

Takuji Masunaga

List of Publications by Year in descending order

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Version: 2024-02-01

9
papers

149
citations

1478505
6
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1474206
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g-index

9
all docs

9
docs citations

9
times ranked

180
citing authors

#	ARTICLE	IF	CITATIONS
1	Epidermal Basement Membrane: Its Molecular Organization and Blistering Disorders. <i>Connective Tissue Research</i> , 2006, 47, 55-66.	2.3	38
2	Differences in recurrent COL7A1 mutations in dystrophic epidermolysis bullosa: ethnic-specific and worldwide recurrent mutations. <i>Archives of Dermatological Research</i> , 2004, 295, 442-447.	1.9	33
3	Recurrent COL7A1 Mutations in Japanese Patients with Dystrophic Epidermolysis Bullosa: Positional Effects of Premature Termination Codon Mutations on Clinical Severity. <i>Journal of Investigative Dermatology</i> , 1999, 112, 991-993.	0.7	32
4	Pyloric atresia-junctional epidermolysis bullosa syndrome showing novel 594insC/Q425P mutations in integrin beta4 gene (ITGB4). <i>Experimental Dermatology</i> , 2004, 13, 61-64.	2.9	13
5	Genotype-phenotype correlations in six Japanese patients with recessive dystrophic epidermolysis bullosa with the recurrent p.Glu2857X mutation. <i>Journal of Dermatological Science</i> , 2008, 52, 13-20.	1.9	12
6	Compound heterozygosity for novel splice site mutations of <i>ITGA6</i> in lethal junctional epidermolysis bullosa with pyloric atresia. <i>Journal of Dermatology</i> , 2017, 44, 160-166.	1.2	9
7	Splicing abnormality of integrin β 4 gene (ITGB4) due to nucleotide substitutions far from splice site underlies pyloric atresia-junctional epidermolysis bullosa syndrome. <i>Journal of Dermatological Science</i> , 2015, 78, 61-66.	1.9	6
8	Splice site mutation in <i>COL7A1</i> resulting in aberrant in-frame transcripts identified in a case of recessive dystrophic epidermolysis bullosa, pretibial. <i>Journal of Dermatology</i> , 2018, 45, 742-745.	1.2	4
9	Japanese recurrent mutation c.6216+5G>T in COL7A1 leads to a mild phenotype of dystrophic epidermolysis bullosa. <i>Journal of Dermatological Science</i> , 2015, 80, 220-223.	1.9	2