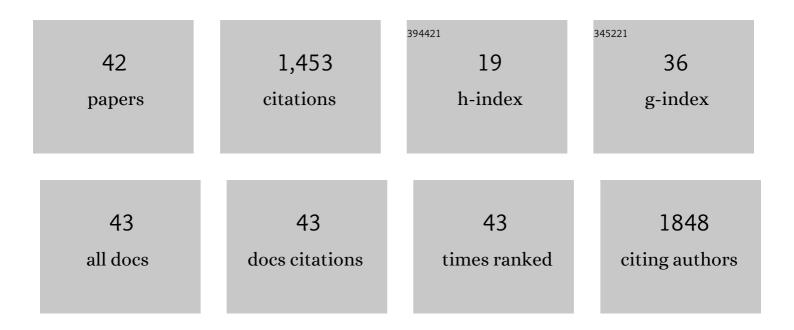
Sarah E Bondos

List of Publications by Year in descending order

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#	Article	lF	CITATIONS
1	Intrinsically disordered proteins play diverse roles in cell signaling. Cell Communication and Signaling, 2022, 20, 20.	6.5	68
2	On the roles of intrinsically disordered proteins and regions in cell communication and signaling. Cell Communication and Signaling, 2021, 19, 88.	6.5	57
3	Context-dependent HOX transcription factor function in health and disease. Progress in Molecular Biology and Translational Science, 2020, 174, 225-262.	1.7	4
4	Evolution of the activation domain in a Hox transcription factor. International Journal of Developmental Biology, 2018, 62, 745-753.	0.6	4
5	Generating Novel Materials Using the Intrinsically Disordered Protein Ubx. Methods in Enzymology, 2018, 611, 583-605.	1.0	0
6	Mechanically-Tunable, Protein-Based Materials Can be Functionalized with Other Proteins and with DNA. Biophysical Journal, 2016, 110, 338a.	0.5	0
7	Culture of Tumorigenic Cells on Protein Fibers Reveals Metastatic Cell Behaviors. Biomacromolecules, 2016, 17, 3790-3799.	5.4	4
8	Separating full-length protein from aggregating proteolytic products using filter flow-through purification. Analytical Biochemistry, 2016, 514, 8-11.	2.4	0
9	Functionalization of Ultrabithorax Materials with Vascular Endothelial Growth Factor Enhances Angiogenic Activity. Biomacromolecules, 2016, 17, 3558-3569.	5.4	7
10	Rethinking gene regulatory networks in light of alternative splicing, intrinsically disordered protein domains, and post-translational modifications. Frontiers in Cell and Developmental Biology, 2015, 3, 8.	3.7	96
11	Identification of Multiple Dityrosine Bonds in Materials Composed of the <i>Drosophila</i> Protein Ultrabithorax. Advanced Functional Materials, 2015, 25, 5988-5998.	14.9	7
12	Materials composed of the <i>Drosophila</i> Hox protein Ultrabithorax are biocompatible and nonimmunogenic. Journal of Biomedical Materials Research - Part A, 2015, 103, 1546-1553.	4.0	6
13	CDK8-Cyclin C Mediates Nutritional Regulation of Developmental Transitions through the Ecdysone Receptor in Drosophila. PLoS Biology, 2015, 13, e1002207.	5.6	38
14	The Effect of Protein Fusions on the Production and Mechanical Properties of Proteinâ€Based Materials. Advanced Functional Materials, 2015, 25, 1442-1450.	14.9	8
15	Flexibility and Disorder in Gene Regulation: LacI/GalR and Hox Proteins. Journal of Biological Chemistry, 2015, 290, 24669-24677.	3.4	19
16	Intrinsically disordered proteins and multicellular organisms. Seminars in Cell and Developmental Biology, 2015, 37, 44-55.	5.0	128
17	Materials composed of the <i>Drosophila melanogaster</i> protein ultrabithorax are cytocompatible. Journal of Biomedical Materials Research - Part A, 2014, 102, 97-104.	4.0	7
18	Measuring Hox-DNA Binding by Electrophoretic Mobility Shift Analysis. Methods in Molecular Biology, 2014, 1196, 211-230.	0.9	3

