Dieter Deforce

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2698983/publications.pdf

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369 papers 11,779 citations

53 h-index 80 g-index

388 all docs 388 docs citations

times ranked

388

17957 citing authors

#	Article	IF	CITATIONS
1	The Chara Genome: Secondary Complexity and Implications for Plant Terrestrialization. Cell, 2018, 174, 448-464.e24.	13.5	420
2	Genome-wide association study of obsessive-compulsive disorder. Molecular Psychiatry, 2013, 18, 788-798.	4.1	312
3	Reduced amounts and abnormal forms of phospholipase C zeta (PLCÂ) in spermatozoa from infertile men. Human Reproduction, 2009, 24, 2417-2428.	0.4	257
4	Partitioning the Heritability of Tourette Syndrome and Obsessive Compulsive Disorder Reveals Differences in Genetic Architecture. PLoS Genetics, 2013, 9, e1003864.	1.5	241
5	Dissection of the phytohormonal regulation of trichome formation and biosynthesis of the antimalarial compound artemisinin in <i>Artemisia annua</i>) plants. New Phytologist, 2011, 189, 176-189.	3 . 5	192
6	Development and evaluation of different normalization strategies for gene expression studies in Candida albicans biofilms by real-time PCR. BMC Molecular Biology, 2006, 7, 25.	3.0	139
7	Development and validation of an LC-MS/MS method for the simultaneous determination of deoxynivalenol, zearalenone, T-2-toxin and some masked metabolites in different cereals and cereal-derived food. Food Additives and Contaminants - Part A Chemistry, Analysis, Control, Exposure and Risk Assessment. 2012. 29. 819-835.	1.1	129
8	Real-time PCR expression profiling of genes encoding potential virulence factors in Candida albicans biofilms: identification of model-dependent and -independent gene expression. BMC Microbiology, 2010, 10, 114.	1.3	127
9	A Plant-Derived Ligand Favoring Monomeric Glucocorticoid Receptor Conformation with Impaired Transactivation Potential Attenuates Collagen-Induced Arthritis. Journal of Immunology, 2008, 180, 2608-2615.	0.4	125
10	Joint GC–MS and LC–MS platforms for comprehensive plant metabolomics: Repeatability and sample pre-treatment. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2009, 877, 3572-3580.	1,2	122
11	Ancient Dispersal of the Human Fungal Pathogen Cryptococcus gattii from the Amazon Rainforest. PLoS ONE, 2013, 8, e71148.	1.1	122
12	Alternative Routes to Induce NaÃ-ve Pluripotency in Human Embryonic Stem Cells. Stem Cells, 2015, 33, 2686-2698.	1.4	118
13	Tracking the progression of the human inner cell mass during embryonic stem cell derivation. Nature Biotechnology, 2012, 30, 278-282.	9.4	109
14	TRAPID: an efficient online tool for the functional and comparative analysis of de novo RNA-Seq transcriptomes. Genome Biology, 2013, 14, R134.	13.9	108
15	Differential transcriptome analysis of glandular and filamentous trichomes in Artemisia annua. BMC Plant Biology, 2013, 13, 220.	1.6	106
16	Simultaneous, quantitative determination of opiates, amphetamines, cocaine and benzoylecgonine in oral fluid by liquid chromatography quadrupole-time-of-flight mass spectrometry. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2002, 779, 321-330.	1.2	102
17	Complete Nucleotide Sequence of CTX-M-15-Plasmids from Clinical Escherichia coli Isolates: Insertional Events of Transposons and Insertion Sequences. PLoS ONE, 2010, 5, e11202.	1.1	101
18	Current and New Approaches in GMO Detection: Challenges and Solutions. BioMed Research International, 2015, 2015, 1-22.	0.9	98

