



Anti-inflammatory effect of functionalized sulfasalazine boron nitride nanocages on cardiovascular disease and breast cancer: An in-silico simulation



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ABSTRACT

The objective of this research work is to investigate the ability of sulfasalazine (as an anti-cytokine drug) functionalized $B_{16}N_{16}$, $B_{15}GeN_{16}$ and $B_{15}SiN_{16}$ nanocages to treat inflammatory cardiovascular disease and breast cancer in comparison with the pure sulfasalazine (SSZ). Density functional theory (DFT) calculations at PBE1 functional were used to investigate the structural, electronic and spectral properties of sulfasalazine decorated $B_{16}N_{16}$, $B_{15}GeN_{16}$ and $B_{15}SiN_{16}$ nanocages. The most stable state was obtained on adsorption of SSZ over $B_{16}N_{16}$, $B_{15}GeN_{16}$ and $B_{15}SiN_{16}$ nanocages via its pyridine ring. Also, the adsorption of SSZ through SO_2 group over $B_{16}N_{16}$, $B_{15}GeN_{16}$ and $B_{15}SiN_{16}$ nanocages causes the lower binding energy and the increment of dipole moment as both factors can lead to increased sensitivity of the $B_{16}GeN_{16}$ nanocage to the drug. Molecular docking simulation illustrates that the interaction of SSZ via its pyridine ring with $B_{15}GeN_{16}$ gives the best binding affinity and inhibition potential of HER2 (human epidermal growth factor receptor 2) and TNF- α (tumor necrosis factor-alpha) whereas interaction of $-SO_2$ group with $B_{16}N_{16}$ gives the best binding affinity and inhibition potential of COX-2 (cyclooxygenase-2) and IL-1 (Interleukin-1) receptors. The predicted results demonstrated that $SSZ/B_{16}N_{16}$ and $SSZ/B_{15}GeN_{16}$ complexes can serve as a promising and preventive agent for inflammatory cardiovascular disease and breast cancer.

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Abbreviations: SSZ, Sulfasalazine; DFT, Density Functional theory; PBE, Perdew Burke Ernzerhof; HER2, Human epidermal growth factor receptor 2; TNF, α , tumor necrosis factor, alpha; COX, 2, cyclooxygenase, 2; IL, 1, Interleukin, 1; PCM, Polarisable continuum model; PDOS, Projected density of states; MPA, Mulliken population analysis; MEP, molecular electrostatic potential; FMO, Frontier molecular orbital; BSSE, Basis set superposition error; TDDFT, Time, dependent density functional theory.

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1. Introduction

Sulfasalazine, 5-[4-(2-pyridylsulfamoyl) phenylazo] salicylic acid (SSZ) is an Azulfidine drug which comes from the sulfa family drugs that is typically used in the treatments for rheumatoid arthritis, ulcerative colitis, and Crohn's disease [1-3]. SSZ is a co-drug formulation consisting of sulfapyridine which is a sulfa antibiotic that is linked with 5-aminosalicylic acid (5-ASA) by an azo-bond. The administered SSZ is often metabolised into the constituents by bacteria in colon where sulfapyridine is further

absorbed and metabolised while 5-ASA remains in the colon [4,5]. SSZ has poor pharmacokinetic behaviour, low oral bioavailability and many side effects. Thus, targeted SSZ administration with optimal amount is necessary for effective and safe SSZ-based treatments [6].

The last two decades have brought about a revolutionising advancement in the field of nanomaterials. Researchers and engineers have been actively involved in finding various applications of nanomaterials in the fields of medical, environment and electronics. The field of nanotechnology has garnered great interest ever since the famous lecture “There is plenty of room at the bottom” by Professor R. Feynman. As large numbers of organic as well as inorganic materials are readily available, the field of molecular electronics has become one of major field of research by many researchers [7-12]. The use of nanotechnology in development of safer and more effective therapeutic nanoparticles enabled researchers to use targeted medications with boron nitride (BN) nanostructures as drug carriers to control drug release in target tissue [13-16]. BN nanostructures are isoelectronic analogues of carbon nanostructures that have been recently investigated by various researchers due to their unique chemical and physical properties.

In contrast to carbon nanostructures which are made up of covalent bonded carbon atoms, BN nanostructures are ionic bonded comprising of negatively charged N and positively charged B atoms. The bonded boron and nitrogen atoms have empty *p* orbitals and lone pair electrons respectively which provide additional covalent interactions with external molecules [110]. Owing to their ionic and covalent characters, these structures tend to offer a unique set of physisorption and chemisorption properties for various external molecules as compared with carbon nanostructures [109]. The configuration of boron nitride nanostructures includes single-walled boron nitride nanotubes (SWBNNTs), multi-walled boron nitride nanotubes (MWBNTs) and boron nitride fullerenes [17-19]. BN nanostructures are hydrophobic, tuneable in size and shape, stable, biodegradable, and non-toxic. Due to this, they are potential candidates for various biomedical applications after specific functionalisation processes [20]. BN functionalisation involves doping or adsorbing certain elements on its surface to modify its chemical, optical, and electronic properties for proper interaction with specific external molecules. Various heteroatoms such as C, O, Si, Al, Au, Ca, Ga, Pt, etc., have been used for BN functionalisation to produce various tailored properties for applications such as catalysis, sensors, drug delivery, nanostructure fabrication and nanoelectronics [21,105,106]. It is also an excellent material for the storage of hydrogen gas [22].

Several cytotoxicity studies on BN nanostructures confirmed their low cytotoxicity and indicated the usage of BN as a potential drug delivery system [23]. In 2009, the suitability of BN nanotubes as bio-carrier for DNA oligomers owing to its non-toxic nature toward biological systems was reported by Chen *et al.* [24]. Celaya *et al.* evaluated the adsorption of melphalan anticancer drug through oxygen atom on B₁₂N₁₂, B₁₂C₆N₆ and B₆C₁₂N₆ cages for both vacuum and aqueous environment conditions by DFT calculations [25]. The obtained results revealed that the value of recovery time for melphalan desorption on the systems under study are suitable for drug delivery purposes. Recently, Guo and co-workers theoretically and experimentally reported the adsorption and detection of melphalan anti-ovarian cancer drug by the B₁₂N₁₂ and the doped XB₁₁N₁₂ nanostructures by using B3LYP and B3PW91 functionals [26]. It was found that GeB₁₁N₁₂ fullerene can be regarded as a suitable biosensor for detection of melphalan and as an acceptable drug delivery vehicle to decrease the metabolic activity of the OVCAR-3 cancer cells [107,108]. The high sensitivity of the electronic features of BN nanocages towards adsorption behaviour of various drugs such as amantadine [27],

curcumin [28], exemestane [29], ifosfamide [30], hydroxyurea [31], sarin and chlorosarin [32], sulfasalazine [33], thiotepa [34], etc., makes them as potential drug carriers for such drug molecules. B₁₆N₁₆ also interacts with glycine molecule through the amine group forming a stable structure thus denoting that this nanocage can be utilized for drug delivery and biomedical applications [35].

