

Mario Medvedovic

List of Publications by Year in descending order

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Version: 2024-02-01

123
papers

6,529
citations

66343

42
h-index

76900

74
g-index

132
all docs

132
docs citations

132
times ranked

10858
citing authors

#	ARTICLE	IF	CITATIONS
1	ALOHA: Aggregated local extrema splines for high-throughput dose-response analysis. <i>Computational Toxicology</i> , 2022, 21, 100196.	3.3	0
2	Phase II Clinical Trial of Neoadjuvant and Adjuvant Pembrolizumab in Resectable Locally Advanced Head and Neck Squamous Cell Carcinoma. <i>Clinical Cancer Research</i> , 2022, 28, 1345-1352.	7.0	38
3	SigCom LINCS: data and metadata search engine for a million gene expression signatures. <i>Nucleic Acids Research</i> , 2022, 50, W697-W709.	14.5	26
4	Melanoma Cell Intrinsic GABAA Receptor Enhancement Potentiates Radiation and Immune Checkpoint Inhibitor Response by Promoting Direct and T Cell-Mediated Antitumor Activity. <i>International Journal of Radiation Oncology Biology Physics</i> , 2021, 109, 1040-1053.	0.8	18
5	Monocyte and bone marrow macrophage transcriptional phenotypes in systemic juvenile idiopathic arthritis reveal TRIM8 as a mediator of IFN- γ hyper-responsiveness and risk for macrophage activation syndrome. <i>Annals of the Rheumatic Diseases</i> , 2021, 80, 617-625.	0.9	31
6	Hexavalent chromium promotes differential binding of CTCF to its cognate sites in Euchromatin. <i>Epigenetics</i> , 2021, 16, 1-16.	2.7	3
7	Connectivity Map Analysis of a Single-Cell RNA-Sequencing -Derived Transcriptional Signature of mTOR Signaling. <i>International Journal of Molecular Sciences</i> , 2021, 22, 4371.	4.1	8
8	Tobacco smoking induces metabolic reprogramming of renal cell carcinoma. <i>Journal of Clinical Investigation</i> , 2021, 131, .	8.2	14
9	LINCS Data Portal 2.0: next generation access point for perturbation-response signatures. <i>Nucleic Acids Research</i> , 2020, 48, D431-D439.	14.5	112
10	Dioxin Disrupts Dynamic DNA Methylation Patterns in Genes That Govern Cardiomyocyte Maturation. <i>Toxicological Sciences</i> , 2020, 178, 325-337.	3.1	7
11	Predicting mechanism of action of cellular perturbations with pathway activity signatures. <i>Bioinformatics</i> , 2020, 36, 4781-4788.	4.1	6
12	Low-Dose Bisphenol A in a Rat Model of Endometrial Cancer: A CLARITY-BPA Study. <i>Environmental Health Perspectives</i> , 2020, 128, 127005.	6.0	15
13	piNET: a versatile web platform for downstream analysis and visualization of proteomics data. <i>Nucleic Acids Research</i> , 2020, 48, W85-W93.	14.5	18
14	Mass spectrometry proteomics reveals a function for mammalian CALCOCO1 in MTOR-regulated selective autophagy. <i>Autophagy</i> , 2020, 16, 2219-2237.	9.1	37
15	Comparability of the small RNA secretome across human biofluids concomitantly collected from healthy adults. <i>PLoS ONE</i> , 2020, 15, e0229976.	2.5	11
16	Interferon-complement loop in transplant-associated thrombotic microangiopathy. <i>Blood Advances</i> , 2020, 4, 1166-1177.	5.2	41
17	Comprehensive mapping of the methylation landscape of 16 CpG-dense regions in oral and pharyngeal squamous cell carcinoma. <i>Epigenomics</i> , 2019, 11, 987-1002.	2.1	3
18	GREIN: An Interactive Web Platform for Re-analyzing GEO RNA-seq Data. <i>Scientific Reports</i> , 2019, 9, 7580.	3.3	126

