

Maia Kivisaar

List of Publications by Year in descending order

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Version: 2024-02-01

60
papers

1,712
citations

279798

23
h-index

315739

38
g-index

62
all docs

62
docs citations

62
times ranked

1439
citing authors

#	ARTICLE	IF	CITATIONS
1	Tryptone in Growth Media Enhances <i>Pseudomonas putida</i> Biofilm. <i>Microorganisms</i> , 2022, 10, 618.	3.6	2
2	<i>Pseudomonas putida</i> Biofilm Depends on the vWFA-Domain of LapA in Peptides-Containing Growth Medium. <i>International Journal of Molecular Sciences</i> , 2022, 23, 5898.	4.1	3
3	Monitoring the growth, survival and phenol utilization of the fluorescent-tagged <i>Pseudomonas oleovorans</i> immobilized and free cells. <i>Bioresource Technology</i> , 2021, 338, 125568.	9.6	7
4	Pseudouridines of tRNA Anticodon Stem-Loop Have Unexpected Role in Mutagenesis in <i>Pseudomonas</i> sp.. <i>Microorganisms</i> , 2021, 9, 25.	3.6	8
5	Integration Host Factor IHF facilitates homologous recombination and mutagenic processes in <i>Pseudomonas putida</i> . <i>DNA Repair</i> , 2020, 85, 102745.	2.8	7
6	Mutation and Recombination Rates Vary Across Bacterial Chromosome. <i>Microorganisms</i> , 2020, 8, 25.	3.6	17
7	Microbial Metabolic Potential of Phenol Degradation in Wastewater Treatment Plant of Crude Oil Refinery: Analysis of Metagenomes and Characterization of Isolates. <i>Microorganisms</i> , 2020, 8, 652.	3.6	16
8	Narrative of a versatile and adept species <i>Pseudomonas putida</i> . <i>Journal of Medical Microbiology</i> , 2020, 69, 324-338.	1.8	27
9	Seasonal bacterial community dynamics in a crude oil refinery wastewater treatment plant. <i>Applied Microbiology and Biotechnology</i> , 2019, 103, 9131-9141.	3.6	11
10	Involvement of transcription-coupled repair factor Mfd and DNA helicase UvrD in mutational processes in <i>Pseudomonas putida</i> . <i>DNA Repair</i> , 2018, 72, 18-27.	2.8	11
11	The Effect of Cellular Redox Status on the Evolvability of New Catabolic Pathways. <i>MBio</i> , 2018, 9, .	4.1	3
12	Colonization efficiency of <i>Pseudomonas putida</i> is influenced by Fis-controlled transcription of <i>nuoA-N</i> operon. <i>PLoS ONE</i> , 2018, 13, e0201841.	2.5	4
13	Ongoing evolution of <i>Pseudomonas aeruginosa</i> PAO1 sublines complicates studies of DNA damage repair and tolerance. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2017, 797-799, 26-37.	1.0	15
14	DNA Polymerases ImuC and DinB Are Involved in DNA Alkylation Damage Tolerance in <i>Pseudomonas aeruginosa</i> and <i>Pseudomonas putida</i> . <i>PLoS ONE</i> , 2017, 12, e0170719.	2.5	20
15	Contribution of increased mutagenesis to the evolution of pollutants-degrading indigenous bacteria. <i>PLoS ONE</i> , 2017, 12, e0182484.	2.5	9
16	The promoter region of <i>lapA</i> and its transcriptional regulation by Fis in <i>Pseudomonas putida</i> . <i>PLoS ONE</i> , 2017, 12, e0185482.	2.5	13
17	A novel papillation assay for the identification of genes affecting mutation rate in <i>Pseudomonas putida</i> and other pseudomonads. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2016, 790, 41-55.	1.0	3
18	LapF and Its Regulation by Fis Affect the Cell Surface Hydrophobicity of <i>Pseudomonas putida</i> . <i>PLoS ONE</i> , 2016, 11, e0166078.	2.5	20

