

# Haloom Rafehi

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6680369/publications.pdf>

Version: 2024-02-01

31  
papers

1,355  
citations

516710

16  
h-index

434195

31  
g-index

32  
all docs

32  
docs citations

32  
times ranked

2897  
citing authors

#	ARTICLE	IF	CITATIONS
1	A family study implicates <i>GBE1</i> in the etiology of autism spectrum disorder. <i>Human Mutation</i> , 2022, 43, 16-29.	2.5	2
2	Heterozygous <i>PNPT1</i> Variants Cause Spinocerebellar Ataxia Type 25. <i>Annals of Neurology</i> , 2022, 92, 122-137.	5.3	8
3	Infanticide vs. inherited cardiac arrhythmias. <i>Europace</i> , 2021, 23, 441-450.	1.7	21
4	The clinical utility of exome sequencing and extended bioinformatic analyses in adolescents and adults with a broad range of neurological phenotypes: an Australian perspective. <i>Journal of the Neurological Sciences</i> , 2021, 420, 117260.	0.6	16
5	Sex-Specific Control of Human Heart Maturation by the Progesterone Receptor. <i>Circulation</i> , 2021, 143, 1614-1628.	1.6	42
6	Rapid Diagnosis of Spinocerebellar Ataxia 36 in a Three-Generation Family Using Short-Read Whole-Genome Sequencing Data. <i>Movement Disorders</i> , 2020, 35, 1675-1679.	3.9	12
7	Familial adult myoclonic epilepsy type 1 <i>SAMD12</i> TTTCA repeat expansion arose 17,000 years ago and is present in Sri Lankan and Indian families. <i>European Journal of Human Genetics</i> , 2020, 28, 973-978.	2.8	23
8	Bioinformatics-Based Identification of Expanded Repeats: A Non-reference Intronic Pentamer Expansion in <i>RFC1</i> Causes <i>CANVAS</i> . <i>American Journal of Human Genetics</i> , 2019, 105, 151-165.	6.2	170
9	Familial early onset Parkinson's disease caused by a homozygous frameshift variant in <i>PARK7</i> : Clinical features and literature update. <i>Parkinsonism and Related Disorders</i> , 2019, 64, 308-311.	2.2	7
10	Unstable TTTTA/TTTCA expansions in <i>MARCH6</i> are associated with Familial Adult Myoclonic Epilepsy type 3. <i>Nature Communications</i> , 2019, 10, 4919.	12.8	111
11	Galectin-3 deficiency ameliorates fibrosis and remodeling in dilated cardiomyopathy mice with enhanced <i>Mst1</i> signaling. <i>American Journal of Physiology - Heart and Circulatory Physiology</i> , 2019, 316, H45-H60.	3.2	47
12	Systems approach to the pharmacological actions of HDAC inhibitors reveals EP300 activities and convergent mechanisms of regulation in diabetes. <i>Epigenetics</i> , 2017, 12, 991-1003.	2.7	20
13	Multicellular Transcriptional Analysis of Mammalian Heart Regeneration. <i>Circulation</i> , 2017, 136, 1123-1139.	1.6	222
14	Metabolism and chromatin dynamics in health and disease. <i>Molecular Aspects of Medicine</i> , 2017, 54, 1-15.	6.4	14
15	MeCP2 interacts with chromosomal microRNAs in brain. <i>Epigenetics</i> , 2017, 12, 1028-1037.	2.7	16
16	NET silencing by <i>let-7i</i> in postural tachycardia syndrome. <i>JCI Insight</i> , 2017, 2, e90183.	5.0	17
17	Pharmacological Histone Deacetylation Distinguishes Transcriptional Regulators. <i>Current Topics in Medicinal Chemistry</i> , 2017, 17, 1611-1622.	2.1	9
18	HDAC Inhibition in Vascular Endothelial Cells Regulates the Expression of ncRNAs. <i>Non-coding RNA</i> , 2016, 2, 4.	2.6	14

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19	Current perspectives in Set7 mediated stem cell differentiation. Non-coding RNA, 2016, 2, 14.	2.6	2
20	Improving understanding of chromatin regulatory proteins and potential implications for drug discovery. Expert Review of Proteomics, 2016, 13, 435-445.	3.0	2
21	RNA sequencing supports distinct reactive oxygen species-mediated pathways of apoptosis by high and low size mass fractions of Bay leaf ( <i>Lauris nobilis</i> ) in HT-29 cells. Food and Function, 2015, 6, 2507-2524.	4.6	8
22	HDAC inhibition attenuates cardiac hypertrophy by acetylation and deacetylation of target genes. Epigenetics, 2015, 10, 418-430.	2.7	111
23	Vascular histone deacetylation by pharmacological HDAC inhibition. Genome Research, 2014, 24, 1271-1284.	5.5	79
24	Mechanisms of Action of Phenolic Compounds in Olive. Journal of Dietary Supplements, 2012, 9, 96-109.	2.6	47
25	Epigenetic mechanisms in the pathogenesis of diabetic foot ulcers. Journal of Diabetes and Its Complications, 2012, 26, 554-561.	2.3	29
26	A pipeline for the identification and characterization of chromatin modifications derived from ChIP-Seq datasets. Biochimie, 2012, 94, 2353-2359.	2.6	6
27	Investigation of the biological properties of Cinnulin PF in the context of diabetes: mechanistic insights by genome-wide mRNA-Seq analysis. Pathobiology of Aging & Age Related Diseases, 2012, 2, 11905.	1.1	3
28	Investigation into the biological properties of the olive polyphenol, hydroxytyrosol: mechanistic insights by genome-wide mRNA-Seq analysis. Genes and Nutrition, 2012, 7, 343-355.	2.5	40
29	Controversies surrounding the clinical potential of cinnamon for the management of diabetes. Diabetes, Obesity and Metabolism, 2012, 14, 493-499.	4.4	59
30	Clonogenic Assay: Adherent Cells. Journal of Visualized Experiments, 2011, , .	0.3	145
31	Genetic and epigenetic events in diabetic wound healing. International Wound Journal, 2011, 8, 12-21.	2.9	53