

Xifeng Wu

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/83402/xifeng-wu-publications-by-year.pdf>
Version: 2024-04-09

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.
The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

398 papers	20,262 citations	72 h-index	126 g-index
410 ext. papers	23,365 ext. citations	7.3 avg, IF	6.18 L-index

#	Paper	IF	Citations
398	A whole-exome case-control association study to characterize the contribution of rare coding variation to pancreatic cancer risk.. <i>Human Genetics and Genomics Advances</i> , 2022 , 3, 100078	0.8	
397	"Sugar-Sweetened Beverages" Is an Independent Risk From Pancreatic Cancer: Based on Half a Million Asian Cohort Followed for 25 Years.. <i>Frontiers in Oncology</i> , 2022 , 12, 835901	5.3	0
396	Predictive Utility of Mortality by Aging Measures at Different Hierarchical Levels and the Response to Modifiable Life Style Factors: Implications for Geroprotective Programs.. <i>Frontiers in Medicine</i> , 2022 , 9, 831260	4.9	1
395	Cross-Cancer Genome-Wide Association Study of Endometrial Cancer and Epithelial Ovarian Cancer Identifies Genetic Risk Regions Associated with Risk of Both Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021 , 30, 217-228	4	7
394	Land use mix and leukocyte telomere length in Mexican Americans. <i>Scientific Reports</i> , 2021 , 11, 19742	4.9	0
393	Different Transmission Dynamics of Coronavirus Disease 2019 (COVID-19) and Influenza Suggest the Relative Efficiency of Isolation/Quarantine and Social Distancing Against COVID-19 in China. <i>Clinical Infectious Diseases</i> , 2021 , 73, e4305-e4311	11.6	8
392	Smoking and nasopharyngeal cancer: individual data meta-analysis of six prospective studies on 334'935 men. <i>International Journal of Epidemiology</i> , 2021 , 50, 975-986	7.8	2
391	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. <i>American Journal of Human Genetics</i> , 2021 , 108, 564-582	11	7
390	Human ribonuclease 1 serves as a secretory ligand of ephrin A4 receptor and induces breast tumor initiation. <i>Nature Communications</i> , 2021 , 12, 2788	17.4	3
389	Identification of 22 susceptibility loci associated with testicular germ cell tumors. <i>Nature Communications</i> , 2021 , 12, 4487	17.4	5
388	Population-based targeted sequencing of 54 candidate genes identifies as a susceptibility gene for high-grade serous ovarian cancer. <i>Journal of Medical Genetics</i> , 2021 , 58, 305-313	5.8	12
387	Circulating adipokine concentrations and risk of five obesity-related cancers: A Mendelian randomization study. <i>International Journal of Cancer</i> , 2021 , 148, 1625-1636	7.5	11
386	Integration of multiomic annotation data to prioritize and characterize inflammation and immune-related risk variants in squamous cell lung cancer. <i>Genetic Epidemiology</i> , 2021 , 45, 99-114	2.6	2
385	Immune checkpoint-related serum proteins and genetic variants predict outcomes of localized prostate cancer, a cohort study. <i>Cancer Immunology, Immunotherapy</i> , 2021 , 70, 701-712	7.4	22
384	Trend Analysis and Intervention Effect Starting Point Detection of COVID-19 Epidemics Using Recalibrated Time Series Models - Worldwide, 2020. <i>China CDC Weekly</i> , 2021 , 3, 417-422	4	1
383	Genetic determinants of multiple myeloma risk within the Wnt/beta-catenin signaling pathway. <i>Cancer Epidemiology</i> , 2021 , 73, 101972	2.8	
382	The impact of health education videos on general public's mental health and behavior during COVID-19. <i>Global Health Research and Policy</i> , 2021 , 6, 37	7.6	2

381	High NAFLD fibrosis score in non-alcoholic fatty liver disease as a predictor of carotid plaque development: a retrospective cohort study based on regular health check-up data in China. <i>Annals of Medicine</i> , 2021 , 53, 1621-1631	1.5	1
380	Cohort profile: The National Colorectal Cancer Cohort (NCRCC) study in China.. <i>BMJ Open</i> , 2021 , 11, e051397	1.3	0
379	Perinatal depressive and anxiety symptoms of pregnant women during the coronavirus disease 2019 outbreak in China. <i>American Journal of Obstetrics and Gynecology</i> , 2020 , 223, 240.e1-240.e9	6.4	223
378	Genetic variants in epithelial-mesenchymal transition genes as predictors of clinical outcomes in localized prostate cancer. <i>Carcinogenesis</i> , 2020 , 41, 1057-1064	4.6	
377	Patterns of racial/ethnic disparities in baseline health-related quality of life and relationship with overall survival in patients with colorectal cancer. <i>Quality of Life Research</i> , 2020 , 29, 2977-2986	3.7	3
376	Coinherited genetics of multiple myeloma and its precursor, monoclonal gammopathy of undetermined significance. <i>Blood Advances</i> , 2020 , 4, 2789-2797	7.8	8
375	A Public Health Perspective on Preventing and Controlling the Spread of Coronavirus Disease 2019. <i>China CDC Weekly</i> , 2020 , 2, 237-240	4	
374	Comprehensive T cell repertoire characterization of non-small cell lung cancer. <i>Nature Communications</i> , 2020 , 11, 603	17.4	67
373	HIF3A DNA methylation, obesity and weight gain, and breast cancer risk among Mexican American women. <i>Obesity Research and Clinical Practice</i> , 2020 , 14, 548-553	5.4	4
372	Genetic associations of T cell cancer immune response-related genes with T cell phenotypes and clinical outcomes of early-stage lung cancer 2020 , 8,		5
371	Validation of plasma metabolites associated with breast cancer risk among Mexican Americans. <i>Cancer Epidemiology</i> , 2020 , 69, 101826	2.8	0
370	Household transmission of COVID-19-a systematic review and meta-analysis. <i>Journal of Infection</i> , 2020 , 81, 979-997	18.9	68
369	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019 , 10, 431	17.4	45
368	Elevated Platelet Count Appears to Be Causally Associated with Increased Risk of Lung Cancer: A Mendelian Randomization Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019 , 28, 935-942	4	12
367	Genetic interaction analysis among oncogenesis-related genes revealed novel genes and networks in lung cancer development. <i>Oncotarget</i> , 2019 , 10, 1760-1774	3.3	12
366	Nanotrap-enabled quantification of KRAS-induced peptide hydroxylation in blood for cancer early detection. <i>Nano Research</i> , 2019 , 12, 1445-1452	10	3
365	Circulating obesity-driven biomarkers are associated with risk of clear cell renal cell carcinoma: a two-stage, case-control study. <i>Carcinogenesis</i> , 2019 , 40, 1191-1197	4.6	8
364	Cross-Cancer Pleiotropic Associations with Lung Cancer Risk in African Americans. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019 , 28, 715-723	4	8

363	Breast cancer risk in relation to plasma metabolites among Hispanic and African American women. <i>Breast Cancer Research and Treatment</i> , 2019 , 176, 687-696	4.4	7
362	Metabolic hormones and breast cancer risk among Mexican American Women in the Mano a Mano Cohort Study. <i>Scientific Reports</i> , 2019 , 9, 9989	4.9	6
361	Investigation of Leukocyte Telomere Length and Genetic Variants in Chromosome 5p15.33 as Prognostic Markers in Lung Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019 , 28, 1228-1237	4	5
360	Sex specific associations in genome wide association analysis of renal cell carcinoma. <i>European Journal of Human Genetics</i> , 2019 , 27, 1589-1598	5.3	15
359	Soluble immune checkpoint-related proteins as predictors of tumor recurrence, survival, and T cell phenotypes in clear cell renal cell carcinoma patients 2019 , 7, 334		51
358	The influence of obesity-related factors in the etiology of renal cell carcinoma-A mendelian randomization study. <i>PLoS Medicine</i> , 2019 , 16, e1002724	11.6	38
357	A 5-microRNA signature identified from serum microRNA profiling predicts survival in patients with advanced stage non-small cell lung cancer. <i>Carcinogenesis</i> , 2019 , 40, 643-650	4.6	23
356	Systematic analyses of regulatory variants in DNase I hypersensitive sites identified two novel lung cancer susceptibility loci. <i>Carcinogenesis</i> , 2019 , 40, 432-440	4.6	3
355	Genetic associations of T cell cancer immune response with tumor aggressiveness in localized prostate cancer patients and disease reclassification in an active surveillance cohort. <i>Oncotmunology</i> , 2019 , 8, e1483303	7.2	6
354	Mendelian Randomization and mediation analysis of leukocyte telomere length and risk of lung and head and neck cancers. <i>International Journal of Epidemiology</i> , 2019 , 48, 751-766	7.8	14
353	Sleep duration and risk of cancer in the Mexican American Mano-a-Mano Cohort. <i>Sleep Health</i> , 2019 , 5, 78-83	4	9
352	Reply to Q Mosaic loss of chromosome Y in leukocytes matters Q <i>Nature Genetics</i> , 2019 , 51, 7-9	36.3	6
351	Is folic acid safe for non-muscle-invasive bladder cancer patients? An evidence-based cohort study. <i>American Journal of Clinical Nutrition</i> , 2018 , 107, 208-216	7	12
350	Genetic variants in cytokine signaling pathways and clinical outcomes in early-stage lung cancer patients. <i>Journal of Thoracic and Cardiovascular Surgery</i> , 2018 , 155, 2635-2645.e15	1.5	2
349	Germline genetic variants in somatically significantly mutated genes in tumors are associated with renal cell carcinoma risk and outcome. <i>Carcinogenesis</i> , 2018 , 39, 752-757	4.6	9
348	Adult height is associated with increased risk of ovarian cancer: a Mendelian randomisation study. <i>British Journal of Cancer</i> , 2018 , 118, 1123-1129	8.7	10
347	Determinants and prognostic value of quality of life in patients with pancreatic ductal adenocarcinoma. <i>European Journal of Cancer</i> , 2018 , 92, 20-32	7.5	14
346	Genome-wide interaction study of smoking behavior and non-small cell lung cancer risk in Caucasian population. <i>Carcinogenesis</i> , 2018 , 39, 336-346	4.6	11

