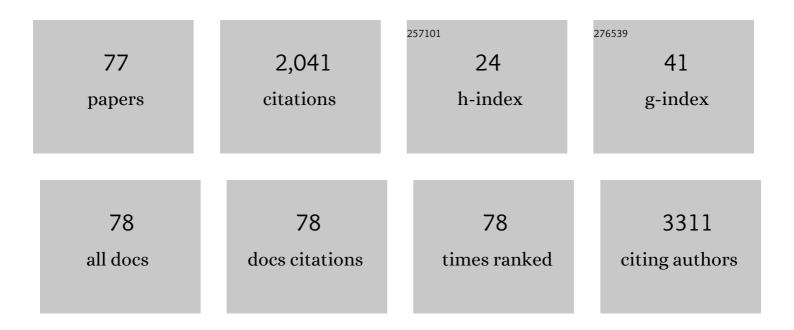
## Valentina Indio

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Filaggrin Loss-of-Function Mutations Are Risk Factors for Severe Food Allergy in Children with Atopic Dermatitis. Journal of Clinical Medicine, 2021, 10, 233.	1.0	14
2	Targeted therapy in <i>SDH-</i> deficient GIST. Therapeutic Advances in Medical Oncology, 2021, 13, 175883592110232.	1.4	16
3	The Identity of PDGFRA D842V-Mutant Gastrointestinal Stromal Tumors (GIST). Cancers, 2021, 13, 705.	1.7	13
4	Non-Coding RNAs in the Transcriptional Network That Differentiates Skeletal Muscles of Sedentary from Long-Term Endurance- and Resistance-Trained Elderly. International Journal of Molecular Sciences, 2021, 22, 1539.	1.8	15
5	Gene Expression Landscape of SDH-Deficient Gastrointestinal Stromal Tumors. Journal of Clinical Medicine, 2021, 10, 1057.	1.0	9
6	Case Report: The Complete Remission of a Mixed Germ Cell Tumor With Somatic Type Malignancy of Sarcoma Type With a GCT-Oriented Therapy: Clinical Findings and Genomic Profiling. Frontiers in Oncology, 2021, 11, 633543.	1.3	2
7	Torque teno mini virus as a cause of childhood acute promyelocytic leukemia lacking PML/RARA fusion. Blood, 2021, 138, 1773-1777.	0.6	16
8	Genetic Factors Associated With Pain Severity, Daily Opioid Dose Requirement, and Pain Response Among Advanced Cancer Patients Receiving Supportive Care. Journal of Pain and Symptom Management, 2021, 62, 785-795.	0.6	5
9	iPSC-Derived Gaucher Macrophages Display Growth Impairment and Activation of Inflammation-Related Cell Death. Cells, 2021, 10, 2822.	1.8	6
10	Role of Mir-192-5p during Response to Azacitidine and Lenalidomide Therapy in Myelodysplastic Syndromes. Blood, 2021, 138, 3673-3673.	0.6	0
11	SDHA Germline Variants in Adult Patients With SDHA-Mutant Gastrointestinal Stromal Tumor. Frontiers in Oncology, 2021, 11, 778461.	1.3	4
12	Whole exome sequencing reveals mutations in FAT1 tumor suppressor gene clinically impacting on peripheral T-cell lymphoma not otherwise specified. Modern Pathology, 2020, 33, 179-187.	2.9	37
13	Gene duplication, rather than epigenetic changes, drives FGF4 overexpression in KIT/PDGFRA/SDH/RAS-P WT GIST. Scientific Reports, 2020, 10, 19829.	1.6	10
14	Phenotypic and Functional Characterization of NK Cells in αβT-Cell and B-Cell Depleted Haplo-HSCT to Cure Pediatric Patients with Acute Leukemia. Cancers, 2020, 12, 2187.	1.7	19
15	Impact of Chemotherapy in the Adjuvant Setting of Early Stage Uterine Leiomyosarcoma: A Systematic Review and Updated Meta-Analysis. Cancers, 2020, 12, 1899.	1.7	26
16	Inhibition of Methyltransferase DOT1L Sensitizes to Sorafenib Treatment AML Cells Irrespective of MLL-Rearrangements: A Novel Therapeutic Strategy for Pediatric AML. Cancers, 2020, 12, 1972.	1.7	19
17	MicroRNA profiling of blastic plasmacytoid dendritic cell neoplasm and myeloid sarcoma. Hematological Oncology, 2020, 38, 831-833.	0.8	1
18	Genomic Database Analysis of Uterine Leiomyosarcoma Mutational Profile. Cancers, 2020, 12, 2126.	1.7	44

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19	Direct Antiviral Treatments for Hepatitis C Virus Have Off-Target Effects of Oncologic Relevance in Hepatocellular Carcinoma. Cancers, 2020, 12, 2674.	1.7	13
20	Targeted Deep Sequencing Uncovers Cryptic KIT Mutations in KIT/PDGFRA/SDH/RAS-P Wild-Type GIST. Frontiers in Oncology, 2020, 10, 504.	1.3	16
