

# Douglas A Bell

## List of Publications by Year in descending order

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163  
papers

13,411  
citations

20817

60  
h-index

22832

112  
g-index

167  
all docs

167  
docs citations

167  
times ranked

14580  
citing authors

#	ARTICLE	IF	CITATIONS
1	Circulating MicroRNAs, Polychlorinated Biphenyls, and Environmental Liver Disease in the Anniston Community Health Survey. <i>Environmental Health Perspectives</i> , 2022, 130, 17003.	6.0	12
2	Epigenome-wide association study of bronchopulmonary dysplasia (BPD) in preterm infants: Results from the Discovery-BPD program. <i>FASEB Journal</i> , 2022, 36, .	0.5	0
3	Epigenome-wide association study of bronchopulmonary dysplasia in preterm infants: results from the discovery-BPD program. <i>Clinical Epigenetics</i> , 2022, 14, 57.	4.1	12
4	Germline and Somatic Genetic Variants in the p53 Pathway Interact to Affect Cancer Risk, Progression, and Drug Response. <i>Cancer Research</i> , 2021, 81, 1667-1680.	0.9	32
5	Polychlorinated biphenyl exposure and DNA methylation in the Anniston Community Health Survey. <i>Epigenetics</i> , 2020, 15, 337-357.	2.7	10
6	Mining a human transcriptome database for chemical modulators of NRF2. <i>PLoS ONE</i> , 2020, 15, e0239367.	2.5	19
7	Single-Cell Analyses Identify Dysfunctional CD16+ CD8 <sup>+</sup> Cells in Smokers. <i>Cell Reports Medicine</i> , 2020, 1, 100054.	6.5	21
8	Analysis of genome-wide methylation using reduced representation bisulfite sequencing (RRBS) technology. , 2020, , 141-156.		1
9	Dioxin-like compound exposures and DNA methylation in the Anniston Community Health Survey Phase II. <i>Science of the Total Environment</i> , 2020, 742, 140424.	8.0	6
10	Mining a human transcriptome database for chemical modulators of NRF2. , 2020, 15, e0239367.		0
11	Mining a human transcriptome database for chemical modulators of NRF2. , 2020, 15, e0239367.		0
12	Mining a human transcriptome database for chemical modulators of NRF2. , 2020, 15, e0239367.		0
13	Mining a human transcriptome database for chemical modulators of NRF2. , 2020, 15, e0239367.		0
14	Mining a human transcriptome database for chemical modulators of NRF2. , 2020, 15, e0239367.		0
15	Mining a human transcriptome database for chemical modulators of NRF2. , 2020, 15, e0239367.		0
16	The discovery BPD (D-BPD) program: study protocol of a prospective translational multicenter collaborative study to investigate determinants of chronic lung disease in very low birth weight infants. <i>BMC Pediatrics</i> , 2019, 19, 227.	1.7	5
17	Microbiota-derived acetate protects against respiratory syncytial virus infection through a GPR43-type 1 interferon response. <i>Nature Communications</i> , 2019, 10, 3273.	12.8	234
18	Epigenome-wide meta-analysis of DNA methylation and childhood asthma. <i>Journal of Allergy and Clinical Immunology</i> , 2019, 143, 2062-2074.	2.9	147

