



# Synthesis, Characterization, Cytotoxic Activity Studies of N1-phenylbenzene-1,2-diamine @CuhNFs and 1,2-phenylenediamine@CuhNFs, and Molecular Docking Calculations of Their Ligands

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## Abstract

In recent years, hybrid nanoflowers (hNFs), the newest class of nanoparticles, have been highly preferred due to their excellent activity and stability. In this study, hybrid nanoflower synthesis was carried out using N1-phenylbenzene-1,2-diamine and 1,2-phenylenediamine as the organic part and copper(II) metal ions as the inorganic part. In the first stage, the characterization of the synthesized hybrid nanoflowers was carried out using various techniques. For the characterization of the synthesized hNFs, structure elucidation was performed using Scanning Electron Microscopy (SEM), Energy Dispersive X-ray spectroscopy (EDX), Fourier transform infrared spectrometry (FT-IR), X-ray diffraction (XRD) spectroscopy and elemental mapping. In the other study stage, the cytotoxic effects of hybrid nanoflowers were evaluated using A549 and MCF7 cell lines. When 1,2-phenylenediamine and N1-phenylbenzene-1,2-diamine were converted into CuhNFs, it was effective in MCF7 and A549 cell lines. Docking studies were performed using the Prime MM-GBSA method to estimate binding affinities and determine the binding mode. ADME analysis was performed using the Schrödinger 2021-2 QikProp wizard. Support was obtained from molecular docking to confirm the potential of N1-phenylbenzene-1,2-diamine and 1,2-phenylenediamine compounds for both breast and lung cancer. Molecular docking studies can provide information about binding interactions between compounds with identified targets, which may explain their inhibitory activity. A better result can be obtained by examining the binding patterns in the active binding region of the compounds through molecular docking.

## Graphical Abstract

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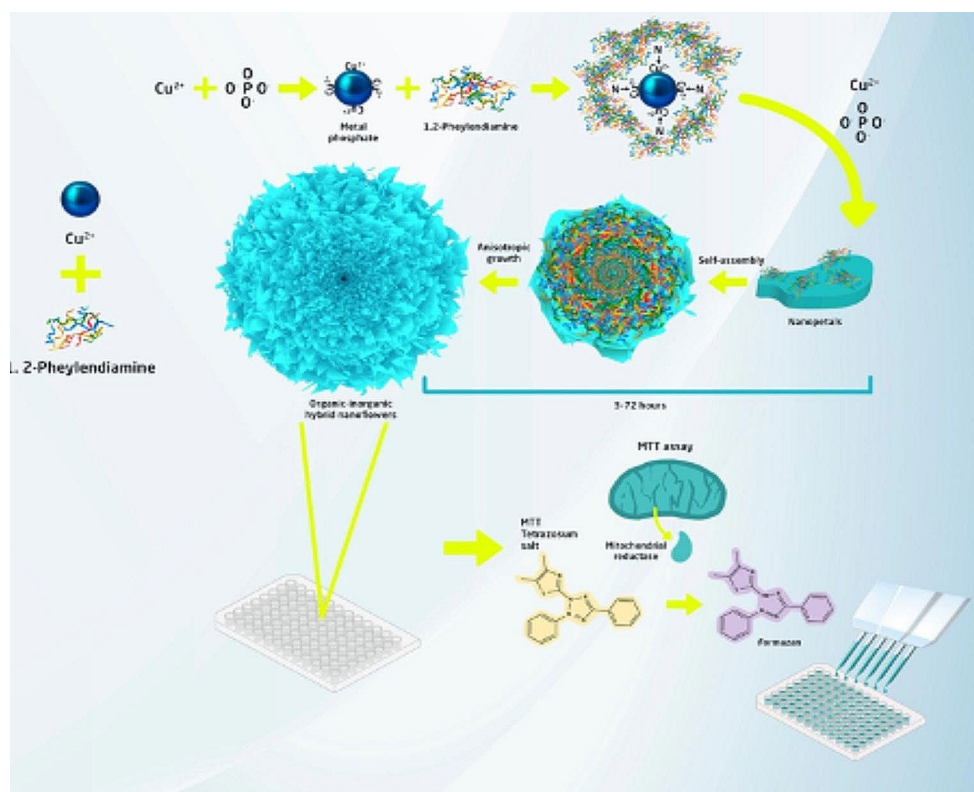
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**Keywords** ADME · Anticancer · Molecular docking · Nanoflower

## 1 Introduction

Nanotechnology is promising in fields like biocatalysis and the development of biosensors [1–5]. Functionalized nanostructured materials are used as carriers for nanobiocatalyst, which is a nanocarrier-enzyme composite material. These include organic-inorganic hybrids, mesoporous/nanoporous carriers, nanoparticles, magnetic nanoparticles, carbon nanotubes, and nanoflowers (HNFs) [6]. Nanoparticles attract much attention among many nanomaterials because they have many unique properties. The large surface area due to its morphology is just one of its unique features, which makes its activity and stability excellent [7–9].

Metal-containing nanoparticles are used in many scientific applications due to their small size and metal content [10–12]. Hybrid nanoflowers (hNFs), among the nanoparticles developed in recent years, have a flower-like structure and have many superior properties compared to other nanoparticles due to this morphology [13].

Hybrid nanoflowers were discovered accidentally by Zare and his group in 2012 through the self-assembling and co-precipitation of enzymes [14]. HNFs have been widely investigated since 2012 due to their good stability, enhanced activity, and easy and environmentally friendly preparation

[15]. Hybrid nanoflowers are flower-like three-dimensional (3D) nanostructures of organic and inorganic components [16]. Hybrid nanoflowers are flower-like, providing a large surface-to-volume ratio and excellent catalytic activity. Recent studies have also preferred it due to its high stability [17].

Various metal ions such as  $\text{Ca}^{2+}$ ,  $\text{Cu}^{2+}$ ,  $\text{Co}^{2+}$ ,  $\text{Fe}^{2+}$ ,  $\text{Mn}^{2+}$ , and  $\text{Zn}^{2+}$  are used as inorganic ingredients to form hybrid nanoflowers. Various biomolecules such as amino acids, DNA, enzymes, and organic molecules are preferred as organic ingredients [18–20].

There are many studies on hybrid nanoflowers today. For example, Güven et al. (2022) synthesized organic@inorganic hybrid copper nanoflowers (Cu NFs) using cherry stem for the first time. Cherry stem extract was obtained using ethanol and water solvents. Various techniques have been used to characterize Cu NFs by synthesis under different pH conditions. Cu NFs were revealed to have catalytic, antioxidant, and antimicrobial activities [21]. Koca and colleagues (2024) synthesized hNF using algae extract and Cu metal ion. They evaluated the effect of ambient pH and algal extract concentration on their morphology. Dye degradation and antioxidant activities of synthesized hNFs were determined under optimum conditions [22].

