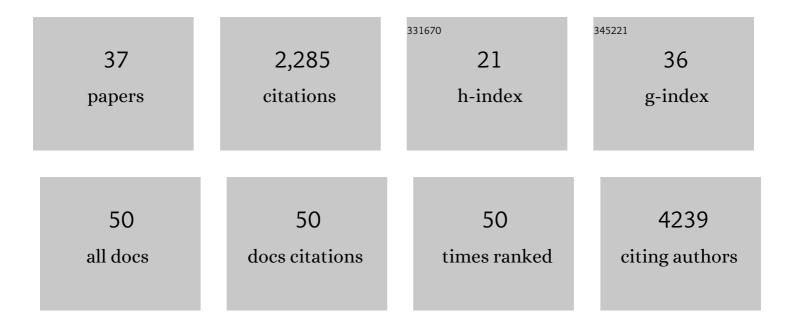
Anna Joanna Jasinska

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3304172/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Enhancer Evolution across 20 Mammalian Species. Cell, 2015, 160, 554-566.	28.9	671
2	The cerebellum ages slowly according to the epigenetic clock. Aging, 2015, 7, 294-306.	3.1	162
3	Systems Biology of the Vervet Monkey. ILAR Journal, 2013, 54, 122-143.	1.8	120
4	The genome of the vervet (<i>Chlorocebus aethiops sabaeus</i>). Genome Research, 2015, 25, 1921-1933.	5.5	114
5	Ancient hybridization and strong adaptation to viruses across African vervet monkey populations. Nature Genetics, 2017, 49, 1705-1713.	21.4	107
6	New applications and developments in the use of multiplex ligationâ€dependent probe amplification. Electrophoresis, 2008, 29, 4627-4636.	2.4	87
7	Factors Associated with Siman Immunodeficiency Virus Transmission in a Natural African Nonhuman Primate Host in the Wild. Journal of Virology, 2014, 88, 5687-5705.	3.4	77
8	Identification of brain transcriptional variation reproduced in peripheral blood: an approach for mapping brain expression traits. Human Molecular Genetics, 2009, 18, 4415-4427.	2.9	72
9	Genetic variation and gene expression across multiple tissues and developmental stages in a nonhuman primate. Nature Genetics, 2017, 49, 1714-1721.	21.4	57
10	A genetic linkage map of the vervet monkey (Chlorocebus aethiops sabaeus). Mammalian Genome, 2007, 18, 347-360.	2.2	55
11	Large-scale meta-analysis of mutations identified in panels of breast/ovarian cancer-related genes — Providing evidence of cancer predisposition genes. Gynecologic Oncology, 2019, 153, 452-462.	1.4	52
12	A quantitative trait locus for variation in dopamine metabolism mapped in a primate model using reference sequences from related species. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 15811-15816.	7.1	51
13	A non-human primate system for large-scale genetic studies of complex traits. Human Molecular Genetics, 2012, 21, 3307-3316.	2.9	51
14	Seroprevalence of Zika Virus in Wild African Green Monkeys and Baboons. MSphere, 2017, 2, .	2.9	50
15	Zoonotic Potential of Simian Arteriviruses. Journal of Virology, 2016, 90, 630-635.	3.4	48
16	Local Virus Extinctions following a Host Population Bottleneck. Journal of Virology, 2015, 89, 8152-8161.	3.4	46
17	Sequencing strategies and characterization of 721 vervet monkey genomes for future genetic analyses of medically relevant traits. BMC Biology, 2015, 13, 41.	3.8	45
18	Neurodegenerative disease biomarkers Aβ _{1–40} , Aβ _{1–42} , tau, and pâ€ŧau ₁₈₁ in the vervet monkey cerebrospinal fluid: RelationÂto normal aging, genetic influences, and cerebral amyloid angiopathy. Brain and Behavior, 2018, 8, e00903.	2.2	45

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19	Transmission of Staphylococcus aureus from Humans to Green Monkeys in The Gambia as Revealed by Whole-Genome Sequencing. Applied and Environmental Microbiology, 2016, 82, 5910-5917.	3.1	30
20	Epigenetic clock and methylation studies in the rhesus macaque. GeroScience, 2021, 43, 2441-2453.	4.6	28
21	Polymorphisms in the <i>GRIA1</i> gene region in psychotic bipolar disorder. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2009, 150B, 24-32.	1.7	27
22	Arteriviruses, Pegiviruses, and Lentiviruses Are Common among Wild African Monkeys. Journal of Virology, 2016, 90, 6724-6737.	3.4	26
23	DNA methylation age analysis of rapamycin in common marmosets. GeroScience, 2021, 43, 2413-2425.	4.6	26
24	ACE2 and TMPRSS2 variation in savanna monkeys (Chlorocebus spp.): Potential risk for zoonotic/anthroponotic transmission of SARS-CoV-2 and a potential model for functional studies. PLoS ONE, 2020, 15, e0235106.	2.5	21
25	CCR5 as a Coreceptor for Human Immunodeficiency Virus and Simian Immunodeficiency Viruses: A Prototypic Love-Hate Affair. Frontiers in Immunology, 2022, 13, 835994.	4.8	20
26	Epigenetic clock and methylation studies in vervet monkeys. GeroScience, 2022, 44, 699-717.	4.6	18
27	Methodological Issues in Molecular Genetic Studies of Mental Disorders. Annual Review of Clinical Psychology, 2009, 5, 49-69.	12.3	12
28	Shifts in microbial diversity, composition, and functionality in the gut and genital microbiome during a natural SIV infection in vervet monkeys. Microbiome, 2020, 8, 154.	11.1	11
29	Resources for functional genomic studies of health and development in nonhuman primates. American Journal of Physical Anthropology, 2020, 171, 174-194.	2.1	7
30	The complex genetic basis of simple behavior. Journal of Biology, 2009, 8, 71.	2.7	5
31	Large Comparative Analyses of Primate Body Site Microbiomes Indicate that the Oral Microbiome Is Unique among All Body Sites and Conserved among Nonhuman Primates. Microbiology Spectrum, 2022, 10, e0164321.	3.0	5
32	Characterization of Expression Quantitative Trait Loci in Pedigrees from Colombia and Costa Rica Ascertained for Bipolar Disorder. PLoS Genetics, 2016, 12, e1006046.	3.5	4
33	Transcriptomic Analysis of Cell-free Fetal RNA in the Amniotic Fluid of Vervet Monkeys (Chlorocebus) Tj ETQq1 1	0.784314	l rgβT /Overlo
34	Biological Resources for Genomic Investigation in the Vervet Monkey (Chlorocebus). , 2019, , 16-28.		3
35	Immunosuppressive effect and global dysregulation of blood transcriptome in response to psychosocial stress in vervet monkeys (Chlorocebus sabaeus). Scientific Reports, 2020, 10, 3459.	3.3	2
36	Expression characteristics of triplet repeat-containing RNAs and triplet repeat-interacting proteins in human tissues. Acta Biochimica Polonica, 2008, 55, 1-8.	0.5	2

#	Article	IF	CITATIONS
37	Expression characteristics of triplet repeat-containing RNAs and triplet repeat-interacting proteins in human tissues Acta Biochimica Polonica, 2008, 55, 1-8.	0.5	1