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19	The Intrinsically Disordered Regions of the Drosophila melanogaster Hox Protein Ultrabithorax Select Interacting Proteins Based on Partner Topology. PLoS ONE, 2014, 9, e108217.	2.5	16
20	What's in a name? Why these proteins are intrinsically disordered. Intrinsically Disordered Proteins, 2013, 1, e24157.	1.9	226
21	Identifying Solubility-Promoting Buffers for Intrinsically Disordered Proteins Prior to Purification. , 2012, 896, 415-427.		10
22	Ultrabithorax, an Intrinsically Disordered Protein, Selects Protein Interactions by Topology. Biophysical Journal, 2012, 102, 633a.	0.5	0
23	Roles for Intrinsic Disorder and Fuzziness in Generating Context-specific Function in Ultrabithorax, a Hox Transcription Factor. Advances in Experimental Medicine and Biology, 2012, 725, 86-105.	1.6	10
24	On the Design of Composite Protein–Quantum Dot Biomaterials via Self-Assembly. Biomacromolecules, 2011, 12, 3629-3637.	5.4	26
25	Generating Context-Specific Functions with Intrinsically Disordered Domains. Biophysical Journal, 2011, 100, 185a.	O.5	0
26	Dynamic protein–DNA recognition: beyond what can be seen. Trends in Biochemical Sciences, 2011, 36, 415-423.	7.5	137
27	Media composition influences yeast one- and two-hybrid results. Biological Procedures Online, 2011, 13, 6.	2.9	4
28	Functionalization and Patterning of Proteinâ€Based Materials Using Active Ultrabithorax Chimeras. Advanced Functional Materials, 2011, 21, 2633-2640.	14.9	13
29	Size Dictates Mechanical Properties for Protein Fibers Self-Assembled by the <i>Drosophila</i> Hox Transcription Factor Ultrabithorax. Biomacromolecules, 2010, 11, 3644-3651.	5.4	15
30	Reverse Regulation: Controlling Intrinsically Disordered Domains with Structured Elements. Biophysical Journal, 2010, 98, 258a.	0.5	0
31	The Drosophila Transcription Factor Ultrabithorax Self-Assembles into Protein-Based Biomaterials with Multiple Morphologies. Biomacromolecules, 2009, 10, 829-837.	5.4	19
32	Internal Regulatory Interactions Determine DNA Binding Specificity by a Hox Transcription Factor. Journal of Molecular Biology, 2009, 390, 760-774.	4.2	36
33	Team-teaching a current events-based biology course for nonmajors. Biochemistry and Molecular Biology Education, 2008, 36, 22-27.	1.2	8
34	Multiple Intrinsically Disordered Sequences Alter DNA Binding by the Homeodomain of the Drosophila Hox Protein Ultrabithorax. Journal of Biological Chemistry, 2008, 283, 20874-20887.	3.4	81
35	Variations on a Theme: Hox and Wnt Combinatorial Regulation During Animal Development. Science's STKE: Signal Transduction Knowledge Environment, 2006, 2006, pe38-pe38.	3.9	22
36	Methods for Measuring Protein Aggregation. Current Analytical Chemistry, 2006, 2, 157-170.	1.2	31

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37	Physical and Genetic Interactions Link Hox Function with Diverse Transcription Factors and Cell Signaling Proteins. Molecular and Cellular Proteomics, 2006, 5, 824-834.	3.8	47
38	Hox Transcription Factor Ultrabithorax Ib Physically and Genetically Interacts with Disconnected Interacting Protein 1, a Double-stranded RNA-binding Protein. Journal of Biological Chemistry, 2004, 279, 26433-26444.	3.4	25
39	Detection and prevention of protein aggregation before, during, and after purification. Analytical Biochemistry, 2003, 316, 223-231.	2.4	201
40	Transcription Activation by Ultrabithorax Ib Protein Requires a Predicted α-Helical Regionâ€. Biochemistry, 2002, 41, 2774-2785.	2.5	21
41	Combinatorial Transcriptional Regulation: The Interaction of Transcription Factors and Cell Signaling Molecules with Homeodomain Proteins in Drosophila Development. Critical Reviews in Eukaryotic Gene Expression, 2001, 11, 28.	0.9	23
42	High-pressure denaturation of apomyoglobin. BBA - Proteins and Proteomics, 2000, 1480, 353-364.	2.1	25