Juan J Lozano

List of Publications by Year in descending order

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176	10,363 citations	57 h-index	94 g-index
papers	Citations	II-IIIdex	g-mdex
179 all docs	179 docs citations	179 times ranked	16994 citing authors

#	Article	IF	Citations
1	Epithelial-mesenchymal transition can suppress major attributes of human epithelial tumor-initiating cells. Journal of Clinical Investigation, 2012, 122, 1849-1868.	3.9	401
2	Fecal MicroRNAs as Novel Biomarkers for Colon Cancer Screening. Cancer Epidemiology Biomarkers and Prevention, 2010, 19, 1766-1774.	1.1	310
3	Epigenetic Silencing of miR-137 Is an Early Event in Colorectal Carcinogenesis. Cancer Research, 2010, 70, 6609-6618.	0.4	275
4	Prospective multicenter clinical trial of immunosuppressive drug withdrawal in stable adult liver transplant recipients. Hepatology, 2013, 58, 1824-1835.	3.6	269
5	DNA Damage Regulates Alternative Splicing through Inhibition of RNA Polymerase II Elongation. Cell, 2009, 137, 708-720.	13.5	267
6	Patients with drug-free long-term graft function display increased numbers of peripheral B cells with a memory and inhibitory phenotype. Kidney International, 2010, 78, 503-513.	2.6	249
7	Using transcriptional profiling to develop a diagnostic test of operational tolerance in liver transplant recipients. Journal of Clinical Investigation, 2008, 118, 2845-57.	3.9	249
8	Blood metabolomics uncovers inflammation-associated mitochondrial dysfunction as a potential mechanism underlying ACLF. Journal of Hepatology, 2020, 72, 688-701.	1.8	223
9	Transcriptional analysis of the intestinal mucosa of patients with ulcerative colitis in remission reveals lasting epithelial cell alterations. Gut, 2013, 62, 967-976.	6.1	208
10	Intra-graft expression of genes involved in iron homeostasis predicts the development of operational tolerance in human liver transplantation. Journal of Clinical Investigation, 2012, 122, 368-382.	3.9	183
11	Gene Discovery in Bladder Cancer Progression using cDNA Microarrays. American Journal of Pathology, 2003, 163, 505-516.	1.9	177
12	Liver progenitor cell markers correlate with liver damage and predict short-term mortality in patients with alcoholic hepatitis. Hepatology, 2012, 55, 1931-1941.	3.6	177
13	Generation of Hepatic Stellate Cells from Human Pluripotent Stem Cells Enables InÂVitro Modeling of Liver Fibrosis. Cell Stem Cell, 2018, 23, 101-113.e7.	5.2	170
14	Transcriptome analysis identifies TNF superfamily receptors as potential therapeutic targets in alcoholic hepatitis. Gut, 2013, 62, 452-460.	6.1	167
15	Profiling Bladder Cancer Using Targeted Antibody Arrays. American Journal of Pathology, 2006, 168, 93-103.	1.9	162
16	Tumour initiating cells and IGF/FGF signalling contribute to sorafenib resistance in hepatocellular carcinoma. Gut, 2017, 66, 530-540.	6.1	161
17	The biliary epithelium gives rise to liver progenitor cells. Hepatology, 2014, 60, 1367-1377.	3. 6	158
18	Circulating MicroRNAs as Biomarkers of Colorectal Cancer: Results From a Genome-Wide Profiling and Validation Study. Clinical Gastroenterology and Hepatology, 2013, 11, 681-688.e3.	2.4	157

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19	Role of UEV-1, an Inactive Variant of the E2 UbiquitinConjugating Enzymes, in In Vitro Differentiation and Cell Cycle Behavior of HT-29-M6 Intestinal Mucosecretory Cells. Molecular and Cellular Biology, 1998, 18, 576-589.	1.1	142
20	Ghrelin attenuates hepatocellular injury and liver fibrogenesis in rodents and influences fibrosis progression in humans. Hepatology, 2010, 51, 974-985.	3.6	141
