

Nathalie Juge

List of Publications by Year in descending order

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117
papers

10,954
citations

34105

52
h-index

31849

101
g-index

119
all docs

119
docs citations

119
times ranked

13644
citing authors

#	ARTICLE	IF	CITATIONS
1	Introduction to the human gut microbiota. <i>Biochemical Journal</i> , 2017, 474, 1823-1836.	3.7	1,988
2	The composition of the gut microbiota throughout life, with an emphasis on early life. <i>Microbial Ecology in Health and Disease</i> , 2015, 26, 26050.	3.5	766
3	Molecular basis for chemoprevention by sulforaphane: a comprehensive review. <i>Cellular and Molecular Life Sciences</i> , 2007, 64, 1105-1127.	5.4	619
4	Mucin glycan foraging in the human gut microbiome. <i>Frontiers in Genetics</i> , 2015, 6, 81.	2.3	612
5	Deglycosylation by small intestinal epithelial cell β -glucosidases is a critical step in the absorption and metabolism of dietary flavonoid glycosides in humans. <i>European Journal of Nutrition</i> , 2003, 42, 29-42.	3.9	579
6	The Evolution of Host Specialization in the Vertebrate Gut Symbiont <i>Lactobacillus reuteri</i> . <i>PLoS Genetics</i> , 2011, 7, e1001314.	3.5	270
7	Utilisation of Mucin Glycans by the Human Gut Symbiont <i>Ruminococcus gnavus</i> Is Strain-Dependent. <i>PLoS ONE</i> , 2013, 8, e76341.	2.5	250
8	Plant protein inhibitors of cell wall degrading enzymes. <i>Trends in Plant Science</i> , 2006, 11, 359-367.	8.8	229
9	Microbial adhesins to gastrointestinal mucus. <i>Trends in Microbiology</i> , 2012, 20, 30-39.	7.7	219
10	Hydrolytic fate of deoxynivalenol-3-glucoside during digestion. <i>Toxicology Letters</i> , 2011, 206, 264-267.	0.8	216
11	Molecular Characterization of Host-Specific Biofilm Formation in a Vertebrate Gut Symbiont. <i>PLoS Genetics</i> , 2013, 9, e1004057.	3.5	162
12	Strain-specific diversity of mucus-binding proteins in the adhesion and aggregation properties of <i>Lactobacillus reuteri</i> . <i>Microbiology (United Kingdom)</i> , 2010, 156, 3368-3378.	1.8	157
13	Unique Organization of Extracellular Amylases into Amylosomes in the Resistant Starch-Utilizing Human Colonic <i>Firmicutes</i> Bacterium <i>Ruminococcus bromii</i> . <i>MBio</i> , 2015, 6, e01058-15.	4.1	145
14	Discovery of intramolecular trans-sialidases in human gut microbiota suggests novel mechanisms of mucosal adaptation. <i>Nature Communications</i> , 2015, 6, 7624.	12.8	143
15	The mucin-degradation strategy of <i>Ruminococcus gnavus</i> : The importance of intramolecular trans-sialidases. <i>Gut Microbes</i> , 2016, 7, 302-312.	9.8	127
16	Mechanistic Insights Into the Cross-Feeding of <i>Ruminococcus gnavus</i> and <i>Ruminococcus bromii</i> on Host and Dietary Carbohydrates. <i>Frontiers in Microbiology</i> , 2018, 9, 2558.	3.5	125
17	Sialidases from gut bacteria: a mini-review. <i>Biochemical Society Transactions</i> , 2016, 44, 166-175.	3.4	121
18	Experimental models to study intestinal microbesâ€™ mucus interactions in health and disease. <i>FEMS Microbiology Reviews</i> , 2019, 43, 457-489.	8.6	114