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19	OSC2 and CYP716A14v2 Catalyze the Biosynthesis of Triterpenoids for the Cuticle of Aerial Organs of <i>Artemisia annua</i> Plant Cell, 2015, 27, 286-301.	3.1	96
20	Forensic STR analysis using massive parallel sequencing. Forensic Science International: Genetics, 2012, 6, 810-818.	1.6	94
21	Chromosomal mosaicism in human blastocysts: the ultimate challenge of preimplantation genetic testing?. Human Reproduction, 2018, 33, 1342-1354.	0.4	94
22	Mass spectrometry-based proteomics as a tool to identify biological matrices in forensic science. International Journal of Legal Medicine, 2013, 127, 287-298.	1.2	91
23	TBX2 is a neuroblastoma core regulatory circuitry component enhancing MYCN/FOXM1 reactivation of DREAM targets. Nature Communications, 2018, 9, 4866.	5. 8	91
24	Integrative Proteomic Profiling Reveals PRC2-Dependent Epigenetic Crosstalk Maintains Ground-State Pluripotency. Cell Stem Cell, 2019, 24, 123-137.e8.	5 . 2	90
25	The transcriptome of Mycobacterium tuberculosis in a lipid-rich dormancy model through RNAseq analysis. Scientific Reports, 2017, 7, 17665.	1.6	88
26	Analysis of DNA Adducts in DNA Hydrolysates by Capillary Zone Electrophoresis and Capillary Zone Electrophoresisâ^Electrospray Mass Spectrometry. Analytical Chemistry, 1996, 68, 3575-3584.	3.2	87
27	Citrullinated vimentin as an important antigen in immune complexes from synovial fluid of rheumatoid arthritis patients with antibodies against citrullinated proteins. Arthritis Research and Therapy, 2010, 12, R132.	1.6	87
28	Diagnostic value of anti-human citrullinated fibrinogen ELISA and comparison with four other anti-citrullinated protein assays. Arthritis Research and Therapy, 2006, 8, R122.	1.6	86
29	Latent sensitivity to Fas-mediated apoptosis after CD40 ligation may explain activity of CD154 gene therapy in chronic lymphocytic leukemia. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 3854-3859.	3.3	78
30	Chronically inflamed synovium from spondyloarthropathy and rheumatoid arthritis investigated by protein expression profiling followed by tandem mass spectrometry. Proteomics, 2005, 5, 2247-2257.	1.3	77
31	Progranulin expression correlates with denseâ€core amyloid plaque burden in Alzheimer disease mouse models. Journal of Pathology, 2009, 219, 173-181.	2.1	75
32	My-Forensic-Loci-queries (MyFLq) framework for analysis of forensic STR data generated by massive parallel sequencing. Forensic Science International: Genetics, 2014, 9, 1-8.	1.6	75
33	Association between serotonergic candidate genes and specific phenotypes of obsessive compulsive disorder. Journal of Affective Disorders, 2006, 91, 39-44.	2.0	73
34	Presence and potential of cell free DNA in different types of forensic samples. Forensic Science International: Genetics, 2013, 7, 316-320.	1.6	72
35	Differential proteome analysis of normal and osteoarthritic chondrocytes reveals distortion of vimentin network in osteoarthritis. Osteoarthritis and Cartilage, 2008, 16, 163-173.	0.6	70
36	Quantitation of artemisinin and its biosynthetic precursors in Artemisia annua L. by high performance liquid chromatography–electrospray quadrupole time-of-flight tandem mass spectrometry. Journal of Chromatography A, 2006, 1118, 180-187.	1.8	69

