

Cristina Rada

List of Publications by Year in descending order

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39
papers

4,095
citations

201674

27
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315739

38
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41
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docs citations

41
times ranked

3794
citing authors

#	ARTICLE	IF	CITATIONS
1	Immunoglobulin Isotype Switching Is Inhibited and Somatic Hypermutation Perturbed in UNG-Deficient Mice. <i>Current Biology</i> , 2002, 12, 1748-1755.	3.9	648
2	Mismatch Recognition and Uracil Excision Provide Complementary Paths to Both Ig Switching and the A/T-Focused Phase of Somatic Mutation. <i>Molecular Cell</i> , 2004, 16, 163-171.	9.7	428
3	Hot Spot Focusing of Somatic Hypermutation in MSH2-Deficient Mice Suggests Two Stages of Mutational Targeting. <i>Immunity</i> , 1998, 9, 135-141.	14.3	354
4	DNA deaminases induce break-associated mutation showers with implication of APOBEC3B and 3A in breast cancer kataegis. <i>ELife</i> , 2013, 2, e00534.	6.0	322
5	Comparison of the Differential Context-dependence of DNA Deamination by APOBEC Enzymes: Correlation with Mutation Spectra in Vivo. <i>Journal of Molecular Biology</i> , 2004, 337, 585-596.	4.2	306
6	The topography of mutational processes in breast cancer genomes. <i>Nature Communications</i> , 2016, 7, 11383.	12.8	235
7	The in vivo pattern of AID targeting to immunoglobulin switch regions deduced from mutation spectra in <i>msh2^{-/-}ung^{-/-}</i> mice. <i>Journal of Experimental Medicine</i> , 2006, 203, 2085-2094.	8.5	162
8	Somatic hypermutation at A-T pairs: polymerase error versus dUTP incorporation. <i>Nature Reviews Immunology</i> , 2005, 5, 171-178.	22.7	132
9	Interaction between Antibody-Diversification Enzyme AID and Spliceosome-Associated Factor CTNNB1. <i>Molecular Cell</i> , 2008, 31, 474-484.	9.7	127
10	AID-GFP chimeric protein increases hypermutation of Ig genes with no evidence of nuclear localization. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 7003-7008.	7.1	119
11	Monitoring and interpreting the intrinsic features of somatic hypermutation. <i>Immunological Reviews</i> , 1998, 162, 107-116.	6.0	117
12	Active demethylation in mouse zygotes involves cytosine deamination and base excision repair. <i>Epigenetics and Chromatin</i> , 2013, 6, 39.	3.9	98
13	The 5' boundary of somatic hypermutation in a <i>VH</i> gene is in the leader intron. <i>European Journal of Immunology</i> , 1994, 24, 1453-1457.	2.9	91
14	SMUG1 is able to excise uracil from immunoglobulin genes: insight into mutation versus repair. <i>EMBO Journal</i> , 2006, 25, 585-595.	7.8	90
15	Altering the spectrum of immunoglobulin V gene somatic hypermutation by modifying the active site of AID. <i>Journal of Experimental Medicine</i> , 2010, 207, 141-153.	8.5	90
16	Germline ablation of SMUG1 DNA glycosylase causes loss of 5-hydroxymethyluracil- and UNG-backup uracil-excision activities and increases cancer predisposition of <i>Ung^{-/-}Msh2^{-/-}</i> mice. <i>Nucleic Acids Research</i> , 2012, 40, 6016-6025.	14.5	89
17	The stability of AID and its function in class-switching are critically sensitive to the identity of its nuclear-export sequence. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 6736-6741.	7.1	77
18	AID upmutants isolated using a high-throughput screen highlight the immunity/cancer balance limiting DNA deaminase activity. <i>Nature Structural and Molecular Biology</i> , 2009, 16, 769-776.	8.2	72