345	Cancer risk associated with chronic diseases and disease markers: prospective cohort study. <i>BMJ, The</i> , 2018 , 360, k134	5.9	57
344	Phase I study of nab-paclitaxel, gemcitabine, and bevacizumab in patients with advanced cancers. <i>British Journal of Cancer</i> , 2018 , 118, 1419-1424	8.7	4
343	Angiogenin/Ribonuclease 5 Is an EGFR Ligand and a Serum Biomarker for Erlotinib Sensitivity in Pancreatic Cancer. <i>Cancer Cell</i> , 2018 , 33, 752-769.e8	24.3	31
342	The somatic mutation landscape of premalignant colorectal adenoma. <i>Gut</i> , 2018 , 67, 1299-1305	19.2	33
341	Novel genetic variants in the P38MAPK pathway gene ZAK and susceptibility to lung cancer. <i>Molecular Carcinogenesis</i> , 2018 , 57, 216-224	5	5
340	Analysis of Genomes and Transcriptomes of Hepatocellular Carcinomas Identifies Mutations and Gene Expression Changes in the Transforming Growth Factor- β Pathway. <i>Gastroenterology</i> , 2018 , 154, 195-210	13.3	68
339	Serum MicroRNA-150 Predicts Prognosis for Early-Stage Non-Small Cell Lung Cancer and Promotes Tumor Cell Proliferation by Targeting Tumor Suppressor Gene SRCIN1. <i>Clinical Pharmacology and Therapeutics</i> , 2018 , 103, 1061-1073	6.1	28
338	Socio-demographic, Clinical, and Genetic Determinants of Quality of Life in Lung Cancer Patients. <i>Scientific Reports</i> , 2018 , 8, 10640	4.9	6
337	Variants in genes encoding small GTPases and association with epithelial ovarian cancer susceptibility. <i>PLoS ONE</i> , 2018 , 13, e0197561	3.7	9
336	rs495139 in the TYMS-ENOSF1 Region and Risk of Ovarian Carcinoma of Mucinous Histology. <i>International Journal of Molecular Sciences</i> , 2018 , 19,	6.3	3
335	Identification of susceptibility pathways for the role of chromosome 15q25.1 in modifying lung cancer risk. <i>Nature Communications</i> , 2018 , 9, 3221	17.4	29
334	Serum microRNAs as predictors of risk for non-muscle invasive bladder cancer. <i>Oncotarget</i> , 2018 , 9, 14895-14908	3.3	14908
333	Associations of blood mitochondrial DNA copy number with social-demographics and cancer risk: results from the Mano-A-Mano Mexican American Cohort. <i>Oncotarget</i> , 2018 , 9, 25491-25502	3.3	5
332	Genome-wide association study of familial lung cancer. <i>Carcinogenesis</i> , 2018 , 39, 1135-1140	4.6	24
331	Prevalence of Aflatoxin-Associated Mutation in Hepatocellular Carcinoma in Hispanics in South Texas. <i>Cancer Prevention Research</i> , 2018 , 11, 103-112	3.2	13
330	Dietary patterns and risk of recurrence and progression in non-muscle-invasive bladder cancer. <i>International Journal of Cancer</i> , 2018 , 142, 1797-1804	7.5	10
329	Global and targeted circulating microRNA profiling of colorectal adenoma and colorectal cancer. <i>Cancer</i> , 2018 , 124, 785-796	6.4	36
328	Fine mapping of MHC region in lung cancer highlights independent susceptibility loci by ethnicity. <i>Nature Communications</i> , 2018 , 9, 3927	17.4	24

327	Consensus report of the 8 and 9th Weinman Symposia on Gene x Environment Interaction in carcinogenesis: novel opportunities for precision medicine. <i>Cell Death and Differentiation</i> , 2018 , 25, 1885-1904 ^{12.7} ¹⁷		
326	Serum miR-331-3p predicts tumor recurrence in esophageal adenocarcinoma. <i>Scientific Reports</i> , 2018 , 8, 14006	4.9	18
325	Global and Targeted miRNA Expression Profiling in Clear Cell Renal Cell Carcinoma Tissues Potentially Links miR-155-5p and miR-210-3p to both Tumorigenesis and Recurrence. <i>American Journal of Pathology</i> , 2018 , 188, 2487-2496	5.8	23
324	Prognostic role of elevated mir-24-3p in breast cancer and its association with the metastatic process. <i>Oncotarget</i> , 2018 , 9, 12868-12878	3.3	32
323	Cohort Profile: The Mexican American Mano a Mano Cohort. <i>International Journal of Epidemiology</i> , 2017 , 46, e3	7.8	26
322	Genetic polymorphisms in genes related to risk-taking behaviours predicting body mass index trajectory among Mexican American adolescents. <i>Pediatric Obesity</i> , 2017 , 12, 356-362	4.6	9
321	D-mannose: a novel prognostic biomarker for patients with esophageal adenocarcinoma. <i>Carcinogenesis</i> , 2017 , 38, 162-167	4.6	9
320	Genetic Variants in Epigenetic Pathways and Risks of Multiple Cancers in the GAME-ON Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017 , 26, 816-825	4	7
319	Cancer Epidemiology 2017 , 1-9		
318	Cohort Profile: The Taiwan MJ Cohort: half a million Chinese with repeated health surveillance data. <i>International Journal of Epidemiology</i> , 2017 , 46, 1744-1744g	7.8	34
317	Genetic variants of PTPN2 are associated with lung cancer risk: a re-analysis of eight GWASs in the TRICL-ILCCO consortium. <i>Scientific Reports</i> , 2017 , 7, 825	4.9	6
316	Low serum testosterone is associated with tumor aggressiveness and poor prognosis in prostate cancer. <i>Oncology Letters</i> , 2017 , 13, 1949-1957	2.6	12
315	Associations between language acculturation, age of immigration, and obesity in the Mexican American Mano A Mano cohort. <i>Obesity Research and Clinical Practice</i> , 2017 , 11, 544-557	5.4	7
314	Preleukaemic clonal haemopoiesis and risk of therapy-related myeloid neoplasms: a case-control study. <i>Lancet Oncology</i> , 2017 , 18, 100-111	21.7	189
313	Personalized Prognostic Prediction Models for Breast Cancer Recurrence and Survival Incorporating Multidimensional Data. <i>Journal of the National Cancer Institute</i> , 2017 , 109,	9.7	23
312	Potential Susceptibility Loci Identified for Renal Cell Carcinoma by Targeting Obesity-Related Genes. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017 , 26, 1436-1442	4	0
311	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. <i>Nature Communications</i> , 2017 , 8, 15724	17.4	50
310	Prognostic significance of promoter CpG island methylation of obesity-related genes in patients with nonmetastatic renal cell carcinoma. <i>Cancer</i> , 2017 , 123, 3617-3627	6.4	17

