List of Publications by Year in descending order

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		70961	35952
162	10,317	41	97
papers	citations	h-index	g-index
173 all docs	173 docs citations	173 times ranked	15938 citing authors

#	Article	IF	CITATIONS
1	A new class of membrane-bound chemokine with a CX3C motif. Nature, 1997, 385, 640-644.	13.7	1,855
2	A family of human receptors structurally related to Drosophila Toll. Proceedings of the National Academy of Sciences of the United States of America, 1998, 95, 588-593.	3.3	1,574
3	A gp130–Src–YAP module links inflammation to epithelial regeneration. Nature, 2015, 519, 57-62.	13.7	528
4	Hepatocyte Necrosis Induced by Oxidative Stress and IL-1α Release Mediate Carcinogen-Induced Compensatory Proliferation and Liver Tumorigenesis. Cancer Cell, 2008, 14, 156-165.	7.7	441
5	Fibroblast-specific protein 1 identifies an inflammatory subpopulation of macrophages in the liver. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 308-313.	3.3	300
6	A Mouse Macrophage Lipidome. Journal of Biological Chemistry, 2010, 285, 39976-39985.	1.6	260
7	Analysis of 94 Candidate Genes and 12 Endophenotypes for Schizophrenia From the Consortium on the Genetics of Schizophrenia. American Journal of Psychiatry, 2011, 168, 930-946.	4.0	241
8	Human cardiac organoids for the modelling of myocardial infarction and drug cardiotoxicity. Nature Biomedical Engineering, 2020, 4, 446-462.	11.6	232
9	Cooperative NCoR/SMRT interactions establish a corepressor-based strategy for integration of inflammatory and anti-inflammatory signaling pathways. Genes and Development, 2009, 23, 681-693.	2.7	215
10	CD38-NAD+Axis Regulates Immunotherapeutic Anti-Tumor T Cell Response. Cell Metabolism, 2018, 27, 85-100.e8.	7.2	197
11	Microarray platforms – comparisons and contrasts. Pharmacogenomics, 2004, 5, 487-502.	0.6	175
12	High-Throughput Isolation of Caenorhabditis elegans Deletion Mutants. Genome Research, 1999, 9, 859-867.	2.4	165
13	Comprehensive Multi-omics Analysis Reveals Mitochondrial Stress as a Central Biological Hub for Spaceflight Impact. Cell, 2020, 183, 1185-1201.e20.	13.5	161
14	Gene expression modulation is associated with gene amplification, supernumerary chromosomes and chromosome loss in antimony-resistant Leishmania infantum. Nucleic Acids Research, 2009, 37, 1387-1399.	6.5	153
15	Mechanisms Establishing TLR4-Responsive Activation States of Inflammatory Response Genes. PLoS Genetics, 2011, 7, e1002401.	1.5	146
16	Microplastics in the marine environment: A review of their sources, distribution processes, uptake and exchange in ecosystems. Case Studies in Chemical and Environmental Engineering, 2020, 2, 100010.	2.9	136
17	Nucleosome landscape and control of transcription in the human malaria parasite. Genome Research, 2010, 20, 228-238.	2.4	126
18	Time- and Oil-Dependent Transcriptomic and Physiological Responses to <i>Deepwater Horizon</i> Oil in Mahi-Mahi (<i>Coryphaena hippurus</i>) Embryos and Larvae. Environmental Science & Technology, 2016, 50, 7842-7851.	4.6	123

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19	The Long Non-Coding HOTAIR Is Modulated by Cyclic Stretch and WNT/β-CATENIN in Human Aortic Valve Cells and Is a Novel Repressor of Calcification Genes. PLoS ONE, 2014, 9, e96577.	1.1	101
20	Lis1 regulates asymmetric division in hematopoietic stem cells and in leukemia. Nature Genetics, 2014, 46, 245-252.	9.4	97
21	Larval Red Drum (<i>Sciaenops ocellatus</i>) Sublethal Exposure to Weathered Deepwater Horizon Crude Oil: Developmental and Transcriptomic Consequences. Environmental Science & Technology, 2017, 51, 10162-10172.	4.6	91
