

Jerzy Ostrowski

List of Publications by Year in descending order

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131
papers

4,031
citations

136950

32
h-index

144013

57
g-index

131
all docs

131
docs citations

131
times ranked

7020
citing authors

#	ARTICLE	IF	CITATIONS
1	Prediction of fetal blood group antigens from maternal plasma using Ion AmpliSeq HD technology. <i>Transfusion</i> , 2022, 62, 458-468.	1.6	3
2	Oncology Drug Repurposing for Sepsis Treatment. <i>Biomedicines</i> , 2022, 10, 921.	3.2	5
3	Combination of HLA-DQ2/-DQ8 Haplotypes and a Single MSH5 Gene Variant in a Polish Population of Patients with Type 1 Diabetes as a First Line Screening for Celiac Disease?. <i>Journal of Clinical Medicine</i> , 2022, 11, 2223.	2.4	1
4	Discovery of indazole-pyridinone derivatives as a novel class of potent and selective MNK1/2 kinase inhibitors that protecting against endotoxin-induced septic shock. <i>European Journal of Medicinal Chemistry</i> , 2021, 213, 113057.	5.5	5
5	Higher genome variability within metabolism genes associates with recurrent <i>Clostridium difficile</i> infection. <i>BMC Microbiology</i> , 2021, 21, 36.	3.3	7
6	Peripheral Blood Cells from Patients with Hodgkinâ€™s and Diffuse Large B Cell Lymphomas May Be a Better Source of Candidate Diagnostic miRNAs Than Circulating miRNAs. <i>BioMed Research International</i> , 2021, 2021, 1-9.	1.9	0
7	Cytotoxic Efficacy and Resistance Mechanism of a TRAIL and VEGFA-Peptide Fusion Protein in Colorectal Cancer Models. <i>International Journal of Molecular Sciences</i> , 2021, 22, 3160.	4.1	2
8	Multi-Organ Transcriptome Dynamics in a Mouse Model of Cecal Ligation and Puncture-Induced Polymicrobial Sepsis. <i>Journal of Inflammation Research</i> , 2021, Volume 14, 2377-2388.	3.5	14
9	The gastric microbiota in patients with Crohnâ€™s disease; a preliminary study. <i>Scientific Reports</i> , 2021, 11, 17866.	3.3	4
10	Selective Extracellular Signal-Regulated Kinase 1/2 (ERK1/2) Inhibition by the SCH772984 Compound Attenuates In Vitro and In Vivo Inflammatory Responses and Prolongs Survival in Murine Sepsis Models. <i>International Journal of Molecular Sciences</i> , 2021, 22, 10204.	4.1	8
11	Limited Practical Utility of Liquid Biopsy in the Treated Patients with Advanced Breast Cancer. <i>Diagnostics</i> , 2020, 10, 523.	2.6	2
12	In search for interplay between stool microRNAs, microbiota and short chain fatty acids in Crohnâ€™s disease - a preliminary study. <i>BMC Gastroenterology</i> , 2020, 20, 307.	2.0	12
13	PARP Inhibition Increases the Reliance on ATR/CHK1 Checkpoint Signaling Leading to Synthetic Lethalityâ€™ An Alternative Treatment Strategy for Epithelial Ovarian Cancer Cells Independent from HR Effectiveness. <i>International Journal of Molecular Sciences</i> , 2020, 21, 9715.	4.1	20
14	The Search of miRNA Related to Invasive Growth of Nonfunctioning Gonadotrophic Pituitary Tumors. <i>International Journal of Endocrinology</i> , 2020, 2020, 1-8.	1.5	4
15	DNA Methylation Influences miRNA Expression in Gonadotroph Pituitary Tumors. <i>Life</i> , 2020, 10, 59.	2.4	4
16	Challenges in Cancer Biomarker Discovery Exemplified by the Identification of Diagnostic MicroRNAs in Prostate Tissues. <i>BioMed Research International</i> , 2020, 2020, 1-4.	1.9	9
17	The composition and richness of the gut microbiota differentiate the top Polish endurance athletes from sedentary controls. <i>Gut Microbes</i> , 2020, 11, 1374-1384.	9.8	48
18	Differences between Well-Differentiated Neuroendocrine Tumors and Ductal Adenocarcinomas of the Pancreas Assessed by Multi-Omics Profiling. <i>International Journal of Molecular Sciences</i> , 2020, 21, 4470.	4.1	10

