

Hideaki E Kato

List of Publications by Year in descending order

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Version: 2024-02-01

44
papers

4,182
citations

279798

23
h-index

345221

36
g-index

47
all docs

47
docs citations

47
times ranked

5533
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Structural insights into μ -opioid receptor activation. <i>Nature</i> , 2015, 524, 315-321. | 27.8 | 743 |
| 2 | Crystal structure of the channelrhodopsin light-gated cation channel. <i>Nature</i> , 2012, 482, 369-374. | 27.8 | 503 |
| 3 | Cortical layer-specific critical dynamics triggering perception. <i>Science</i> , 2019, 365, . | 12.6 | 447 |
| 4 | Structure of a Signaling Cannabinoid Receptor 1-G Protein Complex. <i>Cell</i> , 2019, 176, 448-458.e12. | 28.9 | 323 |
| 5 | Structure of the neurotensin receptor 1 in complex with β -arrestin 1. <i>Nature</i> , 2020, 579, 303-308. | 27.8 | 260 |
| 6 | Structural basis for the drug extrusion mechanism by a MATE multidrug transporter. <i>Nature</i> , 2013, 496, 247-251. | 27.8 | 225 |
| 7 | Structural basis for Na ⁺ transport mechanism by a light-driven Na ⁺ pump. <i>Nature</i> , 2015, 521, 48-53. | 27.8 | 224 |
| 8 | Conformational transitions of a neurotensin receptor-Gi1 complex. <i>Nature</i> , 2019, 572, 80-85. | 27.8 | 199 |
| 9 | Structural basis for dynamic mechanism of proton-coupled symport by the peptide transporter POT. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 11343-11348. | 7.1 | 197 |
| 10 | Outward- and inward-facing structures of a putative bacterial transition-metal transporter with homology to ferroportin. <i>Nature Communications</i> , 2015, 6, 8545. | 12.8 | 103 |
| 11 | Structural insights into ligand recognition by the lysophosphatidic acid receptor LPA6. <i>Nature</i> , 2017, 548, 356-360. | 27.8 | 101 |
| 12 | Disruption of ATM in p53-null cells causes multiple functional abnormalities in cellular response to ionizing radiation. <i>Oncogene</i> , 1999, 18, 7002-7009. | 5.9 | 100 |
| 13 | Crystal structure of the natural anion-conducting channelrhodopsin GtACR1. <i>Nature</i> , 2018, 561, 343-348. | 27.8 | 93 |
| 14 | Atomistic design of microbial opsin-based blue-shifted optogenetics tools. <i>Nature Communications</i> , 2015, 6, 7177. | 12.8 | 78 |
| 15 | Structural basis for channel conduction in the pump-like channelrhodopsin ChRmine. <i>Cell</i> , 2022, 185, 672-689.e23. | 28.9 | 72 |
| 16 | Structural mechanisms of selectivity and gating in anion channelrhodopsins. <i>Nature</i> , 2018, 561, 349-354. | 27.8 | 67 |
| 17 | Water-Containing Hydrogen-Bonding Network in the Active Center of Channelrhodopsin. <i>Journal of the American Chemical Society</i> , 2014, 136, 3475-3482. | 13.7 | 59 |
| 18 | Effective Application of Bicelles for Conformational Analysis of G Protein-Coupled Receptors by Hydrogen/Deuterium Exchange Mass Spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2015, 26, 808-817. | 2.8 | 50 |