309	Diabetes with early kidney involvement may shorten life expectancy by 16 years. <i>Kidney International</i> , 2017 , 92, 388-396	9.9	51
308	Large-scale association analysis identifies new lung cancer susceptibility loci and heterogeneity in genetic susceptibility across histological subtypes. <i>Nature Genetics</i> , 2017 , 49, 1126-1132	36.3	246
307	A Serological Biopsy Using Five Stomach-Specific Circulating Biomarkers for Gastric Cancer Risk Assessment: A Multi-Phase Study. <i>American Journal of Gastroenterology</i> , 2017 , 112, 704-715	0.7	55
306	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017 , 49, 680-691	36.3	190
305	Functional variants in DCAF4 associated with lung cancer risk in European populations. <i>Carcinogenesis</i> , 2017 , 38, 541-551	4.6	13
304	Plasma MicroRNA signature predicting weight gain among Mexican-American women. <i>Obesity</i> , 2017 , 25, 958-964	8	8
303	High baseline levels of interleukin-8 in leukocytes and urine predict tumor recurrence in non-muscle invasive bladder cancer patients receiving bacillus Calmette-Guerin therapy: A long-term survival analysis. <i>Oncotarget</i> , 2017 , 8, e1265719	7.2	13
302	Associations between genetic variants in mRNA splicing-related genes and risk of lung cancer: a pathway-based analysis from published GWASs. <i>Scientific Reports</i> , 2017 , 7, 44634	4.9	6
301	Measurement of DNA damage in peripheral blood by the γ H2AX assay as predictor of colorectal cancer risk. <i>DNA Repair</i> , 2017 , 53, 24-30	4.3	14
300	Energy stress-induced lncRNA FILNC1 represses c-Myc-mediated energy metabolism and inhibits renal tumor development. <i>Nature Communications</i> , 2017 , 8, 783	17.4	120
299	Common, germline genetic variations in the novel tumor suppressor and risk of developing different types of cancer. <i>Oncotarget</i> , 2017 , 8, 74936-74946	3.3	14
298	Polymorphisms in genes related to epithelial-mesenchymal transition and risk of non-small cell lung cancer. <i>Carcinogenesis</i> , 2017 , 38, 1029-1035	4.6	17
297	Hypertension Susceptibility Loci are Associated with Anthracycline-related Cardiotoxicity in Long-term Childhood Cancer Survivors. <i>Scientific Reports</i> , 2017 , 7, 9698	4.9	18
296	A miR-SNP biomarker linked to an increased lung cancer survival by miRNA-mediated down-regulation of FZD4 expression and Wnt signaling. <i>Scientific Reports</i> , 2017 , 7, 9029	4.9	12
295	Methylation of subtelomeric repeat D4Z4 in peripheral blood leukocytes is associated with biochemical recurrence in localized prostate cancer patients. <i>Carcinogenesis</i> , 2017 , 38, 821-826	4.6	5
294	Genetic variants in telomere-maintenance genes are associated with ovarian cancer risk and outcome. <i>Journal of Cellular and Molecular Medicine</i> , 2017 , 21, 510-518	5.6	6
293	Genetic Variants Related to Longer Telomere Length are Associated with Increased Risk of Renal Cell Carcinoma. <i>European Urology</i> , 2017 , 72, 747-754	10.2	27
292	Circulating metabolite profiles to predict overall survival in advanced non-small cell lung cancer patients receiving first-line chemotherapy. <i>Lung Cancer</i> , 2017 , 114, 70-78	5.9	8

291	Global and targeted serum metabolic profiling of colorectal cancer progression. <i>Cancer</i> , 2017 , 123, 4066-4074	38	
290	Susceptibility loci of CNOT6 in the general mRNA degradation pathway and lung cancer risk-A re-analysis of eight GWASs. <i>Molecular Carcinogenesis</i> , 2017 , 56, 1227-1238	5	6
289	Predictors of health-related quality of life and association with survival may identify colorectal cancer patients at high risk of poor prognosis. <i>Quality of Life Research</i> , 2017 , 26, 319-330	3.7	15
288	Glycemic index, glycemic load and carbohydrate intake in association with risk of renal cell carcinoma. <i>Carcinogenesis</i> , 2017 , 38, 1129-1135	4.6	6
287	Copy number alterations detected as clonal hematopoiesis of indeterminate potential. <i>Blood Advances</i> , 2017 , 1, 1031-1036	7.8	21
286	Obesity, metabolic factors and risk of different histological types of lung cancer: A Mendelian randomization study. <i>PLoS ONE</i> , 2017 , 12, e0177875	3.7	56
285	Social-demographics, health behaviors, and telomere length in the Mexican American Mano a Mano Cohort. <i>Oncotarget</i> , 2017 , 8, 96553-96567	3.3	11
284	Genetic variants in the inflammation pathway as predictors of recurrence and progression in non-muscle invasive bladder cancer treated with Bacillus Calmette-Guérin. <i>Oncotarget</i> , 2017 , 8, 88782-88791	3.3	3
283	LINE-1 methylation in peripheral blood leukocytes and clinical characteristics and prognosis of prostate cancer patients. <i>Oncotarget</i> , 2017 , 8, 94020-94027	3.3	6
282	Epidemiology of Renal Cell Carcinoma 2017 , 1-18		6
281	Genetic variants of the Wnt signaling pathway as predictors of aggressive disease and reclassification in men with early stage prostate cancer on active surveillance. <i>Carcinogenesis</i> , 2016 , 37, 965-971	4.6	4
280	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016 , 7, 11843	17.4	59
279	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. <i>Nature Communications</i> , 2016 , 7, 10933	17.4	70
278	Personalized Risk Assessment in Never, Light, and Heavy Smokers in a prospective cohort in Taiwan. <i>Scientific Reports</i> , 2016 , 6, 36482	4.9	21
277	Different dietary patterns and reduction of lung cancer risk: A large case-control study in the U.S. <i>Scientific Reports</i> , 2016 , 6, 26760	4.9	10
276	Lower mitochondrial DNA copy number in peripheral blood leukocytes increases the risk of endometrial cancer. <i>Molecular Carcinogenesis</i> , 2016 , 55, 1111-7	5	11
275	Acculturation and Diabetes Risk in the Mexican American Mano a Mano Cohort. <i>American Journal of Public Health</i> , 2016 , 106, 547-9	5.1	18
274	Novel fluorescence in-situ hybridization-based definition of bacille Calmette-Guérin (BCG) failure for use in enhancing recruitment into clinical trials of intravesical therapies. <i>BJU International</i> , 2016 , 117, 754-60	5.6	26

273	Germline Genetic Variants in the Wnt/ β Catenin Pathway as Predictors of Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016 , 25, 540-6	4	9
272	Genomic DNA Hypomethylation and Risk of Renal Cell Carcinoma: A Case-Control Study. <i>Clinical Cancer Research</i> , 2016 , 22, 2074-82	12.9	18
271	Genetically predicted longer telomere length is associated with increased risk of B-cell lymphoma subtypes. <i>Human Molecular Genetics</i> , 2016 , 25, 1663-76	5.6	39
270	Cohort Profile: The MD Anderson Cancer Patients and Survivors Cohort (MDA-CPSC). <i>International Journal of Epidemiology</i> , 2016 , 45, 713-713f	7.8	10
269	Identification of a novel susceptibility locus at 13q34 and refinement of the 20p12.2 region as a multi-signal locus associated with bladder cancer risk in individuals of European ancestry. <i>Human Molecular Genetics</i> , 2016 , 25, 1203-14	5.6	20
268	Polymorphisms of the centrosomal gene (FGFR1OP) and lung cancer risk: a meta-analysis of 14,463 cases and 44,188 controls. <i>Carcinogenesis</i> , 2016 , 37, 280-289	4.6	6
267	Pathway analysis of bladder cancer genome-wide association study identifies novel pathways involved in bladder cancer development. <i>Genes and Cancer</i> , 2016 , 7, 229-239	2.9	11
266	Assessment of variation in immunosuppressive pathway genes reveals TGFBR2 to be associated with risk of clear cell ovarian cancer. <i>Oncotarget</i> , 2016 , 7, 69097-69110	3.3	4
265	Inherited variants affecting RNA editing may contribute to ovarian cancer susceptibility: results from a large-scale collaboration. <i>Oncotarget</i> , 2016 , 7, 72381-72394	3.3	11
264	Two-stage induced differentiation of OCT4+/Nanog+ stem-like cells in lung adenocarcinoma. <i>Oncotarget</i> , 2016 , 7, 68360-68370	3.3	8
263	Metabolomics profiling in plasma samples from glioma patients correlates with tumor phenotypes. <i>Oncotarget</i> , 2016 , 7, 20486-95	3.3	35
262	Heritability of prostate cancer: a tale of rare variants and common single nucleotide polymorphisms. <i>Annals of Translational Medicine</i> , 2016 , 4, 206	3.2	10
261	Mobile Phone Use and its Association With Sitting Time and Meeting Physical Activity Recommendations in a Mexican American Cohort. <i>JMIR MHealth and UHealth</i> , 2016 , 4, e54	5.5	3
260	Genetic variations in apoptosis pathway and the risk of ovarian cancer. <i>Oncotarget</i> , 2016 , 7, 56737-56745	5.3	1
259	Clonal Hematopoiesis Increases Risk of Therapy-Related Myeloid Neoplasms. <i>Blood</i> , 2016 , 128, 38-38	2.2	
258	Association between Genetic Variants in DNA Double-Strand Break Repair Pathways and Risk of Radiation Therapy-Induced Pneumonitis and Esophagitis in Non-Small Cell Lung Cancer. <i>Cancers</i> , 2016 , 8,	6.6	8
257	MiRNA-Related Genetic Variations Associated with Radiotherapy-Induced Toxicities in Patients with Locally Advanced Non-Small Cell Lung Cancer. <i>PLoS ONE</i> , 2016 , 11, e0150467	3.7	5
256	The Role of Physical Activity in Harm Reduction among Betel Quid Chewers from a Prospective Cohort of 419,378 Individuals. <i>PLoS ONE</i> , 2016 , 11, e0152246	3.7	3

