

Richard A Mcindoe

List of Publications by Year in descending order

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Version: 2024-02-01

39
papers

2,539
citations

393982

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329751

37
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41
all docs

41
docs citations

41
times ranked

3861
citing authors

#	ARTICLE	IF	CITATIONS
1	A novel network based linear model for prioritization of synergistic drug combinations. PLoS ONE, 2022, 17, e0266382.	1.1	7
2	dkNET Hypothesis Center: A Hub for FAIR Data, Online Resources and Hypothesis Generation. FASEB Journal, 2022, 36, .	0.2	0
3	Persistent STAT5 activation reprograms the epigenetic landscape in CD4 ⁺ T cells to drive polyfunctionality and antitumor immunity. Science Immunology, 2020, 5, .	5.6	40
4	Frequent HPV-independent p16/INK4A overexpression in head and neck cancer. Oral Oncology, 2018, 83, 32-37.	0.8	39
5	The Geropathology Research Network: An Interdisciplinary Approach for Integrating Pathology Into Research on Aging. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2016, 71, 431-434.	1.7	16
6	The same self-peptide selects conventional and regulatory CD4 ⁺ T cells with identical antigen receptors. Nature Communications, 2014, 5, 5061.	5.8	16
7	Thymus-derived regulatory T cells contribute to tolerance to commensal microbiota. Nature, 2013, 497, 258-262.	13.7	333
8	Research Resource: dkCOIN, the National Institute of Diabetes, Digestive and Kidney Diseases (NIDDK) Consortium Interconnectivity Network: A Pilot Program to Aggregate Research Resources Generated by Multiple Research Consortia. Molecular Endocrinology, 2012, 26, 1675-1681.	3.7	3
9	Gene Expression Profiling of Early-Phase Sjögren's Syndrome in C57BL/6.NOD-Aec1Aec2 Mice Identifies Focal Adhesion Maturation Associated with Infiltrating Leukocytes. , 2011, 52, 5647.		20
10	Hepatic Gene Expression Profiling Reveals Key Pathways Involved in Leptin-Mediated Weight Loss in ob/ob Mice. PLoS ONE, 2010, 5, e12147.	1.1	21
11	ParaSAM: a parallelized version of the significance analysis of microarrays algorithm. Bioinformatics, 2010, 26, 1465-1467.	1.8	1
12	A modified hyperplane clustering algorithm allows for efficient and accurate clustering of extremely large datasets. Bioinformatics, 2009, 25, 1152-1157.	1.8	18
13	Mouse Models of Diabetic Nephropathy. Journal of the American Society of Nephrology: JASN, 2009, 20, 2503-2512.	3.0	582
14	Differential gene expressions in the lacrimal gland during development and onset of keratoconjunctivitis sicca in Sjögren's syndrome (SJS)-like disease of the C57BL/6.NOD-Aec1Aec2 mouse. Experimental Eye Research, 2009, 88, 398-409.	1.2	35
15	Differential gene expression in the salivary gland during development and onset of xerostomia in Sjögren's syndrome-like disease of the C57BL/6.NOD-Aec1Aec2 mouse. Arthritis Research and Therapy, 2009, 11, R56.	1.6	46
16	ParaKMeans: Implementation of a parallelized K-means algorithm suitable for general laboratory use. BMC Bioinformatics, 2008, 9, 200.	1.2	19
17	The IL-10 and IFN- γ pathways are essential to the potent immunosuppressive activity of cultured CD8 ⁺ NKT-like cells. Genome Biology, 2008, 9, R119.	13.9	11
18	Recipes for Creating Animal Models of Diabetic Cardiovascular Disease. Circulation Research, 2007, 100, 1415-1427.	2.0	206

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19	Early pathogenic events associated with Sjögren's syndrome (SjS)-like disease of the nod mouse using microarray analysis. <i>Laboratory Investigation</i> , 2006, 86, 1243-1260.	1.7	81
20	Intergeneration CAG expansion and contraction in a Chinese HD family. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2006, 141B, 242-244.	1.1	9
21	Alterations of renal phenotype and gene expression profiles due to protein overload in NOD-related mouse strains. <i>BMC Nephrology</i> , 2005, 6, 17.	0.8	7
22	caBIONet--A .NET wrapper to access and process genomic data stored at the National Cancer Institute's Center for Bioinformatics databases. <i>Bioinformatics</i> , 2005, 21, 3456-3458.	1.8	6
23	Gene Expression Profiles Define a Key Checkpoint for Type 1 Diabetes in NOD Mice. <i>Diabetes</i> , 2004, 53, 366-375.	0.3	29
24	Molecular Pathways Altered by Insulin B9-23 Immunization. <i>Annals of the New York Academy of Sciences</i> , 2004, 1037, 175-185.	1.8	6
25	Gene expression associated with interferon alfa antiviral activity in an HCV replicon cell line. <i>Hepatology</i> , 2003, 37, 1180-1188.	3.6	96
26	Gene Expression Profiling during All-trans Retinoic Acid-Induced Cell Differentiation of Acute Promyelocytic Leukemia Cells. <i>Journal of Molecular Diagnostics</i> , 2003, 5, 212-221.	1.2	40
27	Microarray Analysis of Gene Expression in the Kidneys of New- and Post-Onset Diabetic NOD Mice. <i>Diabetes</i> , 2003, 52, 2151-2159.	0.3	59
28	MADGE: scalable distributed data management software for cDNA microarrays. <i>Bioinformatics</i> , 2003, 19, 87-89.	1.8	3
29	Rapid decrease of RNA level of a novel mouse mitochondria solute carrier protein (Mscp) gene at 4-5 weeks of age. <i>Mammalian Genome</i> , 2001, 12, 830-836.	1.0	4
30	Linkage analysis of 150 high-risk prostate cancer families at 1q24-25. , 2000, 18, 251-275.		43
31	WECH. <i>Immunity</i> , 2000, 12, 141-150.	6.6	35
32	Confirmation of Prostate Cancer Susceptibility Genes Using High-Risk Families. <i>Journal of the National Cancer Institute Monographs</i> , 1999, 1999, 81-87.	0.9	9
33	Evidence for a Rare Prostate Cancer Susceptibility Locus at Chromosome 1p36. <i>American Journal of Human Genetics</i> , 1999, 64, 776-787.	2.6	292
34	Linkage Analysis of 49 High-Risk Families Does Not Support a Common Familial Prostate Cancer Susceptibility Gene at 1q24-25. <i>American Journal of Human Genetics</i> , 1997, 61, 347-353.	2.6	114
35	Data-adaptive algorithms for calling alleles in repeat polymorphisms. <i>Electrophoresis</i> , 1997, 18, 1-5.	1.3	9
36	Detection of a Large RIII-Derived Chromosomal Segment on Chromosome 10 in the H-2 Congenic Strain B10.RIII(71NS)/Sn. <i>Genomics</i> , 1996, 31, 266-269.	1.3	9

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37	An analysis of the dynamic range and linearity of an infrared DNA sequencer. <i>Electrophoresis</i> , 1996, 17, 652-658.	1.3	13
38	Ancestral polymorphisms of MHC class II genes: Divergent allele advantage. <i>Immunologic Research</i> , 1990, 9, 115-122.	1.3	125
39	The origin of MHC class II gene polymorphism within the genus <i>Mus</i> . <i>Nature</i> , 1988, 332, 651-654.	13.7	130