Manuel Salto-Tellez

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1718329/publications.pdf

Version: 2024-02-01

123 papers 8,041 citations

35 h-index 82 g-index

127 all docs

 $\begin{array}{c} 127 \\ \text{docs citations} \end{array}$

times ranked

127

15375 citing authors

#	Article	IF	CITATIONS
1	Severity of gastric intestinal metaplasia predicts the risk of gastric cancer: a prospective multicentre cohort study (GCEP). Gut, 2022, 71, 854-863.	6.1	57
2	Identification of a prognostic signature in colorectal cancer using combinatorial algorithmâ€driven analysis. Journal of Pathology: Clinical Research, 2022, , .	1.3	1
3	Potential quality pitfalls of digitalized whole slide image of breast pathology in routine practice. Modern Pathology, 2022, 35, 903-910.	2.9	8
4	Swarm learning for decentralized artificial intelligence in cancer histopathology. Nature Medicine, 2022, 28, 1232-1239.	15.2	77
5	General Roadmap and Core Steps for the Development of Al Tools in Digital Pathology. Diagnostics, 2022, 12, 1272.	1.3	4
6	Digital pathology and artificial intelligence will be key to supporting clinical and academic cellular pathology through COVID-19 and future crises: the PathLAKE consortium perspective. Journal of Clinical Pathology, 2021, 74, 443-447.	1.0	49
7	Identifying mismatch repairâ€deficient colon cancer: nearâ€perfect concordance between immunohistochemistry and microsatellite instability testing in a large, populationâ€based series. Histopathology, 2021, 78, 401-413.	1.6	55
8	Diagnosis of digestive system tumours. International Journal of Cancer, 2021, 148, 1040-1050.	2.3	36
9	In-depth Clinical and Biological Exploration of DNA Damage Immune Response as a Biomarker for Oxaliplatin Use in Colorectal Cancer. Clinical Cancer Research, 2021, 27, 288-300.	3.2	13
10	QuPath: The global impact of an open source digital pathology system. Computational and Structural Biotechnology Journal, 2021, 19, 852-859.	1.9	49
11	Evolutionary genetic algorithm identifies <i>IL2RB</i> as a potential predictive biomarker for immune-checkpoint therapy in colorectal cancer. NAR Genomics and Bioinformatics, 2021, 3, Iqab016.	1.5	10
12	Colonic epithelial cathelicidin (<scp>LL</scp> â€37) expression intensity is associated with progression of colorectal cancer and presence of <scp>CD8</scp> ⁺ T cell infiltrate. Journal of Pathology: Clinical Research, 2021, 7, 495-506.	1.3	8
13	The clinical and molecular significance associated with STING signaling in breast cancer. Npj Breast Cancer, 2021, 7, 81.	2.3	21
14	A Means of Assessing Deep Learning-Based Detection of ICOS Protein Expression in Colon Cancer. Cancers, 2021, 13, 3825.	1.7	17
15	Orthogonal <i>MET</i> analysis in a populationâ€representative stage Il–III colon cancer cohort: prognostic and potential therapeutic implications. Molecular Oncology, 2021, 15, 3317-3328.	2.1	3
16	HistoClean: Open-source software for histological image pre-processing and augmentation to improve development of robust convolutional neural networks. Computational and Structural Biotechnology Journal, 2021, 19, 4840-4853.	1.9	5
17	Metastasis and Immune Evasion from Extracellular cGAMP Hydrolysis. Cancer Discovery, 2021, 11, 1212-1227.	7.7	139
18	PD-L1 Multiplex and Quantitative Image Analysis for Molecular Diagnostics. Cancers, 2021, 13, 29.	1.7	11

