## Mathias Rask-Andersen

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6227813/publications.pdf

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47 papers

4,709 citations

304743 22 h-index 214800 47 g-index

49 all docs 49 docs citations

49 times ranked 9398 citing authors

#	Article	IF	CITATIONS
1	Trends in GPCR drug discovery: new agents, targets and indications. Nature Reviews Drug Discovery, 2017, 16, 829-842.	46.4	1,773
2	Trends in the exploitation of novel drug targets. Nature Reviews Drug Discovery, 2011, 10, 579-590.	46.4	720
3	The Druggable Genome: Evaluation of Drug Targets in Clinical Trials Suggests Major Shifts in Molecular Class and Indication. Annual Review of Pharmacology and Toxicology, 2014, 54, 9-26.	9.4	262
4	Advances in kinase targeting: current clinical use and clinical trials. Trends in Pharmacological Sciences, 2014, 35, 604-620.	8.7	178
5	Contribution of genetics to visceral adiposity and its relation to cardiovascular and metabolic disease. Nature Medicine, 2019, 25, 1390-1395.	30.7	172
6	Genome-wide association study of body fat distribution identifies adiposity loci and sex-specific genetic effects. Nature Communications, 2019, 10, 339.	12.8	163
7	Gene-environment interaction study for BMI reveals interactions between genetic factors and physical activity, alcohol consumption and socioeconomic status. PLoS Genetics, 2017, 13, e1006977.	3 <b>.</b> 5	125
8	Many obesity-associated SNPs strongly associate with DNA methylation changes at proximal promoters and enhancers. Genome Medicine, 2015, 7, 103.	8.2	124
9	Genome-wide association analysis of 350 000 Caucasians from the UK Biobank identifies novel loci for asthma, hay fever and eczema. Human Molecular Genetics, 2019, 28, 4022-4041.	2.9	110
10	Molecular mechanisms underlying anorexia nervosa: Focus on human gene association studies and systems controlling food intake. Brain Research Reviews, 2010, 62, 147-164.	9.0	106
11	A debate on current eating disorder diagnoses in light of neurobiological findings: is it time for a spectrum model?. BMC Psychiatry, 2012, 12, 76.	2.6	90
12	Solute carriers as drug targets: Current use, clinical trials and prospective. Molecular Aspects of Medicine, 2013, 34, 702-710.	6.4	89
13	Epigenome-wide association study reveals differential DNA methylation in individuals with a history of myocardial infarction. Human Molecular Genetics, 2016, 25, ddw302.	2.9	88
14	Soluble ligands as drug targets. Nature Reviews Drug Discovery, 2020, 19, 695-710.	46.4	63
15	The relative contribution of DNA methylation and genetic variants on protein biomarkers for human diseases. PLoS Genetics, 2017, 13, e1007005.	3.5	54
16	Orphan Drugs and Their Impact on Pharmaceutical Development. Trends in Pharmacological Sciences, 2018, 39, 525-535.	8.7	43
17	Improved power and precision with whole genome sequencing data in genome-wide association studies of inflammatory biomarkers. Scientific Reports, 2019, 9, 16844.	3.3	43
18	Genetic and expression studies of SMN2 gene in Russian patients with spinal muscular atrophy type II and III. BMC Medical Genetics, 2011, 12, 96.	2.1	36