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37	Synovial detection and autoantibody reactivity of processed citrullinated isoforms of vimentin in inflammatory arthritides. Rheumatology, 2008, 47, 597-604.	0.9	69
38	The HCV serum proteome: a search for fibrosis protein markers. Journal of Viral Hepatitis, 2009, 16, 418-429.	1.0	68
39	Transgenerational Inheritance of DNA Hypomethylation in <i>Daphnia magna</i> in Response to Salinity Stress. Environmental Science & Environmental Sci	4.6	67
40	Association between the dopamine D2 receptor Taql A2 allele and low activity COMT allele with obsessive–compulsive disorder in males. European Neuropsychopharmacology, 2006, 16, 446-450.	0.3	66
41	<i>In Vitro</i> Inhibition of the Transcription Factor NFâ€PB and Cyclooxygenase by Bamboo Extracts. Phytotherapy Research, 2014, 28, 224-230.	2.8	66
42	Laser capture microdissection in forensic research: a review. International Journal of Legal Medicine, 2010, 124, 513-521.	1.2	65
43	Assessment of Microbial Diversity in Biofilms Recovered from Endotracheal Tubes Using Culture Dependent and Independent Approaches. PLoS ONE, 2012, 7, e38401.	1.1	65
44	Selective serotonin reuptake inhibitors as a novel class of immunosuppressants. International Immunopharmacology, 2014, 20, 148-156.	1.7	65
45	Genetically Modified Micro-Organisms for Industrial Food Enzyme Production: An Overview. Foods, 2020, 9, 326.	1.9	65
46	The relevance of citrullinated vimentin in the production of antibodies against citrullinated proteins and the pathogenesis of rheumatoid arthritis. Rheumatology, 2011, 50, 830-837.	0.9	64
47	Proteomic Analysis of the Excretory-Secretory Products from Larval Stages of Ascaris suum Reveals High Abundance of Clycosyl Hydrolases. PLoS Neglected Tropical Diseases, 2013, 7, e2467.	1.3	63
48	CRISPR/Cas9 mediated knockout of rb1 and rbl1 leads to rapid and penetrant retinoblastoma development in Xenopus tropicalis. Scientific Reports, 2016, 6, 35264.	1.6	63
49	Transcriptome Analysis of Bombyx mori Larval Midgut during Persistent and Pathogenic Cytoplasmic Polyhedrosis Virus Infection. PLoS ONE, 2015, 10, e0121447.	1.1	63
50	Direct comparison of distinct naive pluripotent states in human embryonic stem cells. Nature Communications, 2017, 8, 15055.	5.8	60
51	Prediction of Response to Paroxetine and Venlafaxine by Serotonin-Related Genes in Obsessive-Compulsive Disorder in a Randomized, Double-Blind Trial. Journal of Clinical Psychiatry, 2007, 68, 747-753.	1.1	59
52	Tryptic peptide analysis of protein binders in works of art by liquid chromatography–tandem mass spectrometry. Analytica Chimica Acta, 2010, 658, 156-162.	2.6	58
53	Unbreakable Codes in Electrospun Fibers: Digitally Encoded Polymers to Stop Medicine Counterfeiting. Advanced Materials, 2010, 22, 2657-2662.	11.1	58
54	Histone proteolysis: A proposal for categorization into â€~clipping' and â€~degradation'. BioEssays, 2015, 70-79.	, 37, 1.2	58