#	Article	IF	CITATIONS
19	Comparison of Molecular Assays for HPV Testing in Oropharyngeal Squamous Cell Carcinomas: A Population-Based Study in Northern Ireland. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 31-38.	1.1	14
20	Systematic evaluation of PAXgene® tissue fixation for the histopathological and molecular study of lung cancer. Journal of Pathology: Clinical Research, 2020, 6, 40-54.	1.3	8
21	Alcohol intake, tobacco smoking, and esophageal adenocarcinoma survival: a molecular pathology epidemiology cohort study. Cancer Causes and Control, 2020, 31, 1-11.	0.8	16
22	Immune status is prognostic for poor survival in colorectal cancer patients and is associated with tumour hypoxia. British Journal of Cancer, 2020, 123, 1280-1288.	2.9	45
23	A robust multiplex immunofluorescence and digital pathology workflow for the characterisation of the tumour immune microenvironment. Molecular Oncology, 2020, 14, 2384-2402.	2.1	71
24	Improving the Diagnostic Accuracy of the PD-L1 Test with Image Analysis and Multiplex Hybridization. Cancers, 2020, 12, 1114.	1.7	34
25	The adaptive immune and immune checkpoint landscape of neoadjuvant treated esophageal adenocarcinoma using digital pathology quantitation. BMC Cancer, 2020, 20, 500.	1.1	20
26	Lowâ€contact and highâ€interconnectivity pathology (LC&HI Path): postâ€COVID19â€pandemic practice of pathology. Histopathology, 2020, 77, 518-524.	1.6	9
27	Gastrointestinal tissueâ€based molecular biomarkers: a practical categorisation based on the 2019 World Health Organization classification of epithelial digestive tumours. Histopathology, 2020, 77, 340-350.	1.6	26
28	NUQA: Estimating Cancer Spatial and Temporal Heterogeneity and Evolution through Alignment-Free Methods. Molecular Biology and Evolution, 2019, 36, 2883-2889.	3.5	1
29	Defining the molecular evolution of extrauterine high grade serous carcinoma. Gynecologic Oncology, 2019, 155, 305-317.	0.6	17
30	Training and accreditation standards for pathologists undertaking clinical trial work. Journal of Pathology: Clinical Research, 2019, 5, 100-107.	1.3	10
31	Cancer taxonomy: pathology beyond pathology. European Journal of Cancer, 2019, 115, 57-60.	1.3	10
32	Recommendations for determining HPV status in patients with oropharyngeal cancers under TNM8 guidelines: a two-tier approach. British Journal of Cancer, 2019, 120, 827-833.	2.9	51
33	Immune activation by DNA damage predicts response to chemotherapy and survival in oesophageal adenocarcinoma. Gut, 2019, 68, 1918-1927.	6.1	18
34	Invited reviewâ€"next-generation sequencing: a modern tool in cytopathology. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2019, 475, 3-11.	1.4	31
35	High PTGS2 expression in postâ€neoadjuvant chemotherapyâ€treated oesophageal adenocarcinoma is associated with improved survival: a populationâ€based cohort study. Histopathology, 2019, 74, 587-596.	1.6	1
36	Critical Appraisal of Programmed Death Ligand 1 Reflex Diagnostic Testing: Current Standards and Future Opportunities. Journal of Thoracic Oncology, 2019, 14, 45-53.	0.5	42

