Synthesis, molecular docking, and preliminary cytotoxicity study of some novel 2-(naphthalen-1-yl)-methylimidazo[2,1-*b*][1,3,4]thiadiazoles

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1. Introduction

Cancer is a large group of diseases and can start in any tissue or organ. Cancer worldwide is the second leading cause of death, after heart disease, and is estimated to be the cause of 99 lakh deaths by 2020 [1]. In children under the age of 19, at least three lakh new cancer cases have been diagnosed [1]. In therapeutic chemistry, heteroatoms such as N and S have gained importance because of their many pharmacological functions [2-22]. 2-Amino-1,3,4-thiadiazoles (I) have been tested as carcinostatic agents against several tissues of implanted animals [3]. The imidazo ring [2,1-b] was fused with 2-amino-1,3,4-thiadiazole, which resulted in imidazo[2,1-b][1,3,4]thiadiazole with antibacterial [4-6], antifungal [7], anti-inflammatory [8,9], anticancer [10], analgesic [11], and antitubercular properties [12,13]. We have reported imidazo[2,1-b][1,3,4]thiadiazoles (II) as cytotoxic agent [14-19]. Recent studies have reported experiments on imidazo[2,1-

ABSTRACT

A series of 2-(naphthalen-1-yl)-methyl-6-arylimidazo[2,1-*b*][1,3,4]thiadiazole derivatives was prepared and studied for cytotoxicity against murine leukemia L1210, human cervix carcinoma HeLa, and human T-lymphocyte CEM cell lines. The preliminary study showed that compounds **5g**, **6g**, **7a-c**, **7e**, and **8e** were more potent among the tested compounds. The pharmacokinetic properties of all compounds were then investigated with FAF-Drugs, a tool for prediction of ADME and toxicity. Finally, in order to support *in vitro* studies, molecular docking studies were performed by using AutoDock Vina with a Lamarckian genetic algorithm to determine whether or not the synthesized compounds could be used as inhibitors for the protein structure 1m17 (EGFR). The docking scores of many compounds were found to be higher than [6,7-bis(2-methoxy-ethoxy)quinazoline-4-yl]-(3-ethynyl phenyl)amine, an inhibitor of the 1m17 EGFR receptor. Among the selected compounds **7b**, **7c**, **7e**, **7f**, **7g**, and **8g** showed better stability in the molecular dynamics simulation study.

b][1,3,4] thiadiazoles (III) as non-steroidal ecdysone agonists [20]. Our group combined indolinone with imidazothiadiazole in combination and investigated these chemicals as BCl₂ inhibitors (IV) [21,22]. Another group tested benzenesulfonamide containing imidazothiadiazole as inhibitors of carbonic anhydrase isoforms [23].

Epidermal growth factor receptor (EGFR) is a protein of transmembrane receptor tyrosine kinase protein expressed in other common epithelial, mesenchymal, and neurogenic tissues. Excessive expression of EGFR was observed in many hard tissues, including lung [24], head-neck [25], ovarian [26], and colon tissue. In addition, EGFR is considered one of the most important targets in the development of new anticancer agents [27]. Therefore, EGFR inhibition was studied using the X-ray crystal form structure (PDB ID: 1m17) of the molecular docking simulation method. The molecular docking was supported with molecular dynamics simulation study to understand the ligands stability. The purpose of this work is to carry out a cytotoxicity study against human and murine cancer cells and to determine the drug properties of newly synthesized compounds (V) through pharmacokinetic and toxicity analysis in accordance with Lipinski and FAF-Drugs standardsFigure 1 (Fig. 1).

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2. Experimental

2.1. Chemicals and reagents

Chemicals and reagents used in synthesis were AR grade and procured from local suppliers. The Jasco 430 + was used to record IR sepctra of all the compounds. The Bruker device (400 MHz) was used to record ¹H NMR spectra in CDCl₃/DMSO-d₆ and were reported in hertz (Hz). Various α -bromo ketones were prepared as per literature [28–35].

2.2. Synthetic procedures and spectral data

2.2.1. Preparation of 5-(naphthalen-1-ylmethyl)-1,3,4-thiadiazol-2-amine (3)

Equimolar quantity (0.3 mol) of naphthyl acetic acid (1) and thiosemicarbazide (2) treated in concentrated sulphuric acid (31.5 mL) at 60–70 °C for 8 h. When the reaction conents come to room temperature, poured to cold water & made basic with concentrated aqueous ammonia to pH-7. Filtered the precipitated mass and washed with H₂O to remove excess of ammonia. Re-crystallized from chloroform-ethanol mixture. Yield: 70%; m p: 268-270 °C; FT-IR (cm⁻¹): 3273, 3078, 2964, 2916, 1634, 1523, 1498, 1523, 1498, 1331, 1060. ¹H NMR (400 MHz, DMSO-d₆) δ : 8.14-8.12 (m, 1H, Ar), 8.96-7.94 (m, 1H, Ar), 7.88-7.86 (m, 1H, Ar), 7.56-7.53 (m, 2H, Ar), 7.52-7.48 (m, 2H, Ar), 6.95 (s, br, 2H, NH₂), 4.61 (s, 2H, -CH₂-).

2.2.2. Preparation of 2-(naphthalen-1-ylmethyl)-6-arylimidazo[2,1b][1,3,4]thiadiazole (5a-g)

The derivatives were obtained by the following general procedure. First, 0.03 mol of 2-amino-5-(naphthalen-1-ylmethyl)-1,3,4-thiadiazole (3) and appropriate α -bromo ketone (4) was condensed for 12 h in ethyl alcohol. Solvent was removed and the separated solid was filtered, washed with ethyl alcohol, and neutralized to pH 7.0 using aqueous solution of Na₂CO₃. The precipitated product was dried and purified from ethyl alcohol.

2.2.2.1. 2-(naphthalen-1-ylmethyl)-6-phenylimidazo[2,1-

b][1,3,4]*thiadiazole* (5*a*). FT-IR (cm⁻¹): 3066, 3037, 2947, 2919, 1607, 1525, 1474, 1441, 1258, 1061. ¹H NMR (400 MHz, DMSO-d₆) δ :8.61(s, 1H, Im-H), 8.14 (d, 1H, J=8 Hz), 7.99-7.93 (m, 2H, Ar), 7.81-7.79 (m, 2H, Ar), 7.66 (d, 1H, J=6 Hz), 7.60-7.52 (m, 3H, Ar), 7.37 (t, 2H, J=15.2 Hz), 7.26 (t, 1H, J= 15.0 Hz), 4.93 (s, 2H, -CH₂-). 2.2.2.2. 6-(4-chlorophenyl)-2-(naphthalen-1-ylmethyl)imidazo[2,1-

b][1,3,4]thiadiazole (5b) FT-IR (cm⁻¹): 3093, 3027, 2944, 1592, 1531, 1412, 1065, 999. ¹H NMR (400 MHz, DMSO-d₆) δ : 8.67 (s, 1H, Im-H), 8.10 (d, 1H, J = 8 Hz), 7.99-7.92 (m, 2H, Ar), 7.83 (d, 2H, J = 8.4 Hz), 7.65 (d, 1H, J = 6.4 Hz), 7.60-7.51 (m, 3H, Ar), 7.45 (d, 2H, J = 8.4 Hz), 4.93 (s, 2H, -CH₂-).

