

# Minimum information about a microarray experiment (MIAME) microarray data

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Citation Report

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1805	Probing the Reproducibility of Leaf Growth and Molecular Phenotypes: A Comparison of Three <i>Arabidopsis</i> Accessions Cultivated in Ten Laboratories Å. <i>Plant Physiology</i> , 2010, 152, 2142-2157.	2.3	137
1806	Genomic Profiling of Tumor Necrosis Factor Alpha (TNF-Î±) Receptor and Interleukin-1 Receptor Knockout Mice Reveals a Link between TNF-Î± Signaling and Increased Severity of 1918 Pandemic Influenza Virus Infection. <i>Journal of Virology</i> , 2010, 84, 12576-12588.	1.5	59
1807	Genetics and genomic approaches to improve grape quality for winemaking. , 2010, , 316-364.		5
1808	Comparative Transcriptome and Secretome Analysis of Wood Decay Fungi <i>Postia placenta</i> and <i>Phanerochaete chrysosporium</i>. <i>Applied and Environmental Microbiology</i> , 2010, 76, 3599-3610.	1.4	237
1809	Evaluation of One- and Two-Color Gene Expression Arrays for Microbial Comparative Genome Hybridization Analyses in Routine Applications. <i>Journal of Clinical Microbiology</i> , 2010, 48, 3105-3110.	1.8	10
1810	Bovine pretransfer endometrium and embryo transcriptome fingerprints as predictors of pregnancy success after embryo transfer. <i>Physiological Genomics</i> , 2010, 42, 201-218.	1.0	94
1811	Mouse Resource Browser—a database of mouse databases. <i>Database: the Journal of Biological Databases and Curation</i> , 2010, 2010, baq010-baq010.	1.4	3
1812	Identification of adrenal genes regulated in a potassium-dependent manner. <i>Journal of Molecular Endocrinology</i> , 2010, 45, 193-206.	1.1	12
1813	Gene Expression Profiling of Mouse Oocytes and Preimplantation Embryos. <i>Methods in Enzymology</i> , 2010, 477, 457-480.	0.4	7
1814	Sources of Bias in Specimens for Research About Molecular Markers for Cancer. <i>Journal of Clinical Oncology</i> , 2010, 28, 698-704.	0.8	187
1815	The 2010 <i>Nucleic Acids Research</i> Database Issue and online Database Collection: a community of data resources. <i>Nucleic Acids Research</i> , 2010, 38, D1-D4.	6.5	86
1816	Gene expression profile in flow-associated pulmonary arterial hypertension with neointimal lesions. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2010, 298, L483-L491.	1.3	39
1817	Standards Affecting the Consistency of Gene Expression Arrays in Clinical Applications. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 1000-1003.	1.1	9
1818	Improving bioscience research reporting: The ARRIVE guidelines for reporting animal research. <i>Journal of Pharmacology and Pharmacotherapeutics</i> , 2010, 1, 94-99.	0.2	2,743
1820	Biomedical Informatics for Cancer Research. , 2010, , .		13
1821	EuroDia: a beta-cell gene expression resource. <i>Database: the Journal of Biological Databases and Curation</i> , 2010, 2010, baq024-baq024.	1.4	9
1822	l-Glutamine as a nitrogen source for <i>Corynebacterium glutamicum</i> : derepression of the AmtR regulon and implications for nitrogen sensing. <i>Microbiology (United Kingdom)</i> , 2010, 156, 3180-3193.	0.7	27



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1824	Automated microscopy for high-content RNAi screening. <i>Journal of Cell Biology</i> , 2010, 188, 453-461.	2.3	135
1825	Members of the LATERAL ORGAN BOUNDARIES DOMAIN Transcription Factor Family Are Involved in the Regulation of Secondary Growth in <i>Populus</i> . <i>Plant Cell</i> , 2010, 22, 3662-3677.	3.1	114
1826	Research Resource: Genome-Wide Profiling of Methylated Promoters in Endometriosis Reveals a Subtelomeric Location of Hypermethylation. <i>Molecular Endocrinology</i> , 2010, 24, 1872-1885.	3.7	90
1827	Decision support based on genomics: integration of data- and knowledge-driven reasoning. <i>International Journal of Biomedical Engineering and Technology</i> , 2010, 3, 287.	0.2	3
1828	Meta-analysis and Network Analysis of Five Ovarian Cancer Gene Expression Dataset. , 2010, , .		2
