

# Anna Vulpetti

## List of Publications by Year in descending order

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55  
papers

3,369  
citations

172386

29  
h-index

161767

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56  
all docs

56  
docs citations

56  
times ranked

4492  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of compounds with binding affinity to proteins via magnetization transfer from bulk water. <i>Journal of Biomolecular NMR</i> , 2000, 18, 65-68.	1.6	412
2	Comparability of Mixed IC50 Data – A Statistical Analysis. <i>PLoS ONE</i> , 2013, 8, e61007.	1.1	211
3	The Experimental Uncertainty of Heterogeneous Public <i>i&gt;K&lt;/i&gt;&lt;sub&gt;i&lt;/sub&gt; Data. <i>Journal of Medicinal Chemistry</i>, 2012, 55, 5165-5173.</i>	2.9	183
4	Fluorine as a Hydrogen–Bond Acceptor: Experimental Evidence and Computational Calculations. <i>Chemistry - A European Journal</i> , 2014, 20, 11058-11068.	1.7	153
5	3-Aminopyrazole Inhibitors of CDK2/Cyclin A as Antitumor Agents. 1. Lead Finding. <i>Journal of Medicinal Chemistry</i> , 2004, 47, 3367-3380.	2.9	150
6	Potent and Selective Aurora Inhibitors Identified by the Expansion of a Novel Scaffold for Protein Kinase Inhibition. <i>Journal of Medicinal Chemistry</i> , 2005, 48, 3080-3084.	2.9	147
7	Assessment of Docking Poses:– Interactions-Based Accuracy Classification (IBAC) versus Crystal Structure Deviations. <i>Journal of Chemical Information and Computer Sciences</i> , 2004, 44, 871-881.	2.8	116
8	Ligand-Based Fluorine NMR Screening: Principles and Applications in Drug Discovery Projects. <i>Journal of Medicinal Chemistry</i> , 2019, 62, 2218-2244.	2.9	115
9	Fluorine local environment: from screening to drug design. <i>Drug Discovery Today</i> , 2012, 17, 890-897.	3.2	113
10	Design and NMR-Based Screening of LEF, a Library of Chemical Fragments with Different Local Environment of Fluorine. <i>Journal of the American Chemical Society</i> , 2009, 131, 12949-12959.	6.6	112
11	Identification of <i>N&lt;/i&gt;,1,4,4-Tetramethyl-8-[[4-(4-methylpiperazin-1-yl)phenyl]amino]-4,5-dihydro-1 <i>H&lt;/i&gt;-pyrazolo[4,3-<i>h&lt;/i&gt;]quinazoline-3-carbonyl (PHA-848125), a Potent, Orally Available Cyclin Dependent Kinase Inhibitor. <i>Journal of Medicinal Chemistry</i>, 2009, 52, 5152-5163.</i></i></i>	2.9	111
12	3-Aminopyrazole Inhibitors of CDK2/Cyclin A as Antitumor Agents. 2. Lead Optimization. <i>Journal of Medicinal Chemistry</i> , 2005, 48, 2944-2956.	2.9	98
13	Predicting Polypharmacology by Binding Site Similarity: From Kinases to the Protein Universe. <i>Journal of Chemical Information and Modeling</i> , 2010, 50, 1418-1431.	2.5	93
14	Discovery of LOU064 (Remibrutinib), a Potent and Highly Selective Covalent Inhibitor of Bruton’s Tyrosine Kinase. <i>Journal of Medicinal Chemistry</i> , 2020, 63, 5102-5118.	2.9	92
15	Fluorine–Protein Interactions and <sup>19</sup> F–...NMR Isotropic Chemical Shifts: An Empirical Correlation with Implications for Drug Design. <i>ChemMedChem</i> , 2011, 6, 104-114.	1.6	90
16	Sequence and structural analysis of kinase ATP pocket residues. <i>Il Farmaco</i> , 2004, 59, 759-765.	0.9	87
17	Weak Intermolecular Hydrogen Bonds with Fluorine: Detection and Implications for Enzymatic/Chemical Reactions, Chemical Properties, and Ligand/Protein Fluorine NMR Screening. <i>Chemistry - A European Journal</i> , 2016, 22, 7592-7601.	1.7	71
18	Intermolecular and Intramolecular Hydrogen Bonds Involving Fluorine Atoms: Implications for Recognition, Selectivity, and Chemical Properties. <i>ChemMedChem</i> , 2012, 7, 262-272.	1.6	70