2.2.2.3. 6-(4-bromophenyl)-2-(naphthalen-1-ylmethyl)imidazo[2,1-

b][1,3,4]thiadiazole (5c) FT-IR (cm⁻¹): 3120, 3065, 2931, 1585, 1522, 1397, 1071, 1009. ¹H NMR (400 MHz, DMSO-d₆) δ : 8.67 (s, 1H, Im-H), 8.13 (d, 1H, J=8 Hz), 7.99-7.92 (m, 2H, Ar), 7.76 (d, 2H, J=8.2 Hz), 7.65 (d, 1H, J=7.2 Hz), 7.59-7.51 (m, 5H, Ar), 4.93 (s, 2H, -CH₂-).

2.2.2.4. 6-(4-methylphenyl)-2-(naphthalen-1-ylmethyl)imidazo[2,1-

b][1,3,4]*thiadia-zole* (5*d*) FT-IR (cm⁻¹): 3125, 3019, 2927, 2851, 1621, 1545, 1481, 1476, 1302, 1192, 997. ¹H NMR (400 MHz, DMSO-d₆) δ : 8.54 (s, 1H, Im-H), 8.13 (d, 1H, J=8.0 Hz), 7.99-7.92 (m, 2H, Ar), 7.70 (d, 2H, J=8.4 Hz), 7.65 (d, 1H, J=8.0 Hz), 7.89-7.52 (m, 3H, Ar), 7.19 (d, 2H, J=8.0 Hz), 4.92 (s, 2H, -CH₂-), 2.29 (s, 3H, -CH₃).

2.2.2.5. 6-(4-methoxyphenyl)-2-(naphthalen-1-ylmethyl)imidazo[2,1-

b][1,3,4]thiadiazole (5e) FT-IR (cm⁻¹): 3045, 3008, 2959, 2936, 2834, 1612, 1543, 1486, 1468, 1293, 1249, 1173, 1027. ¹H NMR (400 MHz, DMSO-d₆) δ : 8.49 (s, 1H, Im-H), 8.12 (d, 1H, J=8.0 Hz), 7.97-7.92 (m, 2H, Ar), 7.73 (d, 2H, J=8.4 Hz), 7.63 (d, 1H, J=6.4 Hz), 7.59-7.51

(m, 3H, Ar), 6.93 (d, 2H, J=8.4 Hz), 4.91 (s, 2H, -CH₂-), 3.75 (s, 3H, -OCH₃).

2.2.2.6. 2-(naphthalen-1-ylmethyl)-6-(4-nitrophenyl)imidazo[2,1-

b][1,3,4]thiadiazole (5f) FT-IR (cm⁻¹): 3096, 3060, 3003, 2933, 1599, 1504, 1340, 1181, 1108. ¹H NMR (400 MHz, DMSO-d₆) δ : 8.92 (s, 1H, Im-H), 8.25 (d, 2H, J=8.4 Hz), 8.13 (d, 1H, J=8.0 Hz), 8.07 (d, 2H, J=8.4 Hz), 7.99-7.93 (m, 2H, Ar), 7.66 (d, 1H, J=6.4 Hz), 7.59-7.52 (m, 3H, Ar), 4.95 (s, 2H, -CH₂-).

2.2.2.7. 3-[2-(naphthalen-1-ylmethyl)imidazo[2,1-b][1,3,4]thiadiazol-6yl]-2H-chromen-2-one **(5g)** FT-IR (cm⁻¹): 3068, 3047, 2969, 2945, 1716, 1513, 1472, 1452, 1472, 1452, 1472, 1394, 1212, 1112. ¹H NMR (400 MHz, DMSO-d₆) δ : 8.62 (s, 1H, Im-H), 8.55 (s, 1H, Coum-H), 8.13 (d, 1H, 8.0 Hz), 7.99-7.94 (m, 2H, Ar), 7.85-7.83 (m, 1H, Ar), 7.67-7.65 (m, 1H, Ar), 7.61-7.52 (m, 4H, Ar), 7.45 (d, 1H, J=8.4 Hz), 7.38 (t, 1H, J=16 Hz), 4.95 (s, 2H, -CH₂-).

2.2.3. Preparation of 5-bromo-2-(naphthalen-1-ylmethyl)-6-phenylimidazo[2,1-b][1,3,4] thiadiazole (6a-e)

First, 0.01 mol of bromine was added to a well stirred mixture of powdered anhydrous sodium acetate and 0.01 mol of 2-(naph-thalen-1-ylmethyl)-6-arylimidazo[2,1-b][1,3,4]thiadiazole at room temperature. The agitation was continued for 1 h and poured into cold water. The solid that formed was filtered, washed with water, and purified. 2.2.3.1. 5-bromo-2-(naphthalen-1-ylmethyl)-6-phenylimidazo[2,1-

 $b][1,3,4] \ thiadiazole \ (6a) \ \ FT-IR \ (cm^{-1}): 3058, 2962, 2938, 1599, 1528, 1510, 1472, 1440, 1334, 1114, 1071, 966. ^1H \ \ NMR \ (400 \ \ MHz, DMSO-d_6) \ \delta: 8.14 \ (d, 1H, J=8.0 \ \ Hz), 7.97-7.90 \ (m, 4H, Ar), 7.67 \ (d, 1H, J=6.4 \ \ Hz), 7.60-7.52 \ (m, 3H, Ar), 7.46 \ (t, 2H, J=15.4 \ \ \ Hz), 7.36-7.32 \ (m, 1H, Ar), 4.97 \ (s, 2H, -CH_2-).$

2.2.3.2. 5-bromo-6-(4-chlorophenyl)-2-(naphthalen-1-

ylmethyl)imidazo[2,1-b][1,3,4] thiadiazole (**6b**) FT-IR (cm⁻¹): 3065, 2932, 2891, 1605, 1535, 1529, 1468, 1368, 1109, 1057. ¹H NMR (400 MHz, DMSO-d₆) δ : 8.13 (d, 1H, J=8.0 Hz), 8.00-7.92 (m, 4H, Ar), 7.66 (d, 1H, J=7.2 Hz), 7.61-7.50 (m, 5H, Ar), 4.98 (s, 2H, -CH₂-).

2.2.3.3. 5-bromo-6-(4-bromophenyl)-2-(naphthalen-1-

ylmethyl)imidazo[2,1-b][1,3,4] thiadiazole (6c) FT-IR (cm⁻¹): 3063, 3010, 2936, 1903, 1520, 1475, 1395, 1327, 1096, 1006, 967. ¹H NMR (400 MHz, DMSO-d₆) δ : 8.15 (d, 2H, J=8 Hz), 8.01-7.99 (m, 1H, Ar), 7.97 (d, 1H, J=8.8 Hz), 7.89 (d, 2H, J=8.8 Hz), 7.67 (d, 3H, J=8.8 Hz), 7.62-7.53 (m, 3H, Ar), 4.99 (s, 2H, -CH₂-).