255	Mutational Profiles Reveal an Aberrant TGF- β -CEA Regulated Pathway in Colon Adenomas. <i>PLoS ONE</i> , 2016 , 11, e0153933	3.7	13
254	Adult body mass index and risk of ovarian cancer by subtype: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2016 , 45, 884-95	7.8	45
253	PALB2, CHEK2 and ATM rare variants and cancer risk: data from COGS. <i>Journal of Medical Genetics</i> , 2016 , 53, 800-811	5.8	121
252	Genome-wide association study confirms lung cancer susceptibility loci on chromosomes 5p15 and 15q25 in an African-American population. <i>Lung Cancer</i> , 2016 , 98, 33-42	5.9	31
251	Glycemic Index, Glycemic Load, and Lung Cancer Risk in Non-Hispanic Whites. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016 , 25, 532-9	4	28
250	Assessing the genetic architecture of epithelial ovarian cancer histological subtypes. <i>Human Genetics</i> , 2016 , 135, 741-56	6.3	18
249	Mosaic loss of chromosome Y is associated with common variation near TCL1A. <i>Nature Genetics</i> , 2016 , 48, 563-8	36.3	87
248	Association of vitamin D levels and risk of ovarian cancer: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2016 , 45, 1619-1630	7.8	77
247	Genetic Risk Can Be Decreased: Quitting Smoking Decreases and Delays Lung Cancer for Smokers With High and Low CHRNA5 Risk Genotypes - A Meta-Analysis. <i>EBioMedicine</i> , 2016 , 11, 219-226	8.8	27
246	Gene-environment interaction of genome-wide association study-identified susceptibility loci and meat-cooking mutagens in the etiology of renal cell carcinoma. <i>Cancer</i> , 2016 , 122, 108-15	6.4	18
245	Biomarkers for Assessing Risk of Cancer 2015 , 317-330.e3		
244	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015 , 47, 164-71	36.3	177
243	ABO blood types and cancer risk--a cohort study of 339,432 subjects in Taiwan. <i>Cancer Epidemiology</i> , 2015 , 39, 150-6	2.8	22
242	Network-Based Integration of GWAS and Gene Expression Identifies a HOX-Centric Network Associated with Serous Ovarian Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 1574-84	4	24
241	Genetic Variants in the Wnt/ β -Catenin Signaling Pathway as Indicators of Bladder Cancer Risk. <i>Journal of Urology</i> , 2015 , 194, 1771-6	2.5	30
240	Evaluating the ovarian cancer gonadotropin hypothesis: a candidate gene study. <i>Gynecologic Oncology</i> , 2015 , 136, 542-8	4.9	12
239	Depressive symptoms and short telomere length are associated with increased mortality in bladder cancer patients. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 336-43	4	31
238	Mitochondrial DNA Content as Risk Factor for Bladder Cancer and Its Association with Mitochondrial DNA Polymorphisms. <i>Cancer Prevention Research</i> , 2015 , 8, 607-13	3.2	15

237	The ability of bilirubin in identifying smokers with higher risk of lung cancer: a large cohort study in conjunction with global metabolomic profiling. <i>Clinical Cancer Research</i> , 2015 , 21, 193-200	12.9	36
236	Genetic Variations in Glutathione Pathway Genes Predict Cancer Recurrence in Patients Treated with Transurethral Resection and Bacillus Calmette-Guerin Instillation for Non-muscle Invasive Bladder Cancer. <i>Annals of Surgical Oncology</i> , 2015 , 22, 4104-10	3.1	24
235	CYP2A6 reduced activity gene variants confer reduction in lung cancer risk in African American smokers--findings from two independent populations. <i>Carcinogenesis</i> , 2015 , 36, 99-103	4.6	33
234	Cis-eQTL analysis and functional validation of candidate susceptibility genes for high-grade serous ovarian cancer. <i>Nature Communications</i> , 2015 , 6, 8234	17.4	40
233	Identification of lung cancer histology-specific variants applying Bayesian framework variant prioritization approaches within the TRICL and ILCCO consortia. <i>Carcinogenesis</i> , 2015 , 36, 1314-26	4.6	8
232	Common variants at the CHEK2 gene locus and risk of epithelial ovarian cancer. <i>Carcinogenesis</i> , 2015 , 36, 1341-53	4.6	20
231	Association of leukocyte telomere length in peripheral blood leukocytes with endometrial cancer risk in Caucasian Americans. <i>Carcinogenesis</i> , 2015 , 36, 1327-32	4.6	12
230	Mitochondrial DNA copy number in peripheral blood leukocytes and the risk of clear cell renal cell carcinoma. <i>Carcinogenesis</i> , 2015 , 36, 249-55	4.6	21
229	Prognostic significance of pretreatment serum levels of albumin, LDH and total bilirubin in patients with non-metastatic breast cancer. <i>Carcinogenesis</i> , 2015 , 36, 243-8	4.6	83
228	Genetic variations in base excision repair pathway and risk of bladder cancer: a case-control study in the United States. <i>Molecular Carcinogenesis</i> , 2015 , 54, 50-7	5	9
227	Epigenetic analysis of microRNA genes in tumors from surgically resected lung cancer patients and association with survival. <i>Molecular Carcinogenesis</i> , 2015 , 54 Suppl 1, E45-51	5	11
226	Genetic variation in the TNF/TRAF2/ASK1/p38 kinase signaling pathway as markers for postoperative pulmonary complications in lung cancer patients. <i>Scientific Reports</i> , 2015 , 5, 12068	4.9	10
225	Acculturation, sociodemographic and lifestyle factors associated with compliance with physical activity recommendations in the Mexican-American Mano A Mano cohort. <i>BMJ Open</i> , 2015 , 5, e008302	3	9
224	Epithelial-Mesenchymal Transition (EMT) Gene Variants and Epithelial Ovarian Cancer (EOC) Risk. <i>Genetic Epidemiology</i> , 2015 , 39, 689-97	2.6	18
223	Determination of proline in human serum by a robust LC-MS/MS method: application to identification of human metabolites as candidate biomarkers for esophageal cancer early detection and risk stratification. <i>Biomedical Chromatography</i> , 2015 , 29, 570-7	1.7	14
222	Multilevel-analysis identify a cis-expression quantitative trait locus associated with risk of renal cell carcinoma. <i>Oncotarget</i> , 2015 , 6, 4097-109	3.3	1
221	Common Genetic Variation In Cellular Transport Genes and Epithelial Ovarian Cancer (EOC) Risk. <i>PLoS ONE</i> , 2015 , 10, e0128106	3.7	15
220	The Relationship between Native American Ancestry, Body Mass Index and Diabetes Risk among Mexican-Americans. <i>PLoS ONE</i> , 2015 , 10, e0141260	3.7	19

219	A genome-wide association study of marginal zone lymphoma shows association to the HLA region. <i>Nature Communications</i> , 2015 , 6, 5751	17.4	44
218	Characterization of large structural genetic mosaicism in human autosomes. <i>American Journal of Human Genetics</i> , 2015 , 96, 487-97	11	77
217	Identification of Serum Markers of Esophageal Adenocarcinoma by Global and Targeted Metabolic Profiling. <i>Clinical Gastroenterology and Hepatology</i> , 2015 , 13, 1730-1737.e9	6.9	25
216	Mitochondrial DNA copy number in peripheral blood leukocytes and the aggressiveness of localized prostate cancer. <i>Oncotarget</i> , 2015 , 6, 41988-96	3.3	20
215	Common Genetic Variation in Circadian Rhythm Genes and Risk of Epithelial Ovarian Cancer (EOC). <i>Journal of Genetics and Genome Research</i> , 2015 , 2,		22
214	Mutagen Sensitivity 2015 , 1-4		
213	Genome-wide association study identifies multiple loci associated with bladder cancer risk. <i>Human Molecular Genetics</i> , 2014 , 23, 1387-98	5.6	101
212	Telomere length in peripheral blood leukocytes and lung cancer risk: a large case-control study in Caucasians. <i>Cancer Research</i> , 2014 , 74, 2476-86	10.1	67
211	A genome-wide association study of renal cell carcinoma among African Americans. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014 , 23, 209-14	4	17
210	The 19q12 bladder cancer GWAS signal: association with cyclin E function and aggressive disease. <i>Cancer Research</i> , 2014 , 74, 5808-18	10.1	19
209	The prostate cancer susceptibility variant rs2735839 near KLK3 gene is associated with aggressive prostate cancer and can stratify gleason score 7 patients. <i>Clinical Cancer Research</i> , 2014 , 20, 5133-5139	12.9	25
208	Genome-wide association study identifies multiple susceptibility loci for diffuse large B cell lymphoma. <i>Nature Genetics</i> , 2014 , 46, 1233-8	36.3	108
207	Genome-wide association study identifies five susceptibility loci for follicular lymphoma outside the HLA region. <i>American Journal of Human Genetics</i> , 2014 , 95, 462-71	11	74
206	Joint association of genome-wide association study-identified susceptibility loci and dietary patterns in risk of renal cell carcinoma among non-Hispanic whites. <i>American Journal of Epidemiology</i> , 2014 , 180, 499-507	3.8	16
205	Increased leukocyte mitochondrial DNA copy number is associated with oral premalignant lesions: an epidemiology study. <i>Carcinogenesis</i> , 2014 , 35, 1760-4	4.6	15
204	Intratumor heterogeneity in localized lung adenocarcinomas delineated by multiregion sequencing. <i>Science</i> , 2014 , 346, 256-9	33.3	659
203	Rare variants of large effect in BRCA2 and CHEK2 affect risk of lung cancer. <i>Nature Genetics</i> , 2014 , 46, 736-41	36.3	228
202	Lung Cancer Susceptibility and Risk Assessment Models 2014 , 25-47		