In the current study, for the synthesis of hNFs, 1,2-phenylenediamine, and N1-phenylbenzene-1,2-diamine were used as the organic part, while copper metal ion was used as the inorganic part. This was the first time hNFs were synthesized using the mentioned organic molecules. The characterization of 1,2-phenylenediamine@CuhNFs and N1-phenylbenzene-1,2-diamine@CuhNFs was conducted with FESEM, EDX, XRD, FT-IR, and elemental mapping. In addition, the cytotoxic effects of 1,2-phenylenediamine@CuhNFs and N1-phenylbenzene-1,2-diamine@CuhNFs were performed on A549 and MCF7 cells. The current study aims to create novel nanoflowers N1-phenylbenzene-1,2-diamine and 1,2-phenylenediamine compounds and analyze their activity on anticancer. Support was obtained from molecular docking to confirm the potential of these compounds for both breast and lung cancer. Molecular docking studies can provide information about binding interactions between compounds with identified targets, which may explain their inhibitory activity. A better result can be obtained by examining the binding patterns in the active binding site of the compounds through molecular docking.

## 2 Materials and methods

### 2.1 Chemicals and Reagents

$\text{CuSO}_4 \cdot 5\text{H}_2\text{O}$ ,  $\text{KH}_2\text{PO}_4$ ,  $\text{CaCl}_2 \cdot 2\text{H}_2\text{O}$ ,  $\text{MgCl}_2 \cdot 6\text{H}_2\text{O}$ , NaCl, KCl, and  $\text{Na}_2\text{HPO}_4$  were provided from Sigma-Aldrich. N1-phenylbenzene-1,2-diamine and 1,2-phenylenediamine were purchased from BLD pharm and Merck, respectively.

### 2.2 Synthesis of N1-phenylbenzene-1,2-diamine@CuhNFs and 1,2-phenylenediamine@CuhNFs

N1-phenylbenzene-1,2-diamine@CuhNFs and 1,2-phenylenediamine@CuhNFs were synthesized using a method reported by Somturk et al. [23–25]. Three hundred thirty-three microliters of copper sulfate solution and also 1 mL of organic molecules (0.02 mg/mL) are added into 50 mL of PBS buffer (pH:7.4). After the mixture is gently vortexed, it is incubated at +4 °C for three days. The mixture is centrifuged at 10,000 rpm for 30 min, and the pellet part is left to dry.

### 2.3 Characterization of N1-phenylbenzene-1,2-diamine@CuhNFs and 1,2-phenylenediamine@CuhNFs

Using the SEM device, it was realized that the synthesized N1-phenylbenzene-1,2-diamine@CuhNFs and 1,2-phenylenediamine@CuhNFs had a flower structure. Additionally,

the presence of metal ions in the hNFs was elucidated using EDX. While XRD was used to elucidate the crystal structures of hNFs, the FTIR spectrum was used to determine the presence of phosphorus-dependent oxygen peaks. The average size of hNFs was determined using the Image ProPlus 6.0 program.

### 2.4 Cytotoxicity Assessment

MTT ([3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide]) test was used to reveal the cytotoxic effects of hNFs. For this, A549 and MCF7 cell lines were used. After the cells were increased, hybrid nanoflowers synthesized in the 15,625–1000  $\mu\text{g}/\text{mL}$  concentration range were added to the wells and left for one night. Staining was performed, and cell viability was determined by measuring absorbance in a microplate reader. Statistical analysis was performed by one-way ANOVA.

### 2.5 Computational Studies

The results of the computational studies were obtained with the Schrödinger 2021-2 program [26]. The interactions of compounds N1-phenylbenzene-1,2-diamine and 1,2-phenylenediamine, which was determined to be effective as an anticancer according to in vitro results, on both EGFR and Human Estrogen Receptor were investigated via molecular docking. ADME properties were calculated to evaluate the pharmacokinetic behavior of the compounds.

#### 2.5.1 Molecular Docking Studies

Molecular docking analyses were performed to investigate the active interaction modes of the compounds 1,2-phenylenediamine and N1-phenylbenzene-1,2-diamine interacting with the crystal structures (PDB IDs: 1M17 [27], 3ERT [28]) obtained from the protein database. Targets important in cancer signaling pathways were identified for use in this study. Since the human lung cancer cell line and the human breast cancer cell line studied in vitro, targets are selected accordingly. In this study, EGFR was preferred for lung cancer, and estrogen receptor for breast cancer. In addition, in silico interactions of the designed compounds are essential because these targets prevent cancer cells from spreading throughout the body and metastasis. Ligand preparation, receptor preparation, and ligand-receptor in silico interaction were calculated following the procedure applied in previous studies [29–31].

Possible conformations of ligands are calculated with the Schrödinger 2021-2 LigPrep “Ligand preparation wizard” [32]. The crystal structure was prepared with the “Protein Preparation Wizard” of Schrödinger 2021-2 Protein

Preparation [33] software. OPLS\_2005 force field was used to perform optimization and minimization operations to prepare targets. While the prepared ligands and targets were labeled, their free binding energies were calculated with the MM/GBSA VSGB solvent model at the Schrödinger Prime wizard [34]. This analysis was applied to determine the free binding energies of the interaction results of 1,2-phenylenediamine and N1-phenylbenzene-1,2-diamine compounds with the target.

### 2.5.2 ADME Prediction

Absorption, distribution, metabolism, and excretion (ADME) properties were estimated with Schrödinger's QikProp [35] wizard. The compounds were analyzed according to ADME predictions using the QikProp [35] tool of the Schrödinger program. QikProp values of these compounds, such as molecular weight, dipole moment, SASA, FISA, PISA, QPolarize, QPlogPo/w, QPlogHERG, QPPCaco, QPlogBB, #metab, QPlogKhsa, human absorption percentage, PSA, violations of Lipinski rules (rule of five, rule of three) were calculated.

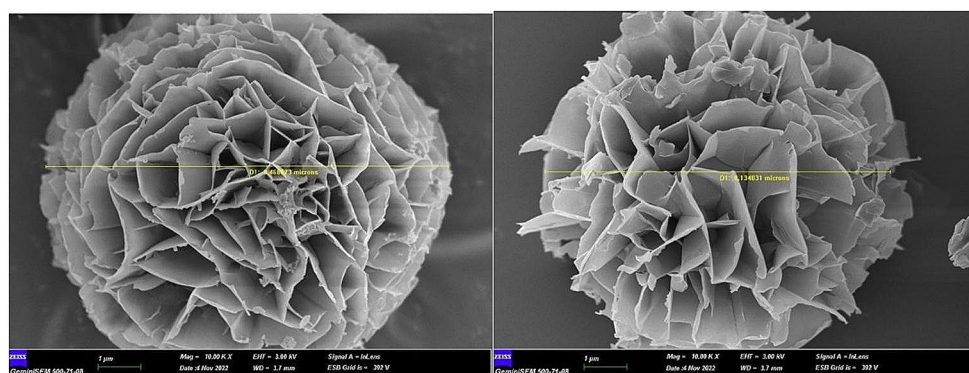
## 3 Results and Discussion

### 3.1 Characterization of hNFs

#### 3.1.1 SEM Analysis

The morphologies of the synthesized hNFs were determined by FESEM analysis, and the images were determined to have flower-like structures. The average dimensions for CuhNFs were  $\sim 8.13 \mu\text{m}$  and  $\sim 9.46 \mu\text{m}$ , respectively (Fig. 1). As seen in the figure, it has been proven that the structures have a flower-like structure. This flower-shaped morphology causes the surface area to expand, which increases its stability and activity.