21	The Emerging Role of the FCF/FGFR Pathway in Gastrointestinal Stromal Tumor. International Journal of Molecular Sciences, 2020, 21, 3313.	1.8	22
22	Skeletal Muscle Gene Expression in Long-Term Endurance and Resistance Trained Elderly. International Journal of Molecular Sciences, 2020, 21, 3988.	1.8	17
23	Genetic aberrations and molecular biology of cardiac sarcoma. Therapeutic Advances in Medical Oncology, 2020, 12, 175883592091849.	1.4	13
24	Gene Expression Profiling of PDGFRA Mutant GIST Reveals Immune Signatures as a Specific Fingerprint of D842V Exon 18 Mutation. Frontiers in Immunology, 2020, 11, 851.	2.2	10
25	Effect of Lactobacillus acidophilus D2/CSL (CECT 4529) supplementation in drinking water on chicken crop and caeca microbiome. PLoS ONE, 2020, 15, e0228338.	1.1	25
26	The Pediatric Acute Leukemia Fusion Oncogene ETO2â€GLIS2 Increases Selfâ€Renewal and Alters Differentiation in a Human Induced Pluripotent Stem Cellsâ€Derived Model. HemaSphere, 2020, 4, e319.	1.2	8
27	Paratesticular Mesenchymal Malignancies: A Single-Center Case Series, Clinical Management, and Review of Literature. Integrative Cancer Therapies, 2020, 19, 153473541990055.	0.8	5
28	Genetics and treatment of gastrointestinal stromal tumors with immune checkpoint inhibitors: what do we know?. Pharmacogenomics, 2020, 21, 231-234.	0.6	6
29	Immune microenvironment profiling of gastrointestinal stromal tumors (CIST) shows gene expression patterns associated to immune checkpoint inhibitors response. Oncolmmunology, 2019, 8, e1617588.	2.1	41
30	Influence of Atopic Dermatitis on Cow's Milk Allergy in Children. Medicina (Lithuania), 2019, 55, 460.	0.8	16
31	<p>Mechanisms of resistance to a PI3K inhibitor in gastrointestinal stromal tumors: an <em>omic</em> approach to identify novel druggable targets</p> . Cancer Management and Research, 2019, Volume 11, 6229-6244.	0.9	2
32	Comparative Assessment of Antitumor Effects and Autophagy Induction as a Resistance Mechanism by Cytotoxics and EZH2 Inhibition in INI1-Negative Epithelioid Sarcoma Patient-Derived Xenograft. Cancers, 2019, 11, 1015.	1.7	21
33	BCOR involvement in cancer. Epigenomics, 2019, 11, 835-855.	1.0	113
34	Gain of FGF4 is a frequent event in KIT/PDGFRA/SDH/RASâ€P WT GIST. Genes Chromosomes and Cancer, 2019, 58, 636-642.	1.5	22
35	Molecular modelling evaluation of exon 18 His845_Asn848delinsPro PDGFRα mutation in a metastatic GIST patient responding to imatinib. Scientific Reports, 2019, 9, 2172.	1.6	5
36	Blastic plasmacytoid dendritic cell neoplasm: genomics mark epigenetic dysregulation as a primary therapeutic target. Haematologica, 2019, 104, 729-737.	1.7	58

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37	An exploratory study by DMET array identifies a germline signature associated with imatinib response in gastrointestinal stromal tumor. Pharmacogenomics Journal, 2019, 19, 390-400.	0.9	20
38	Successful multidisciplinary clinical approach and molecular characterization by whole transcriptome sequencing of a cardiac myxofibrosarcoma: A case report. World Journal of Clinical Cases, 2019, 7, 3018-3026.	0.3	7
39	LncRNAs as novel players in hepatocellular carcinoma recurrence. Oncotarget, 2018, 9, 35085-35099.	0.8	46
40	18F-FDG-PET/CT imaging in cardiac tumors: illustrative clinical cases and review of the literature. Therapeutic Advances in Medical Oncology, 2018, 10, 175883591879356.	1.4	28