201	Integration of Platinum Nanoparticles with a Volumetric Bar-Chart Chip for Biomarker Assays. <i>Angewandte Chemie</i> , 2014 , 126, 12659-12663	3.6	11
200	Inflammation-related genetic variations and survival in patients with advanced non-small cell lung cancer receiving first-line chemotherapy. <i>Clinical Pharmacology and Therapeutics</i> , 2014 , 96, 360-369	6.1	13
199	Genetic variants of the Wnt signaling pathway as predictors of recurrence and survival in early-stage non-small cell lung cancer patients. <i>Carcinogenesis</i> , 2014 , 35, 1284-91	4.6	19
198	High serum iron is associated with increased cancer risk. <i>Cancer Research</i> , 2014 , 74, 6589-97	10.1	55
197	A validated miRNA profile predicts response to therapy in esophageal adenocarcinoma. <i>Cancer</i> , 2014 , 120, 3635-41	6.4	42
196	Genetic and intermediate phenotypic susceptibility markers of gastric cancer in Hispanic Americans: a case-control study. <i>Cancer</i> , 2014 , 120, 3040-8	6.4	32
195	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. <i>Human Molecular Genetics</i> , 2014 , 23, 6616-33	5.6	77
194	Development and validation of risk models and molecular diagnostics to permit personalized management of cancer. <i>Cancer</i> , 2014 , 120, 11-9	6.4	9
193	Mutagen sensitivity as measured by induced chromatid breakage as a marker of cancer risk. <i>Methods in Molecular Biology</i> , 2014 , 1105, 183-92	1.4	
192	Testicular germ cell tumor susceptibility associated with the UCK2 locus on chromosome 1q23. <i>Human Molecular Genetics</i> , 2013 , 22, 2748-53	5.6	53
191	Pathway-based serum microRNA profiling and survival in patients with advanced stage non-small cell lung cancer. <i>Cancer Research</i> , 2013 , 73, 4801-9	10.1	78
190	Long telomeres in peripheral blood leukocytes are associated with an increased risk of soft tissue sarcoma. <i>Cancer</i> , 2013 , 119, 1885-91	6.4	34
189	Genetic variations in regulator of G-protein signaling (RGS) confer risk of bladder cancer. <i>Cancer</i> , 2013 , 119, 1643-51	6.4	11
188	Polymorphisms in the XRCC1 gene modify survival of bladder cancer patients treated with chemotherapy. <i>International Journal of Cancer</i> , 2013 , 133, 2004-9	7.5	26
187	Meta-analysis identifies four new loci associated with testicular germ cell tumor. <i>Nature Genetics</i> , 2013 , 45, 680-5	36.3	132
186	MicroRNA-related genetic variants associated with clinical outcomes in early-stage non-small cell lung cancer patients. <i>Cancer Research</i> , 2013 , 73, 1867-75	10.1	38
185	Ionizing radiation-induced H2AX activity in whole blood culture and the risk of lung cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013 , 22, 443-51	4	14
184	Association of mitochondrial DNA copy number in peripheral blood leukocytes with risk of esophageal adenocarcinoma. <i>Carcinogenesis</i> , 2013 , 34, 2521-4	4.6	34

183	Energy balance, polymorphisms in the mTOR pathway, and renal cell carcinoma risk. <i>Journal of the National Cancer Institute</i> , 2013 , 105, 424-32	9.7	25
182	Genome-wide association study of genetic predictors of overall survival for non-small cell lung cancer in never smokers. <i>Cancer Research</i> , 2013 , 73, 4028-38	10.1	40
181	EH2AX level in peripheral blood lymphocytes as a risk predictor for bladder cancer. <i>Carcinogenesis</i> , 2013 , 34, 2543-7	4.6	17
180	Fine-mapping of the 5p15.33, 6p22.1-p21.31, and 15q25.1 regions identifies functional and histology-specific lung cancer susceptibility loci in African-Americans. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013 , 22, 251-60	4	27
179	Risk assessment of esophageal adenocarcinoma using EH2AX assay. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013 , 22, 1797-804	4	10
178	Common variation at 2q22.3 (ZEB2) influences the risk of renal cancer. <i>Human Molecular Genetics</i> , 2013 , 22, 825-31	5.6	49
177	Reduced mitochondrial DNA copy number in peripheral blood leukocytes increases the risk of soft tissue sarcoma. <i>Carcinogenesis</i> , 2013 , 34, 1039-43	4.6	33
176	Short telomere lengths in peripheral blood leukocytes are associated with an increased risk of oral premalignant lesion and oral squamous cell carcinoma. <i>Cancer</i> , 2013 , 119, 4277-83	6.4	27
175	Re: short telomere length, cancer survival, and cancer risk in 47 102 individuals. <i>Journal of the National Cancer Institute</i> , 2013 , 105, 1157	9.7	12
174	MicroRNA expression signatures during malignant progression from Barrett's esophagus to esophageal adenocarcinoma. <i>Cancer Prevention Research</i> , 2013 , 6, 196-205	3.2	77
173	Application of multi-SNP approaches Bayesian LASSO and AUC-RF to detect main effects of inflammatory-gene variants associated with bladder cancer risk. <i>PLoS ONE</i> , 2013 , 8, e83745	3.7	12
172	MicroRNA in the pathogenesis and prognosis of esophageal cancer. <i>Current Pharmaceutical Design</i> , 2013 , 19, 1292-300	3.3	78
171	GWAS-identified colorectal cancer susceptibility loci associated with clinical outcomes. <i>Carcinogenesis</i> , 2012 , 33, 1327-31	4.6	39
170	Genetic variants on 15q25.1, smoking, and lung cancer: an assessment of mediation and interaction. <i>American Journal of Epidemiology</i> , 2012 , 175, 1013-20	3.8	108
169	Predictors of survival in never-smokers with non-small cell lung cancer: a large-scale, two-phase genetic study. <i>Clinical Cancer Research</i> , 2012 , 18, 5983-91	12.9	11
168	Germline prognostic markers for urinary bladder cancer: obstacles and opportunities. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2012 , 30, 524-32	2.8	19
167	Association of polymorphisms in oxidative stress genes with clinical outcomes for bladder cancer treated with Bacillus Calmette-Guérin. <i>PLoS ONE</i> , 2012 , 7, e38533	3.7	23
166	A genome-wide association study identifies a novel susceptibility locus for renal cell carcinoma on 12p11.23. <i>Human Molecular Genetics</i> , 2012 , 21, 456-62	5.6	74

165	Detectable clonal mosaicism and its relationship to aging and cancer. <i>Nature Genetics</i> , 2012 , 44, 651-8	36.3	409
164	Intake of red meat and heterocyclic amines, metabolic pathway genes and bladder cancer risk. <i>International Journal of Cancer</i> , 2012 , 131, 1892-903	7.5	19
163	Identification of polymorphisms in ultraconserved elements associated with clinical outcomes in locally advanced colorectal adenocarcinoma. <i>Cancer</i> , 2012 , 118, 6188-98	6.4	10
162	Hepatocellular carcinoma risk prediction model for the general population: the predictive power of transaminases. <i>Journal of the National Cancer Institute</i> , 2012 , 104, 1599-611	9.7	81
161	Systematic evaluation of apoptotic pathway gene polymorphisms and lung cancer risk. <i>Carcinogenesis</i> , 2012 , 33, 1699-706	4.6	20
160	Influence of common genetic variation on lung cancer risk: meta-analysis of 14 900 cases and 29 485 controls. <i>Human Molecular Genetics</i> , 2012 , 21, 4980-95	5.6	159
159	Genetic variants in telomere-maintenance genes and bladder cancer risk. <i>PLoS ONE</i> , 2012 , 7, e30665	3.7	15
158	Genetic variations in the transforming growth factor beta pathway as predictors of bladder cancer risk. <i>PLoS ONE</i> , 2012 , 7, e51758	3.7	18
157	Association study of nicotinic acetylcholine receptor genes identifies a novel lung cancer susceptibility locus near CHRNA1 in African-Americans. <i>Oncotarget</i> , 2012 , 3, 1428-38	3.3	9
156	Genetic susceptibility to bladder cancer risk and outcome. <i>Personalized Medicine</i> , 2011 , 8, 365-374	2.2	35
155	Minimum amount of physical activity for reduced mortality and extended life expectancy: a prospective cohort study. <i>Lancet, The</i> , 2011 , 378, 1244-53	4.0	1123
154	Genetic variations in the regulator of G-protein signaling genes are associated with survival in late-stage non-small cell lung cancer. <i>PLoS ONE</i> , 2011 , 6, e21120	3.7	18
153	Genome-wide association study of renal cell carcinoma identifies two susceptibility loci on 2p21 and 11q13.3. <i>Nature Genetics</i> , 2011 , 43, 60-5	36.3	199
152	Genetic variation in DNA-repair pathways and response to radiochemotherapy in esophageal adenocarcinoma: a retrospective cohort study of the Eastern Cooperative Oncology Group. <i>BMC Cancer</i> , 2011 , 11, 176	4.8	19
151	A genome-wide association study identifies a locus on chromosome 14q21 as a predictor of leukocyte telomere length and as a marker of susceptibility for bladder cancer. <i>Cancer Prevention Research</i> , 2011 , 4, 514-21	3.2	60
150	A genetic variant near the PMAIP1/Noxa gene is associated with increased bleomycin sensitivity. <i>Human Molecular Genetics</i> , 2011 , 20, 820-6	5.6	10
149	Genetic variations in the transforming growth factor-beta pathway as predictors of survival in advanced non-small cell lung cancer. <i>Carcinogenesis</i> , 2011 , 32, 1050-6	4.6	30
148	Genome-wide association study of survival in non-small cell lung cancer patients receiving platinum-based chemotherapy. <i>Journal of the National Cancer Institute</i> , 2011 , 103, 817-25	9.7	73