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19	ChIAPoP: a new tool for ChIA-PET data analysis. <i>Nucleic Acids Research</i> , 2019, 47, e37-e37.	14.5	9
20	Balancing yield, purity and practicality: a modified differential ultracentrifugation protocol for efficient isolation of small extracellular vesicles from human serum. <i>RNA Biology</i> , 2019, 16, 5-12.	3.1	52
21	An Organoid-Based Preclinical Model of Human Gastric Cancer. <i>Cellular and Molecular Gastroenterology and Hepatology</i> , 2019, 7, 161-184.	4.5	97
22	The N ⁶ -Methyladenosine mRNA Methylase METTL3 Controls Cardiac Homeostasis and Hypertrophy. <i>Circulation</i> , 2019, 139, 533-545.	1.6	279
23	MicroRNA networks associated with active systemic juvenile idiopathic arthritis regulate CD163 expression and anti-inflammatory functions in macrophages through two distinct mechanisms. <i>Journal of Leukocyte Biology</i> , 2018, 103, 71-85.	3.3	19
24	Differential expression and prognostic value of long non-coding RNA in HPV-negative head and neck squamous cell carcinoma. <i>Head and Neck</i> , 2018, 40, 1555-1564.	2.0	28
25	Loss of microRNA-128 promotes cardiomyocyte proliferation and heart regeneration. <i>Nature Communications</i> , 2018, 9, 700.	12.8	124
26	The Library of Integrated Network-Based Cellular Signatures NIH Program: System-Level Cataloging of Human Cells Response to Perturbations. <i>Cell Systems</i> , 2018, 6, 13-24.	6.2	327
27	Ultra-Deep Genomic Sequencing of HCV NS5A Resistance-Associated Substitutions in HCV/HIV Coinfected Patients. <i>Digestive Diseases and Sciences</i> , 2018, 63, 645-652.	2.3	1
28	Identification of sex-specific DNA methylation changes driven by specific chemicals in cord blood in a Faroese birth cohort. <i>Epigenetics</i> , 2018, 13, 290-300.	2.7	62
29	Chromium disrupts chromatin organization and CTCF access to its cognate sites in promoters of differentially expressed genes. <i>Epigenetics</i> , 2018, 13, 363-375.	2.7	21
30	Sustainable data and metadata management at the BD2K-LINCS Data Coordination and Integration Center. <i>Scientific Data</i> , 2018, 5, 180117.	5.3	22
31	Data Portal for the Library of Integrated Network-based Cellular Signatures (LINCS) program: integrated access to diverse large-scale cellular perturbation response data. <i>Nucleic Acids Research</i> , 2018, 46, D558-D566.	14.5	143
32	Interleukin-22 levels are increased in gastrointestinal graft-versus-host disease in children. <i>Haematologica</i> , 2018, 103, e480-e482.	3.5	7
33	NBCe1 Na ⁺ -HCO ₃ ⁻ cotransporter ablation causes reduced apoptosis following cardiac ischemia-reperfusion injury <i>in vivo</i> . <i>World Journal of Cardiology</i> , 2018, 10, 97-109.	1.5	15
34	CD44 variant isoform 9 emerges in response to injury and contributes to the regeneration of the gastric epithelium. <i>Journal of Pathology</i> , 2017, 242, 463-475.	4.5	41
35	Genetic susceptibility to toxicologic lung responses among inbred mouse strains following exposure to carbon nanotubes and profiling of underlying gene networks. <i>Toxicology and Applied Pharmacology</i> , 2017, 327, 59-70.	2.8	8
36	RNA SEQ Analysis Indicates that the AE3 Cl ⁻ /HCO ₃ ⁻ Exchanger Contributes to Active Transport-Mediated CO ₂ Disposal in Heart. <i>Scientific Reports</i> , 2017, 7, 7264.	3.3	5