41	Mutational burden of resectable pancreatic cancer, as determined by whole transcriptome and whole exome sequencing, predicts a poor prognosis. International Journal of Oncology, 2018, 52, 1972-1980.	1.4	8
42	Integrated Molecular Characterization of Gastrointestinal Stromal Tumors (GIST) Harboring the Rare D842V Mutation in PDGFRA Gene. International Journal of Molecular Sciences, 2018, 19, 732.	1.8	29
43	Identification of an Actionable Mutation of KIT in a Case of Extraskeletal Myxoid Chondrosarcoma. International Journal of Molecular Sciences, 2018, 19, 1855.	1.8	4
44	Novel intraâ€genic large deletions of <i>CTNNB1</i> gene identified in WT desmoidâ€ŧype fibromatosis. Genes Chromosomes and Cancer, 2018, 57, 495-503.	1.5	18
45	Negative Prognostic Relevance of a Specific 3-Gene Cluster in Myelodysplastic Syndromes during Azacitidine and Lenalidomide Therapy. Blood, 2018, 132, 4347-4347.	0.6	0
46	Genome-Wide Analysis Identifies MEN1 and MAX Mutations and a Neuroendocrine-Like Molecular Heterogeneity in Quadruple WT GIST. Molecular Cancer Research, 2017, 15, 553-562.	1.5	53
47	Hh/Gli antagonist in acute myeloid leukemia with CBFA2T3-GLIS2 fusion gene. Journal of Hematology and Oncology, 2017, 10, 26.	6.9	34
48	<scp>HSPA</scp> 8 as a novel fusion partner of <scp>NR</scp> 4 <scp>A</scp> 3 in extraskeletal myxoid chondrosarcoma. Genes Chromosomes and Cancer, 2017, 56, 582-586.	1.5	38
49	Adaptive Immunity in Fibrosarcomatous Dermatofibrosarcoma Protuberans and Response to Imatinib Treatment. Journal of Investigative Dermatology, 2017, 137, 484-493.	0.3	29
50	Personalization of regorafenib treatment in metastatic gastrointestinal stromal tumours in real-life clinical practice. Therapeutic Advances in Medical Oncology, 2017, 9, 731-739.	1.4	20
51	Identification of SRF-E2F1 fusion transcript in EWSR-negative myoepithelioma of the soft tissue. Oncotarget, 2017, 8, 60036-60045.	0.8	17
52	Identification of a cytogenetic and molecular subgroup of acute myeloid leukemias showing sensitivity to L-Asparaginase. Oncotarget, 2017, 8, 109915-109923.	0.8	27
53	Genomic complexity and dynamics of clonal evolution in childhood acute myeloid leukemia studied with whole-exome sequencing. Oncotarget, 2016, 7, 56746-56757.	0.8	23
54	Evolution of Dermatofibrosarcoma Protuberans to DFSP-Derived Fibrosarcoma: An Event Marked by Epithelial–Mesenchymal Transition–like Process and 22q Loss. Molecular Cancer Research, 2016, 14, 820-829.	1.5	25

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55	Integrating miRNA and gene expression profiling analysis revealed regulatory networks in gastrointestinal stromal tumors. Epigenomics, 2016, 8, 1347-1366.	1.0	23
56	Efficacy and Biological Activity of Imatinib in Metastatic Dermatofibrosarcoma Protuberans (DFSP). Clinical Cancer Research, 2016, 22, 837-846.	3.2	78
57	Copy number gain of chromosome 3q is a recurrent event in patients with intraductal papillary mucinous neoplasm (IPMN) associated with disease progression. Oncotarget, 2016, 7, 74797-74806.	0.8	7
58	Characterization of pancreatic ductal adenocarcinoma using whole transcriptome sequencing and copy number analysis by single-nucleotide polymorphism array. Molecular Medicine Reports, 2015, 12, 7479-7484.	1.1	20