147	Novel genetic variants in the chromosome 5p15.33 region associate with lung cancer risk. <i>Carcinogenesis</i> , 2011 , 32, 1493-9	4.6	52
146	Prospective analysis of DNA damage and repair markers of lung cancer risk from the Prostate, Lung, Colorectal and Ovarian (PLCO) Cancer Screening Trial. <i>Carcinogenesis</i> , 2011 , 32, 69-73	4.6	22
145	Variants in inflammation genes are implicated in risk of lung cancer in never smokers exposed to second-hand smoke. <i>Cancer Discovery</i> , 2011 , 1, 420-9	24.4	25
144	A genome-wide association study of bladder cancer identifies a new susceptibility locus within SLC14A1, a urea transporter gene on chromosome 18q12.3. <i>Human Molecular Genetics</i> , 2011 , 20, 4282-9	5.6	82
143	Common genetic variants in cell cycle pathway are associated with survival in stage III-IV non-small-cell lung cancer. <i>Carcinogenesis</i> , 2011 , 32, 1867-71	4.6	10
142	A multi-stage genome-wide association study of bladder cancer identifies multiple susceptibility loci. <i>Nature Genetics</i> , 2010 , 42, 978-84	36.3	408
141	Genetic variants in inflammation-related genes are associated with radiation-induced toxicity following treatment for non-small cell lung cancer. <i>PLoS ONE</i> , 2010 , 5, e12402	3.7	75
140	Replication of lung cancer susceptibility loci at chromosomes 15q25, 5p15, and 6p21: a pooled analysis from the International Lung Cancer Consortium. <i>Journal of the National Cancer Institute</i> , 2010 , 102, 959-71	9.7	153
139	Nicotinic acetylcholine receptor region on chromosome 15q25 and lung cancer risk among African Americans: a case-control study. <i>Journal of the National Cancer Institute</i> , 2010 , 102, 1199-205	9.7	54
138	Genome-wide catalogue of chromosomal aberrations in barrett's esophagus and esophageal adenocarcinoma: a high-density single nucleotide polymorphism array analysis. <i>Cancer Prevention Research</i> , 2010 , 3, 1176-86	3.2	67
137	Genetic variations of the PI3K-AKT-mTOR pathway and clinical outcome in muscle invasive and metastatic bladder cancer patients. <i>Carcinogenesis</i> , 2010 , 31, 1387-91	4.6	52
136	Genetic variations in the sonic hedgehog pathway affect clinical outcomes in non-muscle-invasive bladder cancer. <i>Cancer Prevention Research</i> , 2010 , 3, 1235-45	3.2	42
135	MicroRNA-related genetic variations as predictors for risk of second primary tumor and/or recurrence in patients with early-stage head and neck cancer. <i>Carcinogenesis</i> , 2010 , 31, 2118-23	4.6	45
134	Genetic variations in regulator of G-protein signaling genes as susceptibility loci for second primary tumor/recurrence in head and neck squamous cell carcinoma. <i>Carcinogenesis</i> , 2010 , 31, 1755-61	4.6	20
133	Genetic variations in microRNA-related genes are associated with survival and recurrence in patients with renal cell carcinoma. <i>Carcinogenesis</i> , 2010 , 31, 1805-12	4.6	110
132	Gamma-radiation sensitivity and polymorphisms in RAD51L1 modulate glioma risk. <i>Carcinogenesis</i> , 2010 , 31, 1762-9	4.6	10
131	Genetic variants in MicroRNA biosynthesis pathways and binding sites modify ovarian cancer risk, survival, and treatment response. <i>Cancer Research</i> , 2010 , 70, 9765-76	10.1	105
130	Energy balance, the PI3K-AKT-mTOR pathway genes, and the risk of bladder cancer. <i>Cancer Prevention Research</i> , 2010 , 3, 505-17	3.2	27

129	Genetic variants and risk of lung cancer in never smokers: a genome-wide association study. <i>Lancet Oncology, The</i> , 2010 , 11, 321-30	21.7	190
128	The D2 dopamine receptor gene and nicotine dependence among bladder cancer patients and controls. <i>Behavior Genetics</i> , 2010 , 40, 49-58	3.2	6
127	Genetic variation in MicroRNA genes and risk of oral premalignant lesions. <i>Molecular Carcinogenesis</i> , 2010 , 49, 183-9	5	80
126	Pharmacogenetics of Lung Cancer 2010 , 87-106		
125	Genetic variations in PI3K-AKT-mTOR pathway and bladder cancer risk. <i>Carcinogenesis</i> , 2009 , 30, 2047-52	4.6	70
124	Genetic variations in the PI3K/PTEN/AKT/mTOR pathway are associated with clinical outcomes in esophageal cancer patients treated with chemoradiotherapy. <i>Journal of Clinical Oncology</i> , 2009 , 27, 857-71	2.2	142
123	Polymorphisms in DNA repair genes, smoking, and bladder cancer risk: findings from the international consortium of bladder cancer. <i>Cancer Research</i> , 2009 , 69, 6857-64	10.1	94
122	Constitutive short telomere length of chromosome 17p and 12q but not 11q and 2p is associated with an increased risk for esophageal cancer. <i>Cancer Prevention Research</i> , 2009 , 2, 459-65	3.2	61
121	Genetic susceptibility to esophageal cancer: the role of the nucleotide excision repair pathway. <i>Carcinogenesis</i> , 2009 , 30, 785-92	4.6	72
120	Sensitivity to NNKOAc is associated with renal cancer risk. <i>Carcinogenesis</i> , 2009 , 30, 706-10	4.6	25
119	Dietary intake of vegetables and fruits and the modification effects of GSTM1 and NAT2 genotypes on bladder cancer risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009 , 18, 2090-7	4	38
118	Family history and risk of renal cell carcinoma: results from a case-control study and systematic meta-analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009 , 18, 801-7	4	75
117	Novel susceptibility loci for second primary tumors/recurrence in head and neck cancer patients: large-scale evaluation of genetic variants. <i>Cancer Prevention Research</i> , 2009 , 2, 617-24	3.2	52
116	MicroRNA expression signatures in Barrett's esophagus and esophageal adenocarcinoma. <i>Clinical Cancer Research</i> , 2009 , 15, 5744-52	12.9	110
115	Risk of urinary bladder cancer: a case-control analysis of industry and occupation. <i>BMC Cancer</i> , 2009 , 9, 443	4.8	21
114	Genome-wide profiling of chromosomal alterations in renal cell carcinoma using high-density single nucleotide polymorphism arrays. <i>International Journal of Cancer</i> , 2009 , 125, 2342-8	7.5	66
113	Cyclooxygenase-2 gene polymorphisms reduce the risk of oral premalignant lesions. <i>Cancer</i> , 2009 , 115, 1498-506	6.4	18
112	Genome-wide association studies of bladder cancer risk: a field synopsis of progress and potential applications. <i>Cancer and Metastasis Reviews</i> , 2009 , 28, 269-80	9.6	29

