

Torgny Karlsson

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1483066/publications.pdf>

Version: 2024-02-01

20
papers

1,003
citations

687363

13
h-index

794594

19
g-index

22
all docs

22
docs citations

22
times ranked

2353
citing authors

#	ARTICLE	IF	CITATIONS
1	Contribution of rare whole-genome sequencing variants to plasma protein levels and the missing heritability. <i>Nature Communications</i> , 2022, 13, 2532.	12.8	9
2	Oral Contraceptives, Hormone Replacement Therapy, and Stroke Risk. <i>Stroke</i> , 2022, 53, 3107-3115.	2.0	20
3	Investigating the Effect of Estradiol Levels on the Risk of Breast, Endometrial, and Ovarian Cancer. <i>Journal of the Endocrine Society</i> , 2022, 6, .	0.2	10
4	Modification of Heritability for Educational Attainment and Fluid Intelligence by Socioeconomic Deprivation in the UK Biobank. <i>American Journal of Psychiatry</i> , 2021, 178, 625-634.	7.2	15
5	Genome-wide Association Study of Estradiol Levels and the Causal Effect of Estradiol on Bone Mineral Density. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2021, 106, e4471-e4486.	3.6	41
6	Characterization of the human <i>ABO</i> genotypes and their association to common inflammatory and cardiovascular diseases in the UK Biobank. <i>American Journal of Hematology</i> , 2021, 96, 1350-1362.	4.1	8
7	1414Hormone replacement therapy, oral contraceptives and stroke risk. <i>International Journal of Epidemiology</i> , 2021, 50, .	1.9	0
8	Time-Dependent Effects of Oral Contraceptive Use on Breast, Ovarian, and Endometrial Cancers. <i>Cancer Research</i> , 2021, 81, 1153-1162.	0.9	42
9	Causal effects of inflammatory protein biomarkers on inflammatory diseases. <i>Science Advances</i> , 2021, 7, eabl4359.	10.3	18
10	Genome-wide association analysis of 350,000 Caucasians from the UK Biobank identifies novel loci for asthma, hay fever and eczema. <i>Human Molecular Genetics</i> , 2019, 28, 4022-4041.	2.9	110
11	Associations of autozygosity with a broad range of human phenotypes. <i>Nature Communications</i> , 2019, 10, 4957.	12.8	84
12	Contribution of genetics to visceral adiposity and its relation to cardiovascular and metabolic disease. <i>Nature Medicine</i> , 2019, 25, 1390-1395.	30.7	172
13	Genome-wide association study of body fat distribution identifies adiposity loci and sex-specific genetic effects. <i>Nature Communications</i> , 2019, 10, 339.	12.8	163
14	Improved power and precision with whole genome sequencing data in genome-wide association studies of inflammatory biomarkers. <i>Scientific Reports</i> , 2019, 9, 16844.	3.3	43
15	Genetic variants influencing phenotypic variance heterogeneity. <i>Human Molecular Genetics</i> , 2018, 27, 799-810.	2.9	30
16	Breast-feeding and risk of asthma, hay fever, and eczema. <i>Journal of Allergy and Clinical Immunology</i> , 2018, 141, 1157-1159.e9.	2.9	17
17	Tea and coffee consumption in relation to DNA methylation in four European cohorts. <i>Human Molecular Genetics</i> , 2017, 26, 3221-3231.	2.9	25
18	Gene-environment interaction study for BMI reveals interactions between genetic factors and physical activity, alcohol consumption and socioeconomic status. <i>PLoS Genetics</i> , 2017, 13, e1006977.	3.5	125

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19	RNA-sequence data normalization through in silico prediction of reference genes: the bacterial response to DNA damage as case study. BioData Mining, 2017, 10, 30.	4.0	15
20	The relative contribution of DNA methylation and genetic variants on protein biomarkers for human diseases. PLoS Genetics, 2017, 13, e1007005.	3.5	54