21	Applicability, safety, and biological activity of regulatory T cell therapy in liver transplantation. American Journal of Transplantation, 2020, 20, 1125-1136.	2.6	139
22	Curcumin Modulates DNA Methylation in Colorectal Cancer Cells. PLoS ONE, 2013, 8, e57709.	1.1	135
23	Epithelial-to-Mesenchymal Transition Mediates Docetaxel Resistance and High Risk of Relapse in Prostate Cancer. Molecular Cancer Therapeutics, 2014, 13, 1270-1284.	1.9	131
24	Evidence of Chronic Allograft Injury in Liver Biopsies From Long-term Pediatric Recipients of Liver Transplants. Gastroenterology, 2018, 155, 1838-1851.e7.	0.6	125
25	Defective HNF4alpha-dependent gene expression as a driver of hepatocellular failure in alcoholic hepatitis. Nature Communications, 2019, 10, 3126.	5.8	124
26	Identification of inflammatory mediators in patients with Crohn's disease unresponsive to anti-TNFÎ \pm therapy. Gut, 2015, 64, 233-242.	6.1	123
27	Hnf $1\hat{l}\pm$ (MODY3) Controls Tissue-Specific Transcriptional Programs and Exerts Opposed Effects on Cell Growth in Pancreatic Islets and Liver. Molecular and Cellular Biology, 2009, 29, 2945-2959.	1.1	122
28	Identification of blood serum microâ€RNAs associated with idiopathic and <i>LRRK2</i> Parkinson's disease. Journal of Neuroscience Research, 2014, 92, 1071-1077.	1.3	122
29	Comparison of Transcriptional and Blood Cell-Phenotypic Markers Between Operationally Tolerant Liver and Kidney Recipients. American Journal of Transplantation, 2011, 11, 1916-1926.	2.6	120
30	Colorectal Cancers with Microsatellite Instability Display Unique miRNA Profiles. Clinical Cancer Research, 2011, 17, 6239-6249.	3.2	112
31	Liver-specific deletion of prohibitin 1 results in spontaneous liver injury, fibrosis, and hepatocellular carcinoma in mice. Hepatology, 2010, 52, 2096-2108.	3.6	107
32	Molecular interplay between \hat{l} "5/ \hat{l} "6 desaturases and long-chain fatty acids in the pathogenesis of non-alcoholic steatohepatitis. Gut, 2014, 63, 344-355.	6.1	107
33	Comparative Analysis of Chloroplast Genomes: Functional Annotation, Genome-Based Phylogeny, and Deduced Evolutionary Patterns. Genome Research, 2002, 12, 567-583.	2.4	106
34	Upregulation of miR-142-3p in Peripheral Blood Mononuclear Cells of Operationally Tolerant Patients with a Renal Transplant. Journal of the American Society of Nephrology: JASN, 2012, 23, 597-606.	3.0	105
35	Integrative microRNA profiling in alcoholic hepatitis reveals a role for microRNA-182 in liver injury and inflammation. Gut, 2016, 65, 1535-1545.	6.1	103
36	MERTK as negative regulator of human T cell activation. Journal of Leukocyte Biology, 2015, 97, 751-760.	1.5	99

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37	Parp-2 is required to maintain hematopoiesis following sublethal \hat{I}^3 -irradiation in mice. Blood, 2013, 122, 44-54.	0.6	96
38	PARP-2 sustains erythropoiesis in mice by limiting replicative stress in erythroid progenitors. Cell Death and Differentiation, 2015, 22, 1144-1157.	5.0	95
39	Fusion of the Human Gene for the Polyubiquitination Coeffector UEV1 with Kua, a Newly Identified Gene. Genome Research, 2000, 10, 1743-1756.	2.4	91
40	Using microRNA profiling in urine samples to develop a non-invasive test for bladder cancer. International Journal of Cancer, 2013, 133, n/a-n/a.	2.3	88
41	Signaling and Immunoresolving Actions of Resolvin D1 in Inflamed Human Visceral Adipose Tissue. Journal of Immunology, 2016, 197, 3360-3370.	0.4	87
42	IL-2 therapy restores regulatory T-cell dysfunction induced by calcineurin inhibitors. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 7083-7088.	3.3	87
43	Whole-exome sequencing identifies rare pathogenic variants in new predisposition genes for familial colorectal cancer. Genetics in Medicine, 2015, 17, 131-142.	1.1	82
