Maha Habash

List of Publications by Year in Descending Order

15 papers	229 citations	10 h-index	15 g-index
16 ext. papers	256 ext. citations	3.2 avg, IF	3 L-index

#	Paper	IF	Citations
15	Research advances in kinase enzymes and inhibitors for cardiovascular disease treatment. *Future Science OA*, **2017**, 3, FSO204	2.7	9
14	Docking-based comparative intermolecular contacts analysis and in silico screening reveal new potent acetylcholinesterase inhibitors. *Medicinal Chemistry Research*, **2017**, 26, 2768-2784	2.2	12
13	In Vitro and In Vivo Evaluation of Casein as a Drug Carrier for Enzymatically Triggered Dissolution Enhancement from Solid Dispersions. *AAPS PharmSciTech*, **2017**, 18, 1750-1759	3.9	3
12	Identification of novel inhibitors for Pim-1 kinase using pharmacophore modeling based on a novel method for selecting pharmacophore generation subsets. *Journal of Computer-Aided Molecular Design*, **2016**, 30, 39-68	4.2	10
11	Phytochemical Investigation and Antimicrobial Activities of Jordanian Psidium guajava Raw Fruit Peel Extract using Soxhlet and Microwave Extraction Methods. *Research Journal of Medicinal Plant*, **2016**, 10, 443-449	0.3	1
10	Discovery of potent adenosine A2a antagonists as potential anti-Parkinson disease agents. Non-linear QSAR analyses integrated with pharmacophore modeling. *Chemico-Biological Interactions*, **2016**, 254, 93-101	5	14
9	Ligand-based modeling followed by in vitro bioassay yielded new potent glucokinase activators. *Journal of Molecular Graphics and Modelling*, **2015**, 56, 91-102	2.8	15
8	Synthesis and evaluation of novel diphenylthiazole derivatives as potential anti-inflammatory agents. *Medicinal Chemistry Research*, **2015**, 24, 3681-3695	2.2	5
7	Combining docking-based comparative intermolecular contacts analysis and k-nearest neighbor correlation for the discovery of new check point kinase 1 inhibitors. *Journal of Computer-Aided Molecular Design*, **2015**, 29, 561-81	4.2	17
6	Preparation and in vitro characterization of glibenclamide-loaded alginate hexyl-amide beads: a novel drug delivery system to improve the dissolution rate. *Pharmaceutical Development and Technology*, **2014**, 19, 881-90	3.4	6
5	The use of docking-based comparative intermolecular contacts analysis to identify optimal docking conditions within glucokinase and to discover of new GK activators. *Journal of Computer-Aided Molecular Design*, **2014**, 28, 509-47	4.2	30
4	Elaborate ligand-based modeling reveals new human neutrophil elastase inhibitors. *Medicinal Chemistry Research*, **2014**, 23, 3876-3896	2.2	11
3	Elaborate ligand-based modeling coupled with QSAR analysis and in silico screening reveal new potent acetylcholinesterase inhibitors. *Journal of Computer-Aided Molecular Design*, **2013**, 27, 1075-92	4.2	28
2	Docking-based comparative intermolecular contacts analysis as new 3-D QSAR concept for validating docking studies and in silico screening: NMT and GP inhibitors as case studies. *Journal of Chemical Information and Modeling*, **2011**, 51, 647-69	6.1	57
1	Ligand-based modelling followed by synthetic exploration unveil novel glycogen phosphorylase inhibitory leads. *Bioorganic and Medicinal Chemistry*, **2011**, 19, 4746-71	3.4	11