

# Shinsuke Fujiwara

## List of Publications by Year in descending order

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72  
papers

1,614  
citations

361413

20  
h-index

330143

37  
g-index

72  
all docs

72  
docs citations

72  
times ranked

1420  
citing authors

#	ARTICLE	IF	CITATIONS
1	Substrate Specificity of an Aminopropyltransferase and the Biosynthesis Pathway of Polyamines in the Hyperthermophilic Crenarchaeon <i>Pyrobaculum caldifontis</i> . <i>Catalysts</i> , 2022, 12, 567.	3.5	0
2	Optimization of reaction condition of recombinase polymerase amplification to detect SARS-CoV-2 DNA and RNA using a statistical method. <i>Biochemical and Biophysical Research Communications</i> , 2021, 567, 195-200.	2.1	10
3	Leucine-Responsive Regulatory Protein in Acetic Acid Bacteria Is Stable and Functions at a Wide Range of Intracellular pH Levels. <i>Journal of Bacteriology</i> , 2021, 203, e0016221.	2.2	1
4	The therapeutic and nutraceutical potential of agmatine, and its enhanced production using <i>Aspergillus oryzae</i> . <i>Amino Acids</i> , 2020, 52, 181-197.	2.7	34
5	Genes regulated by branched-chain polyamine in the hyperthermophilic archaeon <i>Thermococcus kodakarensis</i> . <i>Amino Acids</i> , 2020, 52, 287-299.	2.7	4
6	Branched-chain polyamine stabilizes RNA polymerase at elevated temperatures in hyperthermophiles. <i>Amino Acids</i> , 2020, 52, 275-285.	2.7	5
7	Alteration of enzymes and their application to nucleic acid amplification (Review). <i>International Journal of Molecular Medicine</i> , 2020, 46, 1633-1643.	4.0	6
8	Alteration of enzymes and their application to nucleic acid amplification (Review). <i>International Journal of Molecular Medicine</i> , 2020, 46, 1633-1643.	4.0	6
9	Pol B, a Family B DNA Polymerase, in <i>Thermococcus kodakarensis</i> ; is Important for DNA Repair, but not DNA Replication. <i>Microbes and Environments</i> , 2019, 34, 316-326.	1.6	15
10	The C-terminal flexible region of branched-chain polyamine synthase facilitates substrate specificity and catalysis. <i>FEBS Journal</i> , 2019, 286, 3926-3940.	4.7	3
11	Accurate fidelity analysis of the reverse transcriptase by a modified next-generation sequencing. <i>Enzyme and Microbial Technology</i> , 2018, 115, 81-85.	3.2	14
12	Leucine responsive regulatory protein is involved in methionine metabolism and polyamine homeostasis in acetic acid bacterium <i>Komagataeibacter europaeus</i> . <i>Journal of Bioscience and Bioengineering</i> , 2018, 125, 67-75.	2.2	7
13	Thermostable DNA helicase improves the sensitivity of digital PCR. <i>Biochemical and Biophysical Research Communications</i> , 2018, 495, 2189-2194.	2.1	4
14	High sensitive RNA detection by one-step RT-PCR using the genetically engineered variant of DNA polymerase with reverse transcriptase activity from hyperthermophilies. <i>Journal of Bioscience and Bioengineering</i> , 2018, 125, 275-281.	2.2	12
15	Branched-Chain Polyamine Found in Hyperthermophiles Induces Unique Temperature-Dependent Structural Changes in Genome-Size DNA. <i>ChemPhysChem</i> , 2018, 19, 2284-2284.	2.1	0
16	Agmatine Production by <i>Aspergillus oryzae</i> Is Elevated by Low pH during Solid-State Cultivation. <i>Applied and Environmental Microbiology</i> , 2018, 84, .	3.1	13
17	Branched-Chain Polyamine Found in Hyperthermophiles Induces Unique Temperature-Dependent Structural Changes in Genome-Size DNA. <i>ChemPhysChem</i> , 2018, 19, 2299-2304.	2.1	22
18	Identification of Branched-Chain Polyamines in Hyperthermophiles. <i>Methods in Molecular Biology</i> , 2018, 1694, 81-94.	0.9	3