44	Integrative miRNA and Gene Expression Profiling Analysis of Human Quiescent Hepatic Stellate Cells. Scientific Reports, 2015, 5, 11549.	1.6	79
45	Mutational Signatures in Cancer (MuSiCa): a web application to implement mutational signatures analysis in cancer samples. BMC Bioinformatics, 2018, 19, 224.	1.2	77
46	HCV-Induced Immune Responses Influence the Development of Operational Tolerance After Liver Transplantation in Humans. Science Translational Medicine, 2014, 6, 242ra81.	5.8	74
47	Genomic resources for a commercial flatfish, the Senegalese sole (Solea senegalensis): EST sequencing, oligo microarray design, and development of the bioinformatic platform Soleamold. BMC Genomics, 2008, 9, 508.	1.2	70
48	Characterization of $\hat{l}^3\hat{l}$ T cell subsets in organ transplantation. Transplant International, 2010, 23, 1045-1055.	0.8	68
49	S-adenosylmethionine Levels Regulate the Schwann Cell DNA Methylome. Neuron, 2014, 81, 1024-1039.	3.8	67
50	A five-gene expression signature to predict progression in T1G3 bladder cancer. European Journal of Cancer, 2016, 64, 127-136.	1.3	67
51	Genome-wide differences between microsatellite stable and unstable colorectal tumors. Carcinogenesis, 2006, 27, 419-428.	1.3	66
52	DNA Microarray Expression Profiling of Bladder Cancer Allows Identification of Noninvasive Diagnostic Markers. Journal of Urology, 2009, 182, 741-748.	0.2	65
53	ATGâ€Fresenius Treatment and Lowâ€Dose Tacrolimus: Results of a Randomized Controlled Trial in Liver Transplantation. American Journal of Transplantation, 2010, 10, 2296-2304.	2.6	65
54	PTOV1, a novel protein overexpressed in prostate cancer containing a new class of protein homology blocks. Oncogene, 2001, 20, 1455-1464.	2.6	61

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55	3D-QSAR methods on the basis of ligand-receptor complexes. Application of COMBINE and GRID/GOLPE methodologies to a series of CYP1A2 ligands. Journal of Computer-Aided Molecular Design, 2000, 14, 341-353.	1.3	59
56	Comparative Transcriptional and Phenotypic Peripheral Blood Analysis of Kidney Recipients Under Cyclosporin A or Sirolimus Monotherapy. American Journal of Transplantation, 2010, 10, 2604-2614.	2.6	59
57	Aberrant Gene Promoter Methylation Associated with Sporadic Multiple Colorectal Cancer. PLoS ONE, 2010, 5, e8777.	1.1	59
58	TL1A/TNFSF15 directly induces proinflammatory cytokines, including TNF \hat{l}_{\pm} , from CD3+CD161+ T cells to exacerbate gut inflammation. Mucosal Immunology, 2013, 6, 886-899.	2.7	59
59	Predictive Value of MicroRNAs in the Progression of Barrett Esophagus to Adenocarcinoma in a Long-Term Follow-up Study. Annals of Surgery, 2013, 257, 886-893.	2.1	59
60	The Fanconi anemia DNA damage repair pathway in the spotlight for germline predisposition to colorectal cancer. European Journal of Human Genetics, 2016, 24, 1501-1505.	1.4	59
61	Efficacy and Safety of Immunosuppression Withdrawal in Pediatric Liver Transplant Recipients: Moving Toward Personalized Management. Hepatology, 2021, 73, 1985-2004.	3.6	57
62	MicroRNAs for Detection of Pancreatic Neoplasia. Annals of Surgery, 2017, 265, 1226-1234.	2.1	56
63	The specialized proresolving lipid mediator maresin 1 protects hepatocytes from lipotoxic and hypoxiaâ€induced endoplasmic reticulum stress. FASEB Journal, 2017, 31, 5384-5398.	0.2	56
64	Gene Expression Profiling and Secretome Analysis Differentiate Adult-Derived Human Liver Stem/Progenitor Cells and Human Hepatic Stellate Cells. PLoS ONE, 2014, 9, e86137.	1.1	55