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19	Smoking-associated AHRR demethylation in cord blood DNA: impact of CD235a+ nucleated red blood cells. <i>Clinical Epigenetics</i> , 2019, 11, 87.	4.1	18
20	Associations between Maternal Tobacco Smoke Exposure and the Cord Blood CD4+ DNA Methylome. <i>Environmental Health Perspectives</i> , 2019, 127, 47009.	6.0	13
21	Crohn's disease <i>IRGM</i> risk alleles are associated with altered gene expression in human tissues. <i>American Journal of Physiology - Renal Physiology</i> , 2019, 316, G95-G105.	3.4	17
22	Sulforaphane enriched transcriptome of lung mitochondrial energy metabolism and provided pulmonary injury protection via Nrf2 in mice. <i>Toxicology and Applied Pharmacology</i> , 2019, 364, 29-44.	2.8	35
23	A hypermorphic antioxidant response element is associated with increased MS4A6A expression and Alzheimer's disease. <i>Redox Biology</i> , 2018, 14, 686-693.	9.0	21
24	Identification of Smoking-Associated Differentially Methylated Regions Using Reduced Representation Bisulfite Sequencing and Cell type-Specific Enhancer Activation and Gene Expression. <i>Environmental Health Perspectives</i> , 2018, 126, 047015.	6.0	26
25	A distinct class of antioxidant response elements is consistently activated in tumors with NRF2 mutations. <i>Redox Biology</i> , 2018, 19, 235-249.	9.0	37
26	Activation of Nrf2 in the liver is associated with stress resistance mediated by suppression of the growth hormone-regulated STAT5b transcription factor. <i>PLoS ONE</i> , 2018, 13, e0200004.	2.5	36
27	Blood monocyte transcriptome and epigenome analyses reveal loci associated with human atherosclerosis. <i>Nature Communications</i> , 2017, 8, 393.	12.8	51
28	Tobacco exposure-related alterations in DNA methylation and gene expression in human monocytes: the Multi-Ethnic Study of Atherosclerosis (MESA). <i>Epigenetics</i> , 2017, 12, 1092-1100.	2.7	29
29	Distinct Epigenetic Effects of Tobacco Smoking in Whole Blood and among Leukocyte Subtypes. <i>PLoS ONE</i> , 2016, 11, e0166486.	2.5	113
30	Potential therapeutic targets in Nrf2-dependent protection against neonatal respiratory distress disease predicted by cDNA microarray analysis and bioinformatics tools. <i>Current Opinion in Toxicology</i> , 2016, 1, 125-133.	5.0	9
31	An African-specific polymorphism in the <i>TP53</i> gene impairs p53 tumor suppressor function in a mouse model. <i>Genes and Development</i> , 2016, 30, 918-930.	5.9	277
32	A Polymorphic Antioxidant Response Element Links NRF2/sMAF Binding to Enhanced MAPT Expression and Reduced Risk of Parkinsonian Disorders. <i>Cell Reports</i> , 2016, 15, 830-842.	6.4	40
33	Determinants of host susceptibility to murine respiratory syncytial virus (RSV) disease identify a role for the innate immunity scavenger receptor MARCO gene in human infants. <i>EBioMedicine</i> , 2016, 11, 73-84.	6.1	24
34	The importance of p53 pathway genetics in inherited and somatic cancer genomes. <i>Nature Reviews Cancer</i> , 2016, 16, 251-265.	28.4	131
35	Global Analysis of Methylation Profiles From High Resolution CpG Data. <i>Genetic Epidemiology</i> , 2015, 39, 53-64.	1.3	19
36	Beyond antioxidant genes in the ancient Nrf2 regulatory network. <i>Free Radical Biology and Medicine</i> , 2015, 88, 452-465.	2.9	74

