

# Morten Sørli

## List of Publications by Year in descending order

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103  
papers

6,603  
citations

81900

39  
h-index

64796

79  
g-index

106  
all docs

106  
docs citations

106  
times ranked

4615  
citing authors

#	ARTICLE	IF	CITATIONS
1	An Oxidative Enzyme Boosting the Enzymatic Conversion of Recalcitrant Polysaccharides. <i>Science</i> , 2010, 330, 219-222.	12.6	1,059
2	Production of Chitooligosaccharides and Their Potential Applications in Medicine. <i>Marine Drugs</i> , 2010, 8, 1482-1517.	4.6	496
3	Cleavage of cellulose by a CBM33 protein. <i>Protein Science</i> , 2011, 20, 1479-1483.	7.6	317
4	The chitinolytic machinery of <i>Serratia marcescens</i> – a model system for enzymatic degradation of recalcitrant polysaccharides. <i>FEBS Journal</i> , 2013, 280, 3028-3049.	4.7	244
5	Structural and functional characterization of a conserved pair of bacterial cellulose-oxidizing lytic polysaccharide monooxygenases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 8446-8451.	7.1	241
6	Costs and benefits of processivity in enzymatic degradation of recalcitrant polysaccharides. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 18089-18094.	7.1	238
7	NMR structure of a lytic polysaccharide monooxygenase provides insight into copper binding, protein dynamics, and substrate interactions. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 18779-18784.	7.1	236
8	Endo/exo mechanism and processivity of family 18 chitinases produced by <i>Serratia marcescens</i> . <i>FEBS Journal</i> , 2006, 273, 491-503.	4.7	235
9	Structural and Functional Characterization of a Lytic Polysaccharide Monooxygenase with Broad Substrate Specificity. <i>Journal of Biological Chemistry</i> , 2015, 290, 22955-22969.	3.4	157
10	Lytic Polysaccharide Monooxygenases in Enzymatic Processing of Lignocellulosic Biomass. <i>ACS Catalysis</i> , 2019, 9, 4970-4991.	11.2	145
11	Aromatic Residues in the Catalytic Center of Chitinase A from <i>Serratia marcescens</i> Affect Processivity, Enzyme Activity, and Biomass Converting Efficiency. <i>Journal of Biological Chemistry</i> , 2009, 284, 10610-10617.	3.4	142
12	Kinetics of H <sub>2</sub> O <sub>2</sub> -driven degradation of chitin by a bacterial lytic polysaccharide monooxygenase. <i>Journal of Biological Chemistry</i> , 2018, 293, 523-531.	3.4	130
13	Interactions of a fungal lytic polysaccharide monooxygenase with $\beta$ -glucan substrates and cellobiose dehydrogenase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 5922-5927.	7.1	126
14	Comparative Study of Two Chitin-Active and Two Cellulose-Active AA10-Type Lytic Polysaccharide Monooxygenases. <i>Biochemistry</i> , 2014, 53, 1647-1656.	2.5	124
15	Polysaccharide degradation by lytic polysaccharide monooxygenases. <i>Current Opinion in Structural Biology</i> , 2019, 59, 54-64.	5.7	105
16	Methylation of the N-terminal histidine protects a lytic polysaccharide monooxygenase from auto-oxidative inactivation. <i>Protein Science</i> , 2018, 27, 1636-1650.	7.6	91
17	Hallmarks of Processivity in Glycoside Hydrolases from Crystallographic and Computational Studies of the <i>Serratia marcescens</i> Chitinases. <i>Journal of Biological Chemistry</i> , 2012, 287, 36322-36330.	3.4	89
18	Towards a molecular-level theory of carbohydrate processivity in glycoside hydrolases. <i>Current Opinion in Biotechnology</i> , 2014, 27, 96-106.	6.6	89