22	Molecular characterization and modular analysis of human MyD88. Oncogene, 1996, 13, 2467-75.	2.6	90
23	Protein microarrays: challenges and promises. Pharmacogenomics, 2002, 3, 527-536.	0.6	80
24	Disruption of the Ugt1 Locus in Mice Resembles Human Crigler-Najjar Type I Disease. Journal of Biological Chemistry, 2008, 283, 7901-7911.	1.6	77
25	MAPK Reliance via Acquired CDK4/6 Inhibitor Resistance in Cancer. Clinical Cancer Research, 2018, 24, 4201-4214.	3.2	77
26	Primary cilia defects causing mitral valve prolapse. Science Translational Medicine, 2019, 11, .	5.8	76
27	β1 integrin is a crucial regulator of pancreatic β-cell expansion. Development (Cambridge), 2013, 140, 3360-3372.	1.2	75
28	Probiotics Can Induce Follicle Maturational Competence: The Danio rerioCase1. Biology of Reproduction, 2012, 86, 65.	1.2	71
29	Cloning and Characterization of a New Type of Mouse Chemokine. Genomics, 1998, 47, 163-170.	1.3	70
30	Systems analysis of the prostate transcriptome in African–American men compared with European–American men. Pharmacogenomics, 2016, 17, 1129-1143.	0.6	66
31	The stretch responsive microRNA miRâ€148aâ€3p is a novel repressor of <i>IKBKB</i> , NFâ€ĤB signaling, and inflammatory gene expression in human aortic valve cells. FASEB Journal, 2015, 29, 1859-1868.	0.2	65
32	CRTH2 antagonism significantly ameliorates airway hyperreactivity and downregulates inflammation-induced genes in a mouse model of airway inflammation. American Journal of Physiology - Lung Cellular and Molecular Physiology, 2008, 295, L767-L779.	1.3	60
33	A small molecule CRTH2 antagonist inhibits FITC-induced allergic cutaneous inflammation. International Immunology, 2009, 21, 81-93.	1.8	58
34	Tetraspanin 3 Is Required for the Development and Propagation of Acute Myelogenous Leukemia. Cell Stem Cell, 2015, 17, 152-164.	5.2	58
35	Cyclic stretch of embryonic cardiomyocytes increases proliferation, growth, and expression while repressing Tgf-1² signaling. Journal of Molecular and Cellular Cardiology, 2015, 79, 133-144.	0.9	56
36	Variation of the genetic expression pattern after exposure to estradiol-17β and 4-nonylphenol in male zebrafish (Danio rerio). General and Comparative Endocrinology, 2008, 158, 138-144.	0.8	55

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37	Genomics pipelines and data integration: challenges and opportunities in the research setting. Expert Review of Molecular Diagnostics, 2017, 17, 225-237.	1.5	54
38	Analysis of Endocrine Disruption in Southern California Coastal Fish Using an Aquatic Multispecies Microarray. Environmental Health Perspectives, 2009, 117, 223-230.	2.8	52
39	Systems Analysis of the Liver Transcriptome in Adult Male Zebrafish Exposed to the Plasticizer (2-Ethylhexyl) Phthalate (DEHP). Scientific Reports, 2018, 8, 2118.	1.6	48
40	Multi-omics analysis of multiple missions to space reveal a theme of lipid dysregulation in mouse liver. Scientific Reports, 2019, 9, 19195.	1.6	46
41	KDEL-Retained Antigen in B Lymphocytes Induces a Proinflammatory Response: A Possible Role for Endoplasmic Reticulum Stress in Adaptive T Cell Immunity. Journal of Immunology, 2008, 181, 256-264.	0.4	43
42	Genetic Structure and Chromosomal Mapping of MyD88. Genomics, 1997, 45, 332-339.	1.3	42
43	The mouse Wnt-10B gene isolated from helper T cells is widely expressed and a possible oncogene in BR6 mouse mammary tumorigenesis. Gene, 1996, 172, 199-205.	1.0	41
44	Role of cannabinoidergic mechanisms in ethanol self-administration and ethanol seeking in rat adult offspring following perinatal exposure to Δ9-tetrahydrocannabinol. Toxicology and Applied Pharmacology, 2007, 223, 73-85.	1.3	41