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37	Interactions of Chromatin Context, Binding Site Sequence Content, and Sequence Evolution in Stress-Induced p53 Occupancy and Transactivation. <i>PLoS Genetics</i> , 2015, 11, e1004885.	3.5	50
38	DNA Methylation of the Aryl Hydrocarbon Receptor Repressor Associations With Cigarette Smoking and Subclinical Atherosclerosis. <i>Circulation: Cardiovascular Genetics</i> , 2015, 8, 707-716.	5.1	107
39	Linking polymorphic p53 response elements with gene expression in airway epithelial cells of smokers and cancer risk. <i>Human Genetics</i> , 2014, 133, 1467-1476.	3.8	3
40	A genetic model of differential susceptibility to human respiratory syncytial virus (RSV) infection. <i>FASEB Journal</i> , 2014, 28, 1947-1956.	0.5	24
41	Maternal Smoking and DNA Methylation in Newborns: <i>In Utero</i> Effect or Epigenetic Inheritance?. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1007-1017.	2.5	108
42	A Polymorphic p53 Response Element in KIT Ligand Influences Cancer Risk and Has Undergone Natural Selection. <i>Cell</i> , 2013, 155, 410-422.	28.9	115
43	Novel Hematopoietic Target Genes in the NRF2-Mediated Transcriptional Pathway. <i>Oxidative Medicine and Cellular Longevity</i> , 2013, 2013, 1-12.	4.0	75
44	Abstract 3647: Dose-dependent alteration of CpG methylation in AHRR and GFI1 in mononuclear cell DNA of smokers. , 2013, , .		1
45	CSF1 Is a Novel p53 Target Gene Whose Protein Product Functions in a Feed-Forward Manner to Suppress Apoptosis and Enhance p53-Mediated Growth Arrest. <i>PLoS ONE</i> , 2013, 8, e74297.	2.5	20
46	450K Epigenome-Wide Scan Identifies Differential DNA Methylation in Newborns Related to Maternal Smoking during Pregnancy. <i>Environmental Health Perspectives</i> , 2012, 120, 1425-1431.	6.0	654
47	Targeted Deletion of <i>Nrf2</i> Impairs Lung Development and Oxidant Injury in Neonatal Mice. <i>Antioxidants and Redox Signaling</i> , 2012, 17, 1066-1082.	5.4	92
48	Identification of novel NRF2-regulated genes by ChIP-Seq: influence on retinoid X receptor alpha. <i>Nucleic Acids Research</i> , 2012, 40, 7416-7429.	14.5	459
49	Formation of stress-specific p53 binding patterns is influenced by chromatin but not by modulation of p53 binding affinity to response elements. <i>Nucleic Acids Research</i> , 2011, 39, 3053-3063.	14.5	13
50	Human single-nucleotide polymorphisms alter p53 sequence-specific binding at gene regulatory elements. <i>Nucleic Acids Research</i> , 2011, 39, 178-189.	14.5	28
51	Nrf2-regulated PPAR $\gamma$ Expression Is Critical to Protection against Acute Lung Injury in Mice. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2010, 182, 170-182.	5.6	184
52	Abstract B51: Discovery of novel genomic targets in the NRF2-mediated antioxidant response pathway by ChIP-seq and ChIP-chip. , 2010, , .		2
53	Genetic Variation and Antioxidant Response Gene Expression in the Bronchial Airway Epithelium of Smokers at Risk for Lung Cancer. <i>PLoS ONE</i> , 2010, 5, e11934.	2.5	55
54	Abstract 1099: In vitro and in cellulo measurement of p53-binding activities reveals involvement of chromatin in p53-binding pattern formation. , 2010, , .		0

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55	Probing the Functional Impact of Sequence Variation on p53-DNA Interactions Using a Novel Microsphere Assay for Protein-DNA Binding with Human Cell Extracts. <i>PLoS Genetics</i> , 2009, 5, e1000462.	3.5	39
56	Discovery and verification of functional single nucleotide polymorphisms in regulatory genomic regions: Current and developing technologies. <i>Mutation Research - Reviews in Mutation Research</i> , 2008, 659, 147-157.	5.5	142
57	Noncanonical DNA Motifs as Transactivation Targets by Wild Type and Mutant p53. <i>PLoS Genetics</i> , 2008, 4, e1000104.	3.5	91
58	Genetic determinants in the metabolism of bladder carcinogens in relation to risk of bladder cancer. <i>Carcinogenesis</i> , 2008, 29, 1386-1393.	2.8	52
59	Divergent Evolution of Human p53 Binding Sites: Cell Cycle Versus Apoptosis. <i>PLoS Genetics</i> , 2007, 3, e127.	3.5	88
60	Identification of polymorphic antioxidant response elements in the human genome. <i>Human Molecular Genetics</i> , 2007, 16, 1188-1200.	2.9	147
61	Identification of polymorphic antioxidant response elements in the human genome. <i>Human Molecular Genetics</i> , 2007, 16, 2780-2780.	2.9	2
62	Identification and functional characterization of polymorphisms in human cyclooxygenase-1 (PTGS1). <i>Pharmacogenetics and Genomics</i> , 2007, 17, 145-160.	1.5	52
63	Reply to the letter to the Editor: "N-Acetyltransferases and the susceptibility to benzidine-induced bladder carcinogenesis". <i>International Journal of Cancer</i> , 2007, 121, 1637-1639.	5.1	3
64	Alcohol Dehydrogenase Genetic Polymorphisms, Low-to-Moderate Alcohol Consumption, and Risk of Breast Cancer. <i>Alcoholism: Clinical and Experimental Research</i> , 2007, 31, 467-476.	2.4	38
65	Variation in fiberoptic bead-based oligonucleotide microarrays: dispersion characteristics among hybridization and biological replicate samples. <i>Biology Direct</i> , 2006, 1, 18.	4.6	22
66	Generalization of DNA microarray dispersion properties: microarray equivalent of t-distribution. <i>Biology Direct</i> , 2006, 1, 27.	4.6	15
67	Polymorphisms of the DNA repair genes XPD (Lys751Gln) and XRCC1 (Arg399Gln and Arg194Trp): relationship to breast cancer risk and familial predisposition to breast cancer. <i>Breast Cancer Research and Treatment</i> , 2006, 95, 73-80.	2.5	44
68	NAT2 slow acetylation and bladder cancer in workers exposed to benzidine. <i>International Journal of Cancer</i> , 2006, 118, 161-168.	5.1	62
69	Variation in genes relevant to aromatic hydrocarbon metabolism and the risk of adult brain tumors. <i>Neuro-Oncology</i> , 2006, 8, 145-155.	1.2	34
70	Lack of Associations among Cancer and Albumin Adducts, ras p21 Oncoprotein Levels, and CYP1A1, CYP2D6, NAT1, and NAT2 in a Nested Case-Control Study of Lung Cancer within the Physicians' Health Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 1417-1419.	2.5	16
71	CYP1A1 and CYP1B1 genotypes, haplotypes, and TCDD-induced gene expression in subjects from Seveso, Italy. <i>Toxicology</i> , 2005, 207, 191-202.	4.2	61
72	Single nucleotide polymorphism in transcriptional regulatory regions and expression of environmentally responsive genes. <i>Toxicology and Applied Pharmacology</i> , 2005, 207, 84-90.	2.8	100

