

Ryan N Doan

List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

1,590
citations

623734

14
h-index

580821

25
g-index

26
all docs

26
docs citations

26
times ranked

3547
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Rates and Patterns of Clonal Oncogenic Mutations in the Normal Human Brain. <i>Cancer Discovery</i> , 2022, 12, 172-185. | 9.4 | 19 |
| 2 | The landscape of somatic mutation in cerebral cortex of autistic and neurotypical individuals revealed by ultra-deep whole-genome sequencing. <i>Nature Neuroscience</i> , 2021, 24, 176-185. | 14.8 | 73 |
| 3 | MIPP-Seq: ultra-sensitive rapid detection and validation of low-frequency mosaic mutations. <i>BMC Medical Genomics</i> , 2021, 14, 47. | 1.5 | 12 |
| 4 | Landmarks of human embryonic development inscribed in somatic mutations. <i>Science</i> , 2021, 371, 1249-1253. | 12.6 | 65 |
| 5 | Comprehensive identification of somatic nucleotide variants in human brain tissue. <i>Genome Biology</i> , 2021, 22, 92. | 8.8 | 26 |
| 6 | Toward a better definition of focal cortical dysplasia: An iterative histopathological and genetic agreement trial. <i>Epilepsia</i> , 2021, 62, 1416-1428. | 5.1 | 54 |
| 7 | Rewiring of human neurodevelopmental gene regulatory programs by human accelerated regions. <i>Neuron</i> , 2021, 109, 3239-3251.e7. | 8.1 | 91 |
| 8 | Pathologic characterization of canine multiple system degeneration in the Ibizan hound. <i>Veterinary Pathology</i> , 2021, , 030098582110430. | 1.7 | 1 |
| 9 | Accurate detection of mosaic variants in sequencing data without matched controls. <i>Nature Biotechnology</i> , 2020, 38, 314-319. | 17.5 | 54 |
| 10 | Polymicrogyria is Associated With Pathogenic Variants in PTEN. <i>Annals of Neurology</i> , 2020, 88, 1153-1164. | 5.3 | 14 |
| 11 | Homozygous deletions implicate non-coding epigenetic marks in Autism spectrum disorder. <i>Scientific Reports</i> , 2020, 10, 14045. | 3.3 | 12 |
| 12 | Recessive gene disruptions in autism spectrum disorder. <i>Nature Genetics</i> , 2019, 51, 1092-1098. | 21.4 | 109 |
| 13 | Identification of a Candidate Mutation in the COL1A2 Gene of a Chow Chow With Osteogenesis Imperfecta. <i>Journal of Heredity</i> , 2018, 109, 308-314. | 2.4 | 13 |
| 14 | Postnatal changes in epigenetic modifications of neutrophils of foals are associated with increased ROS function and regulation of neutrophil function. <i>Developmental and Comparative Immunology</i> , 2018, 87, 182-187. | 2.3 | 6 |
| 15 | Evolutionary Changes in Transcriptional Regulation: Insights into Human Behavior and Neurological Conditions. <i>Annual Review of Neuroscience</i> , 2018, 41, 185-206. | 10.7 | 18 |
| 16 | Sodium Channel SCN3A (NaV1.3) Regulation of Human Cerebral Cortical Folding and Oral Motor Development. <i>Neuron</i> , 2018, 99, 905-913.e7. | 8.1 | 109 |
| 17 | Co-segregation of sex chromosomes in the male black widow spider <i>Latrodectus mactans</i> (Araneae, Tj ETQq1 1 0.784314 rgBT /Over 2.2 83 | | |
| 18 | Identification of a novel CNTNAP1 mutation causing arthrogyrosis multiplex congenita with cerebral and cerebellar atrophy. <i>European Journal of Medical Genetics</i> , 2017, 60, 245-249. | 1.3 | 20 |

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|----|--|------|-----------|
| 19 | Cell-Type-Specific Alternative Splicing Governs Cell Fate in the Developing Cerebral Cortex. <i>Cell</i> , 2016, 166, 1147-1162.e15. | 28.9 | 276 |
| 20 | Mutations in Human Accelerated Regions Disrupt Cognition and Social Behavior. <i>Cell</i> , 2016, 167, 341-354.e12. | 28.9 | 280 |
| 21 | Nutritional programming of accelerated puberty in heifers: alterations in DNA methylation in the arcuate nucleus ^{<xref ref-type="fn" rid="afn1">â€‹</xref>} , ^{<xref ref-type="fn" rid="afn2">â€‹</xref>} . <i>Biology of Reproduction</i> , 2016, 96, 174-184. | 2.7 | 14 |
| 22 | Single-cell analysis reveals transcriptional heterogeneity of neural progenitors in human cortex. <i>Nature Neuroscience</i> , 2015, 18, 637-646. | 14.8 | 247 |
| 23 | Identification of Genomic Loci Associated with <i>Rhodococcus equi</i> Susceptibility in Foals. <i>PLoS ONE</i> , 2014, 9, e98710. | 2.5 | 11 |
| 24 | Identification of copy number variants in horses. <i>Genome Research</i> , 2012, 22, 899-907. | 5.5 | 49 |