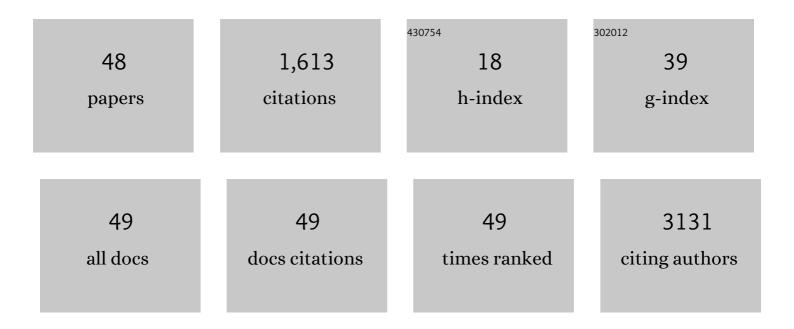
## Alessandra Tessitore

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6062792/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	The Inflammatory Microenvironment in Hepatocellular Carcinoma: A Pivotal Role for Tumor-Associated Macrophages. BioMed Research International, 2013, 2013, 1-15.	0.9	332
2	TrkA alternative splicing. Cancer Cell, 2004, 6, 347-360.	7.7	194
3	Smart Hydrogel Particles:  Biomarker Harvesting:  One-Step Affinity Purification, Size Exclusion, and Protection against Degradation. Nano Letters, 2008, 8, 350-361.	4.5	182
4	An Initial Characterization of the Serum Phosphoproteome. Journal of Proteome Research, 2009, 8, 5523-5531.	1.8	86
5	MicroRNAs in the DNA Damage/Repair Network and Cancer. International Journal of Genomics, 2014, 2014, 1-10.	0.8	80
6	MicroRNA expression analysis in high fat diet-induced NAFLD-NASH-HCC progression: study on C57BL/6J mice. BMC Cancer, 2016, 16, 3.	1.1	77
7	Next-generation sequencing: recent applications to the analysis of colorectal cancer. Journal of Translational Medicine, 2017, 15, 246.	1.8	76
8	Cancer secretome and inflammation: The bright and the dark sides of NF-κB. Seminars in Cell and Developmental Biology, 2018, 78, 51-61.	2.3	72
9	Two gamma-interferon-activation sites (GAS) on the promoter of the human intercellular adhesion molecule (ICAM-1) gene are required for induction of transcription by IFN-gamma. FEBS Journal, 1998, 258, 968-975.	0.2	35
10	MRE11 inhibition highlights a replication stress-dependent vulnerability of MYCN-driven tumors. Cell Death and Disease, 2018, 9, 895.	2.7	35
11	Serum Biomarkers Identification by Mass Spectrometry in High-Mortality Tumors. International Journal of Proteomics, 2013, 2013, 1-15.	2.0	33
12	Targeting the NF-κB pathway in prostate cancer: a promising therapeutic approach?. Current Drug Targets, 2016, 17, 311-320.	1.0	32
13	Neuroprotective effects of human amniotic fluid stem cells-derived secretome in an ischemia/reperfusion model. Stem Cells Translational Medicine, 2021, 10, 251-266.	1.6	31
14	Long-term abuse of a high-carbohydrate diet is as harmful as a high-fat diet for development and progression of liver injury in a mouse model of NAFLD/NASH. Nutrition, 2020, 75-76, 110782.	1.1	29
15	Therapeutic Use of MicroRNAs in Cancer. Anti-Cancer Agents in Medicinal Chemistry, 2015, 16, 7-19.	0.9	25
16	Development of hepatocellular cancer induced by long term low fat-high carbohydrate diet in a NAFLD/NASH mouse model. Oncotarget, 2017, 8, 53482-53494.	0.8	25
17	The prevalent KRAS exon 2 c.35 G>A mutation in metastatic colorectal cancer patients: A biomarker of worse prognosis and potential benefit of bevacizumab-containing intensive regimens?. Critical Reviews in Oncology/Hematology, 2015, 93, 190-202.	2.0	24
18	Applications of Next Generation Sequencing to the Analysis of Familial Breast/Ovarian Cancer. High-Throughput, 2020, 9, 1.	4.4	22