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19	Freeing <i>Pseudomonas putida</i> of its proviral load strengthens endurance to environmental stresses. <i>Environmental Microbiology</i> , 2015, 17, 76-90.	3.8	62
20	NHEJ enzymes LigD and Ku participate in stationary-phase mutagenesis in <i>Pseudomonas putida</i> . <i>DNA Repair</i> , 2015, 31, 11-18.	2.8	17
21	NER enzymes maintain genome integrity and suppress homologous recombination in the absence of exogenously induced DNA damage in <i>Pseudomonas putida</i> . <i>DNA Repair</i> , 2015, 25, 15-26.	2.8	6
22	<i>Pseudomonas putida</i> Fis Binds to the <i>lapF</i> Promoter In Vitro and Represses the Expression of <i>LapF</i> . <i>PLoS ONE</i> , 2014, 9, e115901.	2.5	12
23	Fis overexpression enhances <i>Pseudomonas putida</i> biofilm formation by regulating the ratio of <i>LapA</i> and <i>LapF</i> . <i>Microbiology (United Kingdom)</i> , 2014, 160, 2681-2693.	1.8	27
24	<i>Pseudomonas putida</i> AlkA and AlkB Proteins Comprise Different Defense Systems for the Repair of Alkylation Damage to DNA – In Vivo, In Vitro, and In Silico Studies. <i>PLoS ONE</i> , 2013, 8, e76198.	2.5	12
25	Homologous recombination is facilitated in starving populations of <i>Pseudomonas putida</i> by phenol stress and affected by chromosomal location of the recombination target. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2012, 737, 12-24.	1.0	19
26	Fis regulates the competitiveness of <i>Pseudomonas putida</i> on barley roots by inducing biofilm formation. <i>Microbiology (United Kingdom)</i> , 2012, 158, 708-720.	1.8	38
27	Mutation Frequency and Spectrum of Mutations Vary at Different Chromosomal Positions of <i>Pseudomonas putida</i> . <i>PLoS ONE</i> , 2012, 7, e48511.	2.5	23
28	Evolution of catabolic pathways and their regulatory systems in synthetic nitroaromatic compounds degrading bacteria. <i>Molecular Microbiology</i> , 2011, 82, 265-268.	2.5	16
29	The impact of ColRS two-component system and TtgABC efflux pump on phenol tolerance of <i>Pseudomonas putida</i> becomes evident only in growing bacteria. <i>BMC Microbiology</i> , 2010, 10, 110.	3.3	26
30	Mechanisms of stationary-phase mutagenesis in bacteria: mutational processes in pseudomonads. <i>FEMS Microbiology Letters</i> , 2010, 312, 1-14.	1.8	42
31	Elevated Mutation Frequency in Surviving Populations of Carbon-Starved <i>rpoS</i> -Deficient <i>Pseudomonas putida</i> Is Caused by Reduced Expression of Superoxide Dismutase and Catalase. <i>Journal of Bacteriology</i> , 2009, 191, 3604-3614.	2.2	14
32	Degradation of nitroaromatic compounds: a model to study evolution of metabolic pathways. <i>Molecular Microbiology</i> , 2009, 74, 777-781.	2.5	26
33	Fis negatively affects binding of Tn4652 transposase by out-competing IHF from the left end of Tn4652. <i>Microbiology (United Kingdom)</i> , 2009, 155, 1203-1214.	1.8	14
34	ColRS two-component system prevents lysis of subpopulation of glucose-grown <i>Pseudomonas putida</i> . <i>Environmental Microbiology</i> , 2008, 10, 2886-2893.	3.8	11
35	Dual role of NER in mutagenesis in <i>Pseudomonas putida</i> . <i>DNA Repair</i> , 2008, 7, 20-30.	2.8	24
36	Target Site Selection of <i>Pseudomonas putida</i> Transposon Tn 4652. <i>Journal of Bacteriology</i> , 2007, 189, 3918-3921.	2.2	11