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19	Differential effects of fluoxetine and venlafaxine on memory recognition: Possible mechanisms of action. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2012, 38, 159-167.	4.8	32
20	Determination of the obesity-associated gene variants within the entire FTO gene by ultra-deep targeted sequencing in obese and lean children. International Journal of Obesity, 2013, 37, 424-431.	3.4	32
21	Genetic variants influencing phenotypic variance heterogeneity. Human Molecular Genetics, 2018, 27, 799-810.	2.9	30
22	Tea and coffee consumption in relation to DNA methylation in four European cohorts. Human Molecular Genetics, 2017, 26, 3221-3231.	2.9	25
23	Association of TMEM18 variants with BMI and waist circumference in children and correlation of mRNA expression in the PFC with body weight in rats. European Journal of Human Genetics, 2012, 20, 192-197.	2.8	24
24	Ghrelin effects expression of several genes associated with depression-like behavior. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2015, 56, 227-234.	4.8	23
25	Functional coupling analysis suggests link between the obesity gene FTO and the BDNF-NTRK2 signaling pathway. BMC Neuroscience, 2011, 12, 117.	1.9	22
26	Scrutinizing the FTO locus: compelling evidence for a complex, long-range regulatory context. Human Genetics, 2015, 134, 1183-1193.	3.8	22
27	meQTL and ncRNA functional analyses of 102 GWAS-SNPs associated with depression implicate HACE1 and SHANK2 genes. Clinical Epigenetics, 2020, 12, 99.	4.1	19
28	Personality, lifestyle and job satisfaction: causal association between neuroticism and job satisfaction using Mendelian randomisation in the UK biobank cohort. Translational Psychiatry, 2020, 10, 11.	4.8	19
29	Causal effects of inflammatory protein biomarkers on inflammatory diseases. Science Advances, 2021, 7, eabl4359.	10.3	18
30	Postprandial alterations in whole-blood DNA methylation are mediated by changes in white blood cell composition. American Journal of Clinical Nutrition, 2016, 104, 518-525.	4.7	17
31	Epigenome-wide DNA methylation study of IgE concentration in relation to self-reported allergies. Epigenomics, 2017, 9, 407-418.	2.1	17
32	Breast-feeding and risk of asthma, hay fever, and eczema. Journal of Allergy and Clinical Immunology, 2018, 141, 1157-1159.e9.	2.9	17
33	The MAP2K5-linked SNP rs2241423 is associated with BMI and obesity in two cohorts of Swedish and Greek children. BMC Medical Genetics, 2012, 13, 36.	2.1	16
34	A combined genome-wide association and molecular study of age-related hearing loss in H. sapiens. BMC Medicine, 2021, 19, 302.	5.5	16
35	Obsessive-compulsivity and working memory are associated with differential prefrontal cortex and insula activation in adolescents with a recent diagnosis of an eating disorder. Psychiatry Research - Neuroimaging, 2014, 224, 246-253.	1.8	15
36	Reduced vasopressin receptors activation mediates the anti-depressant effects of fluoxetine and venlafaxine in bulbectomy model of depression. Psychopharmacology, 2016, 233, 1077-1086.	3.1	15

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37	Modification of Heritability for Educational Attainment and Fluid Intelligence by Socioeconomic Deprivation in the UK Biobank. American Journal of Psychiatry, 2021, 178, 625-634.	7.2	15
38	Exposure to a high-fat high-sugar diet causes strong up-regulation of proopiomelanocortin and differentially affects dopamine D1 and D2 receptor gene expression in the brainstem of rats. Neuroscience Letters, 2014, 559, 18-23.	2.1	14
39	Association of the LINGO2-related SNP rs10968576 with body mass in a cohort of elderly Swedes. Molecular Genetics and Genomics, 2015, 290, 1485-1491.	2.1	13
40	Adolescents newly diagnosed with eating disorders have structural differences in brain regions linked with eating disorder symptoms. Nordic Journal of Psychiatry, 2017, 71, 188-196.	1.3	13
41	CDKAL1-Related Single Nucleotide Polymorphisms Are Associated with Insulin Resistance in a Cross-Sectional Cohort of Greek Children. PLoS ONE, 2014, 9, e93193.	2.5	8
42	The STK33-Linked SNP rs4929949 Is Associated with Obesity and BMI in Two Independent Cohorts of Swedish and Greek Children. PLoS ONE, 2013, 8, e71353.	2.5	7
43	The role of DNA methylation in the pathogenesis of disease: what can epigenome-wide association studies tell?. Epigenomics, 2016, 8, 5-7.	2.1	6
44	Insulin receptorâ€ike ectodomain genes and splice variants are found in both arthropods and human brain cDNA. Journal of Systematics and Evolution, 2013, 51, 664-670.	3.1	4
45	Determination of obesity associated gene variants related to TMEM18 through ultra-deep targeted re-sequencing in a case-control cohort for pediatric obesity. Genetical Research, 2015, 97, e16.	0.9	4
46	Accelerated epigenetic aging in suicide attempters uninfluenced by high intent-to-die and choice of lethal methods. Translational Psychiatry, 2022, 12, .	4.8	3
47	HPA-axis dysregulation is not associated with accelerated epigenetic aging in patients with hypersexual disorder. Psychoneuroendocrinology, 2022, 141, 105765.	2.7	2