1829	Molecular remodeling of ion channels, exchangers and pumps in atrial and ventricular myocytes in ischemic cardiomyopathy. <i>Channels</i> , 2010, 4, 101-107.	1.5	27
1830	Two-Way Communication Between Endometrial Stromal Cells and Monocytes. <i>Reproductive Sciences</i> , 2010, 17, 125-136.	1.1	12
1831	Genomic Technologies for Systems Biology. , 2010, , 15-44.		0
1832	Novel Omics Technologies in Nutraceutical and Functional Food Research. , 2010, , 11-22.		2
1833	Early Transcriptional Events During Osteogenic Differentiation of Human Bone Marrow Stromal Cells Induced by Lim Mineralization Protein 3. <i>Gene Expression</i> , 2010, 15, 27-42.	0.5	18
1834	Improving Bioscience Research Reporting: The ARRIVE Guidelines for Reporting Animal Research. <i>PLoS Biology</i> , 2010, 8, e1000412.	2.6	5,621
1835	Gene Expression Atlas at the European Bioinformatics Institute. <i>Nucleic Acids Research</i> , 2010, 38, D690-D698.	6.5	216
1836	Interpretation of Toxicogenomics Data. , 2010, , 663-683.		0
1837	Translating Gene Expression Into Clinical Care: Sarcomas As a Paradigm. <i>Journal of Clinical Oncology</i> , 2010, 28, 1796-1805.	0.8	42
1838	A Primer on Metagenomics. <i>PLoS Computational Biology</i> , 2010, 6, e1000667.	1.5	523
1839	Genomic markers for decision making: what is preventing us from using markers?. <i>Nature Reviews Clinical Oncology</i> , 2010, 7, 90-97.	12.5	13
1840	Robin: An Intuitive Wizard Application for R-Based Expression Microarray Quality Assessment and Analysis. <i>Plant Physiology</i> , 2010, 153, 642-651.	2.3	96

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1842	Best use of experimental data in cancer informatics. <i>Future Oncology</i> , 2010, 6, 1551-1562.	1.1	3
1843	GENOMIC APPROACHES TO ENDOTHELIAL CELL PHENOTYPING. , 2010, , 179-211.		0
1844	Screening for biomarkers of spermatogonia within the human testis: a whole genome approach. <i>Human Reproduction</i> , 2010, 25, 1104-1112.	0.4	95
1845	Gene Expression Profiling of Mouse Embryos with Microarrays. <i>Methods in Enzymology</i> , 2010, 477, 511-541.	0.4	5
1846	Systems biology: opening new avenues in clinical research. <i>Nephrology Dialysis Transplantation</i> , 2010, 25, 1015-1018.	0.4	36
1847	Toxicity Testing in the 21st Century: A Vision and a Strategy. <i>Journal of Toxicology and Environmental Health - Part B: Critical Reviews</i> , 2010, 13, 51-138.	2.9	724
1849	Improved Reporting of Statistical Design and Analysis: Guidelines, Education, and Editorial Policies. <i>Methods in Molecular Biology</i> , 2010, 620, 563-598.	0.4	13
1850	Epigenetic Databases and Computational Methodologies in the Analysis of Epigenetic Datasets. <i>Advances in Genetics</i> , 2010, 71, 259-295.	0.8	4
1852	Biomedical informatics and translational medicine. <i>Journal of Translational Medicine</i> , 2010, 8, 22.	1.8	84
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1854	Inductive Databases and Constraint-Based Data Mining. , 2010, , .		6
1855	GlycoExtractor: A Web-Based Interface for High Throughput Processing of HPLC-Glycan Data. <i>Journal of Proteome Research</i> , 2010, 9, 2037-2041.	1.8	21
1856	An Introduction to Microarray Data Analysis and Visualization. <i>Methods in Enzymology</i> , 2010, 470, 19-50.	0.4	12
1857	Genome-wide transcription profiling of human sepsis: a systematic review. <i>Critical Care</i> , 2010, 14, R237.	2.5	144
1858	High-throughput Gene and Protein Expression Analysis in Pulp Biologic Research: Review. <i>Journal of Endodontics</i> , 2010, 36, 179-189.	1.4	16
1859	Expectations, validity, and reality in gene expression profiling. <i>Journal of Clinical Epidemiology</i> , 2010, 63, 950-959.	2.4	55
1860	Expectations, validity, and reality in omics. <i>Journal of Clinical Epidemiology</i> , 2010, 63, 945-949.	2.4	51

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1863	Gene-expression profiles, tumor microenvironment, and cancer stem cells in breast cancer: Latest advances towards an integrated approach. <i>Cancer Treatment Reviews</i> , 2010, 36, 477-484.	3.4	23