3

#	Article	IF	Citations
37	Artificial intelligence—the third revolution in pathology. Histopathology, 2019, 74, 372-376.	1.6	107
38	Practical guide for the comparison of two next-generation sequencing systems for solid tumour analysis in a universal healthcare system. Journal of Clinical Pathology, 2019, 72, 225-231.	1.0	7
39	A digital pathology demonstration of an "immune hot" ICOS+/CD45RO+ immunephenotype and the impact on survival in patients with esophageal adenocarcinoma Journal of Clinical Oncology, 2019, 37, 4062-4062.	0.8	0
40	Abstract LB-088: Exploratory multiplex tissue image analysis of the impact of heterogeneity in the microenvironment of primary colorectal cancer on apoptosis markers in patients. , 2019, , .		0
41	Abstract 2787: Artificial intelligence approach identifies IL2RB as a common prognostic and potential predictive biomarker associated with immune checkpoints in colorectal cancer. , 2019, , .		0
42	Validation of the systematic scoring of immunohistochemically stained tumour tissue microarrays using <i>QuPath</i> digital image analysis. Histopathology, 2018, 73, 327-338.	1.6	63
43	HER2 testing of gastro-oesophageal adenocarcinoma: a commentary and guidance document from the Association of Clinical Pathologists Molecular Pathology and Diagnostics Committee. Journal of Clinical Pathology, 2018, 71, 388-394.	1.0	14
44	More Than a Decade of Molecular Diagnostic Cytopathology Leading Diagnostic and Therapeutic Decision-Making. Archives of Pathology and Laboratory Medicine, 2018, , .	1.2	6
45	Integrated tumor identification and automated scoring minimizes pathologist involvement and provides new insights to key biomarkers in breast cancer. Laboratory Investigation, 2018, 98, 15-26.	1.7	81
46	More Than a Decade of Molecular Diagnostic Cytopathology Leading Diagnostic and Therapeutic Decision-Making. Archives of Pathology and Laboratory Medicine, 2018, 142, 443-445.	1.2	26
47	A gene signature associated with PTEN activation defines good prognosis intermediate risk prostate cancer cases. Journal of Pathology: Clinical Research, 2018, 4, 103-113.	1.3	20
48	Time for change: a new training programme for morpho-molecular pathologists?. Journal of Clinical Pathology, 2018, 71, 285-290.	1.0	21
49	Automated Tumour Recognition and Digital Pathology Scoring Unravels New Role for PD-L1 in Predicting Good Outcome in ER-/HER2+ Breast Cancer. Journal of Oncology, 2018, 2018, 1-14.	0.6	44
50	Impact of Variable RNA-Sequencing Depth on Gene Expression Signatures and Target Compound Robustness: Case Study Examining Brain Tumor (Glioma) Disease Progression. JCO Precision Oncology, 2018, 2, 1-17.	1.5	3
51	Glucose transporter 1 expression as a marker of prognosis in oesophageal adenocarcinoma. Oncotarget, 2018, 9, 18518-18528.	0.8	13
52	Vitamin D receptor as a marker of prognosis in oesophageal adenocarcinoma: a prospective cohort study. Oncotarget, 2018, 9, 34347-34356.	0.8	7
53	Sex hormone receptor expression and survival in esophageal adenocarcinoma: a prospective cohort study. Oncotarget, 2018, 9, 35300-35312.	0.8	6
54	Abstract 1555: A gene signature associated with PTEN activation defines good outcomes in intermediate-risk prostate cancer cases. , 2018 , , .		0