In this work, we aim to study the potential of B₁₆N₁₆ fullerene as drug delivery vehicle for sulfasalazine. For this aim, a series of computational studies were performed to investigate on the ionic and the covalent interactions of B₁₆N₁₆ fullerene with SSZ and the Si and Ge doping effects on such interactions via the use of B₁₅SiN₁₆ and B₁₅GeN₁₆ fullerenes. In addition, molecular docking was performed to find the theoretical binding energies as these calculations aid in determining the strength and extent of interaction between the macromolecules and the ligand. Docking is helpful in solving two purposes: Firstly, it describes the accurate ligand position with respect to the binding site; secondly it helps in effective scoring [36,37]. As molecular docking is effective in commenting about the interaction and binding energies of SSZ with the pro-inflammatory cytokines such as TNF-alpha, IL-1, and COX-2, etc [38,39]. Therefore, it has been employed in the current study to get detailed information about the drug's interaction and activity with the pro-inflammatory cytokines.

2. Methodology and Computational details

Geometry optimizations of B₁₆N₁₆, B₁₅GeN₁₆, B₁₅SiN₁₆, SSZ and their adsorbed systems in different orientations have been fully relaxed and determined by the PBE1PBE functional with 6-311G** basis set in Gaussian 09 package [40]. Gaussian is a computer-based simulation platform that uses the fundamentals of quantum mechanics to generate optimized structures of molecules, energies, infrared and raman spectra, NMR, MEP plots, HOMO, LUMO etc. From amongst the various functionals offered by Gaussian, PBE1PBE method was selected as in the previously published literature this functional has given good reliability and accurate results while calculating molecular binding energies of various drugs on BN nanostructures [41-44]. 6-311G** basis set has been chosen to come to an appropriate negotiation between accuracy and computational time and effort. Polarizable continuum model (PCM) was carried out to evaluate the role of solvent (water) and to simulate in vivo conditions during SSZ adsorption on B₁₆N₁₆ fullerenes [45]. Projected density of states (PDOS), Mulliken population analysis (MPA), molecular electrostatic potential (MEP) and frontier molecular orbital (FMO) calculations were performed by DFT formalism as implemented in Gaussian 09 package. Basis set superposition error (BSSE) for the binding energy was determined using counter poise correction method to obtain the BSSE energy (E_{BSSE}). Time-dependent density functional theory (TDDFT) calculations and thermodynamic parameters have also been performed by incorporating the PBE1PBE/6-311G** level of theory [46]. Adsorption energies (E_{ads}) of SSZ on the surfaces of B₁₆N₁₆, B₁₅GeN₁₆ and B₁₅SiN₁₆ nanocages were determined as follows:

$$E_{ads} = E_{SSZ/nanocage} - (E_{nanocage} + E_{SSZ}) + E_{BSSE} \quad (1)$$

where $E_{SSZ/nanocage}$, $E_{nanocage}$ and E_{SSZ} are the energies of adsorbed complex, nanocage and SSZ respectively.

Physicochemical properties of SSZ, nanocages and SSZ/nanocage complexes were predicted based on the following quantum molecular descriptors:

$$\mu = -\frac{1}{2}(I + A) \quad (2)$$

$$\chi = -\mu \quad (3)$$

$$\eta = \frac{1}{2}(I - A) \quad (4)$$

$$S = \frac{1}{2\eta} \quad (5)$$

$$\omega = \frac{\mu^2}{2\eta} \quad (6)$$

where μ is chemical potential, I is ionization potential, A is electron affinity, χ is electronegativity, η is global hardness, S is global softness and ω is electrophilicity index. The values of ionization potential and electron affinity were approximated as negative orbital energies of the highest occupied molecular orbital (HOMO), $-E_{HOMO}$, and the lowest unoccupied molecular orbital (LUMO), $-E_{LUMO}$ respectively based on Koopmans' approximation [47]. The fractional change of transferred electron, ΔN , from nanocage to SSZ was calculated based on the following equation:

$$\Delta N = \frac{\mu_{nanocage} - \mu_{SSZ}}{\eta_{SSZ} + \eta_{nanocage}} \quad (7)$$

Stability of adsorption complexes were determined based on the total change of stabilisation energies of adsorption complexes, $(\Delta E_{SE})_{SSZ/nanocage}$:

$$\begin{aligned} (\Delta E_{SE})_{SSZ/nanocage} &= \Delta E_{SSZ(nanocage)} + \Delta E_{nanocage(SSZ)} \\ &= \frac{-(\mu_{nanocage} - \mu_{SSZ})^2}{2(\eta_{SSZ} + \eta_{nanocage})} \end{aligned} \quad (8)$$

where $\Delta E_{SSZ(nanocage)} = \Delta N(\mu_{SSZ} + \frac{1}{2}\eta_{SSZ}\Delta N)$ and $\Delta E_{nanocage(SSZ)} = \Delta N(-\mu_{nanocage} + \frac{1}{2}\eta_{nanocage}\Delta N)$ are the respective energy changes of SSZ and nanocage due to adsorption [39].

The electron localisation functional plots were also determined. It was introduced for the first time by Becke and Edgecombe to comment on the regions in atoms and molecules where electrons are localized. An alternative approach was suggested by Savin et al. that justified the determination of ELF plots by utilizing DFT [48]. Ever since this interpretation, ELF plots have been widely studied to comment on the type of chemical bonds in molecules [49-51]. A similar approach has been utilized here to determine ELF. The projected density of states which is the relative contribution of a particular atom/orbital to the total DOS is also determined for all the structures under consideration.

In silico Molecular docking study was carried out using the Auto Dock software (4.2) [52]. The target proteins viz. TNF-alpha (PDB ID: 2AZ5), IL1A (PDB ID: 2L5X) and cyclooxygenase-2 (COX-2) (PDB ID: 1CX2), Human estrogen receptor alpha ligand (PDB ID: 2IOK) and HER2 kinase (PDB ID: 3RCD) were retrieved from protein data bank (pdb) database. The protein was prepared by elimination of water molecules and non-polar hydrogen atoms, adding Kollman atom partial charges and polar hydrogen atoms. A grid box of 60x60x60 with point spacing of 0.375 Å was set for creating the auto-grid module [53,54]. 150 GA runs were accomplished for each docking. Maestro 11.0 Schrodinger suit was applied for visualization of 2D and 3D presentations.

3. Results and discussion

3.1. Structural analysis of SSZ, B₁₆N₁₆, B₁₅SiN₁₆ and B₁₅GeN₁₆

The optimised molecular structures and geometries of SSZ are shown in Fig. 1 and the related data is tabulated in Table 2. With the change in environment from vacuum to aqueous, S-O₁ bond

length and C₃-S-O₁ bond angle of SSZ were slightly increased from 1.446 Å to 1.451 Å and 108.1° to 108.77° respectively. This could be attributed to the hydrogen bonding between lone pair electrons of O atom within S-O bond of SSZ with H atom of water molecules. Dipole moment of SSZ increased from 6.022 Debye to 8.213 Debye due to the protonation of its amine group and the deprotonation of its carboxyl group in forming zwitterionic form. However, its zwitterionic form appears to be more electrophilic than its neutral form as indicated by the larger values of χ and ω .

