

# Rosa Maria Pascale

## List of Publications by Year in descending order

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77  
papers

3,143  
citations

147801

31  
h-index

168389

53  
g-index

77  
all docs

77  
docs citations

77  
times ranked

4424  
citing authors

#	ARTICLE	IF	CITATIONS
1	Nuclear localization dictates hepatocarcinogenesis suppression by glycine N-methyltransferase. <i>Translational Oncology</i> , 2022, 15, 101239.	3.7	4
2	S-Adenosylmethionine: From the Discovery of Its Inhibition of Tumorigenesis to Its Use as a Therapeutic Agent. <i>Cells</i> , 2022, 11, 409.	4.1	18
3	Blood Cell Count Indexes of Systemic Inflammation in Carotid Artery Disease: Current Evidence and Future Perspectives. <i>Current Pharmaceutical Design</i> , 2021, 27, 2170-2179.	1.9	9
4	HCC in the Era of Direct-Acting Antiviral Agents (DAAs): Surgical and Other Curative or Palliative Strategies in the Elderly. <i>Cancers</i> , 2021, 13, 3025.	3.7	6
5	Cholesterol biosynthesis supports the growth of hepatocarcinoma lesions depleted of fatty acid synthase in mice and humans. <i>Gut</i> , 2020, 69, 177-186.	12.1	121
6	The Warburg Effect 97 Years after Its Discovery. <i>Cancers</i> , 2020, 12, 2819.	3.7	153
7	Laboratory test alterations in patients with COVID-19 and non COVID-19 interstitial pneumonia: a preliminary report. <i>Journal of Infection in Developing Countries</i> , 2020, 14, 685-690.	1.2	47
8	Combined CDK4/6 and Pan-mTOR Inhibition Is Synergistic Against Intrahepatic Cholangiocarcinoma. <i>Clinical Cancer Research</i> , 2019, 25, 403-413.	7.0	56
9	SNAIL Promotes the Cholangiocellular Phenotype, but not Epithelialâ€Mesenchymal Transition, in a Murine Hepatocellular Carcinoma Model. <i>Cancer Research</i> , 2019, 79, 5563-5574.	0.9	12
10	Experimental Models to Define the Genetic Predisposition to Liver Cancer. <i>Cancers</i> , 2019, 11, 1450.	3.7	15
11	Post-translational inhibition of YAP oncogene expression by 4-hydroxynonenal in bladder cancer cells. <i>Free Radical Biology and Medicine</i> , 2019, 141, 205-219.	2.9	13
12	Alterations of Methionine Metabolism as Potential Targets for the Prevention and Therapy of Hepatocellular Carcinoma. <i>Medicina (Lithuania)</i> , 2019, 55, 296.	2.0	29
13	The mTORC2â€Akt1 Cascade Is Crucial for c-Myc to Promote Hepatocarcinogenesis in Mice and Humans. <i>Hepatology</i> , 2019, 70, 1600-1613.	7.3	70
14	Functional role of SGK3 in PI3K/Pten driven liver tumor development. <i>BMC Cancer</i> , 2019, 19, 343.	2.6	17
15	TEA Domain Transcription Factor 4 Is the Major Mediator of Yes-Associated Protein Oncogenic Activity in Mouse and Human Hepatoblastoma. <i>American Journal of Pathology</i> , 2019, 189, 1077-1090.	3.8	25
16	MicroRNA-203 impacts on the growth, aggressiveness and prognosis of hepatocellular carcinoma by targeting <i>MAT2A</i> and <i>MAT2B</i> genes. <i>Oncotarget</i> , 2019, 10, 2835-2854.	1.8	18
17	Deregulation of methionine metabolism as determinant of progression and prognosis of hepatocellular carcinoma. <i>Translational Gastroenterology and Hepatology</i> , 2018, 3, 36-36.	3.0	23
18	Alterations of methionine metabolism in hepatocarcinogenesis: the emergent role of glycine N-methyltransferase in liver injury. <i>Annals of Gastroenterology</i> , 2018, 31, 552-560.	0.6	15