#	Article	IF	CITATIONS
55	Abstract 4049: Assessment of immune biomarkers by digital pathological analysis across a large colorectal cancer patient cohort predicts patient outcome and may provide a clinically relevant therapeutic index for immunotherapeutic treatment stratification., 2018,,.		0
56	Abstract 3142: Tumor-infiltrating lymphocytes and CD4/FOXP3 ratios reliably predict survival using digital image analysis. , 2018, , .		0
57	Abstract B035: Radio-resistance of PTEN-deficient prostate tumors is enhanced by treatment-induced chemokine signaling and is associated with biochemical recurrence and development of metastasis., 2018,,.		0
58	Tissue-based next generation sequencing: application in a universal healthcare system. British Journal of Cancer, 2017, 116, 553-560.	2.9	38
59	BCL-2 system analysis identifies high-risk colorectal cancer patients. Gut, 2017, 66, 2141-2148.	6.1	40
60	Evaluation of PTGS2 Expression, PIK3CA Mutation, Aspirin Use and Colon Cancer Survival in a Population-Based Cohort Study. Clinical and Translational Gastroenterology, 2017, 8, e91.	1.3	56
61	Natural killer-like signature observed post therapy in locally advanced rectal cancer is a determinant of pathological response and improved survival. Modern Pathology, 2017, 30, 1287-1298.	2.9	23
62	Statin use, candidate mevalonate pathway biomarkers, and colon cancer survival in a population-based cohort study. British Journal of Cancer, 2017, 116, 1652-1659.	2.9	37
63	Molecular profiling of signet ring cell colorectal cancer provides a strong rationale for genomic targeted and immune checkpoint inhibitor therapies. British Journal of Cancer, 2017, 117, 203-209.	2.9	38
64	Standardising RNA profiling based biomarker application in cancerâ€"The need for robust control of technical variables. Biochimica Et Biophysica Acta: Reviews on Cancer, 2017, 1868, 258-272.	3.3	16
65	Targeting c-MET in gastrointestinal tumours: rationale, opportunities and challenges. Nature Reviews Clinical Oncology, 2017, 14, 562-576.	12.5	150
66	QuPath: Open source software for digital pathology image analysis. Scientific Reports, 2017, 7, 16878.	1.6	3,854
67	Activation of STING-Dependent Innate Immune Signaling By S-Phase-Specific DNA Damage in Breast Cancer. Journal of the National Cancer Institute, 2017, 109, djw199.	3.0	338
68	A Stepwise Integrated Approach to Personalized Risk Predictions in Stage III Colorectal Cancer. Clinical Cancer Research, 2017, 23, 1200-1212.	3.2	21
69	Morphomolecular pathology: setting the framework for a new generation of pathologists. British Journal of Cancer, 2017, 117, 1581-1582.	2.9	16
70	Rare cancers: the greatest inequality in cancer research and oncology treatment. British Journal of Cancer, 2017, 117, 1255-1257.	2.9	8
71	RNAscope <i>in situ</i> hybridization confirms mRNA integrity in formalin-fixed, paraffin-embedded cancer tissue samples. Oncotarget, 2017, 8, 93392-93403.	0.8	41
72	PD-L1 expression and response to neo-adjuvant chemotherapy in esophageal adenocarcinoma Journal of Clinical Oncology, 2017, 35, 4023-4023.	0.8	1