45	Amplification of WHSC1L1 regulates expression and estrogenâ€independent activation of ERα in SUMâ€44 breast cancer cells and is associated with ERα overâ€expression in breast cancer. Molecular Oncology, 2016, 10, 850-865.	2.1	41
46	Prominence of IL6, IGF, TLR, and Bioenergetics Pathway Perturbation in Lung Tissues of Scleroderma Patients With Pulmonary Fibrosis. Frontiers in Immunology, 2020, 11, 383.	2.2	40
47	Endothelium-Derived Netrin-4 Supports Pancreatic Epithelial Cell Adhesion and Differentiation through Integrins $\hat{1}\pm2\hat{1}^21$ and $\hat{1}\pm3\hat{1}^21$. PLoS ONE, 2011, 6, e22750.	1.1	39
48	Serine Proteolytic Pathway Activation Reveals an Expanded Ensemble of Wound Response Genes in Drosophila. PLoS ONE, 2013, 8, e61773.	1.1	39
49	Molecular staging of marine medaka: A model organism for marine ecotoxicity study. Marine Pollution Bulletin, 2011, 63, 309-317.	2.3	38
50	Transcriptional analysis of endocrine disruption using zebrafish and massively parallel sequencing. Journal of Molecular Endocrinology, 2014, 52, R241-R256.	1.1	38
51	Spaceflight influences gene expression, photoreceptor integrity, and oxidative stress-related damage in the murine retina. Scientific Reports, 2019, 9, 13304.	1.6	38
52	A developmental hepatotoxicity study of dietary bisphenol A in Sparus aurata juveniles. Comparative Biochemistry and Physiology Part - C: Toxicology and Pharmacology, 2014, 166, 1-13.	1.3	37
53	Therapeutic target discovery usingCaenorhabditis elegans. Pharmacogenomics, 2000, 1, 203-217.	0.6	36
54	Antagonism of CRTH2 ameliorates chronic epicutaneous sensitization-induced inflammation by multiple mechanisms. International Immunology, 2009, 21, 1-17.	1.8	34

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55	Teleost fish (Solea solea): A novel model for ecotoxicological assay of contaminated sediments. Aquatic Toxicology, 2012, 109, 133-142.	1.9	34
56	Environmental perfluorooctane sulfonate exposure drives T cell activation in bottlenose dolphins. Journal of Applied Toxicology, 2017, 37, 1108-1116.	1.4	34
57	Novel transcriptome assembly and comparative toxicity pathway analysis in mahi-mahi (Coryphaena) Tj ETQq1	1 0.78431 1.6	4 rgBT /Overld
58	SLC36A1-mTORC1 signaling drives acquired resistance to CDK4/6 inhibitors. Science Advances, 2019, 5, eaax6352.	4.7	31
59	A Post-Developmental Genetic Screen for Zebrafish Models of Inherited Liver Disease. PLoS ONE, 2015, 10, e0125980.	1.1	30
60	Repression of caspase-3 and RNA-binding protein HuR cleavage by cyclooxygenase-2 promotes drug resistance in oral squamous cell carcinoma. Oncogene, 2017, 36, 3137-3148.	2.6	29
61	The Plasticizer Bisphenol A Perturbs the Hepatic Epigenome: A Systems Level Analysis of the miRNome. Genes, 2017, 8, 269.	1.0	28
62	Thioredoxin-1 improves the immunometabolic phenotype of antitumor T cells. Journal of Biological Chemistry, 2019, 294, 9198-9212.	1.6	28
63	Presentation of Telomerase Reverse Transcriptase, a Self-Tumor Antigen, is Down-regulated by Histone Deacetylase Inhibition. Cancer Research, 2008, 68, 8085-8093.	0.4	27
64	Molecular Analysis of Endocrine Disruption in Hornyhead Turbot at Wastewater Outfalls in Southern California Using a Second Generation Multi-Species Microarray. PLoS ONE, 2013, 8, e75553.	1.1	27
65	A Degenerative Retinal Process in HIV-Associated Non-Infectious Retinopathy. PLoS ONE, 2013, 8, e74712.	1.1	26
66	Switching-On Survival and Repair Response Programs in Islet Transplants by Bone Marrow–Derived Vasculogenic Cells. Diabetes, 2008, 57, 2402-2412.	0.3	25
67	miRmapper: A Tool for Interpretation of miRNA–mRNA Interaction Networks. Genes, 2018, 9, 458.	1.0	25
