

Kyle R Taylor

List of Publications by Year in descending order

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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	Comprehensive genetic testing for hereditary hearing loss using massively parallel sequencing. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 21104-21109.	7.1	294
2	Advancing genetic testing for deafness with genomic technology. Journal of Medical Genetics, 2013, 50, 627-634.	3.2	104
3	DFNA8/12 caused by TECTA mutations is the most identified subtype of nonsyndromic autosomal dominant hearing loss. Human Mutation, 2011, 32, 825-834.	2.5	73
4	A Contemporary Review of AudioGene audioprofiling: A machine-based candidate gene prediction tool for autosomal dominant nonsyndromic hearing loss. Laryngoscope, 2009, 119, 2211-2215.	2.0	35
5	AudioGene: Predicting Hearing Loss Genotypes from Phenotypes to Guide Genetic Screening. Human Mutation, 2013, 34, n/a-n/a.	2.5	31
6	A relapsing fever group Borrelia sp. is widely distributed among wild deer in Japan. Ticks and Tick-borne Diseases, 2018, 9, 465-470.	2.7	16
7	Using the Phenome and Genome to Improve Genetic Diagnosis for Deafness. Otolaryngology - Head and Neck Surgery, 2012, 147, 975-977.	1.9	9
8	Audioprofile Surfaces. Annals of Otology, Rhinology and Laryngology, 2016, 125, 361-368.	1.1	8
9	Surgical management of an odontogenic tumor in a banded Gila monster (<i>Heloderma suspectum</i>) Tj ETQq1 1 0.784314 rgBT /Over 6.7	0.7	6
10	Suspected myelodysplastic/myeloproliferative neoplasm in a feline leukemia virus-negative cat. Veterinary Clinical Pathology, 2016, 45, 584-593.	0.7	6
11	Cordova: Web-based management of genetic variation data. Bioinformatics, 2014, 30, 3438-3439.	4.1	3
12	Vitronectin binding protein, BOM1093, confers serum resistance on Borrelia miyamotoi. Scientific Reports, 2021, 11, 5462.	3.3	2
13	Sequencing and disease variation detection tools and techniques. , 2011, , .		1
14	AudioGene: Computer-based prediction of genetic factors involved in non-syndromic hearing impairment. , 2011, , .		0