2.2.3.4. 5-bromo-6-(4-methylphenyl)-2-(naphthalen-1-

ylmethyl)imidazo[2,1-b][1,3,4] thiadiazole (6d) FT-IR (cm⁻¹): 3076, 3001, 2941, 1515, 1484, 1405, 1321, 1096, 1016. ¹H NMR (400 MHz, DMSO-d₆) δ : 8.13 (d, 1H, J=8.4 Hz), 7.98-7.94 (m, 2H, Ar), 7.81 (d, 2H, J=8.0 Hz), 7.66-7.52 (m, 4H, Ar), 7.26 (d, 2H, J=8.0 Hz), 4.97 (s, 2H, -CH₂-), 2.32 (s, 3H, -CH₃).

2.2.3.5. 5-bromo-6-(4-methoxyphenyl)-2-(naphthalen-1-

ylmethyl)imidazo[2,1-b][1,3,4] thiadiazole (6e) FT-IR (cm⁻¹): 3127, 3045, 3010, 2960, 2935, 1611, 1543, 1484, 1396, 1300, 1249, 1173, 1027.

2.2.3.6. 5-bromo-2-(naphthalen-1-ylmethyl)-6-(4-nitrophenyl)imidazo[2,1b][1,3,4] thiadiazole (6f) FT-IR (cm⁻¹): 3107, 3025, 3005, 2959, 2829, 1625, 1491, 1416, 1315, 1198. ¹H NMR (400 MHz, DMSO-d₆) δ : 8.33 (d, 2H, J=8.8 Hz), 8.22 (d, 2H, J=9.2 Hz), 8.15 (d, 1H, J=8.4 Hz), 8.00-7.94 (m, 2H, Ar), 7.67 (d, 1H, J=7.2 Hz), 7.62-7.53 (m, 3H, Ar), 5.00 (s, 2H, -CH₂-).

2.2.3.7. 3-[5-bromo-2-(naphthalen-1-ylmethyl)imidazo[2,1-

b][1,3,4]thiadiazol-6-yl]-2H chromen-2-one (6g) FT-IR (cm⁻¹): 3105, 3067, 3005, 2954, 2831, 1629, 1521, 1491, 1399, 1312, 1213, 1198, 1001. ¹H NMR (400 MHz, DMSO-d₆) δ : 8.29 (s, 1H, Coum-H), 8.14 (d, 1H, J=8.4 Hz), 8.00-7.95 (m, 2H, Ar), 7.83-7.80 (m, 1H, Ar), 7.68-7.65 (m, 2H, Ar), 7.63-7.53 (m, 3H, Ar), 7.46 (d, 1H, J=8.4 Hz), 7.38-7.34 (m, 1H, Ar), 4.96 (s, 2H, -CH₂-).

2.2.4. Synthesis of 2-(naphthalen-1-ylmethyl)-6-arylimidazo[2,1-

b][1,3,4]thiadiazole-5-carbaldehyde (7a-g)

First, added 4 mmol of 2-(4-methoxybenzyl)-6-aryl-imidazo[2,1-*b*][1,3,4]thiadiazole to freshly obtained Vilsmeier-Haack mixture at room temperature with stirring. Continued stirring for 4 h at 80-90°C. Later the reaction mixture was poured into cold water and neutralized to pH 7.0 with an aqueous solution of Na_2CO_3 . The product was filtered and re-crystallized from ethyl alcohol.

2.2.4.1. 2-(naphthalen-1-ylmethyl)-6-phenylimidazo[2,1-

b][1,3,4]*thiadiazole-5-carbaldehyde* (7a) FT-IR (cm⁻¹): 3091, 3015, 2929, 2899, 1675, 1563, 1519, 1487, 1455, 1349, 1338, 1091. ¹H NMR (400 MHz, DMSO-d₆) δ : 10.05 (s, 1H, -CHO), 8.06 (m, 1H, Ar), 7.94-7.90 (m, 2H, Ar), 7.82-7.70 (m, 2H, Ar), 7.58-7.45 (m, 7H, Ar), 4.91 (s, 2H, -CH₂-).

2.2.4.2. 6-(4-chlorophenyl)-2-(naphthalen-1-ylmethyl)imidazo[2,1-

b][1,3,4] thiadiazole-5-carbaldehyde (7b) FT-IR (cm⁻¹): 3125, 3009, 2899, 1675, 1599, 1529, 1491, 1355, 1339, 1111. ¹H NMR (400 MHz, DMSO-d₆) δ : 10.09 (s, 1H, -CHO), 8.03 (d, 1H, J=8.4 Hz), 7.96-7.90 (m, 2H, Ar), 7.83 (d, 2H, J=8.4 Hz), 7.58-7.51 (m, 4H, Ar), 7.49 (d, 2H, J=8.8 Hz), 4.91 (s, 2H, -CH₂-).

2.2.4.3. 6-(4-bromophenyl)-2-(naphthalen-1-ylmethyl)imidazo[2,1-

b][1,3,4] thiadiazole-5-carbaldehyde (7c) FT-IR (cm⁻¹): 3151, 3015, 2915, 2892, 1681, 1598, 1515, 1478, 1450, 1350, 1325, 1091. ¹H NMR (400 MHz, DMSO-d₆) & 10.07 (s, 1H, -CHO), 8.05 (d, 1H, J=8.4 Hz), 7.93 (m, 2H, Ar), 7.75 (d, 2H, J=8.4 Hz), 7.62 (d, 2H, J=8.4 Hz), 7.58-7.49 (m, 4H, Ar), 4.90 (s, 2H, -CH₂-).

2.2.4.4. 6-(4-methylphenyl)-2-(naphthalen-1-ylmethyl)imidazo[2,1b][1,3,4] thiadiazole-5-carbaldehyde (7d) FT-IR (cm⁻¹): 3125, 3025, 2975, 2891, 1665, 1609, 1509, 1465, 1448, 1313, 1121. ¹H NMR (400 MHz, DMSO-d₆) δ : 9.94 (s, 1H, -CHO), 8.15 (d, 1H, J=8.4 Hz), 8.00-7.94 (m, 2H, Ar), 7.80 (d, 2H, J=6.8 Hz), 7.61-7.52 (m, 3H, Ar), 7.43 (d, 2H, J=8.4 Hz), 5.02 (s, 2H, -CH₂-), 2.35 (s, 3H, -CH₃).

2.2.4.5. 6-(4-methoxyphenyl)-2-(naphthalen-1-ylmethyl)imidazo[2,1b][1,3,4] thiadiazole-5-carbaldehyde (7e) FT-IR (cm⁻¹): 3057, 3005, 2936, 2836, 1661, 1607, 1524, 1482, 1395, 1253, 1176, 1110. ¹H NMR (400 MHz, DMSO-d₆) δ : 10.00 (s, 1H, -CHO), 7.94 (d, 2H, J=8.0 Hz), 7.88-7.84 (m, 1H, Ar), 7.75 (d, 2H, J=8.4 Hz), 7.51-7.42 (m, 4H, Ar), 6.97 (d, 2H, J=8.8 Hz), 4.85 (s, 2H, -CH₂-), 3.78 (s, 3H, -OCH₃).

