

# Itziar Salaverria

## List of Publications by Year in descending order

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79  
papers

6,905  
citations

71102

41  
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66911

78  
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79  
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79  
docs citations

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times ranked

8184  
citing authors

#	ARTICLE	IF	CITATIONS
1	Non-coding recurrent mutations in chronic lymphocytic leukaemia. <i>Nature</i> , 2015, 526, 519-524.	27.8	749
2	Landscape of somatic mutations and clonal evolution in mantle cell lymphoma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 18250-18255.	7.1	488
3	Diffuse large B-cell lymphoma subgroups have distinct genetic profiles that influence tumor biology and improve gene-expression-based survival prediction. <i>Blood</i> , 2005, 106, 3183-3190.	1.4	348
4	Translocations activating IRF4 identify a subtype of germinal center-derived B-cell lymphoma affecting predominantly children and young adults. <i>Blood</i> , 2011, 118, 139-147.	1.4	281
5	Molecular Subsets of Mantle Cell Lymphoma Defined by the <i>IGHV</i> Mutational Status and SOX11 Expression Have Distinct Biologic and Clinical Features. <i>Cancer Research</i> , 2012, 72, 5307-5316.	0.9	231
6	A recurrent 11q aberration pattern characterizes a subset of MYC-negative high-grade B-cell lymphomas resembling Burkitt lymphoma. <i>Blood</i> , 2014, 123, 1187-1198.	1.4	185
7	CCND2 rearrangements are the most frequent genetic events in cyclin D1 <sup>hi</sup> mantle cell lymphoma. <i>Blood</i> , 2013, 121, 1394-1402.	1.4	183
8	Genome-wide association study identifies multiple risk loci for chronic lymphocytic leukemia. <i>Nature Genetics</i> , 2013, 45, 868-876.	21.4	179
9	Follicular lymphomas with and without translocation t(14;18) differ in gene expression profiles and genetic alterations. <i>Blood</i> , 2009, 114, 826-834.	1.4	177
10	Specific Secondary Genetic Alterations in Mantle Cell Lymphoma Provide Prognostic Information Independent of the Gene Expression <sup>hi</sup> -Based Proliferation Signature. <i>Journal of Clinical Oncology</i> , 2007, 25, 1216-1222.	1.6	166
11	Patient age at diagnosis is associated with the molecular characteristics of diffuse large B-cell lymphoma. <i>Blood</i> , 2012, 119, 1882-1887.	1.4	163
12	Uniparental disomies, homozygous deletions, amplifications, and target genes in mantle cell lymphoma revealed by integrative high-resolution whole-genome profiling. <i>Blood</i> , 2009, 113, 3059-3069.	1.4	162
13	Follicular Lymphomas in Children and Young Adults. <i>American Journal of Surgical Pathology</i> , 2013, 37, 333-343.	3.7	149
14	Genomic profiling reveals different genetic aberrations in systemic ALK <sup>hi</sup> positive and ALK <sup>hi</sup> negative anaplastic large cell lymphomas. <i>British Journal of Haematology</i> , 2008, 140, 516-526.	2.5	145
15	Follicular lymphoma grade 3B is a distinct neoplasm according to cytogenetic and immunohistochemical profiles. <i>Haematologica</i> , 2011, 96, 1327-1334.	3.5	142
16	Gray zone lymphoma: chromosomal aberrations with immunophenotypic and clinical correlations. <i>Modern Pathology</i> , 2011, 24, 1586-1597.	5.5	137
17	Mantle cell lymphoma: from pathology and molecular pathogenesis to new therapeutic perspectives. <i>Haematologica</i> , 2006, 91, 11-6.	3.5	135
18	Pediatric-type nodal follicular lymphoma: an indolent clonal proliferation in children and adults with high proliferation index and no BCL2 rearrangement. <i>Blood</i> , 2012, 120, 2395-2404.	1.4	132

