

# Christos K Kontos

## List of Publications by Year in descending order

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119  
papers

2,481  
citations

172457

29  
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276875

41  
g-index

123  
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123  
docs citations

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times ranked

3143  
citing authors

#	ARTICLE	IF	CITATIONS
1	Evaluation of PD-L1 Expression and Associated Tumor-Infiltrating Lymphocytes in Laryngeal Squamous Cell Carcinoma. <i>Clinical Cancer Research</i> , 2016, 22, 704-713.	7.0	173
2	<i>Phosphatidylinositol 3-Kinase Catalytic Subunit 1</i> Gene Amplification Contributes to the Pathogenesis of Mantle Cell Lymphoma. <i>Clinical Cancer Research</i> , 2009, 15, 5724-5732.	7.0	99
3	Epigenetic regulation of miR-21 in colorectal cancer. <i>Epigenetics</i> , 2014, 9, 129-141.	2.7	98
4	Apoptosis-related BCL2-family Members: Key Players in Chemotherapy. <i>Anti-Cancer Agents in Medicinal Chemistry</i> , 2014, 14, 353-374.	1.7	85
5	Kallikrein-related peptidases (KLKs): a gene family of novel cancer biomarkers. <i>Clinical Chemistry and Laboratory Medicine</i> , 2012, 50, 1877-1891.	2.3	74
6	Quantitative expression analysis and prognostic significance of L-DOPA decarboxylase in colorectal adenocarcinoma. <i>British Journal of Cancer</i> , 2010, 102, 1384-1390.	6.4	50
7	High miR-96 levels in colorectal adenocarcinoma predict poor prognosis, particularly in patients without distant metastasis at the time of initial diagnosis. <i>Tumor Biology</i> , 2016, 37, 11815-11824.	1.8	44
8	Circular RNAs: A New Piece in the Colorectal Cancer Puzzle. <i>Cancers</i> , 2020, 12, 2464.	3.7	42
9	Circular RNAs: Emerging Regulators of the Major Signaling Pathways Involved in Cancer Progression. <i>Cancers</i> , 2021, 13, 2744.	3.7	42
10	Quantitative expression analysis and prognostic significance of the novel apoptosis-related gene <i>BCL2L12</i> in colon cancer. <i>Biological Chemistry</i> , 2008, 389, 1467-1475.	2.5	40
11	miR-15a-5p, A Novel Prognostic Biomarker, Predicting Recurrent Colorectal Adenocarcinoma. <i>Molecular Diagnosis and Therapy</i> , 2017, 21, 453-464.	3.8	40
12	BCL2L12 is a Novel Biomarker for the Prediction of Short-Term Relapse in Nasopharyngeal Carcinoma. <i>Molecular Medicine</i> , 2011, 17, 163-171.	4.4	39
13	The Novel Member of the <i>BCL2</i> Gene Family, <i>BCL2L12</i> , Is Substantially Elevated in Chronic Lymphocytic Leukemia Patients, Supporting Its Value As a Significant Biomarker. <i>Oncologist</i> , 2011, 16, 1280-1291.	3.7	39
14	Enhanced miR-182 transcription is a predictor of poor overall survival in colorectal adenocarcinoma patients. <i>Clinical Chemistry and Laboratory Medicine</i> , 2014, 52, 1217-27.	2.3	39
15	miR-224 overexpression is a strong and independent prognosticator of short-term relapse and poor overall survival in colorectal adenocarcinoma. <i>International Journal of Oncology</i> , 2015, 46, 849-859.	3.3	38
16	Kallikrein-related peptidases (KLKs) in gastrointestinal cancer: Mechanistic and clinical aspects. <i>Thrombosis and Haemostasis</i> , 2013, 110, 450-457.	3.4	37
17	Myocyte Damage and Loss of Myofibers is the Potential Mechanism of Iron Overload Toxicity in Congestive Cardiac Failure in Thalassemia. Complete Reversal of the Cardiomyopathy and Normalization of Iron Load by Deferiprone. <i>Hemoglobin</i> , 2008, 32, 17-28.	0.8	35
18	The role of circular RNAs in therapy resistance of patients with solid tumors. <i>Personalized Medicine</i> , 2020, 17, 469-490.	1.5	35