**Fig. 1** SEM images of a) 1,2-phenylenediamine@CuhNFs (10.000 KX), b) N1-phenylbenzene-1,2-diamine@CuhNFs (10.000 KX)



#### 3.1.2 EDX Analysis

Elemental analysis of N1-phenylbenzene-1,2-diamine@CuhNFs and 1,2-phenylenediamine@CuhNFs were carried out with EDX. The EDX spectrum of CuhNFs is seen in Fig. 2. The presence of Cu metal ion in the copper phosphate nanocrystal can be seen in the figure.

#### 3.1.3 Elemental Mapping

It is seen in Figs. 3 and 4 that the metal ions contained in hNFs are distributed homogeneously.

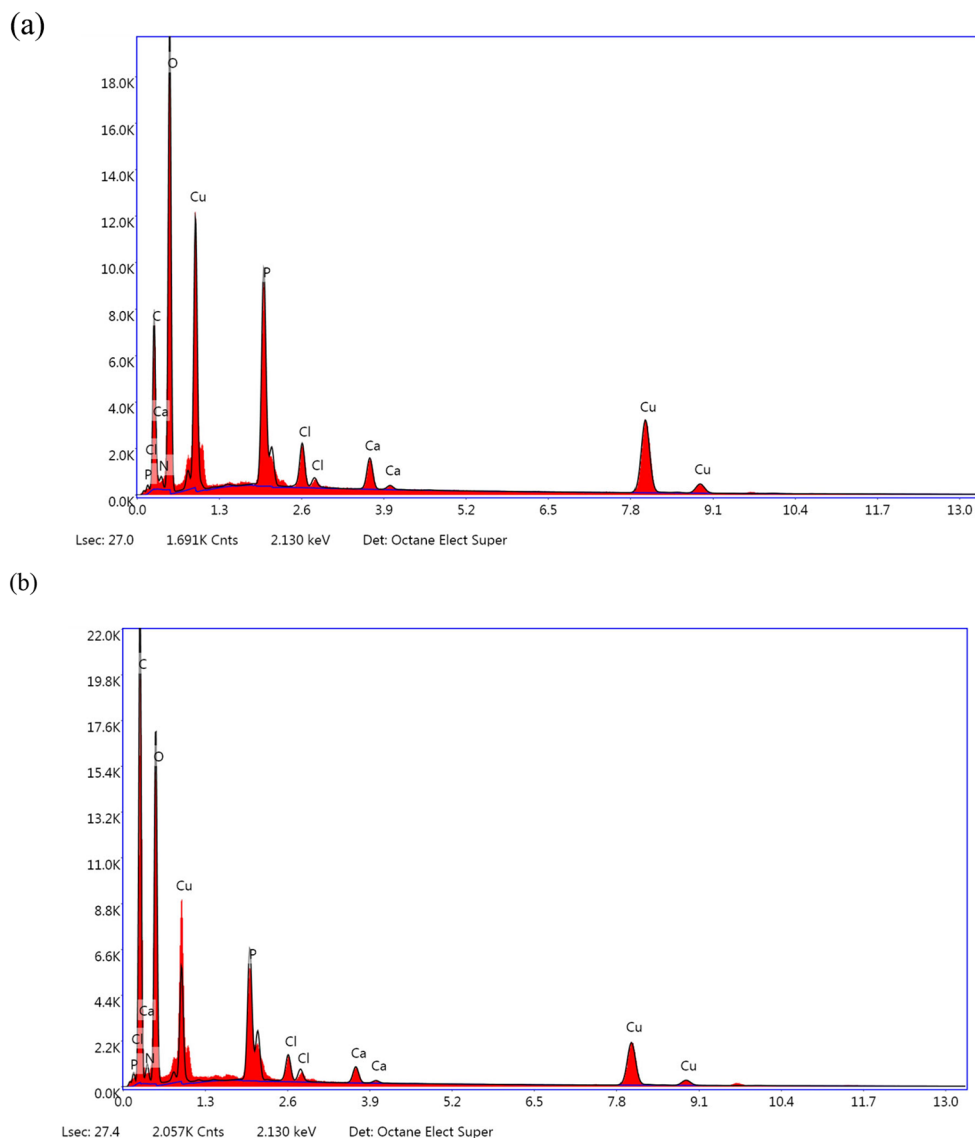
#### 3.1.4 XRD Analysis

The crystal structure of the 1,2-phenylenediamine@CuhNFs and N1-phenylbenzene-1,2-diamine@CuhNFs were carried out with XRD (Fig. 5). The XRD peaks were assigned as for 1,2-phenylenediamine@CuhNFs and N1-phenylbenzene-1,2-diamine@CuhNFs, XRD:  $8.93^\circ$ ,  $12.82^\circ$ ,  $18.47^\circ$ ,  $20.54^\circ$ ,  $22.78^\circ$ ,  $26.66^\circ$ ,  $29.56^\circ$ ,  $33.80^\circ$ ,  $35.31^\circ$ ,  $36.04^\circ$ ,  $37.28^\circ$ ,  $38.44^\circ$ ,  $41.98^\circ$ ,  $47.05^\circ$ ,  $53.55^\circ$ ,  $55.29^\circ$ ,  $57.17^\circ$ ,  $61.79^\circ$ ,  $65.19^\circ$ ,  $68.43^\circ$ ,  $72.68^\circ$  in Fig. 5 compared with JCPDS (00–022–0548).

#### 3.1.5 FT-IR Analysis

The chemical structure of 1,2-phenylenediamine@CuhNFs and N1-phenylbenzene-1,2-diamine@CuhNFs were illuminated using the FT-IR spectrum (Fig. 6). The spectrum data of 1,2-phenylenediamine@CuhNFs was seen as follows; For hNFs, FT-IR ( $\text{cm}^{-1}$ ): 3305 (N–H and O–H stretching), 2975 (Ar–H and C–H stretching), 2912 (C–H, stretching), 1645, 1535, 1420, 1250, 1045 (P=O), 1055, 994 (P–O), 970, 683 (O=P=O), 634, 566 (O=P=O) and N1-phenylbenzene-1,2-diamine@CuhNFs were seen as follows; For hNFs, FT-IR ( $\text{cm}^{-1}$ ): 3351 (N–H and O–H stretching), 2976 (Ar–H and C–H stretching), 2914 (C–H, stretching), 1631, 1542, 1430, 1350, 1059 (P=O), 1071, 994 (P–O), 952, 664(O=P=O), 637, 563 (O=P=O).