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19	Pathway discovery in mantle cell lymphoma by integrated analysis of high-resolution gene expression and copy number profiling. <i>Blood</i> , 2010, 116, 953-961.	1.4	122
20	Genome-wide analysis of pediatric-type follicular lymphoma reveals low genetic complexity and recurrent alterations of TNFRSF14 gene. <i>Blood</i> , 2016, 128, 1101-1111.	1.4	115
21	Pediatric follicular lymphoma - a clinico-pathological study of a population-based series of patients treated within the Non-Hodgkin's Lymphoma - Berlin-Frankfurt-Munster (NHL-BFM) multicenter trials. <i>Haematologica</i> , 2010, 95, 253-259.	3.5	107
22	The Gray Zone Between Burkitt's Lymphoma and Diffuse Large B-Cell Lymphoma From a Genetics Perspective. <i>Journal of Clinical Oncology</i> , 2011, 29, 1835-1843.	1.6	104
23	Decoding the DNA Methylome of Mantle Cell Lymphoma in the Light of the Entire B Cell Lineage. <i>Cancer Cell</i> , 2016, 30, 806-821.	16.8	103
24	The complex landscape of genetic alterations in mantle cell lymphoma. <i>Seminars in Cancer Biology</i> , 2011, 21, 322-334.	9.6	100
25	Clinical, pathological and genetic features of primary mediastinal large B-cell lymphomas and mediastinal gray zone lymphomas in children. <i>Haematologica</i> , 2011, 96, 262-268.	3.5	92
26	Clinicopathologic Significance and Prognostic Value of Chromosomal Imbalances in Diffuse Large B-Cell Lymphomas. <i>Journal of Clinical Oncology</i> , 2004, 22, 3498-3506.	1.6	87
27	Leukemic involvement is a common feature in mantle cell lymphoma. <i>Cancer</i> , 2007, 109, 2473-2480.	4.1	82
28	Distinct molecular profile of IRF4-rearranged large B-cell lymphoma. <i>Blood</i> , 2020, 135, 274-286.	1.4	81
29	Chromosomal alterations detected by comparative genomic hybridization in subgroups of gene expression-defined Burkitt's lymphoma. <i>Haematologica</i> , 2008, 93, 1327-1334.	3.5	80
30	MicroRNA Expression, Chromosomal Alterations, and Immunoglobulin Variable Heavy Chain Hypermutations in Mantle Cell Lymphomas. <i>Cancer Research</i> , 2009, 69, 7071-7078.	0.9	78
31	CCND2 and CCND3 hijack immunoglobulin light-chain enhancers in cyclin D1 <sup>hi</sup> mantle cell lymphoma. <i>Blood</i> , 2019, 133, 940-951.	1.4	77
32	Comprehensive characterization of complex structural variations in cancer by directly comparing genome sequence reads. <i>Nature Biotechnology</i> , 2014, 32, 1106-1112.	17.5	74
33	Burkitt-like lymphoma with 11q aberration: a germinal center-derived lymphoma genetically unrelated to Burkitt lymphoma. <i>Haematologica</i> , 2019, 104, 1822-1829.	3.5	71
34	Mutations of MAP2K1 are frequent in pediatric-type follicular lymphoma and result in ERK pathway activation. <i>Blood</i> , 2017, 130, 323-327.	1.4	69
35	Loss of major histocompatibility class II expression in non-immune-privileged site diffuse large B-cell lymphoma is highly coordinated and not due to chromosomal deletions. <i>Blood</i> , 2005, 107, 1101-1107.	1.4	68
36	Incidence and prognostic impact of secondary cytogenetic aberrations in a series of 145 patients with mantle cell lymphoma. <i>Genes Chromosomes and Cancer</i> , 2010, 49, 439-451.	2.8	68