2.2.4.6. 2-(naphthalen-1-ylmethyl)-6-(4-nitrophenyl)imidazo[2,1b][1,3,4]thiadiazole-5 carbaldehyde (**7f**) FT-IR (cm⁻¹): 3066, 3005, 2975, 2850, 1677, 1600, 1523, 1478, 1448, 1342, 1313, 1108. ¹H NMR (400 MHz, DMSO-d₆) δ : 10.07 (s, 1H, -CHO), 8.34 (d, 2H, J=9.2 Hz), 8.25 (d, 2H, J=9.2 Hz), 8.15 (d, 1H, J=8.0 Hz), 8.00-7.95 (m, 2H, Ar), 7.69 (d, 1H, J=6.4 Hz), 7.61-7.53 (m, 3H, Ar), 5.05 (s, 2H, -CH₂-).

2.2.4.7. 2-(naphthalen-1-ylmethyl)-6-(2-oxo-2H-chromen-3-

yl)imidazo[2,1-b] [1,3,4]thiadiazole-5-carbaldehyde (**7g**) FT-IR (cm⁻¹): 3106, 3009, 2991, 2897, 1683, 1591, 1501, 1467, 1325, 1125. ¹H NMR (400 MHz, DMSO-d₆) δ : 10.01 (s, 1H, -CHO), 8.49 (s, 1H, Coum-H), 8.13 (d, 1H, J=8.0 Hz), 8.01-7.95 (m, 2H, Ar), 7.87 (d, 1H, J=8.0 Hz), 7.71-7.67 (m, 2H, Ar), 7.62-7.53 (m, 3H, Ar), 7.48 (d, 1H, J=7.6 Hz), 7.38 (t, 1H, J=15.4 Hz), 5.04 (s, 2H, -CH₂-).

2.2.5. Procedure for the preparation of 2-(naphthalen-1-ylmethyl)-6arylimidazo [2,1-b][1,3,4]thiadiazol-5-yl thiocyanate (8a-g)

To a suspension of 8 mmol of potassium thiocyanate and 4 mmol of 2-(4-methoxybenzyl)-6-aryl-imidazo[2,1-*b*][1,3,4]thiadiazole, 4 mmol of bromine (in 10 mL of glacial acetic acid) was added under ice at 0-5°C. The reaction mixture was stirred for 3 h at room temperature, then poured into ice-cold water and filtered. The product was re-crystal-lized from ethyl alcohol.

2.2.5.1. 2-(naphthalen-1-ylmethyl)-6-phenylimidazo[2,1-

b][1,3,4]thiadiazol-5-yl thiocyanate (8a) FT-IR (cm⁻¹): 3091, 3065, 3048, 2935, 2841, 2156, 1598, 1510, 1469, 1340, 1261, 1107. ¹H NMR (400 MHz, DMSO-d₆) δ : 8.19 (d, 1H, J=8.0 Hz), 8.00-7.91 (m,

4H, Ar), 7.70 (d, 1H, J=6.0 Hz), 7.61-7.50 (m, 5H, Ar), 7.46-7.42 (m, 1H, Ar), 5.06 (s, 2H, -CH₂-).

2.2.5.2. 6-(4-chlorophenyl)-2-(naphthalen-1-ylmethyl)imidazo[2,1-

b][1,3,4]*thiadiazol-5-yl thiocyanate* (*8b*) FT-IR (cm⁻¹): 3101, 3055, 2948, 2839, 2158, 1601, 1508, 1469, 1345, 1262, 1099. ¹H NMR (400 MHz, DMSO-d₆) & 8.16 (d, 1H, J = 8.0 Hz), 8.00-7.92 (m, 4H, Ar), 7.70 (d, 1H, 8.0 Hz), 7.60-7.53 (m, 5H, Ar), 5.06 (s, 2H, -CH₂-).

2.2.5.3. 6-(4-bromophenyl)-2-(naphthalen-1-ylmethyl)imidazo[2,1-

b][1,3,4]thiadiazol-5-yl thiocyanate (8c) FT-IR (cm⁻¹): 3101, 3005, 2905, 2839, 2166, 1599, 1515, 1475, 1339, 1275, 1111. ¹H NMR (400 MHz, DMSO-d₆) δ : 8.07 (d, 1H, J = 8.0 Hz), 7.95-7.90 (m, 2H, Ar), 7.83 (d, 2H, J = 8.4 Hz), 7.64-7.49 (m, 6H, Ar), 4.87 (s, 2H, -CH₂-).

2.2.5.4. 6-(4-methylphenyl)-2-(naphthalen-1-ylmethyl)imidazo[2,1-

b][1,3,4]*thiadiazo*1-5-*y*1 *thiocyanate* (8*d*) FT-IR (cm⁻¹): 3046, 3017, 2923, 2868, 2159, 1513, 1469, 1326, 1292, 1264, 1186, 1116. ¹H NMR (400 MHz, DMSO-d₆) & 8.04 (d, 1H, J=8.4 Hz), 7.95-7.92 (m, 2H, Ar), 7.83 (d, 2H, J=8.0 Hz), 7.61-7.50 (m, 4H, Ar), 7.33 (d, 2H, J=8.0 Hz), 4.89 (s, 2H, -CH₂-), 2.17 (s, 3H, -CH₃).

2.2.5.5. 6-(4-methoxyphenyl)-2-(naphthalen-1-ylmethyl)imidazo[2,1-

b][1,3,4] thiadiazol-5-yl thiocyanate (8e) FT-IR (cm⁻¹): 3049, 3005, 2945, 2878, 2161, 1598, 1491, 1475, 1255, 1111. ¹H NMR (400 MHz, DMSO-d₆) δ : 7.97 (d, 1H, J=8.4 Hz), 7.88-7.80 (m, 4H, Ar), 7.53-7.42 (m, 4H, Ar), 6.96 (d, 2H, J=8.8 Hz), 4.81 (s, 2H, -CH₂-), 3.80 (s, 3H, -OCH₃).

2.2.5.7. 2-(naphthalen-1-ylmethyl)-6-(4-nitrophenyl)imidazo[2,1-

b][1,3,4]*thiadiazol-5-yl thiocyanate* (*8f*) FT-IR (cm⁻¹): 3091, 3065, 3048, 2935, 2841, 2156, 1598, 1510, 1469, 1340, 1261, 1107.

2.2.5.8. 2-(naphthalen-1-ylmethyl)-6-(2-oxo-2H-chromen-3-

yl)imidazo[2,1-b][1,3,4]thiadiazol-5-yl thiocyanate (8g) FT-IR (cm⁻¹): 3071, 3049, 2977, 2928, 2158, 1708, 1606, 1483, 1372, 1163, 1045. ¹H NMR (400 MHz, DMSO-d₆) δ: 8.45 (s, 1H, Ar), 8.21-8.19 (m, 1H, Ar), 8.01-7.96 (m, 2H, Ar), 7.88-7.86 (m, 1H, Ar), 7.72-7.69 (m, 2H, Ar), 7.59-7.55 (m, 3H, Ar), 7.51 (d, 1H, J=8.4 Hz), 7.42 (t, 1H, J=16 Hz), 5.09 (s, 2H, -CH₂-).

