Harikishore Amaravadhi

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

23 268 11 15 g-index

28 345 5 avg, IF L-index

| # | Paper | IF | Citations |
|----|---|------------------|-----------|
| 23 | Targeting Mycobacterial F-ATP Synthase C-Terminal (Subunit Interaction Motif on Rotary Subunit (<i>Antibiotics</i> , 2021 , 10, | 4.9 | 2 |
| 22 | Targeting C-terminal Helical bundle of NCOVID19 Envelope (E) protein. <i>International Journal of Biological Macromolecules</i> , 2021 , 175, 131-139 | 7.9 | 3 |
| 21 | A systematic assessment of mycobacterial F -ATPase subunit & role in latent ATPase hydrolysis. <i>FEBS Journal</i> , 2021 , 288, 818-836 | 5.7 | 6 |
| 20 | Targeting the menaquinol binding loop of mycobacterial cytochrome bd oxidase. <i>Molecular Diversity</i> , 2021 , 25, 517-524 | 3.1 | 7 |
| 19 | Antiviral activity against Middle East Respiratory Syndrome coronavirus by Montelukast, an anti-asthma drug. <i>Antiviral Research</i> , 2021 , 185, 104996 | 10.8 | 2 |
| 18 | Deciphering the Role of Ion Channels in Early Defense Signaling against Herbivorous Insects. <i>Cells</i> , 2021 , 10, | 7.9 | 1 |
| 17 | Discovery of a Novel Mycobacterial F-ATP Synthase Inhibitor and its Potency in Combination with Diarylquinolines. <i>Angewandte Chemie</i> , 2020 , 132, 13397-13406 | 3.6 | 3 |
| 16 | Discovery of a Novel Mycobacterial F-ATP Synthase Inhibitor and its Potency in Combination with Diarylquinolines. <i>Angewandte Chemie - International Edition</i> , 2020 , 59, 13295-13304 | 16.4 | 15 |
| 15 | Disrupting coupling within mycobacterial F-ATP synthases subunit Itauses dysregulated energy production and cell wall biosynthesis. <i>Scientific Reports</i> , 2019 , 9, 16759 | 4.9 | 22 |
| 14 | Combination of pharmacophore hypothesis and molecular docking to identify novel inhibitors of HCV NS5B polymerase. <i>Molecular Diversity</i> , 2015 , 19, 529-39 | 3.1 | 3 |
| 13 | Ursolic acid exerts anti-cancer activity by suppressing vaccinia-related kinase 1-mediated damage repair in lung cancer cells. <i>Scientific Reports</i> , 2015 , 5, 14570 | 4.9 | 24 |
| 12 | Immunophilins: Structures, Mechanisms and Ligands. Current Molecular Pharmacology, 2015 , 9, 37-47 | 3.7 | 25 |
| 11 | Luteolin suppresses cancer cell proliferation by targeting vaccinia-related kinase 1. <i>PLoS ONE</i> , 2014 , 9, e109655 | 3.7 | 29 |
| 10 | Crystal structure of Plasmodium vivax FK506-binding protein 25 reveals conformational changes responsible for its noncanonical activity. <i>Proteins: Structure, Function and Bioinformatics</i> , 2014 , 82, 1235 | 5-4 2 | 5 |
| 9 | Suprafenacine, an indazole-hydrazide agent, targets cancer cells through microtubule destabilization. <i>PLoS ONE</i> , 2014 , 9, e110955 | 3.7 | 8 |
| 8 | Revisiting de novo drug design: receptor based pharmacophore screening. <i>Current Topics in Medicinal Chemistry</i> , 2014 , 14, 1890-8 | 3 | 14 |
| 7 | Lipoxygenase Inhibitory Effects of Dibenzylbutane Lignans from the Seeds of Myristica fragrans (Nutmeg). <i>Bulletin of the Korean Chemical Society</i> , 2014 , 35, 3095-3098 | 1.2 | 1 |

LIST OF PUBLICATIONS

| 6 | Adamantyl derivative as a potent inhibitor of Plasmodium FK506 binding protein 35. <i>ACS Medicinal Chemistry Letters</i> , 2013 , 4, 1097-101 | 4.3 | 13 |
|---|--|-----|----|
| 5 | The flavonoid myricetin reduces nocturnal melatonin levels in the blood through the inhibition of serotonin N-acetyltransferase. <i>Biochemical and Biophysical Research Communications</i> , 2013 , 440, 312-6 | 3.4 | 13 |
| 4 | NMR solution structure of C2 domain of MFG-E8 and insights into its molecular recognition with phosphatidylserine. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2013 , 1828, 1083-93 | 3.8 | 18 |
| 3 | Small molecule Plasmodium FKBP35 inhibitor as a potential antimalaria agent. <i>Scientific Reports</i> , 2013 , 3, 2501 | 4.9 | 21 |
| 2 | Solution structure of FK506-binding protein 12 from Aedes aegypti. <i>Proteins: Structure, Function and Bioinformatics</i> , 2012 , 80, 2476-81 | 4.2 | 5 |
| 1 | Crystallographic structure of the tetratricopeptide repeat domain of Plasmodium falciparum FKBP35 and its molecular interaction with Hsp90 C-terminal pentapeptide. <i>Protein Science</i> , 2009 , 18, 2115-24 | 6.3 | 25 |