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19	The Dual Nature of the Wheat Xylanase Protein Inhibitor XIP-I. <i>Journal of Biological Chemistry</i> , 2004, 279, 36029-36037.	3.4	111
20	The Crystal Structure of Feruloyl Esterase A from <i>Aspergillus niger</i> Suggests Evolutive Functional Convergence in Feruloyl Esterase Family. <i>Journal of Molecular Biology</i> , 2004, 338, 495-506.	4.2	110
21	Understanding the Structural Basis for Substrate and Inhibitor Recognition in Eukaryotic GH11 Xylanases. <i>Journal of Molecular Biology</i> , 2008, 375, 1293-1305.	4.2	108
22	Purification and biochemical characterization of a novel α -amylase from <i>Bacillus licheniformis</i> NH1. <i>Process Biochemistry</i> , 2008, 43, 499-510.	3.7	107
23	Interactions defining the specificity between fungal xylanases and the xylanase-inhibiting protein XIP-I from wheat. <i>Biochemical Journal</i> , 2002, 365, 773-781.	3.7	105
24	Structural basis for adaptation of lactobacilli to gastrointestinal mucus. <i>Environmental Microbiology</i> , 2014, 16, 888-903.	3.8	102
25	XIP-I, a xylanase inhibitor protein from wheat: a novel protein function. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2004, 1696, 203-211.	2.3	91
26	Effects of Human Milk Oligosaccharides on the Adult Gut Microbiota and Barrier Function. <i>Nutrients</i> , 2020, 12, 2808.	4.1	86
27	Structural insights into bacterial recognition of intestinal mucins. <i>Current Opinion in Structural Biology</i> , 2014, 28, 23-31.	5.7	83
28	Elucidation of a sialic acid metabolism pathway in mucus-foraging <i>Ruminococcus gnavus</i> unravels mechanisms of bacterial adaptation to the gut. <i>Nature Microbiology</i> , 2019, 4, 2393-2404.	13.3	83
29	Overexpression, Purification, and Characterization of Recombinant Barley α -Amylases 1 and 2 Secreted by the Methylophilic Yeast <i>Pichia pastoris</i> . <i>Protein Expression and Purification</i> , 1996, 8, 204-214.	1.3	79
30	Both binding sites of the starch-binding domain of <i>Aspergillus niger</i> glucoamylase are essential for inducing a conformational change in amylose 1. Edited by R. Huber. <i>Journal of Molecular Biology</i> , 2001, 313, 1149-1159.	4.2	79
31	Sporulation capability and amylosome conservation among diverse human colonic and rumen isolates of the keystone starch-degrader <i>Ruminococcus bromii</i> . <i>Environmental Microbiology</i> , 2018, 20, 324-336.	3.8	79
32	Comparative Characterization of Complete and Truncated Forms of <i>Lactobacillus amylovorus</i> α -Amylase and Role of the C-Terminal Direct Repeats in Raw-Starch Binding. <i>Applied and Environmental Microbiology</i> , 2000, 66, 3350-3356.	3.1	78
33	High-Level Production of Recombinant Fungal Endo- β -1,4-xylanase in the Methylophilic Yeast <i>Pichia pastoris</i> . <i>Protein Expression and Purification</i> , 2000, 19, 179-187.	1.3	75
34	Potential role of glycosidase inhibitors in industrial biotechnological applications. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2004, 1696, 275-287.	2.3	74
35	Emergence of a subfamily of xylanase inhibitors within glycoside hydrolase family 18. <i>FEBS Journal</i> , 2005, 272, 1745-1755.	4.7	74
36	Unravelling the specificity and mechanism of sialic acid recognition by the gut symbiont <i>Ruminococcus gnavus</i> . <i>Nature Communications</i> , 2017, 8, 2196.	12.8	74