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55	Suboptimal culture conditions induce more deviations in gene expression in male than female bovine blastocysts. BMC Genomics, 2016, 17, 72.	1.2	58
56	Application of a new anti-zearalenone monoclonal antibody in different immunoassay formats. Analytical and Bioanalytical Chemistry, 2009, 395, 1301-1307.	1.9	57
57	A Single Early Activation of Invariant NK T Cells Confers Long-Term Protection against Collagen-Induced Arthritis in a Ligand-Specific Manner. Journal of Immunology, 2007, 179, 2300-2309.	0.4	56
58	DNA typing of fingerprints using capillary electrophoresis: Effect of dactyloscopic powders. Electrophoresis, 1999, 20, 2870-2876.	1.3	54
59	Growth Differentiation Factor 15, a Marker of Lung Involvement in Systemic Sclerosis, Is Involved in Fibrosis Development but Is not Indispensable for Fibrosis Development. Arthritis and Rheumatology, 2014, 66, 418-427.	2.9	54
60	Forensic SNP Genotyping using Nanopore MinION Sequencing. Scientific Reports, 2017, 7, 41759.	1.6	54
61	Performance of four modern whole genome amplification methods for copy number variant detection in single cells. Scientific Reports, 2017, 7, 3422.	1.6	54
62	Comparative analysis of naive, primed and ground state pluripotency in mouse embryonic stem cells originating from the same genetic background. Scientific Reports, 2018, 8, 5884.	1.6	54
63	Analysis of nucleic acid constituents by on-line capillary electrophoresis-mass spectrometry. Electrophoresis, 2005, 26, 1221-1253.	1.3	53
64	Illumina mate-paired DNA sequencing-library preparation using Cre-Lox recombination. Nucleic Acids Research, 2012, 40, e24-e24.	6.5	52
65	Batrachochytrium dendrobatidis zoospore secretions rapidly disturb intercellular junctions in frog skin. Fungal Genetics and Biology, 2012, 49, 830-837.	0.9	51
66	16S rRNA Amplicon Sequencing Demonstrates that Indoor-Reared Bumblebees (Bombus terrestris) Harbor a Core Subset of Bacteria Normally Associated with the Wild Host. PLoS ONE, 2015, 10, e0125152.	1.1	51
67	Quantitative Bias in Illumina TruSeq and a Novel Post Amplification Barcoding Strategy for Multiplexed DNA and Small RNA Deep Sequencing. PLoS ONE, 2011, 6, e26969.	1.1	51
68	InÂvivo study of Dicer-2-mediated immune response of the small interfering RNA pathway upon systemic infections of virulent and avirulent viruses in Bombus terrestris. Insect Biochemistry and Molecular Biology, 2016, 70, 127-137.	1.2	50
69	Colony contact contributes to the diversity of gut bacteria in bumblebees (<i>Bombus terrestris</i> lnsect Science, 2017, 24, 270-277.	1.5	50
70	Automatic detection of spermatozoa for laser capture microdissection. International Journal of Legal Medicine, 2009, 123, 169-175.	1.2	49
71	Molecular diagnostics for congenital hearing loss including 15 deafness genes using a next generation sequencing platform. BMC Medical Genomics, 2012, 5, 17.	0.7	49
72	Minimizing technical variation during sample preparation prior toÂlabel-free quantitative mass spectrometry. Analytical Biochemistry, 2015, 490, 14-19.	1.1	48

