## Jean-Philippe Merlio

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Exploring <i>hTERT</i> promoter methylation in cutaneous T ell lymphomas. Molecular Oncology, 2022, 16, 1931-1946.	4.6	12
2	Telomeric Repeat-Containing RNA (TERRA): A Review of the Literature and First Assessment in Cutaneous T-Cell Lymphomas. Genes, 2022, 13, 539.	2.4	6
3	Discoidin Domain Receptor 2 orchestrates melanoma resistance combining phenotype switching and proliferation. Oncogene, 2022, 41, 2571-2586.	5.9	6
4	Integrative diagnosis of primary cutaneous large B-cell lymphomas supports the relevance of cell of origin profiling. PLoS ONE, 2022, 17, e0266978.	2.5	4
5	Clinical impact of STK11 mutation in advanced-stage non-small cell lung cancer. European Journal of Cancer, 2022, 172, 85-95.	2.8	20
6	Cutaneous Lymphocyte Antigen Is a PotentialÂTherapeutic Target in Cutaneous T-Cell Lymphoma. Journal of Investigative Dermatology, 2022, 142, 3243-3252.e10.	0.7	6
7	Diagnosis and treatment of lymphomas in the era of epigenetics. Blood Reviews, 2021, 48, 100782.	5.7	7
8	Xenograft and cell culture models of Sézary syndrome reveal cell of origin diversity and subclonal heterogeneity. Leukemia, 2021, 35, 1696-1709.	7.2	16
9	C6 Ceramide (d18:1/6:0) as a Novel Treatment of Cutaneous T Cell Lymphoma. Cancers, 2021, 13, 270.	3.7	8
10	Lack of clinical relevance of blood clonality in primary cutaneous marginal zone B-cell lymphoma. European Journal of Dermatology, 2021, 31, 94-96.	0.6	3
11	Lymphomatoid papulosis types D and E: a multicentre series of the French Cutaneous Lymphomas Study Group. Clinical and Experimental Dermatology, 2021, 46, 1441-1451.	1.3	6
12	Cytokines, Genetic Lesions and Signaling Pathways in Anaplastic Large Cell Lymphomas. Cancers, 2021, 13, 4256.	3.7	1
13	CRISPR-Cas9 globin editing can induce megabase-scale copy-neutral losses of heterozygosity in hematopoietic cells. Nature Communications, 2021, 12, 4922.	12.8	44
14	MSI-High RAS-BRAF wild-type colorectal adenocarcinomas with MLH1 loss have a high frequency of targetable oncogenic gene fusions whose diagnoses are feasible using methods easy-to-implement in pathology laboratories. Human Pathology, 2021, 114, 99-109.	2.0	16
15	A novel 3D culture model recapitulates primary FL B-cell features and promotes their survival. Blood Advances, 2021, 5, 5372-5386.	5.2	18
16	Single-cell trajectories in Sézary syndrome. Blood, 2021, 138, 1384-1386.	1.4	1
17	Targeting Epigenetic Modifiers Can Reduce the Clonogenic Capacities of Sézary Cells. Frontiers in Oncology, 2021, 11, 775253.	2.8	3
18	Positive Association Between Location of Melanoma, Ultraviolet Signature, Tumor Mutational Burden, and Response to Anti–PD-1 Therapy. JCO Precision Oncology, 2021, 5, 1821-1829.	3.0	17

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19	Challenges in Assessing MYC Rearrangement in Primary Cutaneous Diffuse Large B-Cell Lymphoma, Leg-Type. American Journal of Surgical Pathology, 2020, 44, 424-427.	3.7	3
20	hMZF-2, the Elusive Transcription Factor. Frontiers in Genetics, 2020, 11, 581115.	2.3	1
21	Reliable blood cancer cells' telomere length evaluation by qPCR. Cancer Medicine, 2020, 9, 3153-3162.	2.8	13
22	Outcomes of Patients With Advanced NSCLC From the Intergroupe Francophone de Cancérologie Thoracique Biomarkers France Study by KRAS Mutation Subtypes. JTO Clinical and Research Reports, 2020, 1, 100052.	1.1	9