65	microRNA profiling in duodenal ulcer disease caused by Helicobacter pylori infection in a Western population. Clinical Microbiology and Infection, 2012, 18, E273-E282.	2.8	53
66	Plasma MicroRNA Signature Validation for Early Detection of Colorectal Cancer. Clinical and Translational Gastroenterology, 2019, 10, e00003.	1.3	53
67	Ductular Reaction Cells Display an Inflammatory Profile and Recruit Neutrophils in Alcoholic Hepatitis. Hepatology, 2019, 69, 2180-2195.	3.6	52
68	Viral and immune factors associated with successful treatment withdrawal in HBeAg-negative chronic hepatitis B patients. Journal of Hepatology, 2021, 74, 1064-1074.	1.8	52
69	Mitochondrial dysfunction governs immunometabolism in leukocytes of patients with acute-on-chronic liver failure. Journal of Hepatology, 2022, 76, 93-106.	1.8	51
70	Nearâ€tetraploid cancer cells show chromosome instability triggered by replication stress and exhibit enhanced invasiveness. FASEB Journal, 2018, 32, 3502-3517.	0.2	50
71	Identification and Validation of MicroRNA Profiles in Fecal Samples for Detection of Colorectal Cancer. Gastroenterology, 2020, 158, 947-957.e4.	0.6	48
72	Immunosuppression Withdrawal in Liver Transplant Recipients on Sirolimus. Hepatology, 2020, 72, 569-583.	3.6	45

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73	Assessing the role of amino acids in systemic inflammation and organ failure in patients with ACLF. Journal of Hepatology, 2021, 74, 1117-1131.	1.8	45
74	Usefulness of Transcriptional Blood Biomarkers as a Non-invasive Surrogate Marker of Mucosal Healing and Endoscopic Response in Ulcerative Colitis. Journal of Crohn's and Colitis, 2017, 11, 1335-1346.	0.6	44
75	Iron Deficiency Impairs Intra-Hepatic Lymphocyte Mediated Immune Response. PLoS ONE, 2015, 10, e0136106.	1.1	44
76	Transcriptional regulation by Poly(ADP-ribose) polymerase-1 during T cell activation. BMC Genomics, 2008, 9, 171.	1.2	42
77	HER3 is required for the maintenance of neuregulinâ€dependent and â€independent attributes of malignant progression in prostate cancer cells. International Journal of Cancer, 2009, 125, 2565-2575.	2.3	41
78	MiRComb: An R Package to Analyse miRNA-mRNA Interactions. Examples across Five Digestive Cancers. PLoS ONE, 2016, 11, e0151127.	1.1	41
79	Hepatoma Cells From Mice Deficient in Glycine N-Methyltransferase Have Increased RAS Signaling and Activation of Liver Kinase B1. Gastroenterology, 2012, 143, 787-798.e13.	0.6	40
80	Novel Circulating miRNA Signatures for Early Detection of Pancreatic Neoplasia. Clinical and Translational Gastroenterology, 2019, 10, e00029.	1.3	40
81	Urine cell-based DNA methylation classifier for monitoring bladder cancer. Clinical Epigenetics, 2018, 10, 71.	1.8	39
82	Characterization of Blood Immune Cells in Patients With Decompensated Cirrhosis Including ACLF. Frontiers in Immunology, 2020, 11, 619039.	2.2	39
83	Molecular Characterization of Acute Cellular Rejection Occurring During Intentional Immunosuppression Withdrawal in Liver Transplantation. American Journal of Transplantation, 2016, 16, 484-496.	2.6	38
84	HuR/ELAVL1 drives malignant peripheral nerve sheath tumor growth and metastasis. Journal of Clinical Investigation, 2020, 130, 3848-3864.	3.9	38
85	Three-dimensional modelling of human cytochrome P450 1A2 and its interaction with caffeine and MeIQ. Journal of Computer-Aided Molecular Design, 1997, 11, 395-408.	1.3	34
86	Molecular modelling of the differential interaction between several non-steroidal anti-inflammatory drugs and human prostaglandin endoperoxide H synthase-2 (h-PGHS-2). Journal of Molecular Graphics and Modelling, 2002, 20, 329-343.	1.3	34
87	Gene expression profiling distinguishes JAK2V617F-negative from JAK2V617F-positive patients in essential thrombocythemia. Leukemia, 2008, 22, 1368-1376.	3.3	34
