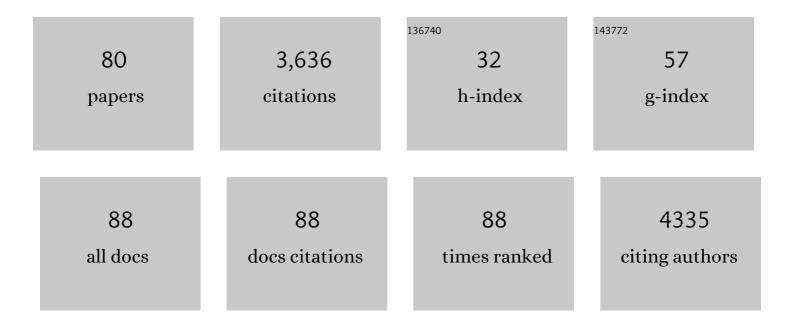
Thomas B Thompson

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Growth differentiation factor 9:bone morphogenetic protein 15 heterodimers are potent regulators of ovarian functions. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, E776-85.	3.3	251
2	The Structure of the Follistatin:Activin Complex Reveals Antagonism of Both Type I and Type II Receptor Binding. Developmental Cell, 2005, 9, 535-543.	3.1	247
3	Structures of an ActRIIB:activin A complex reveal a novel binding mode for TGF-beta ligand:receptor interactions. EMBO Journal, 2003, 22, 1555-1566.	3.5	188
4	Structural Studies of the Parainfluenza Virus 5 Hemagglutinin-Neuraminidase Tetramer in Complex with Its Receptor, Sialyllactose. Structure, 2005, 13, 803-815.	1.6	187
5	The Structure of Apolipoprotein A-I in High Density Lipoproteins. Journal of Biological Chemistry, 2007, 282, 22249-22253.	1.6	176
6	Circulating Growth Differentiation Factor 11/8 Levels Decline With Age. Circulation Research, 2016, 118, 29-37.	2.0	161
7	Biochemistry and Biology of GDF11 and Myostatin. Circulation Research, 2016, 118, 1125-1142.	2.0	155
8	The structure of myostatin:follistatin 288: insights into receptor utilization and heparin binding. EMBO Journal, 2009, 28, 2662-2676.	3.5	148
9	The 1.5-Ã Resolution Crystal Structure of Bacterial Luciferase in Low Salt Conditions. Journal of Biological Chemistry, 1996, 271, 21956-21968.	1.6	122
10	Structural basis for potency differences between GDF8 and GDF11. BMC Biology, 2017, 15, 19.	1.7	90
11	Neural Network Prediction of the HIV-1 Protease Cleavage Sites. Journal of Theoretical Biology, 1995, 177, 369-379.	0.8	88
12	Structure of Myostatin·Follistatin-like 3. Journal of Biological Chemistry, 2012, 287, 1043-1053.	1.6	76
13	Three-Dimensional Structure of ATP:Corrinoid Adenosyltransferase fromSalmonella typhimuriumin Its Free State, Complexed with MgATP, or Complexed with Hydroxycobalamin and MgATPâ€,â€j. Biochemistry, 2001, 40, 361-374.	1.2	72
14	Evolution of Enzymatic Activity in the Enolase Superfamily:  Structure of o-Succinylbenzoate Synthase from Escherichia coli in Complex with Mg2+ and o-Succinylbenzoate,. Biochemistry, 2000, 39, 10662-10676.	1.2	71
15	Beta A versus beta B: is it merely a matter of expression?. Molecular and Cellular Endocrinology, 2004, 225, 9-17.	1.6	70
16	Molecular Biology of Inhibin Action. Seminars in Reproductive Medicine, 2004, 22, 269-276.	0.5	66
17	The Structure of FSTL3·Activin A Complex. Journal of Biological Chemistry, 2008, 283, 32831-32838.	1.6	63
18	The DAN family: Modulators of TGF $\hat{\epsilon}\hat{\mathbf{f}}^2$ signaling and beyond. Protein Science, 2014, 23, 999-1012.	3.1	62

