Celine V Cluzeau

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Gastrointestinal Tract Pathology in a BALB/c Niemann–Pick Disease Type C1 Null Mouse Model. Digestive Diseases and Sciences, 2018, 63, 870-880.	2.3	10
2	Modeling Niemann-Pick disease type C1 in zebrafish: a robust platform for <i>in vivo</i> screening of candidate therapeutic compounds. DMM Disease Models and Mechanisms, 2018, 11, .	2.4	38
3	Modeling Smith-Lemli-Opitz syndrome with induced pluripotent stem cells reveals a causal role for Wnt/β-catenin defects in neuronal cholesterol synthesis phenotypes. Nature Medicine, 2016, 22, 388-396.	30.7	46
4	Human and mouse neuroinflammation markers in Niemannâ€Pick disease, type C1. Journal of Inherited Metabolic Disease, 2014, 37, 83-92.	3.6	71
5	Microarray expression analysis and identification of serum biomarkers for Niemann–Pick disease, type C1. Human Molecular Genetics, 2012, 21, 3632-3646.	2.9	84
6	Quantitative Proteomic Analysis of Niemann-Pick Disease, Type C1 Cerebellum Identifies Protein Biomarkers and Provides Pathological Insight. PLoS ONE, 2012, 7, e47845.	2.5	59
7	Interleukin-36–Receptor Antagonist Deficiency and Generalized Pustular Psoriasis. New England Journal of Medicine, 2011, 365, 620-628.	27.0	836
8	Only four genes (EDA1, EDAR, EDARADD, and WNT10A) account for 90% of hypohidrotic/anhidrotic ectodermal dysplasia cases. Human Mutation, 2011, 32, 70-72.	2.5	240
9	Mutations in <i>EDARADD</i> account for a small proportion of hypohidrotic ectodermal dysplasia cases. British Journal of Dermatology, 2010, 162, 1044-1048.	1.5	27
10	Unusual presentation of a severe autosomal recessive anhydrotic ectodermal dysplasia with a novel mutation in the <i>EDAR</i> gene. American Journal of Medical Genetics, Part A, 2008, 146A, 2657-2662.	1.2	22
11	Autosomal dominant anhidrotic ectodermal dysplasias at the EDARADD locus. Human Mutation, 2007, 28, 703-709.	2.5	80
12	Father-to-daughter transmission of Cornelia de Lange syndrome caused by a mutation in the 5′ untranslated region of theNIPBL Gene. Human Mutation, 2006, 27, 731-735.	2.5	58