68	Microplastic toxicity: A review of the role of marine sentinel species in assessing the environmental and public health impacts. Case Studies in Chemical and Environmental Engineering, 2021, 3, 100073.	2.9	25
69	Acute stress enhances the expression of neuroprotection- and neurogenesis-associated genes in the hippocampus of a mouse restraint model. Oncotarget, 2016, 7, 8455-8465.	0.8	24
70	Gene expression and immunohistochemical analyses identify SOX2 as major risk factor for overall survival and relapse in Ewing sarcoma patients. EBioMedicine, 2019, 47, 156-162.	2.7	23
71	Differential Gene Expression in Liver, Gill, and Olfactory Rosettes of Coho Salmon (Oncorhynchus) Tj ETQq1 1 0	.784314 rg	gBT_/Overlock 22
72	A Novel CLCN5 Mutation Associated WithÂFocal Segmental Glomerulosclerosis andÂPodocyte Injury. Kidney International Reports, 2018, 3, 1443-1453.	0.4	22

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73	Insulin receptor substrate 1 is a substrate of the Pim protein kinases. Oncotarget, 2016, 7, 20152-20165.	0.8	22
74	Microarray Technologies – An Overview. Pharmacogenomics, 2002, 3, 293-297.	0.6	21
75	Microarray technology – an intellectual property retrospective. Pharmacogenomics, 2003, 4, 623-632.	0.6	21
76	Biomarkers of hippocampal gene expression in a mouse restraint chronic stress model. Pharmacogenomics, 2015, 16, 471-482.	0.6	21
77	Microarrays - The Challenge of Preparing Brain Tissue Samples. Addiction Biology, 2005, 10, 5-13.	1.4	20
78	Biological effects of marine contaminated sediments on Sparus aurata juveniles. Aquatic Toxicology, 2011, 104, 308-316.	1.9	20
79	NASA GeneLab RNA-seq consensus pipeline: Standardized processing of short-read RNA-seq data. IScience, 2021, 24, 102361.	1.9	20
80	Microarray Technologies 2003 - An Overview. Pharmacogenomics, 2003, 4, 251-256.	0.6	19
81	Frequency of mitochondrial 12S ribosomal RNA variants in an adult cystic fibrosis population. Pharmacogenetics and Genomics, 2008, 18, 1095-1102.	0.7	19
82	Interferon Regulatory Factorâ€5 in Resident Macrophage Promotes Polycystic Kidney Disease. Kidney360, 2020, 1, 179-190.	0.9	19
83	Nutritional challenges and countermeasures for space travel. Nutrition Bulletin, 2020, 45, 98-105.	0.8	19
84	Antifibrotic factor KLF4 is repressed by the miR-10/TFAP2A/TBX5 axis in dermal fibroblasts: insights from twins discordant for systemic sclerosis. Annals of the Rheumatic Diseases, 2022, 81, 268-277.	0.5	19
85	A Human Skin Model Recapitulates Systemic Sclerosis Dermal Fibrosis and Identifies COL22A1 as a TGFÎ ² Early Response Gene that Mediates Fibroblast to Myofibroblast Transition. Genes, 2019, 10, 75.	1.0	18
86	NASA GeneLab Platform Utilized for Biological Response to Space Radiation in Animal Models. Cancers, 2020, 12, 381.	1.7	18
87	Genomic and phenotypic response of hornyhead turbot exposed to municipal wastewater effluents. Aquatic Toxicology, 2013, 140-141, 174-184.	1.9	17
88	Next-generation antibody discovery platforms. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 18245-18246.	3.3	16
89	Spatial N-glycomics of the human aortic valve in development and pediatric endstage congenital aortic valve stenosis. Journal of Molecular and Cellular Cardiology, 2021, 154, 6-20.	0.9	16
90	Applications of Microarrays and Biochips in Pharmacogenomics. Methods in Molecular Biology, 2008, 448, 21-30.	0.4	16

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91	A CCR2+ myeloid cell niche required for pancreatic \hat{I}^2 cell growth. JCI Insight, 2017, 2, .	2.3	16
92	Advances in pharmacogenomics technologies. Pharmacogenomics, 2010, 11, 481-485.	0.6	15