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37	Follicular lymphoma t(14;18)-negative is genetically a heterogeneous disease. <i>Blood Advances</i> , 2020, 4, 5652-5665.	5.2	67
38	Recurrent loss of heterozygosity in 1p36 associated with TNFRSF14 mutations in IRF4 translocation negative pediatric follicular lymphomas. <i>Haematologica</i> , 2013, 98, 1237-1241.	3.5	65
39	Assessment of SOX11 Expression in Routine Lymphoma Tissue Sections. <i>American Journal of Surgical Pathology</i> , 2014, 38, 86-93.	3.7	58
40	Genetic Predisposition to Chronic Lymphocytic Leukemia Is Mediated by a BMF Super-Enhancer Polymorphism. <i>Cell Reports</i> , 2016, 16, 2061-2067.	6.4	58
41	Genomic imbalances and patterns of karyotypic variability in mantle-cell lymphoma cell lines. <i>Leukemia Research</i> , 2006, 30, 923-934.	0.8	45
42	MAPK and JAK-STAT pathways dysregulation in plasmablastic lymphoma. <i>Haematologica</i> , 2021, 106, 2682-2693.	3.5	44
43	A comprehensive flow-cytometry-based immunophenotypic characterization of Burkitt-like lymphoma with 11q aberration. <i>Modern Pathology</i> , 2018, 31, 732-743.	5.5	42
44	Clinicopathological characteristics and genomic profile of primary sinonasal tract diffuse large B cell lymphoma (<scp>DLBCL</scp>) reveals gain at 1q31 and <scp>RGS</scp>1 encoding protein; high <scp>RGS</scp>1 immunohistochemical expression associates with poor overall survival in <scp>DLBCL</scp> not otherwise specified (<scp>NOS</scp>). <i>Histopathology</i> , 2017, 70, 595-621.	2.9	41
45	NOTCH1, TP53, and MAP2K1 Mutations in Splenic Diffuse Red Pulp Small B-cell Lymphoma Are Associated With Progressive Disease. <i>American Journal of Surgical Pathology</i> , 2016, 40, 192-201.	3.7	40
46	Epigenetic Activation of SOX11 in Lymphoid Neoplasms by Histone Modifications. <i>PLoS ONE</i> , 2011, 6, e21382.	2.5	38
47	A unique case of follicular lymphoma provides insights to the clonal evolution from follicular lymphoma in situ to manifest follicular lymphoma. <i>Blood</i> , 2011, 118, 3442-3444.	1.4	36
48	CREBBP gene mutations are frequently detected in in situ follicular neoplasia. <i>Blood</i> , 2018, 132, 2687-2690.	1.4	36
49	Loss of protein expression and recurrent DNA hypermethylation of the GNG7 gene in squamous cell carcinoma of the head and neck. <i>Journal of Applied Genetics</i> , 2012, 53, 167-174.	1.9	35
50	microRNA Expression Profiles Identify Subtypes of Mantle Cell Lymphoma with Different Clinicobiological Characteristics. <i>Clinical Cancer Research</i> , 2013, 19, 3121-3129.	7.0	35
51	Genomic complexity and IGHV mutational status are key predictors of outcome of chronic lymphocytic leukemia patients with TP53 disruption. <i>Haematologica</i> , 2014, 99, e231-e234.	3.5	33
52	Unbalanced expression of licensing DNA replication factors occurs in a subset of mantle cell lymphomas with genomic instability. <i>International Journal of Cancer</i> , 2006, 119, 2768-2774.	5.1	32
53	Dual PI3K/mTOR inhibition is required to effectively impair microenvironment survival signals in mantle cell lymphoma. <i>Oncotarget</i> , 2014, 5, 6788-6800.	1.8	32
54	High resolution copy number analysis of <i>IRF4</i> translocationâ€positive diffuse large Bâ€cell and follicular lymphomas. <i>Genes Chromosomes and Cancer</i> , 2013, 52, 150-155.	2.8	30