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19	Function of a thermophilic archaeal chaperonin is enhanced by electrostatic interactions with its targets. <i>Journal of Bioscience and Bioengineering</i> , 2017, 124, 283-288.	2.2	2
20	High sensitive one-step RT-PCR using MMLV reverse transcriptase, DNA polymerase with reverse transcriptase activity, and DNA/RNA helicase. <i>Biochemical and Biophysical Research Communications</i> , 2017, 487, 128-133.	2.1	5
21	Active site geometry of a novel aminopropyltransferase for biosynthesis of hyperthermophile-specific branched-chain polyamine. <i>FEBS Journal</i> , 2017, 284, 3684-3701.	4.7	10
22	Identification of a novel acetylated form of branched-chain polyamine from a hyperthermophilic archaeon <i>Thermococcus kodakarensis</i> . <i>Bioscience, Biotechnology and Biochemistry</i> , 2017, 81, 1845-1849.	1.3	4
23	Next-generation sequencing-based analysis of reverse transcriptase fidelity. <i>Biochemical and Biophysical Research Communications</i> , 2017, 492, 147-153.	2.1	20
24	Gene regulation of two ferredoxin:NADP+ oxidoreductases by the redox-responsive regulator SurR in <i>Thermococcus kodakarensis</i> . <i>Extremophiles</i> , 2017, 21, 903-917.	2.3	6
25	Enhanced detection of RNA by MMLV reverse transcriptase coupled with thermostable DNA polymerase and DNA/RNA helicase. <i>Enzyme and Microbial Technology</i> , 2017, 96, 111-120.	3.2	13
26	Polyamines in brown rice vinegar function as potent attractants for the spotted wing drosophila. <i>Journal of Bioscience and Bioengineering</i> , 2017, 123, 78-83.	2.2	10
27	Functional Distribution of Archaeal Chaperonins. <i>Heat Shock Proteins</i> , 2017, , 113-128.	0.2	0
28	Naturally occurring branched-chain polyamines induce a crosslinked meshwork structure in a giant DNA. <i>Journal of Chemical Physics</i> , 2016, 145, 235103.	3.0	17
29	Application of a Euryarchaeota-Specific Helicase from <i>Thermococcus kodakarensis</i> for Noise Reduction in PCR. <i>Applied and Environmental Microbiology</i> , 2016, 82, 3022-3031.	3.1	11
30	Effective Trapping of Fruit Flies with Cultures of Metabolically Modified Acetic Acid Bacteria. <i>Applied and Environmental Microbiology</i> , 2015, 81, 2265-2273.	3.1	10
31	Change in the plasmid copy number in acetic acid Bacteria in response to growth phase and acetic acid Concentration. <i>Journal of Bioscience and Bioengineering</i> , 2015, 119, 661-668.	2.2	9
32	A Mutant Chaperonin That Is Functional at Lower Temperatures Enables Hyperthermophilic Archaea To Grow under Cold-Stress Conditions. <i>Journal of Bacteriology</i> , 2015, 197, 2642-2652.	2.2	10
33	Long-Chain and Branched Polyamines in Thermophilic Microbes. , 2015, , 15-25.		6
34	Identification of a Novel Aminopropyltransferase Involved in the Synthesis of Branched-Chain Polyamines in Hyperthermophiles. <i>Journal of Bacteriology</i> , 2014, 196, 1866-1876.	2.2	37
35	Enhanced production of branched-chain amino acids by <i>Gluconacetobacter europaeus</i> with a specific regional deletion in a leucine responsive regulator. <i>Journal of Bioscience and Bioengineering</i> , 2014, 118, 607-615.	2.2	13
36	Cysteine desulphurase plays an important role in environmental adaptation of the hyperthermophilic archaeon <i>Thermococcus kodakarensis</i> . <i>Molecular Microbiology</i> , 2014, 93, 331-345.	2.5	11

