Hedi Hegyi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11812355/publications.pdf

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16 papers	1,775 citations	16 h-index	940533 16 g-index
16	16	16	2453
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Increased structural disorder of proteins encoded on human sex chromosomes. Molecular BioSystems, 2012, 8, 229-236.	2.9	22
2	The relationship between proteome size, structural disorder and organism complexity. Genome Biology, 2011, 12, R120.	9.6	161
3	Verification of alternative splicing variants based on domain integrity, truncation length and intrinsic protein disorder. Nucleic Acids Research, 2011, 39, 1208-1219.	14.5	46
4	Intrinsic Structural Disorder Confers Cellular Viability on Oncogenic Fusion Proteins. PLoS Computational Biology, 2009, 5, e1000552.	3.2	74
5	Identification and correction of abnormal, incomplete and mispredicted proteins in public databases. BMC Bioinformatics, 2008, 9, 353.	2.6	55
6	Intrinsically Disordered Proteins Display No Preference for Chaperone Binding In Vivo. PLoS Computational Biology, 2008, 4, e1000017.	3.2	48
7	Comprehensive viral oligonucleotide probe design using conserved protein regions. Nucleic Acids Research, 2008, 36, e3-e3.	14.5	21
8	The implications of alternative splicing in the ENCODE protein complement. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 5495-5500.	7.1	206
9	Panmicrobial Oligonucleotide Array for Diagnosis of Infectious Diseases. Emerging Infectious Diseases, 2007, 13, 73-81.	4.3	298
10	Structural disorder promotes assembly of protein complexes. BMC Structural Biology, 2007, 7, 65.	2.3	96
11	Molecular Fossils in the Human Genome: Identification and Analysis of the Pseudogenes in Chromosomes 21 and 22. Genome Research, 2002, 12, 272-280.	5.5	167
12	SNPs on human chromosomes 21 and 22 – analysis in terms of protein features and pseudogenes. Pharmacogenomics, 2002, 3, 393-402.	1.3	16
13	Structural genomics analysis: Characteristics of atypical, common, and horizontally transferred folds. Proteins: Structure, Function and Bioinformatics, 2002, 47, 126-141.	2.6	33
14	Annotation Transfer for Genomics: Measuring Functional Divergence in Multi-Domain Proteins. Genome Research, 2001, 11, 1632-1640.	5 . 5	69
15	The relationship between protein structure and function: a comprehensive survey with application to the yeast genome. Journal of Molecular Biology, 1999, 288, 147-164.	4.2	384
16	Comparing genomes in terms of protein structure: surveys of a finite parts list. FEMS Microbiology Reviews, 1998, 22, 277-304.	8.6	79