The optimised molecular structures of B₁₆N₁₆, B₁₅SiN₁₆ and B₁₅GeN₁₆ are shown in Fig. 2 while the data related to their geometries are shown in Table 1, 2 and 3. Owing to its centrosymmetry, B₁₆N₁₆ is isoelectric and hence has zero dipole moment. This can also be visualized in its uniform electron density distribution via MEP plot. It was found that the both the B₁₅GeN₁₆ and B₁₅SiN₁₆ nanocages showcase a local structural deformation around the Ge and Si sites, which is in agreement with the results obtained in previously published experimental and theoretical studies [55-60]. The newly formed Ge-N and Si-N bonds are about 1.893 Å and 1.774 Å in vacuum environment and 1.900 Å and 1.778 Å in solvent environment as the Ge and Si atoms are projected out of the nanocage surfaces [61]. The substitution of one of the B atoms of B₁₆N₁₆ with heavier atoms such as Si or Ge resulted in centrosymmetry thereby resulting in non-zero dipole moment. The degree of asymmetry increases as Si is replaced with a larger Ge atom thereby causing B₁₅GeN₁₆ to be the most polarised among the considered nanocage samples. B₁₅GeN₁₆ and B₁₅SiN₁₆ had smaller E_g and η ; larger ω as compared with the pristine nanocage. This indicates that Ge and Si doped BN nanocages are electronically more sensitive, chemically reactive and electrophilic, thus, making them a facile molecular carrier.

The transition from vacuum to aqueous environment resulted in the slight protrusion of Si and Ge atoms from their respective nanocages as indicated by the slight increase in Si-N and Ge-N bond lengths and the decrease in N-Si-N and N-Ge-N bond angles. The localised protrusions also altered the geometries of B₁₅SiN₁₆ and B₁₅GeN₁₆ by slightly increasing Si-N-B and Ge-N-B bond angles thus giving flatter B-N arrangements surrounding the Si or Ge. This will result in a more uneven electron density distribution which tends to localise at the vicinity of Si or Ge atoms resulting in further increase in dipole moment. The minor Si and Ge protrusions from nanocage also indicate their interaction with water molecules as Si and Ge are slightly reactive and oxidisable by water [62]. This interaction resulted in making the nanocage electronically more sensitive, chemically softer but less electrophilic as evidenced by the reduced values of E_g, η and ω in aqueous environment.

3.2. Adsorption of SSZ on B₁₆N₁₆

Adsorption of SSZ on B₁₆N₁₆ fullerene occurred through its sulfonyl, phenyl, pyridine and carboxyl groups to form four adsorption configurations: Complex A, B, C and D as shown in Fig. 3. In vacuum, Complex A has E_{ads} of -0.224 eV during SSZ adsorption via its sulfonyl group on B₁₆N₁₆ surface. The B₁-N₁ bond length of B₁₆N₁₆ decreases from 1.469 Å to 1.459 Å and B₃-N₂ bond length increases from 1.454 Å to 1.507 Å under the influence of SSZ drug. The results agree with calculated experimental results (B-N: 1.44 Å) [63]. Significant alternations in the bond angle of B₁₆N₁₆ are noticed with B₁-N₁-B₂ angle decreasing to 78.04°, N₁-B₁-N₂ angle increasing to 100° and B₃-N₁-B₁ angle reaching 119.87° with their original bond angles to be 78.20°, 99.51° and 114.45° respectively. Complex B also had small value of E_{ads} (-0.404 eV) during SSZ adsorption via its phenyl group on B₁₆N₁₆ surface. Both the B-N bond lengths of B₁₆N₁₆ increase under the influence of SSZ through sulfonyl group with the B₁-N₁-B₂ bond angle increasing to a value of 81.61°, N₁-B₁-N₂ angle decreasing to a value of

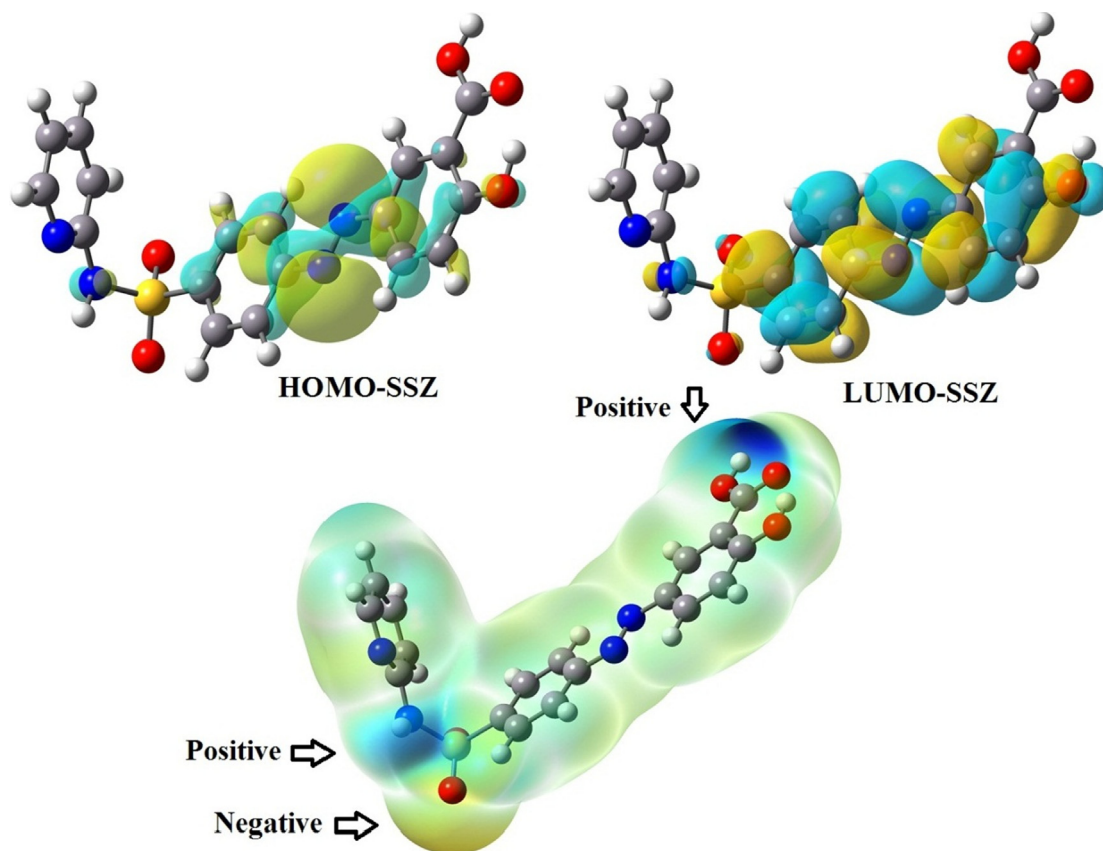


Fig. 1. Optimised structure and FMO plot of SSZ.

94.15° and the $B_3-N_1-B_1$ angles showcasing only a slight decrease of 0.40° from its original value. The adsorptions in Complex A and B though having small values of E_{ads} are chemisorptive since these interactions altered the electronic and the chemical properties of adsorption complexes as shown in the changed HOMO-LUMO energies and quantum molecular descriptors.

The complex C has the largest value of adsorption energy (-1.067 eV) among the four complexes thus implying the strongest interaction between SSZ via its pyridine ring and boron atom of $B_{16}N_{16}$ in this complex. In another study, the interaction of amantadine drug via its N atom with the B atom of $B_{12}N_{12}$ nanocage also had a strong adsorption energy of about -1.93 eV [18]. In our previous research, celecoxib through pyrazole ring had mediocre adsorption energy (-0.95 eV) with the $B_{12}N_{12}$ nanocage in comparison with its sulfonyl (-1.04 eV) and amine (-1.31 eV) groups [64]. As a result of the strong adsorption, B-N bond lengths of nanocage were increased significantly in complex C to values of 1.553 Å and 1.456 Å. The $B_1-N_1-B_2$ and $B_3-N_1-B_1$ bond angles showcase a hike in values to 82.78° and 114.51°. $N_1-B_1-N_2$ angle on the other hand decreases to 92.33°.