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73	Invariant natural killer T cells are natural regulators of murine spondylarthritis. Arthritis and Rheumatism, 2010, 62, 988-999.	6.7	47
74	Viral Small-RNA Analysis of Bombyx mori Larval Midgut during Persistent and Pathogenic Cytoplasmic Polyhedrosis Virus Infection. Journal of Virology, 2015, 89, 11473-11486.	1.5	47
75	Statistical framework for detection of genetically modified organisms based on Next Generation Sequencing. Food Chemistry, 2016, 192, 788-798.	4.2	47
76	Autocrine embryotropins revisited: how do embryos communicate with each other <i>in vitro</i> when cultured in groups?. Biological Reviews, 2017, 92, 505-520.	4.7	47
77	A multi-omics analysis of the grapevine pathogen Lasiodiplodia theobromae reveals that temperature affects the expression of virulence- and pathogenicity-related genes. Scientific Reports, 2019, 9, 13144.	1.6	47
78	Laser capture microdissection: Should an ultraviolet or infrared laser be used?. Analytical Biochemistry, 2013, 439, 88-98.	1.1	45
79	The Determination of Cocaine, Benzoylecgonine, and Cocaethylene in Small-Volume Oral Fluid Samples by Liquid Chromatography-Quadrupole-Time-of-Flight Mass Spectrometry. Journal of Analytical Toxicology, 2004, 28, 655-659.	1.7	44
80	Passive immunization to reduce Campylobacter jejuni colonization and transmission in broiler chickens. Veterinary Research, 2014, 45, 27.	1.1	44
81	Identification of histone H3 clipping activity in human embryonic stem cells. Stem Cell Research, 2014, 13, 123-134.	0.3	44
82	Host metabolites stimulate the bacterial proton motive force to enhance the activity of aminoglycoside antibiotics. PLoS Pathogens, 2019, 15, e1007697.	2.1	44
83	Monitoring ALS1 and ALS3 Gene Expression During InÂVitro Candida albicans Biofilm Formation Under Continuous Flow Conditions. Mycopathologia, 2009, 167, 9-17.	1.3	43
84	Safety assessment of the butyrate-producing Butyricicoccus pullicaecorum strain 25-3T, a potential probiotic for patients with inflammatory bowel disease, based on oral toxicity tests and whole genome sequencing. Food and Chemical Toxicology, 2014, 72, 129-137.	1.8	43
85	Proteins involved in embryo-maternal interaction around the signalling of maternal recognition of pregnancy in the horse. Scientific Reports, 2018, 8, 5249.	1.6	43
86	A Doubling of Microphytobenthos Biomass Coincides with a Tenfold Increase in Denitrifier and Total Bacterial Abundances in Intertidal Sediments of a Temperate Estuary. PLoS ONE, 2015, 10, e0126583.	1.1	43
87	LC–MS metabolic profiling of Arabidopsis thaliana plant leaves and cell cultures: Optimization of pre-LC–MS procedure parameters. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2008, 871, 37-43.	1.2	42
88	Targeting aminopeptidase N, a newly identified receptor for F4ac fimbriae, enhances the intestinal mucosal immune response. Mucosal Immunology, 2012, 5, 635-645.	2.7	42
89	Perfrin, a novel bacteriocin associated with netB positive Clostridium perfringens strains from broilers with necrotic enteritis. Veterinary Research, 2014, 45, 40.	1.1	42
90	Whole genome amplification with SurePlex results in better copy number alteration detection using sequencing data compared to the MALBAC method. Scientific Reports, 2015, 5, 11711.	1.6	42

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91	Dendritic cells from spondylarthritisâ€prone HLA–B27–transgenic rats display altered cytoskeletal dynamics, class II major histocompatibility complex expression, and viability. Arthritis and Rheumatism, 2009, 60, 2622-2632.	6.7	41
92	Shallow whole genome sequencing is well suited for the detection of chromosomal aberrations in human blastocysts. Fertility and Sterility, 2015, 104, 1276-1285.e1.	0.5	40
93	Practical Tools to Implement Massive Parallel Pyrosequencing of PCR Products in Next Generation Molecular Diagnostics. PLoS ONE, 2011, 6, e25531.	1.1	40
94	Cutting Edge: The Chemokine Receptor CXCR3 Retains Invariant NK T Cells in the Thymus. Journal of Immunology, 2009, 183, 2213-2216.	0.4	39
95	Transcriptional response to fluconazole and amphotericin B in Candida albicans biofilms. Research in Microbiology, 2010, 161, 284-292.	1.0	39
96	Transport of Hop Bitter Acids across Intestinal Caco-2 Cell Monolayers. Journal of Agricultural and Food Chemistry, 2010, 58, 4132-4140.	2.4	38
97	An innovative and integrated approach based on DNA walking to identify unauthorised GMOs. Food Chemistry, 2014, 147, 60-69.	4.2	38
98	Removing the Hidden Data Dependency of DIA with Predicted Spectral Libraries. Proteomics, 2020, 20, e1900306.	1.3	38
99	Shorter CAG repeats in the androgen receptor gene may enhance hyperandrogenicity in polycystic ovary syndrome. Gynecological Endocrinology, 2008, 24, 669-673.	0.7	37
100	Do all anti-citrullinated protein/peptide antibody tests measure the same? Evaluation of discrepancy between anti-citrullinated protein/peptide antibody tests in patients with and without rheumatoid arthritis. Annals of the Rheumatic Diseases, 2008, 67, 542-546.	0.5	37
101	Antioxidant Flavone Glycosides from the Leaves of <i>Fargesia robusta</i> . Journal of Natural Products, 2010, 73, 1573-1577.	1.5	37
102	Optimization and diagnostic performance of a single multiparameter lineblot in the serological workup of systemic sclerosis. Journal of Immunological Methods, 2012, 379, 53-60.	0.6	37
103	Divergence between the Highly Virulent Zoonotic Pathogen Helicobacter heilmannii and Its Closest Relative, the Low-Virulence "Helicobacter ailurogastricus―sp. nov. Infection and Immunity, 2016, 84, 293-306.	1.0	37
104	Plasma Functionalization of Polycaprolactone Nanofibers Changes Protein Interactions with Cells, Resulting in Increased Cell Viability. ACS Applied Materials & Samp; Interfaces, 2018, 10, 41962-41977.	4.0	37
105	Optimization of solid-phase extraction for a liquid chromatographic–tandem mass spectrometric general unknown screening procedure by means of computational techniques. Journal of Chromatography A, 2004, 1056, 57-65.	1.8	36
106	Caspase-14-Deficient Mice Are More Prone to the Development of Parakeratosis. Journal of Investigative Dermatology, 2013, 133, 742-750.	0.3	35
107	Impact of sugar syrup and pollen diet on the bacterial diversity in the gut of indoor-reared bumblebees (Bombus terrestris). Apidologie, 2016, 47, 548-560.	0.9	35
108	The Ontogeny of Cytochrome P450 Enzyme Activity and Protein Abundance in Conventional Pigs in Support of Preclinical Pediatric Drug Research. Frontiers in Pharmacology, 2018, 9, 470.	1.6	35

