



## Design and Synthesis of Fused 1,2,3-Triazolo-Pyrano-Quinazoline Using Copper(I) Catalysis: *In silico* Molecular Docking, *in vitro* Tyrosine Inhibition and ADMET Studies

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In this work, the synthesis of novel 1,2,3-triazolo-pyrano-quinazoline conjugates (**6a-n**) using well-known copper-catalyzed CuAAC and C-H arylation cascade reactions is carried out. The anticancer activity of these conjugates was evaluated against two human cancer cell lines, MCF-7 and HepG-2. The results showed that conjugate **6e** exhibited more potent activity compared to the standard drug erlotinib, while compounds **6b**, **6d** and **6f** displayed slightly lower activity compared to the standard drug. These four potent compounds (**6b**, **6d**, **6e** and **6f**) were assessed in a cell survival assay employing the normal breast cell line MCF-10A. None of them showed significant cytotoxicity, with IC<sub>50</sub> values larger than 98.20 μM. *In vitro* tyrosine kinase EGFR inhibitory of four potent compounds were evaluated and results indicate that compound **6e** exhibited higher EGFR inhibitory activity compared to the standard drug erlotinib. On the other hand, compounds **6b** and **6f** displayed lower activity compared to both the standard drug and compound **6e**. Furthermore, the molecular docking studies were also performed on four potent conjugates and the results showed that these conjugates had more EGFR-binding interactions as compared to the standard drug erlotinib. Moreover, the *in silico* pharmacokinetic outline of the potent conjugates **6b**, **6d**, **6e** and **6f** was estimated by using SWISS/ADME and pkCSM and all the four conjugates followed Lipinski rule of five, Ghose, Veber, Egan and Muegge rules without any deviation.

**Keywords:** 1,2,3-Triazolo-pyrano-quinazoline conjugates, Anticancer activity, Molecular docking studies, Pharmacokinetic profile.