

INNOVATIVE 7-O-SUBSTITUTED CHRYSIN DERIVATIVES AS VEGFR-2 INHIBITORS: A NATURAL PRODUCT-BASED ANTICANCER THERAPY

Kalyani R. Thombre¹, Krishna R. Gupta¹, Milind J. Umekar²

¹Department of Pharmaceutical Chemistry, Smt. Kishoritai Bhoyar College of Pharmacy, Kamptee, Nagpur, India

²Department of Pharmaceutics, Smt. Kishoritai Bhoyar College of Pharmacy, Kamptee, Nagpur, India

Corresponding author: kr1903@gmail.com

Doi: <https://doi.org/10.5281/zenodo.17909205>

Received: 10 September 2025

Accepted: 14 September 2025

Abstract:

Introduction

VEGFR-2, critical target in cancer therapy, facilitating tumor angiogenesis, yet existing inhibitors face toxicity and resistance issues. Chrysin, flavonoid with anticancer properties, has VEGFR-2 characteristics but suffers from poor pharmacokinetics. A series of 7-O-substituted chrysin derivatives was designed to enhance binding and drug-like properties by incorporating key hydrogen-bonding groups and hydrophobic elements, informed by VEGFR-2 structural analysis.

To design and assess novel chrysin derivatives through computational predictions and molecular docking; synthesize and characterize selected derivatives; and evaluate their antioxidant and anticancer activities in vitro, to identify effective candidates that exhibit favorable pharmacokinetics and safety.

Methods

Chrysin derivatives featuring alkylamino and ester substituents were designed using molecular docking against VEGFR-2. ADMET profiling was conducted to anticipate pharmacokinetics and toxicity. Selected derivatives were synthesized via alkylation and esterification, and characterized using UV, IR, NMR, and mass spectrometry. Antioxidant activity evaluated using DPPH assay, while anticancer efficacy against MCF-7 cells assessed through cell viability assays, comparing IC₅₀ values with those of ascorbic acid and sorafenib.

Results

Docking studies indicated strong binding affinities, particularly for ester derivatives. ADMET predictions suggested favorable drug-like characteristics. C7 demonstrated remarkable antioxidant activity, exceeding both chrysin and ascorbic acid. In anticancer tests, C7 and C8 displayed significant cytotoxicity (IC₅₀ = 1.0 and 1.5 μM), outperforming chrysin and nearing sorafenib efficacy.

Conclusion

The improved bioactivity and predicted safety of C7 and C8 highlight efficacy of rational structural modifications in optimizing natural products. Their dual antioxidant and anticancer properties underscore their potential as lead compounds for VEGFR-2-targeted breast cancer treatments. Chrysin derivatives, particularly C7 and C8, exhibit promising VEGFR-2 inhibitory activity and therapeutic potential, supporting their further investigation as multifunctional anticancer agents. Building on these findings, future efforts will focus on developing suitable formulations to improve their solubility, stability, and bioavailability

Keywords: Chrysin derivatives, Antioxidant, Anticancer, VEGFR-2 inhibition

References

1. Kalyani, T., Krishna, R., Sudhanshu, S., Pavan, S., Aparna, R., Amaanullah, S., & Mimind, U. (2024). The natural flavone quercetin: An extensive analysis of its pharmacological mechanisms and medicinal prospects. *Indian Journal of Pharmacology and Pharmacology*, 2024, 23734. <https://doi.org/10.18231/j.ijpp.2024.032>
2. De Souza Farias, S. A., Da Costa, K. S., & Martins, J. B. L. (2021). Analysis of conformational, structural, magnetic, and electronic properties related to antioxidant activity: Revisiting flavan, anthocyanidin, flavanone, flavonol, isoflavone, flavone, and flavan-3-ol. *ACS Omega*, 6, 8908–8918. <https://doi.org/10.1021/acsomega.0c06156>
3. Falbo, F., & Aiello, F. (2023). Chrysin: A polyedric flavone as a tool to explore new phytotherapeutic