Theoretical FT-IR spectrum reveals that the -N-H, -C=O, -N=N, -B-N, -C=C- and O=S=O bonds in the most stable complex (complex C) lead to strong peaks at 3224, 1780, 1579, 1462, 1444, and 1380 cm^{-1} respective while theoretical peaks in the pure drug appeared at 3593, 1781, 1582, 1451, and 1372 cm^{-1} which are close to previous experimental researches [65,66]. Theoretical FT-IR spectrum of the pure $B_{16}N_{16}$ exhibited peaks as B-N at 1411 cm^{-1} and B-N-B at 805 cm^{-1} which is similar with the calculated results by experimental FTIR spectrum [67]. In complex D, SSZ is adsorbed via its carboxyl group on the boron atom of $B_{16}N_{16}$ nanocage with a binding energy of -0.694 eV. The B-N bond lengths show significant variations (B_1-N_1 becomes 1.529 Å and B_3-N_2 becomes 1.452 Å). Both the $B_1-N_1-B_2$ and $B_3-N_1-B_1$ angles

increase significantly. On the contrary $N_1-B_1-N_2$ showcases a sharp decrease under the influence of SSZ due to SSZ chemisorption on the surface of $B_{16}N_{16}$. The larger adsorption energies of complex B, C and D are generally more stable due to nanocage interaction with the delocalised electrons of phenyl, pyridine, and carbonyl groups while S in sulfonyl group is a strong electron withdrawing group thus causing weak interaction between sulfonyl group and B in nanocage.

The presence of polar bonds between the SSZ and the fullerene illustrates notable increments in the DM values because of the strong polarization of the resulting adsorption systems [61]. In the solvent medium, further increment of DM and polarisation in all the complexes was noticed [53,68]. Polarisation of pure SSZ was obvious in aqueous environment in which the electronic properties of its four adsorption groups were altered. In aqueous environment, E_{ads} of complex A decreased to -0.187 eV. The B-N bond lengths show a reverse trend in the solvent environment when compared to vacuum environment. The bond angles of $B_{16}N_{16}$ follow similar trend in both the environments. Similar trend was also reported for complex B where its adsorption energy decreases to -0.392 eV in solvent medium. The B-N bond length behaves in a similar manner in both the mediums. The $B_1-N_1-B_2$ and $B_3-N_1-B_1$ angles increase to 81.93° and 119.94° respectively while $N_1-B_1-N_2$ angle decreases to 93.53°. In contrast, complex C and D had their adsorption energies increased to -1.079 eV and -0.698 eV respectively. Both the B-N bond lengths of $B_{16}N_{16}$ fullerene increase while the bond angles show same trend in both the environments.

3.3. Adsorption of SSZ on $B_{15}SiN_{16}$ and $B_{15}GeN_{16}$

Fig. 4 shows two stable adsorption configurations of SSZ on $B_{15}GeN_{16}$ and $B_{15}SiN_{16}$ via pyridine and sulfonyl groups. SSZ inter-

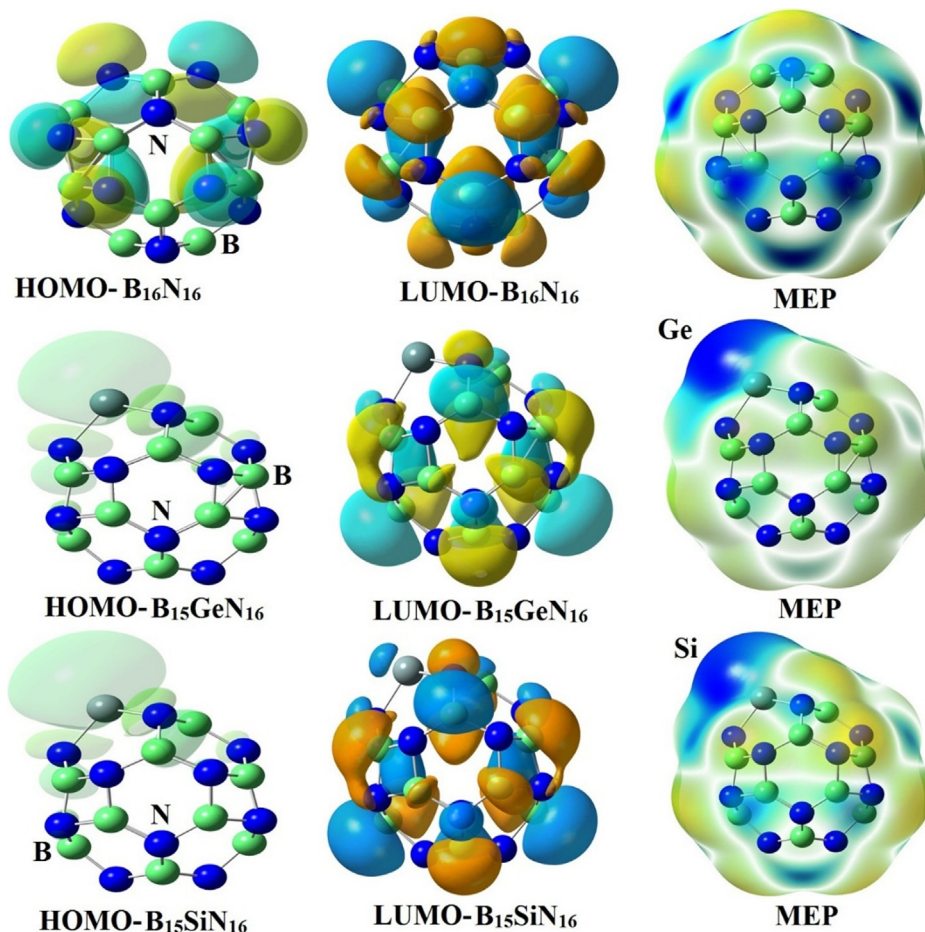


Fig. 2. Optimised structures, FMO and MEP plots of $B_{16}N_{16}$, $B_{15}GeN_{16}$ and $B_{15}SiN_{16}$ fullerenes.

Table 1

Computational results for SSZ- $B_{16}N_{16}$ adsorption complexes in vacuum and aqueous environments using PBE1PBE/6-311G** method.