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19	Mutational Effects on Transglycosylating Activity of Family 18 Chitinases and Construction of a Hypertransglycosylating Mutant. <i>Biochemistry</i> , 2011, 50, 5693-5703.	2.5	82
20	Measuring Processivity. <i>Methods in Enzymology</i> , 2012, 510, 69-95.	1.0	80
21	Antifungal effect of chito-oligosaccharides with different degrees of polymerization. <i>European Journal of Plant Pathology</i> , 2015, 141, 147-158.	1.7	67
22	Electron Inventory, Kinetic Assignment (En), Structure, and Bonding of Nitrogenase Turnover Intermediates with C <sub>2</sub> H <sub>2</sub> and CO. <i>Journal of the American Chemical Society</i> , 2005, 127, 15880-15890.	13.7	65
23	Degradation of Chitosans with a Family 46 Chitosanase from <i>Streptomyces coelicolor</i> A3(2). <i>Biomacromolecules</i> , 2010, 11, 2487-2497.	5.4	63
24	Unraveling the roles of the reductant and free copper ions in LPMO kinetics. <i>Biotechnology for Biofuels</i> , 2021, 14, 28.	6.2	62
25	Kinetic insights into the role of the reductant in H <sub>2</sub> O <sub>2</sub> -driven degradation of chitin by a bacterial lytic polysaccharide monooxygenase. <i>Journal of Biological Chemistry</i> , 2019, 294, 1516-1528.	3.4	60
26	An in vitro investigation of endocrine disrupting effects of the mycotoxin alternariol. <i>Toxicology and Applied Pharmacology</i> , 2013, 271, 64-71.	2.8	59
27	Single-turnover of Nitric-oxide Synthase in the Presence of 4-Amino-tetrahydrobiopterin. <i>Journal of Biological Chemistry</i> , 2003, 278, 48602-48610.	3.4	58
28	Biotransformation of zearalenone and zearalenols to their major glucuronide metabolites reduces estrogenic activity. <i>Toxicology in Vitro</i> , 2015, 29, 575-581.	2.4	58
29	Synthesis and Structure of the Ruthenium(II) Complexes [( $\eta$ -C <sub>5</sub> Me <sub>5</sub> )Ru(NO)(bipy)] <sup>2+</sup> and [( $\eta$ -C <sub>5</sub> Me <sub>5</sub> )Ru(NO)(dppz)] <sup>2+</sup> . DNA Cleavage by an Organometallic dppz Complex (bipy = 2,2'-Bipyridine; dppz) <a href="#">10.1002/anie.201507843</a>	15.7	14
30	Human Chitotriosidase-Catalyzed Hydrolysis of Chitosan. <i>Biochemistry</i> , 2012, 51, 487-495.	2.5	53
31	Characterization of an Intermediate in the Reduction of Acetylene by the Nitrogenase $\hat{I}$ -Gln195MoFe Protein by Q-band EPR and <sup>13</sup> C,1H ENDOR. <i>Journal of the American Chemical Society</i> , 2000, 122, 5582-5587.	13.7	50
32	Expression and Characterization of Endochitinase C from <i>Serratia marcescens</i> BJL200 and Its Purification by a One-Step General Chitinase Purification Method. <i>Bioscience, Biotechnology and Biochemistry</i> , 2008, 72, 715-723.	1.3	50
33	Inhibition of angiogenesis by chito-oligosaccharides with specific degrees of acetylation and polymerization. <i>Carbohydrate Polymers</i> , 2012, 89, 511-518.	10.2	49
34	Inhibition of Fungal Plant Pathogens by Synergistic Action of Chito-Oligosaccharides and Commercially Available Fungicides. <i>PLoS ONE</i> , 2014, 9, e93192.	2.5	49
35	The Predominant Molecular State of Bound Enzyme Determines the Strength and Type of Product Inhibition in the Hydrolysis of Recalcitrant Polysaccharides by Processive Enzymes. <i>Journal of Biological Chemistry</i> , 2015, 290, 11678-11691.	3.4	47
36	Natural substrate assay for chitinases using high-performance liquid chromatography: A comparison with existing assays. <i>Analytical Biochemistry</i> , 2007, 363, 128-134.	2.4	43