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19	Mutational Mosaics of Cell-Free DNA from Pancreatic Cyst Fluids. <i>Digestive Diseases and Sciences</i> , 2020, 65, 2294-2301.	2.3	6
20	Transcription Factor Prospero Homeobox 1 (PROX1) as a Potential Angiogenic Regulator of Follicular Thyroid Cancer Dissemination. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5619.	4.1	18
21	The Role of Aberrant DNA Methylation in Misregulation of Gene Expression in Gonadotroph Nonfunctioning Pituitary Tumors. <i>Cancers</i> , 2019, 11, 1650.	3.7	12
22	Clinical significance of endoscopic findings in the upper gastrointestinal tract in Crohn's disease. <i>Scandinavian Journal of Gastroenterology</i> , 2019, 54, 1075-1080.	1.5	5
23	Common functional alterations identified in blood transcriptome of autoimmune cholestatic liver and inflammatory bowel diseases. <i>Scientific Reports</i> , 2019, 9, 7190.	3.3	18
24	Molecular Signature of Prospero Homeobox 1 (PROX1) in Follicular Thyroid Carcinoma Cells. <i>International Journal of Molecular Sciences</i> , 2019, 20, 2212.	4.1	6
25	Mutation Profiling of Premalignant Colorectal Neoplasia. <i>Gastroenterology Research and Practice</i> , 2019, 2019, 1-9.	1.5	5
26	Whole-exome sequencing identifies novel pathogenic variants across the <i>ATP7B</i> gene and some modifiers of Wilson's disease phenotype. <i>Liver International</i> , 2019, 39, 177-186.	3.9	38
27	Redefining the Practical Utility of Blood Transcriptome Biomarkers in Inflammatory Bowel Diseases. <i>Journal of Crohn's and Colitis</i> , 2019, 13, 626-633.	1.3	22
28	Prediction of fetal blood group and platelet antigens from maternal plasma using next-generation sequencing. <i>Transfusion</i> , 2019, 59, 1102-1107.	1.6	31
29	Transcriptomes of human mesenchymal cells isolated from the right ventricle and epicardial fat differ strikingly both directly after isolation and long-term culture. <i>ESC Heart Failure</i> , 2019, 6, 351-361.	3.1	4
30	Functional analyses of a low-penetrance risk variant rs6702619/1p21.2 associating with colorectal cancer in Polish population. <i>Acta Biochimica Polonica</i> , 2019, 66, 365-370.	0.5	9
31	Candidate diagnostic miRNAs that can detect cancer in prostate biopsy. <i>Prostate</i> , 2018, 78, 178-185.	2.3	24
32	Challenges in Stratifying the Molecular Variability of Patient-Derived Colon Tumor Xenografts. <i>BioMed Research International</i> , 2018, 2018, 1-9.	1.9	18
33	RNA sequencing reveals widespread transcriptome changes in a renal carcinoma cell line. <i>Oncotarget</i> , 2018, 9, 8597-8613.	1.8	22
34	Mass Spectrometry-Based Comprehensive Analysis of Pancreatic Cyst Fluids. <i>BioMed Research International</i> , 2018, 2018, 1-12.	1.9	9
35	Exome scale map of genetic alterations promoting metastasis in colorectal cancer. <i>BMC Genetics</i> , 2018, 19, 85.	2.7	22
36	Noninvasive prenatal HPA typing in HPA negative pregnancies selected in the Polish PREVFNAIT screening program. <i>Transfusion</i> , 2018, 58, 2705-2711.	1.6	8

