Travis Hughes

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11776425/publications.pdf

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		759233	1058476	
15	1,034 citations	12	14	
papers	citations	h-index	g-index	
16	16	16	1979	
all docs	docs citations	times ranked	citing authors	

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