Deborah R Winter

List of Publications by Year in descending order

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42 papers

27,186 citations

201674 27 h-index 276875 41 g-index

53 all docs

53 docs citations

53 times ranked 53469 citing authors

#	Article	IF	CITATIONS
1	Stromal Transcription Factor 21 Regulates Development of the Renal Stroma via Interaction with Wnt $\hat{\mathbb{C}}^2$ -Catenin Signaling. Kidney360, 2022, 3, 1228-1241.	2.1	5
2	Activation of the 15-lipoxygenase pathway in aspirin-exacerbated respiratory disease. Journal of Allergy and Clinical Immunology, 2021, 147, 600-612.	2.9	43
3	Critical role of synovial tissue–resident macrophage niche in joint homeostasis and suppression of chronic inflammation. Science Advances, 2021, 7, .	10.3	27
4	Transcriptional profiling of pediatric cholestatic livers identifies three distinct macrophage populations. PLoS ONE, 2021, 16, e0244743.	2.5	20
5	The lung microenvironment shapes a dysfunctional response of alveolar macrophages in aging. Journal of Clinical Investigation, 2021, 131, .	8.2	86
6	Neuroimmune interactions and osteoarthritis pain: focus on macrophages. Pain Reports, 2021, 6, e892.	2.7	26
7	Microglia Adopt Longitudinal Transcriptional Changes After Traumatic Brain Injury. Journal of Surgical Research, 2020, 246, 113-122.	1.6	18
8	A Genome-Based Model to Predict the Virulence of Pseudomonas aeruginosa Isolates. MBio, 2020, 11 , .	4.1	12
9	A Novel Microglia-Specific Transcriptional Signature Correlates With Behavioral Deficits in Neuropsychiatric Lupus. Frontiers in Immunology, 2020, 11, 230.	4.8	27
10	A Novel Role for 15-Lipoxygenase Metabolites in Aspirin Exacerbated Respiratory Disease. Journal of Allergy and Clinical Immunology, 2020, 145, AB242.	2.9	2
11	SATO106â€NOVEL SUBCLASS OF INTRAVASCULAR NON-CLASSICAL SYNOVIAL MONOCYTES ARE CRITICAL FOR RHEUMATOID ARTHRITIS., 2019,,.	2	0
12	OP0294â€TRANSCRIPTIONAL PROFILING OF RA PATIENTS SYNOVIAL TISSUE REVEALS TARGETS FOR PRECISION MEDICINE. , 2019, , .	1	0
13	Lipocalin-2 is a pathogenic determinant and biomarker of neuropsychiatric lupus. Journal of Autoimmunity, 2019, 96, 59-73.	6.5	43
14	Single-Cell Transcriptomic Analysis of Human Lung Provides Insights into the Pathobiology of Pulmonary Fibrosis. American Journal of Respiratory and Critical Care Medicine, 2019, 199, 1517-1536.	5.6	866
15	A Beginner's Guide to Analysis of RNA Sequencing Data. American Journal of Respiratory Cell and Molecular Biology, 2018, 59, 145-157.	2.9	78
16	Transcriptional Profiling of Synovial Macrophages Using Minimally Invasive Ultrasoundâ€Guided Synovial Biopsies in Rheumatoid Arthritis. Arthritis and Rheumatology, 2018, 70, 841-854.	5.6	44
17	Neuropsychiatric Systemic Lupus Erythematosus Is Dependent on Sphingosine-1-Phosphate Signaling. Frontiers in Immunology, 2018, 9, 2189.	4.8	44
18	Thinking BIG rheumatology: how to make functional genomics data work for you. Arthritis Research and Therapy, 2018, 20, 29.	3.5	4

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19	Mef2C restrains microglial inflammatory response and is lost in brain ageing inÂan IFN-l-dependent manner. Nature Communications, 2017, 8, 717.	12.8	157
20	Bim suppresses the development of SLE by limiting myeloid inflammatory responses. Journal of Experimental Medicine, 2017, 214, 3753-3773.	8.5	27
21	Disease Specific Signatures Identified by RNAâ€seq of Sorted Lung Cellular Populations. FASEB Journal, 2017, 31, 656.4.	0.5	0
22	Microbiota Diurnal Rhythmicity Programs Host Transcriptome Oscillations. Cell, 2016, 167, 1495-1510.e12.	28.9	591
23	Extracellular Matrix Proteolysis by MT1-MMP Contributes to Influenza-Related Tissue Damage and Mortality. Cell Host and Microbe, 2016, 20, 458-470.	11.0	82
24	The Spectrum and Regulatory Landscape of Intestinal Innate Lymphoid Cells Are Shaped by the Microbiome. Cell, 2016, 166, 1231-1246.e13.	28.9	465
25	Co-ChIP enables genome-wide mapping of histone mark co-occurrence at single-molecule resolution. Nature Biotechnology, 2016, 34, 953-961.	17.5	81
26	Microglia development follows a stepwise program to regulate brain homeostasis. Science, 2016, 353, aad8670.	12.6	911
27	The role of the local environment and epigenetics in shaping macrophage identity and their effect on tissue homeostasis. Nature Immunology, 2016, 17, 18-25.	14.5	315
28	Distinct Murine Mucosal Langerhans Cell Subsets Develop from Pre-dendritic Cells and Monocytes. Immunity, 2015, 43, 369-381.	14.3	78
29	DCs are ready to commit. Nature Immunology, 2015, 16, 683-685.	14.5	4
30	Making the case for chromatin profiling: a new tool to investigate the immune-regulatory landscape. Nature Reviews Immunology, 2015, 15, 585-594.	22.7	32
31	From mass cytometry to cancer prognosis. Nature Biotechnology, 2015, 33, 931-932.	17.5	4
32	Transcriptional Heterogeneity and Lineage Commitment in Myeloid Progenitors. Cell, 2015, 163, 1663-1677.	28.9	875
33	Tissue-Resident Macrophage Enhancer Landscapes Are Shaped by the Local Microenvironment. Cell, 2014, 159, 1312-1326.	28.9	1,705
34	A Negative Feedback Loop of Transcription Factors Specifies Alternative Dendritic Cell Chromatin States. Molecular Cell, 2014, 56, 749-762.	9.7	58
35	Chromatin state dynamics during blood formation. Science, 2014, 345, 943-949.	12.6	699
36	The role of chromatin dynamics in immune cell development. Immunological Reviews, 2014, 261, 9-22.	6.0	57

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37	DNase-seq predicts regions of rotational nucleosome stability across diverse human cell types. Genome Research, 2013, 23, 1118-1129.	5.5	22
38	An integrated encyclopedia of DNA elements in the human genome. Nature, 2012, 489, 57-74.	27.8	15,516
39	A User's Guide to the Encyclopedia of DNA Elements (ENCODE). PLoS Biology, 2011, 9, e1001046.	5.6	1,257
40	Open chromatin defined by DNasel and FAIRE identifies regulatory elements that shape cell-type identity. Genome Research, 2011, 21, 1757-1767.	5.5	449
41	Transcription Initiation Patterns Indicate Divergent Strategies for Gene Regulation at the Chromatin Level. PLoS Genetics, 2011, 7, e1001274.	3.5	124
42	An "Electronic Fluorescent Pictograph―Browser for Exploring and Analyzing Large-Scale Biological Data Sets. PLoS ONE, 2007, 2, e718.	2.5	2,236