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19	A thumbwheel mechanism for APOA1 activation of LCAT activity in HDL[S]. Journal of Lipid Research, 2018, 59, 1244-1255.	2.0	59
20	Structure of the human myostatin precursor and determinants of growth factor latency. EMBO Journal, 2018, 37, 367-383.	3.5	58
21	Structural and Functional Analysis of Tetracenomycin F2 Cyclase from Streptomyces glaucescens. Journal of Biological Chemistry, 2004, 279, 37956-37963.	1.6	54
22	Members of the DAN Family Are BMP Antagonists That Form Highly Stable Noncovalent Dimers. Journal of Molecular Biology, 2012, 424, 313-327.	2.0	54
23	Structure of Protein Related to Dan and Cerberus: Insights into the Mechanism of Bone Morphogenetic Protein Antagonism. Structure, 2013, 21, 1417-1429.	1.6	54
24	A consensus model of human apolipoprotein A-I in its monomeric and lipid-free state. Nature Structural and Molecular Biology, 2017, 24, 1093-1099.	3.6	54
25	Three-Dimensional Structure of Adenosylcobinamide Kinase/Adenosylcobinamide Phosphate Guanylyltransferase fromSalmonella typhimuriumDetermined to 2.3 à Resolutionâ€,‡. Biochemistry, 1998, 37, 7686-7695.	1.2	46
26	Structural characterization of an activin class ternary receptor complex reveals a third paradigm for receptor specificity. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 15505-15513.	3.3	46
27	Activin A forms a non-signaling complex with ACVR1 and type II Activin/BMP receptors via its finger 2 tip loop. ELife, 2020, 9, .	2.8	45
28	Cytochrome b5 reductase–cytochrome b5 as an active P450 redox enzyme system in Phanerochaete chrysosporium: Atypical properties and in vivo evidence of electron transfer capability to CYP63A2. Archives of Biochemistry and Biophysics, 2011, 509, 26-32.	1.4	40
29	BMP Antagonist Gremlin 2 Limits Inflammation After Myocardial Infarction. Circulation Research, 2016, 119, 434-449.	2.0	40
30	The Structure of Dimeric Apolipoprotein A-IV and Its Mechanism of Self-Association. Structure, 2012, 20, 767-779.	1.6	39
31	Structure of Gremlin-2 in Complex with GDF5 Gives Insight into DAN-Family-Mediated BMP Antagonism. Cell Reports, 2016, 16, 2077-2086.	2.9	37
32	Myostatin Stimulates, Not Inihibits, C2C12 Myoblast Proliferation. Endocrinology, 2014, 155, 670-675.	1.4	35
33	Alternative Binding Modes Identified for Growth and Differentiation Factor-associated Serum Protein (GASP) Family Antagonism of Myostatin. Journal of Biological Chemistry, 2015, 290, 7506-7516.	1.6	35
34	Structural perspective of BMP ligands and signaling. Bone, 2020, 140, 115549.	1.4	35
35	Three-Dimensional Structure of Adenosylcobinamide Kinase/Adenosylcobinamide Phosphate Guanylyltransferase (CobU) Complexed with GMP:Â Evidence for a Substrate-Induced Transferase Active Siteâ€,‡. Biochemistry, 1999, 38, 12995-13005.	1.2	33
36	Structure of Neuroblastoma Suppressor of Tumorigenicity 1 (NBL1). Journal of Biological Chemistry, 2015, 290, 4759-4771.	1.6	32

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37	Activins bind and signal via bone morphogenetic protein receptor type II (BMPR2) in immortalized gonadotrope-like cells. Cellular Signalling, 2013, 25, 2717-2726.	1.7	30
38	Molecular characterization of latent GDF8 reveals mechanisms of activation. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E866-E875.	3.3	30
39	Characterization of Follistatin-Type Domains and Their Contribution to Myostatin and Activin A Antagonism. Molecular Endocrinology, 2012, 26, 1167-1178.	3.7	28
40	Analysis of the Adenosylcobinamide Kinase/Adenosylcobinamide-phosphate Guanylyltransferase (CobU) Enzyme of Salmonella typhimurium LT2. Journal of Biological Chemistry, 2000, 275, 27576-27586.	1.6	26
41	The Structure of Human Apolipoprotein A-IV as Revealed by Stable Isotope-assisted Cross-linking, Molecular Dynamics, and Small Angle X-ray Scattering. Journal of Biological Chemistry, 2014, 289, 5596-5608.	1.6	26
42	MuSK is a BMP co-receptor that shapes BMP responses and calcium signaling in muscle cells. Science Signaling, 2016, 9, ra87.	1.6	26
43	Structural biology of the TGFÎ ² family. Experimental Biology and Medicine, 2019, 244, 1530-1546.	1.1	26
44	Mutations in GDF11 and the extracellular antagonist, Follistatin, as a likely cause of Mendelian forms of orofacial clefting in humans. Human Mutation, 2019, 40, 1813-1825.	1.1	26
45	Structure of AMH bound to AMHR2 provides insight into a unique signaling pair in the TGF-β family. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	26
46	The orphan ligand, activin C, signals through activin receptor-like kinase 7. ELife, 0, 11, .	2.8	21
47	Development of a Small-Molecule Screening Method for Inhibitors of Cellular Response to Myostatin and Activin A. Journal of Biomolecular Screening, 2013, 18, 837-844.	2.6	20
48	Myostatin Attenuation In Vivo Reduces Adiposity, but Activates Adipogenesis. Endocrinology, 2016, 157, 282-291.	1.4	17
49	Coordinated Proliferation and Differentiation of Human-Induced Pluripotent Stem Cell-Derived Cardiac Progenitor Cells Depend on Bone Morphogenetic Protein Signaling Regulation by GREMLIN 2. Stem Cells and Development, 2017, 26, 678-693.	1.1	17
50	Structural Basis for a Functional Antagonist in the Transforming Growth Factor Î ² Superfamily. Journal of Biological Chemistry, 2005, 280, 40177-40186.	1.6	16
51	An Evaluation of the Crystal Structure of C-terminal Truncated Apolipoprotein A-I in Solution Reveals Structural Dynamics Related to Lipid Binding. Journal of Biological Chemistry, 2016, 291, 5439-5451.	1.6	16
52	Crystal structure of the WFIKKN2 follistatin domain reveals insight into how it inhibits growth differentiation factor 8 (GDF8) and GDF11. Journal of Biological Chemistry, 2019, 294, 6333-6343.	1.6	13
53	Analysis of the Interaction between Heparin and Follistatin and Heparin and Follistatin–Ligand Complexes Using Surface Plasmon Resonance. Biochemistry, 2012, 51, 6797-6803.	1.2	12
54	Myostatin regulates pituitary development and hepatic IGF1. American Journal of Physiology - Endocrinology and Metabolism, 2019, 316, E1036-E1049.	1.8	12