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73	Glutathione S-transferase polymorphisms and survival from head and neck cancer. <i>Head and Neck</i> , 2005, 27, 232-242.	2.0	32
74	Functionally distinct polymorphic sequences in the human genome that are targets for p53 transactivation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 6431-6436.	7.1	80
75	Recombinant CYP3A4*17 Is Defective in Metabolizing the Hypertensive Drug Nifedipine, and the CYP3A4*17 Allele May Occur on the Same Chromosome as CYP3A5*3, Representing a New Putative Defective CYP3A Haplotype. <i>Journal of Pharmacology and Experimental Therapeutics</i> , 2005, 313, 302-309.	2.5	65
76	Functional Diversity in the Gene Network Controlled by the Master Regulator p53 in Humans. <i>Cell Cycle</i> , 2005, 4, 1026-1029.	2.6	28
77	Expression-based discovery of variation in the human glutathione S-transferase M3 promoter and functional analysis in a glioma cell line using allele-specific chromatin immunoprecipitation. <i>Cancer Research</i> , 2005, 65, 99-104.	0.9	26
78	Pooled Analysis of Alcohol Dehydrogenase Genotypes and Head and Neck Cancer: A HuGE Review. <i>American Journal of Epidemiology</i> , 2004, 159, 1-16.	3.4	198
79	N-acetyltransferase 2 (NAT2) genotypes, cigarette smoking, and the risk of breast cancer. <i>Cancer Detection and Prevention</i> , 2004, 28, 187-193.	2.1	30
80	Urinary mutagenesis and fried red meat intake: Influence of cooking temperature, phenotype, and genotype of metabolizing enzymes in a controlled feeding study. <i>Environmental and Molecular Mutagenesis</i> , 2004, 43, 53-74.	2.2	38
81	Carotenoids/vitamin C and smoking-related bladder cancer. <i>International Journal of Cancer</i> , 2004, 110, 417-423.	5.1	74
82	Polychlorinated biphenyls, cytochrome P450 1A1 (CYP1A1) polymorphisms, and breast cancer risk among African American women and white women in North Carolina: a population-based case-control study. <i>Breast Cancer Research</i> , 2004, 7, R12-8.	5.0	69
83	Cigarette smoking, cytochrome P4501A1 polymorphisms, and breast cancer among African-American and white women. <i>Breast Cancer Research</i> , 2004, 6, R460-73.	5.0	69
84	Bilirubin UDP-glucuronosyltransferase 1A1 (UGT1A1) gene promoter polymorphisms and HPRT, glycoporphin A, and micronuclei mutant frequencies in human blood. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2004, 560, 1-10.	1.7	9
85	Sequence Context at Human Single Nucleotide Polymorphisms: Overrepresentation of CpG Dinucleotide at Polymorphic Sites and Suppression of Variation in CpG Islands. <i>Journal of Molecular Biology</i> , 2003, 327, 303-308.	4.2	54
86	Glutathione-S-transferase genotypes, smoking, and their association with markers of inflammation, hemostasis, and endothelial function: the atherosclerosis risk in communities (ARIC) study. <i>Atherosclerosis</i> , 2003, 171, 265-272.	0.8	60
87	Permanent hair dyes and bladder cancer: risk modification by cytochrome P4501A2 and N-acetyltransferases 1 and 2. <i>Carcinogenesis</i> , 2003, 24, 483-489.	2.8	111
88	Single amino acid mutations, but not common polymorphisms, decrease the activity of CYP1B1 against (-)benzo[a]pyrene-7R-trans-7,8-dihydrodiol. <i>Carcinogenesis</i> , 2003, 24, 1247-1255.	2.8	34
89	Diet, GSTM1 and GSTT1 and head and neck cancer. <i>Carcinogenesis</i> , 2003, 25, 735-740.	2.8	29
90	Title is missing!. <i>Epidemiology</i> , 2003, 14, 321-327.	2.7	14