111	Genetic variation in the prostate stem cell antigen gene PSCA confers susceptibility to urinary bladder cancer. <i>Nature Genetics</i> , 2009 , 41, 991-5	36.3	270
110	Pharmacogenomics of platinum-based chemotherapy in NSCLC. <i>Expert Opinion on Drug Metabolism and Toxicology</i> , 2009 , 5, 745-55	5.5	51
109	Genome-wide association studies in bladder cancer: first results and potential relevance. <i>Current Opinion in Urology</i> , 2009 , 19, 540-6	2.8	19
108	Genome-wide association scan of tag SNPs identifies a susceptibility locus for lung cancer at 15q25.1. <i>Nature Genetics</i> , 2008 , 40, 616-22	36.3	1036
107	Common 5p15.33 and 6p21.33 variants influence lung cancer risk. <i>Nature Genetics</i> , 2008 , 40, 1407-9	36.3	465
106	Epidemiology and genetic susceptibility to bladder cancer. <i>BJU International</i> , 2008 , 102, 1207-15	5.6	56
105	Modulation of DNA damage/DNA repair capacity by XPC polymorphisms. <i>DNA Repair</i> , 2008 , 7, 141-8	4.3	64
104	An expanded risk prediction model for lung cancer. <i>Cancer Prevention Research</i> , 2008 , 1, 250-4	3.2	114
103	Genetic polymorphisms in double-strand break DNA repair genes associated with risk of oral premalignant lesions. <i>European Journal of Cancer</i> , 2008 , 44, 1603-11	7.5	24
102	Molecular epidemiology, cancer-related symptoms, and cytokines pathway. <i>Lancet Oncology</i> , 2008 , 9, 777-85	21.7	79
101	BPDE induced lymphocytic chromosome 3p deletions may predict renal cell carcinoma risk. <i>Journal of Urology</i> , 2008 , 179, 2416-21	2.5	15
100	Profiling of genetic variations in inflammation pathway genes in relation to bladder cancer predisposition. <i>Clinical Cancer Research</i> , 2008 , 14, 2236-44	12.9	44
99	Evaluation of genetic variants in microRNA-related genes and risk of bladder cancer. <i>Cancer Research</i> , 2008 , 68, 2530-7	10.1	215
98	Case-control analysis of nucleotide excision repair pathway and the risk of renal cell carcinoma. <i>Carcinogenesis</i> , 2008 , 29, 2112-9	4.6	16
97	Genetic susceptibility to renal cell carcinoma: the role of DNA double-strand break repair pathway. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008 , 17, 2366-73	4	33
96	The CHRNA5-A3 region on chromosome 15q24-25.1 is a risk factor both for nicotine dependence and for lung cancer. <i>Journal of the National Cancer Institute</i> , 2008 , 100, 1552-6	9.7	191
95	Mitochondrial DNA content: its genetic heritability and association with renal cell carcinoma. <i>Journal of the National Cancer Institute</i> , 2008 , 100, 1104-12	9.7	206
94	Familial aggregation of common sequence variants on 15q24-25.1 in lung cancer. <i>Journal of the National Cancer Institute</i> , 2008 , 100, 1326-30	9.7	126

93	Benzo(a)pyrene diol epoxide-induced chromosome 9p21 aberrations are associated with increased risk of bladder cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008 , 17, 2445-50	4	12
92	Smoking-related genomic signatures in non-small cell lung cancer. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2008 , 178, 1164-72	10.2	33
91	Germline genetic variations in drug action pathways predict clinical outcomes in advanced lung cancer treated with platinum-based chemotherapy. <i>Pharmacogenetics and Genomics</i> , 2008 , 18, 955-65	1.9	40
90	Genetic susceptibility to bladder cancer with an emphasis on gene-gene and gene-environmental interactions. <i>Current Opinion in Urology</i> , 2008 , 18, 493-8	2.8	21
89	Plasma vitamins E and A and risk of bladder cancer: a case-control analysis. <i>Cancer Causes and Control</i> , 2008 , 19, 981-92	2.8	26
88	Genetic variants in cell cycle control pathway confer susceptibility to bladder cancer. <i>Cancer</i> , 2008 , 112, 2467-74	6.4	45
87	Genetic variations in cell-cycle pathway and the risk of oral premalignant lesions. <i>Cancer</i> , 2008 , 113, 2488-95	6.4	17
86	Pharmacogenetics in Cancer Chemotherapy 2008 , 113-128		1
85	Strategies to Identify Pharmacogenomic Biomarkers: Candidate Gene, Pathway-Based, and Genome-Wide Approaches 2008 , 353-370		2
84	Differential induction in telomerase activity among bladder cancer patients and controls on gamma-radiation. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007 , 16, 606-9	4	9
83	Dietary isothiocyanates, GSTM1, GSTT1, NAT2 polymorphisms and bladder cancer risk. <i>International Journal of Cancer</i> , 2007 , 120, 2208-13	7.5	62
82	Interplay between mutagen sensitivity and epidemiological factors in modulating lung cancer risk. <i>International Journal of Cancer</i> , 2007 , 120, 2687-95	7.5	22
81	ATM sequence variants associate with susceptibility to non-small cell lung cancer. <i>International Journal of Cancer</i> , 2007 , 121, 2254-9	7.5	27
80	Irradiation-induced telomerase activity and the risk of lung cancer: a pilot case-control study. <i>Cancer</i> , 2007 , 109, 1157-63	6.4	2
79	Risk assessment of renal cell carcinoma using alkaline comet assay. <i>Cancer</i> , 2007 , 110, 282-8	6.4	13
78	Urinary tract diseases and bladder cancer risk: a case-control study. <i>Cancer Causes and Control</i> , 2007 , 18, 839-45	2.8	20
77	Genotypes, haplotypes and diplotypes of XPC and risk of bladder cancer. <i>Carcinogenesis</i> , 2007 , 28, 698-703	4.3	36
76	Mutagen sensitivity and genetic variants in nucleotide excision repair pathway: genotype-phenotype correlation. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007 , 16, 2065-71	4	41

75	Mutagen sensitivity: a genetic predisposition factor for cancer. <i>Cancer Research</i> , 2007 , 67, 3493-5	10.1	64
74	Genetic variants in cell cycle control pathway confer susceptibility to lung cancer. <i>Clinical Cancer Research</i> , 2007 , 13, 5974-81	12.9	57
73	High-order interactions among genetic polymorphisms in nucleotide excision repair pathway genes and smoking in modulating bladder cancer risk. <i>Carcinogenesis</i> , 2007 , 28, 2160-5	4.6	61
72	Projecting individualized probabilities of developing bladder cancer in white individuals. <i>Journal of Clinical Oncology</i> , 2007 , 25, 4974-81	2.2	63
71	High-order interactions among genetic variants in DNA base excision repair pathway genes and smoking in bladder cancer susceptibility. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007 , 16, 84-91	4	81
70	Nucleotide excision repair pathway genes and oral premalignant lesions. <i>Clinical Cancer Research</i> , 2007 , 13, 3753-8	12.9	49
69	Myeloperoxidase G-463A polymorphism and lung cancer: a HuGE genetic susceptibility to environmental carcinogens pooled analysis. <i>Genetics in Medicine</i> , 2007 , 9, 67-73	8.1	42
68	A risk model for prediction of lung cancer. <i>Journal of the National Cancer Institute</i> , 2007 , 99, 715-26	9.7	299
67	Telomere dysfunction in peripheral lymphocytes as a potential predisposition factor for renal cancer. <i>Journal of Urology</i> , 2007 , 178, 1492-6	2.5	66
66	Genetic polymorphism in bladder cancer. <i>Frontiers in Bioscience - Landmark</i> , 2007 , 12, 192-213	2.8	24
65	Deficiency of cell cycle checkpoints and DNA repair system predispose individuals to esophageal cancer. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2006 , 602, 143-50	3.3	8
64	Bladder cancer risk as modified by family history and smoking. <i>Cancer</i> , 2006 , 107, 705-11	6.4	29
63	Matrix metalloproteinase polymorphisms and bladder cancer risk. <i>Cancer Research</i> , 2006 , 66, 11644-8	10.1	61
62	Cyclin D1 gene polymorphism as a risk factor for oral premalignant lesions. <i>Carcinogenesis</i> , 2006 , 27, 2034-7	4.6	41
61	Mutagen sensitivity has high heritability: evidence from a twin study. <i>Cancer Research</i> , 2006 , 66, 5993-6	10.1	68
60	Mutagen sensitivity and neoplastic progression in patients with Barrett's esophagus: a prospective analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006 , 15, 1935-40	4	29
59	Genetic variations in radiation and chemotherapy drug action pathways predict clinical outcomes in esophageal cancer. <i>Journal of Clinical Oncology</i> , 2006 , 24, 3789-98	2.2	153
58	Personal permanent hair dye use is not associated with bladder cancer risk: evidence from a case-control study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006 , 15, 1746-9	4	32