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37	IDH1/2 Mutations Predict Shorter Survival in Chondrosarcoma. <i>Journal of Cancer</i> , 2018, 9, 998-1005.	2.5	50
38	A preliminary evaluation of next-generation sequencing as a screening tool for targeted genotyping of erythrocyte and platelet antigens in blood donors. <i>Blood Transfusion</i> , 2018, 16, 285-292.	0.4	23
39	PALB2 mutations in BRCA1/2-mutation negative breast and ovarian cancer patients from Poland. <i>BMC Medical Genomics</i> , 2017, 10, 14.	1.5	17
40	A Strong Neutrophil Elastase Proteolytic Fingerprint Marks the Carcinoma Tumor Proteome. <i>Molecular and Cellular Proteomics</i> , 2017, 16, 213-227.	3.8	17
41	A heterozygous mutation in GOT1 is associated with familial macro-aspartate aminotransferase. <i>Journal of Hepatology</i> , 2017, 67, 1026-1030.	3.7	18
42	A novel approach to genome-wide association analysis identifies genetic associations with primary biliary cholangitis and primary sclerosing cholangitis in Polish patients. <i>BMC Medical Genomics</i> , 2017, 10, 2.	1.5	18
43	Clinical Applicability of Whole-Exome Sequencing Exemplified by a Study in Young Adults with the Advanced Cryptogenic Cholestatic Liver Diseases. <i>Gastroenterology Research and Practice</i> , 2017, 2017, 1-8.	1.5	5
44	Downregulation of PTPRH (Sap-1) in colorectal tumors. <i>International Journal of Oncology</i> , 2017, 51, 841-850.	3.3	6
45	Antimelanomic Effects of High- and Low-Molecular Weight Bioactive Subfractions Isolated from the Mossy Maze Mushroom, <i>Cerrena unicolor</i> (Agaricomycetes). <i>International Journal of Medicinal Mushrooms</i> , 2017, 19, 619-628.	1.5	7
46	SEL120-34A is a novel CDK8 inhibitor active in AML cells with high levels of serine phosphorylation of STAT1 and STAT5 transactivation domains. <i>Oncotarget</i> , 2017, 8, 33779-33795.	1.8	70
47	Can genetic testing be useful for defining the risk of cancer?. <i>Nowotwory</i> , 2017, 66, 422-426.	0.3	0
48	Effect of <i>Saccharomyces boulardii</i> and Mode of Delivery on the Early Development of the Gut Microbial Community in Preterm Infants. <i>PLoS ONE</i> , 2016, 11, e0150306.	2.5	29
49	Genetic architecture differences between pediatric and adult-onset inflammatory bowel diseases in the Polish population. <i>Scientific Reports</i> , 2016, 6, 39831.	3.3	33
50	Limited prolonged effects of rifaximin treatment on irritable bowel syndrome-related differences in the fecal microbiome and metabolome. <i>Gut Microbes</i> , 2016, 7, 397-413.	9.8	68
51	Prolonged transfer of feces from the lean mice modulates gut microbiota in obese mice. <i>Nutrition and Metabolism</i> , 2016, 13, 57.	3.0	55
52	Expression changes of cell-cell adhesion-related genes in colorectal tumors. <i>Oncology Letters</i> , 2015, 9, 2463-2470.	1.8	60
53	Regulation of the expression of claudin 23 by the enhancer of zeste 2 polycomb group protein in colorectal cancer. <i>Molecular Medicine Reports</i> , 2015, 12, 728-736.	2.4	23
54	Beads-free protein immunoprecipitation for a mass spectrometry-based interactome and posttranslational modifications analysis. <i>Proteome Science</i> , 2015, 13, 23.	1.7	14