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37	Rhinovirus infection results in stronger and more persistent genomic dysregulation: Evidence for altered innate immune response in asthmatics at baseline, early in infection, and during convalescence. <i>PLoS ONE</i> , 2017, 12, e0178096.	2.5	21
38	GRcalculator: an online tool for calculating and mining dose-response data. <i>BMC Cancer</i> , 2017, 17, 698.	2.6	64
39	Comprehensive microRNA-sequencing of exosomes derived from head and neck carcinoma cells <i>in vitro</i> reveals common secretion profiles and potential utility as salivary biomarkers. <i>Oncotarget</i> , 2017, 8, 82459-82474.	1.8	80
40	Reprogramming of the Epigenome by MLL1 Links Early-Life Environmental Exposures to Prostate Cancer Risk. <i>Molecular Endocrinology</i> , 2016, 30, 856-871.	3.7	68
41	In utero exposure of rats to high-fat diets perturbs gene expression profiles and cancer susceptibility of prepubertal mammary glands. <i>Journal of Nutritional Biochemistry</i> , 2016, 29, 73-82.	4.2	26
42	Bisphenol A Disrupts HNF4 α -Regulated Gene Networks Linking to Prostate Preneoplasia and Immune Disruption in Noble Rats. <i>Endocrinology</i> , 2016, 157, 207-219.	2.8	22
43	The genetic fingerprint of susceptibility for transplant-associated thrombotic microangiopathy. <i>Blood</i> , 2016, 127, 989-996.	1.4	152
44	DNA methylome changes by estradiol benzoate and bisphenol A links early-life environmental exposures to prostate cancer risk. <i>Epigenetics</i> , 2016, 11, 674-689.	2.7	59
45	The Development of Spasmolytic Polypeptide/TFF2-Expressing Metaplasia (SPEM) During Gastric Repair Is Absent in the Aged Stomach. <i>Cellular and Molecular Gastroenterology and Hepatology</i> , 2016, 2, 605-624.	4.5	79
46	Model reduction and parameter estimation of nonlinear dynamical biochemical reaction networks. <i>IET Systems Biology</i> , 2016, 10, 10-16.	1.5	12
47	Loss of β kinase β promotes myofibroblast transformation and senescence through activation of the ROS-TGF β autocrine loop. <i>Protein and Cell</i> , 2016, 7, 338-350.	11.0	10
48	Ah Receptor Activation by Dioxin Disrupts Activin, BMP, and WNT Signals During the Early Differentiation of Mouse Embryonic Stem Cells and Inhibits Cardiomyocyte Functions. <i>Toxicological Sciences</i> , 2016, 149, 346-357.	3.1	54
49	IL-31-Driven Skin Remodeling Involves Epidermal Cell Proliferation and Thickening That Lead to Impaired Skin-Barrier Function. <i>PLoS ONE</i> , 2016, 11, e0161877.	2.5	59
50	Disruption of Ah Receptor Signaling during Mouse Development Leads to Abnormal Cardiac Structure and Function in the Adult. <i>PLoS ONE</i> , 2015, 10, e0142440.	2.5	42
51	Ah Receptor Signaling Controls the Expression of Cardiac Development and Homeostasis Genes. <i>Toxicological Sciences</i> , 2015, 147, 425-435.	3.1	38
52	Sex- and tissue-specific methylome changes in brains of mice perinatally exposed to lead. <i>NeuroToxicology</i> , 2015, 46, 92-100.	3.0	52
53	Identification of Secretoglobin <i>Scgb2a1</i> as a target for developmental reprogramming by BPA in the rat prostate. <i>Epigenetics</i> , 2015, 10, 127-134.	2.7	53
54	Exposure of Human Prostataspheres to Bisphenol A Epigenetically Regulates SNORD Family Noncoding RNAs via Histone Modification. <i>Endocrinology</i> , 2015, 156, 3984-3995.	2.8	45

