Andrea Mafficini

List of Publications by Year in descending order

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79 papers

4,329 citations

147801 31 h-index 110387 64 g-index

80 all docs

80 docs citations

80 times ranked 7196 citing authors

#	Article	IF	CITATIONS
1	Whole-genome landscape of pancreatic neuroendocrine tumours. Nature, 2017, 543, 65-71.	27.8	716
2	Exome sequencing identifies frequent inactivating mutations in BAP1, ARID1A and PBRM1 in intrahepatic cholangiocarcinomas. Nature Genetics, 2013, 45, 1470-1473.	21.4	564
3	Targeted nextâ€generation sequencing of cancer genes dissects the molecular profiles of intraductal papillary neoplasms of the pancreas. Journal of Pathology, 2014, 233, 217-227.	4.5	308
4	DNA Qualification Workflow for Next Generation Sequencing of Histopathological Samples. PLoS ONE, 2013, 8, e62692.	2.5	209
5	Lung neuroendocrine tumours: deep sequencing of the four World Health Organization histotypes reveals chromatinâ€remodelling genes as major players and a prognostic role for <i><scp>TERT</scp></i> , <i><scp>RB1</scp></i> , <i><scp>MEN1</scp></i> and <scp><i>KMT2D</i></scp> . lournal of Pathology, 2017, 241, 488-500.	4.5	179
6	Multigene mutational profiling of cholangiocarcinomas identifies actionable molecular subgroups. Oncotarget, 2014, 5, 2839-2852.	1.8	171
7	Genetics and Epigenetics of Gastroenteropancreatic Neuroendocrine Neoplasms. Endocrine Reviews, 2019, 40, 506-536.	20.1	146
8	A Cross-Species Analysis in Pancreatic Neuroendocrine Tumors Reveals Molecular Subtypes with Distinctive Clinical, Metastatic, Developmental, and Metabolic Characteristics. Cancer Discovery, 2015, 5, 1296-1313.	9.4	145
9	Identification of proteins released by pancreatic cancer cells by multidimensional protein identification technology: a strategy for identification of novel cancer markers. FASEB Journal, 2005, 19, 1125-1127.	0.5	122
10	Mixed Adenoneuroendocrine Carcinomas of the Gastrointestinal Tract: Targeted Next-Generation Sequencing Suggests a Monoclonal Origin of the Two Components. Neuroendocrinology, 2014, 100, 310-316.	2.5	115
11	Molecular Typing of Lung Adenocarcinoma on Cytological Samples Using a Multigene Next Generation Sequencing Panel. PLoS ONE, 2013, 8, e80478.	2.5	96
12	Pancreatic undifferentiated carcinoma with osteoclastâ€like giant cells is genetically similar to, but clinically distinct from, conventional ductal adenocarcinoma. Journal of Pathology, 2017, 243, 148-154.	4.5	79
13	Genomic landscape of pancreatic neuroendocrine tumours: the International Cancer Genome Consortium. Journal of Endocrinology, 2018, 236, R161-R167.	2.6	79
14	Gene Expression Profiling of Lung Atypical Carcinoids and Large Cell Neuroendocrine Carcinomas Identifies Three Transcriptomic Subtypes with Specific Genomic Alterations. Journal of Thoracic Oncology, 2019, 14, 1651-1661.	1.1	73
15	Tumor Mutational Burden as a Potential Biomarker for Immunotherapy in Pancreatic Cancer: Systematic Review and Still-Open Questions. Cancers, 2021, 13, 3119.	3.7	69
16	BRCA somatic and germline mutation detection in paraffin embedded ovarian cancers by next-generation sequencing. Oncotarget, 2016, 7, 1076-1083.	1.8	68
17	High-throughput mutation profiling improves diagnostic stratification of sporadic medullary thyroid carcinomas. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2014, 465, 73-78.	2.8	66
18	Most high-grade neuroendocrine tumours of the lung are likely to secondarily develop from pre-existing carcinoids: innovative findings skipping the current pathogenesis paradigm. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2018, 472, 567-577.	2.8	64