1864	The filamentous fungal gene expression database (FFGED). <i>Fungal Genetics and Biology</i> , 2010, 47, 199-204.	0.9	32
1865	Genomics in multiple sclerosis. <i>Clinical Neurology and Neurosurgery</i> , 2010, 112, 621-624.	0.6	13
1866	A Call to Standardize Teratoma Assays Used to Define Human Pluripotent Cell Lines. <i>Cell Stem Cell</i> , 2010, 6, 412-414.	5.2	121
1867	Differentially expressed genes associated with <i>Staphylococcus aureus</i> mastitis in dairy goats. <i>Veterinary Immunology and Immunopathology</i> , 2010, 135, 208-217.	0.5	31
1868	Genomic and transcriptomic studies in <i>Mycobacterium avium</i> subspecies paratuberculosis. <i>Veterinary Immunology and Immunopathology</i> , 2010, 138, 303-311.	0.5	8
1869	ATX expression and LPA signalling are vital for the development of the nervous system. <i>Developmental Biology</i> , 2010, 339, 451-464.	0.9	133
1870	Gene-expression profiling in pancreatic cancer. <i>Expert Review of Molecular Diagnostics</i> , 2010, 10, 591-601.	1.5	30
1871	Systems Biology Approaches and Tools for Analysis of Interactomes and Multi-target Drugs. <i>Methods in Molecular Biology</i> , 2010, 662, 29-58.	0.4	46
1872	Hepatic gene expression in flounder chronically exposed to multiply polluted estuarine sediment: Absence of classical exposure "biomarker" signals and induction of inflammatory, innate immune and apoptotic pathways. <i>Aquatic Toxicology</i> , 2010, 96, 234-245.	1.9	47
1873	Influence of ovarian stage on transcript profiles in fathead minnow ( <i>Pimephales promelas</i> ) ovary tissue. <i>Aquatic Toxicology</i> , 2010, 98, 354-366.	1.9	40
1874	Temporal patterns in the transcriptomic response of rainbow trout, <i>Oncorhynchus mykiss</i> , to crude oil. <i>Aquatic Toxicology</i> , 2010, 99, 320-329.	1.9	35
1875	Multiple sclerosis etiology: beyond genes and environment. <i>Expert Review of Clinical Immunology</i> , 2010, 6, 481-490.	1.3	23
1876	Gene expression in human sepsis: what have we learned?. <i>Critical Care</i> , 2010, 15, 121.	2.5	11
1877	Bioinformatic and Computational Analysis for Genomic Medicine. , 2010, , 111-130.		0
1878	Reciprocal Communication Between Endometrial Stromal Cells and Macrophages. <i>Reproductive Sciences</i> , 2010, 17, 809-822.	1.1	29
1879	Recommendations for Biomarker Identification and Qualification in Clinical Proteomics. <i>Science Translational Medicine</i> , 2010, 2, 46ps42.	5.8	273

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1881	Statistical Methods in Molecular Biology. Methods in Molecular Biology, 2010, , .	0.4	16
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1883	Biological Databases at DNA Data Bank of Japan in the Era of Next-Generation Sequencing Technologies. Advances in Experimental Medicine and Biology, 2010, 680, 125-135.	0.8	14
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1886	Database resources of the National Center for Biotechnology Information. Nucleic Acids Research, 2010, 38, D5-D16.	6.5	417
1887	DNA Microarrays in Biological Discovery and Patient Care. , 2010, , 73-88.		5
1888	Heart Failure in the Era of Genomic Medicine. , 2010, , 313-325.		0
1889	Analytical Methodology Standards for Metabolomics. , 2010, , 4-8.		0
1890	Screening of genes involved in chromosome segregation during meiosis I: toward the identification of genes responsible for infertility in humans. Journal of Human Genetics, 2010, 55, 293-299.	1.1	20
1891	Proteome Bioinformatics. Methods in Molecular Biology, 2010, , .	0.4	11
1892	Introduction to Marine Genomics. , 2010, , .		6
1893	A standard variation file format for human genome sequences. Genome Biology, 2010, 11, R88.	13.9	79
1894	What is a biomarker? Research investments and lack of clinical integration necessitate a review of biomarker terminology and validation schema. Scandinavian Journal of Clinical and Laboratory Investigation, 2010, 70, 6-14.	0.6	83