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37	Oxidative DNA Damage Defense Systems in Avoidance of Stationary-Phase Mutagenesis in <i>Pseudomonas putida</i> . <i>Journal of Bacteriology</i> , 2007, 189, 5504-5514.	2.2	27
38	Study of factors which negatively affect expression of the phenol degradation operon <i>pheBA</i> in <i>Pseudomonas putida</i> . <i>Microbiology (United Kingdom)</i> , 2007, 153, 1860-1871.	1.8	13
39	Study of involvement of <i>ImuB</i> and <i>DnaE2</i> in stationary-phase mutagenesis in <i>Pseudomonas putida</i> . <i>DNA Repair</i> , 2007, 6, 863-868.	2.8	31
40	Involvement of DNA mismatch repair in stationary-phase mutagenesis during prolonged starvation of <i>Pseudomonas putida</i> . <i>DNA Repair</i> , 2006, 5, 505-514.	2.8	18
41	The ColRS Two-Component System Regulates Membrane Functions and Protects <i>Pseudomonas putida</i> against Phenol. <i>Journal of Bacteriology</i> , 2006, 188, 8109-8117.	2.2	53
42	A DNA Polymerase V Homologue Encoded by TOL Plasmid pWW0 Confers Evolutionary Fitness on <i>Pseudomonas putida</i> under Conditions of Environmental Stress. <i>Journal of Bacteriology</i> , 2005, 187, 5203-5213.	2.2	41
43	Simultaneous Degradation of Atrazine and Phenol by <i>Pseudomonas</i> sp. Strain ADP: Effects of Toxicity and Adaptation. <i>Applied and Environmental Microbiology</i> , 2004, 70, 1907-1912.	3.1	104
44	Involvement of Error-Prone DNA Polymerase IV in Stationary-Phase Mutagenesis in <i>Pseudomonas putida</i> . <i>Journal of Bacteriology</i> , 2004, 186, 2735-2744.	2.2	62
45	IHF is the limiting host factor in transposition of <i>Pseudomonas putida</i> transposon Tn4652 in stationary phase. <i>Molecular Microbiology</i> , 2004, 51, 1773-1785.	2.5	21
46	The ColR-ColS two-component signal transduction system is involved in regulation of Tn4652 transposition in <i>Pseudomonas putida</i> under starvation conditions. <i>Molecular Microbiology</i> , 2004, 54, 795-807.	2.5	50
47	Stationary phase mutagenesis: mechanisms that accelerate adaptation of microbial populations under environmental stress. <i>Environmental Microbiology</i> , 2003, 5, 814-827.	3.8	72
48	Different Spectra of Stationary-Phase Mutations in Early-Arising versus Late-Arising Mutants of <i>Pseudomonas putida</i> : Involvement of the DNA Repair Enzyme MutY and the Stationary-Phase Sigma Factor RpoS. <i>Journal of Bacteriology</i> , 2002, 184, 6957-6965.	2.2	47
49	Involvement of σ^S in Starvation-Induced Transposition of <i>Pseudomonas putida</i> Transposon Tn 4652. <i>Journal of Bacteriology</i> , 2001, 183, 5445-5448.	2.2	97
50	Growth medium composition-determined regulatory mechanisms are superimposed on CatR-mediated transcription from the <i>pheBA</i> and <i>catBCA</i> promoters in <i>Pseudomonas putida</i> . <i>Microbiology (United Kingdom)</i> , 2000, 144, 1071-1076.	2.2	10
51	Effects of Combination of Different σ^{70} Hexamers and Downstream Sequences on Stationary-Phase-Specific Sigma Factor σ^S -Dependent Transcription in <i>Pseudomonas putida</i> . <i>Journal of Bacteriology</i> , 2000, 182, 6707-6713.	2.2	35
52	Transcription from Fusion Promoters Generated during Transposition of Transposon Tn 4652 Is Positively Affected by Integration Host Factor in <i>Pseudomonas putida</i> . <i>Journal of Bacteriology</i> , 2000, 182, 589-598.	2.2	21
53	Critical nucleotides in the interaction of CatR with the <i>pheBA</i> promoter: conservation of the CatR-mediated regulation mechanisms between the <i>pheBA</i> and <i>catBCA</i> operons. <i>Microbiology (United Kingdom)</i> , 1999, 143, 1071-1076.	2.2	17
54	Regulation of the Transposase of Tn 4652 by the Transposon-Encoded Protein TnpC. <i>Journal of Bacteriology</i> , 1999, 181, 6312-6318.	2.2	17

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55	Expression of the Transposase Gene <i>tnpA</i> of Tn 4652 Is Positively Affected by Integration Host Factor. <i>Journal of Bacteriology</i> , 1998, 180, 2822-2829.	2.2	49
56	Identification and Characterization of IS 1411 , a New Insertion Sequence Which Causes Transcriptional Activation of the Phenol Degradation Genes in <i>Pseudomonas putida</i> . <i>Journal of Bacteriology</i> , 1998, 180, 5306-5312.	2.2	59
57	In-vivo-generated fusion promoters in <i>Pseudomonas putida</i> . <i>Gene</i> , 1993, 127, 23-29.	2.2	27
58	Sequence of the gene (<i>pheA</i>) encoding phenol monooxygenase from <i>Pseudomonas</i> sp. EST1001: expression in <i>Escherichia coli</i> and <i>Pseudomonas putida</i> . <i>Gene</i> , 1991, 102, 13-18.	2.2	105
59	Sequence of the plasmid-encoded catechol 1,2-dioxygenase-expressing gene, <i>pheB</i> , of phenol-degrading <i>Pseudomonas</i> sp. strain EST1001. <i>Gene</i> , 1991, 98, 15-20.	2.2	60
60	Selection of independent plasmids determining phenol degradation in <i>Pseudomonas putida</i> and the cloning and expression of genes encoding phenol monooxygenase and catechol 1,2-dioxygenase. <i>Plasmid</i> , 1990, 24, 25-36.	1.4	67