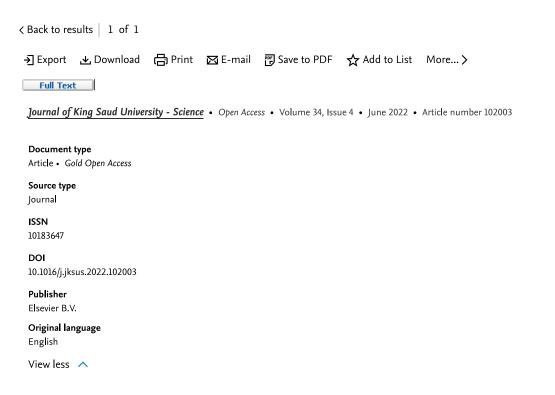


 $Q \equiv$



Design, synthesis, in vitro antiproliferative effect and in situ molecular docking studies of a series of new benzoquinoline derivatives

```
<u>Umar, Abdulrashid</u> a ⋈ ; <u>Faidallah, Hassan M.</u> b ⋈ ; <u>Ahmed, Qamar Uddin</u> c ⋈ ; <u>Alamry, Khalid.A.</u> a ⋈ ; <u>Mukhtar, Sayeed</u> d ⋈ ; <u>Alsharif, Meshari A.</u> e ⋈ ; <u>Azmi, Syed Najmul Hejaz</u> f ⋈ ; <u>Parveen, Humaira</u> d ⋈ ; <u>Zakaria, Zainul Amiruddin</u> g ⋈ ; <u>Hussien, Mostafa A.</u> a ⋈ Gave all to author list
```

- ^a Department of Chemistry, Faculty of Science, King Abdulaziz University, P.O. Box 80203, Jeddah, 21589, Saudi Arabia
- ^b Department of Chemistry, Faculty of Science, Alexandria University, Alexandria, 21526, Egypt
- ^c Drug Discovery And Synthetic Chemistry Research Group, Department of Pharmaceutical Chemistry, Kulliyyah of Pharmacy, International Islamic University Malaysia, Pahang DM, Kuantan, 25200, Malaysia
- d Department of Chemistry, Faculty of Science, University of Tabuk, Tabuk, 71491, Saudi Arabia View additional affiliations ✓

Full text options ∨

Abstract

Author keywords

Reaxys Chemistry database information

SciVal Topics

Metrics

Cited by 0 documents

Inform me when this document is cited in Scopus:

Set citation alert >

Related documents

Molecular docking suggests repurposing of brincidofovir as a potential drug targeting SARS-CoV-2 ACE2 receptor and main protease

Hussien, M.A., Abdelaziz, A.E.M. (2020) Network Modeling Analysis in Health Informatics and Bioinformatics

Synthesis of new 3-(hydroxymethyl)-2-phenyl-2,3 dihydroquinolinone and in-silico evaluation of COVID-19 main protease inhibitor

Nepolraj, A., Shupeniuk, V.I., Sathiyaseelan, M. (2021) Vietnam Journal of Chemistry

A simple and efficient synthesis of benzofuroquinolines via the decarboxylative cross-coupling

Rathod, P.K., Jonnalagadda, S., Panaganti, L. *(2021) Tetrahedron Letters*

View all related documents based on references

Find more related documents in Scopus based on:

Authors > Keywords >

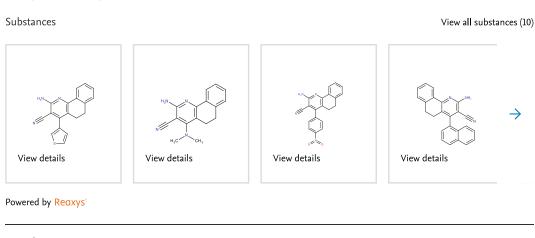
Abstract

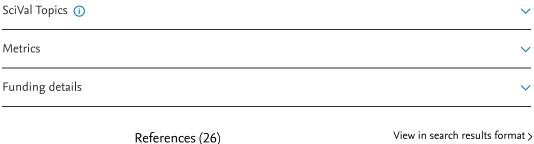
Quinoline derivatives have been reported to possess multi-therapeutic potential owing to the manifestations of different pharmacological effects. The current research work describes about the design and synthesis of a series of novel benzoquinoline analogues with an objective to evaluate their antiproliferative structure-activity relationship against colon, breast and hepatocellular cancers. Upon synthesis, all derivatives' chemical structures were elucidated through FTIR, ¹HNMR and ¹³CNMR spectroscopic analysis. All derivatives were investigated for their in vitro anti-proliferative property against three different cancer cell lines (viz., colon carcinoma HT29, Caucasian breast adenocarcinoma MCF7, hepatocellular carcinoma HepG2) and a normal non-transformed human foreskin fibroblast Hs27 cell line. All derivatives demonstrated varied degrees of strong anticancer effect against all of the cell lines with the 2-Amino-4-(4-nitrophenyl)-5,6-dihydrobenzo[h]quinoline-3-carbonitrile (CNMP, 2) exhibited the most potent antiproliferative effect viz. LC₅₀ 21.23 μM for breast, 8.24 μM for colon, and 26.15 μM for the hepatocellular, respectively. Molecular docking studies against all the the target crystal structures of cancer proteins (1HK7, 3EQM, 3IG7 and 4FM9) revealed significant binding affinities via hydrophobic and H-bonding interactions with all the compounds in conformity with the wet lab results. CNMP showed the highest binding energy of -7.55 in the HT29, -6.9 (both in MCF7 HepG2) kcal/mol. Based on the results obtained from wet lab and dry lab experiments, it can be proposed that CNMP might prove to be a potential lead structure for the design and synthesis of more potent anticancer candidates. © 2022 The Author(s)

