## Luca Pinzi

## List of Publications by Year in descending order

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623574 642610 1,503 26 14 23 citations h-index g-index papers 28 28 28 1492 docs citations all docs times ranked citing authors

#	Article	IF	Citations
1	Dual Targeting Strategies on Histone Deacetylase 6 (HDAC6) and Heat Shock Protein 90 (Hsp90). Current Medicinal Chemistry, 2022, 29, 1474-1502.	1.2	11
2	On the development of B-Raf inhibitors acting through innovative mechanisms. F1000Research, 2022, 11, 237.	0.8	1
3	Inhibitors of histone deacetylase 6 based on a novel 3-hydroxy-isoxazole zinc binding group. Journal of Enzyme Inhibition and Medicinal Chemistry, 2021, 36, 2080-2086.	2.5	5
4	Drug Repurposing and Polypharmacology to Fight SARS-CoV-2 Through Inhibition of the Main Protease. Frontiers in Pharmacology, 2021, 12, 636989.	1.6	28
5	Prediction of activity and selectivity profiles of human Carbonic Anhydrase inhibitors using machine learning classification models. Journal of Cheminformatics, 2021, 13, 18.	2.8	12
6	Antifungal Activity and DNA Topoisomerase Inhibition of Hydrolysable Tannins from Punica granatum L International Journal of Molecular Sciences, 2021, 22, 4175.	1.8	21
7	LigAdvisor: a versatile and user-friendly web-platform for drug design. Nucleic Acids Research, 2021, 49, W326-W335.	6.5	12
8	Chemoinformatics Analyses of Tau Ligands Reveal Key Molecular Requirements for the Identification of Potential Drug Candidates against Tauopathies. Molecules, 2021, 26, 5039.	1.7	6
9	Hydroxamic Acid Derivatives: From Synthetic Strategies to Medicinal Chemistry Applications. ACS Omega, 2021, 6, 21843-21849.	1.6	42
10	Design and Synthesis of Hsp90 Inhibitors with Bâ€Raf and PDHK1 Multiâ€Target Activity. ChemistryOpen, 2021, 10, 1177-1185.	0.9	5
11	Identification of Target Associations for Polypharmacology from Analysis of Crystallographic Ligands of the Protein Data Bank. Journal of Chemical Information and Modeling, 2020, 60, 372-390.	2.5	19
12	Repositioning Natural Products in Drug Discovery. Molecules, 2020, 25, 1154.	1.7	34
13	Design of Dual Inhibitors of Histone Deacetylase 6 and Heat Shock Protein 90. ACS Omega, 2020, 5, 11473-11480.	1.6	27
14	In Silico Repositioning of Cannabigerol as a Novel Inhibitor of the Enoyl Acyl Carrier Protein (ACP) Reductase (InhA). Molecules, 2019, 24, 2567.	1.7	22
15	Refinement and Rescoring of Virtual Screening Results. Frontiers in Chemistry, 2019, 7, 498.	1.8	48
16	Molecular Docking: Shifting Paradigms in Drug Discovery. International Journal of Molecular Sciences, 2019, 20, 4331.	1.8	890
17	Evaluation of Amides, Carbamates, Sulfonamides, and Ureas of 4-Prop-2-ynylidenecycloalkylamine as Potent, Selective, and Bioavailable Negative Allosteric Modulators of Metabotropic Glutamate Receptor 5. Journal of Medicinal Chemistry, 2019, 62, 1246-1273.	2.9	5
18	Identification of 4â€arylâ€1 <i>H</i> à€pyrrole[2,3â€b]pyridine derivatives for the development of new Bâ€Raf inhibitors. Chemical Biology and Drug Design, 2018, 92, 1382-1386.	1.5	3

#	Article	IF	CITATIONS
19	Structure–Activity Relationships of Hexahydrocyclopenta[ <i>&gt;c</i> ) quinoline Derivatives as Allosteric Inhibitors of CDK2 and EGFR. ChemMedChem, 2018, 13, 2627-2634.	1.6	23
20	Identification of small-molecule EGFR allosteric inhibitors by high-throughput docking. Future Medicinal Chemistry, 2018, 10, 1545-1553.	1.1	21
21	Selection of protein conformations for structure-based polypharmacology studies. Drug Discovery Today, 2018, 23, 1889-1896.	3.2	22
22	On the Integration of In Silico Drug Design Methods for Drug Repurposing. Frontiers in Pharmacology, 2017, 8, 298.	1.6	178
23	Computational polypharmacology comes of age. Frontiers in Pharmacology, 2015, 6, 157.	1.6	61
24	Identification of potential biological targets of oxindole scaffolds via in silico repositioning strategies. F1000Research, 0, 11, 217.	0.8	0
25	Identification of potential biological targets of oxindole scaffolds via in silico repositioning strategies. F1000Research, 0, 11, 217.	0.8	0
26	On the development of B-Raf inhibitors acting through innovative mechanisms. F1000Research, 0, 11, 237.	0.8	4