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55	Combination Testing Using a Single MSH5 Variant alongside HLA Haplotypes Improves the Sensitivity of Predicting Coeliac Disease Risk in the Polish Population. <i>PLoS ONE</i> , 2015, 10, e0139197.	2.5	6
56	Epigenetic-Mediated Downregulation of α -Protocadherin in Colorectal Tumours. <i>Gastroenterology Research and Practice</i> , 2015, 2015, 1-9.	1.5	14
57	Transcriptional changes between uninfamed ulcerative colitis and familial adenomatous polyposis pouch mucosa can be attributed to an altered immune response. <i>Acta Biochimica Polonica</i> , 2015, 62, 69-75.	0.5	7
58	New recurrent BRCA1/2 mutations in Polish patients with familial breast/ovarian cancer detected by next generation sequencing. <i>BMC Medical Genomics</i> , 2015, 8, 19.	1.5	34
59	PDL1 expression is an independent prognostic factor in localized GIST. <i>Oncolmmunology</i> , 2015, 4, e1002729.	4.6	75
60	Limited predictive value of achieving beneficial plasma (Z)-endoxifen threshold level by CYP2D6 genotyping in tamoxifen-treated Polish women with breast cancer. <i>BMC Cancer</i> , 2015, 15, 570.	2.6	17
61	Extracellular matrix and cytochrome P450 gene expression can distinguish steatohepatitis from steatosis in mice. <i>Journal of Cellular and Molecular Medicine</i> , 2014, 18, 1762-1772.	3.6	16
62	Mitochondrial-related proteomic changes during obesity and fasting in mice are greater in the liver than skeletal muscles. <i>Functional and Integrative Genomics</i> , 2014, 14, 245-259.	3.5	25
63	Histone H3 lysine 27 acetylation is altered in colon cancer. <i>Clinical Proteomics</i> , 2014, 11, 24.	2.1	72
64	CYP1A1 Ile462Val polymorphism and colorectal cancer risk in Polish patients. <i>Medical Oncology</i> , 2014, 31, 72.	2.5	7
65	Obesity increases histone H3 lysine 9 and 18 acetylation at <i>Tnfa</i> and <i>Ccl2</i> genes in mouse liver. <i>International Journal of Molecular Medicine</i> , 2014, 34, 1647-1654.	4.0	51
66	The N-Reductive System Composed of Mitochondrial Amidoxime Reducing Component (mARC), Cytochrome b5 (CYB5B) and Cytochrome b5 Reductase (CYB5R) Is Regulated by Fasting and High Fat Diet in Mice. <i>PLoS ONE</i> , 2014, 9, e105371.	2.5	21
67	Association of the BRCA1 promoter polymorphism rs11655505 with the risk of familial breast and/or ovarian cancer. <i>Familial Cancer</i> , 2013, 12, 691-698.	1.9	2
68	Heterogeneous Nuclear Ribonucleoprotein (HnRNP) K Genome-wide Binding Survey Reveals Its Role in Regulating 3'-End RNA Processing and Transcription Termination at the Early Growth Response 1 (EGR1) Gene through XRN2 Exonuclease*. <i>Journal of Biological Chemistry</i> , 2013, 288, 24788-24798.	3.4	28
69	Common low-penetrance risk variants associated with breast cancer in Polish women. <i>BMC Cancer</i> , 2013, 13, 510.	2.6	19
70	Pre-analytical-related variability influencing serum peptide profiles demonstrated in a mass spectrometry-based search for colorectal and prostate cancer biomarkers. <i>Acta Biochimica Polonica</i> , 2013, 60, 417-25.	0.5	8
71	Pooled Sample-Based GWAS: A Cost-Effective Alternative for Identifying Colorectal and Prostate Cancer Risk Variants in the Polish Population. <i>PLoS ONE</i> , 2012, 7, e35307.	2.5	34
72	Identification of a Late Onset Alzheimer's Disease Candidate Risk Variant at 9q21.33 in Polish Patients. <i>Journal of Alzheimer's Disease</i> , 2012, 32, 157-168.	2.6	18