Author keywords

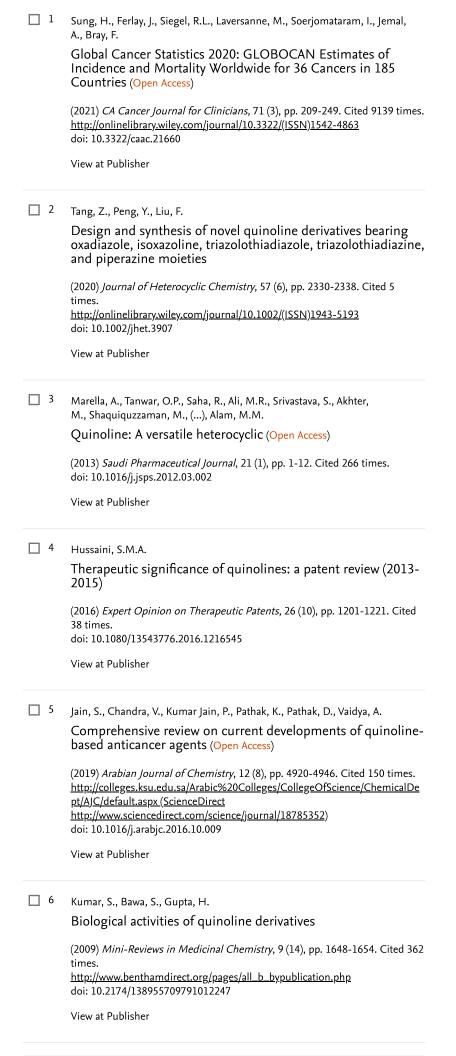
Benzoquinoline derivatives; Computational studies; Cytotoxicity; Molecular docking; Quinolines

Reaxys Chemistry database information (i)

