Jordi Bella

List of Publications by Year in descending order

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159585 223800 5,529 49 30 46 h-index citations g-index papers 49 49 49 6111 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Hydroxyapatite-decorated Fmoc-hydrogel as a bone-mimicking substrate for osteoclast differentiation and culture. Acta Biomaterialia, 2022, 138, 144-154.	8.3	15
2	Incorporation of Natural and Recombinant Collagen Proteins within Fmoc-Based Self-Assembling Peptide Hydrogels. Gels, 2022, 8, 254.	4.5	6
3	Role of OSCAR Signaling in Osteoclastogenesis and Bone Disease. Frontiers in Cell and Developmental Biology, 2021, 9, 641162.	3.7	31
4	Fibrillar Collagens. Sub-Cellular Biochemistry, 2017, 82, 457-490.	2.4	117
5	The open architecture of HD-PTP phosphatase provides new insights into the mechanism of regulation of ESCRT function. Scientific Reports, 2017, 7, 9151.	3.3	22
6	Structural Basis for Selective Interaction between the ESCRT Regulator HD-PTP and UBAP1. Structure, 2016, 24, 2115-2126.	3.3	22
7	Collagen structure: new tricks from a very old dog. Biochemical Journal, 2016, 473, 1001-1025.	3.7	182
8	Analysis of flexible multidomain glycoproteins with SAXS, analytical ultracentrifugation, and torsion-angle molecular dynamics. Acta Crystallographica Section A: Foundations and Advances, 2015, 71, s45-s45.	0.1	0
9	Cadherin flexibility provides a key difference between desmosomes and adherens junctions. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 5395-5400.	7.1	37
10	A first census of collagen interruptions: Collagen's own stutters and stammers. Journal of Structural Biology, 2014, 186, 438-450.	2.8	16
11	Collagen-Like Proteins in Pathogenic E. coli Strains. PLoS ONE, 2012, 7, e37872.	2.5	32
12	A new method for describing the helical conformation of collagen: Dependence of the triple helical twist on amino acid sequence. Journal of Structural Biology, 2010, 170, 377-391.	2.8	63
13	Decorin Core Protein (Decoron) Shape Complements Collagen Fibril Surface Structure and Mediates Its Binding. PLoS ONE, 2009, 4, e7028.	2.5	126
14	Quantitative analysis and prediction of curvature in leucineâ€rich repeat proteins. Proteins: Structure, Function and Bioinformatics, 2009, 77, 342-358.	2.6	16
15	The leucine-rich repeat structure. Cellular and Molecular Life Sciences, 2008, 65, 2307-2333.	5.4	392
16	LRRCE: a leucine-rich repeat cysteine capping motif unique to the chordate lineage. BMC Genomics, 2008, 9, 599.	2.8	39
17	A Role for Soluble <i>N</i> -Ethylmaleimide-sensitive Factor Attachment Protein Receptor Complex Dimerization during Neurosecretion. Molecular Biology of the Cell, 2008, 19, 3379-3389.	2.1	12
18	Collagens at a glance. Journal of Cell Science, 2007, 120, 1955-1958.	2.0	653

