

# Artika P Nath

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5118676/publications.pdf>

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9  
papers

590  
citations

1163117

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1474206

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g-index

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12  
docs citations

12  
times ranked

1775  
citing authors

#	ARTICLE	IF	CITATIONS
1	The Biomarker GlycA Is Associated with Chronic Inflammation and Predicts Long-Term Risk of Severe Infection. <i>Cell Systems</i> , 2015, 1, 293-301.	6.2	179
2	Frequent transmission of the <i>Mycobacterium tuberculosis</i> Beijing lineage and positive selection for the EsxW Beijing variant in Vietnam. <i>Nature Genetics</i> , 2018, 50, 849-856.	21.4	167
3	Blood-Informative Transcripts Define Nine Common Axes of Peripheral Blood Gene Expression. <i>PLoS Genetics</i> , 2013, 9, e1003362.	3.5	53
4	Comparative analysis reveals a role for TGF- $\beta$ 2 in shaping the residency-related transcriptional signature in tissue-resident memory CD8+ T cells. <i>PLoS ONE</i> , 2019, 14, e0210495.	2.5	49
5	An interaction map of circulating metabolites, immune gene networks, and their genetic regulation. <i>Genome Biology</i> , 2017, 18, 146.	8.8	46
6	Identification of expression quantitative trait loci associated with schizophrenia and affective disorders in normal brain tissue. <i>PLoS Genetics</i> , 2018, 14, e1007607.	3.5	34
7	Multivariate Genome-wide Association Analysis of a Cytokine Network Reveals Variants with Widespread Immune, Haematological, and Cardiometabolic Pleiotropy. <i>American Journal of Human Genetics</i> , 2019, 105, 1076-1090.	6.2	31
8	Elevated serum alpha-1 antitrypsin is a major component of GlycA-associated risk for future morbidity and mortality. <i>PLoS ONE</i> , 2019, 14, e0223692.	2.5	14
9	Geographical, Environmental and Pathophysiological Influences on the Human Blood Transcriptome. <i>Current Genetic Medicine Reports</i> , 2013, 1, 203-211.	1.9	8