

Thomas B Hansen

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

13,582
citations

489802

18
h-index

759306

22
g-index

25
all docs

25
docs citations

25
times ranked

13556
citing authors

#	ARTICLE	IF	CITATIONS
1	Best practice standards for circular RNA research. <i>Nature Methods</i> , 2022, 19, 1208-1220.	9.0	58
2	Profiling of Plasma Extracellular Vesicle Transcriptome Reveals That circRNAs Are Prevalent and Differ between Multiple Sclerosis Patients and Healthy Controls. <i>Biomedicines</i> , 2021, 9, 1850.	1.4	8
3	RNA-Seq profiling of leukocytes reveals a sex-dependent global circular RNA upregulation in multiple sclerosis and 6 candidate biomarkers. <i>Human Molecular Genetics</i> , 2020, 29, 3361-3372.	1.4	21
4	Biosynthesis of Circular RNA ciRS-7/CDR1as Is Mediated by Mammalian-wide Interspersed Repeats. <i>IScience</i> , 2020, 23, 101345.	1.9	25
5	Spatial expression analyses of the putative oncogene ciRS-7 in cancer reshape the microRNA sponge theory. <i>Nature Communications</i> , 2020, 11, 4551.	5.8	72
6	circZNF827 nucleates a transcription inhibitory complex to balance neuronal differentiation. <i>ELife</i> , 2020, 9, .	2.8	33
7	The biogenesis, biology and characterization of circular RNAs. <i>Nature Reviews Genetics</i> , 2019, 20, 675-691.	7.7	2,832
8	CircSMARCA5 Regulates VEGFA mRNA Splicing and Angiogenesis in Glioblastoma Multiforme Through the Binding of SRSF1. <i>Cancers</i> , 2019, 11, 194.	1.7	146
9	Noncoding AUG circRNAs constitute an abundant and conserved subclass of circles. <i>Life Science Alliance</i> , 2019, 2, e201900398.	1.3	56
10	Characterization of Circular RNA Concatemers. <i>Methods in Molecular Biology</i> , 2018, 1724, 143-157.	0.4	7
11	Enhanced Tailored MicroRNA Sponge Activity of RNA Pol II-Transcribed TuD Hairpins Relative to Ectopically Expressed ciRS7-Derived circRNAs. <i>Molecular Therapy - Nucleic Acids</i> , 2018, 13, 365-375.	2.3	10
12	Improved circRNA Identification by Combining Prediction Algorithms. <i>Frontiers in Cell and Developmental Biology</i> , 2018, 6, 20.	1.8	135
13	Insights into circular RNA biology. <i>RNA Biology</i> , 2017, 14, 1035-1045.	1.5	362
14	The invasion of circRNAs. <i>RNA Biology</i> , 2017, 14, 973-974.	1.5	1
15	Comparison of circular RNA prediction tools. <i>Nucleic Acids Research</i> , 2016, 44, e58-e58.	6.5	349
16	Circular RNAs: Identification, biogenesis and function. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2016, 1859, 163-168.	0.9	469
17	Spatio-temporal regulation of circular RNA expression during porcine embryonic brain development. <i>Genome Biology</i> , 2015, 16, 245.	3.8	422
18	Circular RNA and miR-7 in Cancer. <i>Cancer Research</i> , 2013, 73, 5609-5612.	0.4	847

#	ARTICLE	IF	CITATIONS
19	Natural RNA circles function as efficient microRNA sponges. <i>Nature</i> , 2013, 495, 384-388.	13.7	6,415
20	RNA Interference Pathways and Therapeutic Exploitation. <i>Advances in Delivery Science and Technology</i> , 2013, , 1-29.	0.4	0
21	miRNA-dependent gene silencing involving Ago2-mediated cleavage of a circular antisense RNA. <i>EMBO Journal</i> , 2011, 30, 4414-4422.	3.5	841
22	A screen of chemical modifications identifies position-specific modification by UNA to most potently reduce siRNA off-target effects. <i>Nucleic Acids Research</i> , 2010, 38, 5761-5773.	6.5	157
23	A large-scale chemical modification screen identifies design rules to generate siRNAs with high activity, high stability and low toxicity. <i>Nucleic Acids Research</i> , 2009, 37, 2867-2881.	6.5	315