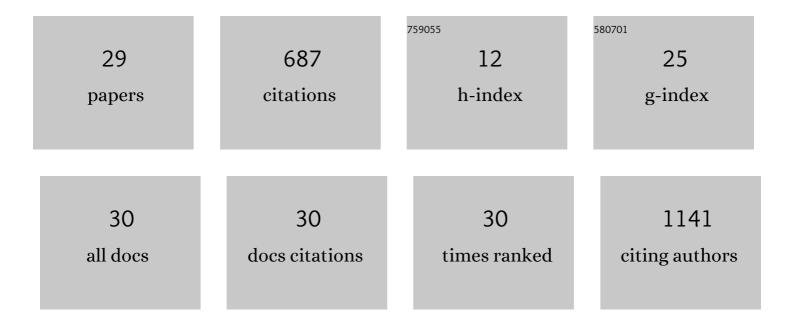
Hideyuki Miyatake

List of Publications by Year in descending order

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Ηισενιικί Μινλτλκέ

#	Article	IF	CITATIONS
1	In Silico and In Cell Hybrid Selection of Nonrapalog Ligands to Allosterically Inhibit the Kinase Activity of mTORC1. Journal of Medicinal Chemistry, 2022, 65, 1329-1341.	2.9	10
2	Mapping of mTOR drug targets: Featured platforms for anti-cancer drug discovery. , 2022, 232, 108012.		12
3	Development of a Non-IgG PD-1/PD-L1 Inhibitor by <i>in Silico</i> Mutagenesis and an In-Cell Protein–Protein Interaction Assay. ACS Chemical Biology, 2021, 16, 316-323.	1.6	7
4	Biomimetic Glucose Triggerâ€Insulin Release System Based on Hydrogel Loading Bidentate β yclodextrin. Advanced Functional Materials, 2021, 31, 2104488.	7.8	15
5	Evaluation of the Binding Kinetics of RHEB with mTORC1 by In-Cell and In Vitro Assays. International Journal of Molecular Sciences, 2021, 22, 8766.	1.8	5
6	Preparation of Biphenyl-Conjugated Bromotyrosine for Inhibition of PD-1/PD-L1 Immune Checkpoint Interactions. International Journal of Molecular Sciences, 2020, 21, 3639.	1.8	8
7	Cell migration and growth induced by photo-immobilised vascular endothelial growth factor (VECF) isoforms. Journal of Materials Chemistry B, 2019, 7, 4272-4279.	2.9	23
8	Crystal structure of phyllogen, a phyllody-inducing effector protein of phytoplasma. Biochemical and Biophysical Research Communications, 2019, 513, 952-957.	1.0	24
9	Enhancement of Binding Affinity of Folate to Its Receptor by Peptide Conjugation. International Journal of Molecular Sciences, 2019, 20, 2152.	1.8	9
10	Escherichia coli expression, purification, and refolding of human folate receptor α (hFRα) and β (hFRβ). Protein Expression and Purification, 2018, 149, 17-22.	0.6	3
11	Bioorthogonal Approaches To Prepare Specifically Modified Functional Proteins. ACS Symposium Series, 2018, , 15-24.	0.5	1
12	Thiophene-Conjugated Ligand Probe for Nonenzymatic Turn-On Electrochemical Protein Detection. Analytical Chemistry, 2018, 90, 11179-11182.	3.2	2
13	IP ₃ -mediated gating mechanism of the IP ₃ receptor revealed by mutagenesis and X-ray crystallography. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 4661-4666.	3.3	70
14	Adhesive Growth Factors Inspired by Underwater Adhesion Proteins. ACS Symposium Series, 2017, , 83-91.	0.5	2
15	A novel sphingomyelin/cholesterol domainâ€specific probe reveals the dynamics of the membrane domains during virus release and in Niemannâ€Pick type C. FASEB Journal, 2017, 31, 1301-1322.	0.2	34
16	Molecular Mechanism of HIV-1 Vpr for Binding to Importin-α. Journal of Molecular Biology, 2016, 428, 2744-2757.	2.0	24
17	A Bioorthogonal Approach for the Preparation of a Titaniumâ€Binding Insulinâ€like Growthâ€Factorâ€1 Derivative by Using Tyrosinase. Angewandte Chemie - International Edition, 2016, 55, 11447-11451.	7.2	26
18	Innenrücktitelbild: A Bioorthogonal Approach for the Preparation of a Titaniumâ€Binding Insulinâ€like Growthâ€Factorâ€1 Derivative by Using Tyrosinase (Angew. Chem. 38/2016). Angewandte Chemie, 2016, 128, 11861-11861.	1.6	0

Ηιδεγυκι Μιγατακέ

#	Article	IF	CITATIONS
19	A Bioorthogonal Approach for the Preparation of a Titaniumâ€Binding Insulinâ€ŀike Growthâ€Factorâ€1 Derivative by Using Tyrosinase. Angewandte Chemie, 2016, 128, 11619-11623.	1.6	2
20	Crystal Structure of Human Importin-α1 (Rch1), Revealing a Potential Autoinhibition Mode Involving Homodimerization. PLoS ONE, 2015, 10, e0115995.	1.1	20
21	Leukocyte cellâ€derived chemotaxin 2 is a zincâ€binding protein. FEBS Letters, 2013, 587, 404-409.	1.3	13
22	A Bilirubin-Inducible Fluorescent Protein from Eel Muscle. Cell, 2013, 153, 1602-1611.	13.5	269
23	Real-Time Control of Nanoscale Protein Assembly for Further Crystallization Using a Solution Circulating Nanoaggregation Control Apparatus. Crystal Growth and Design, 2012, 12, 4466-4472.	1.4	0
24	Structure and characterization of amidase from Rhodococcus sp. N-771: Insight into the molecular mechanism of substrate recognition. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2010, 1804, 184-192.	1.1	50
25	Crystal Structures of the Lumazine Protein from <i>Photobacterium kishitanii</i> in Complexes with the Authentic Chromophore, 6,7-Dimethyl- 8-(1′- <scp>d</scp> -Ribityl) Lumazine, and Its Analogues, Riboflavin and Flavin Mononucleotide, at High Resolution. Journal of Bacteriology, 2010, 192, 127-133.	1.0	24
26	Thermodynamic Characterization of the Interaction between Prefoldin and Group II Chaperonin. Journal of Molecular Biology, 2010, 399, 628-636.	2.0	16
27	Crystallization and preliminary X-ray analysis of the oxygenase component (HpaB) of 4-hydroxyphenylacetate 3-monooxygenase fromThermus thermophilusHB8. Acta Crystallographica Section F: Structural Biology Communications, 2007, 63, 556-559.	0.7	7
28	Development of a fully automated macromolecular crystallization/observation robotic system, HTS-80. Acta Crystallographica Section D: Biological Crystallography, 2005, 61, 658-663.	2.5	10
29	Development of an RHEB-Targeting Peptide To Inhibit mTORC1 Kinase Activity. ACS Omega, 0, , .	1.6	1