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37	Mechanistic basis of substrateâ€™O <sub>2</sub> coupling within a chitin-active lytic polysaccharide monoxygenase: An integrated NMR/EPR study. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 19178-19189.	7.1	42
38	Adherence Inhibition of Enteropathogenic <i>Escherichia coli</i> by Chitooligosaccharides with Specific Degrees of Acetylation and Polymerization. Journal of Agricultural and Food Chemistry, 2013, 61, 2748-2754.	5.2	41
39	Chitin oligosaccharide binding to a family GH19 chitinase from the moss <i>Bryumâ€™coronatum</i> . FEBS Journal, 2011, 278, 3991-4001.	4.7	40
40	Inhibition of a family 18 chitinase by chitooligosaccharides. Carbohydrate Polymers, 2008, 74, 41-49.	10.2	38
41	Identification of a High-Affinity-Binding Oligosaccharide by (+) Nanoelectrospray Quadrupole Time-of-Flight Tandem Mass Spectrometry of a Noncovalent Enzymeâ€™Ligand Complex. Angewandte Chemie - International Edition, 2006, 45, 2429-2434.	13.8	36
42	Determination of substrate binding energies in individual subsites of a family 18 chitinase. FEBS Letters, 2010, 584, 4581-4585.	2.8	36
43	Enzyme processivity changes with the extent of recalcitrant polysaccharide degradation. FEBS Letters, 2014, 588, 4620-4624.	2.8	36
44	Antifungal activity of well-defined chito-oligosaccharide preparations against medically relevant yeasts. PLoS ONE, 2019, 14, e0210208.	2.5	35
45	One-electron oxidation of catecholamines generates free radicals with an in vitro toxicity correlating with their lifetime. Free Radical Biology and Medicine, 2006, 41, 1266-1271.	2.9	34
46	Slow Off-rates and Strong Product Binding Are Required for Processivity and Efficient Degradation of Recalcitrant Chitin by Family 18 Chitinases. Journal of Biological Chemistry, 2015, 290, 29074-29085.	3.4	33
47	Signatures of activation parameters reveal substrate-dependent rate determining steps in polysaccharide turnover by a family 18 chitinase. Carbohydrate Polymers, 2010, 81, 14-20.	10.2	32
48	Evidence of Two Distinct Oxygen Complexes of Reduced Endothelial Nitric Oxide Synthase. Journal of Biological Chemistry, 2004, 279, 19824-19831.	3.4	31
49	Thermodynamic Analysis of Allosamidin Binding to a Family 18 Chitinase. Biochemistry, 2007, 46, 12347-12354.	2.5	31
50	Kinetic Characterization of a Putatively Chitin-Active LPMO Reveals a Preference for Soluble Substrates and Absence of Monoxygenase Activity. ACS Catalysis, 2021, 11, 11685-11695.	11.2	31
51	Fast and Specific Peroxygenase Reactions Catalyzed by Fungal Mono-Copper Enzymes. Biochemistry, 2021, 60, 3633-3643.	2.5	31
52	Variant MoFe proteins of <i>Azotobacter vinelandii</i> : effects of carbon monoxide on electron paramagnetic resonance spectra generated during enzyme turnover. Journal of Biological Inorganic Chemistry, 2005, 10, 394-406.	2.6	30
53	Detection of a New Radical and FeMo-Cofactor EPR Signal during Acetylene Reduction by the Î±-H195Q Mutant of Nitrogenase. Journal of the American Chemical Society, 1999, 121, 9457-9458.	13.7	28
54	The Roles of Three <i>Serratia marcescens</i> Chitinases in Chitin Conversion Are Reflected in Different Thermodynamic Signatures of Allosamidin Binding. Journal of Physical Chemistry B, 2010, 114, 6144-6149.	2.6	28