#	Article	IF	Citations
73	samExploreR: exploring reproducibility and robustness of RNA-seq results based on SAM files. Bioinformatics, 2016, 32, 3345-3347.	1.8	11
74	p16 as a prognostic indicator in ovarian/tubal highâ€grade serous carcinoma. Histopathology, 2016, 68, 615-618.	1.6	8
75	Validation of immunocytochemistry as a morphomolecular technique. Cancer Cytopathology, 2016, 124, 540-545.	1.4	14
76	Quantification of HER2 heterogeneity in breast cancerâ€"implications for identification of sub-dominant clones for personalised treatment. Scientific Reports, 2016, 6, 23383.	1.6	38
77	Immune-Derived PD-L1 Gene Expression Defines a Subgroup of Stage II/III Colorectal Cancer Patients with Favorable Prognosis Who May Be Harmed by Adjuvant Chemotherapy. Cancer Immunology Research, 2016, 4, 582-591.	1.6	35
78	Challenging the Cancer Molecular Stratification Dogma: Intratumoral Heterogeneity Undermines Consensus Molecular Subtypes and Potential Diagnostic Value in Colorectal Cancer. Clinical Cancer Research, 2016, 22, 4095-4104.	3.2	135
79	Building a  Repository of Science': The importance ofÂintegrating biobanks within molecular pathology programmes. European Journal of Cancer, 2016, 67, 191-199.	1.3	31
80	Delivering a researchâ€enabled multistakeholder partnership for enhanced patient care at a population level: The Northern Ireland Comprehensive Cancer Program. Cancer, 2016, 122, 664-673.	2.0	5
81	PTEN mRNA detection by chromogenic, RNA in situ technologies: a reliable alternative to PTEN immunohistochemistry. Human Pathology, 2016, 47, 95-103.	1.1	17
82	EphA2 Expression Is a Key Driver of Migration and Invasion and a Poor Prognostic Marker in Colorectal Cancer. Clinical Cancer Research, 2016, 22, 230-242.	3.2	97
83	EORTC-1203: Integration of trastuzumab (T), with or without pertuzumab (P), into perioperative chemotherapy (CT) of HER-2 positive stomach cancerâ€"INNOVATION trial Journal of Clinical Oncology, 2016, 34, TPS4133-TPS4133.	0.8	2
84	GLOBAL BALLAD: An International Rare Cancers Initiative trial to evaluate the potential benefit of adjuvant chemotherapy for small bowel adenocarcinoma (IRCI 002) Journal of Clinical Oncology, 2016, 34, TPS4154-TPS4154.	0.8	9
85	PTEN deficiency promotes macrophage infiltration and hypersensitivity of prostate cancer to IAP antagonist/radiation combination therapy. Oncotarget, 2016, 7, 7885-7898.	0.8	33
86	Gelsolin-mediated activation of PI3K/Akt pathway is crucial for hepatocyte growth factor-induced cell scattering in gastric carcinoma. Oncotarget, 2016, 7, 25391-25407.	0.8	13
87	Stratified analysis reveals chemokine-like factor (CKLF) as a potential prognostic marker in the MSI-immune consensus molecular subtype CMS1 of colorectal cancer. Oncotarget, 2016, 7, 36632-36644.	0.8	15
88	Caspase modelling to predict personalised risk in stage III colorectal cancer (CRC) patients Journal of Clinical Oncology, 2016, 34, 11592-11592.	0.8	0
89	A systems model of BCL-2 dependent apoptosis to predict stage II CRC patients benefiting from adjuvant chemotherapy and as a prognostic tool for stage III CRC patients with increased risk of recurrence Journal of Clinical Oncology, 2016, 34, 3584-3584.	0.8	1
90	Epidermal growth factor receptor immunohistochemistry: new opportunities in metastatic colorectal cancer. Journal of Translational Medicine, 2015, 13, 217.	1.8	36

#	Article	IF	Citations
91	Automated tumor analysis for molecular profiling in lung cancer. Oncotarget, 2015, 6, 27938-27952.	0.8	43
92	RE: Test of Four Colon Cancer Risk-Scores in Formalin Fixed Paraffin Embedded Microarray Gene Expression Data. Journal of the National Cancer Institute, 2015, 107, djv055-djv055.	3.0	0
93	Integrated molecular pathology: the Belfast model. Drug Discovery Today, 2015, 20, 1451-1454.	3.2	6
94	PICan: An integromics framework for dynamic cancer biomarker discovery. Molecular Oncology, 2015, 9, 1234-1240.	2.1	15
95	Immunohistochemistry should undergo robust validation equivalent to that of molecular diagnostics. Journal of Clinical Pathology, 2015, 68, 766-770.	1.0	37
96	Analysis of wntless (WLS) expression in gastric, ovarian, and breast cancers reveals a strong association with HER2 overexpression. Modern Pathology, 2015, 28, 428-436.	2.9	27
97	MErCuRIC1: A Phase I study of MEK1/2 inhibitor PD-0325901 with cMET inhibitor crizotinib in RASMT and RASWT (with aberrant c-MET) metastatic colorectal cancer (mCRC) patients Journal of Clinical Oncology, 2015, 33, TPS3632-TPS3632.	0.8	3
98	The prognostic value of the stem-like group in colorectal cancer using a panel of immunohistochemistry markers. Oncotarget, 2015, 6, 12763-12773.	0.8	14
99	Comprehensive molecular pathology analysis of small bowel adenocarcinoma reveals novel targets with potential for clinical utility. Oncotarget, 2015, 6, 20863-20874.	0.8	41
100	Molecular classification of non-invasive breast lesions for personalised therapy and chemoprevention. Oncotarget, 2015, 6, 43244-43254.	0.8	8
101	Molecular classification of the invasive front in colorectal cancer Journal of Clinical Oncology, 2015, 33, 3573-3573.	0.8	0
102	Abstract 4792: Comprehensive molecular pathology analysis of small bowel adenocarcinoma reveals novel targets with clinical utility., 2015,,.		0
103	Abstract 4018: The role of c-MET/HGF signaling as a critical mediator of an invasive and resistant phenotype in colorectal cancer. , 2015, , .		1
104	Sphingosine Kinase 1 Promotes Malignant Progression in Colon Cancer and Independently Predicts Survival of Patients With Colon Cancer by Competing Risk Approach in South Asian Population. Clinical and Translational Gastroenterology, 2014, 5, e51.	1.3	34
105	Identification and Validation of an Anthracycline/Cyclophosphamide–Based Chemotherapy Response Assay in Breast Cancer. Journal of the National Cancer Institute, 2014, 106, djt335.	3.0	91
106	The prognostic significance of the aberrant extremes of p53 immunophenotypes in breast cancer. Histopathology, 2014, 65, 340-352.	1.6	59
107	Nextâ€generation sequencing: a change of paradigm in molecular diagnostic validation. Journal of Pathology, 2014, 234, 5-10.	2.1	68
108	AXL Is a Key Regulator of Inherent and Chemotherapy-Induced Invasion and Predicts a Poor Clinical Outcome in Early-Stage Colon Cancer. Clinical Cancer Research, 2014, 20, 164-175.	3.2	95

