Erik A Ehli

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6159432/publications.pdf

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394421 345221 1,545 40 19 36 citations h-index g-index papers 43 43 43 3920 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Genome-wide Association Meta-analysis of Childhood and Adolescent Internalizing Symptoms. Journal of the American Academy of Child and Adolescent Psychiatry, 2022, 61, 934-945.	0.5	26
2	The Prostate Cancer Androgen Receptor Cistrome in African American Men Associates with Upregulation of Lipid Metabolism and Immune Response. Cancer Research, 2022, 82, 2848-2859.	0.9	17
3	Noninvasive Assessment of Epidermal Genomic Markers of UV Exposure in Skin. Journal of Investigative Dermatology, 2021, 141, 124-131.e2.	0.7	6
4	Genetic meta-analysis of twin birth weight shows high genetic correlation with singleton birth weight. Human Molecular Genetics, 2021, 30, 1894-1905.	2.9	6
5	Implementation and implications for polygenic risk scores in healthcare. Human Genomics, 2021, 15, 46.	2.9	36
6	Genetic association study of childhood aggression across raters, instruments, and age. Translational Psychiatry, 2021, 11, 413.	4.8	31
7	Continuity of Genetic Risk for Aggressive Behavior Across the Life-Course. Behavior Genetics, 2021, 51, 592-606.	2.1	13
8	Identical twins carry a persistent epigenetic signature of early genome programming. Nature Communications, 2021, 12, 5618.	12.8	26
9	Integrative transcriptomic analysis for linking acute stress responses to squamous cell carcinoma development. Scientific Reports, 2020, 10, 17209.	3 . 3	4
10	Utilization of archived neonatal dried blood spots for genome-wide genotyping. PLoS ONE, 2020, 15, e0229352.	2.5	7
11	Intergenerational Transmission of Education and ADHD: Effects of Parental Genotypes. Behavior Genetics, 2020, 50, 221-232.	2.1	45
12	Urinary Amine and Organic Acid Metabolites Evaluated as Markers for Childhood Aggression: The ACTION Biomarker Study. Frontiers in Psychiatry, 2020, 11, 165.	2.6	19
13	Utilization of archived neonatal dried blood spots for genome-wide genotyping., 2020, 15, e0229352.		O
14	Utilization of archived neonatal dried blood spots for genome-wide genotyping., 2020, 15, e0229352.		0
15	Utilization of archived neonatal dried blood spots for genome-wide genotyping., 2020, 15, e0229352.		0
16	Utilization of archived neonatal dried blood spots for genome-wide genotyping., 2020, 15, e0229352.		0
17	Inherited genetic susceptibility to acute lymphoblastic leukemia in Down syndrome. Blood, 2019, 134, 1227-1237.	1.4	37
18	Cohabitation is associated with a greater resemblance in gut microbiota which can impact cardiometabolic and inflammatory risk. BMC Microbiology, 2019, 19, 230.	3.3	26

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19	Genetic Similarity Assessment of Twin-Family Populations by Custom-Designed Genotyping Array. Twin Research and Human Genetics, 2019, 22, 210-219.	0.6	11
20	Biological insights into multiple birth: genetic findings from UK Biobank. European Journal of Human Genetics, 2019, 27, 970-979.	2.8	7
21	DNA Methylation Signatures of Breastfeeding in Buccal Cells Collected in Mid-Childhood. Nutrients, 2019, 11, 2804.	4.1	18
22	The Netherlands Twin Register: Longitudinal Research Based on Twin and Twin-Family Designs. Twin Research and Human Genetics, 2019, 22, 623-636.	0.6	112
23	The Nigerian Twin and Sibling Registry: An Update. Twin Research and Human Genetics, 2019, 22, 637-640.	0.6	8
24	Genome-wide analysis of DNA methylation in buccal cells: a study of monozygotic twins and mQTLs. Epigenetics and Chromatin, 2018, 11, 54.	3.9	39
25	Assessing inter-component heterogeneity of biphasic uterine carcinosarcomas. Gynecologic Oncology, 2018, 151, 243-249.	1.4	11
26	Metataxonomic Analysis of Individuals at BMI Extremes and Monozygotic Twins Discordant for BMI. Twin Research and Human Genetics, 2018, 21, 203-213.	0.6	15
27	APOBEC mutation drives early-onset squamous cell carcinomas in recessive dystrophic epidermolysis bullosa. Science Translational Medicine, 2018, 10, .	12.4	91
28	MicroRNAs as biomarkers for psychiatric disorders with a focus on autism spectrum disorder: Current progress in genetic association studies, expression profiling, and translational research. Autism Research, 2017, 10, 1184-1203.	3.8	49
29	Genome-wide identification and differential analysis of translational initiation. Nature Communications, 2017, 8, 1749.	12.8	100
30	A method to customize population-specific arrays for genome-wide association testing. European Journal of Human Genetics, 2017, 25, 267-270.	2.8	29
31	Relative Telomere Repeat Mass in Buccal and Leukocyte-Derived DNA. PLoS ONE, 2017, 12, e0170765.	2.5	22
32	The Genetic Overlap Between Hair and Eye Color. Twin Research and Human Genetics, 2016, 19, 595-599.	0.6	17
33	Genetic and environmental influences interact with age and sex in shaping the human methylome. Nature Communications, 2016, 7, 11115.	12.8	299
34	CNV Concordance in 1,097 MZ Twin Pairs. Twin Research and Human Genetics, 2015, 18, 1-12.	0.6	59
35	Estimation of Genetic Relationships Between Individuals Across Cohorts and Platforms: Application to Childhood Height. Behavior Genetics, 2015, 45, 514-528.	2.1	20
36	Educational Attainment Influences Levels of Homozygosity through Migration and Assortative Mating. PLoS ONE, 2015, 10, e0118935.	2.5	36

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#	Article	lF	CITATION
37	Genetic Basis of a Cognitive Complexity Metric. PLoS ONE, 2015, 10, e0123886.	2.5	22
38	A Genome-wide Association Meta-analysis of Preschool Internalizing Problems. Journal of the American Academy of Child and Adolescent Psychiatry, 2014, 53, 667-676.e7.	0.5	54
39	Metformin and Berberine Prevent Olanzapine-Induced Weight Gain in Rats. PLoS ONE, 2014, 9, e93310.	2.5	57
40	The Young Netherlands Twin Register (YNTR): Longitudinal Twin and Family Studies in Over 70,000 Children. Twin Research and Human Genetics, 2013, 16, 252-267.	0.6	164