59	SDHC methylation in gastrointestinal stromal tumors (GIST): a case report. BMC Medical Genetics, 2015, 16, 87.	2.1	22
60	Identification of the NUP98-PHF23 fusion gene in pediatric cytogenetically normal acute myeloid leukemia by whole-transcriptome sequencing. Journal of Hematology and Oncology, 2015, 8, 69.	6.9	14
61	Whole transcriptome sequencing identifies BCOR internal tandem duplication as a common feature of clear cell sarcoma of the kidney. Oncotarget, 2015, 6, 40934-40939.	0.8	61
62	Whole exome sequencing (WES) on formalin-fixed, paraffin-embedded (FFPE) tumor tissue in gastrointestinal stromal tumors (GIST). BMC Genomics, 2015, 16, 892.	1.2	48
63	Good survival outcome of metastatic SDH-deficient gastrointestinal stromal tumors harboring SDHA mutations. Genetics in Medicine, 2015, 17, 391-395.	1.1	41
64	Molecular characterization of metastatic exon 11 mutant gastrointestinal stromal tumors (GIST) beyond KIT/PDGFRα genotype evaluated by next generation sequencing (NGS). Oncotarget, 2015, 6, 42243-42257.	0.8	20
65	Integrated genomic study of quadruple-WT GIST (KIT/PDGFRA/SDH/RAS pathway wild-type GIST). BMC Cancer, 2014, 14, 685.	1.1	70
66	Whole transcriptome sequencing of a paediatric case of <i>de novo</i> acute myeloid leukaemia with del(5q) reveals <i><scp>RUNX</scp>1</i> â€ <i><scp>USP</scp>42</i> and <i><scp>PRDM</scp>16â€<scp>SKI</scp></i> fusion transcripts. British Journal of Haematology, 2014, 166, 449-452.	1.2	12
67	Activity of sunitinib in extraskeletal myxoid chondrosarcoma. European Journal of Cancer, 2014, 50, 1657-1664.	1.3	74
68	Dystrophin deregulation is associated with tumor progression in KIT/PDGFRA mutant gastrointestinal stromal tumors. Clinical Sarcoma Research, 2014, 4, 9.	2.3	9
69	Analysis of all subunits, SDHA, SDHB, SDHC, SDHD, of the succinate dehydrogenase complex in KIT/PDGFRA wild-type GIST. European Journal of Human Genetics, 2014, 22, 32-39.	1.4	90
70	Expression of IGF-1 receptor in KIT/PDGF receptor-α wild-type gastrointestinal stromal tumors with succinate dehydrogenase complex dysfunction. Future Oncology, 2013, 9, 121-126.	1.1	30
71	The prediction of organelle-targeting peptides in eukaryotic proteins with Grammatical-Restrained Hidden Conditional Random Fields. Bioinformatics, 2013, 29, 981-988.	1.8	17
72	CBFA2T3-GLIS2 fusion transcript is a novel common feature in pediatric, cytogenetically normal AML, not restricted to FAB M7 subtype. Blood, 2013, 121, 3469-3472.	0.6	119

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73	DHH-RHEBL1fusion transcript: a novel recurrent feature in the new landscape of pediatricCBFA2T3-GLIS2-positive acute myeloid leukemia. Oncotarget, 2013, 4, 1712-1720.	0.8	23
74	Gene fusions evidence in a KIT/PDGFRA wild-type GIST without mutations in SDH units identified by a whole transcriptome study Journal of Clinical Oncology, 2013, 31, e21523-e21523.	0.8	0
75	Genome study of PDGFRA D842V mutant GIST using next generation sequencing approach Journal of Clinical Oncology, 2013, 31, 10540-10540.	0.8	Ο
76	SDHA Loss-of-Function Mutations in KIT-PDGFRA Wild-Type Gastrointestinal Stromal Tumors Identified by Massively Parallel Sequencing. Journal of the National Cancer Institute, 2011, 103, 983-987.	3.0	137
77	MemPype: a pipeline for the annotation of eukaryotic membrane proteins. Nucleic Acids Research, 2011, 39, W375-W380.	6.5	28