Optimizing of core structure of 4-hydroxy-2-quinolone as potential anticancer agent.

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Abstract

Cancer is a complex of deadliest heterogeneous diseases associated with a vast incidence and death rates. Genetic mutations and aberrant signaling pathways are considered as one of numerous factors predisposing cancer development. Derivatives of 4-hydroxy-1,6-naphthyridin-2-one-3-carboxamide were designed and synthesized in order to optimize the scaffold of 4-hydroxy-2-quinolone-3-carboxamide. The identity of the synthesized analogues was characterized using FT-IR, 1 H NMR, 13 C NMR, and elemental analysis. Biological investigations against human colon carcinoma cell (HCT-116) and breast cancer cell (MCF-7) lines showed that analogues exhibit potent and selective inhibitory activity. Analogues tailored with benzyl moiety (47) (IC50 HCT-116 = 43.30 μ M, IC50 MCF-7 = 88.98 μ M) exerted promising activity implying that flexibility of carboxamide side chain assists in orientating the core structure deeply

in the binding domain. Derivatives with substituted benzyl motif exemplified by o-F (49) (IC₅₀ HCT-116 = 62.32 μ M, IC₅₀ MCF-7 = 89.27 μ M), m-F (50) (IC₅₀ HCT-116 = 52.34 μ M, IC₅₀ MCF-7 = 37.06 μ M), p-F (51) (IC₅₀ HCT-116 = 50.48 μ M, IC₅₀ MCF-7 = 20.0 μ M), m-CF₃ (53) (IC₅₀ HCT-116 = 88.50 μ M, IC₅₀ MCF-7 = 33.0 μ M), and p-CF₃ (53) (IC₅₀ HCT-116 = 81.04 μ M, IC₅₀ MCF-7 = 54.0 μ M) confirmed that flexibility of the side chain along with introducing -F and/or -CF₃ induce the activity. Results anticipate that electronegativity and/or hydrophobicity improves ligand/receptor binding interaction. The induced-fit docking (IFD) studies against potential receptors such as phosphoinositide-3-kinase (PI3K α), histone deacetylase (HDAC-II), and estrogen receptor (ER) revealed distinguishable binding scores against HDAC-II encouraging us to dip deeply into the molecular level to better retrieve potential receptor (s).

Keywords: Cancer, Docking, 4-Hydroxy-1,6-naphthyridin-2-one-3-carboxamide, HCT-116, MCF-7.