## Jacopo Vertemara

List of Publications by Year in descending order

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933447 996975 16 248 10 15 citations h-index g-index papers 16 16 16 396 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Investigations of the electronic-molecular structure of bio-inorganic systems using modern methods of quantum chemistry. Inorganica Chimica Acta, 2022, 532, 120728.	2.4	1
2	Molecular Dynamics Simulations Reveal Structural Interconnections within Sec14-PH Bipartite Domain from Human Neurofibromin. International Journal of Molecular Sciences, 2022, 23, 5707.	4.1	0
3	DNA binding modes influence Rap1 activity in the regulation of telomere length and MRX functions at DNA ends. Nucleic Acids Research, 2020, 48, 2424-2441.	14.5	7
4	Rational Design of Fe <sub>2</sub> (νâ€PR <sub>2</sub> ) <sub>2</sub> (L) <sub>6</sub> Coordination Compounds Featuring Tailored Potential Inversion. ChemPhysChem, 2020, 21, 2279-2292.	2.1	11
5	Functional and structural insights into the MRX/MRN complex, a key player in recognition and repair of DNA double-strand breaks. Computational and Structural Biotechnology Journal, 2020, 18, 1137-1152.	4.1	31
6	The ATP-bound conformation of the Mrell–Rad50 complex is essential for Tell/ATM activation. Nucleic Acids Research, 2019, 47, 3550-3567.	14.5	35
7	Structurally distinct Mre11 domains mediate MRX functions in resection, end-tethering and DNA damage resistance. Nucleic Acids Research, 2018, 46, 2990-3008.	14.5	34
8	Structural characterization of the nitrogenase molybdenum-iron protein with the substrate acetylene trapped near the active site. Journal of Inorganic Biochemistry, 2018, 180, 129-134.	3.5	21
9	On the photochemistry of Fe <sub>2</sub> (edt)(CO) <sub>4</sub> (PMe <sub>3</sub> ) <sub>2</sub> , a [FeFe]â€hydrogenase model: A DFT/TDDFT investigation. International Journal of Quantum Chemistry, 2018, 118, e25537.	2.0	9
10	Local unwinding of double-strand DNA ends by the MRX complex promotes Exo1 processing activity. Molecular and Cellular Oncology, 2018, 5, e1511208.	0.7	6
11	Evolutionary Analysis Provides Insight Into the Origin and Adaptation of HCV. Frontiers in Microbiology, 2018, 9, 854.	<b>3.</b> 5	15
12	The <scp>MRX</scp> complex regulates Exo1 resection activity by altering <scp>DNA</scp> end structure. EMBO Journal, 2018, 37, .	7.8	21
13	Epigallocatechin-3-gallate and related phenol compounds redirect the amyloidogenic aggregation pathway of ataxin-3 towards non-toxic aggregates and prevent toxicity in neural cells and Caenorhabditis elegans animal model. Human Molecular Genetics, 2017, 26, 3271-3284.	2.9	21
14	Natural Selection at the Brush-Border: Adaptations to Carbohydrate Diets in Humans and Other Mammals. Genome Biology and Evolution, 2015, 7, 2569-2584.	2.5	16
15	Diverse selective regimes shape genetic diversity at <i>ADAR</i> genes and at their coding targets. RNA Biology, 2015, 12, 149-161.	3.1	9
16	Albuminoid Genes: Evolving at the Interface of Dispensability and Selection. Genome Biology and Evolution, 2014, 6, 2983-2997.	2.5	11