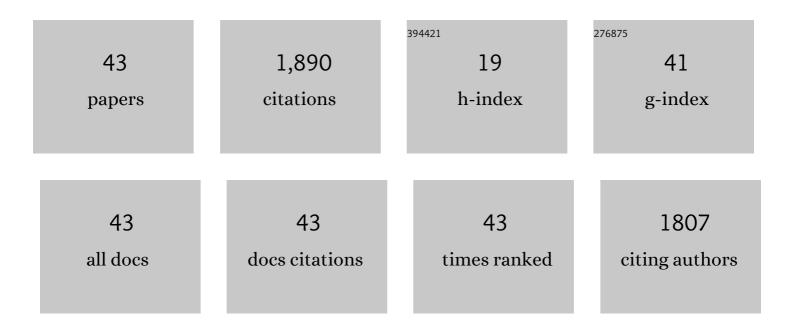
Yoshimitsu Kakuta

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Crystal structure of endocrine-disrupting chemical bisphenol A and estrogen-related receptor γ. Journal of Biochemistry, 2022, 171, 23-25.	1.7	4
2	The crystal structure of mouse SULT2A8 reveals the mechanism of 7α-hydroxyl, bile acid sulfation. Biochemical and Biophysical Research Communications, 2021, 562, 15-20.	2.1	3
3	Minimal protein-only RNase P structure reveals insights into tRNA precursor recognition and catalysis. Journal of Biological Chemistry, 2021, 297, 101028.	3.4	13
4	Non-conventional octameric structure of C-phycocyanin. Communications Biology, 2021, 4, 1238.	4.4	15
5	Pentatricopeptide repeats of protein-only RNase P use a distinct mode to recognize conserved bases and structural elements of pre-tRNA. Nucleic Acids Research, 2020, 48, 11815-11826.	14.5	26
6	Structural insight into the recognition of pathogen-derived phosphoglycolipids by C-type lectin receptor DCAR. Journal of Biological Chemistry, 2020, 295, 5807-5817.	3.4	13
7	Biochemical and structural characterization of a thermostable Dps protein with Hisâ€ŧype ferroxidase centers and outer metalâ€binding sites. FEBS Open Bio, 2020, 10, 1219-1229.	2.3	8
8	Evaluation of the Influence of Halogenation on the Binding of Bisphenol A to the Estrogen-Related Receptor Î ³ . Chemical Research in Toxicology, 2020, 33, 889-902.	3.3	6
9	Structural basis for the broad substrate specificity of the human tyrosylprotein sulfotransferase-1. Scientific Reports, 2017, 7, 8776.	3.3	21
10	A rationally engineered yeast pyruvyltransferase Pvg1p introduces sialylation-like properties in neo-human-type complex oligosaccharide. Scientific Reports, 2016, 6, 26349.	3.3	16
11	Functional implication of archaeal homologues of human RNase P protein pair Pop5 and Rpp30. Journal of Biochemistry, 2016, 159, 31-40.	1.7	7
12	Mutation of the gene encoding the ribonuclease P RNA in the hyperthermophilic archaeon Thermococcus kodakarensis causes decreased growth rate and impaired processing of tRNA precursors. Biochemical and Biophysical Research Communications, 2015, 468, 660-665.	2.1	3
13	Pentatricopeptide repeat motifs in the processing enzyme PRORP1 in Arabidopsis thaliana play a crucial role in recognition of nucleotide bases at TI^C loop in precursor tRNAs. Biochemical and Biophysical Research Communications, 2014, 450, 1541-1546.	2.1	25
14	Tamavidin 2-HOT, a highly thermostable biotin-binding protein. Journal of Biotechnology, 2014, 169, 1-8.	3.8	14
15	The crystal structure of novel chondroitin lyase ODVâ€E66, a baculovirus envelope protein. FEBS Letters, 2013, 587, 3943-3948.	2.8	15
16	Crystal structure of human tyrosylprotein sulfotransferase-2 reveals the mechanism of protein tyrosine sulfation reaction. Nature Communications, 2013, 4, 1572.	12.8	57
17	The Structural Basis for a Coordinated Reaction Catalyzed by a Bifunctional Glycosyltransferase in Chondroitin Biosynthesis. Journal of Biological Chemistry, 2012, 287, 36022-36028.	3.4	14
18	Loss-of-Function Mutation in Bi-Functional Marine Bacterial Sialyltransferase. Bioscience, Biotechnology and Biochemistry, 2012, 76, 1639-1644.	1.3	4