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19	Kallikrein-related peptidase-6 (KLK6) mRNA expression is an independent prognostic tissue biomarker of poor disease-free and overall survival in colorectal adenocarcinoma. <i>Tumor Biology</i> , 2014, 35, 4673-4685.	1.8	34
20	22-gauge core vs 22-gauge aspiration needle for endoscopic ultrasound-guided sampling of abdominal masses. <i>World Journal of Gastroenterology</i> , 2016, 22, 8820.	3.3	34
21	Upregulated miR-16 expression is an independent indicator of relapse and poor overall survival of colorectal adenocarcinoma patients. <i>Clinical Chemistry and Laboratory Medicine</i> , 2017, 55, 737-747.	2.3	33
22	Nature Promises New Anticancer Agents: Interplay with the Apoptosis-related BCL2 Gene Family. <i>Anti-Cancer Agents in Medicinal Chemistry</i> , 2014, 14, 375-399.	1.7	33
23	High BAX/BCL2 mRNA ratio predicts favorable prognosis in laryngeal squamous cell carcinoma, particularly in patients with negative lymph nodes at the time of diagnosis. <i>Clinical Biochemistry</i> , 2016, 49, 890-896.	1.9	32
24	MicroRNA-155-5p Overexpression in Peripheral Blood Mononuclear Cells of Chronic Lymphocytic Leukemia Patients Is a Novel, Independent Molecular Biomarker of Poor Prognosis. <i>Disease Markers</i> , 2017, 2017, 1-10.	1.3	32
25	Expression analysis of mir-17-5p, mir-20a and let-7a microRNAs and their target proteins in CD34+ bone marrow cells of patients with myelodysplastic syndromes. <i>Leukemia Research</i> , 2013, 37, 251-258.	0.8	31
26	Elevated expression of miR-24-3p is a potentially adverse prognostic factor in colorectal adenocarcinoma. <i>Clinical Biochemistry</i> , 2017, 50, 285-292.	1.9	31
27	MicroRNAs: Tiny Regulators of Gene Expression with Pivotal Roles in Normal B-Cell Development and B-Cell Chronic Lymphocytic Leukemia. <i>Cancers</i> , 2021, 13, 593.	3.7	31
28	Molecular cloning of novel alternatively spliced variants of BCL2L12, a new member of the BCL2 gene family, and their expression analysis in cancer cells. <i>Gene</i> , 2012, 505, 153-166.	2.2	30
29	Kallikrein-related peptidase 4 (KLK4) mRNA predicts short-term relapse in colorectal adenocarcinoma patients. <i>Cancer Letters</i> , 2013, 330, 106-112.	7.2	30
30	KLK11 mRNA expression predicts poor disease-free and overall survival in colorectal adenocarcinoma patients. <i>Biomarkers in Medicine</i> , 2014, 8, 671-685.	1.4	30
31	Prognostic and predictive biomarkers in prostate cancer. <i>Expert Review of Molecular Diagnostics</i> , 2015, 15, 1567-1576.	3.1	29
32	Elevated miR-20b-5p expression in peripheral blood mononuclear cells: A novel, independent molecular biomarker of favorable prognosis in chronic lymphocytic leukemia. <i>Leukemia Research</i> , 2018, 70, 1-7.	0.8	29
33	High microRNA-28-5p expression in colorectal adenocarcinoma predicts short-term relapse of node-negative patients and poor overall survival of patients with non-metastatic disease. <i>Clinical Chemistry and Laboratory Medicine</i> , 2018, 56, 990-1000.	2.3	28
34	Identification of a novel tRNA-derived RNA fragment exhibiting high prognostic potential in chronic lymphocytic leukemia. <i>Hematological Oncology</i> , 2019, 37, 498-504.	1.7	28
35	KLKB1 mRNA overexpression: A novel molecular biomarker for the diagnosis of chronic lymphocytic leukemia. <i>Clinical Biochemistry</i> , 2015, 48, 849-854.	1.9	27
36	Quantitative analysis of BCL2 mRNA expression in nasopharyngeal carcinoma: an unfavorable and independent prognostic factor. <i>Tumor Biology</i> , 2010, 31, 391-399.	1.8	26