Property	Vacuum					Aqueous				
	$B_{16}N_{16}$	A	B	C	D	$B_{16}N_{16}$	A	B	C	D
B_1-N_1 (Å)	1.469	1.459	1.533	1.553	1.529	1.470	1.456	1.542	1.560	1.538
B_3-N_2 (Å)	1.454	1.507	1.508	1.456	1.452	1.454	1.516	1.457	1.456	1.456
C_1-N_3 (Å)	-	1.326	1.325	1.356	1.328	-	1.326	1.325	1.355	1.329
$S-O_1$ (Å)	-	1.487	1.496	1.441	1.447	-	1.498	1.507	1.448	1.451
C_2-O_2 (Å)	-	1.222	1.222	1.222	1.270	-	1.224	1.224	1.224	1.272
$B_1-N_1-B_2$ (°)	78.20	78.04	81.61	82.78	82.07	78.25	78.08	81.93	82.94	82.33
$N_1-B_1-N_2$ (°)	99.51	100.0	94.15	92.33	94.04	99.44	99.97	93.53	91.88	93.42
$B_3-N_1-B_1$ (°)	114.45	119.87	114.05	114.51	121.21	114.36	120.13	119.94	121.63	121.17
C_3-S-O_1 (°)	-	107.83	108.30	108.42	107.50	-	107.64	107.58	108.78	107.92
$O_2-C_2-O_3$ (°)	-	121.46	121.47	121.36	121.82	-	122.11	122.11	122.08	122.13
$N_3-C_1-N_4$ (°)	-	112.95	112.99	117.04	113.41	-	113.63	113.60	117.09	113.40
Diameter (Å)	4.917	4.921	5.14	5.22	5.19	4.918	4.921	5.17	5.24	5.20
D (Å)	-	1.742	1.659	1.656	1.590	-	1.679	1.616	1.642	1.572
E_{ads} (eV)	-	-0.224	-0.404	-1.067	-0.694	-	-0.187	-0.392	-1.079	-0.698
E_{HOMO} (eV)	-7.14	-6.97	-6.92	-6.94	-7.14	-7.84	-6.82	-6.82	-6.79	-6.98
E_{LUMO} (eV)	-0.32	-3.06	-3.13	-2.99	-3.18	-0.95	-2.93	-2.96	-2.87	-2.92
E_g (eV)	6.82	3.91	3.79	3.95	3.96	6.89	3.89	3.86	3.92	4.06
ΔE_g (%)	-	42.67	44.43	42.08	41.93	-	43.64	43.98	43.11	41.07
DM (Debye)	0.0	9.89	11.31	9.63	3.63	0.0	13.12	14.65	13.61	4.77
E_F (eV)	-4.48	-5.02	-5.03	-4.97	-5.16	-4.40	-4.88	-4.89	-4.83	-4.95
I (eV)	7.14	6.97	6.92	6.94	7.14	7.84	6.82	6.82	6.79	6.98
A (eV)	0.32	3.06	3.13	2.99	3.18	0.95	2.93	2.96	2.87	2.92
μ (eV)	-3.73	-5.02	-5.03	-4.97	-5.16	-4.40	-4.88	-4.89	-4.83	-4.95
χ (eV)	3.73	5.02	5.03	4.97	5.16	4.40	4.88	4.89	4.83	4.95
η (eV)	3.41	1.96	1.90	1.98	1.98	3.45	1.95	1.93	1.96	2.03
S (eV^{-1})	0.15	0.26	0.26	0.25	0.25	0.15	0.26	0.26	0.26	0.25
ω (eV)	2.04	6.43	6.66	6.24	6.72	2.80	6.11	6.19	5.95	6.04

Table 2Computational results for SSZ-B₁₅GeN₁₆ adsorption complexes in vacuum and aqueous environments using PBE1PBE/6-311G** method.

Property	Vacuum				Aqueous			
	B ₁₅ GeN ₁₆	SSZ	E	F	B ₁₅ GeN ₁₆	SSZ	E	F
B ₁ -N ₁ (Å)	1.469	-	1.472	1.480	1.468	-	1.473	1.482
Ge-N ₂ (Å)	1.893	-	1.872	1.845	1.900	-	1.872	1.846
Ge-N ₃ (Å)	1.881	-	1.858	1.838	1.886	-	1.858	1.831
C ₁ -N ₃ (Å)	-	1.329	1.406	1.328	-	1.329	1.407	1.327
S-O ₁ (Å)	-	1.446	1.449	1.603	-	1.451	1.454	1.582
N ₁ -B ₁ -N ₂ (°)	109.49	-	108.03	107.82	109.51	-	107.99	107.74
Ge-N ₂ -B ₁ (°)	85.75	-	86.09	85.44	85.94	-	86.12	85.38
N ₃ -Ge-N ₂ (°)	105.11	-	106.70	109.59	104.60	-	106.69	110.45
C ₃ -S-O ₁ (°)	-	108.10	109.83	114.30	-	108.77	109.67	106.11
N ₃ -C ₁ -N ₄ (°)	-	113.47	111.43	114.21	-	113.48	112.09	114.50
Diameter (Å)	4.932	-	4.928	4.935	4.935	-	4.929	4.936
D (Å)	-	-	1.826	1.808	-	-	1.829	1.818
E _{ads} (eV)	-	-	-0.357	-0.038	-	-	-0.233	-0.030
E _{HOMO} (eV)	-7.44	-6.77	-4.27	-4.27	-7.22	-6.75	-4.20	-4.43
E _{LUMO} (eV)	-0.90	-2.78	-2.83	-1.68	-0.80	-2.80	-2.82	-1.69
E _g (eV)	6.54	3.99	1.44	2.59	6.42	3.95	1.38	1.61
ΔE _g (%)	-	-	77.98	60.40	-	-	78.50	74.92
DM (Debye)	1.766	6.022	4.620	7.668	3.016	8.213	6.024	15.111
E _F (eV)	-4.17	-4.78	-3.55	-2.98	-4.01	-4.78	-3.51	-3.06
I (eV)	7.44	6.77	4.27	4.27	7.22	6.75	4.20	4.43
A (eV)	0.90	2.78	2.83	1.68	0.80	2.80	2.82	1.69
μ (eV)	-4.17	-4.78	-3.55	-2.98	-4.01	-4.78	-3.51	-3.06
χ (eV)	4.17	4.78	3.55	2.98	4.01	4.78	3.51	3.06
η (eV)	3.27	2.00	0.72	1.30	3.21	1.98	0.69	1.37
S (eV ⁻¹)	0.15	0.25	0.69	0.39	0.16	0.25	0.72	0.36
ω (eV)	2.66	5.71	8.75	3.42	2.50	5.77	8.93	3.42

Table 3Computational results for SSZ-B₁₅SiN₁₆ adsorption complexes in vacuum and aqueous environments using PBE1PBE/6-311G** method.

Property	Vacuum				Aqueous			
	B ₁₅ SiN ₁₆	SSZ	G	H	B ₁₅ SiN ₁₆	SSZ	G	H
B ₁ -N ₁ (Å)	1.470	-	1.473	1.478	1.470	-	1.0.473	1.481
Si-N ₂ (Å)	1.774	-	1.761	1.734	1.778	-	1.760	1.736
Si-N ₃ (Å)	1.772	-	1.759	1.738	1.773	-	1.757	1.734
C ₁ -N ₃ (Å)	-	1.329	1.410	1.328	-	1.329	1.410	1.326
S-O ₁ (Å)	-	1.446	1.449	1.608	-	1.451	1.454	1.588
N ₁ -B ₁ -N ₂ (°)	106.74	-	105.91	105.66	106.79	-	105.85	105.54
Si-N ₂ -B ₁ (°)	84.71	-	84.81	84.21	84.84	-	84.86	84.11
N ₃ -Si-N ₂ (°)	107.64	-	108.50	111.03	107.32	-	108.66	111.63
C ₃ -S-O ₁ (°)	-	108.10	109.59	104.93	-	108.77	109.80	105.96
N ₃ -C ₁ -N ₄ (°)	-	113.47	111.73	113.95	-	113.48	112.09	114.63
Diameter (Å)	4.916	-	4.913	4.918	4.918	-	4.916	4.918
D (Å)	-	-	1.733	1.673	-	-	1.735	1.684
E _{ads} (eV)	-	-	-1.274	-0.602	-	-	-1.174	-0.663
E _{HOMO} (eV)	-7.01	-6.77	-4.29	-4.27	-6.81	-6.75	-4.26	-4.42
E _{LUMO} (eV)	-0.96	-2.78	-2.81	-1.63	-0.85	-2.80	-2.81	-1.69
E _g (eV)	6.05	3.99	1.48	2.64	5.96	3.95	1.45	2.73
ΔE _g (%)	-	-	75.54	56.36	-	-	75.67	54.19
DM (Debye)	0.945	6.022	4.373	7.654	1.653	8.213	6.016	15.12
E _F (eV)	-3.99	-4.78	-3.55	-2.95	-3.83	-4.78	-3.54	-3.06
I (eV)	7.01	6.77	4.29	4.27	6.81	6.75	4.26	4.42
A (eV)	0.96	2.78	2.81	1.63	0.85	2.80	2.81	1.69
μ (eV)	-3.99	-4.78	-3.55	-2.95	-3.83	-4.78	-3.54	-3.06
χ (eV)	3.99	4.78	3.55	2.95	3.83	4.78	3.54	3.06
η (eV)	3.03	2.00	0.74	1.32	2.98	1.98	0.73	1.37
S (eV ⁻¹)	0.17	0.25	0.68	0.38	0.17	0.25	0.69	0.37
ω (eV)	2.62	5.71	8.52	3.30	2.46	5.77	8.62	3.42