23	Circulating Tumor Cell Clusters: United We Stand Divided We Fall. International Journal of Molecular Sciences, 2020, 21, 2653.	4.1	73
24	Mutations of the B-Cell Receptor Pathway Confer Chemoresistance in Primary Cutaneous Diffuse Large B-Cell Lymphoma Leg Type. Journal of Investigative Dermatology, 2019, 139, 2334-2342.e8.	0.7	28
25	Primary cutaneous large Bâ€cell lymphomas: relevance of the 2017 World Health Organization classification: clinicopathological and molecular analyses of 64 cases. Histopathology, 2019, 74, 1067-1080.	2.9	28
26	A Single-Arm Phase II Trial of Lenalidomide in Relapsing or Refractory Primary Cutaneous Large B-Cell Lymphoma, LegÂType. Journal of Investigative Dermatology, 2018, 138, 1982-1989.	0.7	27
27	Double-hit or dual expression of MYC and BCL2 in primary cutaneous large B-cell lymphomas. Modern Pathology, 2018, 31, 1332-1342.	5.5	31
28	Assessment of BRAFV600E mutation in pulmonary Langerhans cell histiocytosis in tissue biopsies and bronchoalveolar lavages by droplet digital polymerase chain reaction. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2018, 472, 247-258.	2.8	10
29	Calcium Independent Effect of Orai1 and STIM1 in Non-Hodgkin B Cell Lymphoma Dissemination. Cancers, 2018, 10, 402.	3.7	7
30	SATB1 Is a Pivotal Epigenetic Biomarker in Cutaneous T-Cell Lymphomas. Journal of Investigative Dermatology, 2018, 138, 1694-1696.	0.7	11
31	Identification of Somatic Mutations in Primary Cutaneous Diffuse Large B-Cell Lymphoma, Leg Type by Massive Parallel Sequencing. Journal of Investigative Dermatology, 2017, 137, 1984-1994.	0.7	93
32	Clinical and molecular characteristics of non-small-cell lung cancer (NSCLC) harboring EGFR mutation: results of the nationwide French Cooperative Thoracic Intergroup (IFCT) program. Annals of Oncology, 2017, 28, 2715-2724.	1.2	72
33	TP53 alterations in primary and secondary Sézary syndrome: A diagnostic tool for the assessment of malignancy in patients with erythroderma. PLoS ONE, 2017, 12, e0173171.	2.5	13
34	Telomerase Activation in Hematological Malignancies. Genes, 2016, 7, 61.	2.4	25
35	Hybridization Capture-Based Next-Generation Sequencing to Evaluate Coding Sequence and Deep Intronic Mutations in the NF1 Gene. Genes, 2016, 7, 133.	2.4	12
36	Intrahepatic Xenograft of Cutaneous T-Cell Lymphoma Cell Lines. American Journal of Pathology, 2016, 186, 1775-1785.	3.8	11

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37	MYD88 Somatic Mutation Is a Diagnostic Criterion in Primary Cutaneous Large B-CellÂLymphoma. Journal of Investigative Dermatology, 2016, 136, 1741-1744.	0.7	46
38	Proliferative Nodules vs Melanoma Arising in Giant Congenital Melanocytic Nevi During Childhood. JAMA Dermatology, 2016, 152, 1147.	4.1	21
39	Sézary Syndrome: Translating Genetic Diversity into Personalized Medicine. Journal of Investigative Dermatology, 2016, 136, 1319-1324.	0.7	16
40	Primary digestive melanoma in association with tubular adenoma: a case report illustrating the distinction from metastatic colonic melanoma. Human Pathology, 2016, 48, 167-171.	2.0	3
41	Routine molecular profiling of patients with advanced non-small-cell lung cancer: results of a 1-year nationwide programme of the French Cooperative Thoracic Intergroup (IFCT). Lancet, The, 2016, 387, 1415-1426.	13.7	790
42	Molecular alterations and tumor suppressive function of the <i>DUSP22 (Dual Specificity) Tj ETQq0 0 0 rgBT /C</i>	verlock 10 1.8	) Tf 50 542 Td 41