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55	Activation of enzymatic chitin degradation by a lytic polysaccharide monooxygenase. <i>Carbohydrate Research</i> , 2015, 407, 166-169.	2.3	28
56	Modulation of Lck Function through Multisite Docking to T Cell-specific Adapter Protein. <i>Journal of Biological Chemistry</i> , 2008, 283, 21909-21919.	3.4	25
57	Key Residues Affecting Transglycosylation Activity in Family 18 Chitinases: Insights into Donor and Acceptor Subsites. <i>Biochemistry</i> , 2018, 57, 4325-4337.	2.5	25
58	Use of Stopped-Flow Spectrophotometry to Establish Midpoint Potentials for Redox Proteins. <i>Analytical Biochemistry</i> , 2000, 287, 118-125.	2.4	24
59	Heat capacity changes in heme protein–ligand interactions. <i>Thermochimica Acta</i> , 2007, 464, 24-28.	2.7	24
60	Can we make Chitosan by Enzymatic Deacetylation of Chitin?. <i>Molecules</i> , 2019, 24, 3862.	3.8	24
61	An investigation of the endocrine disrupting potential of enniatin B using in vitro bioassays. <i>Toxicology Letters</i> , 2015, 233, 84-94.	0.8	23
62	Aromatic-Mediated Carbohydrate Recognition in Processive <i>Serratia marcescens</i> Chitinases. <i>Journal of Physical Chemistry B</i> , 2016, 120, 1236-1249.	2.6	23
63	Substrate positioning in chitinase A, a processive chito-biohydrolase from <i>Serratia marcescens</i> . <i>FEBS Letters</i> , 2011, 585, 2339-2344.	2.8	21
64	Analysis of productive binding modes in the human chitotriosidase. <i>FEBS Letters</i> , 2013, 587, 3508-3513.	2.8	21
65	The kinase I $\kappa$ K and the adaptor TSA $\delta$ change the specificity of the kinase Lck in T cells by promoting the phosphorylation of Tyr <sup>192</sup> . <i>Science Signaling</i> , 2014, 7, ra118.	3.6	21
66	The effect of the carbohydrate binding module on substrate degradation by the human chitotriosidase. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2015, 1854, 1494-1501.	2.3	21
67	Processivity, Substrate Positioning, and Binding: The Role of Polar Residues in a Family 18 Glycoside Hydrolase. <i>Biochemistry</i> , 2015, 54, 7292-7306.	2.5	20
68	Thermodynamic Relationships with Processivity in <i>Serratia marcescens</i> Family 18 Chitinases. <i>Journal of Physical Chemistry B</i> , 2015, 119, 9601-9613.	2.6	20
69	Thermodynamics of tunnel formation upon substrate binding in a processive glycoside hydrolase. <i>Archives of Biochemistry and Biophysics</i> , 2017, 620, 35-42.	3.0	20
70	Analytical Tools for Characterizing Cellulose-Active Lytic Polysaccharide Monooxygenases (LPMOs). <i>Methods in Molecular Biology</i> , 2018, 1796, 219-246.	0.9	19
71	The use of isothermal titration calorimetry to determine the thermodynamics of metal ion binding to low-cost sorbents. <i>Thermochimica Acta</i> , 2010, 501, 119-121.	2.7	18
72	Processivity and substrate-binding in family 18 chitinases. <i>Biocatalysis and Biotransformation</i> , 2012, 30, 353-365.	2.0	18

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73	Treatment of recalcitrant crystalline polysaccharides with lytic polysaccharide monoxygenase relieves the need for glycoside hydrolase processivity. <i>Carbohydrate Research</i> , 2019, 473, 66-71.	2.3	18
74	Analysis of Noncovalent Chitinase-Chito-Oligosaccharide Complexes by Infrared-Matrix Assisted Laser Desorption Ionization and Nanoelectrospray Ionization Mass Spectrometry. <i>Analytical Chemistry</i> , 2011, 83, 4030-4036.	6.5	16
75	Relative quantification of the proteomic changes associated with the mycotoxin zearalenone in the H295R steroidogenesis model. <i>Toxicon</i> , 2011, 58, 533-542.	1.6	16
76	Tetrahydrobiopterin as Combined Electron/Proton Donor in Nitric Oxide Biosynthesis: Cryogenic UV-Vis and EPR Detection of Reaction Intermediates. <i>Methods in Enzymology</i> , 2005, 396, 456-466.	1.0	15
77	Cytosol protein regulation in H295R steroidogenesis model induced by the zearalenone metabolites, $\hat{1}\pm$ - and $\hat{1}^2$ -zearalenol. <i>Toxicon</i> , 2012, 59, 17-24.	1.6	14
78	Human Chitotriosidase: Catalytic Domain or Carbohydrate Binding Module, Who's Leading HCHT's Biological Function. <i>Scientific Reports</i> , 2017, 7, 2768.	3.3	14
79	Human Chitotriosidase Is an Endo-Processive Enzyme. <i>PLoS ONE</i> , 2017, 12, e0171042.	2.5	14
80	Antibiotic saving effect of combination therapy through synergistic interactions between well-characterized chito-oligosaccharides and commercial antifungals against medically relevant yeasts. <i>PLoS ONE</i> , 2019, 14, e0227098.	2.5	13
81	The role of active site aromatic residues in substrate degradation by the human chitotriosidase. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2016, 1864, 242-247.	2.3	12
82	Crystal structure and thermodynamic dissection of chitin oligosaccharide binding to the LysM module of chitinase-A from <i>Pteris ryukyuensis</i> . <i>Biochemical and Biophysical Research Communications</i> , 2017, 494, 736-741.	2.1	12
83	Dissecting factors that contribute to ligand-binding energetics for family 18 chitinases. <i>Thermochimica Acta</i> , 2010, 511, 189-193.	2.7	11
84	Effect of enzyme processivity on the efficacy of a competitive chitinase inhibitor. <i>Carbohydrate Polymers</i> , 2010, 82, 779-785.	10.2	11
85	Enzyme assay for chitinase catalyzed hydrolysis of tetra-N-acetylchitotetraose by isothermal titration calorimetry. <i>Thermochimica Acta</i> , 2007, 454, 144-146.	2.7	10
86	Using chitosan to understand chitinases and the role of processivity in the degradation of recalcitrant polysaccharides. <i>Reactive and Functional Polymers</i> , 2020, 148, 104488.	4.1	10
87	Novel molecular biological tools for the efficient expression of fungal lytic polysaccharide monoxygenases in <i>Pichia pastoris</i> . <i>Biotechnology for Biofuels</i> , 2021, 14, 122.	6.2	10
88	CO exchange of the oxyferrous complexes of endothelial nitric-oxide synthase oxygenase domain in the presence of 4-amino-tetrahydrobiopterin. <i>Journal of Inorganic Biochemistry</i> , 2004, 98, 1217-1222.	3.5	9
89	The SH3 domains of the protein kinases ITK and LCK compete for adjacent sites on T cell-specific adapter protein. <i>Journal of Biological Chemistry</i> , 2019, 294, 15480-15494.	3.4	9
90	Chemoenzymatic Synthesis of Chito-oligosaccharides with Alternating N-Acetylglucosamine and D-Glucosamine. <i>Biochemistry</i> , 2020, 59, 4581-4590.	2.5	9