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37	Occurrence of proteinaceous endoxylanase inhibitors in cereals. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2004, 1696, 193-202.	2.3	73
38	Factors affecting xylanase functionality in the degradation of arabinoxylans. <i>Biotechnology Letters</i> , 2008, 30, 1139-1150.	2.2	72
39	High-level production of recombinant <i>Aspergillus niger</i> cinnamoyl esterase (FAEA) in the methylotrophic yeast <i>Pichia pastoris</i> . <i>FEMS Yeast Research</i> , 2001, 1, 127-132.	2.3	71
40	Differential Epitope Mapping by STD NMR Spectroscopy To Reveal the Nature of Protein-Ligand Contacts. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 15289-15293.	13.8	71
41	Functional expression of human liver cytosolic β -glucosidase in <i>Pichia pastoris</i> . <i>FEBS Journal</i> , 2002, 269, 249-258.	0.2	70
42	Substrate (aglycone) specificity of human cytosolic beta-glucosidase. <i>Biochemical Journal</i> , 2003, 373, 41-48.	3.7	70
43	Crystal Structure of a Mucus-binding Protein Repeat Reveals an Unexpected Functional Immunoglobulin Binding Activity. <i>Journal of Biological Chemistry</i> , 2009, 284, 32444-32453.	3.4	70
44	<i>Lactobacillus reuteri</i> Inhibition of Enteropathogenic <i>Escherichia coli</i> Adherence to Human Intestinal Epithelium. <i>Frontiers in Microbiology</i> , 2016, 7, 244.	3.5	69
45	Specific Characterization of Substrate and Inhibitor Binding Sites of a Glycosyl Hydrolase Family 11 Xylanase from <i>Aspergillus niger</i> . <i>Journal of Biological Chemistry</i> , 2002, 277, 44035-44043.	3.4	67
46	Structural analysis of xylanase inhibitor protein I (XIP-I), a proteinaceous xylanase inhibitor from wheat (<i>Triticum aestivum</i> , var. Soisson). <i>Biochemical Journal</i> , 2003, 372, 399-405.	3.7	65
47	Potential physiological role of plant glycosidase inhibitors. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2004, 1696, 265-274.	2.3	64
48	The pan-genome of <i>Lactobacillus reuteri</i> strains originating from the pig gastrointestinal tract. <i>BMC Genomics</i> , 2015, 16, 1023.	2.8	64
49	Specificity of feruloyl esterases for water-extractable and water-unextractable feruloylated polysaccharides: influence of xylanase. <i>Journal of Cereal Science</i> , 2003, 38, 281-288.	3.7	63
50	Surfactant-mediated solubilisation of amylose and visualisation by atomic force microscopy. <i>Carbohydrate Polymers</i> , 2003, 51, 177-182.	10.2	61
51	Probing the determinants of substrate specificity of a feruloyl esterase, AnFaeA, from <i>Aspergillus niger</i> . <i>FEBS Journal</i> , 2005, 272, 4362-4371.	4.7	59
52	The StcE metalloprotease of enterohaemorrhagic <i>Escherichia coli</i> reduces the inner mucus layer and promotes adherence to human colonic epithelium <i>ex vivo</i> . <i>Cellular Microbiology</i> , 2017, 19, e12717.	2.1	58
53	Formate cross-feeding and cooperative metabolic interactions revealed by transcriptomics in co-cultures of acetogenic and amyolytic human colonic bacteria. <i>Environmental Microbiology</i> , 2019, 21, 259-271.	3.8	58
54	Mucosal glycan degradation of the host by the gut microbiota. <i>Glycobiology</i> , 2021, 31, 691-696.	2.5	53