#	Article	IF	CITATIONS
109	BRCA1 Deficiency Exacerbates Estrogen-Induced DNA Damage and Genomic Instability. Cancer Research, 2014, 74, 2773-2784.	0.4	94
110	Digital pathology and image analysis in tissue biomarker research. Methods, 2014, 70, 59-73.	1.9	162
111	Molecular pathology – The value of an integrative approach. Molecular Oncology, 2014, 8, 1163-1168.	2.1	32
112	Identification of a BRCA1-mRNA Splicing Complex Required for Efficient DNA Repair and Maintenance of Genomic Stability. Molecular Cell, 2014, 54, 445-459.	4.5	146
113	Abstract 2079: EpHA2 is an essential driver of invasion and a novel target in KRAS mutant colorectal cancer., 2014,,.		1
114	Association of a DNA damage response deficiency (DDRD) assay and prognosis in early-stage esophageal adenocarcinoma Journal of Clinical Oncology, 2014, 32, 4015-4015.	0.8	3
115	The prognostic and therapeutic value of EpHA2 in early colorectal cancer (CRC) Journal of Clinical Oncology, 2014, 32, 3581-3581.	0.8	0
116	Abstract 1905: Defining a therapeutic classification of breast cancer by actionable targets. , 2014, , .		0
117	Guidelines and considerations for conducting experiments using tissue microarrays. Histopathology, 2013, 62, 827-839.	1.6	57
118	Immunohistochemistry in the era of personalised medicine. Journal of Clinical Pathology, 2013, 66, 58-61.	1.0	40
119	MicroRNA-34c Inversely Couples the Biological Functions of the Runt-related Transcription Factor RUNX2 and the Tumor Suppressor p53 in Osteosarcoma. Journal of Biological Chemistry, 2013, 288, 21307-21319.	1.6	95
120	Identification and validation of an assay predictive of response and prognosis following anthracycline-based chemotherapy for early breast cancer Journal of Clinical Oncology, 2013, 31, TPS11120-TPS11120.	0.8	0
121	Molecular Pathology in Contemporary Diagnostic Pathology Laboratory. American Journal of Surgical Pathology, 2010, 34, 115-117.	2.1	15
122	Elucidating the molecular physiopathology of acute respiratory distress syndrome in severe acute respiratory syndrome patients. Virus Research, 2009, 145, 260-269.	1.1	85
123	Reliability of Tissue Microarrays in Detecting Protein Expression and Gene Amplification in Breast Cancer. Modern Pathology, 2003, 16, 79-85.	2.9	161