

# Anna Joanna Jasinska

## List of Publications by Year in descending order

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Version: 2024-02-01

37  
papers

2,285  
citations

331670

21  
h-index

345221

36  
g-index

50  
all docs

50  
docs citations

50  
times ranked

4239  
citing authors

#	ARTICLE	IF	CITATIONS
1	Epigenetic clock and methylation studies in vervet monkeys. <i>GeroScience</i> , 2022, 44, 699-717.	4.6	18
2	CCR5 as a Coreceptor for Human Immunodeficiency Virus and Simian Immunodeficiency Viruses: A Prototypic Love-Hate Affair. <i>Frontiers in Immunology</i> , 2022, 13, 835994.	4.8	20
3	Large Comparative Analyses of Primate Body Site Microbiomes Indicate that the Oral Microbiome Is Unique among All Body Sites and Conserved among Nonhuman Primates. <i>Microbiology Spectrum</i> , 2022, 10, e0164321.	3.0	5
4	Epigenetic clock and methylation studies in the rhesus macaque. <i>GeroScience</i> , 2021, 43, 2441-2453.	4.6	28
5	DNA methylation age analysis of rapamycin in common marmosets. <i>GeroScience</i> , 2021, 43, 2413-2425.	4.6	26
6	Shifts in microbial diversity, composition, and functionality in the gut and genital microbiome during a natural SIV infection in vervet monkeys. <i>Microbiome</i> , 2020, 8, 154.	11.1	11
7	ACE2 and TMPRSS2 variation in savanna monkeys ( <i>Chlorocebus</i> spp.): Potential risk for zoonotic/anthroponotic transmission of SARS-CoV-2 and a potential model for functional studies. <i>PLoS ONE</i> , 2020, 15, e0235106.	2.5	21
8	Immunosuppressive effect and global dysregulation of blood transcriptome in response to psychosocial stress in vervet monkeys ( <i>Chlorocebus sabaeus</i> ). <i>Scientific Reports</i> , 2020, 10, 3459.	3.3	2
9	Resources for functional genomic studies of health and development in nonhuman primates. <i>American Journal of Physical Anthropology</i> , 2020, 171, 174-194.	2.1	7
10	Transcriptomic Analysis of Cell-free Fetal RNA in the Amniotic Fluid of Vervet Monkeys ( <i>Chlorocebus</i> ) Tj ETQq0 0 0 r gBT /Overlock 10 Tf	1.0	4
11	Biological Resources for Genomic Investigation in the Vervet Monkey ( <i>Chlorocebus</i> ). , 2019, , 16-28.		3
12	Large-scale meta-analysis of mutations identified in panels of breast/ovarian cancer-related genes â€” Providing evidence of cancer predisposition genes. <i>Gynecologic Oncology</i> , 2019, 153, 452-462.	1.4	52
13	Neurodegenerative disease biomarkers AÎ²<sub>1-40</sub>, AÎ²<sub>1-42</sub>, tau, and p-tau<sub>181</sub> in the vervet monkey cerebrospinal fluid: Relation to normal aging, genetic influences, and cerebral amyloid angiopathy. <i>Brain and Behavior</i> , 2018, 8, e00903.	2.2	45
14	Seroprevalence of Zika Virus in Wild African Green Monkeys and Baboons. <i>MSphere</i> , 2017, 2, .	2.9	50
15	Ancient hybridization and strong adaptation to viruses across African vervet monkey populations. <i>Nature Genetics</i> , 2017, 49, 1705-1713.	21.4	107
16	Genetic variation and gene expression across multiple tissues and developmental stages in a nonhuman primate. <i>Nature Genetics</i> , 2017, 49, 1714-1721.	21.4	57
17	Characterization of Expression Quantitative Trait Loci in Pedigrees from Colombia and Costa Rica Ascertained for Bipolar Disorder. <i>PLoS Genetics</i> , 2016, 12, e1006046.	3.5	4
18	Arteriviruses, Pegviruses, and Lentiviruses Are Common among Wild African Monkeys. <i>Journal of Virology</i> , 2016, 90, 6724-6737.	3.4	26

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19	Transmission of <i>Staphylococcus aureus</i> from Humans to Green Monkeys in The Gambia as Revealed by Whole-Genome Sequencing. <i>Applied and Environmental Microbiology</i> , 2016, 82, 5910-5917.	3.1	30
20	Zoonotic Potential of Simian Arteriviruses. <i>Journal of Virology</i> , 2016, 90, 630-635.	3.4	48
21	Enhancer Evolution across 20 Mammalian Species. <i>Cell</i> , 2015, 160, 554-566.	28.9	671
22	Local Virus Extinctions following a Host Population Bottleneck. <i>Journal of Virology</i> , 2015, 89, 8152-8161.	3.4	46
23	Sequencing strategies and characterization of 721 vervet monkey genomes for future genetic analyses of medically relevant traits. <i>BMC Biology</i> , 2015, 13, 41.	3.8	45
24	The genome of the vervet ( <i>Chlorocebus aethiops sabaeus</i> ). <i>Genome Research</i> , 2015, 25, 1921-1933.	5.5	114
25	The cerebellum ages slowly according to the epigenetic clock. <i>Aging</i> , 2015, 7, 294-306.	3.1	162
26	Factors Associated with Simian Immunodeficiency Virus Transmission in a Natural African Nonhuman Primate Host in the Wild. <i>Journal of Virology</i> , 2014, 88, 5687-5705.	3.4	77
27	Systems Biology of the Vervet Monkey. <i>ILAR Journal</i> , 2013, 54, 122-143.	1.8	120
28	A non-human primate system for large-scale genetic studies of complex traits. <i>Human Molecular Genetics</i> , 2012, 21, 3307-3316.	2.9	51
29	Identification of brain transcriptional variation reproduced in peripheral blood: an approach for mapping brain expression traits. <i>Human Molecular Genetics</i> , 2009, 18, 4415-4427.	2.9	72
30	Polymorphisms in the <i>GRIA1</i> gene region in psychotic bipolar disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2009, 150B, 24-32.	1.7	27
31	The complex genetic basis of simple behavior. <i>Journal of Biology</i> , 2009, 8, 71.	2.7	5
32	Methodological Issues in Molecular Genetic Studies of Mental Disorders. <i>Annual Review of Clinical Psychology</i> , 2009, 5, 49-69.	12.3	12
33	New applications and developments in the use of multiplex ligation-dependent probe amplification. <i>Electrophoresis</i> , 2008, 29, 4627-4636.	2.4	87
34	Expression characteristics of triplet repeat-containing RNAs and triplet repeat-interacting proteins in human tissues.. <i>Acta Biochimica Polonica</i> , 2008, 55, 1-8.	0.5	1
35	Expression characteristics of triplet repeat-containing RNAs and triplet repeat-interacting proteins in human tissues. <i>Acta Biochimica Polonica</i> , 2008, 55, 1-8.	0.5	2
36	A quantitative trait locus for variation in dopamine metabolism mapped in a primate model using reference sequences from related species. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 15811-15816.	7.1	51

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37	A genetic linkage map of the vervet monkey ( <i>Chlorocebus aethiops sabaeus</i> ). <i>Mammalian Genome</i> , 2007, 18, 347-360.	2.2	55