93	Microarrays Technologies 2006: an overview. Pharmacogenomics, 2006, 7, 1153-1158.	0.6	14
94	SUMO Modification of the RNA-Binding Protein La Regulates Cell Proliferation and STAT3 Protein Stability. Molecular and Cellular Biology, 2018, 38, .	1.1	14
95	Developmental and extracellular matrix-remodeling processes in rosiglitazone-exposed neonatal rat cardiomyocytes. Pharmacogenomics, 2014, 15, 759-774.	0.6	13
96	A de novo transcriptome assembly approach elucidates the dynamics of ovarian maturation in the swordfish (Xiphias gladius). Scientific Reports, 2019, 9, 7375.	1.6	12
97	Mammalian and Invertebrate Models as Complementary Tools for Gaining Mechanistic Insight on Muscle Responses to Spaceflight. International Journal of Molecular Sciences, 2021, 22, 9470.	1.8	12
98	Differential DNA Methylation Landscape in Skin Fibroblasts from African Americans with Systemic Sclerosis. Genes, 2021, 12, 129.	1.0	12
99	A Primer on the Current State of Microarray Technologies. Methods in Molecular Biology, 2012, 802, 3-17.	0.4	12
100	Genome-Wide Analysis of Low Dose Bisphenol-A (BPA) Exposure in Human Prostate Cells. Current Genomics, 2019, 20, 260-274.	0.7	12
101	CHARACTERIZATION OF CYTOKINE GENE EXPRESSION ASSOCIATED WITH NONINFECTIOUS HUMAN IMMUNODEFICIENCY VIRUS RETINOPATHY IN HUMAN AUTOPSY EYES. Retina, 2010, 30, 952-957.	1.0	11
102	Clinical Application of a Modular Genomics Technique in Systemic Lupus Erythematosus: Progress towards Precision Medicine. International Journal of Genomics, 2016, 2016, 1-7.	0.8	11
103	Systems analysis of the liver transcriptome in adult male zebrafish exposed to the non-ionic surfactant nonylphenol. General and Comparative Endocrinology, 2019, 271, 1-14.	0.8	11
104	Revamping Space-omics in Europe. Cell Systems, 2020, 11, 555-556.	2.9	11
105	Developmental transcriptomic analyses for mechanistic insights into critical pathways involved in embryogenesis of pelagic mahi-mahi (Coryphaena hippurus). PLoS ONE, 2017, 12, e0180454.	1.1	10
106	Proteomic biomarkers of cognitive impairment in obstructive sleep apnea syndrome. Sleep and Breathing, 2019, 23, 251-257.	0.9	10
107	Development of a microarray assay that measures hybridization stoichiometry in moles. BioTechniques, 2004, 36, 464-470.	0.8	9
108	Application of a targeted endocrine q-PCR panel to monitor the effects of pollution in southern California flatfish. Endocrine Disruptors (Austin, Tex), 2014, 2, e969598.	1.1	8

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109	Molecular Profiling of RNA Tumors Using High-Throughput RNA Sequencing: From Raw Data to Systems Level Analyses. Methods in Molecular Biology, 2019, 1908, 185-204.	0.4	8
110	Transcriptomic analysis of short-term 17α-ethynylestradiol exposure in two Californian sentinel fish species sardine (Sardinops sagax) and mackerel (Scomber japonicus). Environmental Pollution, 2019, 244, 926-937.	3.7	8
111	Induced Torpor as a Countermeasure for Low Dose Radiation Exposure in a Zebrafish Model. Cells, 2021, 10, 906.	1.8	8
112	Nucleobase-containing compounds evoke behavioural, olfactory, and transcriptional responses in model fishes. Facets, 2018, 3, 79-102.	1.1	8
113	Development and application of a microarray meter tool to optimize microarray experiments. BMC Research Notes, 2008, 1, 45.	0.6	7
114	An Analytic Approach Using Candidate Gene Selection and Logic Forest to Identify Gene by Environment Interactions (G × E) for Systemic Lupus Erythematosus in African Americans. Genes, 2018, 9, 496.	1.0	7
115	A high prevalence of biallelic <i>RPE65</i> mutations in Costa Rican children with Leber congenital amaurosis and early-onset retinal dystrophy. Ophthalmic Genetics, 2019, 40, 110-117.	0.5	7