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19	Analysis of pre-mRNA and pre-rRNA processing factor Snu13p structure and mutants. Biochemical and Biophysical Research Communications, 2007, 360, 857-862.	2.1	10
20	A Cys-capping motif unique to small leucine-rich repeat proteins and proteoglycans of the extracellular matrix. BMC Systems Biology, 2007, 1 , .	3.0	1
21	Conformational Effects of Gly–X–Gly Interruptions in the Collagen Triple Helix. Journal of Molecular Biology, 2006, 362, 298-311.	4.2	61
22	Structural correlations in the family of small leucine-rich repeat proteins and proteoglycans. Journal of Structural Biology, 2006, 155, 294-305.	2.8	180
23	On the calculation of the binding force between decorin and collagen. Journal of Biomechanics, 2006, 39, 1159-1160.	2.1	5
24	Calpha-HO = C hydrogen bonds contribute to the specificity of RGD cell-adhesion interactions., $2005, 5, 4$.		24
25	Crystal structure of the dimeric protein core of decorin, the archetypal small leucine-rich repeat proteoglycan. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 15633-15638.	7.1	193
26	Integrin structure: heady advances in ligand binding, but activation still makes the knees wobble. Trends in Biochemical Sciences, 2003, 28, 313-320.	7.5	123
27	Structural Basis of Type VI Collagen Dimer Formation. Journal of Biological Chemistry, 2003, 278, 15326-15332.	3.4	47
28	Structure of an Integrin-Ligand Complex Deduced from Solution X-ray Scattering and Site-directed Mutagenesis. Journal of Biological Chemistry, 2003, 278, 39993-39999.	3.4	93
29	The crystal and molecular structure of a collagen-like peptide with A biologically relevant sequence. Journal of Molecular Biology, 2001, 311, 131-147.	4.2	179
30	Interaction of Coxsackievirus A21 with Its Cellular Receptor, ICAM-1. Journal of Virology, 2001, 75, 2444-2451.	3.4	78
31	ICAM-1 receptors and cold viruses. Pharmacochemistry Library, 2000, , 291-297.	0.1	2
32	Integrin–collagen complex: a metal–glutamate handshake. Structure, 2000, 8, R121-R126.	3.3	36
33	ICAM-1 receptors and cold viruses. Pharmaceutica Acta Helvetiae, 2000, 74, 291-297.	1.2	41
34	Cell Recognition and Entry by Rhino- and Enteroviruses. Virology, 2000, 269, 239-247.	2.4	62
35	Staggered molecular packing in crystals of a collagen-like peptide with a single charged pair. Journal of Molecular Biology, 2000, 301, 1191-1205.	4.2	197
36	The dynamics of receptor recognition by human rhinoviruses: Response. Trends in Microbiology, 2000, 8, 254.	7.7	0

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37	Sequence dependent conformational variations of collagen triple-helical structure. Nature Structural Biology, 1999, 6, 454-457.	9.7	277
38	X-ray Crystallographic Structure of the Norwalk Virus Capsid. Science, 1999, 286, 287-290.	12.6	820
39	The structure of the two amino-terminal domains of human intercellular adhesion molecule-1 suggests how it functions as a rhinovirus receptor. Virus Research, 1999, 62, 107-117.	2.2	19
40	Structural studies of two rhinovirus serotypes complexed with fragments of their cellular receptor. EMBO Journal, 1999, 18, 6249-6259.	7.8	189
41	Review: Rhinoviruses and Their ICAM Receptors. Journal of Structural Biology, 1999, 128, 69-74.	2.8	54
42	A General Phasing Algorithm for Multiple MAD and MIR Data. Acta Crystallographica Section D: Biological Crystallography, 1998, 54, 159-174.	2.5	7
43	X-ray crystallographic determination of a collagen-like peptide with the repeating sequence (Pro-Pro-Gly). Journal of Molecular Biology, 1998, 280, 623-638.	4.2	166
44	Disrupted Collagen Architecture in the Crystal Structure of a Triple-Helical Peptide with a Gly → Ala Substitution. Connective Tissue Research, 1996, 35, 401-406.	2.3	30
45	Crystallographic Evidence for Cα–H···O=C Hydrogen Bonds in a Collagen Triple Helix. Journal of Molecular Biology, 1996, 264, 734-742.	4.2	209
46	Hydration structure of a collagen peptide. Structure, 1995, 3, 893-906.	3.3	570
47	Chain conformation in polyretropeptides: Quantum mechanical and empirical force field calculations on 2,6,8-trioxo-3,5,9-triazadecane, a model compound for poly(retro-glycine). Biopolymers, 1995, 35, 257-269.	2.4	11
48	Glycine residues induce a helical structure in polyamides. Polymer, 1994, 35, 1291-1297.	3.8	34
49	Crystal structure of a helical oligopeptide model of polyglycine II and of other polyamides: Acetyl-(glycyl-β-alanyl)2-NHpropyl. Biopolymers, 1992, 32, 643-648.	2.4	30