**Fig. 2** EDX analysis of (a) 1,2-phenylenediamine@CuhNFs, (b) N1-phenylbenzene-1,2-diamine@CuhNFs



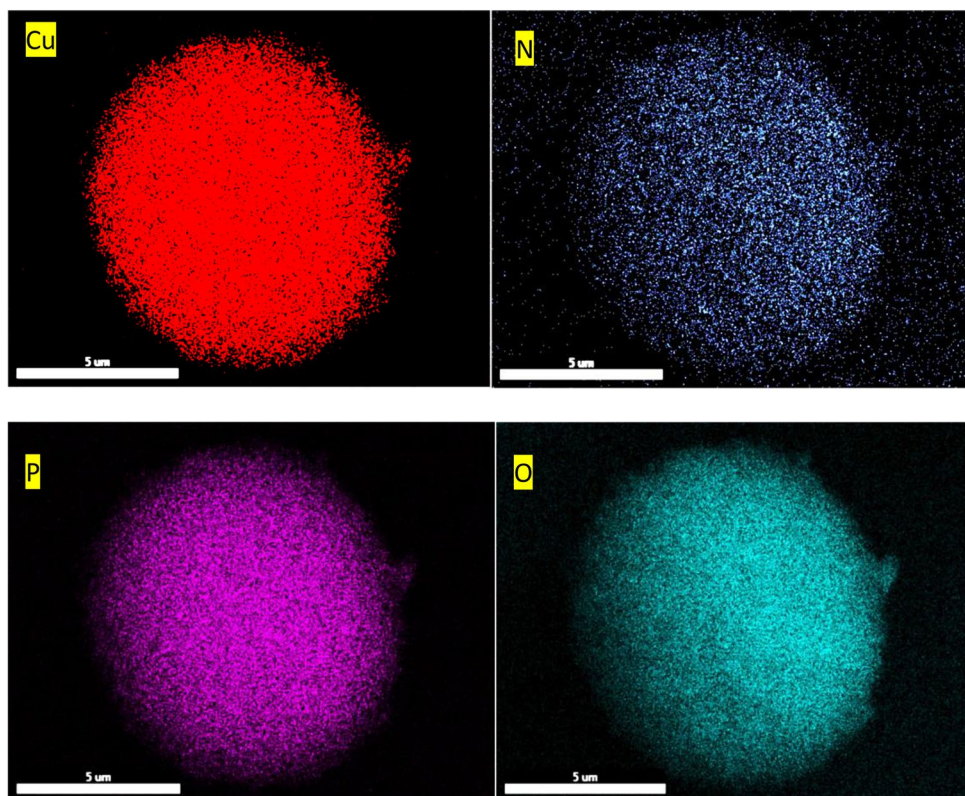
### 3.1.6 Cytotoxicity Assessment

The effect of ligands, 1,2-phenylenediamine@CuhNFs and N1-phenylbenzene-1,2-diamine@CuhNFs on the viability of A549 (lung) and MCF7 (breast) cancer cell lines was shown in Fig. 7. Different concentrations (15,625–1000  $\mu\text{g}/\text{mL}$ ) were applied to cells.

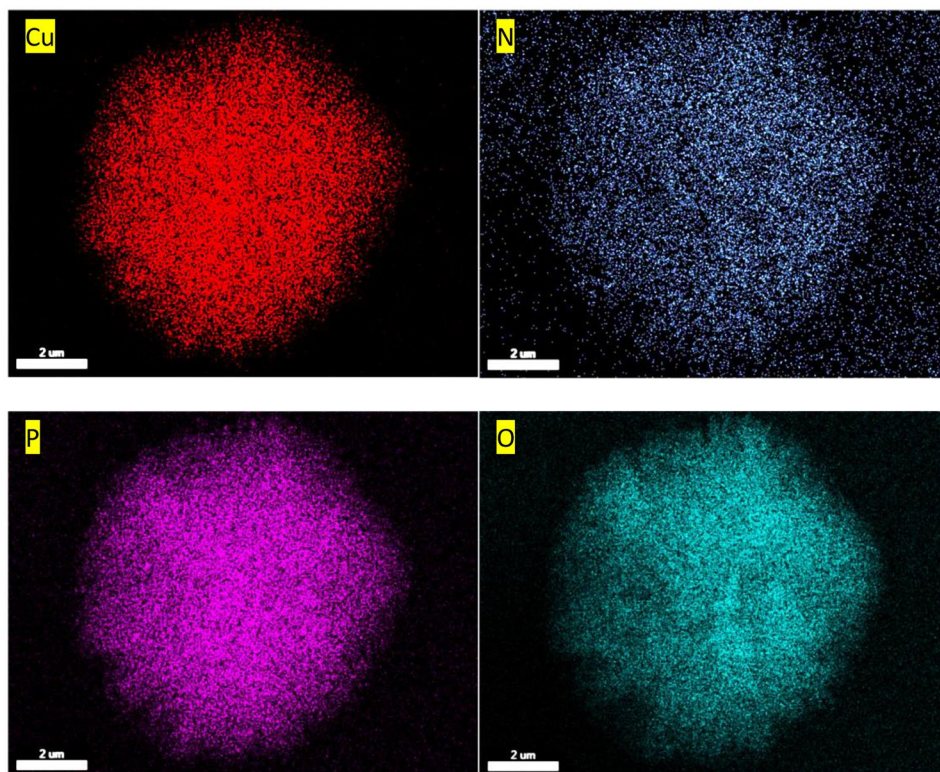
1,2-Phenylenediamine and 1,2-phenylenediamine@CuhNFs were tested in the A549 cell line. It was observed that the anticancer activity was better when the organic molecule was converted into a hybrid nanoflower. While the hybrid nanoflower caused 70% cell death, the organic molecule caused 40%. When the N1-phenylbenzene-1,2-diamine molecule A549 cell line was examined, it was determined that the hybrid nanoflower killed the cells by 90%, while the organic molecule killed by 60%. Although these two organic molecules were effective in the A549 cell line, it

was observed that their effectiveness increased even more when they were converted into a hybrid nanoflower form. When the 1,2-phenylenediamine molecule was examined in the MCF7 cell line, it was observed that its anticancer activity was better when the organic molecule was converted into a hybrid nanoflower. While the hybrid nanoflower caused 70% cell death, it was observed that organic molecules did not affect the cell line. When the N1-phenylbenzene-1,2-diamine molecule was examined in the MCF7 cell line, it was determined that the hybrid nanoflower killed the cells by 80%, while the organic molecule killed by 40%. Although these two organic molecules were effective in the MCF7 cell line, it was observed that their effectiveness increased even more when they were converted into a hybrid nanoflower form. There are many studies in the literature about hybrid nanoflowers. However, there are only a few studies on anticancer. One is a study conducted by Somturk Yilmaz