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73	High frequency of BRCA1 founder mutations in Polish women with nonfamilial breast cancer. <i>Familial Cancer</i> , 2012, 11, 623-628.	1.9	20
74	Comparative kinome analysis to identify putative colon tumor biomarkers. <i>Journal of Molecular Medicine</i> , 2012, 90, 447-456.	3.9	34
75	Lack of evidence for increased level of circulating urothelial cells in the peripheral blood after transurethral resection of bladder tumors. <i>International Urology and Nephrology</i> , 2012, 44, 761-767.	1.4	11
76	Inferring serum proteolytic activity from LC-MS/MS data. <i>BMC Bioinformatics</i> , 2012, 13, S7.	2.6	4
77	Integrating proteomic and transcriptomic high-throughput surveys for search of new biomarkers of colon tumors. <i>Functional and Integrative Genomics</i> , 2011, 11, 215-224.	3.5	97
78	Inferring serum proteolytic activity from LC-MS/MS data. , 2011, , .		0
79	Halogenated imidazole derivatives block RNA polymerase II elongation along mitogen inducible genes. <i>BMC Molecular Biology</i> , 2010, 11, 4.	3.0	7
80	Comprehensive Analysis of the Palindromic Motif TCTCGCGAGA: A Regulatory Element of the HNRNPK Promoter. <i>DNA Research</i> , 2010, 17, 245-260.	3.4	39
81	Modeling Oncogenic Signaling in Colon Tumors by Multidirectional Analyses of Microarray Data Directed for Maximization of Analytical Reliability. <i>PLoS ONE</i> , 2010, 5, e13091.	2.5	320
82	Two-Stage Model-Based Clustering for Liquid Chromatography Mass Spectrometry Data Analysis. <i>Statistical Applications in Genetics and Molecular Biology</i> , 2009, 8, 1-34.	0.6	5
83	Functional features of gene expression profiles differentiating gastrointestinal stromal tumours according to KIT mutations and expression. <i>BMC Cancer</i> , 2009, 9, 413.	2.6	21
84	An integrated LC-ESI-MS platform for quantitation of serum peptide ladders. Application for colon carcinoma study. <i>Proteomics - Clinical Applications</i> , 2009, 3, 932-946.	1.6	45
85	Integrating genomics, proteomics and bioinformatics in translational studies of molecular medicine. <i>Expert Review of Molecular Diagnostics</i> , 2009, 9, 623-630.	3.1	34
86	Tip-I \pm (hp0596 Gene Product) Is a Highly Immunogenic Helicobacter pylori Protein Involved in Colonization of Mouse Gastric Mucosa. <i>Current Microbiology</i> , 2008, 56, 279-286.	2.2	21
87	Large intra- and inter-individual variability of genes expression levels limits potential predictive value of molecular diagnosis of dysplasia in Barrett's esophagus. <i>Journal of Molecular Medicine</i> , 2008, 86, 233-242.	3.9	8
88	Lack of evidence for association of primary sclerosing cholangitis and primary biliary cirrhosis with risk alleles for Crohn's disease in Polish patients. <i>BMC Medical Genetics</i> , 2008, 9, 81.	2.1	27
89	Limited clinical relevance of mitochondrial DNA mutation and gene expression analyses in ovarian cancer. <i>BMC Cancer</i> , 2008, 8, 292.	2.6	30
90	Association between gynecomastia and aromatase (CYP19) polymorphisms. <i>European Journal of Endocrinology</i> , 2008, 158, 721-727.	3.7	36