88	Pregnane X-receptor promotes stem cell-mediated colon cancer relapse. Oncotarget, 2016, 7, 56558-56573.	0.8	34
89	New genes emerging for colorectal cancer predisposition. World Journal of Gastroenterology, 2014, 20, 1961.	1.4	34
90	Analysis of A 6-Mirna Signature in Serum from Colorectal Cancer Screening Participants as Non-Invasive Biomarkers for Advanced Adenoma and Colorectal Cancer Detection. Cancers, 2019, 11, 1542.	1.7	33

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91	Gene expression test for the non-invasive diagnosis of bladder cancer: A prospective, blinded, international and multicenter validation study. European Journal of Cancer, 2016, 54, 131-138.	1.3	32
92	RAC1b overexpression correlates with poor prognosis in KRAS/BRAF WT metastatic colorectal cancer patients treated with first-line FOLFOX/XELOX chemotherapy. European Journal of Cancer, 2014, 50, 1973-1981.	1.3	31
93	Untargeted lipidomics uncovers lipid signatures that distinguish severe from moderate forms of acutely decompensated cirrhosis. Journal of Hepatology, 2021, 75, 1116-1127.	1.8	31
94	Validation Study of a Noninvasive Urine Test for Diagnosis and Prognosis Assessment of Bladder Cancer: Evidence for Improved Models. Journal of Urology, 2014, 191, 261-269.	0.2	30
95	Simvastatin Attenuates Liver Injury in Rodents with Biliary Cirrhosis Submitted to Hemorrhage/Resuscitation. Shock, 2017, 47, 370-377.	1.0	30
96	Aging Influences Hepatic Microvascular Biology and Liver Fibrosis in Advanced Chronic Liver Disease., 2019, 10, 684.		30
97	Prognostic value of <scp>microRNA</scp> expression pattern in upper tract urothelial carcinoma. BJU International, 2014, 113, 813-821.	1.3	29
98	QSAR and conformational analysis of the antiinflammatory agent amfenac and analogues. Journal of Computer-Aided Molecular Design, 1993, 7, 183-198.	1.3	28
99	Metabolomics Discloses Potential Biomarkers for the Noninvasive Diagnosis of Idiopathic Portal Hypertension. American Journal of Gastroenterology, 2013, 108, 926-932.	0.2	28
100	<i>Solute carrier family 2 member $1 < i$ is involved in the development of nonalcoholic fatty liver disease. Hepatology, 2013, 57, 505-514.</i>	3.6	25
101	LPS-TLR4 Pathway Mediates Ductular Cell Expansion in Alcoholic Hepatitis. Scientific Reports, 2016, 6, 35610.	1.6	25
102	Activation of the epidermal growth factor signalling pathway by tissue plasminogen activator in pancreas cancer cells. Gut, 2007, 56, 1266-1274.	6.1	24
103	MicroRNA-200, associated with metastatic breast cancer, promotes traits of mammary luminal progenitor cells. Oncotarget, 2017, 8, 83384-83406.	0.8	23
104	Hypothalamus transcriptome profile suggests an anorexia-cachexia syndrome in the anx/anx mouse model. Physiological Genomics, 2008, 35, 341-350.	1.0	22
105	Protection from hepatic lipid accumulation and inflammation by genetic ablation of 5-lipoxygenase. Prostaglandins and Other Lipid Mediators, 2010, 92, 54-61.	1.0	22
106	Geneâ€expression signature of tumor recurrence in patients with stage II and III colon cancer treated with 5′fluoruracilã€based adjuvant chemotherapy. International Journal of Cancer, 2013, 132, 1090-1097.	2.3	22
107	Using gene expression from urine sediment to diagnose prostate cancer: development of a new multiplex mRNA urine test and validation of current biomarkers. BMC Cancer, 2016, 16, 76.	1.1	22
108	Urinary cell microRNA-based prognostic classifier for non-muscle invasive bladder cancer. Oncotarget, 2017, 8, 18238-18247.	0.8	22

