## Jing Xing

## List of Publications by Citations

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26 papers 556 citations h-index g-index

29 cxt. papers ext. citations 8.7 avg, IF 3.34 L-index

| #  | Paper  | IF               | Citations |
|----|--|------------------|-----------|
| 26 | In silico ADME/T modelling for rational drug design. <i>Quarterly Reviews of Biophysics</i> , <b>2015</b> , 48, 488-515  | 7                | 137       |
| 25 | Artificial intelligence in drug design. Science China Life Sciences, 2018, 61, 1191-1204   | 8.5              | 72        |
| 24 | Discovery of Highly Potent, Selective, and Orally Efficacious p300/CBP Histone Acetyltransferases Inhibitors. <i>Journal of Medicinal Chemistry</i> , <b>2020</b> , 63, 1337-1360  | 8.3              | 42        |
| 23 | Efficient and Practical One-Pot Conversions of N-Tosyltetrahydroisoquinolines into Isoquinolines and of N-Tosyltetrahydro-Etarbolines into Ecarbolines through Tandem Etlimination and Aromatization. <i>European Journal of Organic Chemistry</i> , <b>2010</b> , 2010, 6987-6992 | 3.2              | 33        |
| 22 | Aldehyde Oxidase Mediated Metabolism in Drug-like Molecules: A Combined Computational and Experimental Study. <i>Journal of Medicinal Chemistry</i> , <b>2017</b> , 60, 2973-2982  | 8.3              | 29        |
| 21 | TarPred: a web application for predicting therapeutic and side effect targets of chemical compounds. <i>Bioinformatics</i> , <b>2015</b> , 31, 2049-51   | 7.2              | 29        |
| 20 | Machine-Learning-Assisted Approach for Discovering Novel Inhibitors Targeting Bromodomain-Containing Protein 4. <i>Journal of Chemical Information and Modeling</i> , <b>2017</b> , 57, 1677-1690  | 6.1              | 27        |
| 19 | Discovery and optimization of selective inhibitors of protein arginine methyltransferase 5 by docking-based virtual screening. <i>Organic and Biomolecular Chemistry</i> , <b>2017</b> , 15, 3648-3661   | 3.9              | 23        |
| 18 | Discovery of novel BET inhibitors by drug repurposing of nitroxoline and its analogues. <i>Organic and Biomolecular Chemistry</i> , <b>2017</b> , 15, 9352-9361  | 3.9              | 23        |
| 17 | Discovery of Novel Disruptor of Silencing Telomeric 1-Like (DOT1L) Inhibitors using a Target-Specific Scoring Function for the (S)-Adenosyl-l-methionine (SAM)-Dependent Methyltransferase Family. <i>Journal of Medicinal Chemistry</i> , <b>2017</b> , 60, 2026-2036             | 8.3              | 20        |
| 16 | Discovery of Novel Inhibitors of Indoleamine 2,3-Dioxygenase 1 Through Structure-Based Virtual Screening. <i>Frontiers in Pharmacology</i> , <b>2018</b> , 9, 277  | 5.6              | 20        |
| 15 | Modulating TRADD to restore cellular homeostasis and inhibit apoptosis. <i>Nature</i> , <b>2020</b> , 587, 133-138   | 50.4             | 19        |
| 14 | Discovery of Novel Inhibitors Targeting the Menin-Mixed Lineage Leukemia Interface Using Pharmacophore- and Docking-Based Virtual Screening. <i>Journal of Chemical Information and Modeling</i> , <b>2016</b> , 56, 1847-55   | 6.1              | 19        |
| 13 | Analysis of Infected Host Gene Expression Reveals Repurposed Drug Candidates and Time-Dependent Host Response Dynamics for COVID-19 <b>2020</b> ,  |                  | 10        |
| 12 | An Efficient and General Method for the Stereodivergent Syntheses of Tadalafil-Like Tetracyclic Compounds. <i>European Journal of Organic Chemistry</i> , <b>2010</b> , 2010, 1711-1716  | 3.2              | 9         |
| 11 | Rational design of 5-((1H-imidazol-1-yl)methyl)quinolin-8-ol derivatives as novel bromodomain-containing protein 4 inhibitors. <i>European Journal of Medicinal Chemistry</i> , <b>2019</b> , 163, 281-  | 294 <sup>8</sup> | 9         |
| 10 | Deep Neural Network Classifier for Virtual Screening Inhibitors of (S)-Adenosyl-L-Methionine (SAM)-Dependent Methyltransferase Family. <i>Frontiers in Chemistry</i> , <b>2019</b> , 7, 324  | 5                | 5         |

## LIST OF PUBLICATIONS

| 9 | Mild and Efficient Syntheses of 1-Aryl-3,4-dihydroisoquinolines and 1-Aryl-3,4-dihydro-Larbolines via Regiospecific Eliminations of the Corresponding N-Tosyl-1,2,3,4-tetrahydroisoquinolines and N-Tosyl-1,2,3,4-tetrahydro-Earbolines. <i>Synthetic Communications</i> , <b>2012</b> , 42, 2806-2817 | 1.7  | 5 |  |
|---|--|------|---|--|
| 8 | OCTAD: an open workspace for virtually screening therapeutics targeting precise cancer patient groups using gene expression features. <i>Nature Protocols</i> , <b>2021</b> , 16, 728-753  | 18.8 | 5 |  |
| 7 | Sex differences in viral entry protein expression, host responses to SARS-CoV-2, and in vitro responses to sex steroid hormone treatment in COVID-19 <b>2020</b> ,   |      | 4 |  |
| 6 | Development and evaluation of a novel series of Nitroxoline-derived BET inhibitors with antitumor activity in renal cell carcinoma. <i>Oncogenesis</i> , <b>2018</b> , 7, 83   | 6.6  | 4 |  |
| 5 | MoCL: Data-driven Molecular Fingerprint via Knowledge-aware Contrastive Learning from Molecular Graph. <b>2021</b> , 2021, 3585-3594   |      | 4 |  |
| 4 | Gene expression signatures identify paediatric patients with multiple organ dysfunction who require advanced life support in the intensive care unit. <i>EBioMedicine</i> , <b>2020</b> , 62, 103122   | 8.8  | 3 |  |
| 3 | OCTAD: an open workplace for virtually screening therapeutics targeting precise cancer patient groups using gene expression features   |      | 2 |  |
| 2 | Published anti-SARS-CoV-2 in vitro hits share common mechanisms of action that synergize with antivirals. <i>Briefings in Bioinformatics</i> , <b>2021</b> , 22,   | 13.4 | 2 |  |
| 1 | Published Anti-SARS-CoV-2 In Vitro Hits Share Common Mechanisms of Action that Synergize with Antivirals <b>2021</b> ,   |      | 1 |  |