

# Taylor B Cavazos

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8874200/publications.pdf>

Version: 2024-02-01

9  
papers

495  
citations

1163117

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1474206

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docs citations

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1707  
citing authors

#	ARTICLE	IF	CITATIONS
1	Pan-cancer study detects genetic risk variants and shared genetic basis in two large cohorts. <i>Nature Communications</i> , 2020, 11, 4423.	12.8	142
2	The landscape of host genetic factors involved in immune response to common viral infections. <i>Genome Medicine</i> , 2020, 12, 93.	8.2	65
3	Inclusion of variants discovered from diverse populations improves polygenic risk score transferability. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100017.	1.7	64
4	Cross-cancer evaluation of polygenic risk scores for 16 cancer types in two large cohorts. <i>Nature Communications</i> , 2021, 12, 970.	12.8	50
5	Identification of Novel Susceptibility Loci and Genes for Prostate Cancer Risk: A Transcriptome-Wide Association Study in Over 140,000 European Descendants. <i>Cancer Research</i> , 2019, 79, 3192-3204.	0.9	43
6	Decomposing Oncogenic Transcriptional Signatures to Generate Maps of Divergent Cellular States. <i>Cell Systems</i> , 2017, 5, 105-118.e9.	6.2	40
7	Age-of-onset information helps identify 76 genetic variants associated with allergic disease. <i>PLoS Genetics</i> , 2020, 16, e1008725.	3.5	27
8	A Large-Scale Association Study Detects Novel Rare Variants, Risk Genes, Functional Elements, and Polygenic Architecture of Prostate Cancer Susceptibility. <i>Cancer Research</i> , 2021, 81, 1695-1703.	0.9	15
9	Nurturing diversity and inclusion in AI in Biomedicine through a virtual summer program for high school students. <i>PLoS Computational Biology</i> , 2022, 18, e1009719.	3.2	3