Kati M Donner

List of Publications by Year
in descending order

34	1,833	331670 21	377865 34
papers	citations	h-index	g-index

34	34	34	1808
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Pharmacoepigenetics of hypertension: genome-wide methylation analysis of responsiveness to four classes of antihypertensive drugs using a double-blind crossover study design. Epigenetics, 2022, , 1-14.	2.7	7
2	Missense mutations in small muscle protein X-linked (SMPX) cause distal myopathy with protein inclusions. Acta Neuropathologica, 2021, 142, 375-393.	7.7	6
3	Dominant Distal Myopathy 3 (MPD3) Caused by a Deletion in the HNRNPA1 Gene. Neurology: Genetics, 2021, 7, e632.	1.9	7
4	Genomics of asthma, allergy and chronic rhinosinusitis: novel concepts and relevance in airway mucosa. Clinical and Translational Allergy, 2020, 10, 45.	3.2	26
5	Heme oxygenase-1 repeat polymorphism in septic acute kidney injury. PLoS ONE, 2019, 14, e0217291.	2.5	16
6	Replicated evidence for aminoacylase 3 and nephrin gene variations to predict antihypertensive drug responses. Pharmacogenomics, 2017, 18, 445-458.	1.3	18
7	PRKCDBP (CAVIN3) and CRY2 associate with major depressive disorder. Journal of Affective Disorders, 2017, 207, 136-140.	4.1	20
8	CRY1 and CRY2 genetic variants in seasonality: A longitudinal and cross-sectional study. Psychiatry Research, 2016, 242, 101-110.	3.3	10
9	PTPRD gene associated with blood pressure response to atenolol and resistant hypertension. Journal of Hypertension, 2015, 33, 2278-2285.	0.5	38
10	TET2 and CSMD1 genes affect SBP response to hydrochlorothiazide in never-treated essential hypertensives. Journal of Hypertension, 2015, 33, 1301-1309.	0.5	29
11	Genetic Loci Associated with Allergic Sensitization in Lithuanians. PLoS ONE, 2015, 10, e0134188.	2.5	4
12	SIRT1 Polymorphisms Associate with Seasonal Weight Variation, Depressive Disorders, and Diastolic Blood Pressure in the General Population. PLoS ONE, 2015, 10, e0141001.	2.5	23
13	Pharmacogenomics of Hypertension: A Genomeâ€Wide, Placeboâ€Controlled Crossâ€Over Study, Using Four Classes of Antihypertensive Drugs. Journal of the American Heart Association, 2015, 4, e001521.	3.7	74
14	CRY1, CRY2 and PRKCDBP genetic variants in metabolic syndrome. Hypertension Research, 2015, 38, 186-192.	2.7	35
15	Mutation Update: The Spectra of Nebulin Variants and Associated Myopathies. Human Mutation, 2014, 35, 1418-1426.	2.5	107
16	Generalized glucocorticoid resistance caused by a novel two-nucleotide deletion in the hormone-binding domain of the glucocorticoid receptor gene NR3C1. European Journal of Endocrinology, 2013, 168, K9-K18.	3.7	18
17	Genomic Association Analysis of Common Variants Influencing Antihypertensive Response to Hydrochlorothiazide. Hypertension, 2013, 62, 391-397.	2.7	96
18	CRY2 Genetic Variants Associate with Dysthymia. PLoS ONE, 2013, 8, e71450.	2.5	42

#	Article	IF	Citations
19	STK39 variation predicts the ambulatory blood pressure response to losartan in hypertensive men. Hypertension Research, 2012, 35, 107-114.	2.7	21
20	Abnormal actin binding of aberrant Î²-tropomyosins is a molecular cause of muscle weakness in TPM2-related nemaline and cap myopathy. Biochemical Journal, 2012, 442, 231-239.	3.7	48
21	Common genetic variation of Î²1- and Î²2-adrenergic receptor and response to four classes of antihypertensive treatment. Pharmacogenetics and Genomics, 2010, 20, 342-345.	1.5	33
22	Reninâ€“Angiotensin System and Î±-Adducin Gene Polymorphisms and Their Relation to Responses to Antihypertensive Drugs: Results From the GENRES Study. American Journal of Hypertension, 2009, 22, 169-175.	2.0	37
23	CYP2C9 genotype modifies activity of the reninâ€“angiotensinâ€“aldosterone system in hypertensive men. Journal of Hypertension, 2009, 27, 2001-2009.	0.5	16
24	Identification of a founder mutation in TPM3 in nemaline myopathy patients of Turkish origin. European Journal of Human Genetics, 2008, 16, 1055-1061.	2.8	36
25	Laboratory tests as predictors of the antihypertensive effects of amlodipine, bisoprolol, hydrochlorothiazide and losartan in men: results from the randomized, double-blind, crossover GENRES Study. Journal of Hypertension, 2008, 26, 1250-1256.	0.5	29
26	Developmental and muscle-type-specific expression of mouse nebulin exons 127 and 128. Genomics, 2006, 88, 489-495.	2.9	20
27	Identification of 45 novel mutations in the nebulin gene associated with autosomal recessive nemaline myopathy. Human Mutation, 2006, 27, 946-956.	2.5	112
28	Complete genomic structure of the human nebulin gene and identification of alternatively spliced transcripts. European Journal of Human Genetics, 2004, 12, 744-751.	2.8	101
29	Mutations in the Î²-tropomyosin (TPM2) gene â€“ a rare cause of nemaline myopathy. Neuromuscular Disorders, 2002, 12, 151-158.	0.6	210
30	Mutations in the nebulin gene can cause severe congenital nemaline myopathy. Neuromuscular Disorders, 2002, 12, 674-679.	0.6	85
31	Nebulin mutations in autosomal recessive nemaline myopathy: an update. Neuromuscular Disorders, 2002, 12, 680-686.	0.6	55
32	Nebulin expression in patients with nemaline myopathy. Neuromuscular Disorders, 2001, 11, 154-162.	0.6	39
33	Mutations in the skeletal muscle Î±-actin gene in patients with actin myopathy and nemaline myopathy. Nature Genetics, 1999, 23, 208-212.	21.4	389
34	Refined Localisation of the Genes for Nebulin and Titin on Chromosome 2q Allows the Assignment of Nebulin as a Candidate Gene for Autosomal Recessive Nemaline Myopathy. European Journal of Human Genetics, 1997, 5, 229-234.	2.8	26