## Marie Tosolini

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/133353/publications.pdf

Version: 2024-02-01

52 papers 19,114 citations

218592 26 h-index 206029 48 g-index

58 all docs

58 docs citations

58 times ranked 30564 citing authors

#	Article	IF	CITATIONS
1	Type, Density, and Location of Immune Cells Within Human Colorectal Tumors Predict Clinical Outcome. Science, 2006, 313, 1960-1964.	6.0	5,356
2	ClueGO: a Cytoscape plug-in to decipher functionally grouped gene ontology and pathway annotation networks. Bioinformatics, 2009, 25, 1091-1093.	1.8	5,348
3	Spatiotemporal Dynamics of Intratumoral Immune Cells Reveal the Immune Landscape in Human Cancer. Immunity, 2013, 39, 782-795.	6.6	2,983
4	Clinical Impact of Different Classes of Infiltrating T Cytotoxic and Helper Cells (Th1, Th2, Treg, Th17) in Patients with Colorectal Cancer. Cancer Research, 2011, 71, 1263-1271.	0.4	986
5	Histopathologic-Based Prognostic Factors of Colorectal Cancers Are Associated With the State of the Local Immune Reaction. Journal of Clinical Oncology, 2011, 29, 610-618.	0.8	864
6	In Situ Cytotoxic and Memory T Cells Predict Outcome in Patients With Early-Stage Colorectal Cancer. Journal of Clinical Oncology, 2009, 27, 5944-5951.	0.8	822
7	The tumor microenvironment and Immunoscore are critical determinants of dissemination to distant metastasis. Science Translational Medicine, 2016, 8, 327ra26.	5.8	360
8	Biomolecular Network Reconstruction Identifies T-Cell Homing Factors Associated With Survival in Colorectal Cancer. Gastroenterology, 2010, 138, 1429-1440.	0.6	280
9	Coordination of Intratumoral Immune Reaction and Human Colorectal Cancer Recurrence. Cancer Research, 2009, 69, 2685-2693.	0.4	262
10	Inter―and intraâ€ŧumoural heterogeneity in cancerâ€associated fibroblasts of human pancreatic ductal adenocarcinoma. Journal of Pathology, 2019, 248, 51-65.	2.1	215
11	Functional Network Pipeline Reveals Genetic Determinants Associated with in Situ Lymphocyte Proliferation and Survival of Cancer Patients. Science Translational Medicine, 2014, 6, 228ra37.	5.8	181
12	Single-cell RNA sequencing unveils the shared and the distinct cytotoxic hallmarks of human TCRVÎ'1 and TCRVÎ'2 Î3Î'T lymphocytes. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 11906-11915.	3.3	152
13	Assessment of tumor-infiltrating TCRV $\langle b \rangle \hat{l}^3 \langle b \rangle 9V \langle b \rangle \hat{l}' \langle b \rangle 2 \langle b \rangle \hat{l}^3 \hat{l}' \langle b \rangle $ lymphocyte abundance by deconvolution of human cancers microarrays. Oncolmmunology, 2017, 6, e1284723.	2.1	134
14	Distinctive features of tumor-infiltrating $\hat{I}^3\hat{I}$ T lymphocytes in human colorectal cancer. Oncolmmunology, 2017, 6, e1347742.	2.1	119
15	Single-Cell Signature Explorer for comprehensive visualization of single cell signatures across scRNA-seq datasets. Nucleic Acids Research, 2019, 47, e133-e133.	6.5	96
16	Dendrogenin A drives LXR to trigger lethal autophagy in cancers. Nature Communications, 2017, 8, 1903.	5.8	84
17	Several immune escape patterns in non-Hodgkin's lymphomas. Oncolmmunology, 2015, 4, e1026530.	2.1	82
18	Mechanisms of PD-1/PD-L1 expression and prognostic relevance in non-Hodgkin lymphoma: a summary of immunohistochemical studies. Oncotarget, 2017, 8, 44960-44975.	0.8	82