action on both the nanocages are chemisorptive as indicated by the partial overlapping of N orbital from pyridine group of SSZ with the orbitals of Ge and Si in the doped nanocages as visualized in the ELF plots (Fig. 4). Adsorption energies obtained in Ge doped fullerene are quite less in comparison to pure fullerene. The DM values of complex E and F were respectively increased from 1.766 Debye in pure B₁₅GeN₁₆ to 4.620 and 7.668 Debye in the vacuum environment and from 3.016 Debye to 6.024 and 15.111 Debye in the solvent environment. In complex E, in vacuum, the Ge atom bonds

with the SSZ through nitrogen atom of pyridine ring with adsorption energy of -0.357 eV and the interaction distance of 1.826 Å. The B-N bond length increases while the Ge-N bond length decreases on interaction with SSZ. For solvent medium, the binding energy of complex E decreases significantly to -0.233 eV with B-N and Ge-N bond lengths showing similar trend as in vacuum. In Complex F, under vacuum conditions, Ge bonds with O atom of sulfonfyl group with adsorption energy of -0.038 eV. Both the B-N and Ge-N bond lengths decrease on adsorption of SSZ. The N₁-B₁-N₂

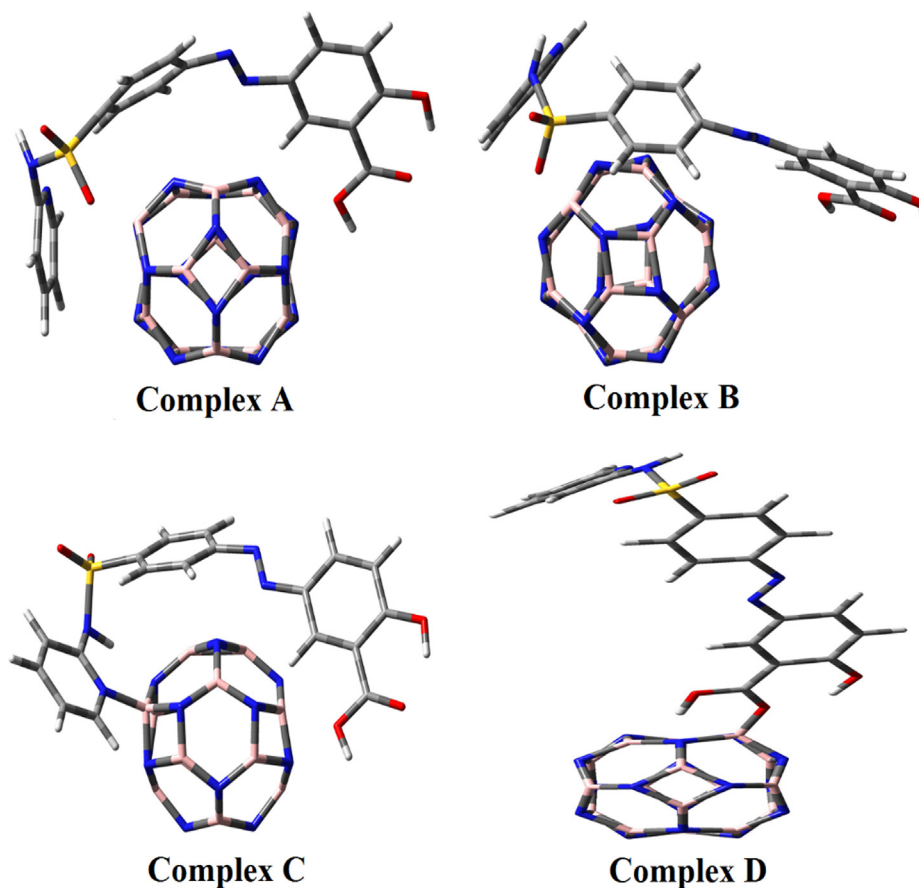


Fig. 3. Adsorption configurations of SSZ on $B_{16}N_{16}$ fullerene.

and Ge-N₂-B₁ angles decrease while N₃-Ge-N₂ increases. In solvent environment, the adsorption energy of complex F lowers to a value of -0.030 eV. The bond angle shows a similar behaviour in both mediums.

In complex G, Si bonds with nitrogen atom of SSZ through covalent bonding with the interaction distance of 1.733 Å [39]. The DM values in complex G and H respectively increased from 0.945 Debye in the pure $B_{15}SiN_{16}$ to 4.373 and 7.654 Debye in the vacuum environment and from 1.653 Debye to 6.016 and 15.120 Debye in the solvent environment. In vacuum, a very high adsorption energy is obtained (-1.274 eV) which is even higher than the SSZ interaction with pure fullerene, implying formation of a more stable complex. This effect is useful as the Si doping enhances the capability of $B_{16}N_{16}$ as drug nano carrier [69]. We observed an increase in the reactivity of the carrier when SSZ is chemisorbed on the surface of $B_{15}SiN_{16}$ fullerene, which is similar with the obtained results in the interaction between Molnupiravir and Si-doped C_{60} fullerene [70]. The B-N bond length increases while that of Si-N decreases. In solvent medium, adsorption energy of complex G decreases to a value of -1.174 eV. The bond lengths and bond angles vary in same fashion in both the mediums. In complex H, the Si atom bonds with O atom of SSZ. In vacuum the binding energy comes out to be -0.602 eV. The B-N bond length increases while Si-N bond length decreases. In solvent medium the binding energy gives an increased value of -0.663 eV. Both the bond angles and bond lengths show same behaviour in both the mediums.