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91	Risk of Atherosclerosis: Interaction of Smoking and Glutathione S-Transferase Genes. <i>Epidemiology</i> , 2003, 14, 321-327.	2.7	48
92	Genetic polymorphisms in GSTM1, -P1, -T1, and CYP2E1 and the risk of adult brain tumors. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2003, 12, 14-22.	2.5	40
93	Pooled Analysis and Meta-analysis of Glutathione S-Transferase M1 and Bladder Cancer: A HuGE Review. <i>American Journal of Epidemiology</i> , 2002, 156, 95-109.	3.4	209
94	Associations between carcinogen-DNA damage, glutathione S-transferase genotypes, and risk of lung cancer in the prospective Physicians' Health Cohort Study. <i>Carcinogenesis</i> , 2002, 23, 1641-1646.	2.8	97
95	XRCC1 polymorphisms and head and neck cancer. <i>Cancer Letters</i> , 2002, 178, 181-186.	7.2	96
96	Symposium Overview: Genetic Polymorphisms in DNA Repair and Cancer Risk. <i>Toxicology and Applied Pharmacology</i> , 2002, 185, 64-73.	2.8	73
97	Genetic variability in susceptibility and response to toxicants. <i>Toxicology Letters</i> , 2001, 120, 269-280.	0.8	148
98	Risk of head and neck cancer and the alcohol dehydrogenase 3 genotype. <i>Carcinogenesis</i> , 2001, 22, 57-61.	2.8	60
99	Inherited polymorphism in the N-acetyltransferase 1(NAT1) and 2(NAT2) genes and susceptibility to gastric and colorectal adenocarcinoma. <i>International Journal of Cancer</i> , 2000, 85, 46-49.	5.1	59
100	A pilot study investigating the role of NAT1 and NAT2 polymorphisms in gastric adenocarcinoma. <i>International Journal of Cancer</i> , 2000, 87, 507-511.	5.1	32
101	Bilirubin UDP-glucuronosyltransferase1A1 gene polymorphisms: Susceptibility to oxidative damage and cancer?. <i>Molecular Carcinogenesis</i> , 2000, 29, 198-204.	2.7	31
102	Preliminary evidence of an association of tumour necrosis factor microsatellites with increased risk of multiple basal cell carcinomas. <i>British Journal of Dermatology</i> , 2000, 142, 441-445.	1.5	43
103	Re: Hemminki,K., Dickey,C., Karlsson,S., Bell,D., Hsu,Y., Tsai,W.-Y., Mooney,L.A., Savela,K. and Perera,F.P. (1997) Aromatic DNA adducts in foundry workers in relation to exposure, lifestyle and CYP1A1 and glutathione transferase M1 genotype. <i>Carcinogenesis</i> , 18, 345-350. <i>Carcinogenesis</i> , 2000, 21, 849-849.	2.8	3
104	Localization, sequence analysis, and ethnic distribution of a 96-bp insertion in the promoter of the human CYP2E1 gene. <i>Mutation Research - Mutation Research Genomics</i> , 2000, 432, 1-5.	1.1	23
105	XPD polymorphisms: effects on DNA repair proficiency. <i>Carcinogenesis</i> , 2000, 21, 551-555.	2.8	407
106	Erratum to "Glutathione S-transferase genotype as a susceptibility factor in smoking-related coronary heart disease". <i>Atherosclerosis</i> , 2000, 150, 447-449.	0.8	0
107	Glutathione S-transferase genotype as a susceptibility factor in smoking-related coronary heart disease. <i>Atherosclerosis</i> , 2000, 149, 451-462.	0.8	114
108	A pilot study investigating the role of NAT1 and NAT2 polymorphisms in gastric adenocarcinoma. <i>International Journal of Cancer</i> , 2000, 87, 507-511.	5.1	0