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109	Forensic tri-allelic SNP genotyping using nanopore sequencing. Forensic Science International: Genetics, 2019, 38, 204-210.	1.6	35
110	Comparison of slab gel electrophoresis and capillary electrophoresis for the detection of the fluorescently labeled polymerase chain reaction products of short tandem repeat fragments. Journal of Chromatography A, 1998, 806, 149-155.	1.8	34
111	Modification of citrulline residues with 2,3â€butanedione facilitates their detection by liquid chromatography/mass spectrometry. Rapid Communications in Mass Spectrometry, 2011, 25, 1536-1542.	0.7	34
112	Pitfalls in histone propionylation during bottomâ€up mass spectrometry analysis. Proteomics, 2015, 15, 2966-2971.	1.3	34
113	The Equine Embryo Influences Immune-Related Gene Expression in the Oviduct1. Biology of Reproduction, 2016, 94, 36.	1.2	34
114	First draft genome assembly of the desert locust, Schistocerca gregaria. F1000Research, 2020, 9, 775.	0.8	34
115	Suspension fluorescence in situ hybridization (S-FISH) combined with automatic detection and laser microdissection for STR profiling of male cells in male/female mixtures. International Journal of Legal Medicine, 2009, 123, 441-447.	1.2	33
116	Fetal Microchimeric Cells in Blood of Women with an Autoimmune Thyroid Disease. PLoS ONE, 2011, 6, e29646.	1.1	33
117	Genome-wide transcription start site profiling in biofilm-grown Burkholderia cenocepacia J2315. BMC Genomics, 2015, 16, 775.	1.2	33
118	An integrated strategy combining DNA walking and NGS to detect GMOs. Food Chemistry, 2017, 232, 351-358.	4.2	33
119	Nanopore Sequencing of a Forensic STR Multiplex Reveals Loci Suitable for Single-Contributor STR Profiling. Genes, 2020, 11, 381.	1.0	33
120	Presence and Analysis of Plasmids in Human and Animal Associated Arcobacter Species. PLoS ONE, 2014, 9, e85487.	1.1	33
121	Characterization of DNA Oligonucleotides by Coupling of Capillary Zone Electrophoresis to Electrospray Ionization Q-TOF Mass Spectrometry. Analytical Chemistry, 1998, 70, 3060-3068.	3.2	32
122	NKT cells: manipulable managers of joint inflammation. Rheumatology, 2006, 46, 565-571.	0.9	32
123	A unique lymphotoxin Âbeta-dependent pathway regulates thymic emigration of VÂ14 invariant natural killer T cells. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 9160-9165.	3.3	32
124	Evaluation of hydrophilic interaction chromatography <i>versus</i> reversedâ€phase chromatography in a plant metabolomics perspective. Journal of Separation Science, 2008, 31, 1609-1614.	1.3	31
125	Improved canine exome designs, featuring ncRNAs and increased coverage of protein coding genes. Scientific Reports, 2015, 5, 12810.	1.6	31
126	Validation of a sensitive DNA walking strategy to characterise unauthorised GMOs using model food matrices mimicking common rice products. Food Chemistry, 2015, 173, 1259-1265.	4.2	31