1895	Identification of Radiation-Induced Expression Changes in Nonimmortalized Human T Cells. Radiation Research, 2010, 175, 172.	0.7	28
1896	PODD - Towards an Extensible, Domain-Agnostic Scientific Data Management System. , 2010, , .		6
1897	Microarray Technology: A Promising Tool in Nutrigenomics. Critical Reviews in Food Science and Nutrition, 2010, 50, 693-698.	5.4	22
1898	Comprehensive data infrastructure for plant bioinformatics. , 2010, , .		4

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1900	Systematic polymorphism discovery after genome-wide identification of potential susceptibility loci in a hereditary rodent model of human hypertension. <i>Blood Pressure</i> , 2011, 20, 222-231.	0.7	10
1901	Tomato Functional Genomics Database: a comprehensive resource and analysis package for tomato functional genomics. <i>Nucleic Acids Research</i> , 2011, 39, D1156-D1163.	6.5	113
1902	Semantic infrastructure to enable collaboration in ontology development. , 2011, , .		5
1903	Comparative transcriptome analysis of yeast strains carrying <i>slt2</i> , <i>rlm1</i> , and <i>pop2</i> deletions. <i>Genome</i> , 2011, 54, 99-109.	0.9	3
1904	Database resources of the National Center for Biotechnology Information. <i>Nucleic Acids Research</i> , 2011, 39, D38-D51.	6.5	582
1905	Bacterial Subfamily of LuxR Regulators That Respond to Plant Compounds. <i>Applied and Environmental Microbiology</i> , 2011, 77, 4579-4588.	1.4	68
1906	Standards, Databases, and Modeling Tools in Systems Biology. <i>Methods in Molecular Biology</i> , 2011, 696, 413-427.	0.4	20
1907	Hepatic in vitro toxicity assessment of PBDE congeners BDE47, BDE153 and BDE154 in Atlantic salmon ( <i>Salmo salar</i> L.). <i>Aquatic Toxicology</i> , 2011, 105, 246-263.	1.9	31
1908	Analyzing Schizophrenia by DNA Microarrays. <i>Biological Psychiatry</i> , 2011, 69, 157-162.	0.7	58
1909	The SEEK. <i>Methods in Enzymology</i> , 2011, 500, 629-655.	0.4	44
1910	Transcriptome Analysis in Endothelial Progenitor Cell Biology. <i>Antioxidants and Redox Signaling</i> , 2011, 15, 1029-1042.	2.5	26
1912	What Information Should Be Required to Support Clinical "Omics" Publications?. <i>Clinical Chemistry</i> , 2011, 57, 688-690.	1.5	58
1913	Data Standards for Omics Data: The Basis of Data Sharing and Reuse. <i>Methods in Molecular Biology</i> , 2011, 719, 31-69.	0.4	73
1914	Omics Data Management and Annotation. <i>Methods in Molecular Biology</i> , 2011, 719, 71-96.	0.4	12
1915	Insights into Global Mechanisms and Disease by Gene Expression Profiling. <i>Methods in Molecular Biology</i> , 2011, 719, 269-298.	0.4	2
1916	Computational Analysis Workflows for Omics Data Interpretation. <i>Methods in Molecular Biology</i> , 2011, 719, 379-397.	0.4	4
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1920	Dynamic changes in gene expression during human early embryo development: from fundamental aspects to clinical applications. Human Reproduction Update, 2011, 17, 272-290.	5.2	121
1922	The gene expression profile of preclinical autoimmune arthritis and its modulation by a tolerogenic disease-protective antigenic challenge. Arthritis Research and Therapy, 2011, 13, R143.	1.6	6
1923	Regulation of chondrocyte gene expression by osteogenic protein-1. Arthritis Research and Therapy, 2011, 13, R55.	1.6	27
1925	Biological Microarrays. Methods in Molecular Biology, 2011, , .	0.4	2
1927	Standards, Tools, and Databases for the Analysis of Yeast Omics Data. Methods in Molecular Biology, 2011, 759, 345-365.	0.4	2
1928	Methoxychlor affects multiple hormone signaling pathways in the largemouth bass ( <i>Micropterus</i> ) Tj ETQq0 0 0 rgBT JOverlock 10 Tf 50	1.9	26
1929	Effects of a short-term exposure to the fungicide prochloraz on endocrine function and gene expression in female fathead minnows ( <i>Pimephales promelas</i> ). Aquatic Toxicology, 2011, 103, 170-178.	1.9	57
1930	Development of a diagnostic multiplex polymerase chain reaction microarray assay to detect and differentiate <i>Brucella</i> spp.. Diagnostic Microbiology and Infectious Disease, 2011, 71, 341-353.	0.8	11