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19	Non-functional pancreatic neuroendocrine tumours: ATRX/DAXX and alternative lengthening of telomeres (ALT) are prognostically independent from ARX/PDX1 expression and tumour size. Gut, 2022, 71, 961-973.	12.1	60
20	High-throughput mutation profiling identifies novel molecular dysregulation in high-grade intraepithelial neoplasia and early gastric cancers. Gastric Cancer, 2014, 17, 442-449.	5. 3	52
21	Development of a semi-conductor sequencing-based panel for genotyping of colon and lung cancer by the Onconetwork consortium. BMC Cancer, 2015, 15, 26.	2.6	49
22	Unmasking the impact of Rictor in cancer: novel insights of mTORC2 complex. Carcinogenesis, 2018, 39, 971-980.	2.8	48
23	Next-Generation Histopathologic Diagnosis: A Lesson From a Hepatic Carcinosarcoma. Journal of Clinical Oncology, 2014, 32, e63-e66.	1.6	47
24	Protein Tyrosine Phosphatase Receptor Type \hat{I}^3 Is a Functional Tumor Suppressor Gene Specifically Downregulated in Chronic Myeloid Leukemia. Cancer Research, 2010, 70, 8896-8906.	0.9	46
25	PD-1, PD-L1, and CD163 in pancreatic undifferentiated carcinoma with osteoclast-like giant cells: expression patterns and clinical implications. Human Pathology, 2018, 81, 157-165.	2.0	44
26	Mutational and copy number asset of primary sporadic neuroendocrine tumors of the small intestine. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2018, 473, 709-717.	2.8	40
27	Genetic alterations analysis in prognostic stratified groups identified TP53 and ARID1A as poor clinical performance markers in intrahepatic cholangiocarcinoma. Scientific Reports, 2018, 8, 7119.	3.3	39
28	Elevated urinary levels of urokinase-type plasminogen activator receptor (uPAR) in pancreatic ductal adenocarcinoma identify a clinically high-risk group. BMC Cancer, 2011, 11, 448.	2.6	35
29	Ampulla of Vater Carcinoma. Annals of Surgery, 2018, 267, 149-156.	4.2	35
30	Reporting Tumor Molecular Heterogeneity in Histopathological Diagnosis. PLoS ONE, 2014, 9, e104979.	2.5	35
31	Splice variants as novel targets in pancreatic ductal adenocarcinoma. Scientific Reports, 2017, 7, 2980.	3.3	34
32	Molecular alterations associated with metastases of solid pseudopapillary neoplasms of the pancreas. Journal of Pathology, 2019, 247, 123-134.	4.5	32
33	Next-generation sequencing for genetic testing of familial colorectal cancer syndromes. Hereditary Cancer in Clinical Practice, 2015, 13, 18.	1.5	31
34	Expression of transmembrane protein tyrosine phosphatase gamma (PTP?) in normal and neoplastic human tissues. Histopathology, 2007, 50, 615-628.	2.9	28
35	DNA methylation patterns identify subgroups of pancreatic neuroendocrine tumors with clinical association. Communications Biology, 2021, 4, 155.	4.4	26
36	Carbon dating cancer: defining the chronology of metastatic progression in colorectal cancer. Annals of Oncology, 2017, 28, 1243-1249.	1.2	25