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37	Different roles of two transcription factor B proteins in the hyperthermophilic archaeon <i>Thermococcus kodakarensis</i> . <i>Extremophiles</i> , 2014, 18, 573-588.	2.3	9
38	Importance and Determinants of Induction of Cold-Induced DEAD RNA Helicase in the Hyperthermophilic Archaeon <i>Thermococcus kodakarensis</i> . <i>Journal of Bacteriology</i> , 2013, 195, 3442-3450.	2.2	20
39	An Efficient Method Using <i>Gluconacetobacter europaeus</i> To Reduce an Unfavorable Flavor Compound, Acetoin, in Rice Vinegar Production. <i>Applied and Environmental Microbiology</i> , 2013, 79, 7334-7342.	3.1	21
40	Indole-3-Glycerol-Phosphate Synthase Is Recognized by a Cold-Inducible Group II Chaperonin in <i>Thermococcus kodakarensis</i> . <i>Applied and Environmental Microbiology</i> , 2012, 78, 3806-3815.	3.1	11
41	Kinetic analysis of reverse transcriptase activity of bacterial family A DNA polymerases. <i>Biochemical and Biophysical Research Communications</i> , 2012, 427, 654-658.	2.1	6
42	Mutations to create thermostable reverse transcriptase with bacterial family A DNA polymerase from <i>Thermotoga petrophila</i> K4. <i>Journal of Bioscience and Bioengineering</i> , 2012, 113, 315-321.	2.2	20
43	Dual Biosynthesis Pathway for Longer-Chain Polyamines in the Hyperthermophilic Archaeon <i>Thermococcus kodakarensis</i> . <i>Journal of Bacteriology</i> , 2010, 192, 4991-5001.	2.2	49
44	Identification of the Phr-dependent heat shock regulon in the hyperthermophilic archaeon, <i>Thermococcus kodakaraensis</i> . <i>Journal of Biochemistry</i> , 2010, 147, 361-370.	1.7	23
45	Efficient in vitro synthesis of cis-polyisoprenes using a thermostable cis-prenyltransferase from a hyperthermophilic archaeon <i>Thermococcus kodakaraensis</i> . <i>Journal of Biotechnology</i> , 2009, 143, 151-156.	3.8	8
46	Effect of Growth Temperature and Growth Phase on the Lipid Composition of the Archaeal Membrane from <i>Thermococcus kodakaraensis</i> . <i>Bioscience, Biotechnology and Biochemistry</i> , 2009, 73, 104-108.	1.3	50
47	Property of cold inducible DEAD-box RNA helicase in hyperthermophilic archaea. <i>Biochemical and Biophysical Research Communications</i> , 2009, 389, 622-627.	2.1	23
48	Agmatine is essential for the cell growth of <i>Thermococcus kodakaraensis</i> . <i>FEMS Microbiology Letters</i> , 2008, 287, 113-120.	1.8	52
49	Functional Characterization of Recombinant Prefoldin Complexes from a Hyperthermophilic Archaeon, <i>Thermococcus</i> sp. Strain KS-1. <i>Journal of Molecular Biology</i> , 2008, 377, 972-983.	4.2	22
50	Expression Profiles and Physiological Roles of Two Types of Prefoldins from the Hyperthermophilic Archaeon <i>Thermococcus kodakaraensis</i> . <i>Journal of Molecular Biology</i> , 2008, 382, 298-311.	4.2	25
51	Efficient synthesis of trans-polyisoprene compounds using two thermostable enzymes in an organic-aqueous dual-liquid phase system. <i>Biochemical and Biophysical Research Communications</i> , 2008, 365, 118-123.	2.1	9
52	Expression Profiles and Physiological Roles of Two Types of Molecular Chaperonins from the Hyperthermophilic Archaeon <i>Thermococcus kodakarensis</i> . <i>Applied and Environmental Microbiology</i> , 2008, 74, 7306-7312.	3.1	28
53	Characterization of MobR, the 3-Hydroxybenzoate-responsive Transcriptional Regulator for the 3-Hydroxybenzoate Hydroxylase Gene of <i>Comamonas testosteroni</i> KH122-3s. <i>Journal of Molecular Biology</i> , 2006, 364, 863-877.	4.2	19
54	Complete genome sequence of the hyperthermophilic archaeon <i>Thermococcus kodakaraensis</i> KOD1 and comparison with <i>Pyrococcus</i> genomes. <i>Genome Research</i> , 2005, 15, 352-363.	5.5	376