116	An Introduction to Systems Analytics and Integration of Big Omics Data. Genes, 2020, 11, 245.	1.0	7
117	A Systems Approach to Interrogate Gene Expression Patterns in African American Men Presenting with Clinically Localized Prostate Cancer. Cancers, 2021, 13, 5143.	1.7	7
118	miRNA and IncRNA Expression Networks Modulate Cell Cycle and DNA Repair Inhibition in Senescent Prostate Cells. Genes, 2022, 13, 208.	1.0	7
119	Ultra-high-throughput sequencing, microarray-based genomic selection and pharmacogenomics. Pharmacogenomics, 2008, 9, 5-9.	0.6	6
120	Evaluation of reproductive endocrine status in hornyhead turbot sampled from Southern California's urbanized coastal environments. Environmental Toxicology and Chemistry, 2012, 31, 2689-2700.	2.2	6
121	Effects of age on growth in Atlantic bluefin tuna (Thunnus thynnus). General and Comparative Endocrinology, 2018, 265, 64-70.	0.8	6
122	Long non-coding RNAs and their potential impact on diagnosis, prognosis, and therapy in prostate cancer: racial, ethnic, and geographical considerations. Expert Review of Molecular Diagnostics, 2021, 21, 1257-1271.	1.5	6
123	Contemporary Approaches to the Discovery and Development of Broad-Spectrum Natural Product Prototypes for the Control of Coronaviruses. Journal of Natural Products, 2021, 84, 3001-3007.	1.5	6
124	Space omics research in Europe: Contributions, geographical distribution and ESA member state funding schemes. IScience, 2022, 25, 103920.	1.9	6
125	Differential transferrin gene expression in Atlantic salmon (Salmo salar L.) freshwater parr and seawater smolts. Journal of Applied Ichthyology, 1996, 12, 43-47.	0.3	5
126	Biological responses of marine flatfish exposed to municipal wastewater effluent. Environmental Toxicology and Chemistry, 2014, 33, 583-591.	2.2	5

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127	Interplay Between MicroRNAs and Targeted Genes in Cellular Homeostasis of Adult Zebrafish (Danio) Tj ETQq1	1 0.784314	4 rgBT /Over
128	GAIL: An interactive webserver for inference and dynamic visualization of gene-gene associations based on gene ontology guided mining of biomedical literature. PLoS ONE, 2019, 14, e0219195.	1.1	5
129	Semi-supervised identification of cancer subgroups using survival outcomes and overlapping grouping information. Statistical Methods in Medical Research, 2019, 28, 2137-2149.	0.7	5
130	Biochip platforms as functional genomics tools for drug discovery. Current Opinion in Drug Discovery & Development, 2005, 8, 347-54.	1.9	5
131	Microarray technology – advances, applications, future prospects. Pharmacogenomics, 2007, 8, 1639-1642.	0.6	4
132	Integrated Genomic and Bioinformatics Approaches to Identify Molecular Links between Endocrine Disruptors and Adverse Outcomes. International Journal of Environmental Research and Public Health, 2022, 19, 574.	1.2	4
133	GPA-MDS: A Visualization Approach to Investigate Genetic Architecture among Phenotypes Using GWAS Results. International Journal of Genomics, 2016, 2016, 1-6.	0.8	3
134	ShinyGPA: An interactive visualization toolkit for investigating pleiotropic architecture using GWAS datasets. PLoS ONE, 2018, 13, e0190949.	1.1	3
135	Application of Ultra-High Throughput Sequencing and Microarray Technologies in Pharmacogenomics Testing. , 2012, , 143-159.		2
136	The Next Generation of Automated Microarray Platforms for a Multiplexed CYP2D6 Assay. Drug Discovery Series, 2006, , 97-108.	0.1	2
137	Isolation of Atlantic salmon (Salmo salar L.) cytochrome c oxidase subunit II gene (coxII). Journal of Applied Ichthyology, 1994, 10, 64-68.	0.3	1
