

Shoji Maeda

List of Publications by Year in descending order

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43
papers

4,614
citations

172457

29
h-index

289244

40
g-index

51
all docs

51
docs citations

51
times ranked

5526
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>Nod2</i> Mutation in Crohn's Disease Potentiates NF- κ B Activity and IL-1 β Processing. <i>Science</i> , 2005, 307, 734-738.	12.6	717
2	Structure of the connexin 26 gap junction channel at 3.5-Å resolution. <i>Nature</i> , 2009, 458, 597-602.	27.8	642
3	Structure of the μ -opioid receptor-Gi protein complex. <i>Nature</i> , 2018, 558, 547-552.	27.8	527
4	Structure of a Signaling Cannabinoid Receptor 1-G Protein Complex. <i>Cell</i> , 2019, 176, 448-458.e12.	28.9	323
5	Structures of the M1 and M2 muscarinic acetylcholine receptor/G-protein complexes. <i>Science</i> , 2019, 364, 552-557.	12.6	244
6	Major virulence factors, VacA and CagA, are commonly positive in <i>Helicobacter pylori</i> isolates in Japan. <i>Gut</i> , 1998, 42, 338-343.	12.1	227
7	Distinct Mechanism of <i>Helicobacter pylori</i> -mediated NF- κ B Activation between Gastric Cancer Cells and Monocytic Cells. <i>Journal of Biological Chemistry</i> , 2001, 276, 44856-44864.	3.4	173
8	Structure of <i>cag</i> pathogenicity island in Japanese <i>Helicobacter pylori</i> isolates. <i>Gut</i> , 1999, 44, 336-341.	12.1	162
9	Development of an antibody fragment that stabilizes GPCR/G-protein complexes. <i>Nature Communications</i> , 2018, 9, 3712.	12.8	157
10	Structure of the gap junction channel and its implications for its biological functions. <i>Cellular and Molecular Life Sciences</i> , 2011, 68, 1115-1129.	5.4	115
11	Roles of Met-34, Cys-64, and Arg-75 in the Assembly of Human Connexin 26. <i>Journal of Biological Chemistry</i> , 2003, 278, 1807-1816.	3.4	96
12	cDNA Microarray Analysis of <i>Helicobacter pylori</i> -Mediated Alteration of Gene Expression in Gastric Cancer Cells. <i>Biochemical and Biophysical Research Communications</i> , 2001, 284, 443-449.	2.1	74
13	Changes with Age in Proteoglycan Synthesis in Cells Cultured In Vitro From the Inner and Outer Rabbit Annulus Fibrosus. <i>Spine</i> , 2000, 25, 166.	2.0	69
14	Functional Impact of Human Collagen $\alpha 2(XI)$ Gene Polymorphism in Pathogenesis of Ossification of the Posterior Longitudinal Ligament of the Spine. <i>Journal of Bone and Mineral Research</i> , 2001, 16, 948-957.	2.8	69
15	High seropositivity of anti-CagA antibody in <i>Helicobacter pylori</i> -infected patients irrelevant to peptic ulcers and normal mucosa in Japan. <i>Digestive Diseases and Sciences</i> , 1997, 42, 1841-1847.	2.3	67
16	Gender-specific haplotype association of collagen $\alpha 2(XI)$ gene in ossification of the posterior longitudinal ligament of the spine. <i>Journal of Human Genetics</i> , 2001, 46, 1-4.	2.3	65
17	<i>Helicobacter pylori</i> specific nested PCR assay for the detection of 23S rRNA mutation associated with clarithromycin resistance. <i>Gut</i> , 1998, 43, 317-321.	12.1	63
18	Structural and functional studies of gap junction channels. <i>Current Opinion in Structural Biology</i> , 2010, 20, 423-430.	5.7	63

