Bence Daniel

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Transient rest restores functionality in exhausted CAR-T cells through epigenetic remodeling. Science, 2021, 372, .	12.6	297
2	Human B Cell Clonal Expansion and Convergent Antibody Responses to SARS-CoV-2. Cell Host and Microbe, 2020, 28, 516-525.e5.	11.0	219
3	The Transcription Factor STAT6 Mediates Direct Repression of Inflammatory Enhancers and Limits Activation of Alternatively Polarized Macrophages. Immunity, 2018, 48, 75-90.e6.	14.3	185
4	Macrophage PPARÎ ³ , a Lipid Activated Transcription Factor Controls the Growth Factor GDF3 and Skeletal Muscle Regeneration. Immunity, 2016, 45, 1038-1051.	14.3	134
5	The Nuclear Receptor PPARÎ ³ Controls Progressive Macrophage Polarization as a Ligand-Insensitive Epigenomic Ratchet of Transcriptional Memory. Immunity, 2018, 49, 615-626.e6.	14.3	128
6	Genome-wide CRISPR screens of TÂcell exhaustion identify chromatin remodeling factors that limit TÂcell persistence. Cancer Cell, 2022, 40, 768-786.e7.	16.8	104
7	Hepatocyte-Macrophage Acetoacetate Shuttle Protects against Tissue Fibrosis. Cell Metabolism, 2019, 29, 383-398.e7.	16.2	87
8	The active enhancer network operated by liganded RXR supports angiogenic activity in macrophages. Genes and Development, 2014, 28, 1562-1577.	5.9	85
9	Epigenetic regulation of T cell exhaustion. Nature Immunology, 2022, 23, 848-860.	14.5	82
10	The transcription factor EGR2 is the molecular linchpin connecting STAT6 activation to the late, stable epigenomic program of alternative macrophage polarization. Genes and Development, 2020, 34, 1474-1492.	5.9	38
11	A growth factor–expressing macrophage subpopulation orchestrates regenerative inflammation via GDF-15. Journal of Experimental Medicine, 2022, 219, .	8.5	31
12	BCL6-dependent TCF-1+ progenitor cells maintain effector and helper CD4+ TÂcell responses to persistent antigen. Immunity, 2022, 55, 1200-1215.e6.	14.3	30
13	Retinoid X receptor suppresses a metastasis-promoting transcriptional program in myeloid cells via a ligand-insensitive mechanism. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 10725-10730.	7.1	24
14	The intriguing complexities of mammalian gene regulation: How to link enhancers to regulated genes. Are we there yet?. FEBS Letters, 2014, 588, 2379-2391.	2.8	21
15	Mapping the Genomic Binding Sites of the Activated Retinoid X Receptor in Murine Bone Marrow-Derived Macrophages Using Chromatin Immunoprecipitation Sequencing. Methods in Molecular Biology, 2014, 1204, 15-24.	0.9	18
16	Identification of a T-bethi Quiescent Exhausted CD8 T Cell Subpopulation That Can Differentiate into TIM3+CX3CR1+ Effectors and Memory-like Cells. Journal of Immunology, 2021, 206, 2924-2936.	0.8	17
17	Labelled regulatory elements are pervasive features of the macrophage genome and are dynamically utilized by classical and alternative polarization signals. Nucleic Acids Research, 2019, 47, 2778-2792.	14.5	14
18	Transcriptional control of transglutaminase 2 expression in mouse apoptotic thymocytes. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2016, 1859, 964-974.	1.9	10

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19	Unraveling the Hierarchy of <i>cis</i> and <i>trans</i> Factors That Determine the DNA Binding by Peroxisome Proliferator-Activated Receptor <i>γ</i> . Molecular and Cellular Biology, 2020, 40, .	2.3	5
20	Global Run-on Sequencing (GRO-Seq). Methods in Molecular Biology, 2021, 2351, 25-39.	0.9	3
21	Charting a shared epigenetic pathway to CD8+ T cell dysfunction in infection and cancer. Molecular Cell, 2021, 81, 2272-2274.	9.7	0