57	Joint effect of mutagen sensitivity and insulin-like growth factors in predicting the risk of developing secondary primary tumors and tumor recurrence in patients with head and neck cancer. <i>Clinical Cancer Research</i> , 2006 , 12, 7194-201	12.9	16
56	Bladder cancer predisposition: a multigenic approach to DNA-repair and cell-cycle-control genes. <i>American Journal of Human Genetics</i> , 2006 , 78, 464-79	11	234
55	Effects of N-acetyl transferase 1 and 2 polymorphisms on bladder cancer risk in Caucasians. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2005 , 581, 97-104	3	83
54	Glutathione peroxidase 1 gene polymorphism and risk of recurrence in patients with superficial bladder cancer. <i>Urology</i> , 2005 , 66, 769-74	1.6	41
53	Mutagen-induced chromatid breakage as a marker of cancer risk. <i>Methods in Molecular Biology</i> , 2005 , 291, 59-67	1.4	7
52	Pharmacogenetics in esophageal cancer. <i>Seminars in Oncology</i> , 2005 , 32, S87-9	5.5	5
51	Predictors of esophageal cancer risk: assessment of susceptibility to DNA damage using comet assay. <i>Genes Chromosomes and Cancer</i> , 2005 , 44, 415-22	5	22
50	Case-control analysis of dietary folate and risk of bladder cancer. <i>Nutrition and Cancer</i> , 2005 , 53, 144-51	2.8	33
49	Opposing effects of emphysema, hay fever, and select genetic variants on lung cancer risk. <i>American Journal of Epidemiology</i> , 2005 , 161, 412-22	3.8	68
48	Nucleotide excision repair gene polymorphisms and recurrence after treatment for superficial bladder cancer. <i>Clinical Cancer Research</i> , 2005 , 11, 1408-15	12.9	83
47	Roles of tumor suppressor and telomere maintenance genes in cancer and aging--an epidemiological study. <i>Carcinogenesis</i> , 2005 , 26, 1741-7	4.6	20
46	Polymorphisms in inflammation genes and bladder cancer: from initiation to recurrence, progression, and survival. <i>Journal of Clinical Oncology</i> , 2005 , 23, 5746-56	2.2	123
45	High mutagen sensitivity in peripheral blood lymphocytes predicts poor overall and disease-specific survival in patients with stage III non-small cell lung cancer treated with radiotherapy and chemotherapy. <i>Clinical Cancer Research</i> , 2005 , 11, 2894-8	12.9	10
44	Cell cycle checkpoints, DNA damage/repair, and lung cancer risk. <i>Cancer Research</i> , 2005 , 65, 349-57	10.1	30
43	Dietary carotenoids and genetic instability modify bladder cancer risk. <i>Journal of Nutrition</i> , 2004 , 134, 3362-9	4.1	31
42	Serum levels of insulin growth factor (IGF-I) and IGF-binding protein predict risk of second primary tumors in patients with head and neck cancer. <i>Clinical Cancer Research</i> , 2004 , 10, 3988-95	12.9	61
41	Genetic susceptibility to tobacco-related cancer. <i>Oncogene</i> , 2004 , 23, 6500-23	9.2	130
40	Focus on bladder cancer. <i>Cancer Cell</i> , 2004 , 6, 111-6	24.3	221

39	Benzo[a]pyrene diol epoxide-induced 9p21 aberrations associated with genetic predisposition to bladder cancer. <i>Genes Chromosomes and Cancer</i> , 2004 , 41, 330-8	5	8
38	Specific chromosome aberrations in peripheral blood lymphocytes are associated with risk of bladder cancer. <i>Genes Chromosomes and Cancer</i> , 2004 , 41, 379-89	5	5
37	Methyl-CpG-binding domain 2: a protective role in bladder carcinoma. <i>Cancer</i> , 2004 , 100, 1853-8	6.4	15
36	An evolutionary perspective on single-nucleotide polymorphism screening in molecular cancer epidemiology. <i>Cancer Research</i> , 2004 , 64, 2251-7	10.1	90
35	XRCC3 genetic polymorphism, smoking, and lung carcinoma risk in minority populations. <i>Cancer</i> , 2003 , 98, 1701-6	6.4	24
34	XPA polymorphism associated with reduced lung cancer risk and a modulating effect on nucleotide excision repair capacity. <i>Carcinogenesis</i> , 2003 , 24, 505-9	4.6	132
33	Sulfotransferase 1A1 (SULT1A1) polymorphism and bladder cancer risk: a case-control study. <i>Cancer Letters</i> , 2003 , 202, 61-9	9.9	50
32	An association between NQO1 genetic polymorphism and risk of bladder cancer. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2003 , 536, 131-7	3	70
31	From genotype to phenotype: correlating XRCC1 polymorphisms with mutagen sensitivity. <i>DNA Repair</i> , 2003 , 2, 901-8	4.3	167
30	Telomere dysfunction: a potential cancer predisposition factor. <i>Journal of the National Cancer Institute</i> , 2003 , 95, 1211-8	9.7	381
29	Genetic instability in bladder cancer assessed by the comet assay. <i>Journal of the National Cancer Institute</i> , 2003 , 95, 540-7	9.7	76
28	Myeloperoxidase promoter region polymorphism and lung cancer risk. <i>Methods in Molecular Medicine</i> , 2003 , 75, 121-33		9
27	Assessment of insulin-like growth factors and mutagen sensitivity as predictors of lung cancer risk. <i>Methods in Molecular Medicine</i> , 2003 , 75, 279-87		
26	Genetic susceptibility to lung cancer: the role of DNA damage and repair. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2003 , 12, 689-98	4	118
25	Serum levels of insulin-like growth factor I and risk of squamous intraepithelial lesions of the cervix. <i>Clinical Cancer Research</i> , 2003 , 9, 3356-61	12.9	27
24	Modification of lung cancer susceptibility by green tea extract as measured by the comet assay. <i>Cancer Detection and Prevention</i> , 2002 , 26, 411-8		24
23	Association between asbestos exposure, cigarette smoking, myeloperoxidase (MPO) genotypes, and lung cancer risk. <i>American Journal of Industrial Medicine</i> , 2002 , 42, 29-37	2.7	40
22	Microsomal epoxide hydrolase polymorphisms and lung cancer risk in non-Hispanic whites. <i>Molecular Carcinogenesis</i> , 2002 , 33, 99-104	5	31

21	A case-control analysis of lymphocytic chromosome 9 aberrations in lung cancer. <i>International Journal of Cancer</i> , 2002 , 102, 536-40	7.5	6
20	BPDE-induced lymphocytic 3p21.3 aberrations may predict head and neck carcinoma risk. <i>Cancer</i> , 2002 , 95, 563-8	6.4	10
19	p53 Genotypes and Haplotypes Associated With Lung Cancer Susceptibility and Ethnicity. <i>Journal of the National Cancer Institute</i> , 2002 , 94, 681-90	9.7	191
18	Sulfotransferase (SULT) 1A1 polymorphism as a predisposition factor for lung cancer: a case-control analysis. <i>Lung Cancer</i> , 2002 , 35, 137-42	5.9	88
17	A myeloperoxidase polymorphism associated with reduced risk of lung cancer. <i>Lung Cancer</i> , 2002 , 37, 35-40	5.9	73
16	Chromosome instability in lymphocytes: a potential indicator of predisposition to oral premalignant lesions. <i>Cancer Research</i> , 2002 , 62, 2813-8	10.1	28
15	Serum insulin-like growth factor (IGF) and IGF-binding protein levels and risk of lung cancer: a case-control study nested in the beta-Carotene and Retinol Efficacy Trial Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2002 , 11, 1413-8	4	31
14	Radiation-induced chromatid breaks as a predictor of breast cancer risk. <i>International Journal of Radiation Oncology Biology Physics</i> , 2001 , 49, 533-7	4	37
13	Analysis of aromatic DNA adducts and 7,8-dihydro-8-oxo-2'-deoxyguanosine in lymphocyte DNA from a case-control study of lung cancer involving minority populations 2000 , 27, 34-46		50
12	Genetic variants of myeloperoxidase and lung cancer risk. <i>Carcinogenesis</i> , 2000 , 21, 1163-1166	4.6	76
11	Ethnic differences in poly(ADP-ribose) polymerase pseudogene genotype distribution and association with lung cancer risk. <i>Carcinogenesis</i> , 1999 , 20, 1465-9	4.6	16
10	Genetic susceptibility to tobacco carcinogenesis. <i>Cancer Investigation</i> , 1999 , 17, 645-59	2.1	48
9	Three measures of mutagen sensitivity in a cancer-free population. <i>Cancer Genetics and Cytogenetics</i> , 1999 , 110, 65-9		16
8	A parallel study of in vitro sensitivity to benzo[a]pyrene diol epoxide and bleomycin in lung carcinoma cases and controls. <i>Cancer</i> , 1998 , 83, 1118-1127	6.4	57
7	Chromosome 5 aberrations and genetic predisposition to lung cancer. <i>International Journal of Cancer</i> , 1998 , 79, 490-3	7.5	21
6	Benzo[a]pyrene diol epoxide and bleomycin sensitivity and susceptibility to cancer of upper aerodigestive tract. <i>Journal of the National Cancer Institute</i> , 1998 , 90, 1393-9	9.7	53
5	Case-control study of the D2 dopamine receptor gene and smoking status in lung cancer patients. <i>Journal of the National Cancer Institute</i> , 1998 , 90, 358-63	9.7	202
4	Lung carcinoma patients with a family history of cancer and lymphocyte primary chromosome 9 aberrations. <i>Cancer</i> , 1997 , 79, 1527-32	6.4	3

- 3 Mutagen sensitivity in humans. A comparison between two nomenclature systems for recording chromatid breaks. *Cancer Genetics and Cytogenetics*, **1996**, 87, 127-32 18
- 2 Mutagen sensitivity as a marker of cancer susceptibility. *Journal of Cellular Biochemistry*, **1996**, 63, 80-84 4.7 13
- 1 Higher lung cancer risk for younger African-Americans with the Pro/Pro p53 genotype. *Carcinogenesis*, **1995**, 16, 2205-8 4.6 124