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37	Immune landscape, evolution, hypoxia-mediated viral mimicry pathways and therapeutic potential in molecular subtypes of pancreatic neuroendocrine tumours. Gut, 2021, 70, 1904-1913.	12.1	24
38	CD71 in Gestational Pathology. Applied Immunohistochemistry and Molecular Morphology, 2016, 24, 215-220.	1.2	23
39	Ultra-Mutation in IDH Wild-Type Glioblastomas of Patients Younger than 55 Years is Associated with Defective Mismatch Repair, Microsatellite Instability, and Giant Cell Enrichment. Cancers, 2019, 11, 1279.	3.7	23
40	Simultaneous detection of lung fusions using a multiplex RT-PCR next generation sequencing-based approach: a multi-institutional research study. BMC Cancer, 2018, 18, 828.	2.6	19
41	Specific expression patterns of epithelial to mesenchymal transition factors in gestational molar disease. Placenta, 2015, 36, 1318-1324.	1.5	18
42	A new monoclonal antibody detects downregulation of protein tyrosine phosphatase receptor type \hat{l}^3 in chronic myeloid leukemia patients. Journal of Hematology and Oncology, 2017, 10, 129.	17.0	17
43	Ki-67 assessment of pancreatic neuroendocrine neoplasms: Systematic review and meta-analysis of manual vs. digital pathology scoring. Modern Pathology, 2022, 35, 712-720.	5.5	17
44	Receptor-type protein tyrosine phosphatase gamma (PTP \hat{I}^3), a new identifier for myeloid dendritic cells and specialized macrophages. Blood, 2006, 108, 4223-4231.	1.4	16
45	Solid Pseudopapillary Neoplasm of the Pancreas and Abdominal Desmoid Tumor in a Patient Carrying Two Different BRCA2 Germline Mutations: New Horizons from Tumor Molecular Profiling. Genes, 2021, 12, 481.	2.4	13
46	Molecular characterization of extrahepatic cholangiocarcinoma: perihilar and distal tumors display divergent genomic and transcriptomic profiles. Expert Opinion on Therapeutic Targets, 2021, 25, 1095-1105.	3.4	13
47	Histo-molecular characterization of pancreatic cancer with microsatellite instability: intra-tumor heterogeneity, B2M inactivation, and the importance of metastatic sites. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2022, 480, 1261-1268.	2.8	12
48	Perineural Invasion is a Strong Prognostic Moderator in Ampulla of Vater Carcinoma. Pancreas, 2019, 48, 70-76.	1.1	11
49	Gene Expression Profiling of Pancreas Neuroendocrine Tumors with Different Ki67-Based Grades. Cancers, 2021, 13, 2054.	3.7	10
50	IDH-wild type glioblastomas featuring at least 30% giant cells are characterized by frequent RB1 and NF1 alterations and hypermutation. Acta Neuropathologica Communications, 2021, 9, 200.	5.2	10
51	Protein Tyrosine Phosphatase Gamma (PTPγ) is a Novel Leukocyte Marker Highly Expressed by CD34+ Precursors. Biomarker Insights, 2007, 2, 117727190700200.	2.5	9
52	Genomic characterization of hepatoid tumors: context matters. Human Pathology, 2021, 118, 30-41.	2.0	9
53	Both HIV- and EIAV-based lentiviral vectors mediate gene delivery to pancreatic cancer cells and human pancreatic primary patient xenografts. Cancer Gene Therapy, 2007, 14, 781-790.	4.6	8
54	Immunohistochemical detection of arginine methylated proteins (MeRP) in archival tissues. Histopathology, 2010, 57, 725-733.	2.9	7