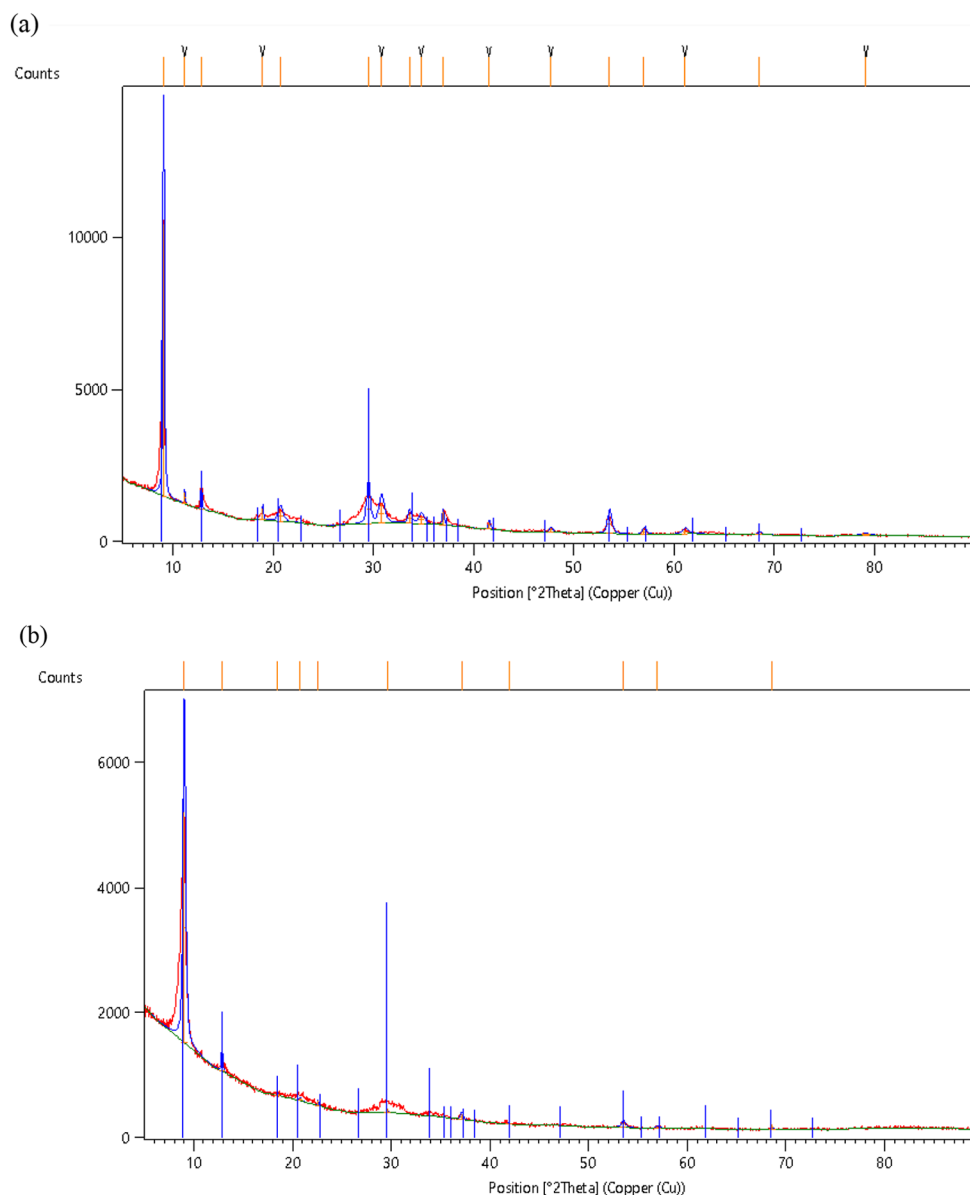
**Fig. 3** Elemental mapping (Cu, N, P, O) of 1,2-phenylenediamine@CuhNFs



**Fig. 4** Elemental mapping (Cu, N, P, O) of N1-phenylbenzene-1,2-diamine@CuhNFs



**Fig. 5** X-ray diffraction analysis of (a) 1,2-phenylenediamine@CuhNFs, (b) N1-phenylbenzene-1,2-diamine@CuhNFs.



and his colleagues in 2024. This study conducted a cytotoxic study on the A49 cell line with different metal ions using *Tribulus Terrestris L.* plant extract. They observed that its anticancer activity increased significantly when the extract was converted into a hybrid nanoflower form [36]. Recent studies have shown that, in addition to proteins and enzymes (containing amino groups), it has been proven in many studies that hybrid nanoflowers can be synthesized with plant extracts and organic molecules (containing carboxyl and hydroxyl groups).