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109	Conformational analysis of the antiinflammatory fenamates: a molecular mechanics and semiempirical molecular orbital study. Computational and Theoretical Chemistry, 1995, 335, 215-227.	1.5	21
110	Transportome Profiling Identifies Profound Alterations in Crohn's Disease Partially Restored by Commensal Bacteria. Journal of Crohn's and Colitis, 2016, 10, 850-859.	0.6	21
111	Up-regulation of EP2 and EP3 receptors in human tolerogenic dendritic cells boosts the immunosuppressive activity of PGE2. Journal of Leukocyte Biology, 2017, 102, 881-895.	1.5	21
112	Molecular profiling of peripheral blood is associated with circulating tumor cells content and poor survival in metastatic castration-resistant prostate cancer. Oncotarget, 2015, 6, 10604-10616.	0.8	21
113	Human amniotic stem cells improve hepatic microvascular dysfunction and portal hypertension in cirrhotic rats. Liver International, 2020, 40, 2500-2514.	1.9	20
114	Ductular reaction promotes intrahepatic angiogenesis through Slit2–Roundabout 1 signaling. Hepatology, 2022, 75, 353-368.	3.6	20
115	Partially Degraded RNA from Bladder Washing is a Suitable Sample for Studying Gene Expression Profiles in Bladder Cancer. European Urology, 2006, 50, 1347-1356.	0.9	19
116	Evaluation of Responsive Gene Expression as a Sensitive and Specific Biomarker in Patients with Ulcerative Colitis. Inflammatory Bowel Diseases, 2013, 19, 221-229.	0.9	19
117	Dual activation of pathways regulated by steroid receptors and peptide growth factors in primary prostate cancer revealed by Factor Analysis of microarray data. BMC Genomics, 2005, 6, 109.	1.2	18
118	Genomic imbalances in Schistosoma-associated and non–Schistosoma-associated bladder carcinoma. An array comparative genomic hybridization analysis. Cancer Genetics and Cytogenetics, 2007, 177, 16-19.	1.0	18
119	Genetic susceptibility variants associated with colorectal cancer prognosis. Carcinogenesis, 2013, 34, 2286-2291.	1.3	18
120	Patterns of somatic uniparental disomy identify novel tumor suppressor genes in colorectal cancer. Carcinogenesis, 2015, 36, 1103-1110.	1.3	18
121	New Rat Model of Advanced NASH Mimicking Pathophysiological Features and Transcriptomic Signature of The Human Disease. Cells, 2019, 8, 1062.	1.8	17
122	Metabolomics discloses potential biomarkers to predict the acute HVPG response to propranolol in patients with cirrhosis. Liver International, 2019, 39, 705-713.	1.9	17
123	Clinical, histological and molecular profiling of different stages of alcohol-related liver disease. Gut, 2022, 71, 1856-1866.	6.1	17
124	Role of UEV-1A, a homologue of the tumor suppressor protein TSG101, in protection from DNA damage. FEBS Letters, 1998, 423, 49-52.	1.3	16
125	The structural and electronical factors that contribute affinity for the time-dependent inhibition of PGHS-1 by indomethacin, diclofenac and fenamates. Journal of Computer-Aided Molecular Design, 1999, 13, 297-313.	1.3	16
126	Comparison of Gene Expression Profiles in Laser-Microdissected, Nonembedded, and OCT-Embedded Tumor Samples by Oligonucleotide Microarray Analysis. Clinical Chemistry, 2003, 49, 2096-2100.	1.5	16

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127	Integrated Analysis of Germline and Tumor DNA Identifies New Candidate Genes Involved in Familial Colorectal Cancer. Cancers, 2019, 11, 362.	1.7	16
128	Prognostic value of circulating microRNAs in upper tract urinary carcinoma. Oncotarget, 2018, 9, 16691-16700.	0.8	16
129	Structure-based QSAR study on differential inhibition of human prostaglandin endoperoxide H synthase-2 (COX-2) by nonsteroidal anti-inflammatory drugs. Journal of Computer-Aided Molecular Design, 2002, 16, 683-709.	1.3	15