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55	Enzymatic and structural characterization of type II isopentenyl diphosphate isomerase from hyperthermophilic archaeon <i>Thermococcus kodakaraensis</i> . <i>Biochemical and Biophysical Research Communications</i> , 2005, 331, 1127-1136.	2.1	17
56	Biophysical analysis of heat-induced structural maturation of glutamate dehydrogenase from a hyperthermophilic archaeon. <i>Journal of Bioscience and Bioengineering</i> , 2004, 97, 305-309.	2.2	7
57	Temperature-dependent modulation of farnesyl diphosphate/geranylgeranyl diphosphate synthase from hyperthermophilic archaea. <i>Biochemical and Biophysical Research Communications</i> , 2004, 325, 1066-1074.	2.1	19
58	Extremophiles: Developments of their special functions and potential resources. <i>Journal of Bioscience and Bioengineering</i> , 2002, 94, 518-525.	2.2	56
59	Two Kinds of Archaeal Chaperonin with Different Temperature Dependency from a Hyperthermophile. <i>Biochemical and Biophysical Research Communications</i> , 2001, 280, 581-587.	2.1	25
60	Comparative analyses of the conformational stability of a hyperthermophilic protein and its mesophilic counterpart. <i>FEBS Journal</i> , 2001, 268, 4144-4150.	0.2	37
61	Conformational Stability of a Hyperthermophilic Protein in Various Conditions for Denaturation. <i>Electrochemistry</i> , 2001, 69, 949-952.	1.4	8
62	Effect of polyamines on histone-induced DNA compaction of hyperthermophilic archaea. <i>Journal of Bioscience and Bioengineering</i> , 2000, 89, 103-106.	2.2	21
63	Sequence and transcriptional studies of five clustered flagellin genes from hyperthermophilic archaeon <i>Pyrococcus kodakaraensis</i> KOD1. <i>FEMS Microbiology Letters</i> , 1999, 178, 183-190.	1.8	15
64	Hyperthermostable Protein Structure Maintained by Intra and Inter-helix Ion-pairs in Archaeal O6-Methylguanine-DNA Methyltransferase. <i>Journal of Molecular Biology</i> , 1999, 292, 707-716.	4.2	97
65	Sequence and transcriptional studies of five clustered flagellin genes from hyperthermophilic archaeon <i>Pyrococcus kodakaraensis</i> KOD1. <i>FEMS Microbiology Letters</i> , 1999, 178, 183-190.	1.8	2
66	Isolation and Characterization of a Second Subunit of Molecular Chaperonin from <i>Pyrococcus kodakaraensis</i> KOD1: Analysis of an ATPase-Deficient Mutant Enzyme. <i>Applied and Environmental Microbiology</i> , 1999, 65, 1801-1805.	3.1	33
67	Comparison of two glutamate producing enzymes from the hyperthermophilic archaeon <i>Pyrococcus</i> sp. KOD1. <i>FEMS Microbiology Letters</i> , 1998, 158, 243-248.	1.8	2
68	In vitro heat effect on heterooligomeric subunit assembly of thermostable indolepyruvate ferredoxin oxidoreductase. <i>FEBS Letters</i> , 1998, 434, 372-376.	2.8	34
69	Ion Pairs Involved in Maintaining a Thermostable Structure of Glutamate Dehydrogenase from a Hyperthermophilic Archaeon. <i>Biochemical and Biophysical Research Communications</i> , 1998, 248, 920-926.	2.1	50
70	Effect of Heat Treatment on Proper Oligomeric Structure Formation of Thermostable Glutamate Dehydrogenase from a Hyperthermophilic Archaeon. <i>Biochemical and Biophysical Research Communications</i> , 1997, 241, 646-652.	2.1	38
71	Thermostable Enzymes of Hyperthermophilic Archaea. <i>Journal of Japan Oil Chemists' Society</i> , 1997, 46, 525-533, 597.	0.3	0
72	Unusual enzyme characteristics of aspartyl-tRNA synthetase from hyperthermophilic archaeon <i>Pyrococcus</i> sp. KOD1. <i>FEBS Letters</i> , 1996, 394, 66-70.	2.8	29