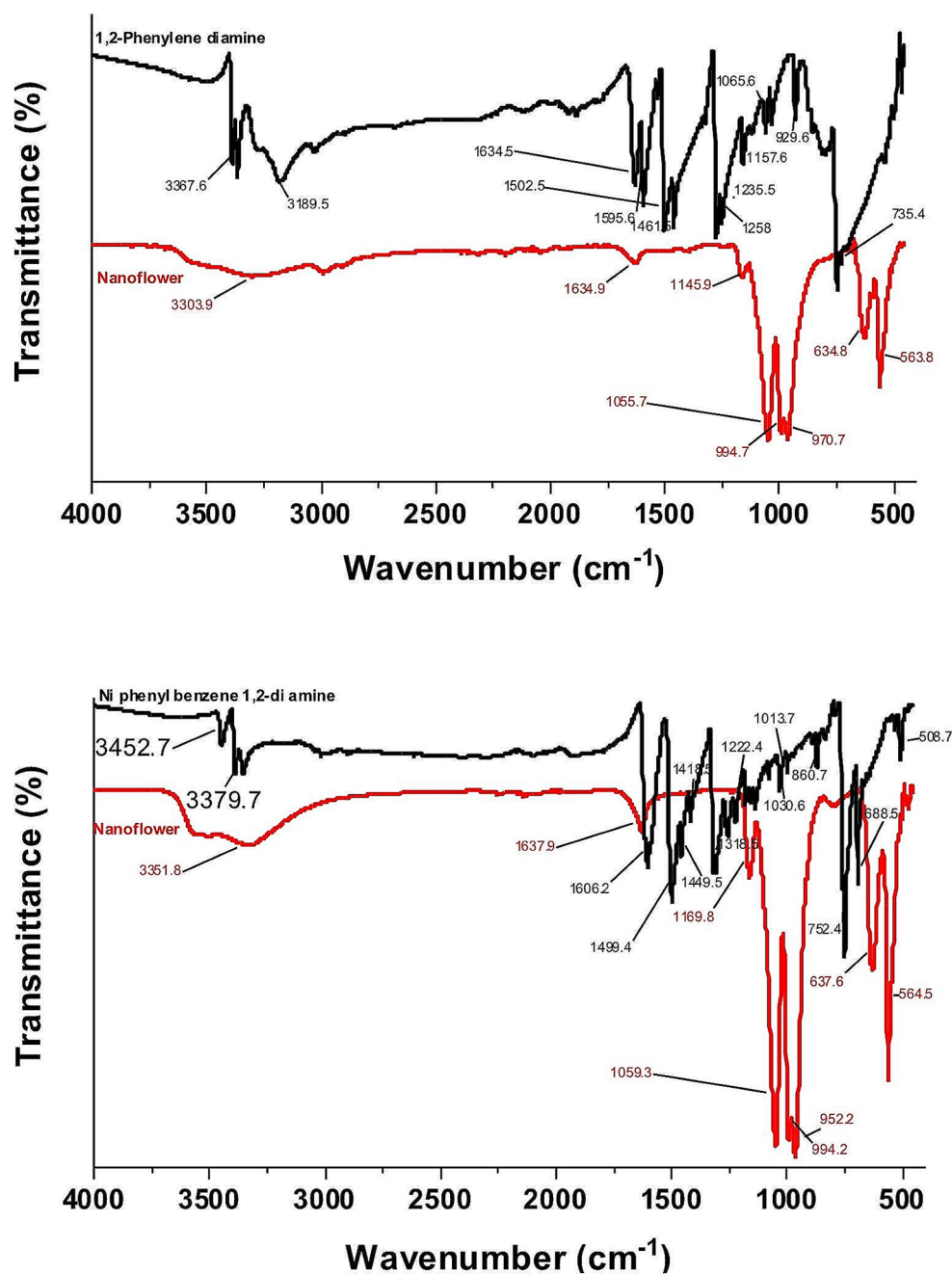
## 3.2 Computational Studies

### 3.2.1 Molecular Docking Studies

Molecular docking studies have been performed to investigate the ligand-protein binding pattern and orientations within binding sites on anticancer targets. However, both ligands docked to the active binding region of the crystal structures of both the breast cancer target (PDB ID:3ERT [28]) and the lung cancer target (PDB ID:1M17 [27]) obtained from the protein database. Table 1 presents the computational results of the interaction of these crystal structures with both ligands.

In silico approaches to support in vitro studies with computational methods, firstly, the results of the

**Fig. 6** FT-IR spectrum of 1,2-phenylenediamine@CuhNFs and N1-phenylbenzene-1,2-diamine@CuhNFs

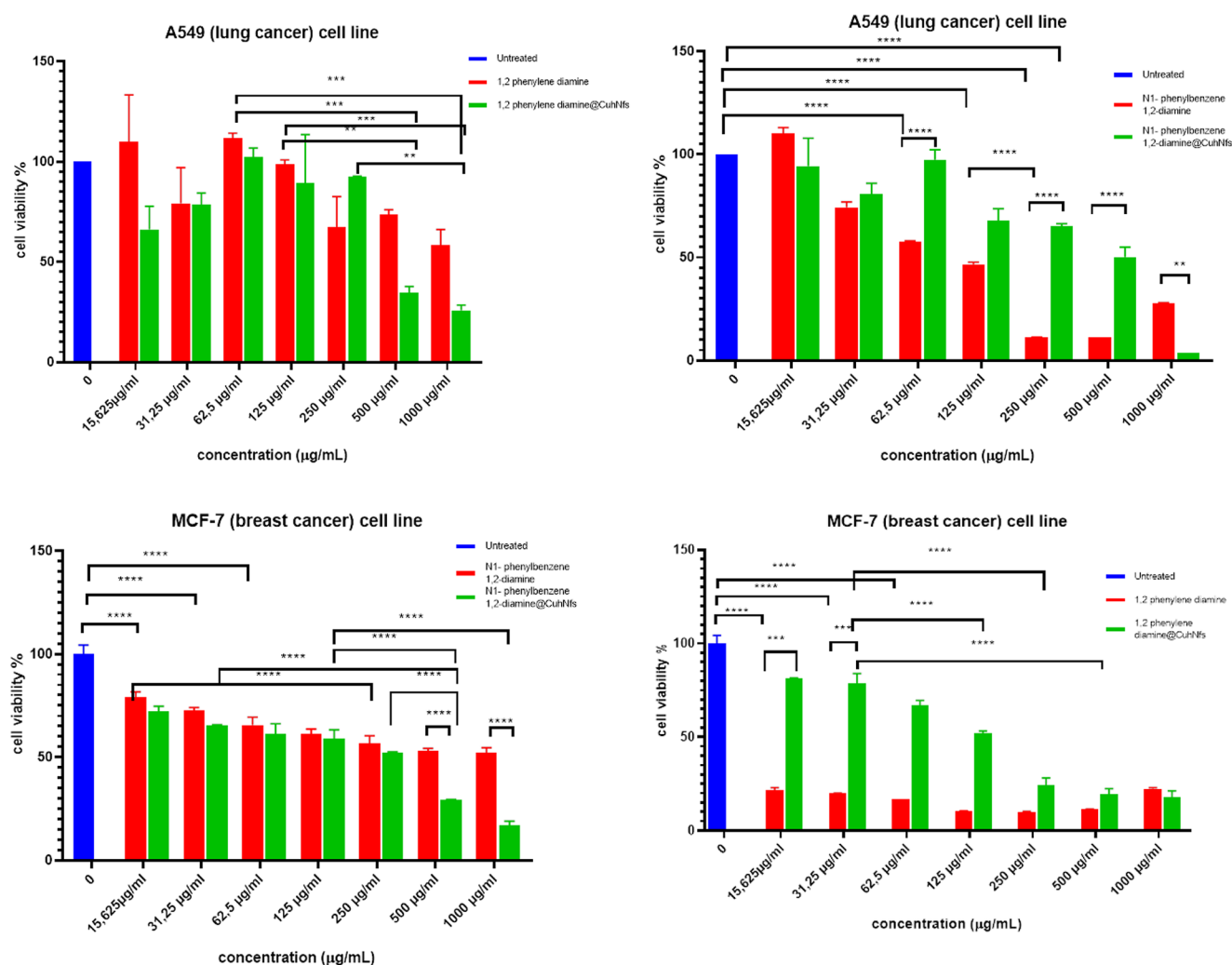


N1-phenylbenzene-1,2-diamine, and 1,2-phenylenediamine compounds interacting with the crystal structure of the human estrogen receptor (PDB ID: 3ERT) were analyzed. In Table 1, the docking score values of N1-phenylbenzene-1,2-diamine, and 1,2-phenylenediamine compounds were calculated as -7.304 kcal/mol and -4.157 kcal/mol, respectively. The result of these calculations was also visualized to analyze the binding mode. Table 1 also presents the result of the binding energy values of the second target, the Epidermal Growth Factor Receptor. Docking score values between EGFR and both ligands were calculated with the Schrödinger 2021-2 Glide program. The binding interaction

values between EGFR and compounds N1-phenylbenzene-1,2-diamine and 1,2-phenylenediamine were calculated as -4.220 kcal/mol, -3.596 kcal/mol, respectively.