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55	Fibrocytes Regulate Wilms Tumor 1â€™Positive Cell Accumulation in Severe Fibrotic Lung Disease. <i>Journal of Immunology</i> , 2015, 195, 3978-3991.	0.8	29
56	SERPINB3/B4 Contributes to Early Inflammation and Barrier Dysfunction in an Experimental Murine Model of Atopic Dermatitis. <i>Journal of Investigative Dermatology</i> , 2015, 135, 160-169.	0.7	61
57	Expression of Signaling Components in Embryonic Eyelid Epithelium. <i>PLoS ONE</i> , 2014, 9, e87038.	2.5	5
58	Formaldehyde-Assisted Isolation of Regulatory Elements (FAIRE) Analysis Uncovers Broad Changes in Chromatin Structure Resulting from Hexavalent Chromium Exposure. <i>PLoS ONE</i> , 2014, 9, e97849.	2.5	9
59	Stratified randomization controls better for batch effects in 450K methylation analysis: a cautionary tale. <i>Frontiers in Genetics</i> , 2014, 5, 354.	2.3	43
60	SPDEF Inhibits Prostate Carcinogenesis by Disrupting a Positive Feedback Loop in Regulation of the Foxm1 Oncogene. <i>PLoS Genetics</i> , 2014, 10, e1004656.	3.5	75
61	Targeting GPR30 with G-1: a new therapeutic target for castration-resistant prostate cancer. <i>Endocrine-Related Cancer</i> , 2014, 21, 903-914.	3.1	45
62	Long-term exposure to low-concentrations of Cr(VI) induce DNA damage and disrupt the transcriptional response to benzo[a]pyrene. <i>Toxicology</i> , 2014, 316, 14-24.	4.2	31
63	Overexpression of Dimethylarginine Dimethylaminohydrolase 1 Attenuates Airway Inflammation in a Mouse Model of Asthma. <i>PLoS ONE</i> , 2014, 9, e85148.	2.5	21
64	Dual Targeting of MEK and PI3K Pathways Attenuates Established and Progressive Pulmonary Fibrosis. <i>PLoS ONE</i> , 2014, 9, e86536.	2.5	24
65	Deciphering gene expression program of MAP3K1 in mouse eyelid morphogenesis. <i>Developmental Biology</i> , 2013, 374, 96-107.	2.0	13
66	Control of Nutrient Stress-Induced Metabolic Reprogramming by PKCÎ¶ in Tumorigenesis. <i>Cell</i> , 2013, 152, 599-611.	28.9	160
67	Genome-Wide Signatures of Transcription Factor Activity: Connecting Transcription Factors, Disease, and Small Molecules. <i>PLoS Computational Biology</i> , 2013, 9, e1003198.	3.2	30
68	Disruption of Aryl Hydrocarbon Receptor Homeostatic Levels during Embryonic Stem Cell Differentiation Alters Expression of Homeobox Transcription Factors that Control Cardiomyogenesis. <i>Environmental Health Perspectives</i> , 2013, 121, 1334-1343.	6.0	45
69	Expressomal approach for comprehensive analysis and visualization of ligand sensitivities of xenoestrogen responsive genes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 16508-16513.	7.1	29
70	c-Myc phosphorylation by PKCÎ¶ represses prostate tumorigenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 6418-6423.	7.1	49
71	Folliculin Contributes to VHL Tumor Suppressing Activity in Renal Cancer through Regulation of Autophagy. <i>PLoS ONE</i> , 2013, 8, e70030.	2.5	23
72	Integrative Assessment of Chlorine-Induced Acute Lung Injury in Mice. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2012, 47, 234-244.	2.9	28

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73	Rapid and Weight-Independent Improvement of Glucose Tolerance Induced by a Peptide Designed to Elicit Apoptosis in Adipose Tissue Endothelium. <i>Diabetes</i> , 2012, 61, 2299-2310.	0.6	20
74	Loss of the miR-144/451 cluster impairs ischaemic preconditioning-mediated cardioprotection by targeting Rac-1. <i>Cardiovascular Research</i> , 2012, 94, 379-390.	3.8	124
75	Identification of a NF- κ B cardioprotective gene program: NF- κ B regulation of Hsp70.1 contributes to cardioprotection after permanent coronary occlusion. <i>Journal of Molecular and Cellular Cardiology</i> , 2011, 51, 82-89.	1.9	32
76	WebGimm: An integrated web-based platform for cluster analysis, functional analysis, and interactive visualization of results. <i>Source Code for Biology and Medicine</i> , 2011, 6, 3.	1.7	6
77	Generalized random set framework for functional enrichment analysis using primary genomics datasets. <i>Bioinformatics</i> , 2011, 27, 70-77.	4.1	18
78	Haplotype Association Mapping of Acute Lung Injury in Mice Implicates Activin A Receptor, Type 1. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2011, 183, 1499-1509.	5.6	38
79	Identification of maternally regulated fetal gene networks in the placenta with a novel embryo transfer system in mice. <i>Physiological Genomics</i> , 2011, 43, 317-324.	2.3	3
80	Epigenetic Changes with Dietary Soy in Cynomolgus Monkeys. <i>PLoS ONE</i> , 2011, 6, e26791.	2.5	48
81	A semi-parametric Bayesian model for unsupervised differential co-expression analysis. <i>BMC Bioinformatics</i> , 2010, 11, 234.	2.6	35
82	Genomics Portals: integrative web-platform for mining genomics data. <i>BMC Genomics</i> , 2010, 11, 27.	2.8	13
83	Genomewide Association Analysis of Respiratory Syncytial Virus Infection in Mice. <i>Journal of Virology</i> , 2010, 84, 2257-2269.	3.4	15
84	MicroRNA-494 Targeting Both Proapoptotic and Antiapoptotic Proteins Protects Against Ischemia/Reperfusion-Induced Cardiac Injury. <i>Circulation</i> , 2010, 122, 1308-1318.	1.6	296
85	Research Resource: Estrogen-Driven Prolactin-Mediated Gene-Expression Networks in Hormone-Induced Prostatic Intraepithelial Neoplasia. <i>Molecular Endocrinology</i> , 2010, 24, 2207-2217.	3.7	14
86	NF- κ B driven cardioprotective gene programs; Hsp70.3 and cardioprotection after late ischemic preconditioning. <i>Journal of Molecular and Cellular Cardiology</i> , 2010, 49, 664-672.	1.9	41
87	Surfactant-Associated Protein B Is Critical to Survival in Nickel-Induced Injury in Mice. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2009, 41, 226-236.	2.9	20
88	Deletion Hotspots in AMACR Promoter CpG Island Are cis-Regulatory Elements Controlling the Gene Expression in the Colon. <i>PLoS Genetics</i> , 2009, 5, e1000334.	3.5	30
89	LRpath: a logistic regression approach for identifying enriched biological groups in gene expression data. <i>Bioinformatics</i> , 2009, 25, 211-217.	4.1	163
90	CLEAN: CLustering Enrichment ANalysis. <i>BMC Bioinformatics</i> , 2009, 10, 234.	2.6	65