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37	Identification of novel alternative splice variants of the BCL2L12 gene in human cancer cells using next-generation sequencing methodology. <i>Cancer Letters</i> , 2016, 373, 119-129.	7.2	26
38	mRNA overexpression of the hypoxia inducible factor 1 alpha subunit gene (HIF1A): An independent predictor of poor overall survival in chronic lymphocytic leukemia. <i>Leukemia Research</i> , 2017, 53, 65-73.	0.8	26
39	Quantitative expression analysis of the apoptosis-related gene, <i>BCL2L12</i> , in head and neck squamous cell carcinoma. <i>Journal of Oral Pathology and Medicine</i> , 2013, 42, 154-161.	2.7	25
40	Copper(II) Inverse-[9-Metallacrown-3] Compounds Accommodating Nitrate or Diclofenac Ligands: Structure, Magnetism, and Biological Activity. <i>European Journal of Inorganic Chemistry</i> , 2016, 2016, 219-231.	2.0	25
41	miR-34a overexpression predicts poor prognostic outcome in colorectal adenocarcinoma, independently of clinicopathological factors with established prognostic value. <i>Clinical Biochemistry</i> , 2017, 50, 918-924.	1.9	25
42	Quantitative expression analysis and prognostic significance of the BCL2-associated Xgene in nasopharyngeal carcinoma: a retrospective cohort study. <i>BMC Cancer</i> , 2013, 13, 293.	2.6	24
43	Progression of mouse skin carcinogenesis is associated with the orchestrated deregulation of miR-200 family members, miR-205 and their common targets. <i>Molecular Carcinogenesis</i> , 2016, 55, 1229-1242.	2.7	24
44	Identification of a novel, internal tRNA-derived RNA fragment as a new prognostic and screening biomarker in chronic lymphocytic leukemia, using an innovative quantitative real-time PCR assay. <i>Leukemia Research</i> , 2019, 87, 106234.	0.8	24
45	Identification of Two Novel Circular RNAs Deriving from BCL2L12 and Investigation of Their Potential Value as a Molecular Signature in Colorectal Cancer. <i>International Journal of Molecular Sciences</i> , 2020, 21, 8867.	4.1	24
46	High clusterin (CLU) mRNA expression levels in tumors of colorectal cancer patients predict a poor prognostic outcome. <i>Clinical Biochemistry</i> , 2020, 75, 62-69.	1.9	23
47	A novel, mitochondrial, internal tRNA-derived RNA fragment possesses clinical utility as a molecular prognostic biomarker in chronic lymphocytic leukemia. <i>Clinical Biochemistry</i> , 2020, 85, 20-26.	1.9	23
48	Effect of Vinca Alkaloids on the Expression Levels of microRNAs Targeting Apoptosis-related Genes in Breast Cancer Cell Lines. <i>Current Pharmaceutical Biotechnology</i> , 2019, 19, 1076-1086.	1.6	23
49	Blood-based analysis of type-2 diabetes mellitus susceptibility genes identifies specific transcript variants with deregulated expression and association with disease risk. <i>Scientific Reports</i> , 2019, 9, 1512.	3.3	21
50	Clinical utility of microRNAs in renal cell carcinoma: current evidence and future perspectives. <i>Expert Review of Molecular Diagnostics</i> , 2018, 18, 981-991.	3.1	20
51	L-DOPA decarboxylase mRNA expression is associated with tumor stage and size in head and neck squamous cell carcinoma: a retrospective cohort study. <i>BMC Cancer</i> , 2012, 12, 484.	2.6	19
52	Quantitative and qualitative analysis of regulatory T cells in B cell chronic lymphocytic leukemia. <i>Leukemia Research</i> , 2017, 60, 74-81.	0.8	18
53	MicroRNA-92a-3p overexpression in peripheral blood mononuclear cells is an independent predictor of prolonged overall survival of patients with chronic lymphocytic leukemia. <i>Leukemia and Lymphoma</i> , 2019, 60, 658-667.	1.3	18
54	Discovery of novel transcripts of the human tissue kallikrein (KLK1) and kallikrein-related peptidase 2 (KLK2) in human cancer cells, exploiting Next-Generation Sequencing technology. <i>Genomics</i> , 2019, 111, 642-652.	2.9	18