3.4. Electronic features

Tables 1 and 2 illustrate the electronic structures of $B_{16}N_{16}$, $B_{15}GeN_{16}$ and $B_{15}SiN_{16}$ nanocages in both vacuum and solvent

environments. The E_H (E_L) value for the $B_{16}N_{16}$ nanocage in both vacuum and solvent environments is found to be -7.14 (-3.18) and -7.84 (-0.95) eV with E_{FL} of -5.16 eV and -4.40 eV. The calculated E_{gap} of the pure $B_{16}N_{16}$ nanocage in both vacuum and solvent environments are 6.82 eV and 6.89 eV by the PBE functional. In studies by Shakerzadeh [71] and Boshra [72], the E_{gap} value of the pure $B_{16}N_{16}$ nanocage in the vacuum environment was obtained as 6.17 eV and 6.25 eV by the B3LYP functional, respectively. Xie and co-workers have shown that $B_{16}N_{16}$ nanocage has an E_{gap} of 6.25 eV [73]. After substituting boron atoms with Ge and Si impurities, the values of E_{gap} were diminished to 6.54 eV and 6.05 eV in vacuum environment and 6.42 eV and 5.96 eV in solvent environment. These results suggested substantial alternations in the electronic structure of $B_{15}GeN_{16}$ nanocage because of higher reactivity of this system than $B_{15}SiN_{16}$ nanocage leading to enhanced sensing capabilities of cage-based biosensors.

During adsorption, interacting B atom forms an additional partial chemisorbed bond with SSZ which led to orbital hybridisation of B from sp^2 to a configuration ranging from sp^2 to sp^3 . This hybridisation resulted in a slight geometry change of interacting B with the nanocages. Also, additional degenerate states in interacting B atoms were also observed in TDOS plots of Figs. 5 and 6. These additional states led to HOMO and LUMO with higher and lower energies respectively, leading to reduced HOMO-LUMO energy gap in adsorption complexes. In terms of adsorption modes, HOMO-LUMO energy gap reduction increases in the following order: carboxyl < pyridine < sulfonyl < phenyl. In terms of nanocage type, HOMO-LUMO energy gap reduction increases in the following order: $B_{16}N_{16}$ < $B_{15}SiN_{16}$ < $B_{15}GeN_{16}$. The change in conduction electron population due to adsorption is dependent on the change in energy gap:

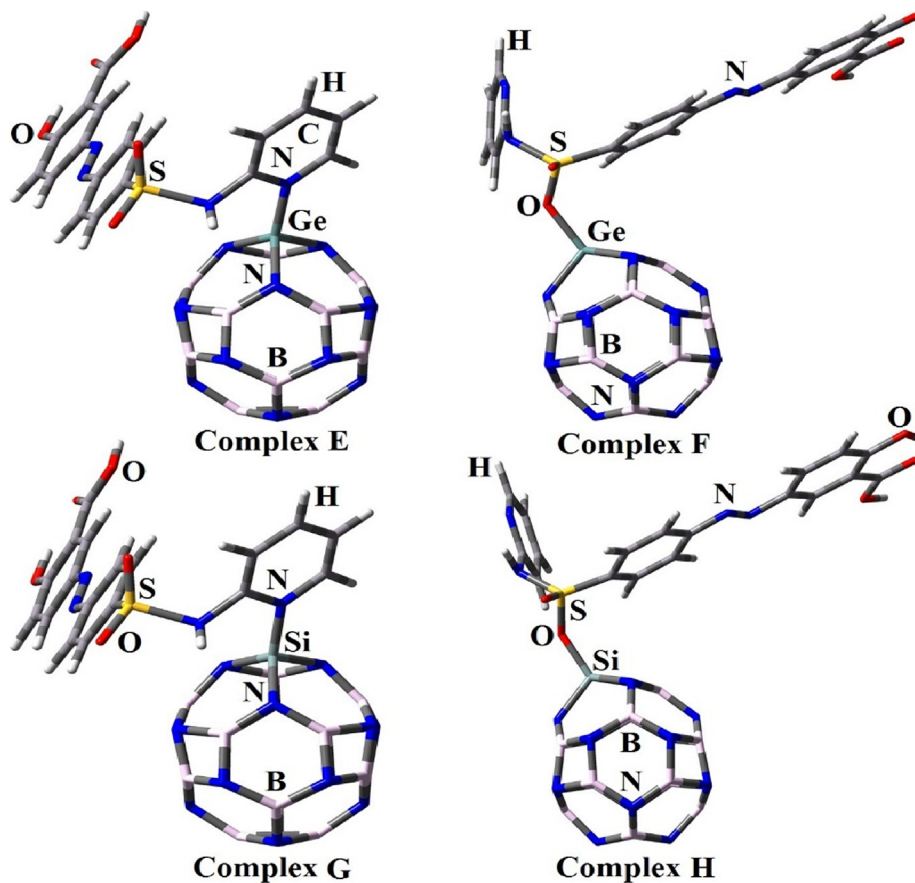


Fig. 4. Adsorption configurations of SSZ on $B_{15}GeN_{16}$ (Complex E and F) and $B_{15}SiN_{16}$ (Complex G and H).

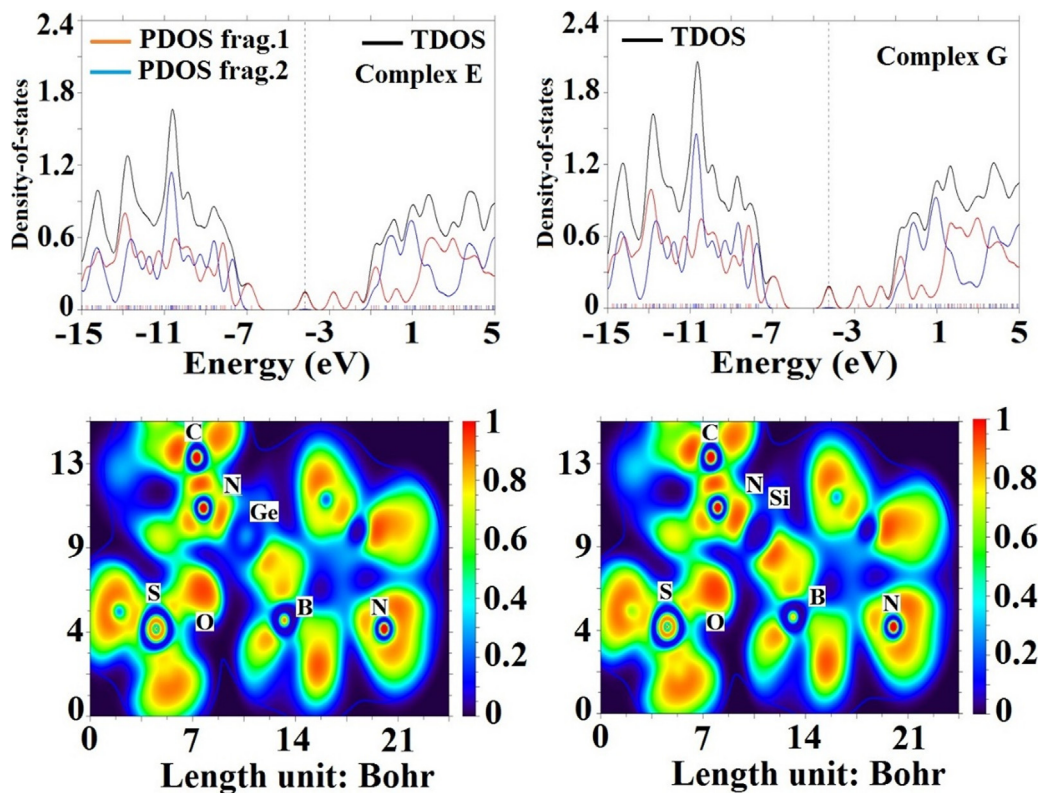


Fig. 5. DOS and ELF plots of SSZ- $B_{15}GeN_{16}$ (Complex E) and SSZ- $B_{15}SiN_{16}$ (Complex G) complexes.