2.6. Bio-evaluation

The cytotoxicities of the compounds illustrated in Scheme 1 were studied against human cervix carcinoma HeLa, human T-lymphocyte CEM, and murine leukemia L1210 cell lines [36]. The IC_{50} values were calculated and expressed in μ M. All experiments were performed in triplicate.

2.7. Computational study

2.7.1. Details of theoretical calculations

Optimized structures of all compounds were obtained using the 6-311G(d) basis set [37] and DFT/B3LYP density functional theory [38,39] in Gaussian 09 [40]. AutoDock Vina [41], which includes Lamarckian genetics, was used in PyRx software [42] as the scoring algorithm for molecular docking virtual screening operations. Some molecular descriptors commonly used in absorption, distribution, metabolism, and elimination (ADME) analysis were calculated using the FAF-Drugs4 webserver [43].

2.7.2. Molecular dynamics simulation studiesScheme 1

Desmond's Schrodinger tool was used to perform molecular dynamics (MD) simulation studies, which led to the understanding of the binding of ligand-protein complex in the simulated physiological solvent-based system [44,45]. EGFR pdbID: 1M17 truncated to remove C and N terminal loops. The fixed structure of the EGFR (residues: 684-951) was used for molecular dynamics. The Molecular Dynamic simulation is performed for 20ns in all chemicals using Desmond under the isothermal-isobaric ensemble at temperatures and pressures set at 300K and 1.013 bar, respectively. TIP4PEW was used as a solvent model. The boundary is set to 8Å in the form of a box that is orthorhombic. NaCl



Scheme 1. Synthesis of 2-(naphthalen-1-yl)-methyl-5,6-disubstituted imidazo[2,1-b][1,3,4]thiadiazole.

salt at 0.15M was added, Cl-counter ions were added to neutralize the system. The recording interval trajectory, every 25 ps and 1000 frames were captured throughout simulation. Root means square deviation (RMSD) calculated by all complex protein combinations. The RMSD shows a moderate change in the location of selected atoms in atoms by comparisons made with the trajectory framework.

3. Results and discussion

3.1. Chemistry

5-(Naphthalen-1-yl)-methyl-1,3,4-thiadiazole-2-amine (3) was prepared by slowly adding naphthalene-1-acetic acid (1) with stirring to a suspension of thiosemicarbazide (2) in sulfuric acid at 60-70°C for 8 h and cooled to room temperature, added to water, and made basic with an ammonia solution. Different 6-substituted 2-(naphthalen-1-yl)-methylimidazo[2,1-b][1,3,4]thiadiazoles (5a-g) were obtained by reacting 3 with α -bromo ketones (4) in ethyl alcohol by heating for 12 h and neutralizing with aqueous sodium carbonate. The electrophilic substitution reaction was performed at the 5th position of the imidazothiadiazoles (5) to get compounds 6a-g, 7a-g, and 8a-gFigure Scheme 1 (Scheme 1).

Infrared spectroscopy exhibited characteristic C-H stretching vibrations in the range of 3129-3000 cm⁻¹. Aliphatic C-H stretching vibrations were seen between 2999 and 2850 cm⁻¹ and C=O stretching vibrations between 1720 and 1640 cm⁻¹ for derivatives **5g**, **6g**, **7a-g**, and **8g**. The derivatives of SCN (**8a-g**) exhibited stretching vibrations between 2170 and 2150 cm⁻¹.

Protons of the imidazole at C₅ were exhibited at 8.92-8.49 ppm for derivatives **5a-g** in their ¹H NMR spectra, which confirms the reaction between aminothiadiazole (**3**) and the respective 2-bromoketone (**4**) derivatives. Substitution at the C₅ position of imidazothiadiazole was confirmed by the absence of the C₅-H proton in the respective derivatives (**6a-g, 7a-g**, and **8a-g**).

The presence of a formyl proton was observed for compounds 7a-g at 10.09 and 9.94 δ ppm for their respective proton NMR, which con-

firms the Vilsmeier-Haack reaction on the imidazothiadiazole derivatives (**5a-g**).

All the compounds containing aromatic proton signals exhibited prominent multiplet signals between 8.55 and 6.97 δ ppm. The methylene proton (C2 of imidazothiadiazole) was seen between 5.05 and 4.85 δ ppm as a singlet. Methoxy protons of derivatives **5g**, **6g**, **7g**, and **8g** were observed as singlets between 3.80 and 3.75 δ ppm. Methyl protons for derivatives **5d**, **6d**, **7d**, and **8d** appeared as singlets between 2.35 and 2.30 δ ppm. Chemical characterization of the synthesized compounds is given in Table 1.

3.2. Cytotoxicity study

Evaluation of the cytotoxic activity of 5-(naph-thalen-1-yl)-methyl-6-aryl-imidazo[2,1-*b*][1,3,4]-thiadiazole derivatives was performed with leukemia murine and human cell lines.

The cytotoxicities of the 27 derivatives of imidazo[2,1-b][1,3,4]thiadiazoles and standards melphalan and levamisole were studied by screening using the CEM, HeLa, and L1210 cells. Among the series of 5a-g against these three cell lines, only 5g exhibited equipotent activity against L1210 [2.1 µM] and HeLa cells [4.0 µM] in comparison to standard melphalan [2.13 µM], and it was very much potent in comparison to levamisole [206 µM to >250 µM]. The remaining derivatives (5a-f) were poor in cytotoxicity for all three cell lines tested. Introduction of bromine at the 5th position of imidazothiadiazole gave 5-bromo derivatives 6a-g in good yield. Among these 5-bromo derivatives, only 6g emerged as the most potent cytotoxic compound against L1210 [3.4 µM] and HeLa cells [4.5 µM]. This clearly reveals that 2H-chromen-2-one-3-yl substitution was preferred over the phenyl group at the 6th position of the parent fused heterocyclic ring. The remaining 5-bromo derivatives failed to emerge as cytotoxic compounds against the tested cell lines. Formyl group substitution was introduced at the 5th position by Vilsmeier-Haack reaction on the imidazothiadiazole fused ring to give compounds 7a-g in good yields. Most of these formvl deriv-

Table 1 Chemical characterization of the synthesized compounds.