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91	The fold recognition of CP2 transcription factors gives new insights into the function and evolution of tumor suppressor protein p53. <i>Cell Cycle</i> , 2008, 7, 2907-2915.	2.6	39
92	Automated reduction and interpretation of multidimensional mass spectra for analysis of complex peptide mixtures. <i>International Journal of Mass Spectrometry</i> , 2007, 260, 20-30.	1.5	16
93	Molecular defense mechanisms of Barrett's metaplasia estimated by an integrative genomics. <i>Journal of Molecular Medicine</i> , 2007, 85, 733-743.	3.9	44
94	The diverse involvement of heterogeneous nuclear ribonucleoprotein K in mitochondrial response to insulin. <i>FEBS Letters</i> , 2006, 580, 1839-1845.	2.8	9
95	<i>Helicobacter pylori</i> protein oxidation influences the colonization process. <i>International Journal of Medical Microbiology</i> , 2006, 296, 321-324.	3.6	24
96	Landscape of the hnRNP K protein's protein interactome. <i>Proteomics</i> , 2006, 6, 2395-2406.	2.2	69
97	Casein kinases phosphorylate multiple residues spanning the entire hnRNP K length. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2006, 1764, 299-306.	2.3	12
98	Three clinical variants of gastroesophageal reflux disease form two distinct gene expression signatures. <i>Journal of Molecular Medicine</i> , 2006, 84, 872-882.	3.9	11
99	<i>Helicobacter pylori</i> VacA cytotoxin interacts with fibronectin and alters HeLa cell adhesion and cytoskeletal organization in vitro. <i>FEMS Immunology and Medical Microbiology</i> , 2005, 44, 143-150.	2.7	31
100	Increased mitochondrial gene expression during L6 cell myogenesis is accelerated by insulin. <i>International Journal of Biochemistry and Cell Biology</i> , 2005, 37, 1815-1828.	2.8	17
101	Heterogeneous Nuclear Ribonucleoprotein K Enhances Insulin-induced Expression of Mitochondrial UCP2 Protein. <i>Journal of Biological Chemistry</i> , 2004, 279, 54599-54609.	3.4	28
102	hnRNP K: One protein multiple processes. <i>BioEssays</i> , 2004, 26, 629-638.	2.5	419
103	Cooperative binding of the hnRNP K three KH domains to mRNA targets. <i>FEBS Letters</i> , 2004, 577, 134-140.	2.8	56
104	Characterization of hnRNP K Protein's RNA Interactions. <i>Journal of Molecular Biology</i> , 2004, 342, 1131-1141.	4.2	46
105	Members of the poly (rC) binding protein family stimulate the activity of the c-myc internal ribosome entry segment in vitro and in vivo. <i>Oncogene</i> , 2003, 22, 8012-8020.	5.9	205
106	Quantitative detection for low levels of <i>Helicobacter pylori</i> infection in experimentally infected mice by real-time PCR. <i>Journal of Microbiological Methods</i> , 2003, 55, 351-359.	1.6	65
107	Homoplasmic MELAS A3243G mtDNA mutation in a colon cancer sample. <i>Mitochondrion</i> , 2003, 3, 119-124.	3.4	34
108	Transient recruitment of the hnRNP K protein to inducibly transcribed gene loci. <i>Nucleic Acids Research</i> , 2003, 31, 3954-3962.	14.5	48