138	Introduction to proteomics: tools for the new biology. Expert Review of Proteomics, 2004, 1, 9-10.	1.3	1
139	Analysis of Endocrine Disruption in Southern California Coastal Fish using an Aquatic Multi-Species Microarray. Nature Precedings, 2009, , .	0.1	1
140	Research Highlights. Pharmacogenomics, 2011, 12, 1637-1639.	0.6	1
141	De Novo Hepatic Transcriptome Assembly and Systems Level Analysis of Three Species of Dietary Fish, Sardinops sagax, Scomber japonicus, and Pleuronichthys verticalis. Genes, 2018, 9, 521.	1.0	1
142	Pipeline for Integrated Microarray Expression Normalization Tool Kit (PIMENTo) for Tumor Microarray Profiling Experiments. Methods in Molecular Biology, 2019, 1908, 153-168.	0.4	1
143	The acute transcriptome response of the midbrain/diencephalon to injury in the adult mummichog (Fundulus heteroclitus). Molecular Brain, 2019, 12, 119.	1.3	1
144	The effects of rosiglitazone on the neonatal rat cardiomyocyte transcriptome: a temporal analysis. Pharmacogenomics, 2019, 20, 1125-1141.	0.6	1

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145	Molecular Profiling of RNA Tumors Using High-Throughput RNA Sequencing: Overview of Library Preparation Methods. Methods in Molecular Biology, 2019, 1908, 169-184.	0.4	1
146	The isolation and structure of liver and globin genes from Atlantic salmon. Coastal and Estuarine Studies, 1993, , 255-273.	0.4	0
147	Novel and rapid personal whole-genome sequencing: recent advances and the promise for translational medicine. Pharmacogenomics, 2008, 9, 667-670.	0.6	0
148	Development of a bioassay to monitor circulating plasma Ki-67. Leukemia Research, 2010, 34, 848-849.	0.4	0
149	PO-136 SOX2 is a novel biomarker for high-risk ewing sarcoma. ESMO Open, 2018, 3, A74.	2.0	Ο
150	Gene Expression Differences Between Young Adults Based on Trauma History and Post-traumatic Stress Disorder. Frontiers in Psychiatry, 2021, 12, 581093.	1.3	0
151	Characterization of Microarray Hybridization Stoichiometry. Drug Discovery Series, 2009, , 87-96.	0.1	Ο
152	Microarrays in Neuroscience. Drug Discovery Series, 2009, , 271-288.	0.1	0
153	Introduction to Large-Scale Gene Expression Data Analysis. Drug Discovery Series, 2009, , 11-24.	0.1	0
154	Effect of bypass kinase pathways on acquired CDK4/6 inhibitor resistance Journal of Clinical Oncology, 2018, 36, 379-379.	0.8	0
155	Abstract P4-04-11: The 8p11-p12 amplicon oncogene ASH2L regulates expression of genes involved in tumorigenic processes and response to palbociclib via promoter H3K4me3. , 2018, , .		Ο
156	Abstract A043: Bypass kinase pathways lead to acquired CDK4/6 inhibitor resistance in prostate cancer. , 2018, , .		0
157	Correction: Nucleobase-containing compounds evoke behavioural, olfactory, and transcriptional responses in model fishes. Facets, 2018, 3, 598-598.	1.1	0
158	META-ANALYSIS OF DOLPHIN AND HUMAN PERIPHERAL BLOOD MONONUCLEAR CELLS REVEALS INFLAMMATORY SIGNATURES ASSOCIATED WITH EXPOSURE TO HIGH LEVELS OF PERFLUOROALKYL SUBSTANCES. International Journal of Advances in Science Engineering and Technology, 2019, 7, 66-72.	1.0	0
159	Cloning and sequencing of the Atlantic salmon (Salmo salar) cytochrome c oxidase subunit III gene (coxIII) and analysis of coxIII expression during parr-smolt transformation. Molecular Marine Biology and Biotechnology, 1994, 3, 210-6.	0.4	0
160	Genomic variation in the mouse. Pharmacogenomics, 2011, 12, 1638-9.	0.6	0
161	Structural variation in the mouse genome. Pharmacogenomics, 2011, 12, 1639.	0.6	0
162	Fusion Genes in Prostate Cancer: A Comparison in Men of African and European Descent. Biology, 2022, 11, 625.	1.3	0