

# Veronique Adoue

## List of Publications by Year in descending order

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22  
papers

1,200  
citations

471509

17  
h-index

677142

22  
g-index

24  
all docs

24  
docs citations

24  
times ranked

3870  
citing authors

#	ARTICLE	IF	CITATIONS
1	The Histone Methyltransferase SETDB1 Controls T <sup>H</sup> Helper Cell Lineage Integrity by Repressing Endogenous Retroviruses. <i>Immunity</i> , 2019, 50, 629-644.e8.	14.3	63
2	Critical role for TRIM28 and HP1 <sup>2/3</sup> in the epigenetic control of T cell metabolic reprogramming and effector differentiation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 25839-25849.	7.1	23
3	Limited Foxp3 <sup>+</sup> Regulatory T Cells Response During Acute <i>Trypanosoma cruzi</i> Infection Is Required to Allow the Emergence of Robust Parasite-Specific CD8 <sup>+</sup> T Cell Immunity. <i>Frontiers in Immunology</i> , 2018, 9, 2555.	4.8	21
4	Association of breast cancer risk with genetic variants showing differential allelic expression: Identification of a novel breast cancer susceptibility locus at 4q21. <i>Oncotarget</i> , 2016, 7, 80140-80163.	1.8	31
5	Peripheral regulatory T lymphocytes recirculating to the thymus suppress the development of their precursors. <i>Nature Immunology</i> , 2015, 16, 628-634.	14.5	144
6	Interrogation of allelic chromatin states in human cells by high-density ChIP-genotyping. <i>Epigenetics</i> , 2014, 9, 1238-1251.	2.7	9
7	Fusion of TTYH1 with the C19MC microRNA cluster drives expression of a brain-specific DNMT3B isoform in the embryonal brain tumor ETMR. <i>Nature Genetics</i> , 2014, 46, 39-44.	21.4	167
8	Allelic expression mapping across cellular lineages to establish impact of non-coding <i>scnp&gt;SNP&lt;/scnp&gt;</i> s. <i>Molecular Systems Biology</i> , 2014, 10, 754.	7.2	21
9	Global identification of conserved post-transcriptional regulatory programs in trypanosomatids. <i>Nucleic Acids Research</i> , 2013, 41, 8591-8600.	14.5	28
10	Analysis of expressed SNPs identifies variable extents of expression from the human inactive X chromosome. <i>Genome Biology</i> , 2013, 14, R122.	9.6	174
11	Promoter polymorphisms in CHI3L1 are associated with asthma. <i>Journal of Allergy and Clinical Immunology</i> , 2012, 130, 533-535.	2.9	10
12	RNA editing of protein sequences: A rare event in human transcriptomes. <i>Rna</i> , 2012, 18, 1586-1596.	3.5	42
13	Deimination and expression of peptidylarginine deiminases during cutaneous wound healing in mice. <i>European Journal of Dermatology</i> , 2011, 21, 376-384.	0.6	25
14	Global Analysis of the Impact of Environmental Perturbation on cis-Regulation of Gene Expression. <i>PLoS Genetics</i> , 2011, 7, e1001279.	3.5	81
15	Identification, Replication, and Fine-Mapping of Loci Associated with Adult Height in Individuals of African Ancestry. <i>PLoS Genetics</i> , 2011, 7, e1002298.	3.5	93
16	Deimination is regulated at multiple levels including auto-deimination of peptidylarginine deiminases. <i>Cellular and Molecular Life Sciences</i> , 2010, 67, 1491-1503.	5.4	41
17	Gene Regulation at a Distance in the Epidermal Keratinocyte: The Paradigm of the PADI Gene Locus. <i>Open Dermatology Journal</i> , 2010, 4, 21-26.	0.3	0
18	Transcriptional regulation of peptidylarginine deiminase expression in human keratinocytes. <i>Journal of Dermatological Science</i> , 2009, 53, 2-9.	1.9	43

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19	Crucial Roles of MZF1 and Sp1 in the Transcriptional Regulation of the Peptidylarginine Deiminase Type I Gene (PADI1) in Human Keratinocytes. <i>Journal of Investigative Dermatology</i> , 2008, 128, 549-557.	0.7	33
20	Long-Range Enhancer Differentially Regulated by c-Jun and JunD Controls Peptidylarginine Deiminase-3 Gene in Keratinocytes. <i>Journal of Molecular Biology</i> , 2008, 384, 1048-1057.	4.2	24
21	Long-Range Enhancer Associated with Chromatin Looping Allows AP-1 Regulation of the Peptidylarginine Deiminase 3 Gene in Differentiated Keratinocyte. <i>PLoS ONE</i> , 2008, 3, e3408.	2.5	47
22	Peptidylarginine deiminases and deimination in biology and pathology: Relevance to skin homeostasis. <i>Journal of Dermatological Science</i> , 2006, 44, 63-72.	1.9	74