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55	Fucosidases from the human gut symbiont <i>Ruminococcus gnavus</i> . <i>Cellular and Molecular Life Sciences</i> , 2021, 78, 675-693.	5.4	52
56	The Crystal Structure of Human Cytosolic β -Glucosidase Unravels the Substrate Aglycone Specificity of a Family 1 Glycoside Hydrolase. <i>Journal of Molecular Biology</i> , 2007, 370, 964-975.	4.2	51
57	Substrate and product hydrolysis specificity in family 11 glycoside hydrolases: an analysis of <i>Penicillium funiculosum</i> and <i>Penicillium griseofulvum</i> xylanases. <i>Applied Microbiology and Biotechnology</i> , 2007, 74, 1001-1010.	3.6	47
58	Domain B protruding at the third beta strand of the alpha/beta barrel in barley alpha-amylase confers distinct isozyme-specific properties. <i>FEBS Journal</i> , 1994, 221, 277-284.	0.2	43
59	A family 11 xylanase from the pathogen <i>Botrytis cinerea</i> is inhibited by plant endoxylanase inhibitors XIP-I and TAXI-I. <i>Biochemical and Biophysical Research Communications</i> , 2005, 337, 160-166.	2.1	43
60	Functional Analysis of Family GH36 β -Galactosidases from <i>Ruminococcus gnavus</i> E1: Insights into the Metabolism of a Plant Oligosaccharide by a Human Gut Symbiont. <i>Applied and Environmental Microbiology</i> , 2012, 78, 7720-7732.	3.1	43
61	<i>Lactobacillus reuteri</i> Surface Mucus Adhesins Upregulate Inflammatory Responses Through Interactions With Innate C-Type Lectin Receptors. <i>Frontiers in Microbiology</i> , 2017, 8, 321.	3.5	43
62	Functional identification of the cDNA coding for a wheat endo-1,4- β -D-xylanase inhibitor1. <i>FEBS Letters</i> , 2002, 519, 66-70.	2.8	42
63	Involvement of Individual Subsites and Secondary Substrate Binding Sites in Multiple Attack on Amylose by Barley β -Amylase. <i>Biochemistry</i> , 2005, 44, 1824-1832.	2.5	42
64	The activity of barley β -amylase on starch granules is enhanced by fusion of a starch binding domain from <i>Aspergillus niger</i> glucoamylase. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2006, 1764, 275-284.	2.3	42
65	Proteinaceous inhibitors of carbohydrate-active enzymes in cereals: implication in agriculture, cereal processing and nutrition. <i>Journal of the Science of Food and Agriculture</i> , 2006, 86, 1573-1586.	3.5	41
66	Use of Atomic Force Microscopy to Study the Multi-Modular Interaction of Bacterial Adhesins to Mucins. <i>International Journal of Molecular Sciences</i> , 2016, 17, 1854.	4.1	39
67	Isozyme hybrids within the protruding third loop domain of the barley β -amylase (β / β) ₈ -barrel implication for BASI sensitivity and substrate affinity. <i>FEBS Letters</i> , 1995, 363, 299-303.	2.8	37
68	Specific inhibition of barley β -amylase β 2 by barley β -amylase/subtilisin inhibitor depends on charge interactions and can be conferred to isozyme β 1 by mutation. <i>FEBS Journal</i> , 2000, 267, 1019-1029.	0.2	37
69	How Sweet Are Our Gut Beneficial Bacteria? A Focus on Protein Glycosylation in <i>Lactobacillus</i> . <i>International Journal of Molecular Sciences</i> , 2018, 19, 136.	4.1	37
70	Variation in the levels of the different xylanase inhibitors in grain and flour of 20 French wheat cultivars. <i>Journal of Cereal Science</i> , 2005, 41, 375-379.	3.7	35
71	Structural basis for the role of serine-rich repeat proteins from <i>Lactobacillus reuteri</i> in gut microbe β host interactions. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E2706-E2715.	7.1	35
72	Molecular determinants of substrate and inhibitor specificities of the <i>Penicillium griseofulvum</i> family 11 xylanases. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2009, 1794, 438-445.	2.3	32