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1932	Expression of HOXB genes is significantly different in acute myeloid leukemia with a partial tandem duplication of MLL vs. a MLL translocation: a cross-laboratory study. Cancer Genetics, 2011, 204, 252-259.	0.2	12
1933	Counting the cost of carbon in bioenergy systems: sources of variation and hidden pitfalls when comparing life cycle assessments. Biofuels, 2011, 2, 693-707.	1.4	17
1934	Minimum information about a bioactive entity (MIABE). Nature Reviews Drug Discovery, 2011, 10, 661-669.	21.5	80
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1936	Data Model Standardization for Synthetic Biomolecular Circuits and Systems. , 2011, , 281-293.		1
1937	Whole blood gene expression analyses in patients with single versus recurrent venous thromboembolism. Thrombosis Research, 2011, 128, 536-540.	0.8	39
1938	Analyses of placental gene expression in pregnancy-related hypertensive disorders. Taiwanese Journal of Obstetrics and Gynecology, 2011, 50, 283-291.	0.5	14
1939	Omics Technologies, Data and Bioinformatics Principles. Methods in Molecular Biology, 2011, 719, 3-30.	0.4	92

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1942	Microarray analysis in gynaecology and its findings: a systematic review. <i>Reproductive BioMedicine Online</i> , 2011, 22, 569-582.	1.1	0
1943	Advancing transcriptome platforms. <i>Nature Precedings</i> , 2011, , .	0.1	0
1944	Expression profiling in pediatric acute leukemias. , 0, , 202-216.		0
1946	Immuno-Modulatory Effects of Phytomedicines Evaluated Using Omics Approaches. , 0, , .		0
1947	Sources of diversity in T cell epitope discovery. <i>Frontiers in Bioscience - Landmark</i> , 2011, 16, 3014.	3.0	11
1948	An Integrated Framework to Model Cellular Phenotype as a Component of Biochemical Networks. <i>Advances in Bioinformatics</i> , 2011, 2011, 1-14.	5.7	0
1949	Tracheal Occlusion Conditioning in Conscious Rats Modulates Gene Expression Profile of Medial Thalamus. <i>Frontiers in Physiology</i> , 2011, 2, 24.	1.3	11
1950	Plant Cell Biology: With Grand Challenges Come Great Possibilities. <i>Frontiers in Plant Science</i> , 2011, 2, 3.	1.7	3
1951	Data Ontology and an Information System Realization for Web-Based Management of Image Measurements. <i>Frontiers in Neuroinformatics</i> , 2011, 5, 25.	1.3	4
1952	Assessing Epigenetic Information. , 2011, , 173-181.		5
1953	Uropathogenic <i>E. coli</i> Induce Different Immune Response in Testicular and Peritoneal Macrophages: Implications for Testicular Immune Privilege. <i>PLoS ONE</i> , 2011, 6, e28452.	1.1	68
1954	Who Shares? Who Doesn't? Factors Associated with Openly Archiving Raw Research Data. <i>PLoS ONE</i> , 2011, 6, e18657.	1.1	161
1955	COLOMBOS: Access Port for Cross-Platform Bacterial Expression Compendia. <i>PLoS ONE</i> , 2011, 6, e20938.	1.1	25
1956	The Two-Component Signal Transduction System CopRS of <i>Corynebacterium glutamicum</i> Is Required for Adaptation to Copper-Excess Stress. <i>PLoS ONE</i> , 2011, 6, e22143.	1.1	34
1957	Rhythmic Diel Pattern of Gene Expression in Juvenile Maize Leaf. <i>PLoS ONE</i> , 2011, 6, e23628.	1.1	12
1958	DBE2 – Management of experimental data for the VANTED system. <i>Journal of Integrative Bioinformatics</i> , 2011, 8, 85-94.	1.0	0

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1960	The modENCODE Data Coordination Center: lessons in harvesting comprehensive experimental details. Database: the Journal of Biological Databases and Curation, 2011, 2011, bar023.	1.4	32
1961	Metadata Management and Semantics in Microarray Repositories. <i>Balkan Journal of Medical Genetics</i> , 2011, 14, 49-64.	0.5	0
1962	Gene expression signatures differ with extent of atherosclerosis in monkey iliac artery. <i>Menopause</i> , 2011, 18, 1087-1095.	0.8	16
1963	Digital Slide Repositories for Publications. <i>American Journal of Surgical Pathology</i> , 2011, 35, 783-786.	2.1	25
