

Kyle R Taylor

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11548339/publications.pdf>

Version: 2024-02-01

14
papers

590
citations

1163117

8
h-index

1199594

12
g-index

14
all docs

14
docs citations

14
times ranked

998
citing authors

#	ARTICLE	IF	CITATIONS
1	Vitronectin binding protein, BOM1093, confers serum resistance on <i>Borrelia miyamotoi</i> . <i>Scientific Reports</i> , 2021, 11, 5462.	3.3	2
2	A relapsing fever group <i>Borrelia</i> sp. is widely distributed among wild deer in Japan. <i>Ticks and Tick-borne Diseases</i> , 2018, 9, 465-470.	2.7	16
3	Suspected myelodysplastic/myeloproliferative neoplasm in a feline leukemia virus-negative cat. <i>Veterinary Clinical Pathology</i> , 2016, 45, 584-593.	0.7	6
4	Audioprofile Surfaces. <i>Annals of Otology, Rhinology and Laryngology</i> , 2016, 125, 361-368.	1.1	8
5	Surgical management of an odontogenic tumor in a banded Gila monster (<i>Heloderma suspectum</i>) Tj ETQq1 1 0.784314 rgBT /Overlo	0.7	8
6	Cordova: Web-based management of genetic variation data. <i>Bioinformatics</i> , 2014, 30, 3438-3439.	4.1	3
7	AudioGene: Predicting Hearing Loss Genotypes from Phenotypes to Guide Genetic Screening. <i>Human Mutation</i> , 2013, 34, n/a-n/a.	2.5	31
8	Advancing genetic testing for deafness with genomic technology. <i>Journal of Medical Genetics</i> , 2013, 50, 627-634.	3.2	104
9	Using the Phenome and Genome to Improve Genetic Diagnosis for Deafness. <i>Otolaryngology - Head and Neck Surgery</i> , 2012, 147, 975-977.	1.9	9
10	AudioGene: Computer-based prediction of genetic factors involved in non-syndromic hearing impairment. , 2011, , .		0
11	Sequencing and disease variation detection tools and techniques. , 2011, , .		1
12	DFNA8/12 caused by TECTA mutations is the most identified subtype of nonsyndromic autosomal dominant hearing loss. <i>Human Mutation</i> , 2011, 32, 825-834.	2.5	73
13	Comprehensive genetic testing for hereditary hearing loss using massively parallel sequencing. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 21104-21109.	7.1	294
14	A Contemporary Review of AudioGene audioprofiling: A machine-based candidate gene prediction tool for autosomal dominant nonsyndromic hearing loss. <i>Laryngoscope</i> , 2009, 119, 2211-2215.	2.0	35