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127	Characterization of Porcine Hepatic and Intestinal Drug Metabolizing CYP450: Comparison with Human Orthologues from A Quantitative, Activity and Selectivity Perspective. Scientific Reports, 2019, 9, 9233.	1.6	31
128	Synthesis and in vitro cytotoxicity of aminocoumarin platinum(II) complexes. Bioorganic and Medicinal Chemistry Letters, 1997, 7, 2165-2168.	1.0	30
129	X linked agammaglobulinaemia and rheumatoid arthritis. Annals of the Rheumatic Diseases, 2005, 64, 1075-1078.	0.5	30
130	T-2 toxin induced Salmonella Typhimurium intoxication results in decreased Salmonella numbers in the cecum contents of pigs, despite marked effects on Salmonella-host cell interactions. Veterinary Research, 2012, 43, 22.	1.1	30
131	Metabolism of Hop-Derived Bitter Acids. Journal of Agricultural and Food Chemistry, 2013, 61, 7916-7924.	2.4	30
132	Reference loci for RT-qPCR analysis of differentiating human embryonic stem cells. BMC Molecular Biology, 2013, 14, 21.	3.0	30
133	Biotech rice: Current developments and future detection challenges in food and feed chain. Trends in Food Science and Technology, 2016, 52, 66-79.	7.8	30
134	Host intestinal biomarker identification in a gut leakage model in broilers. Veterinary Research, 2019, 50, 46.	1.1	30
135	Unique long non-coding RNA expression signature in ETV6/RUNX1-driven B-cell precursor acute lymphoblastic leukemia. Oncotarget, 2016, 7, 73769-73780.	0.8	30
136	Integrated multi-omics reveal polycomb repressive complex 2 restricts human trophoblast induction. Nature Cell Biology, 2022, 24, 858-871.	4.6	30
137	Phenolic Compounds and Antiâ€oxidant Capacity of Twelve Morphologically Heterogeneous Bamboo species. Phytochemical Analysis, 2012, 23, 433-443.	1.2	29
138	Bovine Embryo-Secreted microRNA-30c Is a Potential Non-invasive Biomarker for Hampered Preimplantation Developmental Competence. Frontiers in Genetics, 2019, 10, 315.	1.1	29
139	The transcriptome of the marine calanoid copepod Temora longicornis under heat stress and recovery. Marine Environmental Research, 2019, 143, 10-23.	1.1	29
140	Cov-MS: A Community-Based Template Assay for Mass-Spectrometry-Based Protein Detection in SARS-CoV-2 Patients. Jacs Au, 2021, 1, 750-765.	3.6	29
141	Galectin-3-binding protein: a serological and histological assessment in accordance with hepatitis C-related liver fibrosis. European Journal of Gastroenterology and Hepatology, 2010, 22, 1066-1073.	0.8	28
142	Proteome characterization of human articular chondrocytes leads to novel insights in the function of small heat-shock proteins in chondrocyte homeostasis. Osteoarthritis and Cartilage, 2010, 18, 440-446.	0.6	28
143	In vivo relevance of citrullinated proteins and the challenges in their detection. Proteomics, 2012, 12, 752-760.	1.3	28
144	The Combination of Inhibitors of FGF/MEK/Erk and GSK3 \hat{l}^2 Signaling Increases the Number of OCT3/4- and NANOG-Positive Cells in the Human Inner Cell Mass, But Does Not Improve Stem Cell Derivation. Stem Cells and Development, 2013, 22, 296-306.	1.1	28