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55	Mutational Analysis of the Putative Anti-Müllerian Hormone (AMH) Binding Interface on its Type II Receptor, AMHR2. Endocrinology, 2020, 161, .	1.4	12
56	Role of Conserved Proline Residues in Human Apolipoprotein A-IV Structure and Function. Journal of Biological Chemistry, 2015, 290, 10689-10702.	1.6	11
57	Expression and purification of recombinant protein related to DAN and cerberus (PRDC). Protein Expression and Purification, 2012, 82, 389-395.	0.6	10
58	Improving the diffraction of apoA-IV crystals through extreme dehydration. Acta Crystallographica Section F: Structural Biology Communications, 2012, 68, 105-110.	0.7	10
59	Small-angle X-ray Scattering of Apolipoprotein A-IV Reveals the Importance of Its Termini for Structural Stability. Journal of Biological Chemistry, 2013, 288, 4854-4866.	1.6	10
60	Analysis and identification of the Grem2 heparin/heparan sulfate-binding motif. Biochemical Journal, 2017, 474, 1093-1107.	1.7	10
61	The anti-sigma factor MucA of Pseudomonas aeruginosa: Dramatic differences of a mucA22 vs. a ΔmucA mutant in anaerobic acidified nitrite sensitivity of planktonic and biofilm bacteria in vitro and during chronic murine lung infection. PLoS ONE, 2019, 14, e0216401.	1.1	10
62	Molecular Mechanisms of AMH Signaling. Frontiers in Endocrinology, 0, 13, .	1.5	10
63	Visceral adipose tissue remodeling in pancreatic ductal adenocarcinoma cachexia: the role of activin A signaling. Scientific Reports, 2022, 12, 1659.	1.6	8
64	Fibronectin-based scaffold domain proteins that bind myostatin: a patent evaluation of WO2014043344. Expert Opinion on Therapeutic Patents, 2015, 25, 619-624.	2.4	7
65	Characterization of the different oligomeric states of the DAN family antagonists SOSTDC1 and SOST. Biochemical Journal, 2020, 477, 3167-3182.	1.7	7
66	Structures of activin ligand traps using natural sets of type I and type II TGFÎ ² receptors. IScience, 2022, 25, 103590.	1.9	7
67	Analysis of the loop-helix interaction in bundle motif protein structures. The Protein Journal, 1995, 14, 559-566.	1.1	6
68	Reply to Mottershead et al.: GDF9:BMP15 heterodimers are potent regulators of ovarian functions. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, E2258-E2258.	3.3	6
69	Deletion of Gremlin-2 alters estrous cyclicity and disrupts female fertility in mice. Biology of Reproduction, 2021, 105, 1205-1220.	1.2	6
70	Amino Acid 72 of Mouse and Human GDF9 Mature Domain Is Responsible for Altered Homodimer Bioactivities but Has Subtle Effects on GDF9:BMP15 Heterodimer Activities1. Biology of Reproduction, 2014, 91, 142.	1.2	4
71	Functional recombinant apolipoprotein A5 that is stable at high concentrations at physiological pH. Journal of Lipid Research, 2020, 61, 244-251.	2.0	4
72	Characterization of tolloid-mediated cleavage of the GDF8 procomplex. Biochemical Journal, 2021, 478, 1733-1747.	1.7	4

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73	Heparin-mediated dimerization of follistatin. Experimental Biology and Medicine, 2021, 246, 467-482.	1.1	3
74	New Insight Into Hyperemesis Gravidarum and a Potential Role for GDF15. Endocrinology, 2018, 159, 2698-2700.	1.4	2
75	FSTL3-Neutralizing Antibodies Enhance Glucose-Responsive Insulin Secretion in Dysfunctional Male Mouse and Human Islets. Endocrinology, 2021, 162, .	1.4	2
76	BMP and BMP Regulation: Structure and Function. , 2017, , 73-111.		1
77	Structural biology: Gaining atomic level insight into the biological function of macromolecules. Experimental Biology and Medicine, 2019, 244, 1507-1509.	1.1	1
78	Activins and Inhibins in Female Reproduction. , 2018, , 202-210.		0
79	Front Cover, Volume 40, Issue 10. Human Mutation, 2019, 40, i.	1.1	0
80	Structure of dimeric apoAâ€IV: basis for HDL model. FASEB Journal, 2011, 25, 938.1.	0.2	0