

**Design, Synthesis, and Biological Evaluation of *N*-Substituted-4-Hydroxy-6-Methyl-2-Quinolone-3-Carboxamides as PI3K $\alpha$**

**Inhibitors**

**By**

**Shaima' Emad Hasan**

**Supervisor**

**Dr. Dima A. Sabbah**

**Co-Supervisor**

**Dr. Reema Abu khalaf**

**Abstract**

Phosphatidylinositol 3-kinase alpha (PI3K $\alpha$ ) has been highlighted as a hot target for anticancer drug design. A series of *N*-substituted-4-hydroxy-6-methyl-2-quinolone-3-carboxamides was designed and synthesized as possible PI3K $\alpha$  inhibitors recruiting structure-based drug design and molecular docking. The identity of the core nucleus and synthesized derivatives was characterized using (<sup>1</sup>H and <sup>13</sup>C) NMR analysis technique.

Biological studies in human colon carcinoma (HCT116) cell line showed that the analogues **37**, **39-45** exhibited distinct antiproliferative activity against PI3K $\alpha$ . Compound tailored with hydrazide motif (**42**) exhibited higher inhibitory activity (IC<sub>50</sub> = 0.323 mM) suggesting that H-bond interaction improves the binding affinity. Compound tailored with *o*-COOH and *p*-CH<sub>3</sub> (**45**) showed high activity (IC<sub>50</sub> = 0.567

mM) implying that H-bond and hydrophobic interaction on the *o*- and *p*-positions elicit the activity.

Compounds **39** and **41** displayed comparable antiproliferative activity ( $IC_{50} = 0.849$  and  $0.811$  mM) inferring that similar binding conformation is generated in the kinase domain. The loss of activity of compounds **43** and **44** indicates that *p*-H-bond acceptor and/or donor doesn't mediate ligand/PI3K $\alpha$  complex formation. Conversely, the activity of **40** ( $IC_{50} = 1.54$  mM) suggests that *p*-OCH<sub>3</sub> accommodates a tight hydrophobic cleft and confirms the absence of H-bond acceptor on *p*-position.

The Induced-fit docking studies against PI3K $\alpha$  demonstrated that the derivatives accommodate PI3K $\alpha$  kinase catalytic domain and form H-bonding with the key binding residues. Our results suggest that further optimization of this series would be beneficial for colon cancer treatment.