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19	Focal adhesion kinase activation limits efficacy of Dasatinib in c-Myc driven hepatocellular carcinoma. <i>Cancer Medicine</i> , 2018, 7, 6170-6181.	2.8	11
20	The complex role of bone morphogenetic protein 9 in liver damage and regeneration: New evidence from in vivo and in vitro studies. <i>Liver International</i> , 2018, 38, 1547-1549.	3.9	2
21	Oncogene-dependent addiction to carbohydrate-responsive element binding protein in hepatocellular carcinoma. <i>Cell Cycle</i> , 2018, 17, 1496-1512.	2.6	14
22	A functional mammalian target of rapamycin complex 1 signaling is indispensable for c-Myc driven hepatocarcinogenesis. <i>Hepatology</i> , 2017, 66, 167-181.	7.3	119
23	Pan-mTOR inhibitor MLN0128 is effective against intrahepatic cholangiocarcinoma in mice. <i>Journal of Hepatology</i> , 2017, 67, 1194-1203.	3.7	77
24	Deregulated c-Myc requires a functional HSF1 for experimental and human hepatocarcinogenesis. <i>Oncotarget</i> , 2017, 8, 90638-90650.	1.8	17
25	Inhibition of HSF1 suppresses the growth of hepatocarcinoma cell lines <i>in vitro</i> and AKT-driven hepatocarcinogenesis in mice. <i>Oncotarget</i> , 2017, 8, 54149-54159.	1.8	24
26	Focal loss of long non-coding RNA-PRAL, as determinant of cell function and phenotype of hepatocellular carcinoma. <i>Annals of Translational Medicine</i> , 2016, 4, 183-183.	1.7	5
27	Sulfatase 1: a new Jekyll and Hyde in hepatocellular carcinoma?. <i>Translational Gastroenterology and Hepatology</i> , 2016, 1, 43-43.	3.0	3
28	An infernal cross-talk between oncogenic $\beta$ -catenin and c-Met in hepatocellular carcinoma: Evidence from mouse modeling. <i>Hepatology</i> , 2016, 64, 1421-1423.	7.3	5
29	DNA-PKcs: A promising therapeutic target in human hepatocellular carcinoma?. <i>DNA Repair</i> , 2016, 47, 12-20.	2.8	25
30	Inactivation of fatty acid synthase impairs hepatocarcinogenesis driven by AKT in mice and humans. <i>Journal of Hepatology</i> , 2016, 64, 333-341.	3.7	115
31	Post-translational deregulation of YAP1 is genetically controlled in rat liver cancer and determines the fate and stem-like behavior of the human disease. <i>Oncotarget</i> , 2016, 7, 49194-49216.	1.8	20
32	Epidermal growth factor-like repeats and discoidin I-like domains 3: a multifaceted oncoprotein at the crossroad of MAPK and TGF-beta pathways in human hepatocellular carcinoma. <i>Translational Cancer Research</i> , 2016, 5, 103-109.	1.0	2
33	Phenotype: the result of a complex and still largely unknown interplay between molecules. <i>Translational Cancer Research</i> , 2016, 5, S895-S897.	1.0	0
34	Multifocal hepatocellular carcinoma: intrahepatic metastasis or multicentric carcinogenesis?. <i>Annals of Translational Medicine</i> , 2015, 3, 4.	1.7	39
35	New insights on the role of epigenetic alterations in hepatocellular carcinoma. <i>Journal of Hepatocellular Carcinoma</i> , 2014, 1, 65.	3.7	4
36	Pleiotropic effects of methionine adenosyltransferases deregulation as determinants of liver cancer progression and prognosis. <i>Journal of Hepatology</i> , 2013, 59, 830-841.	3.7	117

