

Ashim Mukherjee

List of Publications by Year in descending order

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Version: 2024-02-01

34
papers

1,031
citations

840776

11
h-index

434195

31
g-index

34
all docs

34
docs citations

34
times ranked

1900
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Regulation of Notch signaling by E3 ubiquitin ligases. FEBS Journal, 2022, 289, 937-954. | 4.7 | 18 |
| 2 | Whole exome sequencing identifies a novel splice-site mutation in IMPG2 gene causing Stargardt-like juvenile macular dystrophy in a north Indian family. Gene, 2022, 816, 146158. | 2.2 | 3 |
| 3 | Somatic Clonal Analyses Using FLP/FRT and MARCM System to Understand Notch Signaling Mechanism and Its Regulation. Methods in Molecular Biology, 2022, , 83-94. | 0.9 | 1 |
| 4 | Deltex cooperates with TRAF6 to promote apoptosis and cell migration through Eiger-independent JNK activation in Drosophila. Cell Biology International, 2021, 45, 686-700. | 3.0 | 10 |
| 5 | Deltex positively regulates Toll signaling in a JNK independent manner in Drosophila. Genes To Cells, 2021, 26, 254-263. | 1.2 | 4 |
| 6 | Maheshvara regulates JAK/STAT signaling by interacting and stabilizing hopscotch transcripts which leads to apoptosis in Drosophila melanogaster. Cell Death and Disease, 2021, 12, 363. | 6.3 | 2 |
| 7 | Signaling cross-talk during development: Context-specific networking of Notch, NF- κ B and JNK signaling pathways in Drosophila. Cellular Signalling, 2021, 82, 109937. | 3.6 | 9 |
| 8 | A Forward Genetic Approach to Mapping a P-Element Second Site Mutation Identifies DCP2 as a Novel Tumor Suppressor in Drosophila melanogaster. G3: Genes, Genomes, Genetics, 2020, 10, 2601-2618. | 1.8 | 0 |
| 9 | Maheshvara, a Conserved RNA Helicase, Regulates Notch Signaling in Drosophila melanogaster. Advances in Experimental Medicine and Biology, 2020, 1227, 69-79. | 1.6 | 2 |
| 10 | Regulation of Notch Signaling in Drosophila melanogaster: The Role of the Heterogeneous Nuclear Ribonucleoprotein Hrp48 and Deltex. Advances in Experimental Medicine and Biology, 2020, 1227, 95-105. | 1.6 | 6 |
| 11 | Regulation of notch signaling by a chromatin modeling protein Hat-trick. Development (Cambridge), 2019, 146, . | 2.5 | 7 |
| 12 | Synergistic interaction of Deltex and Hrp48 leads to JNK activation. Cell Biology International, 2019, 43, 350-357. | 3.0 | 5 |
| 13 | Notch Signaling: From Neurogenesis to Neurodegeneration. , 2019, , 185-221. | | 0 |
| 14 | Pleiotropic Functions of the Chromodomain-Containing Protein Hat-trick During Oogenesis in Drosophila melanogaster. G3: Genes, Genomes, Genetics, 2018, 8, 1067-1077. | 1.8 | 5 |
| 15 | TP53 codon 72 polymorphism and the risk of glaucoma in a north Indian cohort: A genetic association study. Ophthalmic Genetics, 2018, 39, 228-235. | 1.2 | 4 |
| 16 | Notch signals modulate Igl mediated tumorigenesis by the activation of JNK signaling. BMC Research Notes, 2018, 11, 247. | 1.4 | 7 |
| 17 | A loss-of-function homozygous mutation in DDX59 implicates a conserved DEAD-box RNA helicase in nervous system development and function. Human Mutation, 2018, 39, 187-192. | 2.5 | 44 |
| 18 | Regulation of Notch signaling in the developing Drosophila eye by a T-box containing transcription factor, Dorsocross. Genesis, 2018, 56, e23251. | 1.6 | 6 |

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 19 | Deltex interacts with Eiger and consequently influences the cell death in <i>Drosophila melanogaster</i> . <i>Cellular Signalling</i> , 2018, 49, 17-29. | 3.6 | 14 |
| 20 | Regulation of Notch Signaling by the Heterogeneous Nuclear Ribonucleoprotein Hrp48 and Deltex in <i>Drosophila melanogaster</i> . <i>Genetics</i> , 2017, 206, 905-918. | 2.9 | 20 |
| 21 | Whole exome sequencing: Uncovering causal genetic variants for ocular diseases. <i>Experimental Eye Research</i> , 2017, 164, 139-150. | 2.6 | 20 |
| 22 | Interaction of Spoonbill with Prospero in <i>Drosophila</i> : Implications in neuroblast development. <i>Genesis</i> , 2017, 55, e23049. | 1.6 | 2 |
| 23 | Whole exome sequencing unveils a frameshift mutation in CNGB3 for cone dystrophy. <i>Medicine (United States)</i> , 2017, 96, e7490. | 1.0 | 8 |
| 24 | The RNA binding KH domain of Spoonbill depletes pathogenic non-coding spinocerebellar ataxia 8 transcripts and suppresses neurodegeneration in <i>Drosophila</i> . <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2016, 1862, 1732-1741. | 3.8 | 7 |
| 25 | Kinase active Misshapen regulates Notch signaling in <i>Drosophila melanogaster</i> . <i>Experimental Cell Research</i> , 2015, 339, 51-60. | 2.6 | 7 |
| 26 | Regulation of Notch Signaling by an Evolutionary Conserved DEAD Box RNA Helicase, Maheshvara in <i>Drosophila melanogaster</i> . <i>Genetics</i> , 2015, 201, 1071-1085. | 2.9 | 14 |
| 27 | A novel mutation in FRMD7 causes X-linked idiopathic congenital nystagmus in a North Indian family. <i>Neuroscience Letters</i> , 2015, 597, 170-175. | 2.1 | 9 |
| 28 | Chip physically interacts with Notch and their stoichiometry is critical for Notch function in wing development and cell proliferation in <i>Drosophila</i> . <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2015, 1850, 802-812. | 2.4 | 6 |
| 29 | TRAF6 is a novel regulator of Notch signaling in <i>Drosophila melanogaster</i> . <i>Cellular Signalling</i> , 2014, 26, 3016-3026. | 3.6 | 22 |
| 30 | dLin52 is crucial for dE2F and dRBF mediated transcriptional regulation of pro-apoptotic gene hid. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2014, 1839, 800-812. | 1.9 | 1 |
| 31 | MTHFR C677T Predisposes to POAG but Not to PACG in a North Indian Population: A Case Control Study. <i>PLoS ONE</i> , 2014, 9, e103063. | 2.5 | 17 |
| 32 | The <i>Drosophila</i> Importin- β 3 Is Required for Nuclear Import of Notch In Vivo and It Displays Synergistic Effects with Notch Receptor on Cell Proliferation. <i>PLoS ONE</i> , 2013, 8, e68247. | 2.5 | 29 |
| 33 | Regulation of Notch signalling by non-visual β 2-arrestin. <i>Nature Cell Biology</i> , 2005, 7, 1191-1201. | 10.3 | 213 |
| 34 | Protein interaction mapping: A <i>Drosophila</i> case study. <i>Genome Research</i> , 2005, 15, 376-384. | 5.5 | 509 |