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55	Impact of polymorphism of Multidrug Resistance-associated Protein 1 (ABCC1) gene on the severity of cystic fibrosis. Journal of Cystic Fibrosis, 2011, 10, 228-233.	0.7	7
56	Impact of MIF Gene Promoter Polymorphism on F508del Cystic Fibrosis Patients. PLoS ONE, 2014, 9, e114274.	2.5	7
57	Colorectal cancer with microsatellite instability: Right-sided location and signet ring cell histology are associated with nodal metastases, and extranodal extension influences disease-free survival. Pathology Research and Practice, 2021, 224, 153519.	2.3	7
58	Protein Tyrosine Phosphatase Gamma (PTPgamma) is a Novel Leukocyte Marker Highly Expressed by CD34 Precursors. Biomarker Insights, 2007, 2, 218-25.	2.5	7
59	ICAT is a novel Ptf1a interactor that regulates pancreatic acinar differentiation and displays altered expression in tumours. Biochemical Journal, 2013, 451, 395-405.	3.7	6
60	"Pure―hepatoid tumors of the pancreas harboring CTNNB1 somatic mutations: a new entity among solid pseudopapillary neoplasms. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2022, 481, 41-47.	2.8	6
61	Genomic characterization of undifferentiated sarcomatoid carcinoma of the pancreas. Human Pathology, 2022, 128, 124-133.	2.0	6
62	New genomic landscapes and therapeutic targets for biliary tract cancers. Frontiers in Bioscience - Landmark, 2016, 21, 707-718.	3.0	5
63	Fhit down-regulation is an early event in pancreatic carcinogenesis. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2017, 470, 647-653.	2.8	5
64	Molecular Analysis of an Intestinal Neuroendocrine/Non-neuroendocrine Neoplasm (MiNEN) Reveals MLH1 Methylation-driven Microsatellite Instability and a Monoclonal Origin: Diagnostic and Clinical Implications. Applied Immunohistochemistry and Molecular Morphology, 2022, 30, 145-152.	1,2	5
65	Refining targeted therapeutic approaches in pancreatic cancer: from histology and molecular pathology to the clinic. Expert Opinion on Therapeutic Targets, 2022, 26, 1-4.	3.4	5
66	OA06.06 Druggable Alterations Involving Crucial Carcinogenesis Pathways Drive the Prognosis of Squamous Cell Lung Carcinoma (SqCLC). Journal of Thoracic Oncology, 2017, 12, S266-S267.	1.1	4
67	Abstract 3575: The OncoNetwork Consortium: A global collaborative research study on the development and verification of an Ion AmpliSeq RNA gene lung fusion panel. Cancer Research, 2014, 74, 3575-3575.	0.9	4
68	Non-coding regulatory variations: the dark matter of pancreatic cancer genomics. Gut, 2018, 67, 399-400.	12.1	3
69	Juvenile polyposis diagnosed with an integrated histological, immunohistochemical and molecular approach identifying new SMAD4 pathogenic variants. Familial Cancer, 2022, 21, 441-451.	1.9	3
70	Recurrent oligodendroglioma with changed 1p/19q status. Neuropathology, 2022, , .	1.2	3
71	ERG alterations and mTOR pathway activation in primary prostate carcinomas developing castration-resistance. Pathology Research and Practice, 2018, 214, 1675-1680.	2.3	1
72	Pancreatic Cancer Genomics., 2013,, 219-253.		1

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73	P2.04-12 A Genomic Signature [JAK2, JAK3, PIAS4, PTPN2, STAT3, IFNAR2] Predicts Baseline Resistance to Nivolumab in Advanced NSCLC Journal of Thoracic Oncology, 2018, 13, S734-S735.	1.1	O
74	Validation of a tumour mutational burden workflow on routine histological samples of colorectal cancer and assessment of a cohort with synchronous hepatic metastases. Annals of Oncology, 2019, 30, ν 574.	1.2	0
75	Comparative Lesions Analysis Through a Targeted Sequencing Approach. Journal of Visualized Experiments, 2019, , .	0.3	O
76	P2.04-51 A 6-Gene Immune Genomic Signature (IGS) Predicts Resistance to Nivolumab [NIV] in Advanced Pretreated NSCLC: Results of PRINCiPe Trial. Journal of Thoracic Oncology, 2019, 14, S728.	1.1	0
77	Molecular Biology of Neuroendocrine Tumors. , 2021, , 37-53.		O
78	Abstract 4891: Comprehensive genetic profiling of chromosomal translocations in lung cancer tumors: development and validation of a next-generation sequencing panel in an international multicenter study. , 2015, , .		0
79	Abstract 5694: Multi institutional evaluation of a new NGS assay for mutation detection from cfDNA in lung cancer. , 2017 , , .		0