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19	Small-molecule factor D inhibitors targeting the alternative complement pathway. <i>Nature Chemical Biology</i> , 2016, 12, 1105-1110.	3.9	68
20	Tautomer Preference in PDB Complexes and its Impact on Structure-Based Drug Discovery. <i>Journal of Chemical Information and Modeling</i> , 2010, 50, 1062-1074.	2.5	62
21	Optimization of 6,6-dimethyl pyrrolo[3,4-c]pyrazoles: Identification of PHA-793887, a potent CDK inhibitor suitable for intravenous dosing. <i>Bioorganic and Medicinal Chemistry</i> , 2010, 18, 1844-1853.	1.4	58
22	3-Amino-1,4,5,6-tetrahydropyrrolo[3,4-c]pyrazoles: A new class of CDK2 inhibitors. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2006, 16, 1084-1090.	1.0	56
23	NMR-Based Quality Control Approach for the Identification of False Positives and False Negatives in High Throughput Screening. <i>Current Drug Discovery Technologies</i> , 2006, 3, 115-124.	0.6	56
24	Polyfluorinated Amino Acids for Sensitive <sup>19</sup> F NMR-Based Screening and Kinetic Measurements. <i>Journal of the American Chemical Society</i> , 2007, 129, 5665-5672.	6.6	48
25	Design and Generation of Highly Diverse Fluorinated Fragment Libraries and their Efficient Screening with Improved <sup>19</sup> F NMR Methodology. <i>ChemMedChem</i> , 2013, 8, 2057-2069.	1.6	48
26	Structure-Based Approaches to Improve Selectivity: CDK2 <sup>vs</sup> GSK3 <sup>β</sup> Binding Site Analysis. <i>Journal of Chemical Information and Modeling</i> , 2005, 45, 1282-1290.	2.5	45
27	Technical and practical aspects of <sup>19</sup> F NMR-based screening: toward sensitive high-throughput screening with rapid deconvolution. <i>Magnetic Resonance in Chemistry</i> , 2012, 50, 592-597.	1.1	38
28	Identification of Potent Pyrazolo[4,3- <i>h</i> ]quinazoline-3-carboxamides as Multi-Cyclin-Dependent Kinase Inhibitors. <i>Journal of Medicinal Chemistry</i> , 2010, 53, 2171-2187.	2.9	36
29	Combined use of computational chemistry, NMR screening, and X-ray crystallography for identification and characterization of fluorophilic protein environments. <i>Proteins: Structure, Function and Bioinformatics</i> , 2010, 78, 3281-3291.	1.5	30
30	Discovery of Highly Potent and Selective Small-Molecule Reversible Factor D Inhibitors Demonstrating Alternative Complement Pathway Inhibition <i>in Vivo</i> . <i>Journal of Medicinal Chemistry</i> , 2017, 60, 5717-5735.	2.9	27
31	Chemogenomics in drug discovery: computational methods based on the comparison of binding sites. <i>Future Medicinal Chemistry</i> , 2012, 4, 1971-1979.	1.1	26
32	Synthesis and Biological Evaluation of New Triazolo- and Imidazopyridine ROR <sup>β</sup> Inverse Agonists. <i>ChemMedChem</i> , 2016, 11, 2640-2648.	1.6	26
33	Comprehensive and High-Throughput Exploration of Chemical Space Using Broadband <sup>19</sup> F NMR-Based Screening. <i>Angewandte Chemie - International Edition</i> , 2020, 59, 14809-14817. <sup>7.2</sup>		24
34	Subpocket Analysis Method for Fragment-Based Drug Discovery. <i>Journal of Chemical Information and Modeling</i> , 2013, 53, 131-141.	2.5	23
35	<sup>19</sup> F NMR chemical shift prediction with fluorine fingerprint descriptor. <i>Journal of Fluorine Chemistry</i> , 2010, 131, 570-577.	0.9	22
36	Structure-Based Library Design and Fragment Screening for the Identification of Reversible Complement Factor D Protease Inhibitors. <i>Journal of Medicinal Chemistry</i> , 2017, 60, 1946-1958.	2.9	22

