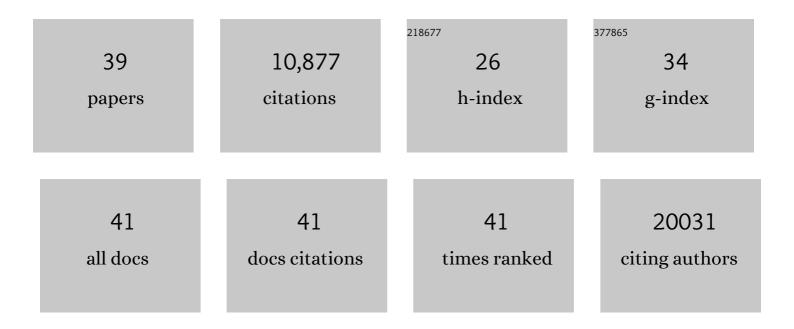
## Luigi Nezi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4631616/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Gut microbiome modulates response to anti–PD-1 immunotherapy in melanoma patients. Science, 2018, 359, 97-103.	12.6	3,126
2	Genomic Classification of Cutaneous Melanoma. Cell, 2015, 161, 1681-1696.	28.9	2,562
3	DNA breaks and chromosome pulverization from errors in mitosis. Nature, 2012, 482, 53-58.	27.8	1,051
4	Oncogene ablation-resistant pancreatic cancer cells depend on mitochondrial function. Nature, 2014, 514, 628-632.	27.8	998
5	The Mad1/Mad2 Complex as a Template for Mad2 Activation in the Spindle Assembly Checkpoint. Current Biology, 2005, 15, 214-225.	3.9	376
6	Dietary fiber and probiotics influence the gut microbiome and melanoma immunotherapy response. Science, 2021, 374, 1632-1640.	12.6	369
7	Passenger deletions generate therapeutic vulnerabilities in cancer. Nature, 2012, 488, 337-342.	27.8	294
8	Gut microbiota signatures are associated with toxicity to combined CTLA-4 and PD-1 blockade. Nature Medicine, 2021, 27, 1432-1441.	30.7	216
9	Reporting guidelines for human microbiome research: the STORMS checklist. Nature Medicine, 2021, 27, 1885-1892.	30.7	170
10	Cross-cohort gut microbiome associations with immune checkpoint inhibitor response in advanced melanoma. Nature Medicine, 2022, 28, 535-544.	30.7	158
11	Accumulation of long-chain fatty acids in the tumor microenvironment drives dysfunction in in intrapancreatic CD8+ T cells. Journal of Experimental Medicine, 2020, 217, .	8.5	142
12	Sister chromatid tension and the spindle assembly checkpoint. Current Opinion in Cell Biology, 2009, 21, 785-795.	5.4	137
13	Syndecan 1 is a critical mediator of macropinocytosis in pancreatic cancer. Nature, 2019, 568, 410-414.	27.8	129
14	Determinants of conformational dimerization of Mad2 and its inhibition by p31comet. EMBO Journal, 2006, 25, 1273-1284.	7.8	124
15	Genomic and immune heterogeneity are associated with differential responses to therapy in melanoma. Npj Genomic Medicine, 2017, 2, .	3.8	120
16	InÂVivo Functional Platform Targeting Patient-Derived Xenografts Identifies WDR5-Myc Association as a Critical Determinant of Pancreatic Cancer. Cell Reports, 2016, 16, 133-147.	6.4	114
17	ILF2 Is a Regulator of RNA Splicing and DNA Damage Response in 1q21-Amplified Multiple Myeloma. Cancer Cell, 2017, 32, 88-100.e6.	16.8	114
18	Synthetic vulnerabilities of mesenchymal subpopulations in pancreatic cancer. Nature, 2017, 542, 362-366.	27.8	105

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19	Telomere Dysfunction Drives Aberrant Hematopoietic Differentiation and Myelodysplastic Syndrome. Cancer Cell, 2015, 27, 644-657.	16.8	85
20	Genetic Events That Limit the Efficacy of MEK and RTK Inhibitor Therapies in a Mouse Model of KRAS-Driven Pancreatic Cancer. Cancer Research, 2015, 75, 1091-1101.	0.9	68
21	Protein-tyrosine Phosphatase PTPD1 Regulates Focal Adhesion Kinase Autophosphorylation and Cell Migration. Journal of Biological Chemistry, 2008, 283, 10919-10929.	3.4	64