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55	The Multifaceted Role and Utility of MicroRNAs in Indolent B-Cell Non-Hodgkin Lymphomas. <i>Biomedicines</i> , 2021, 9, 333.	3.2	18
56	Identification and molecular cloning of novel transcripts of the human kallikrein-related peptidase 10 (KLK10) gene using next-generation sequencing. <i>Biochemical and Biophysical Research Communications</i> , 2017, 487, 776-781.	2.1	17
57	The transcriptome of a "sleeping" invader: de novo assembly and annotation of the transcriptome of aestivating <i>Cornu aspersum</i> . <i>BMC Genomics</i> , 2017, 18, 491.	2.8	17
58	Molecular cloning of novel transcripts of human kallikrein-related peptidases 5, 6, 7, 8 and 9 (KLK5 "Tj ETQq0 0,0rgBT /Overlock 10	3.3	17
59	Heat shock protein beta 3 (HSPB3) is an unfavorable molecular biomarker in colorectal adenocarcinoma. <i>Molecular Carcinogenesis</i> , 2020, 59, 116-125.	2.7	17
60	Multiple Myeloma Bone Disease: Implication of MicroRNAs in Its Molecular Background. <i>International Journal of Molecular Sciences</i> , 2021, 22, 2375.	4.1	17
61	Surrogate Prognostic Biomarkers in OSCC: The Paradigm of PA28 <sup>β</sup> Overexpression. <i>EBioMedicine</i> , 2015, 2, 784-785.	6.1	16
62	The Stat3/5 Signaling Biosignature in Hematopoietic Stem/Progenitor Cells Predicts Response and Outcome in Myelodysplastic Syndrome Patients Treated with Azacitidine. <i>Clinical Cancer Research</i> , 2016, 22, 1958-1968.	7.0	16
63	mRNA overexpression of kallikrein-related peptidase 14 (KLK14) is an independent predictor of poor overall survival in chronic lymphocytic leukemia patients. <i>Clinical Chemistry and Laboratory Medicine</i> , 2016, 54, 315-24.	2.3	15
64	The role of transcription factors in laboratory medicine. <i>Clinical Chemistry and Laboratory Medicine</i> , 2013, 51, 1563-1571.	2.3	14
65	Low mRNA expression levels of kallikrein-related peptidase 4 (KLK4) predict short-term relapse in patients with laryngeal squamous cell carcinoma. <i>Biological Chemistry</i> , 2014, 395, 1051-1062.	2.5	14
66	The outcome of patients with high-risk MDS achieving stable disease after treatment with 5-azacytidine: A retrospective analysis of the Hellenic (Greek) MDS Study Group. <i>Hematological Oncology</i> , 2018, 36, 693-700.	1.7	14
67	Novel alternative splice variants of the human protein arginine methyltransferase 1 (PRMT1) gene, discovered using next-generation sequencing. <i>Gene</i> , 2019, 699, 135-144.	2.2	14
68	Increased expression of phosphorylated NBS1, a key molecule of the DNA damage response machinery, is an adverse prognostic factor in patients with de novo myelodysplastic syndromes. <i>Leukemia Research</i> , 2013, 37, 1576-1582.	0.8	13
69	Treatment with 5-Azacitidine improves clinical outcome in high-risk MDS patients in the "real life" setting: A single center observational study. <i>Hematology</i> , 2016, 21, 34-41.	1.5	13
70	A Cancer-Related microRNA Signature Shows Biomarker Utility in Multiple Myeloma. <i>International Journal of Molecular Sciences</i> , 2021, 22, 13144.	4.1	13
71	Chronic myelomonocytic leukemia treated with 5-azacytidine " results from the Hellenic 5-Azacitidine Registry: proposal of a new risk stratification system. <i>Leukemia and Lymphoma</i> , 2019, 60, 1721-1730.	1.3	12
72	A Molecular Signature of Circulating MicroRNA Can Predict Osteolytic Bone Disease in Multiple Myeloma. <i>Cancers</i> , 2021, 13, 3877.	3.7	12