	R	Code	MF	MW	Yield %	MP°C
	Ph	5a	$C_{21}H_{15}N_3S$	341.43	58	196-198
(5a-g)	4-Cl-Ph 4-Br-Ph 4-CH ₃ -Ph 4-OCH ₂ -Ph	5b 5c 5d 5e	C ₂₁ H ₁₄ ClN ₃ S C ₂₁ H ₁₄ BrN ₃ S C ₂₂ H ₁₇ N ₃ S C ₂₂ H ₁₇ N ₃ S	375.87 420.32 355.46 371.45	61 64 60 58	198-200 198-202 180-182 194-196
Br N-N-R	4-NO ₂ -Ph 2H-chromen-2-one-3-yl Ph	5f 5g 6a	C ₂₁ H ₁₄ N ₄ O ₂ S C ₂₄ H ₁₅ N ₃ O ₂ S C ₂₁ H ₁₄ BrN ₃ S	386.43 409.46 420.32	55 62 63	268-270 220-222 128-130
(6a-g)	4-CI-Ph	6b	C ₂₁ H ₁₃ BrClN ₃ S	454.77	67	154-156
	4-Br-Ph 4-CH ₃ -Ph 4-OCH ₃ -Ph 4-NO ₂ -Ph 2H-chromen-2-one-3-yl	6c 6d 6e 6f 6g	$C_{21}H_{13}N_3S \\ C_{22}H_{16}BrN_3S \\ C_{22}H_{16}BrN_3OS \\ C_{21}H_{13}BrN_4O_2S \\ C_{24}H_{14}BrN_3O_2S \\ C_{24}H_{14}Br$	499.22 434.35 450.53 465.32 488.36	62 61 65 67 58	168-170 152-154 138-140 184-186 226-228
(7a-g)	Рһ	7a	C ₂₂ H ₁₅ N ₃ OS	369.44	52	154-156
	4-Cl-Ph	7b	C22H14ClN3OS	403.88	51	172-174
	4-Br-Ph	7c 7d	C ₂₂ H ₁₄ BrN ₃ OS	448.34	48	138-140
	4-OCH ₃ -Ph	7e	C ₂₃ H ₁₇ N ₃ O ₂ S	399.46	52	90-92
	4-NO ₂ -Ph	7f	C ₂₂ H ₁₄ N ₄ O ₃ S	414.44	53	170-172
	2H-chromen-2-one-3-yl	7g	C ₂₅ H ₁₅ N ₃ O ₃ S	437.47	48	98-101
$(8a-g) \xrightarrow{N_{N}} (S \rightarrow \mathbb{R})$		88	C ₂₂ H ₁₄ N ₄ S ₂	398.50	58	148-150
	4-Cl-Ph	8b	C22H13ClN4S2	432.95	61	144-146
	4-Br-Ph	8c	$C_{22}H_{13}BrN_4S_2$	477.40	60	146-148
	4-CH ₃ -Ph	8d	C ₂₃ H ₁₆ N ₄ S ₂	412.53	60	102-104
	4-OCH ₃ -Ph	8e of	$C_{23}H_{16}N_4OS_2$	428.52	61	160-162
	4-NO ₂ -Pfi 2H-chromen-2-one-3-vl	81 89	$C_{22}H_{13}N_5O_2S_2$ $C_{25}H_{14}N_4O_2S_2$	443.50	57	204-200 258-260
	211 Chromen-2-one-3-yr	05	0251114140202	100.00	57	200-200

atives became good cytotoxins against all three cell lines in the range of 4.4-9.7 μ M in comparison to the standard drug melphalan [1.4-2.13 μ M]. Introduction of the thiocyanate (–SCN) group was carried out at the C₅ position of the imidazothiadiazole fused ring, which resulted in the formation of 5-thiocyanate compounds **8a-g**. Thiocyanate-derivative imidazothiadiazoles were moderately cytotoxic against all three cells, especially against HeLa cells. Compound **8e** (IC₅₀: 9.8 μ M) emerged as having the most potent cytotoxic activity against cervix carcinoma cells.

In general, the 5-thiocyanate derivatives exhibited moderate and the 5-formyl derivatives exhibited good cytotoxicity in comparison to the positive controls, melphalan and levamisole.

Electrophiles such as bromine, formyl, and thiocyanate groups were substituted at the 5^{th} position and aryl groups at the 6^{th} position of the

2-(naphthalen-1-ylmethyl)-imidazothiadiazole fused ring, respectively, to understand the structure-activity relationship (SAR). The presence of the bromine atom did not produce much improvement in cytotoxic activity against all three cell lines tested except for compound **5g** against L1210 leukemia cells [2.1 μ M] and against HeLa cervix carcinoma cells [4.0 μ M]. Compound **5g** emerged as equipotent with the standard drug melphalan. This was due to the presence of 2*H*-1-benzopyran-2-one-3-yl at the C₆ position of the imidazothiadiazole-fused ring. Formyl group substitution at C₅ of the imidazothiadiazole-fused ring provided good cytotoxins as many of these compounds became potent except **7d** and **7f**. The third set of modifications done by thiocyanate substitution at the C₅ position of the imidazothiadiazole-fused ring gave compounds with moderate cytotoxicity against all three cell lines tested. Among

these thiocyanate derivatives, only compound 8e proved to be a good cytotoxin against HeLa cells [9.8 μ M] (Table 2).

3.3. Molecular docking and calculation of physicochemical and ADME properties

Table 3 shows the molecular weight (MW), human oral absorption percentage, indicator of hydrophobicity (logP), topological polar surface area (tPSA), and violation of Lipinski's rule of five (Ro5) [46] calculated for all compounds. The Ro5 was developed by Lipinski *et al.* as a guide to designing molecules that can be used orally and it has had a notable impact on drug discovery strategies. Four key parameters were selected for 90% of the molecules studied that could be taken orally:

- A molecular weight (MW) below 500 Da
- A calculated octanol-water coefficient (CLogP) smaller than 5
- No more than 5 hydrogen bond donors
- No more than 10 hydrogen bond acceptors

The results in Table 3 indicate that other compounds having experimental activity besides the co-ligand violate the Ro5. Another important value is the tPSA (recommended value: $\leq 140 \text{ Å}^2$) [47], and this value was within the recommended range for all compounds except **8f**. On the other hand, Fsp³ (fraction of saturated carbons) is a newer index indicating drug similarity [48]. Lovering *et al.* [49] pointed out that a decreasing Fsp³ value leads to increased CYP inhibition effect. The Fsp³ values of all compounds were found to be lower than those of the co-ligand, melphalan, and levamisole.

Table 2	
Antiproliferative effects of the compounds on L1210, CEM and HeLa cells.	

Compound	IC ₅₀ * (μM)		
	L1210	CEM	HeLa
5a	116 ± 6	138 ± 13	87 ± 2
5b	94 ± 49	108 ± 33	78 ± 13
5c	> 250	$111~\pm~20$	≥ 250
5d	> 250	> 250	> 250
5e	131 ± 20	111 ± 7	120 ± 36
5f	116 ± 15	121 ± 6	117 ± 52
5g	$2.1 \pm 0.8^{**}$	80 ± 5**	$4.0 \pm 3.1^{**}$
6a	≥ 250	136 ± 84	> 250
6b	120 ± 13	82 ± 12	79 ± 14
6c	94 ± 14	71 ± 0	66 ± 18
6d	113 ± 21	77 ± 2	81 ± 8
6e	70 ± 10	51 ± 5	75 ± 2
6f	71 ± 54	64 ± 17	85 ± 4
6g	3.4 ± 1.5	54 ± 6	4.5 ± 0.8
7a	9.7 ± 0.1	7.9 ± 1.6	8.2 ± 0.8
7b	5.0 ± 0.5	5.4 ± 0.3	4.6 ± 0.4
7c	4.8 ± 0.3	4.9 ± 0.4	4.4 ± 0.5
7d	127 ± 4	113 ± 3	86 ± 14
7e	4.9 ± 0.5	5.0 ± 0.0	7.7 ± 0.2
7f	107 ± 21	95 ± 11	74 ± 12
7g	NT	NT	NT
8a	57 ± 39	44 ± 34	13 ± 8
8b	66 ± 0	40 ± 21	13 ± 6
8c	79 ± 31	51 ± 0	27 ± 6
8d	82 ± 11	42 ± 20	25 ± 5
8e	62 ± 35	53 ± 1	9.8 ± 1.9
8f	126 ± 4	103 ± 5	70 ± 5
8g	16 ± 3	70 ± 10	28 ± 22
Melphalan	2.13 ± 0.02	1.4 ± 0.4	NT
Levamisole	206	>250	>250

In addition, Table 3 shows the pharmacological properties of the compounds as well as their binding affinities, and compounds **5g**, **6g**, **7a**, **7b**, **7c**, **7e** and **8e** found to be effective in *in vitro* studies, and exhbited the highest binding affinity results compared to the co-ligand, melphalan, and levamisole. Similarly, *in vitro* study showed that the docking score of compound **5c**, which had the lowest activity, was also low.

