Pascal F Pucholt

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/8780544/publications.pdf

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24	479	759233 12	752698 20
papers	citations	h-index	g-index

26	26	26	750
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Contribution of Rare Genetic Variation to Disease Susceptibility in a Large Scandinavian Myositis Cohort. Arthritis and Rheumatology, 2022, 74, 342-352.	5.6	7
2	Complement C4 Copy Number Variation is Linked to SSA/Ro and SSB/La Autoantibodies in Systemic Inflammatory Autoimmune Diseases. Arthritis and Rheumatology, 2022, 74, 1440-1450.	5.6	17
3	Toll-like receptors revisited; a possible role for TLR1 in lupus nephritis. Annals of the Rheumatic Diseases, 2021, 80, 404-406.	0.9	7
4	Molecular pathways in patients with systemic lupus erythematosus revealed by gene-centred DNA sequencing. Annals of the Rheumatic Diseases, 2021, 80, 109-117.	0.9	35
5	Interaction between the STAT4 rs11889341(T) risk allele and smoking confers increased risk of myocardial infarction and nephritis in patients with systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2021, 80, 1183-1189.	0.9	10
6	Variants in BANK1 are associated with lupus nephritis of European ancestry. Genes and Immunity, 2021, 22, 194-202.	4.1	9
7	OUP accepted manuscript. Rheumatology, 2021, 60, 837-848.	1.9	15
8	Genome-wide association mapping uncovers sex-associated copy number variation markers and female hemizygous regions on the W chromosome in Salix viminalis. BMC Genomics, 2021, 22, 710.	2.8	6
9	P96â€…The regulation and pharmacological modulation of immune complex induced production of type III IFN by plasmacytoid dendritic cells. , 2020, , .		0
10	O23â€…Identification of protein-quantitative trait loci (pQTLs) in the interferon signalling pathway. , 2020, , .		0
11	The regulation and pharmacological modulation of immune complex induced type III IFN production by plasmacytoid dendritic cells. Arthritis Research and Therapy, 2020, 22, 130.	3.5	14
12	miRNA target identification and prediction as a function of time in gene expression data. RNA Biology, 2020, 17, 990-1000.	3.1	2
13	Genome assembly of the basket willow, Salix viminalis, reveals earliest stages of sex chromosome expansion. BMC Biology, 2020, 18, 78.	3.8	39
14	Activation of plasmacytoid dendritic cells and B cells with two structurally different Tollâ€‐like receptor 7 agonists. Scandinavian Journal of Immunology, 2020, 91, e12880.	2.7	5
15	Function of multiple sclerosis-protective HLA class I alleles revealed by genome-wide protein-quantitative trait loci mapping of interferon signalling. PLoS Genetics, 2020, 16, e1009199.	3.5	12
16	Circulating Levels of Interferon Regulatory Factor-5 Associates With Subgroups of Systemic Lupus Erythematosus Patients. Frontiers in Immunology, 2019, 10, 1029.	4.8	11
17	Gains of Chromosome 1p and 15q are Associated with Poor Survival After Cytoreductive Surgery and HIPEC for Treating Colorectal Peritoneal Metastases. Annals of Surgical Oncology, 2019, 26, 4835-4842.	1.5	5
18	Slow evolution of sexâ€‐biased genes in the reproductive tissue of the dioecious plant Salix viminalis. Molecular Ecology, 2018, 27, 694-708.	3.9	37

#	Article	IF	Citations
19	Recent Sex Chromosome Divergence despite Ancient Dioecy in the Willow Salix viminalis. Molecular Biology and Evolution, 2017, 34, 1991-2001.	8.9	57
20	Allelic incompatibility can explain female biased sex ratios in dioecious plants. BMC Genomics, 2017, 18, 251.	2.8	28
21	Sequence and gene expression evolution of paralogous genes in willows. Scientific Reports, 2016, 5, 18662.	3.3	16
22	Genome-wide transcriptional and physiological responses to drought stress in leaves and roots of two willow genotypes. BMC Plant Biology, 2015, 15, 244.	3.6	37
23	Genetic and morphological evidence for introgression between three species of willows. BMC Evolutionary Biology, 2015, 15, 193.	3.2	29
24	Single locus sex determination and female heterogamety in the basket willow (Salix viminalis L.). Heredity, 2015, 114, 575-583.	2.6	76