

Inma Ponte

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6916544/publications.pdf>

Version: 2024-02-01

32
papers

924
citations

430754

18
h-index

454834

30
g-index

33
all docs

33
docs citations

33
times ranked

840
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Towards understanding the Regulation of Histone H1 Somatic Subtypes with OMICs. <i>Journal of Molecular Biology</i> , 2021, 433, 166734. | 2.0 | 3 |
| 2 | A CON-based NMR assignment strategy for pro-rich intrinsically disordered proteins with low signal dispersion: the C-terminal domain of histone H1.0 as a case study. <i>Journal of Biomolecular NMR</i> , 2018, 72, 139-148. | 1.6 | 12 |
| 3 | Post-translational modifications of the intrinsically disordered terminal domains of histone H1: effects on secondary structure and chromatin dynamics. <i>Chromosoma</i> , 2017, 126, 83-91. | 1.0 | 25 |
| 4 | Complex Evolutionary History of the Mammalian Histone H1.1-H1.5 Gene Family. <i>Molecular Biology and Evolution</i> , 2017, 34, msw241. | 3.5 | 20 |
| 5 | The subtype-specific role of histone H1.0 in cancer cell differentiation and intratumor heterogeneity. <i>Translational Cancer Research</i> , 2017, 6, S414-S417. | 0.4 | 3 |
| 6 | Interplay between histone H1 structure and function. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2016, 1859, 444-454. | 0.9 | 36 |
| 7 | Histone H1 Favors Folding and Parallel Fibrillar Aggregation of the 1-42 Amyloid- β Peptide. <i>Langmuir</i> , 2015, 31, 6782-6790. | 1.6 | 13 |
| 8 | Linker histone partial phosphorylation: effects on secondary structure and chromatin condensation. <i>Nucleic Acids Research</i> , 2015, 43, 4463-4476. | 6.5 | 35 |
| 9 | Identification of novel post-translational modifications in linker histones from chicken erythrocytes. <i>Journal of Proteomics</i> , 2015, 113, 162-177. | 1.2 | 28 |
| 10 | Sequence conservation of linker histones between chicken and mammalian species. <i>Data in Brief</i> , 2014, 1, 60-64. | 0.5 | 6 |
| 11 | Contribution of hydrophobic interactions to the folding and fibrillation of histone H1 and its carboxy-terminal domain. <i>Journal of Structural Biology</i> , 2012, 180, 101-109. | 1.3 | 12 |
| 12 | Secondary structure of protamine in sperm nuclei: an infrared spectroscopy study. <i>BMC Structural Biology</i> , 2011, 11, 14. | 2.3 | 28 |
| 13 | An inducible helix-Gly-Gly-helix motif in the N-terminal domain of histone H1e: A CD and NMR study. <i>Protein Science</i> , 2009, 11, 214-220. | 3.1 | 28 |
| 14 | Role of Charge Neutralization in the Folding of the Carboxy-Terminal Domain of Histone H1. <i>Journal of Physical Chemistry B</i> , 2009, 113, 12061-12066. | 1.2 | 25 |
| 15 | Phosphorylation of the carboxy-terminal domain of histone H1: effects on secondary structure and DNA condensation. <i>Nucleic Acids Research</i> , 2008, 36, 4719-4726. | 6.5 | 71 |
| 16 | Macromolecular Crowding Induces a Molten Globule State in the C-Terminal Domain of Histone H1. <i>Biophysical Journal</i> , 2007, 93, 2170-2177. | 0.2 | 51 |
| 17 | Differential affinity of mammalian histone H1 somatic subtypes for DNA and chromatin. <i>BMC Biology</i> , 2007, 5, 22. | 1.7 | 68 |
| 18 | DNA-induced Secondary Structure of the Carboxyl-terminal Domain of Histone H1. <i>Journal of Biological Chemistry</i> , 2005, 280, 32141-32147. | 1.6 | 86 |

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 19 | The preferential binding of histone H1 to DNA scaffold-associated regions is determined by its C-terminal domain. <i>Nucleic Acids Research</i> , 2004, 32, 6111-6119. | 6.5 | 34 |
| 20 | Sequence Complexity of Histone H1 Subtypes. <i>Molecular Biology and Evolution</i> , 2003, 20, 371-380. | 3.5 | 35 |
| 21 | Sequence and analysis of the 5' flanking and 5' untranslated regions of the rat N-methyl-D-aspartate receptor 2A gene. <i>Gene</i> , 2002, 295, 135-142. | 1.0 | 13 |
| 22 | DNA-induced α -Helical Structure in the NH ₂ -terminal Domain of Histone H1. <i>Journal of Biological Chemistry</i> , 2001, 276, 46429-46435. | 1.6 | 57 |
| 23 | Induction of Secondary Structure in a COOH-terminal Peptide of Histone H1 by Interaction with the DNA. <i>Journal of Biological Chemistry</i> , 2001, 276, 30898-30903. | 1.6 | 63 |
| 24 | A helix-turn motif in the C-terminal domain of histone H1. <i>Protein Science</i> , 2000, 9, 627-636. | 3.1 | 38 |
| 25 | Isolation and characterization of a 28-kDa HMG-like protein that binds to A/T-rich distal promoter regions of zein genes. <i>Plant Science</i> , 1998, 135, 31-38. | 1.7 | 3 |
| 26 | Evolution of the vertebrate H1 histone class: evidence for the functional differentiation of the subtypes. <i>Molecular Biology and Evolution</i> , 1998, 15, 702-708. | 3.5 | 72 |
| 27 | Sequence simplicity and evolution of the 3' untranslated region of the histone H1 ^o Gene. <i>Journal of Molecular Evolution</i> , 1996, 43, 125-134. | 0.8 | 4 |
| 28 | Cloning and analysis of the coding region of the histone H1 ^o -encoding gene from rat PC12 cells. <i>Gene</i> , 1995, 166, 313-316. | 1.0 | 5 |
| 29 | Narrow A/T-rich zones present at the distal 5' flanking sequences of the zein genes Zc1 and Zc2 bind a unique 30 kDa HMG-like protein. <i>Plant Molecular Biology</i> , 1994, 26, 1893-1906. | 2.0 | 7 |
| 30 | Transcriptional activation of Histone H1 ^o during neuronal terminal differentiation. <i>Developmental Brain Research</i> , 1994, 80, 35-44. | 2.1 | 10 |
| 31 | DNA sequence of the gene encoding the Zc1 protein from <i>Zea mays</i> W64 A. <i>Nucleic Acids Research</i> , 1990, 18, 6425-6425. | 6.5 | 12 |
| 32 | Sequence analysis of a genomic clone encoding a Zc2 protein from <i>Zea mays</i> W64 A. <i>Nucleic Acids Research</i> , 1990, 18, 6426-6426. | 6.5 | 21 |