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91	Gene position within chromosome territories correlates with their involvement in distinct rearrangement types in thyroid cancer cells. <i>Genes Chromosomes and Cancer</i> , 2009, 48, 222-228.	2.8	28
92	Gene expression profiling of blood to predict the onset of leukemia. <i>Blood Cells, Molecules, and Diseases</i> , 2009, 42, 64-70.	1.4	2
93	Genomewide Analysis of Aryl Hydrocarbon Receptor Binding Targets Reveals an Extensive Array of Gene Clusters that Control Morphogenetic and Developmental Programs. <i>Environmental Health Perspectives</i> , 2009, 117, 1139-1146.	6.0	90
94	Gene Expression Profiling Identifies Lobe-Specific and Common Disruptions of Multiple Gene Networks in Testosterone-Supported, 17 β -Estradiol- or Diethylstilbestrol-Induced Prostate Dysplasia in Noble Rats. <i>Neoplasia</i> , 2008, 10, 20-IN18.	5.3	13
95	Nonredundant Functions of $\hat{1}\hat{2}$ and $\hat{3}\hat{1}$ T Cells in Acrolein-Induced Pulmonary Pathology. <i>Toxicological Sciences</i> , 2008, 105, 188-199.	3.1	11
96	Transcriptome Analyses in Normal Prostate Epithelial Cells Exposed to Low-Dose Cadmium: Oncogenic and Immunomodulations Involving the Action of Tumor Necrosis Factor. <i>Environmental Health Perspectives</i> , 2008, 116, 769-776.	6.0	48
97	Candidate genes controlling pulmonary function in mice: transcript profiling and predicted protein structure. <i>Physiological Genomics</i> , 2007, 31, 410-421.	2.3	45
98	Regeneration of the adult thymus is preceded by the expansion of K5+K8+ epithelial cell progenitors and by increased expression of Trp63, cMyc and Tcf3 transcription factors in the thymic stroma. <i>International Immunology</i> , 2007, 19, 1249-1260.	4.0	38
99	Genomic Profile of Matrix and Vasculature Remodeling in TGF- $\hat{1}$ -Induced Pulmonary Fibrosis. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2007, 37, 309-321.	2.9	60
100	A new method to remove hybridization bias for interspecies comparison of global gene expression profiles uncovers an association between mRNA sequence divergence and differential gene expression in <i>Xenopus</i> . <i>Nucleic Acids Research</i> , 2006, 34, 185-200.	14.5	37
101	The Role of Metallothionein in the Pathogenesis of Acute Lung Injury. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2006, 34, 73-82.	2.9	46
102	Conditional Activation of RET/PTC3 and BRAFV600E in Thyroid Cells Is Associated with Gene Expression Profiles that Predict a Preferential Role of BRAF in Extracellular Matrix Remodeling. <i>Cancer Research</i> , 2006, 66, 6521-6529.	0.9	129
103	Context-specific infinite mixtures for clustering gene expression profiles across diverse microarray dataset. <i>Bioinformatics</i> , 2006, 22, 1737-1744.	4.1	29
104	Gene expression and discovery during lens regeneration in mouse: regulation of epithelial to mesenchymal transition and lens differentiation. <i>Molecular Vision</i> , 2006, 12, 422-40.	1.1	32
105	Modeling variation in tumors in vivo. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 2408-2413.	7.1	10
106	Gene Expression Changes during the Development of Acute Lung Injury Role of Transforming Growth Factor $\hat{1}$. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2005, 172, 1399-1411.	5.6	71
107	Microarray results improve significantly as hybridization approaches equilibrium. <i>BioTechniques</i> , 2004, 36, 790-796.	1.8	76
108	Neural system-enriched gene expression: relationship to biological pathways and neurological diseases. <i>Physiological Genomics</i> , 2004, 18, 167-183.	2.3	15