- applications and drug design. *Archiv der Pharmazie*, 356, 2200347. <https://doi.org/10.1002/ardp.202200347>
4. Xue, X. Y., He, J. L., Song, M. W., Yu, Z., Lin, S. L., Zhang, C. P., Hao, B. D., Ma, Z. H., Zhang, W. H., Zou, Y. Y., Jing, J. Y., & Shi, D. H. (2025). Design, synthesis, bioactivity, X-ray crystallography, and molecular docking studies of chrysin-1,3,5-triazine derivatives as anticancer agents. *Bioorganic Chemistry*, 161, 108486. <https://doi.org/10.1016/j.bioorg.2025.108486>
 5. Massi, A., Bortolini, O., Ragno, D., Bernardi, T., Sacchetti, G., Tacchini, M., & De Risi, C. (2017). Research progress in the modification of quercetin leading to anticancer agents. *Molecules*, 22, 1270. <https://doi.org/10.3390/molecules22081270>
 6. Abdel-Mohsen, H. T., Ibrahim, M. A., Nageeb, A. M., & El Kerdawy, A. M. (2024). Receptor-based pharmacophore modeling, molecular docking, synthesis and biological evaluation of novel VEGFR-2, FGFR-1, and BRAF multi-kinase inhibitors. *BMC Chemistry*, 18, 42. <https://doi.org/10.1186/s13065-024-01135-0>
 7. Zhang, L., Shan, Y., Ji, X., Zhu, M., Li, C., Sun, Y., Si, R., Pan, X., Wang, J., Ma, W., Dai, B., Wang, B., & Zhang, J. (2017). Discovery and evaluation of triple inhibitors of VEGFR-2, TIE-2 and EphB4 as anti-angiogenic and anti-cancer agents. *Oncotarget*, 8, 104745–104760. <https://doi.org/10.18632/oncotarget.20065>
 8. Abdallah, A. E., Mabrouk, R. R., Al Ward, M. M. S., Eissa, S. I., Elkaeed, E. B., Mehany, A. B. M., Abo-Saif, M. A., El-Feky, O. A., Alesawy, M. S., & El-Zahabi, M. A. (2022). Synthesis, biological evaluation, and molecular docking of new series of antitumor and apoptosis inducers designed as VEGFR-2 inhibitors. *Journal of Enzyme Inhibition and Medicinal Chemistry*, 37, 573–591. <https://doi.org/10.1080/14756366.2021.2017911>
 9. Moghadam, E. R., Ang, H. L., Asnaf, S. E., Zabolian, A., Saleki, H., Yavari, M., Esmaeili, H., Zarrabi, A., Ashrafizadeh, M., & Kumar, A. P. (2020). Broad-spectrum preclinical antitumor activity of chrysin: Current trends and future perspectives. *Biomolecules*, 10, 1374. <https://doi.org/10.3390/biom10101374>
 10. Sokal, A., Mruczek, P., Niedoba, M., Dewalska, A., Stocercz, K., & Kadela-Tomanek, M. (2025). Anticancer activity of ether derivatives of chrysin. *Molecules*, 30, 960. <https://doi.org/10.3390/molecules30040960>
 11. Salari, N., Faraji, F., Jafarpour, S., Faraji, F., Rasoulpoor, S., Dokaneheifard, S., & Mohammadi, M. (2022). Anti-cancer activity of chrysin in cancer therapy: A systematic review. *Indian Journal of Surgical Oncology*, 13, 681–690. <https://doi.org/10.1007/s13193-022-01550-6>
 12. 31. During, A., & Larondelle, Y. (2013). The O-methylation of chrysin markedly improves its intestinal anti-inflammatory properties: Structure–activity relationships of flavones. *Biochemical Pharmacology*, 86(12), 1739–1746. <https://doi.org/10.1016/j.bcp.2013.10.003>
 13. Gao, S., Siddiqui, N., Etim, I., Du, T., Zhang, Y., & Liang, D. (2021). Developing nutritional component chrysin as a therapeutic agent: Bioavailability and pharmacokinetics consideration, and ADME mechanisms. *Biomedicine & Pharmacotherapy*, 142, 112080. <https://doi.org/10.1016/j.biopha.2021.112080>
 14. Garg, A., & Chaturvedi, S. (2022). A comprehensive review on chrysin: Emphasis on molecular targets, pharmacological actions and bio-pharmaceutical aspects. *Current Drug Targets*, 23(4), 420–436. <https://doi.org/10.2174/1389450122666210824141044>
 15. Şandor, A., Ionuţ, I., Marc, G., Oniga, I., Eniu, D., & Oniga, O. (2023). Structure–activity relationship studies based on quinazoline derivatives as EGFR kinase inhibitors (2017–present). *Pharmaceuticals*, 16(4), 534. <https://doi.org/10.3390/ph16040534>
 16. Yuan, Y. H., Mao, N. D., Duan, J. L., Zhang, H., Garrido, C., Lirussi, F., Gao, Y., Xie, T., & Ye, X. Y. (2023). Recent progress in discovery of novel AAK1 inhibitors: From pain therapy to potential anti-viral agents. *Journal of Enzyme Inhibition and Medicinal Chemistry*, 38(1), 2279906. <https://doi.org/10.1080/14756366.2023.2279906>
 17. Eldehna, W. M., Abou-Seri, S. M., El Kerdawy, A. M., Ayyad, R. R., Hamdy, A. M., Ghabbour, H. A., Ali, M. M., & Abou El Ella, D. A. (2016). Increasing the binding affinity of VEGFR-2 inhibitors by extending their hydrophobic interaction with the active site: Design, synthesis and biological evaluation of 1-substituted-4-(4-methoxybenzyl)phthalazine derivatives. *European Journal of Medicinal Chemistry*, 113, 50–62. <https://doi.org/10.1016/j.ejmech.2016.02.029>
 18. Rizvi, S. U. F., Siddiqui, H. L., Nisar, M., Khan, N., & Khan, I. (2012). Discovery and molecular docking of quinolyl-thienyl chalcones as anti-angiogenic agents targeting VEGFR-2 tyrosine kinase. *Bioorganic & Medicinal Chemistry Letters*, 22(3), 942–944. <https://doi.org/10.1016/j.bmcl.2011.12.017>

19. Li, X., Cai, Y., Yang, F., & Meng, Q. (2017). Synthesis and molecular docking studies of chrysin derivatives as antibacterial agents. *Medicinal Chemistry Research*, 26(10), 2225–2234. <https://doi.org/10.1007/s00044-017-1952-4>
20. Iqbal, D., Alsaweed, M., Jamal, Q. M. S., Asad, M. R., Rizvi, S. M. D., Rizvi, M. R., Albadrani, H. M., Hamed, M., Jahan, S., & Alyenbaawi, H. (2023). Pharmacophore-based screening, molecular docking, and dynamic simulation of fungal metabolites as inhibitors of multi-targets in neurodegenerative disorders. *Biomolecules*, 13(11), 1613. <https://doi.org/10.3390/biom13111613>
21. The role of pyrogallol as an anti-cancer agent reduces cell proliferation in lung cancer cells via AKT/PI3K signaling pathways—An in vitro and in silico approaches. (2025). *Texila International Journal of Public Health*, 25(1), Article 020. <https://doi.org/10.21522/TIJPH.2013.SE.25.01.Art020>