43	Diagnostic and Prognostic Value of <i>BCL2</i> Rearrangement in 53 Patients With Follicular Lymphoma Presenting as Primary Skin Lesions. American Journal of Clinical Pathology, 2015, 143, 362-373.	0.7	38
44	PLCG1 Gene Mutations Are Uncommon in Cutaneous T-Cell Lymphomas. Journal of Investigative Dermatology, 2015, 135, 2334-2337.	0.7	16
45	Detection of BRAF V600 Mutations in Melanoma: Evaluation of Concordance between the Cobas® 4800 BRAF V600 Mutation Test and the Methods Used in French National Cancer Institute (INCa) Platforms in a Real-Life Setting. PLoS ONE, 2015, 10, e0120232.	2.5	24
46	Multiple genetic alterations in primary cutaneous large B-cell lymphoma, leg type support a common lymphomagenesis with activated B-cell-like diffuse large B-cell lymphoma. Modern Pathology, 2014, 27, 402-411.	5.5	78
47	BRAFV600E mutation analysis by immunohistochemistry in patients with thoracic metastases from colorectal cancer. Pathology, 2014, 46, 311-315.	0.6	10
48	High Frequency and Clinical Prognostic Value of MYD88 L265P Mutation in Primary Cutaneous Diffuse Large B-Cell Lymphoma, Leg-Type. JAMA Dermatology, 2014, 150, 1173.	4.1	110
49	Telomerase functions beyond telomere maintenance in primary cutaneous T-cell lymphoma. Blood, 2014, 123, 1850-1859.	1.4	24
50	Diagnostic value of immunohistochemistry for the detection of the BRAF mutation in primary lung adenocarcinoma Caucasian patients. Annals of Oncology, 2013, 24, 742-748.	1.2	103
51	Tumor Homogeneity between Primary and Metastatic Sites for BRAF Status in Metastatic Melanoma Determined by Immunohistochemical and Molecular Testing. PLoS ONE, 2013, 8, e70826.	2.5	97
52	MYD88 Somatic Mutation Is a Genetic Feature of Primary Cutaneous Diffuse Large B-Cell Lymphoma, Leg Type. Journal of Investigative Dermatology, 2012, 132, 2118-2120.	0.7	85
53	IRF4 Gene Rearrangements Define a Subgroup of CD30-Positive Cutaneous T-Cell Lymphoma: A Study of 54 Cases. Journal of Investigative Dermatology, 2010, 130, 816-825.	0.7	114
54	Genome-Wide Analysis of Cutaneous T-Cell Lymphomas Identifies Three Clinically Relevant Classes. Journal of Investigative Dermatology, 2010, 130, 1707-1718.	0.7	100

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55	CDKN2A–CDKN2B deletion defines an aggressive subset of cutaneous T-cell lymphoma. Modern Pathology, 2010, 23, 547-558.	5.5	80
56	Primary Cutaneous T-Cell Lymphomas Do not Show Specific NAV3 Gene Deletion or Translocation. Journal of Investigative Dermatology, 2008, 128, 2458-2466.	0.7	24
57	Common chromosomal abnormalities in mycosis fungoides transformation. Genes Chromosomes and Cancer, 2007, 46, 828-838.	2.8	44
58	Large cell transformation of mycosis fungoides: tetraploidization within skin tumor large cells. Cancer Genetics and Cytogenetics, 2005, 163, 1-6.	1.0	21
59	Neoplastic Cells Do Not Carry bcl2-JH Rearrangements Detected in a Subset of Primary Cutaneous Follicle Center B-cell Lymphomas. American Journal of Surgical Pathology, 2004, 28, 748-755.	3.7	51
60	Evidence that an Identical T Cell Clone in Skin and Peripheral Blood Lymphocytes is an Independent Prognostic Factor in Primary Cutaneous T Cell Lymphomas. Journal of Investigative Dermatology, 2001, 117, 920-926.	0.7	74
61	Characterization of t(2;5) reciprocal transcripts and genomic breakpoints in CD30+ cutaneous lymphoproliferations. Blood, 1998, 91, 4668-76.	1.4	18