Support was taken from Prime MM/GBSA analysis to investigate the relative binding affinity of the two ligands with both targets. Within the binding site of the 3ERT crystal structure, two compounds (N1-phenylbenzene-1,2-diamine, 1,2-phenylenediamine) showed good binding free energy values ( $\Delta G_{\text{Bind}}$ ) of -55.03 kcal/mol, -32.69 kcal/mol, respectively. The second target, calculating the binding free energy of the ligands with the 1M17 crystal structure, resulted in  $\Delta G_{\text{Bind}}$  values of -50.06 kcal/mol, -29.20 kcal/





**Fig. 7** Cytotoxicity of 1,2-phenylenediamine@CuhNFs and N1-phenylbenzene-1,2-diamine@CuhNFs on A549 and MCF7 cell lines, respectively. Data were presented as mean  $\pm$  SD. \*\*\* $p < 0.001$ , \*\*\*\* $p < 0.0001$

**Table 1** Docking scores of ligands and free energy calculation ( $\Delta G_{\text{Bind}}$ ) using Prime/MM-GBSA.

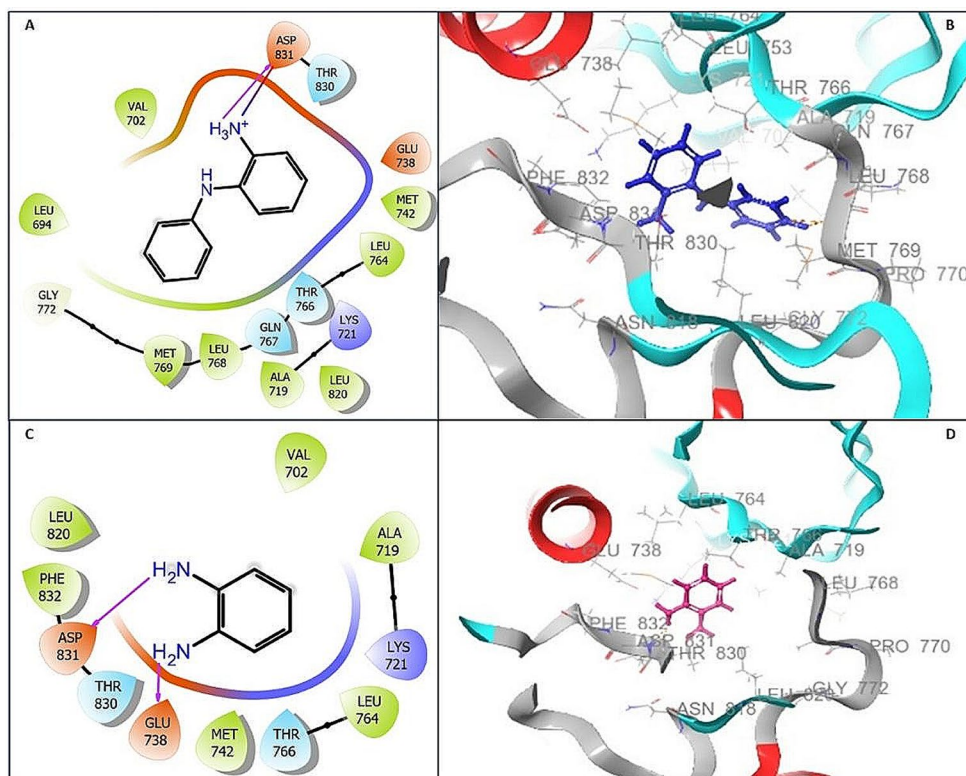
Ligands	PDB ID: 1M17		PDB ID: 3ERT	
	Docking score	$\Delta G_{\text{Bind}}$	Docking score	$\Delta G_{\text{Bind}}$
1,2-Phenyl-enediamine	-3.596 kcal/mol	-29.20 kcal/mol	-4.157 kcal/mol	-32.69 kcal/mol
N1-phenylbenzene-1,2-diamine	-4.220 kcal/mol	-50.06 kcal/mol	-7.304 kcal/mol	-55.03 kcal/mol

mol for N1-phenylbenzene-1,2-diamine, 1,2-phenylenediamine, respectively. All MM-GBSA calculations are summarized in Table 1. Both ligands' calculated binding affinity ( $\Delta G_{\text{Bind}}$ ) is different because they bind to their active sites with different amino acids while binding to both targets. As a result, docking score values and free binding energy values are consistent with biological values.

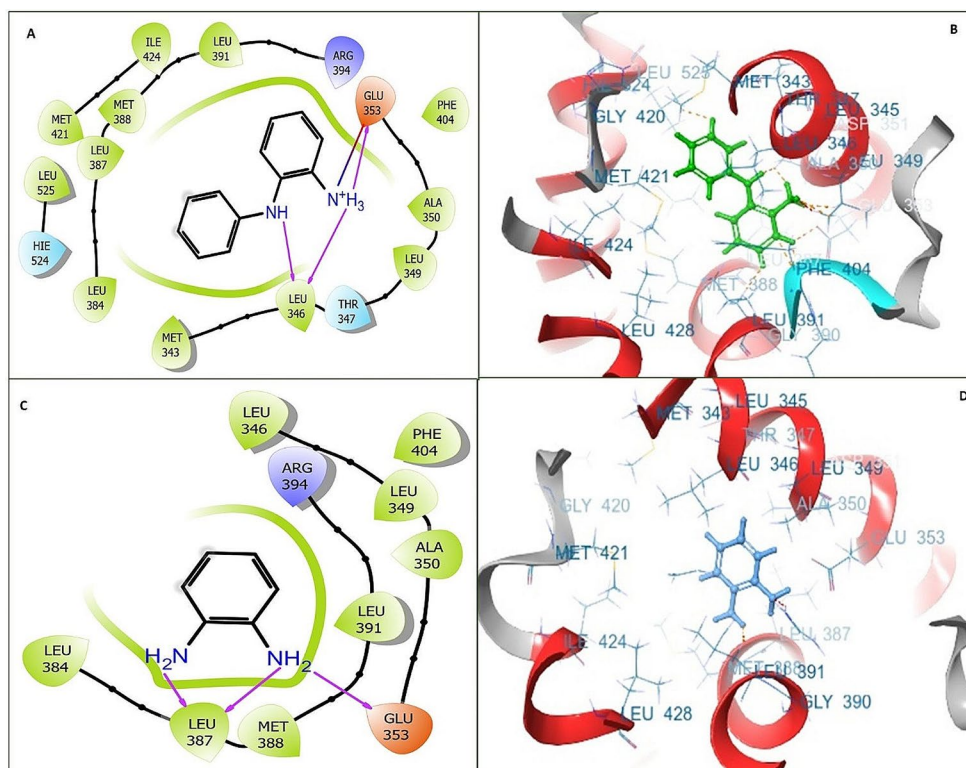
In Figs. 2D, 3D and 8 interaction diagrams of the interactions of N1-phenylbenzene-1,2-diamine, 1,2-phenylenediamine compounds with 1M17, the EGFR crystal structure, are presented. The 2D interaction diagram of the N1-phenylbenzene-1,2-diamine compound located in the active pocket region of the 1M17 crystal structure and the hydrogen bond interaction with Asp831, one of the essential amino acids is presented in Fig. 8(A). The 3D representation of the N1-phenylbenzene-1,2-diamine-1M17 complex is presented in Fig. 8(B). In the representation in Fig. 8(B), it is observed that critical amino acids surround the N1-phenylbenzene-1,2-diamine compound.

