Christian Cole

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3621505/publications.pdf Version: 2024-02-01



CHDISTIAN COLF

#	Article	IF	CITATIONS
1	Research trends in forensic science: A scientometric approach to analyze the content of the <scp>INTERPOL</scp> reviews. Wiley Interdisciplinary Reviews Forensic Science, 2022, 4, e1447.	2.1	5
2	A National Network of Safe Havens: Scottish Perspective. Journal of Medical Internet Research, 2022, 24, e31684.	4.3	6
3	Creation of a universal experimental protocol for the investigation of transfer and persistence of trace evidence: Part 1 - From design to implementation for particulate evidence. Forensic Science International (Online), 2021, 3, 100165.	1.3	1
4	Creation of a universal experimental protocol for the investigation of transfer and persistence of trace evidence: Part 2 – Implementation and preliminary data. Forensic Science International (Online), 2021, 3, 100164.	1.3	0
5	Reviewing Research Trends—A Scientometric Approach Using Gunshot Residue (GSR) Literature as an Example. Publications, 2020, 8, 7.	3.8	9
6	EMSY expression affects multiple components of the skin barrier with relevance to atopic dermatitis. Journal of Allergy and Clinical Immunology, 2019, 144, 470-481.	2.9	23
7	Adenylate cyclase A acting on PKA mediates induction of stalk formation by cyclic diguanylate at the <i>Dictyostelium</i> organizer. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 516-521.	7.1	22
8	A mechanistic target of rapamycin complex 1/2 (mTORC1)/V-Akt murine thymoma viral oncogene homolog 1 (AKT1)/cathepsin H axis controls filaggrin expression and processing in skin, a novel mechanism for skin barrier disruption in patients with atopic dermatitis. Journal of Allergy and Clinical Immunology, 2017, 139, 1228-1241.	2.9	38
9	A Proteomic Approach to Analyze the Aspirin-mediated Lysine Acetylome. Molecular and Cellular Proteomics, 2017, 16, 310-326.	3.8	45
10	Improved annotation with de novo transcriptome assembly in four social amoeba species. BMC Genomics, 2017, 18, 120.	2.8	7
11	How many biological replicates are needed in an RNA-seq experiment and which differential expression tool should you use?. Rna, 2016, 22, 839-851.	3.5	622
12	AlmostSignificant: simplifying quality control of high-throughput sequencing data. Bioinformatics, 2016, 32, 3850-3851.	4.1	6
13	Keratin 12 missense mutation induces the unfolded protein response and apoptosis in Meesmann epithelial corneal dystrophy. Human Molecular Genetics, 2016, 25, 1176-1191.	2.9	22
14	14-3-3-Pred: improved methods to predict 14-3-3-binding phosphopeptides. Bioinformatics, 2015, 31, 2276-2283.	4.1	177
15	JPred4: a protein secondary structure prediction server. Nucleic Acids Research, 2015, 43, W389-W394.	14.5	1,546
16	Loss-of-Function Mutations in CAST Cause Peeling Skin, Leukonychia, Acral Punctate Keratoses, Cheilitis, and Knuckle Pads. American Journal of Human Genetics, 2015, 96, 440-447.	6.2	36
17	Severe dermatitis, multiple allergies, and metabolic wasting syndrome caused by a novel mutation in the N-terminal plakin domain of desmoplakin. Journal of Allergy and Clinical Immunology, 2015, 136, 1268-1276.	2.9	103
18	Statistical models for RNA-seq data derived from a two-condition 48-replicate experiment. Bioinformatics, 2015, 31, 3625-3630.	4.1	76

CHRISTIAN COLE

#	Article	IF	CITATIONS
19	Expanding the Phenotypic Spectrum of Olmsted Syndrome. Journal of Investigative Dermatology, 2015, 135, 2879-2883.	0.7	23
20	Improved Annotation of 3′ Untranslated Regions and Complex Loci by Combination of Strand-Specific Direct RNA Sequencing, RNA-Seq and ESTs. PLoS ONE, 2014, 9, e94270.	2.5	27
21	Filaggrin-stratified transcriptomic analysis of pediatric skin identifies mechanistic pathways in patients with atopic dermatitis. Journal of Allergy and Clinical Immunology, 2014, 134, 82-91.	2.9	118
22	Tmem79/Matt is the matted mouse gene and is a predisposing gene for atopic dermatitis in human subjects. Journal of Allergy and Clinical Immunology, 2013, 132, 1121-1129.	2.9	135
23	Transcription Termination and Chimeric RNA Formation Controlled by Arabidopsis thaliana FPA. PLoS Genetics, 2013, 9, e1003867.	3.5	67
24	Haploinsufficiency for AAGAB causes clinically heterogeneous forms of punctate palmoplantar keratoderma. Nature Genetics, 2012, 44, 1272-1276.	21.4	78
25	Direct sequencing of Arabidopsis thaliana RNA reveals patterns of cleavage and polyadenylation. Nature Structural and Molecular Biology, 2012, 19, 845-852.	8.2	142
26	A Dictyostelium SH2 adaptor protein required for correct DIF-1 signaling and pattern formation. Developmental Biology, 2011, 353, 290-301.	2.0	3
27	Regulation of the miR-212/132 locus by MSK1 and CREB in response to neurotrophins. Biochemical Journal, 2010, 428, 281-291.	3.7	195
28	Filtering of deep sequencing data reveals the existence of abundant Dicer-dependent small RNAs derived from tRNAs. Rna, 2009, 15, 2147-2160.	3.5	525
29	System-Wide Changes to SUMO Modifications in Response to Heat Shock. Science Signaling, 2009, 2, ra24.	3.6	415
30	Side-chain conformational entropy at protein-protein interfaces. Protein Science, 2009, 11, 2860-2870.	7.6	71
31	The Jpred 3 secondary structure prediction server. Nucleic Acids Research, 2008, 36, W197-W201.	14.5	1,308
32	Aminoimidazolylmethyluracil Analogues as Potent Inhibitors of Thymidine Phosphorylase and Their Bioreductive Nitroimidazolyl Prodrugs. Journal of Medicinal Chemistry, 2005, 48, 392-402.	6.4	43
33	3-Substituted-5-aziridinyl-1-methylindole-4,7-diones as NQO1-directed antitumour agents: mechanism of activation and cytotoxicity in vitro. Biochemical Pharmacology, 2003, 66, 1199-1206.	4.4	16
34	Potential Tumor-Selective Nitroimidazolylmethyluracil Prodrug Derivatives:  Inhibitors of the Angiogenic Enzyme Thymidine Phosphorylase. Journal of Medicinal Chemistry, 2003, 46, 207-209.	6.4	44