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73	Cross-inhibitory activity of cereal protein inhibitors against α -amylases and xylanases. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2003, 1650, 136-144.	2.3	31
74	Secretion, purification, and characterisation of barley α -amylase produced by heterologous gene expression in <i>Aspergillus niger</i> . <i>Applied Microbiology and Biotechnology</i> , 1998, 49, 385-392.	3.6	30
75	Structural and molecular insights into novel surface-exposed mucus adhesins from <i>Lactobacillus reuteri</i> human strains. <i>Molecular Microbiology</i> , 2014, 92, 543-556.	2.5	29
76	^{13}C T-cell-deficient mice show alterations in mucin expression, glycosylation, and goblet cells but maintain an intact mucus layer. <i>American Journal of Physiology - Renal Physiology</i> , 2014, 306, G582-G593.	3.4	27
77	The inhibition specificity of recombinant <i>Penicillium funiculosum</i> xylanase B towards wheat proteinaceous inhibitors. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2004, 1701, 121-128.	2.3	26
78	Mining the “glycode” exploring the spatial distribution of glycans in gastrointestinal mucin using force spectroscopy. <i>FASEB Journal</i> , 2013, 27, 2342-2354.	0.5	26
79	Uncovering a novel molecular mechanism for scavenging sialic acids in bacteria. <i>Journal of Biological Chemistry</i> , 2020, 295, 13724-13736.	3.4	26
80	Molecular cloning and primary structure analysis of porcine pancreatic α -amylase. <i>BBA - Proteins and Proteomics</i> , 1999, 1430, 281-289.	2.1	24
81	Comparison of barley malt α -amylase isozymes 1 and 2: construction of cDNA hybrids by in vivo recombination and their expression in yeast. <i>Gene</i> , 1993, 130, 159-166.	2.2	23
82	Behaviour of family 10 and 11 xylanases towards arabinoxylans with varying structure. <i>Journal of the Science of Food and Agriculture</i> , 2006, 86, 1618-1622.	3.5	22
83	Molecular basis for intestinal mucin recognition by galectin-3 and type lectins. <i>FASEB Journal</i> , 2018, 32, 3301-3320.	0.5	21
84	Functional importance of Asp37 from a family 11 xylanase in the binding to two proteinaceous xylanase inhibitors from wheat. <i>FEMS Microbiology Letters</i> , 2004, 239, 9-15.	1.8	19
85	Genome Sequence of the Vertebrate Gut Symbiont <i>Lactobacillus reuteri</i> ATCC 53608. <i>Journal of Bacteriology</i> , 2011, 193, 4015-4016.	2.2	18
86	The Immunomodulatory Properties of β -2,6 Fructans: A Comprehensive Review. <i>Nutrients</i> , 2021, 13, 1309.	4.1	18
87	The Effects of Human Milk Oligosaccharides on Gut Microbiota, Metabolite Profiles and Host Mucosal Response in Patients with Irritable Bowel Syndrome. <i>Nutrients</i> , 2021, 13, 3836.	4.1	17
88	Differential Epitope Mapping by STD NMR Spectroscopy To Reveal the Nature of Protein-Ligand Contacts. <i>Angewandte Chemie</i> , 2017, 129, 15491-15495.	2.0	16
89	Serine-rich repeat protein adhesins from <i>Lactobacillus reuteri</i> display strain specific glycosylation profiles. <i>Glycobiology</i> , 2019, 29, 45-58.	2.5	15
90	Serine-rich repeat proteins from gut microbes. <i>Gut Microbes</i> , 2020, 11, 102-117.	9.8	15

