

# Denzil Bernard

## List of Publications by Year in descending order

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43  
papers

3,408  
citations

236925

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302126

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docs citations

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times ranked

4907  
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#	ARTICLE	IF	CITATIONS
1	Follicular Lymphoma-associated BTK Mutations are Inactivating Resulting in Augmented AKT Activation. <i>Clinical Cancer Research</i> , 2021, 27, 2301-2313.	7.0	16
2	Selective inhibition of cullin 3 neddylation through covalent targeting DCN1 protects mice from acetaminophen-induced liver toxicity. <i>Nature Communications</i> , 2021, 12, 2621.	12.8	15
3	Discovery of Potent Small-Molecule Inhibitors of MLL Methyltransferase. <i>ACS Medicinal Chemistry Letters</i> , 2020, 11, 1348-1352.	2.8	9
4	Structure-Based Discovery of M-89 as a Highly Potent Inhibitor of the Menin-Mixed Lineage Leukemia (Menin-MLL) Protein-Protein Interaction. <i>Journal of Medicinal Chemistry</i> , 2019, 62, 6015-6034.	6.4	20
5	Follicular lymphoma-associated mutations in vacuolar ATPase ATP6V1B2 activate autophagic flux and mTOR. <i>Journal of Clinical Investigation</i> , 2019, 129, 1626-1640.	8.2	23
6	High-Affinity Peptidomimetic Inhibitors of the DCN1-UBC12 Protein-Protein Interaction. <i>Journal of Medicinal Chemistry</i> , 2018, 61, 1934-1950.	6.4	46
7	Design of the First-Class, Highly Potent Irreversible Inhibitor Targeting the Menin-MLL Protein-Protein Interaction. <i>Angewandte Chemie - International Edition</i> , 2018, 57, 1601-1605.	13.8	49
8	Design of the First-Class, Highly Potent Irreversible Inhibitor Targeting the Menin-MLL Protein-Protein Interaction. <i>Angewandte Chemie</i> , 2018, 130, 1617-1621.	2.0	1
9	Targeting the MDM2-p53 Protein-Protein Interaction for New Cancer Therapy: Progress and Challenges. <i>Cold Spring Harbor Perspectives in Medicine</i> , 2017, 7, a026245.	6.2	217
10	Discovery of a Highly Potent, Cell-Permeable Macrocyclic Peptidomimetic (MM-589) Targeting the WD Repeat Domain 5 Protein (WDR5)-Mixed Lineage Leukemia (MLL) Protein-Protein Interaction. <i>Journal of Medicinal Chemistry</i> , 2017, 60, 4818-4839.	6.4	72
11	Discovery of 4-((3 <i>R</i> ,4 <i>S</i> ,5 <i>R</i> )-6-chloro-4-(3-chloro-2-fluorophenyl)-1-ethyl-2-oxodispiro[cyclohexane-1,2]undecane-8-carboxylic acid (AA-115/APG-115): A Potent and Orally Active Murine Double Minute 2 (MDM2) Inhibitor in Clinical Development. <i>Journal of Medicinal Chemistry</i> , 2017, 60, 2819-2839.	6.4	143
12	A potent small-molecule inhibitor of the DCN1-UBC12 interaction that selectively blocks cullin 3 neddylation. <i>Nature Communications</i> , 2017, 8, 1150.	12.8	71
13	Functional Analyses of BTK Mutations in Follicular Lymphoma. <i>Blood</i> , 2017, 130, 647-647.	1.4	0
14	Recurrent Mutations in the MTOR Regulator RRAGC in Follicular Lymphoma. <i>Clinical Cancer Research</i> , 2016, 22, 5383-5393.	7.0	36
15	Functional Analyses of V-AtPase Mutations in Follicular Lymphoma. <i>Blood</i> , 2016, 128, 1762-1762.	1.4	0
16	Activating STAT6 mutations in follicular lymphoma. <i>Blood</i> , 2015, 125, 668-679.	1.4	117
17	Structure-based design of conformationally constrained cyclic peptidomimetics to target the MLL1-WDR5 protein-protein interaction as inhibitors of the MLL1 methyltransferase activity. <i>Chinese Chemical Letters</i> , 2015, 26, 455-458.	9.0	4
18	Small-Molecule Inhibitors of the MDM2-p53 Protein-Protein Interaction (MDM2 Inhibitors) in Clinical Trials for Cancer Treatment. <i>Journal of Medicinal Chemistry</i> , 2015, 58, 1038-1052.	6.4	390

