599.	21.4	592
2	Association analyses based on false discovery rate implicate new loci for coronary artery disease. Nature Genetics, 2017, 49, 1385-1391.	21.4	571
3	Cardiometabolic risk loci share downstream cis- and trans-gene regulation across tissues and diseases. Science, 2016, 353, 827-830.	12.6	241
4	Cross-Tissue Regulatory Gene Networks in Coronary Artery Disease. Cell Systems, 2016, 2, 196-208.	6.2	120
5	A mechanistic framework for cardiometabolic and coronary artery diseases. , 2022, 1, 85-100.		51
6	Variation in the SERPINA6/SERPINA1 locus alters morning plasma cortisol, hepatic corticosteroid binding globulin expression, gene expression in peripheral tissues, and risk of cardiovascular disease. Journal of Human Genetics, 2021, 66, 625-636.	2.3	40
7	Expression Quantitative Trait Loci Acting Across Multiple Tissues Are Enriched in Inherited Risk for Coronary Artery Disease. Circulation: Cardiovascular Genetics, 2015, 8, 305-315.	5.1	39
8	An integrative multiomic network model links lipid metabolism to glucose regulation in coronary artery disease. Nature Communications, 2021, 12, 547.	12.8	35
9	Transcriptome-wide association study of coronary artery disease identifies novel susceptibility genes. Basic Research in Cardiology, 2022, 117, 6.	5.9	22
10	Global analysis of A-to-I RNA editing reveals association with common disease variants. PeerJ, 2018, 6, e4466.	2.0	21
11	Integrative Prioritization of Causal Genes for Coronary Artery Disease. Circulation Genomic and Precision Medicine, 2022, 15, CIRCGEN121003365.	3.6	11
12	Multiple independent mechanisms link gene polymorphisms in the region of ZEB2 with risk of coronary artery disease. Atherosclerosis, 2020, 311, 20-29.	0.8	9
13	<scp>European Society of Cardiology Working Group</scp> on Adult Congenital Heart Disease and <scp>Study Group for Adult Congenital Heart Care in Central and South Eastern European Countries</scp> consensus paper: current status, provision gaps and investment required. European lournal of Heart Failure. 2021. 23. 445-453.	7.1	9
14	Integrative analysis of loss-of-function variants in clinical and genomic data reveals novel genes associated with cardiovascular traits. BMC Medical Genomics, 2019, 12, 108.	1.5	8
15	The HDAC9-associated risk locus promotes coronary artery disease by governing TWIST1. PLoS Genetics, 2022, 18, e1010261.	3.5	2
16	OR09-04 Common Genetic Variants Associated with SERPINA6 Expression in Liver Influence Cortisol-Responsive Transcriptional Networks in Human Adipose Tissue. Journal of the Endocrine Society, 2020, 4, .	0.2	0