The properties are PhysChem and FAFDrugs filters. HBD: hydrogen bond donor; HBA: hydrogen bond acceptor; tPSA: topological polar surface area; LogP: indicator of hydrophobicity.

In molecular docking studies, the X-ray crystal structure (1m17) of EGFR was used as a target and [6,7-bis(2-methoxy-ethoxy)quinazoline-4-yl]-(3-ethynyl phenyl)amine, the inhibitor of this crystal structure, was used as the co-ligand. The 1m17 receptor and all compounds were prepared for docking operations using the protein and ligand preparation wizards in the PyRx package. Docking operations were performed using similar protocols to our previous study [50] for all compounds, the co-ligand molecules, and standards melphalan and levamisole, and the docking scores of all compounds are shown in Table 3. The 3D interactions between the receptor and **5c**, **5g**, **6g**, the co-ligand, and standards melphalan and levamisole are shown in Figure 2Fig. 2.

Fig. 3 Figure 3shows the 3D representation of the co-ligand settling in the active binding sites of the target protein and compounds **5g** and **6g**, which had high docking scores, and compound **5c**, which had a low docking score. Finally, as shown in Fig. 2, all 4 ligands were also observed to be oriented parallel to each other at receptor binding sites.

Molecular Dynamics (MD) simulation studies provide an understanding of the molecular complex of protein-ligand complex interactions and their stability in simulated physiological conditions, employing a Desmond Maestro module. MD simulations were performed on all synthesized compounds. The frame was taken during the 25 ps simulation and kept in trajectory and was followed by the production of 1000 frames in each combination over a simulation time of 25 ns. The 'Ligfit Episode' shows the RMSD (Root Mean Square Deviation) structure of ligand and protein structure. In all compounds, protein-ligand interactions or contacts are observed; in particular the interaction was three types or otherwise four types: hydrogen bonds, hydrophobic intercations, ionic bridges and water (Table 4). Most of the compounds showed strong hydrogen bond interaction with residue MET769, except for the compounds 5d, 5f, 6a-c, 6e and 8c. In addition to Met769, compounds 8a and 8d formed hydrogen bond with residue CYS773, 7d with residue LYS73 and 8g with residue THR830. Concerning hydrophobic interactions, residues LEU694, ALA719, and LEU820 were the commonly involved binding with most of the compounds. Besides these there were some ionic and water bridge interactions observed for most of the compounds (Fig.S1). Among the performed molecular dynamics-based simulation studies, the enzyme complex with the compounds 7b, 7c, 7e, 7f, 7g, and 8g showed more stable binding based on RMSD plot shown inFigure 4, 5 and 6 Figs. 4-6 and observed interactions.

4. Conclusions

Cytotoxic activities for 28 compounds and two standards were tested against L1210, CEM, and HeLa cells. Among these compounds, **5g**, **6g**, **7a**, **7b**, **7c**, and **7e** emerged as the most potent. Overall, formyl substitution was preferred at the 5th position of the imidazo[2,1-*b*][1,3,4]thia-diazole nucleus as most of the formyl compounds were cytotoxic to all three tested cell lines. The order of potency at the 5th position was CHO>SCN>Br>H. The potency of these compounds was supported by a docking study against the 1m17 EGFR receptor. In molecular docking studies, it was observed that all compounds had higher docking scores than [6,7-bis(2-methoxy-ethoxy)quinazo-line-4-yl]-(3-ethynylphenyl)amine, an inhibitor of the 1m17 receptor. Molecu-

Table 3
The pharmacological properties of virtual screening and rationally designed compounds.

Compounds	Binding Affinity (kcal/mol)	MW	logP	tPSA	Rotatable Bonds	Flexibility	HBD	HBA	Lipinski Violation	Solubility (mg/L)	Solubility ForecastIndex	Oral Bioavailability VEBER	Fsp ³
5a	-9.0	341.43	5.72	58.43	3	0.1	0	3	1	1046.22	Reduced	Good	0.05
5b	-8.9	375.87	6.34	58.43	3	0.1	0	3	1	629.48	Reduced	Good	0.05
5c	-8.8	420.32	6.41	58.43	3	0.1	0	3	1	511.31	Reduced	Good	0.05
5d	-9.1	355.46	6.08	58.43	3	0.1	0	3	1	819.85	Reduced	Good	0.09
5e	-8.8	371.45	5.69	67.66	4	0.13	0	4	1	1069.18	Reduced	Good	0.09
5f	-9.1	386.43	5.54	104.25	4	0.13	0	6	1	1093.8	Reduced	Good	0.05
5g	-10.3	409.46	5.71	88.64	3	0.09	0	5	1	812.93	Reduced	Good	0.04
6a	-9.2	420.32	6.74	58.43	3	0.1	0	3	1	415.33	Reduced Solubility	Good	0.05
6b	-8.9	454.77	7.37	58.43	3	0.1	0	3	1	244.05	Reduced	Good	0.05
6c	-8.6	499.22	7.43	58.43	3	0.1	0	3	1	195.83	Reduced	Good	0.05
6d	-9.0	434.35	7.11	58.43	3	0.1	0	3	1	321.02	Reduced Solubility	Good	0.09
6e	-8.8	450.35	6.71	67.66	4	0.13	0	4	1	418	Reduced	Good	0.09
6f	-9.1	465.32	6.57	104.25	4	0.13	0	6	1	422.06	Reduced Solubility	Good	0.05
6g	-10.3	488.36	6.74	88.64	3	0.09	0	5	1	310.69	Reduced Solubility	Good	0.04
7a	-9.0	369.44	5.51	75.5	4	0.13	0	4	1	1184.03	Reduced	Good	0.05
7b	-8.9	403.88	6.14	75.5	4	0.13	0	4	1	703	Reduced Solubility	Good	0.05
7c	-8.7	448.34	6.2	75.5	4	0.13	0	4	1	570.42	Reduced Solubility	Good	0.05
7d	-9.0	383.47	5.88	75.5	4	0.13	0	4	1	916.81	Reduced Solubility	Good	0.09
7e	-8.8	399.46	5.48	84.73	5	0.16	0	5	1	1198.47	Reduced Solubility	Good	0.09
7f	-9.2	414.44	5.34	121.32	5	0.15	0	7	1	1217.01	Reduced Solubility	Good	0.05
7g 8a	- -9.1	- 398.5	- 6.61	- 107.52	4	- 0.13	- 0	- 4	- 1	- 533.37	- Reduced	- Good	- 0.05
8b	-9.7	432.95	7.24	107.52	4	0.13	0	4	1	314.71	Solubility Reduced	Good	0.05
8c	-8.7	477.4	7.3	107.52	4	0.13	0	4	1	253.66	Solubility Reduced	Good	0.05
8d	-9.4	412.53	6.98	107.52	4	0.13	0	4	1	411.89	Solubility Reduced	Good	0.09
8e	-9.0	428.53	6.58	116.75	5	0.16	0	5	1	536.91	Solubility Reduced	Good	0.09
8f	-8.5	443.5	6.44	153.34	5	0.15	0	7	1	543.88	Solubility Reduced	Good	0.05
8g	-9.7	466.53	6.61	137.73	4	0.11	0	6	1	401.48	Solubility Reduced	Good	0.04
Melphalan	-5.8	305.2	-0.69	71.01	8	0.53	3	4	0	125389.7	Solubility Good	Good	0.46
Levamisole	-6.0	204.29	1.84	40.9	1	0.06	0	2	0	19086.31	Solubility Good	Good	0.36
Co-ligand	-7.2	393.44	3.19	74.73	10	0.36	1	7	0	8316.35	Solubility Reduced	Good	0.27
											Solubility		