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55	Update on the molecular pathogenesis and clinical treatment of mantle cell lymphoma: report of the 10th annual conference of the European Mantle Cell Lymphoma Network. <i>Leukemia and Lymphoma</i> , 2011, 52, 2226-2236.	1.3	29
56	Analysis of Aurora-A and hMPS1 mitotic kinases in mantle cell lymphoma. <i>International Journal of Cancer</i> , 2006, 118, 357-363.	5.1	28
57	Multiple recurrent chromosomal breakpoints in mantle cell lymphoma revealed by a combination of molecular cytogenetic techniques. <i>Genes Chromosomes and Cancer</i> , 2008, 47, 1086-1097.	2.8	28
58	Improved classification of leukemic B-cell lymphoproliferative disorders using a transcriptional and genetic classifier. <i>Haematologica</i> , 2017, 102, e360-e363.	3.5	27
59	Diffuse large B-cell lymphomas in adults with aberrant coexpression of CD10, BCL6, and MUM1 are enriched in <i>IRF4</i> rearrangements. <i>Blood Advances</i> , 2022, 6, 2361-2372.	5.2	26
60	Synergistic anti-tumor activity of acadesine (AICAR) in combination with the anti-CD20 monoclonal antibody rituximab in <i>in vivo</i> and <i>in vitro</i> models of mantle cell lymphoma. <i>Oncotarget</i> , 2014, 5, 726-739.	1.8	25
61	Detection of chromothripsis-like patterns with a custom array platform for chronic lymphocytic leukemia. <i>Genes Chromosomes and Cancer</i> , 2015, 54, 668-680.	2.8	23
62	Update on the molecular pathogenesis and clinical treatment of Mantle Cell Lymphoma (MCL): minutes of the 9th European MCL Network conference. <i>Leukemia and Lymphoma</i> , 2010, 51, 1612-1622.	1.3	21
63	Follicular lymphoma grade 3B. <i>Best Practice and Research in Clinical Haematology</i> , 2011, 24, 111-119.	1.7	21
64	High-resolution copy number analysis of paired normal-tumor samples from diffuse large B cell lymphoma. <i>Annals of Hematology</i> , 2016, 95, 253-262.	1.8	19
65	LMO2-negative Expression Predicts the Presence of MYC Translocations in Aggressive B-Cell Lymphomas. <i>American Journal of Surgical Pathology</i> , 2017, 41, 877-886.	3.7	19
66	Frequent polymorphic changes but not mutations of TRAIL receptors DR4 and DR5 in mantle cell lymphoma and other B-cell lymphoid neoplasms. <i>Haematologica</i> , 2004, 89, 1322-31.	3.5	19
67	A unifying hypothesis for PNMZL and PTFL: morphological variants with a common molecular profile. <i>Blood Advances</i> , 2022, 6, 4661-4674.	5.2	19
68	The <i>CBFA2T3/ACSF3</i> locus is recurrently involved in <i>IGH</i> chromosomal translocation t(14;16)(q32;q24) in pediatric B-cell lymphoma with germinal center phenotype. <i>Genes Chromosomes and Cancer</i> , 2012, 51, 338-343.	2.8	18
69	Gene expression profile and genomic changes in disease progression of early-stage chronic lymphocytic leukemia. <i>Haematologica</i> , 2008, 93, 132-136.	3.5	17
70	Response: proliferative versus functional anergy. <i>Blood</i> , 2011, 118, 3442-3442.	1.4	16
71	The molecular hallmarks of primary and secondary vitreoretinal lymphoma. <i>Blood Advances</i> , 2021, , .	5.2	16
72	Cryptic insertions of the immunoglobulin light chain enhancer region near <i>CCND1</i> in t(11;14)-negative mantle cell lymphoma. <i>Haematologica</i> , 2020, 105, e408-e411.	3.5	13

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73	Novel Double Factor PGT strategy analyzing blastocyst stage embryos in a single NGS procedure. PLoS ONE, 2018, 13, e0205692.	2.5	10
74	Follicular Lymphomas with and without Translocation t(14;18) Differ in Gene Expression Profiles and Genetic Alterations.. Blood, 2007, 110, 360-360.	1.4	7
75	Risk of Central Nervous System (CNS) Involvement in Patients with Mantle Cell Lymphoma (MCL): Analysis of Clinico-Biological Factors in a Series of 283 Cases. Blood, 2014, 124, 1677-1677.	1.4	4
76	Chromosomal Imbalances in Germinal Center B-Cell-Like and Activated B-Cell-Like Diffuse Large B-Cell Lymphoma Influence Gene Expression Signatures and Improve Gene Expression-Based Survival Prediction(the First Two Authors Contributed Equally to This Work).. Blood, 2004, 104, 415-415.	1.4	1
77	Checkâ€™ing DLBCL. Oncoscience, 2015, 2, 71-72.	2.2	1
78	The Genomic Landscape of Plasmablastic Lymphoma (PBL) - an L.L.M.P.P. Project. Blood, 2021, 138, 1326-1326.	1.4	1
79	SNP Array Analysis Reveals Copy Number Alterations and Uniparental Disomy in Mantle Cell Lymphomas at High Resolution.. Blood, 2007, 110, 1585-1585.	1.4	0