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91	Thermodynamic Signatures of Substrate Binding for Three Thermobifida fusca Cellulases with Different Modes of Action. <i>Biochemistry</i> , 2019, 58, 1648-1659.	2.5	8
92	Thermodynamic analysis of allosamidin binding to the human chitotriosidase. <i>Thermochimica Acta</i> , 2013, 565, 146-150.	2.7	7
93	Structural and Thermodynamic Signatures of Ligand Binding to the Enigmatic Chitinase D of <i>Serratia proteamaculans</i> . <i>Journal of Physical Chemistry B</i> , 2019, 123, 2270-2279.	2.6	7
94	NMR and Fluorescence Spectroscopies Reveal the Preorganized Binding Site in Family 14 Carbohydrate-Binding Module from Human Chitotriosidase. <i>ACS Omega</i> , 2019, 4, 21975-21984.	3.5	7
95	The directionality of processive enzymes acting on recalcitrant polysaccharides is reflected in the kinetic signatures of oligomer degradation. <i>FEBS Letters</i> , 2015, 589, 1807-1812.	2.8	5
96	Synergistic Antifungal Activity of Chito-Oligosaccharides and Commercial Antifungals on Biofilms of Clinical Candida Isolates. <i>Journal of Fungi (Basel, Switzerland)</i> , 2021, 7, 718.	3.5	5
97	Polar residues lining the binding cleft of a <i>Serratia marcescens</i> family 18 chitinase position the substrate for attack and stabilize associative interactions. <i>Molecular Physics</i> , 2019, 117, 3664-3682.	1.7	4
98	Kinetic relationships with processivity in <i>Serratia marcescens</i> family 18 glycoside hydrolases. <i>Biochemical and Biophysical Research Communications</i> , 2020, 521, 120-124.	2.1	3
99	Genomic and Proteomic Study of <i>Andreprevotia ripae</i> Isolated from an Anthill Reveals an Extensive Repertoire of Chitinolytic Enzymes. <i>Journal of Proteome Research</i> , 2021, 20, 4041-4052.	3.7	3
100	The characterization of the glucono- $\delta$ -lactone-carboxylic acid equilibrium in the products of chitin-active lytic polysaccharide monoxygenases. <i>Journal of Chemical Thermodynamics</i> , 2017, 106, 10-15.	2.0	1
101	The effect of carbohydrate binding modules and linkers on inhibitor binding to family 18 glycoside hydrolases. <i>Journal of Chemical Thermodynamics</i> , 2018, 125, 220-224.	2.0	1
102	Thermodynamic insights into the role of aromatic residues in chito-oligosaccharide binding to the transglycosylating chitinase-D from <i>Serratia proteamaculans</i> . <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2020, 1868, 140414.	2.3	0
103	The Ribonucleotide Reductase system from <i>Bacillus cereus</i> . <i>FASEB Journal</i> , 2007, 21, A640.	0.5	0