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37	Regulation of Amphiregulin Gene Expression by $\beta$ -Catenin Signaling in Human Hepatocellular Carcinoma Cells: A Novel Crosstalk between FGF19 and the EGFR System. <i>PLoS ONE</i> , 2012, 7, e52711.	2.5	45
38	Role of transcriptional and posttranscriptional regulation of methionine adenosyltransferases in liver cancer progression. <i>Hepatology</i> , 2012, 56, 165-175.	7.3	73
39	<i>S</i> -adenosyl methionine regulates ubiquitin-conjugating enzyme 9 protein expression and sumoylation in murine liver and human cancers. <i>Hepatology</i> , 2012, 56, 982-993.	7.3	70
40	An expression signature of phenotypic resistance to hepatocellular carcinoma identified by cross-species gene expression analysis. <i>Cellular Oncology (Dordrecht)</i> , 2012, 35, 163-173.	4.4	30
41	Deregulation of signalling pathways in prognostic subtypes of hepatocellular carcinoma: Novel insights from interspecies comparison. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2012, 1826, 215-237.	7.4	27
42	Mybl2 expression is under genetic control and contributes to determine a hepatocellular carcinoma susceptible phenotype. <i>Journal of Hepatology</i> , 2011, 55, 111-119.	3.7	36
43	Connective tissue growth factor autocriny in human hepatocellular carcinoma: Oncogenic role and regulation by epidermal growth factor receptor/yes-associated protein-mediated activation. <i>Hepatology</i> , 2011, 54, 2149-2158.	7.3	108
44	Activation of v-Myb avian myeloblastosis viral oncogene homolog-like2 (MYBL2)-LIN9 complex contributes to human hepatocarcinogenesis and identifies a subset of hepatocellular carcinoma with mutant p53. <i>Hepatology</i> , 2011, 53, 1226-1236.	7.3	53
45	The degradation of cell cycle regulators by SKP2/CKS1 ubiquitin ligase is genetically controlled in rodent liver cancer and contributes to determine the susceptibility to the disease. <i>International Journal of Cancer</i> , 2010, 126, 1275-1281.	5.1	19
46	Prognostic markers and putative therapeutic targets for hepatocellular carcinoma. <i>Molecular Aspects of Medicine</i> , 2010, 31, 179-193.	6.4	75
47	Genetic and Epigenetic Control of Molecular Alterations in Hepatocellular Carcinoma. <i>Experimental Biology and Medicine</i> , 2009, 234, 726-736.	2.4	46
48	SKP2 and CKS1 Promote Degradation of Cell Cycle Regulators and Are Associated With Hepatocellular Carcinoma Prognosis. <i>Gastroenterology</i> , 2009, 137, 1816-1826.e10.	1.3	95
49	Ras-driven proliferation and apoptosis signaling during rat liver carcinogenesis is under genetic control. <i>International Journal of Cancer</i> , 2008, 123, 2057-2064.	5.1	38
50	Genetic control of resistance to hepatocarcinogenesis by the mouse Hpcr3 locus. <i>Hepatology</i> , 2008, 48, 617-623.	7.3	16
51	Dual-Specificity Phosphatase 1 Ubiquitination in Extracellular Signal-Regulated Kinase-Mediated Control of Growth in Human Hepatocellular Carcinoma. <i>Cancer Research</i> , 2008, 68, 4192-4200.	0.9	107
52	Aberrant iNOS signaling is under genetic control in rodent liver cancer and potentially prognostic for the human disease. <i>Carcinogenesis</i> , 2008, 29, 1639-1647.	2.8	54
53	Interaction of major genes predisposing to hepatocellular carcinoma with genes encoding signal transduction pathways influences tumor phenotype and prognosis. <i>World Journal of Gastroenterology</i> , 2008, 14, 6601.	3.3	12
54	Identification and chromosome mapping of loci predisposing to colorectal cancer that control Wnt/ $\beta$ -catenin pathway and progression of early lesions in the rat. <i>Carcinogenesis</i> , 2007, 28, 2367-2374.	2.8	10