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19	Kinetics of Iterative Carbohydrate Transfer to Polysaccharide Catalyzed by Chondroitin Polymerase on a Highly Sensitive Flowâ€Type 27â€MHz Quartzâ€Crystal Microbalance. Chemistry - A European Journal, 2012, 18, 7388-7393.	3.3	5
20	Crystallographic survey of active sites of an unclassified glutathione transferase from Bombyx mori. Biochimica Et Biophysica Acta - General Subjects, 2011, 1810, 1355-1360.	2.4	9
21	Molecular cloning, expression, and functional analysis of a predicted sulfotransferase STF9 from Mycobacterium avium. Molecular and Cellular Biochemistry, 2011, 350, 155-162.	3.1	7
22	Glucuronyltransferase Activity of KfiC from Escherichia coli Strain K5 Requires Association of KfiA. Journal of Biological Chemistry, 2010, 285, 1597-1606.	3.4	32
23	Crystal structure of <i>α</i> /i>î²â€galactoside <i>î±</i> 2,3â€sialyltransferase from a luminous marine bacterium, <i>Photobacterium phosphoreum</i> . FEBS Letters, 2009, 583, 2083-2087.	2.8	23
24	On the similar spatial arrangement of active site residues in PAPSâ€dependent and phenolic sulfateâ€utilizing sulfotransferases. FEBS Letters, 2009, 583, 3091-3094.	2.8	4
25	Crystal structure of chondroitin polymerase from Escherichia coli K4. Biochemical and Biophysical Research Communications, 2009, 378, 10-14.	2.1	69
26	Structural basis for the broad range substrate specificity of a novel mouse cytosolic sulfotransferase—mSULT1D1. Biochemical and Biophysical Research Communications, 2009, 379, 76-80.	2.1	2
27	Snapshot of a Michaelis complex in a sulfuryl transfer reaction: Crystal structure of a mouse sulfotransferase, mSULT1D1, complexed with donor substrate and accepter substrate. Biochemical and Biophysical Research Communications, 2009, 383, 83-87.	2.1	20
28	Crystal structure of mSULT1D1, a mouse catecholamine sulfotransferase. FEBS Letters, 2008, 582, 3909-3914.	2.8	7
29	ERRγ tethers strongly bisphenol A and 4-α-cumylphenol in an induced-fit manner. Biochemical and Biophysical Research Communications, 2008, 373, 408-413.	2.1	75
30	The Chondroitin Polymerase K4CP and the Molecular Mechanism of Selective Bindings of Donor Substrates to Two Active Sites. Journal of Biological Chemistry, 2008, 283, 32328-32333.	3.4	24
31	2P-019 X-ray crystal structure of mouse sulfotransferase SULT5A1(The 46th Annual Meeting of the) Tj ETQq1 1	0.784314 0.1	rgBT /Overlo
32	Structural Evidence for Endocrine Disruptor Bisphenol A Binding to Human Nuclear Receptor ERRÂ. Journal of Biochemistry, 2007, 142, 517-524.	1.7	206
33	Crystal Structure of Vibrionaceae Photobacterium sp. JT-ISH-224 Â2,6-Sialyltransferase in a Ternary Complex With Donor Product CMP and Acceptor Substrate Lactose: Catalytic Mechanism and Substrate Recognition. Glycobiology, 2007, 18, 66-73.	2.5	58
34	Crystal Structures of Archaeal Ribonuclease P Proteins. Nihon Kessho Gakkaishi, 2007, 49, 255-258.	0.0	0
35	Crystal Structure of a Ribonuclease P Protein Ph1601p from Pyrococcus horikoshii OT3: An Archaeal Homologue of Human Nuclear Ribonuclease P Protein Rpp21,. Biochemistry, 2005, 44, 12086-12093.	2.5	40
36	Crystal structure of the regulatory subunit of archaeal initiation factor 2B (aIF2B) from hyperthermophilic archaeon Pyrococcus horikoshii OT3: a proposed structure of the regulatory subcomplex of eukaryotic IF2B. Biochemical and Biophysical Research Communications, 2004, 319, 725-732.	2.1	24

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#	Article	IF	CITATIONS
37	Structure and Function of Sulfotransferases. Archives of Biochemistry and Biophysics, 2001, 390, 149-157.	3.0	306
38	Substrate Gating Confers Steroid Specificity to Estrogen Sulfotransferase. Journal of Biological Chemistry, 1999, 274, 30019-30022.	3.4	59
39	Conserved structural motifs in the sulfotransferase family. Trends in Biochemical Sciences, 1998, 23, 129-130.	7.5	158
40	A role of Lys614in the sulfotransferase activity of human heparan sulfateN-deacetylase/N-sulfotransferase. FEBS Letters, 1998, 433, 211-214.	2.8	48
41	The Sulfuryl Transfer Mechanism. Journal of Biological Chemistry, 1998, 273, 27325-27330.	3.4	135
42	Crystal structure of estrogen sulphotransferase. Nature Structural and Molecular Biology, 1997, 4, 904-908.	8.2	263
43	Glycine-15 in the Bend between Two α-Helices Can Explain the Thermostability of DNA Binding Protein HU fromBacillus stearothermophilus. Biochemistry, 1996, 35, 1195-1200.	2.5	52