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37	Optimizing a Weakly Binding Fragment into a Potent ROR $\beta$ Inverse Agonist with Efficacy in an in Vivo Inflammation Model. <i>Journal of Medicinal Chemistry</i> , 2018, 61, 6724-6735.	2.9	22
38	6-Substituted Pyrrolo[3,4-c]pyrazoles: An Improved Class of CDK2 Inhibitors. <i>ChemMedChem</i> , 2007, 2, 841-852.	1.6	21
39	Quality Issues with Public Domain Chemogenomics Data. <i>Molecular Informatics</i> , 2013, 32, 898-905.	1.4	21
40	Structure-based drug design to the discovery of new 2-aminothiazole CDK2 inhibitors. <i>Journal of Molecular Graphics and Modelling</i> , 2006, 24, 341-348.	1.3	20
41	Fluorine NMR functional screening: from purified enzymes to human intact living cells. <i>Journal of Biomolecular NMR</i> , 2020, 74, 613-631.	1.6	20
42	Virtual screening to enrich a compound collection with CDK2 inhibitors using docking, scoring, and composite scoring models. <i>Proteins: Structure, Function and Bioinformatics</i> , 2005, 60, 629-643.	1.5	19
43	Hydrogen Bond Acceptor Propensity of Different Fluorine Atom Types: An Analysis of Experimentally and Computationally Derived Parameters. <i>Chemistry - A European Journal</i> , 2021, 27, 8764-8773.	1.7	18
44	Making sure there's a "give" associated with the "take": producing and using open-source software in big pharma. <i>Journal of Cheminformatics</i> , 2011, 3, .	2.8	16
45	Design, Synthesis, and Preclinical Characterization of Selective Factor D Inhibitors Targeting the Alternative Complement Pathway. <i>Journal of Medicinal Chemistry</i> , 2019, 62, 4656-4668.	2.9	16
46	Application of the rule of shielding in the design of novel fluorinated structural motifs and peptidomimetics. <i>Journal of Fluorine Chemistry</i> , 2013, 152, 129-135.	0.9	15
47	Design of Potent and Selective Covalent Inhibitors of Bruton's Tyrosine Kinase Targeting an Inactive Conformation. <i>ACS Medicinal Chemistry Letters</i> , 2019, 10, 1467-1472.	1.3	15
48	Discovery and Design of First Benzylamine-Based Ligands Binding to an Unlocked Conformation of the Complement Factor D. <i>ACS Medicinal Chemistry Letters</i> , 2018, 9, 490-495.	1.3	9
49	Large-scale Evaluation of CavBase for Analyzing the Polypharmacology of Kinase Inhibitors. <i>Molecular Informatics</i> , 2011, 30, 923-925.	1.4	5
50	Fluorine NMR spectroscopy and computational calculations for assessing intramolecular hydrogen bond involving fluorine and for characterizing the dynamic of a fluorinated molecule. <i>Journal of Fluorine Chemistry</i> , 2017, 202, 34-40.	0.9	4
51	Efficient Screening of Target-specific Selected Compounds in Mixtures by $^{19}\text{F}$ NMR Binding Assay with Predicted $^{19}\text{F}$ NMR Chemical Shifts. <i>ChemMedChem</i> , 2022, , .	1.6	4
52	Comprehensive and High-throughput Exploration of Chemical Space Using Broadband $^{19}\text{F}$ -NMR-Based Screening. <i>Angewandte Chemie</i> , 2020, 132, 14919-14927.	1.6	3
53	Pyrazoles as Efficient Adenine-Mimetic Heterocycles for the Discovery of CDK Inhibitors. <i>Enzyme Inhibitors Series</i> , 2006, , 323-347.	0.1	1
54	Assessment of Docking Poses: Interactions-Based Accuracy Classification (IBAC) versus Crystal Structure Deviations.. <i>ChemInform</i> , 2004, 35, no.	0.1	0

#	ARTICLE	IF	CITATIONS
55	Innentitelbild: Comprehensive and High-Throughput Exploration of Chemical Space Using Broadband <sup>19</sup> F-NMR-Based Screening (Angew. Chem. 35/2020). Angewandte Chemie, 2020, 132, 14806-14806.	16	0