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73	High Expression of a tRNA <sup>Pro</sup> Derivative Associates with Poor Survival and Independently Predicts Colorectal Cancer Recurrence. <i>Biomedicines</i> , 2022, 10, 1120.	3.2	12
74	Positive BCL2L12 expression predicts favorable prognosis in patients with laryngeal squamous cell carcinoma. <i>Cancer Biomarkers</i> , 2019, 25, 141-149.	1.7	11
75	A 3â€² tRNAâ€derived fragment produced by tRNA <sup>LeuAAG</sup> and tRNA <sup>LeuTAC</sup> is associated with poor prognosis in Bâ€cell chronic lymphocytic leukemia, independently of classical prognostic factors. <i>European Journal of Haematology</i> , 2021, 106, 821-830.	2.2	11
76	The Role of Circulating MicroRNAs in Patients with Early-Stage Pancreatic Adenocarcinoma. <i>Biomedicines</i> , 2021, 9, 1468.	3.2	11
77	Identification of Novel Circular RNAs of the Human Protein Arginine Methyltransferase 1 (PRMT1) Gene, Expressed in Breast Cancer Cells. <i>Genes</i> , 2022, 13, 1133.	2.4	11
78	The prognostic value of monosomal karyotype (MK) in higherâ€risk patients with myelodysplastic syndromes treated with 5â€Azacitidine: A retrospective analysis of the Hellenic (Greek) Myelodysplastic syndromes Study Group. <i>American Journal of Hematology</i> , 2018, 93, 895-901.	4.1	10
79	BCL2L12 improves risk stratification and prediction of BFM-chemotherapy response in childhood acute lymphoblastic leukemia. <i>Clinical Chemistry and Laboratory Medicine</i> , 2018, 56, 2104-2118.	2.3	10
80	Novel splice variants of the human kallikrein-related peptidases 11 (<i>KLK11</i>) and 12 (<i>KLK12</i>), unraveled by next-generation sequencing technology. <i>Biological Chemistry</i> , 2018, 399, 1065-1071.	2.5	10
81	Identification of novel alternative splice variants of the human L-DOPA decarboxylase (DDC) gene in human cancer cells, using high-throughput sequencing approaches. <i>Gene</i> , 2019, 719, 144075.	2.2	10
82	Revised Exon Structure of L-DOPA Decarboxylase (DDC) Reveals Novel Splice Variants Associated with Colorectal Cancer Progression. <i>International Journal of Molecular Sciences</i> , 2020, 21, 8568.	4.1	10
83	Next-generation sequencing reveals alternative L-DOPA decarboxylase (DDC) splice variants bearing novel exons, in human hepatocellular and lung cancer cells. <i>Gene</i> , 2021, 768, 145262.	2.2	10
84	Molecular Effects of Treatment of Human Colorectal Cancer Cells with Natural and Classical Chemotherapeutic Drugs: Alterations in the Expression of Apoptosis-related BCL2 Family Members, Including BCL2L12. <i>Current Pharmaceutical Biotechnology</i> , 2019, 19, 1064-1075.	1.6	10
85	BCL2L12 protein overexpression is associated with favorable outcome in diffuse large B-cell lymphoma patients in the rituximab era. <i>Leukemia and Lymphoma</i> , 2016, 57, 2199-2203.	1.3	9
86	Discovery and expression analysis of novel transcripts of the human SR-related CTD-associated factor 1 (SCAF1) gene in human cancer cells using Next-Generation Sequencing. <i>Gene</i> , 2018, 670, 155-165.	2.2	9
87	Molecular cloning of novel transcripts of the adaptor-related protein complex 2 alpha 1 subunit (AP2A1) gene, using Next-Generation Sequencing. <i>Gene</i> , 2018, 678, 55-64.	2.2	9
88	tRNA Derivatives in Multiple Myeloma: Investigation of the Potential Value of a tRNA-Derived Molecular Signature. <i>Biomedicines</i> , 2021, 9, 1811.	3.2	8
89	Immunophenotypic Profile of CD34+ Subpopulations and Their Role in the Diagnosis and Prognosis of Patients with Deâ€Novo, Particularly Lowâ€Grade Myelodysplastic Syndromes. <i>Cytometry Part B - Clinical Cytometry</i> , 2019, 96, 73-82.	1.5	7
90	Contribution of miRNAs, tRNAs and tRFs to Aberrant Signaling and Translation Deregulation in Lung Cancer. <i>Cancers</i> , 2020, 12, 3056.	3.7	7