As a result of its in silico interaction with the EGFR target, the second ligand, 1,2-phenylenediamine, interacted with Asp831, a critical amino acid in this target, through hydrogen bonding, as mentioned before. The visual of this interaction is presented in Fig. 8(C). Figure 8(D) presents a 3D visual of the 1,2-phenylenediamine compound.

**Fig. 8** (A) 2D and (B) 3D docking pose displaying interaction of compound N1-phenylbenzene-1,2-diamine in the binding site of Epidermal Growth Factor Receptor (PDB ID: 1M17). (C) 2D and (D) 3D docking poses displaying the interaction of compound 1,2-phenylenediamine in the binding site of the Epidermal Growth Factor Receptor (PDB ID: 1M17)



**Fig. 9** (A) 2D and (B) 3D docking pose displaying interaction of N1-phenylbenzene-1,2-diamine in the binding site of Estrogen Receptor (PDB ID: 3ERT). (C) 2D and (D) 3D docking poses display the interaction of 1,2-phenylenediamine in the binding site of Estrogen Receptor (PDB ID: 3ERT).



The crystal structure of the target used as an estrogen receptor in breast cancer is presented due to the interaction of 3ERT and the ligands specified in Table 1. 2D and 3D interaction diagrams of ligands interacting with the 3ERT

target are presented in Fig. 9. It is presented in Fig. 9(A) that the expected hydrogen bond interaction of the N1-phenylbenzene-1,2-diamine compound with the amino acid residues Glu353 and Leu346, which are essential in this target,

occurs. It is understood from the 3D interaction in Fig. 9(B) that the N1-phenylbenzene-1,2-diamine compound can interact with the 3ERT crystal structure by settling in the active pocket site.

It is understood from the images in Fig. 9(C) and Fig. 9(D) that the 1,2-phenylenediamine compound binds to the estrogen receptor in the correct area, and the expected hydrogen bonds occur. The hydrogen bond interaction of the compound 1,2-phenylenediamine with the amino acid residues Leu387 and Glu353 caused the ligand to settle into the active binding region correctly.

When the molecular docking results were evaluated quantitatively and qualitatively, both ligands were analyzed in silico and in vitro. The binding parameter values presented in Table 1 are pretty striking. Moreover, Figs. 8 and 9 present this to us qualitatively.

### 3.2.2 ADME Profiling Studies

ADME calculation of compounds that may be drug candidates using in silico approaches is a frequently used prediction tool. ADME parameter values were calculated using Schrödinger 2021-2 Qikprop wizard in silico approaches.

**Table 2** In silico ADME profile of the N1-phenylbenzene-1,2-diamine and 1,2-phenylenediamine ligands

ADME Parameters	Ligands		Recommended values [35]
	1,2-phenylenediamine	N1-phenylbenzene-1,2-diamine	
Molecular Weight	108.143	184.20	130–725
Dipole	3.274	2.205	1.0–12.5
SASA	425.183	426.540	300.0–1000.0
FISA	64.766	66.224	7.0–330.0
PISA	360.417	360.316	0.0–450.0
QPolarize	22.687	22.798	130–70.0
QPlogPo/w	2.323	2.332	-2.0 to 6.5
QPlogH-ERG	-5.161	-5.167	Below -5
QPPCaco	2408.376	2332.922	<25 poor >500 great
QPlogBB	-0.172	-0.185	-3 to 1.2
#metab	6	6	1–8
QPlogKhsa	-0.149	-0.141	-1.5 to 1.5
Percent Human Absorption	100	100	>80% is high <25% is low
PSA	36.225	37.107	7–200
Rule of Five	0	0	<4
Rule of Three	0	0	<3

Determining the physicochemical and pharmacokinetic properties of ligands is an essential tool. Therefore, to check the drug-likeness of the N1-phenylbenzene-1,2-diamine and 1,2-phenylenediamine compounds, physicochemical properties were performed using Schrödinger 2021-2 Qikprop (Table 2).

The ADME profile and Lipinski Rule values of both ligands are presented in Table 2. It was determined that the ADME parameters of the compounds were within the acceptable range for human use. The values in Table 2 present the positive pharmacokinetic profile of the compounds N1-phenylbenzene-1,2-diamine and 1,2-phenylenediamine. The results are presented in Table 2. Lipinski's rule of five (Ro5) parameters are calculated to estimate the probability of drug similarity when deciding whether the synthesized compounds will be drug candidates. According to Lipinski's rule of three and five, all values are acceptable. Both ADME pharmacokinetics and Lipinski's Rule values indicate that this compound is expected to have drug-like properties and has the potential to be an orally active drug. Elangovan and colleagues (2023) synthesized and characterized the Schiff base from 3,4,5-trimethoxybenzaldehyde and sulfadiazine. A Swiss-ADME was used for a pharmacological study to determine whether the TMBDA molecule has pharmacological properties. The synthesized compound was found to show good antimicrobial activity compared to standard [37]. Elangovan et al. (2023) reported a Schiff base synthesized from vanillin and *o*-phenylenediamine. Electronic spectra, HOMO-LUMO and MESP, were performed with the gas and various solvent phases [38]. Elangovan and colleagues (2023) synthesized the (E)-3-(((9H-purin-6-yl)imino)methyl)phenol (A3H) molecule and characterized its structure with spectroscopic techniques. The molecule's structure was optimized using the B3LYP/cc-pVDZ basis set. The electronic properties of molecule was analyzed and they found that water has the lowest energy gap value compared to other solvents. Molecular docking techniques determined the binding interaction between the targeted proteins and the ligand [39].

## 4 Conclusion

To summarize, hNF synthesis was conducted using 1,2-phenylenediamine and N1-phenylbenzene-1,2-diamine molecules as the organic fraction and Cu (II) metal ion as the inorganic fractions. A549 and MCF7 cell lines were used to determine the cytotoxic effects of the synthesized hybrid nanoflowers. The compound N1-phenylbenzene-1,2-diamine plays a vital role in the localization of both targets in the binding pocket site according to findings from molecular docking studies. It causes a significant interaction

due to forming a critical hydrogen bond with ASp831 at the EGFR target. In the second target, the estrogen receptor, both ligands settled in the active binding site with the support of hydrogen bonding to interact with Glu353 and Leu346. Due to both ADME profile, docking score, and binding mode values, ligands are suitable for further development as compounds that could be anticancer drug candidates.

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**Data Availability** No datasets were generated or analysed during the current study.

## Declarations

**Ethics Approval and Consent to Participate** Not applicable.

**Consent for Publication** Not applicable.

**Competing Interests** The authors declare no competing interests.

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