22	Extra-mitochondrial localisation of frataxin and its association with IscU1 during enterocyte-like differentiation of the human colon adenocarcinoma cell line Caco-2. Journal of Cell Science, 2005, 118, 3917-3924.	2.0	61
23	Truncating PREX2 mutations activate its GEF activity and alter gene expression regulation in NRAS-mutant melanoma. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E1296-305.	7.1	59
24	Accumulation of Mad2–Cdc20 complex during spindle checkpoint activation requires binding of open and closed conformers of Mad2 in Saccharomyces cerevisiae. Journal of Cell Biology, 2006, 174, 39-51.	5.2	51
25	Fecal Microbiota Transplantation Controls Murine Chronic Intestinal Inflammation by Modulating Immune Cell Functions and Gut Microbiota Composition. Cells, 2019, 8, 517.	4.1	50
26	PAF promotes stemness and radioresistance of glioma stem cells. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E9086-E9095.	7.1	40
27	Role of the Mad2 Dimerization Interface in the Spindle Assembly Checkpoint Independent of Kinetochores. Current Biology, 2012, 22, 1900-1908.	3.9	26
28	TERT promoter mutations and melanoma survival: A comprehensive literature review and meta-analysis. Critical Reviews in Oncology/Hematology, 2021, 160, 103288.	4.4	20
29	Sex Differences in Efficacy and Toxicity of Systemic Cancer Treatments: Role of the Microbiome. Journal of Clinical Oncology, 2019, 37, 439-439.	1.6	16
30	A cell-of-origin epigenetic tracer reveals clinically distinct subtypes of high-grade serous ovarian cancer. Genome Medicine, 2020, 12, 94.	8.2	11
31	The Role of Proteases in Fibronectin Matrix Remodeling in Thyroid Epithelial Cell Monolayer Cultures. Biological Chemistry, 2002, 383, 167-76.	2.5	7
32	Short-term treatment with multi-drug regimens combining BRAF/MEK-targeted therapy and immunotherapy results in durable responses in <i>Braf</i> -mutated melanoma. Oncolmmunology, 2021, 10, 1992880.	4.6	7
33	Characterization of DIP1, a novel nuclear protein in Drosophila melanogaster. Biochemical and Biophysical Research Communications, 2003, 307, 224-228.	2.1	4
34	1072P Primary ipilimumab/nivolumab immunotherapy followed by adjuvant nivolumab in patients with locally advanced or oligometastatic melanoma: Update on outcome. Annals of Oncology, 2021, 32, S889-S890.	1.2	1
35	1147P Primary ipilimumab/nivolumab immunotherapy followed by adjuvant nivolumab in locally advanced or oligometastatic melanoma: Preliminary results. Annals of Oncology, 2020, 31, S765-S766.	1.2	0
36	ILF2 Is a Regulator of RNA Splicing and DNA Damage Response in 1q21-Amplified Multiple Myeloma. Blood, 2014, 124, 30-30.	1.4	0

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#	Article	IF	CITATIONS
37	Abstract 976: Metabolic eradication of treatment resistant cancer stem cells in pancreatic tumors: A clonal tracking-based platform for identifying the best personalized treatment. , 2015, , .		Ο
38	Abstract 1701: Identification of epigenetic modifiers able to suppress growth of pancreatic ductal adenocarcinoma: A patient-orientedin vivofunctional platform. , 2015, , .		0
39	ILF2-YB1 Protein Interaction Modulates RNA Splicing to Induce Resistance to Chemotherapy in High Risk Multiple Myeloma. Blood, 2016, 128, 359-359.	1.4	0