Alessandra Tessitore

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19	Mesenchymal stem cells of Systemic Sclerosis patients, derived from different sources, show a profibrotic microRNA profiling. Scientific Reports, 2019, 9, 7144.	1.6	18
20	Bioinformatics approach to predict target genes for dysregulated microRNAs in hepatocellular carcinoma: study on a chemically-induced HCC mouse model. BMC Bioinformatics, 2015, 16, 408.	1.2	17
21	Circulating MicroRNAs as Prognostic and Therapeutic Biomarkers in Breast Cancer Molecular Subtypes. Journal of Personalized Medicine, 2020, 10, 98.	1.1	16
22	High sensitivity of detection ofTP53 somatic mutations by fluorescence-assisted mismatch analysis. Genes Chromosomes and Cancer, 2002, 35, 86-91.	1.5	15
23	New mutations and protein variants ofNBS1 are identified in cancer cell lines. Genes Chromosomes and Cancer, 2003, 36, 198-204.	1.5	15
24	A Simplified Genomic Profiling Approach Predicts Outcome in Metastatic Colorectal Cancer. Cancers, 2019, 11, 147.	1.7	15
25	KCTD11Tumor Suppressor Gene Expression Is Reduced in Prostate Adenocarcinoma. BioMed Research International, 2014, 2014, 1-9.	0.9	13
26	Effective treatment of a platinum‑resistant cutaneous squamous cell carcinoma case by EGFR pathway inhibition. Molecular and Clinical Oncology, 2018, 9, 30-34.	0.4	13
27	Emerging Role of isomiRs in Cancer: State of the Art and Recent Advances. Genes, 2021, 12, 1447.	1.0	11
28	Evaluating the role of FAMIly history of cancer and diagnosis of multiple neoplasms in cancer patients receiving PD-1/PD-L1 checkpoint inhibitors: the multicenter FAMI-L1 study. OncoImmunology, 2020, 9, 1710389.	2.1	9
29	Identification of a novel HLA-DRB1*11 allele: DRB1*1152. Tissue Antigens, 2006, 67, 180-182.	1.0	8
30	Novel P53 mutations detected by FAMA in colorectal cancers. Annals of Oncology, 2006, 17, vii78-vii83.	0.6	8
31	Low Radiation Environment Switches the Overgrowth-Induced Cell Apoptosis Toward Autophagy. Frontiers in Public Health, 2020, 8, 594789.	1.3	8
32	Serum Low-Molecular-Weight Protein Fractionation for Biomarker Discovery. Methods in Molecular Biology, 2012, 823, 237-249.	0.4	6
33	Prognostic significance of clinicopathological factors in early breast cancer: 20 years of follow-up in a single-center analysis. Oncotarget, 2017, 8, 72031-72043.	0.8	6
34	Neoadjuvant chemotherapy in breast cancer: a dose-dense schedule in real life and putative role of <i>PIK3CA</i> mutations. Oncotarget, 2018, 9, 27380-27396.	0.8	6
35	Prenatal diagnosis of a rhodopsin mutation using chemical cleavage of the mismatch. Prenatal Diagnosis, 2002, 22, 380-384.	1.1	5
36	Reverse-phase protein microarray highlights HER2 signaling activation in immunohistochemistry/FISH/HER2-negative breast cancers. Expert Review of Proteomics, 2013, 10, 223-226.	1.3	5

Alessandra Tessitore

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37	KRAS and 2 rare PI3KCA mutations coexisting in a metastatic colorectal cancer patient with aggressive and resistant disease. Human Pathology, 2018, 74, 178-182.	1.1	5
38	Identification of the uncommon allele HLA-A*7403 in a Caucasian renal transplant cadaveric donor: extension of the exon 4 sequence. Tissue Antigens, 2007, 69, 615-618.	1.0	4
39	SMO-M2 mutation does not support cell-autonomous Hedgehog activity in cerebellar granule cell precursors. Scientific Reports, 2019, 9, 19623.	1.6	4
40	Involvement of an Arachidonic-Acid-Dependent Pathway in the Interferon-beta-Mediated Expression of C202 Gene in Ehrlich-Ascites-Tumor Cells. FEBS Journal, 1996, 235, 91-96.	0.2	3
41	Increased <scp>CD</scp> 1D polymorphism: identification of two novel alleles, <scp>CD</scp> 1D*03 and *04, in individuals from <scp>M</scp> orocco. International Journal of Immunogenetics, 2015, 42, 287-291.	0.8	3
42	MICAâ^—078: A novel allele identified in a Moroccan individual affected by celiac disease. Human Immunology, 2015, 76, 438-441.	1.2	3
43	Codon optimization by 0-1 linear programming. Computers and Operations Research, 2020, 119, 104932.	2.4	3
44	MicroRNAs Expression in Response to rhNGF in Epithelial Corneal Cells: Focus on Neurotrophin Signaling Pathway. International Journal of Molecular Sciences, 2022, 23, 3597.	1.8	2
45	Concurrent RAS and RAS/BRAF V600E Variants in Colorectal Cancer: More Frequent Than Expected? A Case Report. Frontiers in Oncology, 2022, 12, 863639.	1.3	2
46	Aryl hydrocarbon receptor Interacting Protein (AIP) status in a functional adrenal adenoma occurring in a patient with a germline AIP mutation. Endocrine Abstracts, 0, , .	0.0	0
47	Prognostic relevance of KRAS genotype and the prevalent C.35 G > a mutation in metastatic colorectal cancer (MCRC) patients fitting for intensive FIr-B/FOx triplet chemotherapy plus bevacizumab Journal of Clinical Oncology, 2014, 32, e14575-e14575.	0.8	0
48	Ultrasound-Based Method for the Identification of Novel MicroRNA Biomarkers in Prostate Cancer. Genes, 2021, 12, 1726.	1.0	0