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109	Prostate cancer risk and polymorphism in 17 hydroxylase (CYP17) and steroid reductase (SRD5A2). <i>Carcinogenesis</i> , 1999, 20, 1727-1731.	2.8	175
110	Effects of Glutathione Transferase Theta Polymorphism on the Risk Estimates of Dichloromethane to Humans. <i>Toxicology and Applied Pharmacology</i> , 1999, 158, 221-230.	2.8	64
111	Arylamine N-acetyltransferase 1 (NAT1) and 2 (NAT2) genes and risk of urothelial transitional cell carcinoma among Japanese. <i>Pharmacogenetics and Genomics</i> , 1999, 9, 401-404.	5.7	63
112	Relationship between Ambient Air Pollution and DNA Damage in Polish Mothers and Newborns. <i>Environmental Health Perspectives</i> , 1998, 106, 821.	6.0	15
113	Association Between Glutathione S-Transferase M1, P1, and T1 Genetic Polymorphisms and Development of Breast Cancer. <i>Journal of the National Cancer Institute</i> , 1998, 90, 512-518.	6.3	245
114	Genetic susceptibility: significance in risk assessment. <i>Toxicology Letters</i> , 1998, 102-103, 185-189.	0.8	9
115	Human glutathione S-transferase P1 polymorphisms: relationship to lung tissue enzyme activity and population frequency distribution. <i>Carcinogenesis</i> , 1998, 19, 275-280.	2.8	556
116	A pilot study testing the association between N-acetyltransferases 1 and 2 and risk of oral squamous cell carcinoma in Japanese people. <i>Carcinogenesis</i> , 1998, 19, 1803-1807.	2.8	61
117	Catechol-O-methyltransferase and breast cancer risk. <i>Carcinogenesis</i> , 1998, 19, 1943-1947.	2.8	85
118	Polycyclic aromatic hydrocarbon-DNA adducts in human placenta and modulation by CYP1A1 induction and genotype. <i>Carcinogenesis</i> , 1998, 19, 1389-1392.	2.8	92
119	Identification and characterization of variant alleles of human acetyltransferase NAT1 with defective function using p-aminosalicylate as an in-vivo and in-vitro probe. <i>Pharmacogenetics and Genomics</i> , 1998, 8, 55-66.	5.7	122
120	Dichloromethane Metabolism to Formaldehyde and Reaction of Formaldehyde with Nucleic Acids in Hepatocytes of Rodents and Humans with and without Glutathione S-Transferase T1 and M1 Genes. <i>Toxicological Sciences</i> , 1997, 37, 168-180.	3.1	10
121	Genetic Polymorphisms in Human Drug Metabolic Enzymes. <i>Toxicological Sciences</i> , 1997, 40, 1-14.	3.1	1
122	Contribution of genetic and nutritional factors to DNA damage in heavy smokers. <i>Carcinogenesis</i> , 1997, 18, 503-509.	2.8	123
123	Frequencies of the defective CYP2C19 alleles responsible for the mephenytoin poor metabolizer phenotype in various Oriental, Caucasian, Saudi Arabian and American black populations. <i>Pharmacogenetics and Genomics</i> , 1997, 7, 59-64.	5.7	314
124	Genetic Analysis of Complex Diseases. <i>Science</i> , 1997, 275, 1327-1330.	12.6	55
125	Dichloromethane Metabolism to Formaldehyde and Reaction of Formaldehyde with Nucleic Acids in Hepatocytes of Rodents and Humans with and without Glutathione S-Transferase T1 and M1 Genes. <i>Fundamental and Applied Toxicology</i> , 1997, 37, 168-180.	1.8	32
126	Genetic Polymorphisms in Human Drug Metabolic Enzymes., <i>Fundamental and Applied Toxicology</i> , 1997, 40, 1-14.	1.8	48



