# AmÃ©lie C Pinard

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/8082504/publications.pdf

Version: 2024-02-01

| 18 | 890 | 840585 11 | 887953 17 |
|---|---|---|---|
| papers | citations | h-index | g-index |

| 20 | 20 | 20 | 2042 |
|---|---|---|---|
| all docs | docs citations | times ranked | citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | Genetics of Thoracic and Abdominal Aortic Diseases. Circulation Research, 2019, 124, 588-606. | 2.0 | 253 |
| 2 | Comparative genomics of emerging pathogens in the Candida glabrata clade. BMC Genomics, 2013, 14, 623. | 1.2 | 174 |
| 3 | UMDâ€Predictor: A Highâ€Throughput Sequencing Compliant System for Pathogenicity Prediction of any Human cDNA Substitution. Human Mutation, 2016, 37, 439-446. | 1.1 | 104 |
| 4 | A genome-wide approach reveals novel imprinted genes expressed in the human placenta. Epigenetics, 2012, 7, 1079-1090. | 1.3 | 81 |
| 5 | LTBP3 Pathogenic Variants Predispose Individuals to Thoracic Aortic Aneurysms and Dissections. American Journal of Human Genetics, 2018, 102, 706-712. | 2.6 | 51 |
| 6 | Piezo1 is required for outflow tract and aortic valve development.. Journal of Molecular and Cellular Cardiology, 2020, 143, 51-62. | 0.9 | 44 |
| 7 | The revised ghent nosology; reclassifying isolated ectopia lentis. Clinical Genetics, 2015, 87, 284-287. | 1.0 | 41 |
| 8 | The pleiotropy associated with de novo variants in CHD4, CNOT3, and SETD5 extends to moyamoya angiopathy. Genetics in Medicine, 2020, 22, 427-431. | 1.1 | 34 |
| 9 | DIAPH1 Variants in Nonâ€“East Asian Patients With Sporadic Moyamoya Disease. JAMA Neurology, 2021, 78, 993. | 4.5 | 33 |
| 10 | Association of De Novo RNF213 Variants With Childhood Onset Moyamoya Disease and Diffuse Occlusive Vasculopathy. Neurology, 2021, 96, e1783-e1791. | 1.5 | 21 |
| 11 | Rare deleterious variants of NOTCH1, GATA4, SMAD6, and ROBO4 are enriched in BAV with early onset complications but not in BAV with heritable thoracic aortic disease. Molecular Genetics & Genomic Medicine, 2020, 8, e1406. | 0.6 | 17 |
| 12 | Update on the genetic risk for thoracic aortic aneurysms and acute aortic dissections: implications for clinical care. Journal of Cardiovascular Surgery, 2021, 62, 203-210. | 0.3 | 11 |
| 13 | A genome-wide search for new imprinted genes in the human placenta identifies DSCAM as the first imprinted gene on chromosome 21. European Journal of Human Genetics, 2019, 27, 49-60. | 1.4 | 8 |
| 14 | Actionable Genes, Core Databases, and Locus-Specific Databases. Human Mutation, 2016, 37, 1299-1307. | 1.1 | 6 |
| 15 | WES/WGS Reporting of Mutations from Cardiovascular â€œActionableâ€•Genes in Clinical Practice: A Key Role for UMD Knowledgebases in the Era of Big Databases. Human Mutation, 2016, 37, 1308-1317. | 1.1 | 5 |
| 16 | An uncommon cause of tricuspid regurgitation: three-dimensional echocardiographic incremental value, surgical and genetic insights. European Journal of Cardio-thoracic Surgery, 2016, 50, 180-182. | 0.6 | 1 |
| 17 | Preventing Acute Aortic Dissections: The Power of Familial Screening and Risk Assessment. Journal of the American Heart Association, 2022, 11, e025441. | 1.6 | 1 |
| 18 | Analysis of HOXB1 gene in a cohort of patients with sporadic ventricular septal defect. Molecular Biology Reports, 2018, 45, 1507-1513. | 1.0 | 0 |