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109	Multivariate Analysis of Risk Factors for Development of Duodenal Ulcer in <i>Helicobacter pylori</i> -Infected Patients. <i>Digestion</i> , 2003, 67, 25-31.	2.3	11
110	Heterogeneous Nuclear Ribonucleoprotein K Protein Associates with Multiple Mitochondrial Transcripts within the Organelle. <i>Journal of Biological Chemistry</i> , 2002, 277, 6303-6310.	3.4	50
111	The <i>hppA</i> gene of <i>Helicobacter pylori</i> encodes the class C acid phosphatase precursor. <i>FEBS Letters</i> , 2002, 525, 39-42.	2.8	7
112	RACK1 Protein Interacts with <i>Helicobacter pylori</i> VacA Cytotoxin: The Yeast Two-Hybrid Approach. <i>Biochemical and Biophysical Research Communications</i> , 2001, 289, 103-110.	2.1	26
113	Peptide analog of fibronectin that inhibits cell migration and ERK 1/2 activity. <i>Peptides</i> , 2001, 22, 1949-1953.	2.4	7
114	Mutations, methylation and expression of CDKN2a/p16 gene in colorectal cancer and normal colonic mucosa. <i>Cancer Letters</i> , 2001, 163, 17-23.	7.2	20
115	p53 protein accumulation and p53 gene mutation in colorectal cancer. <i>Pathology and Oncology Research</i> , 2000, 6, 275-279.	1.9	16
116	Role of Tyrosine Phosphorylation in the Regulation of the Interaction of Heterogeneous Nuclear Ribonucleoprotein K Protein with Its Protein and RNA Partners. <i>Journal of Biological Chemistry</i> , 2000, 275, 3619-3628.	3.4	74
117	In Search of Immunogenic <i>Helicobacter pylori</i> Proteins by Screening of Expression Library. <i>Digestion</i> , 2000, 61, 14-21.	2.3	10
118	Treatment of mice with EGF and orthovanadate activates cytoplasmic and nuclear MAPK, p70S6k, and p90rsk in the liver. <i>Journal of Hepatology</i> , 2000, 32, 965-974.	3.7	15
119	Stimulation of p85/RING3 kinase in multiple organs after systemic administration of mitogens into mice. <i>Oncogene</i> , 1998, 16, 1223-1227.	5.9	24
120	Diverse molecular interactions of the hnRNP K protein. <i>FEBS Letters</i> , 1997, 403, 113-115.	2.8	151
121	Zik1, a Transcriptional Repressor That Interacts with the Heterogeneous Nuclear Ribonucleoprotein Particle K Protein. <i>Journal of Biological Chemistry</i> , 1996, 271, 27701-27706.	3.4	73
122	The K Protein Domain That Recruits the Interleukin 1-responsive K Protein Kinase Lies Adjacent to a Cluster of c-Src and Vav SH3-binding Sites. <i>Journal of Biological Chemistry</i> , 1995, 270, 26976-26985.	3.4	104
123	Description of an IL-1-Responsive Kinase That Phosphorylates the K Protein. Enhancement of Phosphorylation by Selective DNA and RNA Motifs. <i>Biochemistry</i> , 1995, 34, 5644-5650.	2.5	45
124	Inhibition of acid secretory response and induction of ornithine decarboxylase and its mRNA by TGF β and EGF in isolated rat gastric glands. <i>Regulatory Peptides</i> , 1995, 56, 1-8.	1.9	10
125	Does Transmembrane Potential Control Acid Secretion in the Stomach?. <i>Electromagnetic Biology and Medicine</i> , 1994, 13, 93-97.	0.4	0
126	Purification of DNA-binding proteins using tandem DNA- affinity column. <i>Nucleic Acids Research</i> , 1993, 21, 1045-1046.	14.5	14

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127	Generation of phosphatidic acid and diacylglycerols following ligation of surface immunoglobulin in human B lymphocytes: Potential role in PKC activation. Cellular Immunology, 1992, 141, 373-387.	3.0	7
128	Weak bases increase surface IgM expression in 70Z/3 B lymphoid cell line without increasing \hat{I}^{β} gene expression. Cellular Immunology, 1990, 130, 11-21.	3.0	0
129	Local Anesthetics (Benzyl Alcohol, Lidocaine, Procainamide) Inhibit Aminopyrine Accumulation in Isolated Rat Parietal Cells. Pharmacology, 1989, 39, 265-272.	2.2	1
130	Erythrocyte Porphobilinogen Deaminase Activity in Lives Disease. Gastroenterology, 1987, 92, 845-851.	1.3	2
131	Abnormalities in Liver Function and Morphology and Impaired Aminopyrine Metabolism in Hereditary Hepatic Porphyrias. Gastroenterology, 1983, 85, 1131-1137.	1.3	30