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127	Pilot study of free and conjugated urinary mutagenicity during consumption of pan-fried meats: possible modulation by cruciferous vegetables, glutathione S-transferase-M1, and N-acetyltransferase-2. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 1997, 381, 83-96.	1.0	27
128	Increased risk for myelodysplastic syndromes in individuals with glutathione transferase theta 1 (GSTT1) gene defect. <i>Lancet, The</i> , 1996, 347, 295-297.	13.7	272
129	Increased oxidative DNA damage in livers of 2,3,7,8-tetrachlorodibenzo-p-dioxin treated intact but not ovariectomized rats. <i>Cancer Letters</i> , 1996, 98, 219-225.	7.2	78
130	SHORT COMMUNICATION: Glutathione S-transferase GSTT1 genotypes and susceptibility to cancer: studies of interactions with GSTM1 in lung, oral, gastric and colorectal cancers. <i>Carcinogenesis</i> , 1996, 17, 881-884.	2.8	277
131	The impact of interindividual variation in NAT2 activity on benzidine urinary metabolites and urothelial DNA adducts in exposed workers.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1996, 93, 5084-5089.	7.1	86
132	The role of the CYP2C9-Leu 359 allelic variant in the tolbutamide polymorphism. <i>Pharmacogenetics and Genomics</i> , 1996, 6, 341-349.	5.7	600
133	Xenobiotic Metabolism Genes and the Risk of Recurrent Spontaneous Abortion. <i>Epidemiology</i> , 1996, 7, 206-208.	2.7	20
134	Mutation spectra of chemical fractions of a complex mixture: role of nitroarenes in the mutagenic specificity of municipal waste incinerator emissions. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 1996, 349, 1-20.	1.0	42
135	Microsomal epoxide hydrolase polymorphism as a risk factor for ovarian cancer. , 1996, 17, 160-162.		99
136	Glutathione S-transferase M1 (GSTM1) and T1 (GSTT1) genetic polymorphism and susceptibility to gastric and colorectal adenocarcinoma. <i>Carcinogenesis</i> , 1996, 17, 1855-1859.	2.8	231
137	Isolation and characterization of a novel gene induced by 2, 3, 7, 8-tetrachlorodibenzo-p-dioxin in rat liver. <i>Carcinogenesis</i> , 1996, 17, 2609-2615.	2.8	115
138	Microsomal epoxide hydrolase polymorphism as a risk factor for ovarian cancer. <i>Molecular Carcinogenesis</i> , 1996, 17, 160-162.	2.7	1
139	Nomenclature for N-acetyltransferases. <i>Pharmacogenetics and Genomics</i> , 1995, 5, 1-17.	5.7	369
140	Extended-term cultures of human T-lymphocytes: a practical alternative to primary human lymphocytes for use in genotoxicity testing. <i>Mutagenesis</i> , 1995, 10, 189-201.	2.6	38
141	Occurrence of bcl-2 Oncogene Translocation With Increased Frequency in the Peripheral Blood of Heavy Smokers. <i>Journal of the National Cancer Institute</i> , 1995, 87, 223-224.	6.3	105
142	N-Acetylbenzidine and N,Nâ€²-diacetylbenzidine formation by rat and human liver slices exposed to benzidine. <i>Carcinogenesis</i> , 1995, 16, 1565-1571.	2.8	28
143	Polycyclic Aromatic Hydrocarbon-DNA Adducts in Smokers and Their Relationship to Micronutrient Levels and Glutathione-S-Transferase M1 Genotype. , 1995, , 191-209.		0
144	Polycyclic aromatic hydrocarbonâ€™DNA adducts in smokers and their relationship to micronutrient levels and the glutathione-S-transferase M1 genotype. <i>Carcinogenesis</i> , 1994, 15, 2449-2454.	2.8	84

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145	mutation spectra in salmonella of complex mixtures: Comparison of urban air to benzo[a]pyrene. Environmental and Molecular Mutagenesis, 1994, 24, 262-275.	2.2	52
146	Ethnic variation in the CYP2E1 gene: polymorphism analysis of 695 African-Americans, European-Americans and Taiwanese. Pharmacogenetics and Genomics, 1994, 4, 185-192.	5.7	158
147	L-myc Proto-oncogene alleles and susceptibility to hepatocellular carcinoma. International Journal of Cancer, 1993, 54, 927-930.	5.1	30
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