130	Metabolomics as a diagnostic tool for idiopathic nonâ€cirrhotic portal hypertension. Liver International, 2016, 36, 1051-1058.	1.9	15
131	MiR-93 is related to poor prognosis in pancreatic cancer and promotes tumor progression by targeting microtubule dynamics. Oncogenesis, 2020, 9, 43.	2.1	15
132	Genetic Variants Associated with Colorectal Adenoma Susceptibility. PLoS ONE, 2016, 11, e0153084.	1.1	15
133	Profiling circulating microRNAs in patients with cirrhosis and acute-on-chronic liver failure. JHEP Reports, 2021, 3, 100233.	2.6	14
134	Molecular characterization of chronic liver disease dynamics: From liver fibrosis to acute-on-chronic liver failure. JHEP Reports, 2022, 4, 100482.	2.6	14
135	The RNA-Binding Protein Human Antigen R Controls Global Changes in Gene Expression during Schwann Cell Development. Journal of Neuroscience, 2012, 32, 4944-4958.	1.7	12
136	Aberrant brain microRNA target and miRISC gene expression in the anx/anx anorexia mouse model. Gene, 2012, 497, 181-190.	1.0	12
137	Randomized Controlled Trial Substudy of Cell-specific Mechanisms of Janus Kinase 1 Inhibition With Upadacitinib in the Crohn's Disease Intestinal Mucosa: Analysis From the CELEST Study. Inflammatory Bowel Diseases, 2021, 27, 1999-2009.	0.9	12
138	Perfil de expresión génica en el cáncer de próstata: identificación de marcadores candidatos para el diagnóstico no invasivo. Actas Urológicas Españolas, 2014, 38, 143-149.	0.3	11
139	Rare germline copy number variants in colorectal cancer predisposition characterized by exome sequencing analysis. Journal of Genetics and Genomics, 2018, 45, 41-45.	1.7	11
140	Colorectal cancer genetic variants are also associated with serrated polyposis syndrome susceptibility. Journal of Medical Genetics, 2020, 57, 677-682.	1.5	11
141	Expression profile of circulating microRNAs in the Correa pathway of progression to gastric cancer. United European Gastroenterology Journal, 2018, 6, 691-701.	1.6	10
142	Comparison of biomolecules on the basis of Molecular Interaction Potentials. Journal of the Brazilian Chemical Society, 2002, 13, 795-799.	0.6	10
143	Validation of miR-1228-3p as Housekeeping for MicroRNA Analysis in Liquid Biopsies from Colorectal Cancer Patients. Biomolecules, 2020, 10, 16.	1.8	9
144	Multi-omic modelling of inflammatory bowel disease with regularized canonical correlation analysis. PLoS ONE, 2021, 16, e0246367.	1.1	9

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145	Multiple Sporadic Colorectal Cancers Display a Unique Methylation Phenotype. PLoS ONE, 2014, 9, e91033.	1.1	9
146	Use of alignment-free molecular descriptors in diversity analysis and optimal sampling of molecular libraries. Molecular Diversity, 2000, 6, 135-147.	2.1	8
147	Gene Expression Profiling and Transplantation Tolerance in the Clinic. Transplantation, 2009, 88, S50-S53.	0.5	8
148	Serum transferrin as a biomarker of hepatocyte nuclear factor 4 alpha activity and hepatocyte function in liver diseases. BMC Medicine, 2021, 19, 39.	2.3	8
149	Identification of New Genes Involved in Germline Predisposition to Early-Onset Gastric Cancer. International Journal of Molecular Sciences, 2021, 22, 1310.	1.8	8
150	Deciphering microRNA targets in pancreatic cancer using miRComb R package. Oncotarget, 2018, 9, 6499-6517.	0.8	8
151	Peripheral phenotype and gene expression profiles of combined liver–kidney transplant patients. Liver International, 2016, 36, 401-409.	1.9	7
152	Cell Plasticity-Related Phenotypes and Taxanes Resistance in Castration-Resistant Prostate Cancer. Frontiers in Oncology, 2020, 10, 594023.	1.3	7