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55	Dissection of Signal Transduction Pathways as a Tool for the Development of Targeted Therapies of Hepatocellular Carcinoma. <i>Reviews on Recent Clinical Trials</i> , 2007, 2, 217-236.	0.8	43
56	Altered methionine metabolism and global DNA methylation in liver cancer: Relationship with genomic instability and prognosis. <i>International Journal of Cancer</i> , 2007, 121, 2410-2420.	5.1	73
57	Mapping a Sex Hormone- Sensitive Gene Determining Female Resistance to Liver Carcinogenesis in a Congenic F344.BN-Hcs4 Rat. <i>Cancer Research</i> , 2006, 66, 10384-10390.	0.9	10
58	Role of HSP90, CDC37, and CRM1 as modulators of P16INK4A activity in rat liver carcinogenesis and human liver cancer. <i>Hepatology</i> , 2005, 42, 1310-1319.	7.3	87
59	Chemopreventive N-(4-hydroxyphenyl)retinamide (fenretinide) targets deregulated NF- $\kappa$ B and Mat1A genes in the early stages of rat liver carcinogenesis. <i>Carcinogenesis</i> , 2004, 26, 417-427.	2.8	28
60	Polygenic control of hepatocarcinogenesis in Copenhagen $\tilde{\text{A}}$ - F344 rats. <i>International Journal of Cancer</i> , 2004, 111, 9-16.	5.1	26
61	Phenotypic reversion of rat neoplastic liver nodules is under genetic control. <i>International Journal of Cancer</i> , 2003, 105, 70-75.	5.1	16
62	Down-regulation of c-myc and Cyclin D1 genes by antisense oligodeoxy nucleotides inhibits the expression of E2F1 and in vitro growth of HepG2 and Morris 5123 liver cancer cells. <i>Carcinogenesis</i> , 2003, 25, 333-341.	2.8	59
63	Stearoyl-CoA desaturase 1 (Scd1) gene overexpression is associated with genetic predisposition to hepatocarcinogenesis in mice and rats. <i>Carcinogenesis</i> , 2002, 23, 1933-1936.	2.8	81
64	Chemoprevention of hepatocarcinogenesis. <i>Alcohol</i> , 2002, 27, 193-198.	1.7	95
65	Cell cycle deregulation in liver lesions of rats with and without genetic predisposition to hepatocarcinogenesis. <i>Hepatology</i> , 2002, 35, 1341-1350.	7.3	63
66	Chromosome mapping of multiple loci affecting the genetic predisposition to rat liver carcinogenesis. <i>Cancer Research</i> , 2002, 62, 4459-63.	0.9	17
67	5 $\alpha$ -Methylthioadenosine administration prevents lipid peroxidation and fibrogenesis induced in rat liver by carbon-tetrachloride intoxication. <i>Journal of Hepatology</i> , 2001, 34, 386-394.	3.7	93
68	Frequent loss of heterozygosity at the Hcr1 (hepatocarcinogenesis resistance) locus on chromosome 10 in primary hepatocellular carcinomas from LFF1 rat strain. <i>Hepatology</i> , 2001, 33, 1110-1117.	7.3	18
69	Implication of Bcl-2 family genes in basal and d-amphetamine-induced apoptosis in preneoplastic and neoplastic rat liver lesions. <i>Hepatology</i> , 2000, 31, 956-965.	7.3	12
70	Genetic Alterations in Liver Carcinogenesis: Implications for New Preventive and Therapeutic Strategies. <i>Critical Reviews in Oncogenesis</i> , 2000, 11, 44.	0.4	65
71	Correlation of c-myc overexpression and amplification with progression of preneoplastic liver lesions to malignancy in the poorly susceptible wistar rat strain. , 1999, 25, 21-29.		34
72	Transferrin and transferrin receptor gene expression and iron uptake in hepatocellular carcinoma in the rat. <i>Hepatology</i> , 1998, 27, 452-461.	7.3	22

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73	c-myc amplification in pre-malignant and malignant lesions induced in rat liver by the resistant hepatocyte model. , 1996, 68, 136-142.		29
74	Persistent chemopreventive effect of S-adenosyl-L-methionine on the development of liver putative preneoplastic lesions induced by thiobenzamide in diethylnitrosamine-initiated rats. Carcinogenesis, 1996, 17, 1533-1537.	2.8	24
75	SHORT COMMUNICATION: The BN rat strain carries dominant hepatocarcinogen resistance loci. Carcinogenesis, 1996, 17, 1765-1768.	2.8	27
76	Inhibition of 3-hydroxy-3-methylglutaryl-CoA reductase activity and gene expression by dehydrocpiandrosterone in preneoplastic liver nodules. Carcinogenesis, 1995, 16, 1537-1542.	2.8	31
77	Alterations of ornithine decarboxylase gene during the progression of rat liver carcinogenesis. Carcinogenesis, 1993, 14, 1077-1080.	2.8	26