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91	Effectiveness of 5-Azacytidine in older patients with high-risk myelodysplastic syndromes and oligoblastic acute myeloid leukemia: A retrospective analysis of the Hellenic (Greek) MDS Study Group. <i>Journal of Geriatric Oncology</i> , 2020, 11, 121-124.	1.0	5
92	Complex transcriptional regulation of the BCL2L12 gene: Novel, active promoter in K562 cells. <i>Gene</i> , 2020, 750, 144723.	2.2	4
93	Editorial for the Special Issue "Molecular Biomarkers in Colorectal Adenocarcinoma". <i>International Journal of Molecular Sciences</i> , 2021, 22, 2052.	4.1	4
94	Molecular Biomarkers of Laryngeal Cancer Laryngeal squamous cell carcinoma (LSCC) Tumor markers. <i>Biomarkers in Disease</i> , 2015, , 891-919.	0.1	4
95	Identification of novel alternative transcripts of the human Ribonuclease H <sup>2</sup> (RNASEH2) gene using 3' RACE and high-throughput sequencing approaches. <i>Genomics</i> , 2020, 112, 943-951.	2.9	3
96	Identification and expression analysis of novel splice variants of the human carcinoembryonic antigen-related cell adhesion molecule 19 (CEACAM19) gene using a high-throughput sequencing approach. <i>Genomics</i> , 2020, 112, 4268-4276.	2.9	3
97	Pharmacoepiggenomics circuits induced by a novel retinoid-polyamine conjugate in human immortalized keratinocytes. <i>Pharmacogenomics Journal</i> , 2021, 21, 638-648.	2.0	3
98	Recovery and quantification of a myelin oligodendrocyte glycoprotein peptide from rat plasma after protein precipitation. <i>Analytical Biochemistry</i> , 2017, 538, 71-73.	2.4	1
99	THE tRNA-DERIVED RNA FRAGMENTS (tRFs) BEARING THE GLYCINE ANTICODONS GCC AND CCC AS EMERGING MOLECULAR BIOMARKERS OF UNFAVORABLE PROGNOSIS IN CHRONIC LYMPHOCYTIC LEUKEMIA. <i>Hematological Oncology</i> , 2019, 37, 375-376.	1.7	1
100	Molecular Biomarkers of Laryngeal Cancer. , 2014, , 1-24.		1
101	A Molecular Signature of Three tRNA-Derived RNA Fragments May Discriminate Smoldering from Symptomatic Multiple Myeloma Patients. <i>Blood</i> , 2019, 134, 5528-5528.	1.4	1
102	Let-7a, Mir-17 and Mir-20a Expression Levels in CD34+ Bone Marrow Cells of Patients with Myelodysplastic Syndromes (MDS) Are Associated with Established Prognostic Factors, Supporting Their Implication in the Pathogenesis of the Disease,. <i>Blood</i> , 2011, 118, 3792-3792.	1.4	1
103	Translating transcriptome to immunophenotype in head and neck squamous cell carcinoma (HNSCC) to identify pathways promoting T-cell infiltration.. <i>Journal of Clinical Oncology</i> , 2019, 37, e17542-e17542.	1.6	1
104	Body mass index and relative dose intensity does not affect the response and outcome of high-risk MDS patients treated with azacytidine. Results from the Hellenic (Greek) MDS study group. <i>Leukemia Research</i> , 2018, 71, 55-59.	0.8	0
105	962P Immunotherapeutic implications of mucin 1/16 expression in head and neck squamous cell carcinoma (HNSCC). <i>Annals of Oncology</i> , 2020, 31, S679-S680.	1.2	0
106	Identification of six novel alternative transcripts of the human kallikrein-related peptidase 15 (KLK15), using 3' RACE and high-throughput sequencing. <i>Gene</i> , 2020, 749, 144708.	2.2	0
107	Abstract 2006: Identification and molecular cloning of novel splicing variants of BCL2L12, a new member of the BCL2 family, and their expression analysis in breast, prostate, and colon cancer cells. , 2012, , .		0
108	Abstract LB-72: Expression of the BCL2-like 12 (BCL2L12) gene is associated with prolonged survival of breast adenocarcinoma patients.. , 2013, , .		0