lar dynamics simulation study of the synthesized compounds led to the selection of 6 potent compounds (**7b**, **7c**, **7e**, **7f**, **7g**, and **8g**) with the good enzyme binding property. These in silico study was supported by in vitro results showed that compounds **7b**, **7c**, and **8g** were most cytotoxic among tested compouds against all three cell lines tested. How-

ever further in vivo and other biochemical study required to strengthen these compounds as molecules.

Declarations

Author contribution statement



Fig. 1. Structures of aminothiadiazole and imidazothiadiazoles as anticancer and cytotoxic agents.



Fig. 2. 3D interactions between the 1m17 receptor and compounds a) co-ligand, b) 5c, c) 5g, and d) 6g

CB: Performed the synthetic parts of experiment. DS: Carried out cytotoxicity study. HT and TK: Designed and performed the computational study. SK, SB & RS: performed the molecular dynamics simulation study. SSK: Designed the experiment, analyzed the spectral data, and compiled and wrote the paper.

Credit author statement

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CB: Performed the synthetic parts of experiment.

- D S: Carried out cytotoxicity study.
- HT and TK: Performed the docking computational study.

SK, SB and RS: Performed the Molecular Dynamics Simulation study SSK: Designed the experiment, analyzed the spectral data, analyzed the MDS study, compiled, and wrote the paper.

Declaration of Competing Interest

The authors declare that there is no conflict of interests regarding the publication of the paper.

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Supplementary materials

Supplementary material associated with this article can be found, in the online version, at doi:10.1016/j.molstruc.2021.130174.



Fig. 3. 2D interactions between 1m17 receptor and compounds a) co-ligand b) 5c, c) 5g, and d) 6g.

Table 4

Prominent molecular interactions towards 1m17 EGFR receptor domain.

Compound	Hydrogen bond	Hydrophobic Interactions	Others (Ionic, Water)
5a	Met769	Leu694, Ala719,	Leu694,
		Leu820, Met769	Cys773
5b	Met769	Leu694, Val702,	Leu694,
		Ala719, Leu820,	Cys773
E.	Mat760	Met/69	Mat760
50	Met/09	Leuo94, Ala/19, Mot742, Loui920	Met/69
54	_	Met/42, Letto20	Le11604
50		Met742, Leu820	Met769
5e	Met769	Leu694, Ala719,	Leu694.
		Lys721, Met742,	Lys704,
		Leu820	Pro770,
			Glu780
5f	-	Leu694, Phe699,	Gln767,
		Val702, Ala719,	Met769
		Met742, Leu820	
5g	Met769	Leu694, Val702,	Leu694
		Ala719, Leu820	
6a	-	Leu694, Ala719,	-
		Lys721, Leu768,	
		Leu820	
6b	-	Leu694, Val702,	-
		Leu/68, Leu820	
6c	-	Leu694, Phe699,	Leu694,
(1	M-+7(0	Val/02, Leu820	Met/69
60	Met/69	Leu694, Ala719,	Met/69, Dro 770
6.0		Met/42, Leu820	Pr0770
be	-	Leuo94, Vai/02,	Lys/21,
		Leu/08	GIU/38,
6f	Lvc721		Азрозт
01	Lys721 Leu694 Phe699	Lvs721 Thr766	
	Val702 Leu820	Gln767, Met769	4
	Lvs721	Asp831	
69	Met769		
-0	Leu694, Val702,	Leu694	
	Ala719, Phe771,		
	His781, Leu820		
7a	Met769	Leu694, Val702,	-
		Ala719, Leu820	
7b	Met769	Leu694, Val702,	Lys721,
		Ala719, Lys721,	Met769
		Leu820	
7c	Met769		
	Leu694, Val702,	-	
	Ala719, Leu764,		
	Leu768, Leu820		
7d	Met/69, Lsy/21	Leu694, Val702,	Lys/21,
		Ala/19, Leu820	Met/69,
7.	Mat760	Leu604 Ale710	Asp831
7e	Met/09	Leuo94, Ala/19,	Lys704, Mot760
		Lys/21, Met/42,	Pro770
7f	Met769	Val702 Ala719	Leu694
/1	Metros	Met769 Leu820	Leuost
7g	Thr830	Leu694, Ala719,	Lys721.
. 9		Lvs721, Met742,	Thr830.
		Leu768	Asp831
8a	Met769, Cys773	Leu694, Phe699,	Cys773,
		Leu820	Asp776
8b	Met769	Leu694, Ala719,	Cys773,
		Lys721, Met742,	Asp776
		Leu820	
8c	-	Leu694, Lys721,	Ser696,
		Leu768	Met769
8d	Met769, Cys773	Leu694, Ala719,	Pro770,
		Lys721, Met742,	Cys773,
		LCU02U	Asp//o

Compound	Hydrogen bond	Hydrophobic Interactions	Others (Ionic, Water)
8e	Met769	Leu694, Ala719,	Met769,
		Lys721, Met742	Lys704,
			Pro770,
			Asp776
8f	Met769	Leu694, Phe699,	Leu694,
		Leu820	Met769, sp776
8g	Met769, Thr830	Val702, Ala719,	Arg817,
Ū		Leu768, Leu820	Thr830,
			Asp831
			-



Fig. 4. Root mean square deviation (RMSD) plot of ten different ligands seen during Molecular Dynamics simulation (MD) run for the protein backbone (blue) and the ligands (red) 7b, 7c, 7e, 7f, 7g, and 8g.



Fig. 5. Protein ligand contact for the compounds 7b, 7c, 7e, 7f, 7g, and 8g.



Fig. 6. Protein ligand timeline for the compounds 7b, 7c, 7e, 7f, 7g, and 8g

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