1964	Minimum Information about a Genotyping Experiment (MIGEN). <i>Standards in Genomic Sciences</i> , 2011, 5, 224-229.	1.5	3
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1966	FBXO31 determines poor prognosis in esophageal squamous cell carcinoma. <i>International Journal of Oncology</i> , 2011, 39, 155-9.	1.4	14
1967	Basal-like molecular subtype and HER4 up-regulation and response to neoadjuvant chemotherapy in breast cancer. <i>Oncology Reports</i> , 2011, 26, 1037-45.	1.2	31
1968	Evidence that ACN1 (acetate non-utilizing 1) prevents carbon leakage from peroxisomes during lipid mobilization in <i>Arabidopsis</i> seedlings. <i>Biochemical Journal</i> , 2011, 437, 505-513.	1.7	17
1970	Examples of ecological data synthesis driven by rich metadata, and practical guidelines to use the Ecological Metadata Language specification to this end. <i>International Journal of Metadata, Semantics and Ontologies</i> , 2011, 6, 46.	0.2	6
1971	Review On Feature Selection Techniques And The Impact Of Svm For Cancer Classification Using Gene Expression Profile. <i>International Journal of Computer Science &amp; Engineering Survey</i> , 2011, 2, 16-27.	0.2	40
1972	Genomic expression profiles of blood and placenta reveal significant immune-related pathways and categories in Chinese women with gestational diabetes mellitus. <i>Diabetic Medicine</i> , 2011, 28, 237-246.	1.2	45
1973	Metabolomic approaches reveal that cell wall modifications play a major role in ethylene-mediated resistance against <i>Botrytis cinerea</i> . <i>Plant Journal</i> , 2011, 67, 852-868.	2.8	77
1974	Protective pathways against colitis mediated by appendicitis and appendectomy. <i>Clinical and Experimental Immunology</i> , 2011, 165, 393-400.	1.1	15
1975	Ecological genetics and genomics of plant defences: evidence and approaches. <i>Functional Ecology</i> , 2011, 25, 312-324.	1.7	54
1976	Channeling the data deluge. <i>Nature Methods</i> , 2011, 8, 463-465.	9.0	13
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1979	Transcriptional responses of host peripheral blood cells to tuberculosis infection. <i>Tuberculosis</i> , 2011, 91, 390-399.	0.8	64
1980	Expression of the selenoprotein S (SELS) gene in subcutaneous adipose tissue and SELS genotype are associated with metabolic risk factors. <i>Metabolism: Clinical and Experimental</i> , 2011, 60, 114-120.	1.5	62
1981	Cardiac cell modelling: Observations from the heart of the cardiac physiome project. <i>Progress in Biophysics and Molecular Biology</i> , 2011, 104, 2-21.	1.4	139
1982	Minimum Information about a Cardiac Electrophysiology Experiment (MICEE): Standardised reporting for model reproducibility, interoperability, and data sharing. <i>Progress in Biophysics and Molecular Biology</i> , 2011, 107, 4-10.	1.4	75
1983	Apigenin inhibits hepatoma cell growth through alteration of gene expression patterns. <i>Phytomedicine</i> , 2011, 18, 366-373.	2.3	63
1984	Increased endoplasmic reticulum stress in decidual tissue from pregnancies complicated by fetal growth restriction with and without pre-eclampsia. <i>Placenta</i> , 2011, 32, 823-829.	0.7	82
1985	Amplification-free detection of grapevine viruses using an oligonucleotide microarray. <i>Journal of Virological Methods</i> , 2011, 178, 1-15.	1.0	15
1986	GRISOM Platform: Enabling Distributed Processing and Management of Biological Data Through Fusion of Grid and Web Technologies. <i>IEEE Transactions on Information Technology in Biomedicine</i> , 2011, 15, 83-92.	3.6	3
1987	Estrogens and selective estrogen receptor modulators regulate gene and protein expression in the mesenteric arteries. <i>Vascular Pharmacology</i> , 2011, 55, 42-49.	1.0	8
1988	Comparison of gene expression patterns between avian and human ovarian cancers. <i>Gynecologic Oncology</i> , 2011, 120, 256-264.	0.6	18
1989	Current and future use of genomics data in toxicology: Opportunities and challenges for regulatory applications. <i>Regulatory Toxicology and Pharmacology</i> , 2011, 61, 141-153.	1.3	33
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