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 19 | Structural basis for dynamic mechanism of nitrate/nitrite antiport by NarK. <i>Nature Communications</i> , 2015, 6, 7097. | 12.8 | 50 |
| 20 | Mutant of a Light-Driven Sodium Ion Pump Can Transport Cesium Ions. <i>Journal of Physical Chemistry Letters</i> , 2016, 7, 51-55. | 4.6 | 42 |
| 21 | Molecular Dynamics of Channelrhodopsin at the Early Stages of Channel Opening. <i>PLoS ONE</i> , 2015, 10, e0131094. | 2.5 | 33 |
| 22 | Chimeras of Channelrhodopsin-1 and -2 from <i>Chlamydomonas reinhardtii</i> Exhibit Distinctive Light-induced Structural Changes from Channelrhodopsin-2. <i>Journal of Biological Chemistry</i> , 2015, 290, 11623-11634. | 3.4 | 31 |
| 23 | Role of Asn112 in a Light-Driven Sodium Ion-Pumping Rhodopsin. <i>Biochemistry</i> , 2016, 55, 5790-5797. | 2.5 | 27 |
| 24 | The light-driven sodium ion pump: A new player in rhodopsin research. <i>BioEssays</i> , 2016, 38, 1274-1282. | 2.5 | 23 |
| 25 | Crystal structures of the TRIC trimeric intracellular cation channel orthologues. <i>Cell Research</i> , 2016, 26, 1288-1301. | 12.0 | 21 |
| 26 | Structural and spectral characterizations of C1C2 channelrhodopsin and its mutants by molecular simulations. <i>Chemical Physics Letters</i> , 2013, 556, 266-271. | 2.6 | 18 |
| 27 | Structural Properties of the Human Protease-Activated Receptor 1 Changing by a Strong Antagonist. <i>Structure</i> , 2018, 26, 829-838.e4. | 3.3 | 13 |
| 28 | Exciton Circular Dichroism in Channelrhodopsin. <i>Journal of Physical Chemistry B</i> , 2014, 118, 11873-11885. | 2.6 | 12 |
| 29 | Structure-Function Relationship of Channelrhodopsins. <i>Advances in Experimental Medicine and Biology</i> , 2021, 1293, 35-53. | 1.6 | 12 |
| 30 | Crystallization and preliminary X-ray diffraction analysis of YidC, a membrane-protein chaperone and insertase from <i>Bacillus halodurans</i> . <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2014, 70, 1056-1060. | 0.8 | 11 |
| 31 | Atypical structural snapshots of human cytomegalovirus GPCR interactions with host G proteins. <i>Science Advances</i> , 2022, 8, eabl5442. | 10.3 | 11 |
| 32 | Conformational Plasticity of Human Protease-Activated Receptor 1 upon Antagonist- and Agonist-Binding. <i>Structure</i> , 2019, 27, 1517-1526.e3. | 3.3 | 8 |
| 33 | Crystal structure of channelrhodopsin, a light-gated cation channel - all cations lead through the monomer -. <i>Biophysics (Nagoya-shi, Japan)</i> , 2013, 9, 57-61. | 0.4 | 7 |
| 34 | Development of High Frequency Vestibulo-Ocular Responses to Active Head Shaking. <i>Acta Oto-Laryngologica</i> , 1995, 115, 265-267. | 0.9 | 5 |
| 35 | Structural Basis for Dynamic Mechanism of Proton-Coupled Symport by the Peptide Transporter POT. <i>Seibutsu Butsuri</i> , 2014, 54, 085-090. | 0.1 | 1 |
| 36 | A Clinical and Electroencephalographic Study on Anti-Epileptic Activity of Clonazepam.. <i>Psychiatry and Clinical Neurosciences</i> , 1977, 31, 183-194. | 1.8 | 0 |

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 37 | 1PT128 Crystal Structure of a light-gated cation channel, channelrhodopsin(The 50th Annual Meeting) Tj ETQq1 1 0.784314 rgBT /Over | 0.1 | 0 |
| 38 | 2P109 Hydrogen-bonding network in the active center of a light-gated ion channel, channelrhodopsin(03. Membrane proteins,Poster). Seibutsu Butsuri, 2013, 53, S177. | 0.1 | 0 |
| 39 | Molecular Mechanisms of Membrane Channel and Transporter. Acta Crystallographica Section A: Foundations and Advances, 2014, 70, C38-C38. | 0.1 | 0 |
| 40 | 1P109 Structural changes of channelrhodopsin under various cation conditions(03. Membrane) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 62 Butsuri, 2014, 54, S159. | 0.1 | 0 |
| 41 | Structure-Functional Analysis of Channelrhodopsins. , 2015, , 31-45. | | 0 |
| 42 | Structure Based Engineering of Blue-shifted Optogenetics Tools. Seibutsu Butsuri, 2017, 57, 196-199. | 0.1 | 0 |
| 43 | Crystal Structure of Channelrhodopsin, A Light-Gated Cation Channel. Nihon Kessho Gakkaishi, 2012, 54, 220-225. | 0.0 | 0 |
| 44 | Crystal Structure of Channelrhodopsin, a Light-Gated Cation Channel. Seibutsu Butsuri, 2013, 53, 246-249. | 0.1 | 0 |