153	Differential gene expression profile between progressive and de novo muscle invasive bladder cancer and its prognostic implication. Scientific Reports, 2021, 11, 6132.	1.6	7
154	AMarge. Applied Bioinformatics, 2006, 5, 45-47.	1.7	6
155	Infrequent Loss of Luminal Differentiation in Ductal Breast Cancer Metastasis. PLoS ONE, 2013, 8, e78097.	1.1	6
156	Quantitative analysis of somatically acquired and constitutive uniparental disomy in gastrointestinal cancers. International Journal of Cancer, 2019, 144, 513-524.	2.3	6
157	Copyâ€number intratumor heterogeneity increases the risk of relapse in chemotherapyâ€naive stage <scp>ll</scp> colon cancer. Journal of Pathology, 2022, 257, 68-81.	2.1	6
158	Ability of a urine gene expression classifier to reduce the number of follow-up cystoscopies in bladder cancer patients. Translational Research, 2019, 208, 73-84.	2.2	5
159	Variability in Cerebrospinal Fluid MicroRNAs Through Life. Molecular Neurobiology, 2020, 57, 4134-4142.	1.9	5
160	Defining a Methylation Signature Associated With Operational Tolerance in Kidney Transplant Recipients. Frontiers in Immunology, 2021, 12, 709164.	2.2	5
161	Treatment With Simvastatin and Rifaximin Restores the Plasma Metabolomic Profile in Patients With Decompensated Cirrhosis. Hepatology Communications, 2022, 6, 1100-1112.	2.0	5
162	Differential gene expression induced by growth hormone treatment in the uremic rat growth plate. Growth Hormone and IGF Research, 2008, 18, 353-359.	0.5	4

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163	Pharmacogenomic analyzis of the responsiveness of gastrointestinal tumor cell lines to drug therapy: A transportome approach. Pharmacological Research, 2016, 113, 364-375.	3.1	4
164	Molecular Diversity Sample Generation on the Basis of Quantum-Mechanical Computations and Principal Component Analysis. Combinatorial Chemistry and High Throughput Screening, 2002, 5, 49-57.	0.6	3
165	Urine Gene Expression Profiles in Bladder Pain Syndrome Patients Treated with Triamcinolone. European Urology Focus, 2020, 6, 390-396.	1.6	3
166	QSAR in the nonsteroidal anti-inflammatory agents: The fenamic acids. , 1993, , 560-561.		3
167	Clinicopathological and Molecular Prognostic Classifier for Intermediate/High-Risk Clear Cell Renal Cell Carcinoma. Cancers, 2021, 13, 6338.	1.7	2
168	Coding and non-coding co-expression network analysis identifies key modules and driver genes associated with precursor lesions of gastric cancer. Genomics, 2022, 114, 110370.	1.3	2
169	DNA Damage Regulates Alternative Splicing through Inhibition of RNA Polymerase II Elongation. Cell, 2009, 139, 211.	13.5	1
170	Cross-sectional study of human coding- and non-coding RNAs in progressive stages of Helicobacter pylori infection. Scientific Data, 2020, 7, 296.	2.4	1
171	Identification of circulating microRNAs for the diagnosis of early-stage gastric cancer Journal of Clinical Oncology, 2017, 35, 33-33.	0.8	1
172	Validation of Urine-based Gene Classifiers for Detecting Bladder Cancer in a Chinese Study. Journal of Cancer, 2018, 9, 3208-3215.	1.2	0
173	MicroRNAs Deregulated in Intraductal Papillary Mucinous Neoplasm Converge on Actin Cytoskeleton-Related Pathways That Are Maintained in Pancreatic Ductal Adenocarcinoma. Cancers, 2021, 13, 2369.	1.7	0
174	3D QSAR on Mutagenic Heterocyclic Amines That are Substrates of Cytochrome P450 1A2., 2000,, 321-322.		0
175	Pharmacophore Development for the Interaction of Cytochrome P450 1A2 with Its Substrates and Inhibitors., 2000,, 141-146.		0
176	Gene Expression Profiling Distinguishes Essential Thrombocythemia from Polycythemia Vera Patients and Identifies a Common Expressed Set of Genes in Relation to JAK2V617F Status. Blood, 2008, 112, 2788-2788.	0.6	0