

# Aashish R Jha

## List of Publications by Year in descending order

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Version: 2024-02-01

27  
papers

5,190  
citations

471509

17  
h-index

580821

25  
g-index

33  
all docs

33  
docs citations

33  
times ranked

9349  
citing authors

#	ARTICLE	IF	CITATIONS
1	Human Gut Microbes Associated with Systolic Blood Pressure. International Journal of Hypertension, 2022, 2022, 1-12.	1.3	1
2	Prevalence and determinants of hypertension in underrepresented indigenous populations of Nepal. PLOS Global Public Health, 2022, 2, e0000133.	1.6	3
3	Robust variation in infant gut microbiome assembly across a spectrum of lifestyles. Science, 2022, 376, 1220-1223.	12.6	63
4	The microbiota of healthy dogs demonstrates individualized responses to synbiotic supplementation in a randomized controlled trial. Animal Microbiome, 2021, 3, 36.	3.8	8
5	Heterogeneity of gut microbial responses in healthy household dogs transitioning from an extruded to a mildly cooked diet. PeerJ, 2021, 9, e11648.	2.0	5
6	Characterization of gut microbiomes of household pets in the United States using a direct-to-consumer approach. PLoS ONE, 2020, 15, e0227289.	2.5	32
7	Gut microbiome transition across a lifestyle gradient in Himalaya. PLoS Biology, 2018, 16, e2005396.	5.6	128
8	Shared Genetic Signals of Hypoxia Adaptation in <i>Drosophila</i> and in High-Altitude Human Populations. Molecular Biology and Evolution, 2016, 33, 501-517.	8.9	44
9	The Simons Genome Diversity Project: 300 genomes from 142 diverse populations. Nature, 2016, 538, 201-206.	27.8	1,216
10	Integrative genomics reveals hypoxia inducible genes that are associated with a poor prognosis in neuroblastoma patients. Oncotarget, 2016, 7, 76816-76826.	1.8	33
11	Whole-Genome Resequencing of Experimental Populations Reveals Polygenic Basis of Egg-Size Variation in <i>Drosophila melanogaster</i> . Molecular Biology and Evolution, 2015, 32, 2616-2632.	8.9	80
12	Global diversity, population stratification, and selection of human copy-number variation. Science, 2015, 349, aab3761.	12.6	293
13	Ancient human genomes suggest three ancestral populations for present-day Europeans. Nature, 2014, 513, 409-413.	27.8	1,179
14	miR-9a Minimizes the Phenotypic Impact of Genomic Diversity by Buffering a Transcription Factor. Cell, 2013, 155, 1556-1567.	28.9	99
15	Tim-3 marks human natural killer cell maturation and suppresses cell-mediated cytotoxicity. Blood, 2012, 119, 3734-3743.	1.4	406
16	Associations between Antibodies to a Panel of Plasmodium falciparum Specific Antigens and Response to Sub-Optimal Antimalarial Therapy in Kampala, Uganda. PLoS ONE, 2012, 7, e52571.	2.5	8
17	An Aboriginal Australian Genome Reveals Separate Human Dispersals into Asia. Science, 2011, 334, 94-98.	12.6	675
18	HTLV-1 Tax Specific CD8+ T Cells Express Low Levels of Tim-3 in HTLV-1 Infection: Implications for Progression to Neurological Complications. PLoS Neglected Tropical Diseases, 2011, 5, e1030.	3.0	29

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19	Human Endogenous Retrovirus K106 (HERV-K106) Was Infectious after the Emergence of Anatomically Modern Humans. PLoS ONE, 2011, 6, e20234.	2.5	46
20	A novel human CD4 <sup>+</sup> T <sub>H</sub> 17 cell inducer subset with potent immunostimulatory properties. European Journal of Immunology, 2010, 40, 134-141.	2.9	14
21	IL-2 Immunotherapy to Recently HIV-1 Infected Adults Maintains the Numbers of IL-17 Expressing CD4 <sup>+</sup> T (TH17) Cells in the Periphery. Journal of Clinical Immunology, 2010, 30, 681-692.	3.8	10
22	Actions speak louder than words to prevent language extinctions. Nature, 2010, 464, 1125-1125.	27.8	0
23	Cross-Sectional Dating of Novel Haplotypes of HERV-K 113 and HERV-K 115 Indicate These Proviruses Originated in Africa before Homo sapiens. Molecular Biology and Evolution, 2009, 26, 2617-2626.	8.9	34
24	Interleukin-10-secreting T cells define a suppressive subset within the HIV-1-specific T <sub>H</sub> 17 cell population. European Journal of Immunology, 2009, 39, 1280-1287.	2.9	18
25	Tim-3 expression defines a novel population of dysfunctional T cells with highly elevated frequencies in progressive HIV-1 infection. Journal of Experimental Medicine, 2008, 205, 2763-2779.	8.5	681
26	Suppression of HIV-1 plasma viral load below detection preserves IL-17 producing T cells in HIV-1 infection. Aids, 2008, 22, 990-992.	2.2	66
27	Tim-3 expression defines a novel population of dysfunctional T cells with highly elevated frequencies in progressive HIV-1 infection. Journal of Cell Biology, 2008, 183, i9-i9.	5.2	0