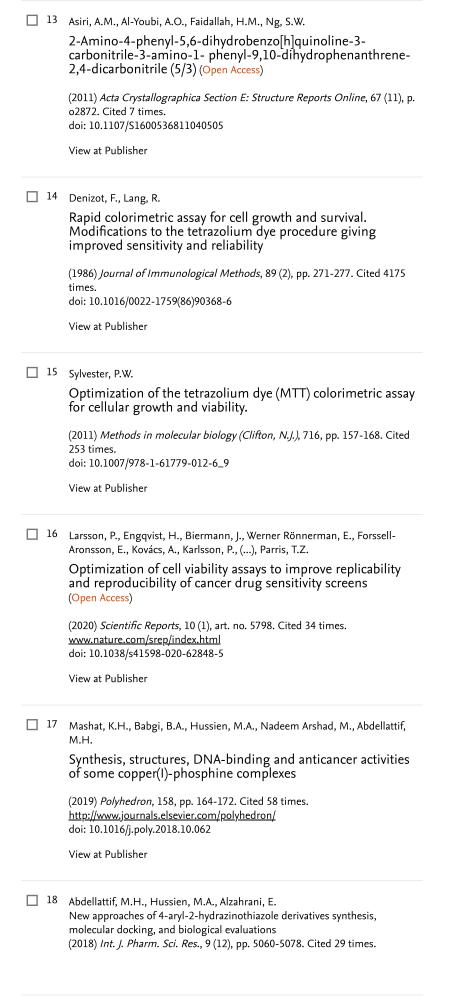


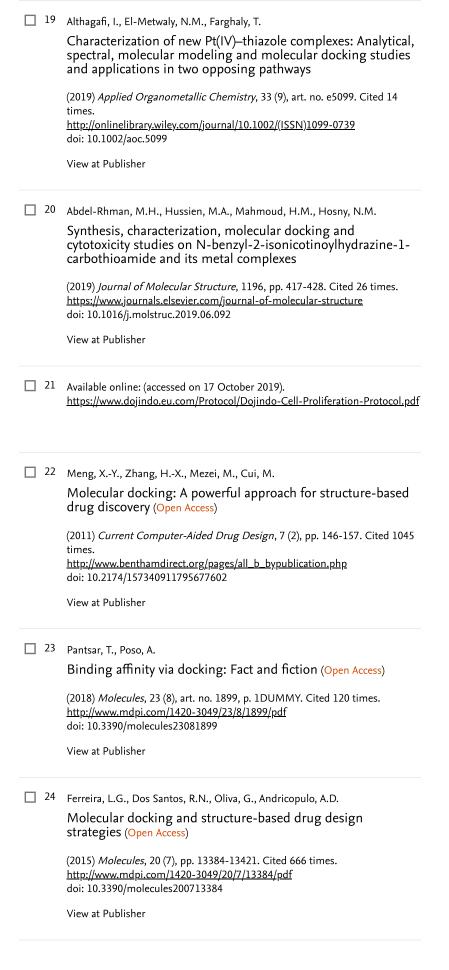






J 7	Sidoryk, K., Switalska, M., Jaromin, A., Cmoch, P., Bujak, I., Kaczmarska, M., Wietrzyk, J., (), Kaczmarek, L. The synthesis of indolo[2,3-b]quinoline derivatives with a guanidine group: Highly selective cytotoxic agents (2015) European Journal of Medicinal Chemistry, 105, pp. 208-219. Cited 28 times. http://www.journals.elsevier.com/european-journal-of-medicinal-chemistry/doi:10.1016/j.ejmech.2015.10.022
	View at Publisher
8	Nqoro, X., Tobeka, N., Aderibigbe, B.A. Quinoline-based hybrid compounds with antimalarial activity (Open Access)
	(2017) Molecules, 22 (12), art. no. 2268. Cited 75 times. http://www.mdpi.com/1420-3049/22/12/2268/pdf doi: 10.3390/molecules22122268 View at Publisher
9	Surabhi, S., Singh, B.K. Computer aided drug design: An overview (2018) J.D.D.T., 8 (5), pp. 504-509. Cited 29 times.
] 10	Vilar, S., Cozza, G., Moro, S. Medicinal chemistry and the Molecular Operating Environment (MOE): Application of QSAR and molecular docking to drug discovery (2008) Current Topics in Medicinal Chemistry, 8 (18), pp. 1555-1572. Cited 490 times. http://docserver.ingentaconnect.com/deliver/connect/ben/15680266/v8n18/s2 _pdf? expires=1235130221&id=48925662&titleid=3902&accname=Elsevier+Bibliogr aphic+Databases&checksum=B4B6CEF4ACF71CFBE5C504C619D08756 doi: 10.2174/156802608786786624 View at Publisher
] 11	Fischer, N., Seo, EJ., Abdelfatah, S., Fleischer, E., Klinger, A., Efferth, T. A novel ligand of the translationally controlled tumor protein (TCTP) identified by virtual drug screening for cancer differentiation therapy (Open Access) (2021) Investigational New Drugs, 39 (4), pp. 914-927. Cited 2 times. www.wkap.nl/journalhome.htm/0167-6997 doi: 10.1007/s10637-020-01042-w View at Publisher
12	Qandalee, M., Alikarami, M., Mighani, H., Asghari, S., Beikjani, S. Synthesis of quinoline derivatives from the reaction of aminobenzophenones and acetylenic esters in the presence of SnO ₂ nanoparticles (2013) <i>Int. Nano Lett.</i> , 3, p. 49. Cited 3 times.





□ 25	Singh, N., Villoutreix, B.O., Ecker, G.F. Rigorous sampling of docking poses unveils binding hypothesis for the halogenated ligands of L-type Amino acid Transporter 1 (LAT1) (Open Access) (2019) Scientific Reports, 9 (1), art. no. 15061. Cited 13 times. www.nature.com/srep/index.html doi: 10.1038/s41598-019-51455-8 View at Publisher
□ 26	Boittier, E.D., Tang, Y.Y., Buckley, M.E., Schuurs, Z.P., Richard, D.J., Gandhi, N.S. Assessing molecular docking tools to guide targeted drug discovery of cd38 inhibitors (Open Access) (2020) International Journal of Molecular Sciences, 21 (15), art. no. 5183, pp. 1-19. Cited 13 times. https://www.mdpi.com/1422-0067/21/15/5183/pdf doi: 10.3390/ijms21155183 View at Publisher
of Pharm Malaysia	ed, Q.U.; Drug Discovery And Synthetic Chemistry Research Group, Department naceutical Chemistry, Kulliyyah of Pharmacy, International Islamic University, Pahang DM, Kuantan, Malaysia; email:quahmed@iium.edu.my right 2022 Elsevier B.V., All rights reserved.

About Scopus

What is Scopus

Content coverage

Scopus blog

Scopus API

Privacy matters

Language

日本語に切り替える

切换到简体中文

切換到繁體中文

Русский язык

Customer Service

Help

Tutorials

Contact us

ELSEVIER

Terms and conditions *¬* Privacy policy *¬*

Copyright © Elsevier B.V 对. All rights reserved. Scopus® is a registered trademark of Elsevier B.V.

We use cookies to help provide and enhance our service and tailor content. By continuing, you agree to the use of cookies.

