

Travis Hughes

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

1,034
citations

759233

12
h-index

1058476

14
g-index

16
all docs

16
docs citations

16
times ranked

1979
citing authors

#	ARTICLE	IF	CITATIONS
1	JAK inhibition in a patient with a STAT1 gain-of-function variant reveals STAT1 dysregulation as a common feature of aplastic anemia. <i>Med</i> , 2022, 3, 42-57.e5.	4.4	11
2	Hypoxic, glycolytic metabolism is a vulnerability of B-acute lymphoblastic leukemia-initiating cells. <i>Cell Reports</i> , 2022, 39, 110752.	6.4	5
3	973. Single-cell RNA Sequencing Analysis of Zika Virus Infection in Human Stem Cell-Derived Cerebral Organoids. <i>Open Forum Infectious Diseases</i> , 2019, 6, S33-S34.	0.9	0
4	Genetic heterogeneity within the HLA region in three distinct clinical subgroups of myasthenia gravis. <i>Clinical Immunology</i> , 2016, 166-167, 81-88.	3.2	38
5	Epigenome-Wide Scan Identifies a Treatment-Responsive Pattern of Altered DNA Methylation Among Cytoskeletal Remodeling Genes in Monocytes and CD4+ T Cells From Patients With Behçet's Disease. <i>Arthritis and Rheumatology</i> , 2014, 66, 1648-1658.	5.6	72
6	Genome-Wide DNA Methylation Patterns in Naive CD4+ T Cells From Patients With Primary Sjögren's Syndrome. <i>Arthritis and Rheumatology</i> , 2014, 66, 731-739.	5.6	147
7	Identification of Multiple Genetic Susceptibility Loci in Takayasu Arteritis. <i>American Journal of Human Genetics</i> , 2013, 93, 298-305.	6.2	143
8	Identification of multiple independent susceptibility loci in the HLA region in Behçet's disease. <i>Nature Genetics</i> , 2013, 45, 319-324.	21.4	130
9	Analysis of autosomal genes reveals gene-sex interactions and higher total genetic risk in men with systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 694-699.	0.9	87
10	Sex-specific differences in the relationship between genetic susceptibility, T cell DNA demethylation and lupus flare severity. <i>Journal of Autoimmunity</i> , 2012, 38, J216-J222.	6.5	71
11	Evidence for gene-gene epistatic interactions among susceptibility loci for systemic lupus erythematosus. <i>Arthritis and Rheumatism</i> , 2012, 64, 485-492.	6.7	53
12	The role of epigenetic variation in the pathogenesis of systemic lupus erythematosus. <i>Arthritis Research and Therapy</i> , 2011, 13, 245.	3.5	34
13	Fine-mapping and transethnic genotyping establish IL2/IL21 genetic association with lupus and localize this genetic effect to IL21. <i>Arthritis and Rheumatism</i> , 2011, 63, 1689-1697.	6.7	49
14	A putative functional variant within the <i>UBAC2</i> gene is associated with increased risk of Behçet's disease. <i>Arthritis and Rheumatism</i> , 2011, 63, 3607-3612.	6.7	39
15	Early disease onset is predicted by a higher genetic risk for lupus and is associated with a more severe phenotype in lupus patients. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 151-156.	0.9	155