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19	Case Study: Discovery of Inhibitors of the MDM2-p53 Protein-Protein Interaction. <i>Methods in Molecular Biology</i> , 2015, 1278, 567-585.	0.9	4
20	Analysis of 54 Follicular Lymphomas By Whole Exome Sequencing Identifies Multiple Novel Recurrently Mutated Pathways. <i>Blood</i> , 2015, 126, 112-112.	1.4	1
21	Design of Chemically Stable, Potent, and Efficacious MDM2 Inhibitors That Exploit the Retro-Mannich Ring-Opening-Cyclization Reaction Mechanism in Spiro-oxindoles. <i>Journal of Medicinal Chemistry</i> , 2014, 57, 10486-10498.	6.4	57
22	SAR405838: An Optimized Inhibitor of MDM2-p53 Interaction That Induces Complete and Durable Tumor Regression. <i>Cancer Research</i> , 2014, 74, 5855-5865.	0.9	261
23	Structure-Based Design of High-Affinity Macrocyclic Peptidomimetics to Block the Menin-Mixed Lineage Leukemia 1 (MLL1) Protein-Protein Interaction. <i>Journal of Medicinal Chemistry</i> , 2013, 56, 1113-1123.	6.4	78
24	High-Affinity, Small-Molecule Peptidomimetic Inhibitors of MLL1/WDR5 Protein-Protein Interaction. <i>Journal of the American Chemical Society</i> , 2013, 135, 669-682.	13.7	157
25	Diastereomeric Spirooxindoles as Highly Potent and Efficacious MDM2 Inhibitors. <i>Journal of the American Chemical Society</i> , 2013, 135, 7223-7234.	13.7	200
26	A Potent Small-Molecule Inhibitor of the MDM2-p53 Interaction (MI-888) Achieved Complete and Durable Tumor Regression in Mice. <i>Journal of Medicinal Chemistry</i> , 2013, 56, 5553-5561.	6.4	229
27	Targeting the MDM2-p53 Protein-Protein Interaction for New Cancer Therapeutics. <i>Topics in Medicinal Chemistry</i> , 2012, , 57-79.	0.8	60
28	AM-8553: A Novel MDM2 Inhibitor with a Promising Outlook for Potential Clinical Development. <i>Journal of Medicinal Chemistry</i> , 2012, 55, 4934-4935.	6.4	32
29	Abstract LB-204: Highly potent and optimized small-molecule inhibitors of MDM2 achieve complete tumor regression in animal models of solid tumors and leukemia.. <i>Cancer Research</i> , 2011, 71, LB-204-LB-204.	0.9	3
30	Case Study: Inhibitors of the MDM2-p53 Protein-Protein Interaction. , 2010, , 273-293.		0
31	Split Renilla Luciferase Protein Fragment-assisted Complementation (SRL-PFAC) to Characterize Hsp90-Cdc37 Complex and Identify Critical Residues in Protein/Protein Interactions. <i>Journal of Biological Chemistry</i> , 2010, 285, 21023-21036.	3.4	33
32	Analysis of the Binding of Mixed Lineage Leukemia 1 (MLL1) and Histone 3 Peptides to WD Repeat Domain 5 (WDR5) for the Design of Inhibitors of the MLL1-WDR5 Interaction. <i>Journal of Medicinal Chemistry</i> , 2010, 53, 5179-5185.	6.4	61
33	Structure-Based Design of Conformationally Constrained, Cell-Permeable STAT3 Inhibitors. <i>ACS Medicinal Chemistry Letters</i> , 2010, 1, 85-89.	2.8	80
34	Temporal activation of p53 by a specific MDM2 inhibitor is selectively toxic to tumors and leads to complete tumor growth inhibition. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 3933-3938.	7.1	641
35	Reactivation of p53 by a specific MDM2 antagonist (MI-43) leads to p21-mediated cell cycle arrest and selective cell death in colon cancer. <i>Molecular Cancer Therapeutics</i> , 2008, 7, 1533-1542.	4.1	87
36	Quantitative Conformationally Sampled Pharmacophore for $\mu$ Opioid Ligands: A Reevaluation of Hydrophobic Moieties Essential for Biological Activity. <i>Journal of Medicinal Chemistry</i> , 2007, 50, 1799-1809.	6.4	36

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37	Position of Coordination of the Lithium Ion Determines the Regioselectivity of Demethylations of 3,4-Dimethoxymorphinans with L-Selectride. <i>Organic Letters</i> , 2005, 7, 2531-2534.	4.6	18
38	Conformationally Sampled Pharmacophore for Peptidic $\mu$ Opioid Ligands. <i>Journal of Medicinal Chemistry</i> , 2005, 48, 7773-7780.	6.4	53
39	Functionalization of the 6,14-Bridge of the Orvinols. 2.1 Preparation of 18- and 19-Hydroxyl-Substituted Thevinols and Their Treatment with Benzyl Bromide. <i>Journal of Organic Chemistry</i> , 2005, 70, 1907-1910.	3.2	9
40	Computer-Aided Drug Design: Structure-Activity Relationships of Delta Opioid Ligands. <i>Drug Design Reviews Online</i> , 2005, 2, 277-291.	0.7	8
41	Rearrangement of 5-Trimethylsilylthebaine on Treatment with L-Selectride: An Efficient Synthesis of (+)-Bractazone. <i>Journal of Organic Chemistry</i> , 2003, 68, 1929-1932.	3.2	10
42	2D Conformationally Sampled Pharmacophore: A Ligand-Based Pharmacophore To Differentiate $\mu$ Opioid Agonists from Antagonists. <i>Journal of the American Chemical Society</i> , 2003, 125, 3101-3107.	13.7	61
43	Basic Principles and Practices of Computer-Aided Drug Design. , 0, , 259-278.		0