

# 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

687335

13  
h-index

794568

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. <i>BioData Mining</i> , 2017, 10, 30.	4.0	15
20	The relative contribution of DNA methylation and genetic variants on protein biomarkers for human diseases. <i>PLoS Genetics</i> , 2017, 13, e1007005.	3.5	54