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91	Deriving Ligand Orientation in Weak Protein-Ligand Complexes by DEER-STD NMR Spectroscopy in the Absence of Protein Chemical Shift Assignment. <i>ChemBioChem</i> , 2019, 20, 340-344.	2.6	14
92	Structure-based mutagenesis of <i>Penicillium griseofulvum</i> xylanase using computational design. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008, 72, 1298-1307.	2.6	13
93	The role of the mucin-glycan foraging <i>Ruminococcus gnavus</i> in the communication between the gut and the brain. <i>Gut Microbes</i> , 2022, 14, 2073784.	9.8	13
94	Multiple evolutionary origins reflect the importance of sialic acid transporters in the colonization potential of bacterial pathogens and commensals. <i>Microbial Genomics</i> , 2021, 7, .	2.0	12
95	Proteinaceous Xylanase Inhibitors: Structure, Function and Evolution. <i>Current Enzyme Inhibition</i> , 2006, 2, 29-35.	0.4	11
96	Mucin-lectin interactions assessed by flow cytometry. <i>Carbohydrate Research</i> , 2010, 345, 1486-1491.	2.3	10
97	Interlaboratory evaluation of plasma N-glycan antennary fucosylation as a clinical biomarker for HNF1A-MODY using liquid chromatography methods. <i>Glycoconjugate Journal</i> , 2021, 38, 375-386.	2.7	10
98	Siglec-7 Mediates Immunomodulation by Colorectal Cancer-Associated <i>Fusobacterium nucleatum</i> ssp. <i>animalis</i> . <i>Frontiers in Immunology</i> , 2021, 12, 744184.	4.8	10
99	The human gut symbiont <i>Ruminococcus gnavus</i> shows specificity to blood group A antigen during mucin glycan foraging: Implication for niche colonisation in the gastrointestinal tract. <i>PLoS Biology</i> , 2021, 19, e3001498.	5.6	10
100	Functional characterization of <i>Penicillium occitanis</i> Pol6 and <i>Penicillium funiculosum</i> GH11 xylanases. <i>Protein Expression and Purification</i> , 2013, 90, 195-201.	1.3	9
101	Structural properties of porcine intestine acylpeptide hydrolase. <i>The Protein Journal</i> , 2003, 22, 183-191.	1.1	8
102	Cloning, sequencing and functional expression of a cDNA encoding porcine pancreatic preprocarboxypeptidase A1. <i>FEBS Journal</i> , 1999, 259, 719-726.	0.2	8
103	Mice deficient in intestinal intraepithelial lymphocytes display an altered intestinal O-glycan profile compared with wild-type littermates. <i>Glycobiology</i> , 2015, 25, 42-54.	2.5	7
104	Membrane-enclosed multienzyme (MEME) synthesis of 2,7-anhydro-sialic acid derivatives. <i>Carbohydrate Research</i> , 2017, 451, 110-117.	2.3	7
105	Structure of the O-Antigen and the Lipid A from the Lipopolysaccharide of <i>Fusobacterium nucleatum</i> ATCC 51191. <i>ChemBioChem</i> , 2021, 22, 1252-1260.	2.6	7
106	Biochemical Basis of Xylooligosaccharide Utilisation by Gut Bacteria. <i>International Journal of Molecular Sciences</i> , 2022, 23, 2992.	4.1	7
107	Isolation and characterisation of a xylanase inhibitor Xip-II gene from durum wheat. <i>Journal of Cereal Science</i> , 2009, 50, 324-331.	3.7	6
108	Development of a novel human intestinal model to elucidate the effect of anaerobic commensals on <i>Escherichia coli</i> infection. <i>DMM Disease Models and Mechanisms</i> , 2022, 15, .	2.4	5

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109	Ascertaining the biochemical function of an essential pectin methylesterase in the gut microbe <i>Bacteroides thetaiotaomicron</i> . <i>Journal of Biological Chemistry</i> , 2020, 295, 18625-18637.	3.4	4
110	Plant proteinaceous inhibitors of carbohydrate-active enzymes. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2004, 1696, 141.	2.3	3
111	Quantification of xylanase inhibitors by immunodetection: the way forward?. <i>Journal of the Science of Food and Agriculture</i> , 2006, 86, 1737-1740.	3.5	3
112	High-level production of recombinant <i>Aspergillus niger</i> cinnamoyl esterase (FAEA) in the methylotrophic yeast <i>Pichia pastoris</i> . <i>FEMS Yeast Research</i> , 2001, 1, 127-132.	2.3	3
113	Development of an exoglycosidase plate-based assay for detecting α 1-3,4 fucosylation biomarker in individuals with HNF1A-MODY. <i>Glycobiology</i> , 2022, 32, 230-238.	2.5	3
114	Interactions of barley α -amylase isozymes with Ca^{2+} , substrates and proteinaceous inhibitors. <i>Biocatalysis and Biotransformation</i> , 2006, 24, 83-93.	2.0	1
115	Enzymes in grain processing. <i>Journal of Cereal Science</i> , 2009, 50, 305.	3.7	0
116	Glycobiology of Host-Microbe Interactions in the Gut. <i>FASEB Journal</i> , 2019, 33, 216.4.	0.5	0
117	Glycans and the Gut Microbiota. , 2022, , .		0