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109	Expression of genes in the TGF- β 2 signaling pathway is significantly deregulated in smooth muscle cells from aorta of aryl hydrocarbon receptor knockout mice. <i>Toxicology and Applied Pharmacology</i> , 2004, 194, 79-89.	2.8	93
110	Different Global Gene Expression Profiles in Benzo[<i>a</i>]Pyrene- and Dioxin-Treated Vascular Smooth Muscle Cells of AHR-Knockout and Wild-Type Mice. <i>Cardiovascular Toxicology</i> , 2004, 4, 47-74.	2.7	49
111	Gene Expression Profiles of Mouse Aorta and Cultured Vascular Smooth Muscle Cells Differ Widely, Yet Show Common Responses to Dioxin Exposure. <i>Cardiovascular Toxicology</i> , 2004, 4, 385-404.	2.7	21
112	Critical regulation of genes for tumor cell migration by AP-1. <i>Clinical and Experimental Metastasis</i> , 2004, 21, 293-304.	3.3	42
113	4-Aminobiphenyl-Induced Liver and Urinary Bladder DNA Adduct Formation in Cyp1a2(-/-) and Cyp1a2(+/-) Mice. <i>Journal of the National Cancer Institute</i> , 2003, 95, 1227-1237.	6.3	61
114	Isozyme-Specific Abnormalities of PKC in Thyroid Cancer: Evidence for Post-Transcriptional Changes in PKC Epsilon. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2002, 87, 2150-2159.	3.6	37
115	Bayesian infinite mixture model based clustering of gene expression profiles. <i>Bioinformatics</i> , 2002, 18, 1194-1206.	4.1	248
116	Expression of ATM in ataxia telangiectasia fibroblasts rescues defects in DNA double-strand break repair in nuclear extracts. <i>Environmental and Molecular Mutagenesis</i> , 2001, 37, 128-140.	2.2	19
117	XPA protein alters the specificity of ultraviolet light-induced mutagenesis in vitro. <i>Environmental and Molecular Mutagenesis</i> , 2001, 37, 329-339.	2.2	5
118	Clustering mutational spectra via classification likelihood and markov chain monte carlo algorithms. <i>Journal of Agricultural, Biological, and Environmental Statistics</i> , 2001, 6, 19-37.	1.4	5
119	The transcriptional signature of dioxin in human hepatoma HepG2 cells. <i>Biochemical Pharmacology</i> , 2000, 60, 1129-1142.	4.4	212
120	Proximity of Chromosomal Loci That Participate in Radiation-Induced Rearrangements in Human Cells. <i>Science</i> , 2000, 290, 138-141.	12.6	450
121	What Is the Potential Measurement Error in Occupational Exposure Studies?. <i>Journal of the Air and Waste Management Association</i> , 2000, 50, 941-947.	1.9	3
122	Mutational specificity in a shuttle vector replicating in chromium(VI)-treated mammalian cells. , 1999, 33, 313-319.		26
123	Longitudinal Estimates of Pulmonary Function in Refractory Ceramic Fiber Manufacturing Workers. <i>American Journal of Respiratory and Critical Care Medicine</i> , 1998, 157, 1226-1233.	5.6	23