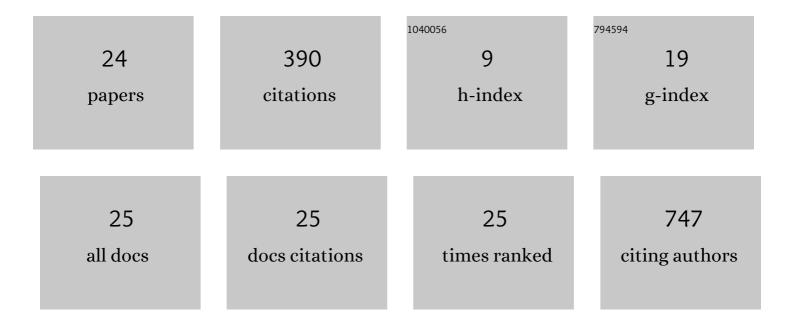
Artem M Kiselev

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5040738/publications.pdf

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#	Article	IF	CITATIONS
1	Characterization of the novel heterozygous SCN5A genetic variant Y739D associated with Brugada syndrome. Biochemistry and Biophysics Reports, 2022, 30, 101249.	1.3	2
2	Genetic Spectrum of Left Ventricular Non-Compaction in Paediatric Patients. Cardiology, 2020, 145, 746-756.	1.4	9
3	The Activity of KIF14, Mieap, and EZR in a New Type of the Invasive Component, Torpedo-Like Structures, Predetermines the Metastatic Potential of Breast Cancer. Cancers, 2020, 12, 1909.	3.7	10
4	Two New Cases of Hypertrophic Cardiomyopathy and Skeletal Muscle Features Associated with ALPK3 Homozygous and Compound Heterozygous Variants. Genes, 2020, 11, 1201.	2.4	20
5	Insights Image for "Dysregulation of Notch signaling in cardiac mesenchymal cells of patients with Tetralogy of Fallot― Pediatric Research, 2020, 88, 139-139.	2.3	Ο
6	New germline mutations in non-BRCA genes among breast cancer women of Mongoloid origin. Molecular Biology Reports, 2020, 47, 5315-5321.	2.3	4
7	Systematic dissection of biases in whole-exome and whole-genome sequencing reveals major determinants of coding sequence coverage. Scientific Reports, 2020, 10, 2057.	3.3	72
8	Dysregulation of Notch signaling in cardiac mesenchymal cells of patients with tetralogy of Fallot. Pediatric Research, 2020, 88, 38-47.	2.3	5
9	Atomic Mechanisms of Timothy Syndrome-Associated Mutations in Calcium Channel Cav1.2. Frontiers in Physiology, 2019, 10, 335.	2.8	11
10	Characterization of a novel SCN5A genetic variant A1294G associated with mixed clinical phenotype. Biochemical and Biophysical Research Communications, 2019, 516, 777-783.	2.1	8
11	New germline BRCA2 gene variant in the Tuvinian Mongol breast cancer patients. Molecular Biology Reports, 2019, 46, 5537-5541.	2.3	6
12	Truncating Variant in Myof Gene Is Associated With Limb-Girdle Type Muscular Dystrophy and Cardiomyopathy. Frontiers in Genetics, 2019, 10, 608.	2.3	10
13	Draft Genome Sequence of <i>Coxiella burnetii</i> Historical Strain Leningrad-2, Isolated from Blood of a Patient with Acute Q Fever in Saint Petersburg, Russia. Genome Announcements, 2018, 6, .	0.8	1
14	De novo mutations in <i>FLNC</i> leading to early-onset restrictive cardiomyopathy and congenital myopathy. Human Mutation, 2018, 39, 1161-1172.	2.5	49
15	Rare Case of Ulnar-Mammary-Like Syndrome With Left Ventricular Tachycardia and Lack of TBX3 Mutation. Frontiers in Genetics, 2018, 9, 209.	2.3	4
16	Structural consequences of mutations associated with idiopathic restrictive cardiomyopathy. Amino Acids, 2017, 49, 1815-1829.	2.7	5
17	The exon junction complex factor Y14 is dynamic in the nucleus of the beetle Tribolium castaneum during late oogenesis. Molecular Cytogenetics, 2017, 10, 41.	0.9	1
18	Progressive cardiac conduction disease associated with a DSP gene mutation. International Journal of Cardiology, 2016, 216, 188-189.	1.7	9

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#	Article	IF	CITATIONS
19	Chicken rRNA Gene Cluster Structure. PLoS ONE, 2016, 11, e0157464.	2.5	24
20	Genetic Spectrum of Idiopathic Restrictive Cardiomyopathy Uncovered by Next-Generation Sequencing. PLoS ONE, 2016, 11, e0163362.	2.5	78
21	Characterization of Tribolium castaneum oocyte nuclear structures using microinjection of a fusion nuclear protein mRNA. Molecular Reproduction and Development, 2015, 82, 628-629.	2.0	5
22	Cysteine-Rich Atrial Secretory Protein from the Snail Achatina achatina: Purification and Structural Characterization. PLoS ONE, 2015, 10, e0138787.	2.5	5
23	Neonatal hypertrophic cardiomyopathy caused by double mutation in RAS pathway genes. International Journal of Cardiology, 2015, 184, 272-273.	1.7	2
24	Variants in the <i>NOTCH1</i> Gene in Patients with Aortic Coarctation. Congenital Heart Disease, 2014, 9, 391-396.	0.2	48