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109	Quantitative and Qualitative Analysis Of Regulatory T Cells (Tregs) Derived From The Peripheral Blood Of Chronic Lymphocytic Leukemia Patients. <i>Blood</i> , 2013, 122, 5280-5280.	1.4	0
110	Quantitative and Qualitative Analysis of Regulatory T Cells (Tregs) in B Cell Chronic Lymphocytic Leukemia (B-CLL). <i>Blood</i> , 2015, 126, 2928-2928.	1.4	0
111	Validation of the Revised International Prognostic Scoring System in 2582 Patients with Myelodysplastic Syndrome: A Multicenter Study By the Hellenic MDS Study Group. <i>Blood</i> , 2016, 128, 2004-2004.	1.4	0
112	KLK14 (kallikrein-related peptidase 14). <i>Atlas of Genetics and Cytogenetics in Oncology and Haematology</i> , 2018, , .	0.1	0
113	Prognostic Significance of Severe Thrombocytopenia in Overall Survival of Patients with Myelodysplastic Syndromes Treated with Azacytidine. a Multicenter Study By the Hellenic MDS Study Group. <i>Blood</i> , 2018, 132, 1822-1822.	1.4	0
114	The Prognostic Significance of Monocytopenia in Patients with Myelodysplastic Syndrome. <i>Blood</i> , 2019, 134, 5427-5427.	1.4	0
115	BOK (BCL2 family apoptosis regulator BOK). <i>Atlas of Genetics and Cytogenetics in Oncology and Haematology</i> , 2019, , .	0.1	0
116	The Clinical Significance of a Novel microRNA Signature in Multiple Myeloma. <i>Blood</i> , 2019, 134, 5529-5529.	1.4	0
117	KLK12 (kallikrein-related peptidase 12). <i>Atlas of Genetics and Cytogenetics in Oncology and Haematology</i> , 2020, , .	0.1	0
118	KLK6 (kallikrein-related peptidase 6). <i>Atlas of Genetics and Cytogenetics in Oncology and Haematology</i> , 2020, , .	0.1	0
119	CEACAM19 (carcinoembryonic antigen related cell adhesion molecule 19). <i>Atlas of Genetics and Cytogenetics in Oncology and Haematology</i> , 2020, , .	0.1	0