

Jeffrey Bryan Carroll

List of Publications by Year in descending order

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Version: 2024-02-01

31
papers

2,391
citations

331670

21
h-index

434195

31
g-index

39
all docs

39
docs citations

39
times ranked

3236
citing authors

#	ARTICLE	IF	CITATIONS
1	Dynamics of huntingtin protein interactions in the striatum identifies candidate modifiers of Huntington disease. <i>Cell Systems</i> , 2022, 13, 304-320.e5.	6.2	15
2	Stool is a sensitive and noninvasive source of DNA for monitoring expansion in repeat expansion disease mouse models. <i>DMM Disease Models and Mechanisms</i> , 2022, 15, .	2.4	1
3	Single-Nucleus RNA-Seq Reveals Dysregulation of Striatal Cell Identity Due to Huntington's Disease Mutations. <i>Journal of Neuroscience</i> , 2021, 41, 5534-5552.	3.6	30
4	Huntingtin structure is orchestrated by HAP40 and shows a polyglutamine expansion-specific interaction with exon 1. <i>Communications Biology</i> , 2021, 4, 1374.	4.4	22
5	Prion protein lowering is a disease-modifying therapy across prion disease stages, strains and endpoints. <i>Nucleic Acids Research</i> , 2020, 48, 10615-10631.	14.5	69
6	Antisense oligonucleotides extend survival of prion-infected mice. <i>JCI Insight</i> , 2019, 4, .	5.0	80
7	HttQ111/+ Huntingtonâ€™s Disease Knock-in Mice Exhibit Brain Region-Specific Morphological Changes and Synaptic Dysfunction. <i>Journal of Huntington's Disease</i> , 2018, 7, 17-33.	1.9	27
8	Early Detection of Apathetic Phenotypes in Huntingtonâ€™s Disease Knock-in Mice Using Open Source Tools. <i>Scientific Reports</i> , 2018, 8, 2304.	3.3	6
9	Transcriptional regulatory networks underlying gene expression changes in Huntington's disease. <i>Molecular Systems Biology</i> , 2018, 14, e7435.	7.2	55
10	Motivational, proteostatic and transcriptional deficits precede synapse loss, gliosis and neurodegeneration in the B6.HttQ111/+ model of Huntingtonâ€™s disease. <i>Scientific Reports</i> , 2017, 7, 41570.	3.3	16
11	High resolution time-course mapping of early transcriptomic, molecular and cellular phenotypes in Huntingtonâ€™s disease CAG knock-in mice across multiple genetic backgrounds. <i>Human Molecular Genetics</i> , 2017, 26, 913-922.	2.9	37
12	Peripheral huntingtin silencing does not ameliorate central signs of disease in the B6.HttQ111/+ mouse model of Huntingtonâ€™s disease. <i>PLoS ONE</i> , 2017, 12, e0175968.	2.5	13
13	HdhQ111 Mice Exhibit Tissue Specific Metabolite Profiles that Include Striatal Lipid Accumulation. <i>PLoS ONE</i> , 2015, 10, e0134465.	2.5	20
14	Treating the whole body in Huntington's disease. <i>Lancet Neurology</i> , The, 2015, 14, 1135-1142.	10.2	126
15	In Vivo Evaluation of Candidate Allele-specific Mutant Huntingtin Gene Silencing Antisense Oligonucleotides. <i>Molecular Therapy</i> , 2014, 22, 2093-2106.	8.2	115
16	Allele-Specific Suppression of Mutant Huntingtin Using Antisense Oligonucleotides: Providing a Therapeutic Option for All Huntington Disease Patients. <i>PLoS ONE</i> , 2014, 9, e107434.	2.5	92
17	A fully humanized transgenic mouse model of Huntington disease. <i>Human Molecular Genetics</i> , 2013, 22, 18-34.	2.9	93
18	HDBuzz: empowering patients through accessible education. <i>Trends in Molecular Medicine</i> , 2012, 18, 1-3.	6.7	6

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19	Potent and Selective Antisense Oligonucleotides Targeting Single-Nucleotide Polymorphisms in the Huntington Disease Gene / Allele-Specific Silencing of Mutant Huntingtin. <i>Molecular Therapy</i> , 2011, 19, 2178-2185.	8.2	246
20	Natural history of disease in the YAC128 mouse reveals a discrete signature of pathology in Huntington disease. <i>Neurobiology of Disease</i> , 2011, 43, 257-265.	4.4	65
21	Mice lacking caspase-2 are protected from behavioral changes, but not pathology, in the YAC128 model of Huntington disease. <i>Molecular Neurodegeneration</i> , 2011, 6, 59.	10.8	30
22	Wild-type HTT modulates the enzymatic activity of the neuronal palmitoyl transferase HIP14. <i>Human Molecular Genetics</i> , 2011, 20, 3356-3365.	2.9	71
23	Cholesterol Defect Is Marked across Multiple Rodent Models of Huntington's Disease and Is Manifest in Astrocytes. <i>Journal of Neuroscience</i> , 2010, 30, 10844-10850.	3.6	136
24	Age-Dependent Alterations of Corticostriatal Activity in the YAC128 Mouse Model of Huntington Disease. <i>Journal of Neuroscience</i> , 2009, 29, 2414-2427.	3.6	160
25	CAG Expansion in the Huntington Disease Gene Is Associated with a Specific and Targetable Predisposing Haplogroup. <i>American Journal of Human Genetics</i> , 2009, 84, 351-366.	6.2	204
26	Automated deformation analysis in the YAC128 Huntington disease mouse model. <i>NeuroImage</i> , 2008, 39, 32-39.	4.2	150
27	Cortical thickness measured from MRI in the YAC128 mouse model of Huntington's disease. <i>NeuroImage</i> , 2008, 41, 243-251.	4.2	115
28	Activated caspase-6 and caspase-6-cleaved fragments of huntingtin specifically colocalize in the nucleus. <i>Human Molecular Genetics</i> , 2008, 17, 2390-2404.	2.9	112
29	Cholesterol biosynthesis pathway is disturbed in YAC128 mice and is modulated by huntingtin mutation. <i>Human Molecular Genetics</i> , 2007, 16, 2187-2198.	2.9	106
30	Phenotypic abnormalities in the YAC128 mouse model of Huntington disease are penetrant on multiple genetic backgrounds and modulated by strain. <i>Neurobiology of Disease</i> , 2007, 26, 189-200.	4.4	97
31	Parvoviral nuclear import: bypassing the host nuclear-transport machinery. <i>Journal of General Virology</i> , 2006, 87, 3209-3213.	2.9	54