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19	PD-1 blockade restores helper activity of tumor-infiltrating, exhausted PD-1hiCD39+ CD4 T cells. JCI Insight, 2021, 6, .	2.3	64
20	Boosting $\hat{I}^3\hat{I}$ T cell-mediated antibody-dependent cellular cytotoxicity by PD-1 blockade in follicular lymphoma. Oncolmmunology, 2019, 8, 1554175.	2.1	53
21	Large-scale microarray profiling reveals four stages of immune escape in non-Hodgkin lymphomas. Oncolmmunology, 2016, 5, e1188246.	2.1	43
22	Microtubule-Driven Stress Granule Dynamics Regulate Inhibitory Immune Checkpoint Expression in T Cells. Cell Reports, 2019, 26, 94-107.e7.	2.9	42
23	Mitochondrial inhibitors circumvent adaptive resistance to venetoclax and cytarabine combination therapy in acute myeloid leukemia. Nature Cancer, 2021, 2, 1204-1223.	5.7	42
24	lonizing radiation induces endothelial transdifferentiation of glioblastoma stem-like cells through the Tie2 signaling pathway. Cell Death and Disease, 2019, 10, 816.	2.7	34
25	GAPDH Overexpression in the T Cell Lineage Promotes Angioimmunoblastic T Cell Lymphoma through an NF-κB-Dependent Mechanism. Cancer Cell, 2019, 36, 268-287.e10.	7.7	34
26	Long non-coding RNA expression profile in cytogenetically normal acute myeloid leukemia identifies a distinct signature and a new biomarker in NPM1-mutated patients. Haematologica, 2017, 102, 1718-1726.	1.7	32
27	Neutral Sphingomyelinase 2 Heightens Anti-Melanoma Immune Responses and Anti–PD-1 Therapy Efficacy. Cancer Immunology Research, 2021, 9, 568-582.	1.6	30
28	Nurse-like cells promote CLL survival through LFA-3/CD2 interactions. Oncotarget, 2017, 8, 52225-52236.	0.8	28
29	FOXP3-enriched infiltrates associated with better outcome in renal allografts with inflamed fibrosis. Nephrology Dialysis Transplantation, 2009, 24, 3847-3854.	0.4	25
30	A Tridimensional Model for NK Cell-Mediated ADCC of Follicular Lymphoma. Frontiers in Immunology, 2019, 10, 1943.	2.2	22
31	MCM8- and MCM9 Deficiencies Cause Lifelong Increased Hematopoietic DNA Damage Driving p53-Dependent Myeloid Tumors. Cell Reports, 2019, 28, 2851-2865.e4.	2.9	20
32	EBV infection determines the immune hallmarks of plasmablastic lymphoma. Oncolmmunology, 2018, 7, e1486950.	2.1	19
33	Profiling Immune Escape in Hodgkin's and Diffuse large B-Cell Lymphomas Using the Transcriptome and Immunostaining. Cancers, 2018, 10, 415.	1.7	19
34	Longitudinal CITE-Seq profiling of chronic lymphocytic leukemia during ibrutinib treatment: evolution of leukemic and immune cells at relapse. Biomarker Research, 2020, 8, 72.	2.8	19
35	Pancreatic cancer intrinsic PI3Kα activity accelerates metastasis and rewires macrophage component. EMBO Molecular Medicine, 2021, 13, e13502.	3.3	19
36	Human Monocyte Recognition of Adenosine-Based Cyclic Dinucleotides Unveils the A2a G <sub>αs</sub> Protein-Coupled Receptor Tonic Inhibition of Mitochondrially Induced Cell Death. Molecular and Cellular Biology, 2015, 35, 479-495.	1.1	18

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#	Article	IF	Citations
37	Colon-specific immune microenvironment regulates cancer progression versus rejection. Oncolmmunology, 2020, 9, 1790125.	2.1	17
38	Baseline SUV <sub>max</sub> is related to tumor cell proliferation and patient outcome in follicular lymphoma. Haematologica, 2022, 107, 221-230.	1.7	17
39	Dual Relief of T-lymphocyte Proliferation and Effector Function Underlies Response to PD-1 Blockade in Epithelial Malignancies. Cancer Immunology Research, 2020, 8, 869-882.	1.6	16
40	PTMselect: optimization of protein modifications discovery by mass spectrometry. Scientific Reports, 2019, 9, 4181.	1.6	14
41	Single-Cell Virtual Cytometer allows user-friendly and versatile analysis and visualization of multimodal single cell RNAseq datasets. NAR Genomics and Bioinformatics, 2020, 2, Iqaa025.	1.5	13
42	ALK-transformed mature T lymphocytes restore early thymus progenitor features. Journal of Clinical Investigation, 2020, 130, 6395-6408.	3.9	12
43	Single-Cell RNAseq Profiling of Human $\hat{I}^{\hat{I}}$ T Lymphocytes in Virus-Related Cancers and COVID-19 Disease. Viruses, 2021, 13, 2212.	1.5	12
44	Phased differentiation of $\hat{l}^3\hat{l}$ and T CD8 tumor-infiltrating lymphocytes revealed by single-cell transcriptomics of human cancers. Oncolmmunology, 2021, 10, 1939518.	2.1	11
45	Lymphoma Heterogeneity Unraveled by Single-Cell Transcriptomics. Frontiers in Immunology, 2021, 12, 597651.	2.2	9
46	Cyclic dinucleotides modulate human Tâ€cell response through monocyte cell death. European Journal of Immunology, 2015, 45, 3313-3323.	1.6	8
47	RND1 regulates migration of human glioblastoma stem-like cells according to their anatomical localization and defines a prognostic signature in glioblastoma. Oncotarget, 2018, 9, 33788-33803.	0.8	7
48	Intratumoral immune reaction: A novel paradigm for cancer Journal of Clinical Oncology, 2011, 29, 471-471.	0.8	1
49	nwCompare and AutoCompare Softwares for Proteomics and Transcriptomics Data Mining – Application to the Exploration of Gene Expression Profiles of Aggressive Lymphomas. , 0, , .		1
50	Simultaneous growth of two cancer cell lines demonstrates variability in growth rates. Open Access Medical Statistics, 2014, , 29.	0.5	0
51	EBV infection promotes tumor infiltrating leucocyte and immune escape in plasmablastic lymphoma according to gene expression profiling. Hematological Oncology, 2017, 35, 297-298.	0.8	0
52	Boosting Gamma Delta T Cells-Mediated ADCC By PD-1 Blockade in Follicular Lymphoma. Blood, 2018, 132, 5381-5381.	0.6	0