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19	Somatic hypermutation: activation-induced deaminase for C/G followed by polymerase \hat{I} for A/T. <i>Journal of Experimental Medicine</i> , 2007, 204, 7-10.	8.5	63
20	Active RNAP pre-initiation sites are highly mutated by cytidine deaminases in yeast, with AID targeting small RNA genes. <i>ELife</i> , 2014, 3, e03553.	6.0	51
21	Cytoplasmic activation-induced cytidine deaminase (AID) exists in stoichiometric complex with translation elongation factor 1 \hat{I} (eEF1A). <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 18366-18371.	7.1	49
22	Intrinsic transcriptional heterogeneity in B cells controls early class switching to IgE. <i>Journal of Experimental Medicine</i> , 2017, 214, 183-196.	8.5	49
23	The 5 \hat{A} hypermutation boundary of α chains is independent of local and neighbouring sequences and related to the distance from the initiation of transcription. <i>European Journal of Immunology</i> , 1997, 27, 3115-3120.	2.9	47
24	Uracil Accumulation and Mutagenesis Dominated by Cytosine Deamination in CpG Dinucleotides in Mice Lacking UNG and SMUG1. <i>Scientific Reports</i> , 2017, 7, 7199.	3.3	43
25	Affinity maturation leads to differential expression of multiple copies of a \hat{I} light-chain transgene. <i>Nature</i> , 1993, 363, 271-273.	27.8	41
26	CTNNBL1 Is a Novel Nuclear Localization Sequence-binding Protein That Recognizes RNA-splicing Factors CDC5L and Prp31. <i>Journal of Biological Chemistry</i> , 2011, 286, 17091-17102.	3.4	41
27	Uracil excision by endogenous SMUG 1 glycosylase promotes efficient I g class switching and impacts on A : T substitutions during somatic mutation. <i>European Journal of Immunology</i> , 2014, 44, 1925-1935.	2.9	40
28	Apobec2 deficiency causes mitochondrial defects and mitophagy in skeletal muscle. <i>FASEB Journal</i> , 2018, 32, 1428-1439.	0.5	20
29	CTNNBL1 facilitates the association of CWC15 with CDC5L and is required to maintain the abundance of the Prp19 spliceosomal complex. <i>Nucleic Acids Research</i> , 2015, 43, 7058-7069.	14.5	19
30	The cytoplasmic AID complex. <i>Seminars in Immunology</i> , 2012, 24, 273-280.	5.6	18
31	Epigenetics: Monoallelic Expression in the Immune System. <i>Current Biology</i> , 2002, 12, R108-R110.	3.9	16
32	Deficiency in spliceosome-associated factor CTNNBL1 does not affect ongoing cell cycling but delays exit from quiescence and results in embryonic lethality in mice. <i>Cell Cycle</i> , 2013, 12, 732-742.	2.6	13
33	The maturation of the antibody response. , 1995, , 57-81.		10
34	Structural and mutational analysis reveals that CTNNBL1 binds NLSs in a manner distinct from that of its closest armadillo \hat{I} -relative, karyopherin \hat{I} . <i>FEBS Letters</i> , 2014, 588, 21-27.	2.8	5
35	Aicardi \hat{I} Gouti \hat{I} res Syndrome associated mutations of RNase H2B impair its interaction with ZMYM3 and the CoREST histone-modifying complex. <i>PLoS ONE</i> , 2019, 14, e0213553.	2.5	5
36	AID and RPA: PKA makes the connection local. <i>Nature Immunology</i> , 2009, 10, 367-369.	14.5	4

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37	Mutagenesis by AID: Being in the Right Place at the Right Time. PLoS Genetics, 2015, 11, e1005489.	3.5	2
38	Harnessing mutation: The best of two worlds. Science, 2016, 353, 1206-1207.	12.6	1
39	The mechanism of somatic hypermutation at A•T pairs remains an open question. Nature Reviews Immunology, 2005, 5, 180-180.	22.7	1