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19	Determination of interstitial collagenase (MMP-1) in patients with rheumatoid arthritis.. <i>Annals of the Rheumatic Diseases</i> , 1995, 54, 970-975.	0.9	60
20	Structural insights into the subtype-selective antagonist binding to the M2 muscarinic receptor. <i>Nature Chemical Biology</i> , 2018, 14, 1150-1158.	8.0	59
21	Conformational Complexity and Dynamics in a Muscarinic Receptor Revealed by NMR Spectroscopy. <i>Molecular Cell</i> , 2019, 75, 53-65.e7.	9.7	59
22	Probing G α i1 protein activation at single- α amino acid resolution. <i>Nature Structural and Molecular Biology</i> , 2015, 22, 686-694.	8.2	58
23	Local membrane charge regulates β 2 adrenergic receptor coupling to Gi3. <i>Nature Communications</i> , 2019, 10, 2234.	12.8	57
24	Cryo-EM structure of the rhodopsin-G α i- β 3 complex reveals binding of the rhodopsin C-terminal tail to the β subunit. <i>ELife</i> , 2019, 8, .	6.0	52
25	Activation of the β 2B adrenoceptor by the sedative sympatholytic dexmedetomidine. <i>Nature Chemical Biology</i> , 2020, 16, 507-512.	8.0	51
26	Ligand Binding of the Second PDZ Domain Regulates Clustering of PSD-95 with the Kv1.4 Potassium Channel. <i>Journal of Biological Chemistry</i> , 2002, 277, 3640-3646.	3.4	49
27	Asparagine 175 of Connexin32 Is a Critical Residue for Docking and Forming Functional Heterotypic Gap Junction Channels with Connexin26. <i>Journal of Biological Chemistry</i> , 2011, 286, 19672-19681.	3.4	43
28	Production of GPCR and GPCR complexes for structure determination. <i>Current Opinion in Structural Biology</i> , 2013, 23, 381-392.	5.7	37
29	Structural mechanism underlying primary and secondary coupling between GPCRs and the Gi/o family. <i>Nature Communications</i> , 2020, 11, 3160.	12.8	36
30	Structure and selectivity engineering of the M μ 1 muscarinic receptor toxin complex. <i>Science</i> , 2020, 369, 161-167.	12.6	35
31	Analysis of apoptotic and antiapoptotic signalling pathways induced by <i>Helicobacter pylori</i> . <i>Journal of Clinical Pathology</i> , 2002, 55, 286-293.	1.9	32
32	Crystallization Scale Preparation of a Stable GPCR Signaling Complex between Constitutively Active Rhodopsin and G-Protein. <i>PLoS ONE</i> , 2014, 9, e98714.	2.5	24
33	A description of the structural determination procedures of a gap junction channel at 3.5- \AA resolution. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2009, 65, 758-766.	2.5	23
34	Transport of organic cation in renal brush-border membrane from rats with renal ischemic injury. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1993, 1150, 103-110.	2.6	19
35	Assessment of gastric carcinoma risk associated with <i>Helicobacter pylori</i> may vary depending on the antigen used: CagA specific enzyme-linked immunoadsorbent assay (ELISA) versus commercially available <i>H. pylori</i> ELISAs. <i>Cancer</i> , 2000, 88, 1530-5.	4.1	19
36	Structural basis for the constitutive activity and immunomodulatory properties of the Epstein-Barr virus-encoded G protein-coupled receptor BILF1. <i>Immunity</i> , 2021, 54, 1405-1416.e7.	14.3	18

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37	Atypical structural snapshots of human cytomegalovirus GPCR interactions with host G proteins. Science Advances, 2022, 8, eabl5442.	10.3	11
38	Histochemical Demonstration of Pyrophosphatase. Biotechnic & Histochemistry, 1956, 31, 13-16.	0.4	10
39	Identification of a surface structure in the fourth component of human complement, C4, which becomes hidden upon activation by C1E% _s . Biochemical Journal, 1993, 289, 503-508.	3.7	5
40	A Description of a Structure Determination Procedure of a Gap Junction Channel at 3.5Å Resolution.. Nihon Kessho Gakkaishi, 2009, 51, 327-333.	0.0	0
41	Structure of Human Gap Junction Channel. Seibutsu Butsuri, 2010, 50, 190-191.	0.1	0
42	Structure of the Gap Junction Channel. Nihon Kessho Gakkaishi, 2010, 52, 25-30.	0.0	0
43	Structural insights into the subtype-selective antagonist binding to the M2 muscarinic receptor. Proceedings for Annual Meeting of the Japanese Pharmacological Society, 2020, 93, 3-P-359.	0.0	0