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145	Refined Candidate Region for F4ab/ac Enterotoxigenic Escherichia coli Susceptibility Situated Proximal to MUC13 in Pigs. PLoS ONE, 2014, 9, e105013.	1.1	28
146	Proteome of equine oviducal fluid: effects of ovulation and pregnancy. Reproduction, Fertility and Development, 2017, 29, 1085.	0.1	28
147	Dual RNA Sequencing of Vitis vinifera during Lasiodiplodia theobromae Infection Unveils Host–Pathogen Interactions. International Journal of Molecular Sciences, 2019, 20, 6083.	1.8	28
148	Analysis of the DNA adducts of phenyl glycidyl ether in a calf thymus DNA hydrolysate by capillary zone electrophoresis-electrospray mass spectrometry: evidence for phosphate alkylation. Carcinogenesis, 1998, 19, 1077-1086.	1.3	27
149	Analysis of melphalan adducts of 2′-deoxynucleotides in calf thymus DNA hydrolysates by capillary high-performance liquid chromatography‰electrospray tandem mass spectrometry. Biomedical Applications, 1999, 736, 43-59.	1.7	27
150	Assessment of nuclear transfer techniques to prevent the transmission of heritable mitochondrial disorders without compromising embryonic development competence in mice. Mitochondrion, 2014, 18, 27-33.	1.6	27
151	Forensic massively parallel sequencing data analysis tool: Implementation of MyFLq as a standalone web- and Illumina BaseSpaceA®-application. Forensic Science International: Genetics, 2015, 15, 2-7.	1.6	27
152	BMP-SMAD Signaling Regulates Lineage Priming, but Is Dispensable for Self-Renewal in Mouse Embryonic Stem Cells. Stem Cell Reports, 2016, 6, 85-94.	2.3	27
153	Comparison of Three Methods to Quantify Laxity in the Canine Hip Joint. Veterinary and Comparative Orthopaedics and Traumatology, 2018, 31, 023-029.	0.2	27
154	Deguelin inhibits expression of lκBα protein and induces apoptosis of B-CLL cells in vitro. Leukemia, 2007, 21, 1610-1618.	3.3	26
155	The novel â€~ <i>Candidatus</i> Amphibiichlamydia ranarum' is highly prevalent in invasive exotic bullfrogs (<i>Lithobates catesbeianus</i>). Environmental Microbiology Reports, 2013, 5, 105-108.	1.0	26
156	Exogenous supplementation of Activin A enhances germ cell differentiation of human embryonic stem cellsâ€. Molecular Human Reproduction, 2015, 21, 410-423.	1.3	26
157	Nanopore sequencing technology: a new route for the fast detection of unauthorized GMO. Scientific Reports, 2018, 8, 7903.	1.6	26
158	Analysis of the DNA Damage Induced by Phenyl Glycidyl Ether Using Capillary Zone Electrophoresis–Electrospray Mass Spectrometry. Analytical Biochemistry, 1998, 258, 331-338.	1.1	25
159	Treatment of human embryos with the TGFÂ inhibitor SB431542 increases epiblast proliferation and permits successful human embryonic stem cell derivation. Human Reproduction, 2014, 29, 41-48.	0.4	25
160	Efficiency of Exome Sequencing for the Molecular Diagnosis of Pseudoxanthoma Elasticum. Journal of Investigative Dermatology, 2015, 135, 992-998.	0.3	25
161	Ecological divergence of wild strawberry DNA methylation patterns at distinct spatial scales. Molecular Ecology, 2020, 29, 4871-4881.	2.0	25
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