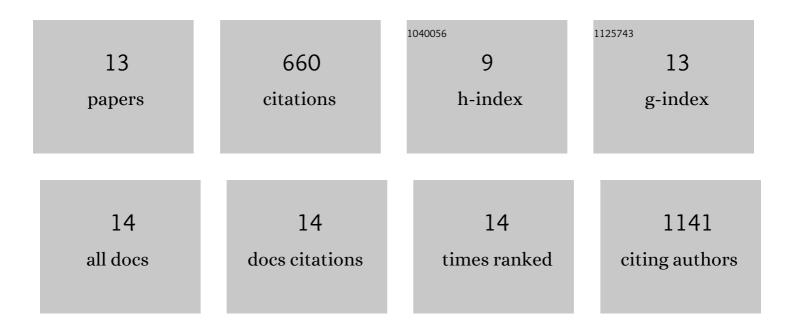


## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4573796/publications.pdf Version: 2024-02-01



DINCL

#	Article	IF	CITATIONS
1	A Liver-Enriched Long Non-Coding RNA, IncLSTR, Regulates Systemic Lipid Metabolism in Mice. Cell Metabolism, 2015, 21, 455-467.	16.2	247
2	Integrative Transcriptome Analyses of Metabolic Responses in Mice Define Pivotal LncRNA Metabolic Regulators. Cell Metabolism, 2016, 24, 627-639.	16.2	107
3	In vivo functional analysis of non-conserved human IncRNAs associated with cardiometabolic traits. Nature Communications, 2020, 11, 45.	12.8	69
4	A Long Non-coding RNA, IncLGR, Regulates Hepatic Glucokinase Expression and Glycogen Storage during Fasting. Cell Reports, 2016, 14, 1867-1875.	6.4	67
5	Comprehensive Characterization of 10,571 Mouse Large Intergenic Noncoding RNAs from Whole Transcriptome Sequencing. PLoS ONE, 2013, 8, e70835.	2.5	51
6	lncRNAKB, a knowledgebase of tissue-specific functional annotation and trait association of long noncoding RNA. Scientific Data, 2020, 7, 326.	5.3	40
7	Host and microbes in a pickle. Nature, 2010, 464, 1287-1288.	27.8	27
8	Identification of human long noncoding RNAs associated with nonalcoholic fatty liver disease and metabolic homeostasis. Journal of Clinical Investigation, 2021, 131, .	8.2	23
9	Comparative Transcriptomics Analyses in Livers of Mice, Humans, and Humanized Mice Define Human-Specific Gene Networks. Cells, 2020, 9, 2566.	4.1	19
10	Identification of Transcriptional Regulators That Bind to Long Noncoding RNAs by RNA Pull-Down and RNA Immunoprecipitation. Methods in Molecular Biology, 2018, 1783, 185-191.	0.9	5
11	Identification of Accessible Hepatic Gene Signatures for Interindividual Variations in Nutrigenomic Response to Dietary Supplementation of Omega-3 Fatty Acids. Cells, 2021, 10, 467.	4.1	2
12	Long noncoding RNAs-a new dimension in the molecular architecture of the bile acid/FXR pathway. Molecular and Cellular Endocrinology, 2021, 525, 111191.	3.2	2
13	In Vivo Functional Analysis of Nonconserved Human IncRNAs Using a Humanized Mouse Model. Methods in Molecular Biology, 2021, 2254, 339-347.	0.9	1