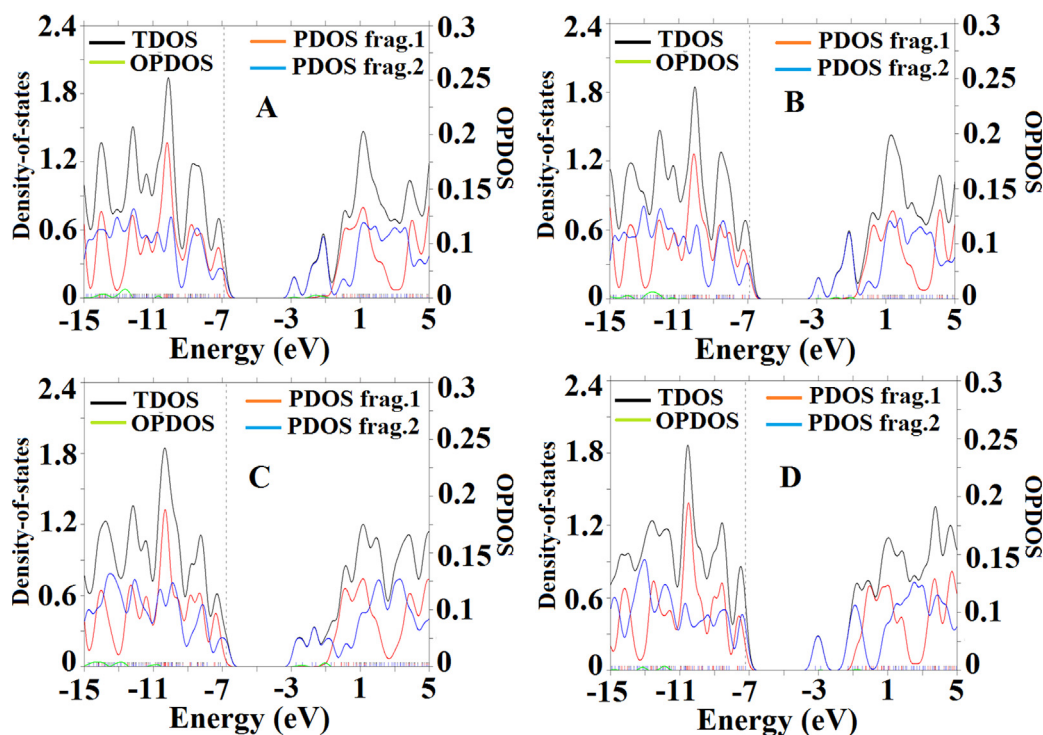


Fig. 6. DOS plots of SSZ-B₁₆N₁₆ complexes.

$$\frac{N_f}{N_i} = \exp\left(-\frac{\Delta E_g}{2kT}\right) \quad (9)$$

where N is conduction electron population, subscripts f and i represent adsorption complex and pristine material, ΔE_g is change in HOMO-LUMO energy gap, k is Boltzmann constant and T is absolute temperature. Based on Eqn. (9), a reduction in HOMO-LUMO energy gap ($\Delta E_g < 0$) will increase N_f thus enhancing electronic conductivity and sensitivity of adsorption complex. Among these nanocages, B₁₅GeN₁₆ has the largest ΔE_g and hence is the most sensitive molecular sensor.

SSZ desorption from BN nanocages could also be assessed based on the following equation [74]:

$$\tau = \nu^{-1} \exp\left(-\frac{E_{ads}}{kT}\right) \quad (10)$$

where τ is recovery time of BN nanocage and ν is attempt frequency [75–77]. In terms of adsorption modes, E_{ads} increases in the following order: sulfonyl < phenyl < carboxyl < pyridine. In terms of nanocage type, E_{ads} increases in the following order: B₁₅-GeN₁₆ < B₁₆N₁₆ < B₁₅SiN₁₆. Based on Eqn. (10), the recovery time of BN nanocage increases in the following order: B₁₅GeN₁₆ < B₁₆N₁₆ < B₁₅SiN₁₆ thus making B₁₅SiN₁₆ a molecular sensor to have best recovery performance which is instrumental for efficient and rapid drug release. Bagheri *et al.* demonstrated the interaction of adrucil with the pure, Al-, and Si-doped phagraphenes and they found that the Si-doped phagraphene could function as a suitable carrier for adrucil delivery due to a short recovery time (0.02 s at 298 K) and moderate adsorption energy (-14.0 kcal/mol) in comparison with the adrucil/phagraphene and the adrucil/Al-doped phagraphene complexes (4.09 * 10¹⁸ s and -41.8 kcal/mol) [26].

Based on Eqn. (7), ΔN values for SSZ/B₁₆N₁₆, SSZ/B₁₅GeN₁₆ and SSZ/B₁₅SiN₁₆ are 0.193 |e|, 0.115 |e| and 0.157 |e| in vacuum environment and 0.07 |e|, 0.148 |e| and 0.191 |e| in aqueous environment. The positive ΔN values indicate that there is net electron transfer from nanocage (electron donor) to SSZ (electron acceptor)

during interaction. Based on Eqn. (8), ΔE_{SE} values for SSZ/B₁₆N₁₆, SSZ/B₁₅GeN₁₆ and SSZ/B₁₅SiN₁₆ are -0.101 eV, -0.035 eV, -0.062 eV in vacuum environment and -0.013 eV, -0.056 eV and -0.09 eV in aqueous environment. All negative ΔE_{SE} values indicate that all adsorption complexes are spontaneous and stable with SSZ/B₁₆N₁₆ being the most stable.

3.5. Molecular docking

The potential anti-inflammatory and anticancer activity of selected compounds was studied using *in silico* docking simulation. To gain more understanding the inhibition mechanism between the selected complexes (Complex B, Complex C, Complex E, Complex F, Complex H, Complex G, SSZ, B₁₅GeN₁₆, B₁₅SiN₁₆, and B₁₆N₁₆) and the active site of receptors, *in silico* molecular docking study was carried out (Fig. 7). According to Table 4, complexes B and E have the lowest binding energy (BE) and appropriate interactions in the binding pocket of COX-2 (-10.8 kcal/mol) and TNF- α (-10.6 kcal/mol) compared to the other targets. Furthermore, it was found that binding energy for the complex B in the binding pocket of proteins (2AZ5, 2L5X, 3RCD and 2I0K) were -9.8, -9.7, -10.1 and -9.9 (kcal/mol) respectively. Fig. 7 depicts the 2D and 3D model of interactions between complex B and target of COX-2 indicating high binding affinity towards the active site of the target [78,79]. Complex B is structurally composed of sulfasalazine and B₁₆N₁₆ nanocage that is connected by sulfonyl group. The analysis of the interactions between complex B and target of COX-2 illustrated that amino acid residues such as Asn581, Gln192, Thr94, His95, Ser553, His90 were involved in the active site of the target through hydrophobic interaction and the complex formed polar interaction with residues like Pro 191, Tyr355, Met522, Leu352, Val523, Trp387, Phe518, Ala516, Ala527, and Pro514. Furthermore, pyridine moiety and carbonyl group of 5-aminosalicylic acid of the complex B formed two hydrogen bonds with residues Asp515 and met522 at the distances of 2.13 Å and 2.68 Å, respectively. The previous studies exhibited that sulfasalazine (1000 mg/day) at a cer-

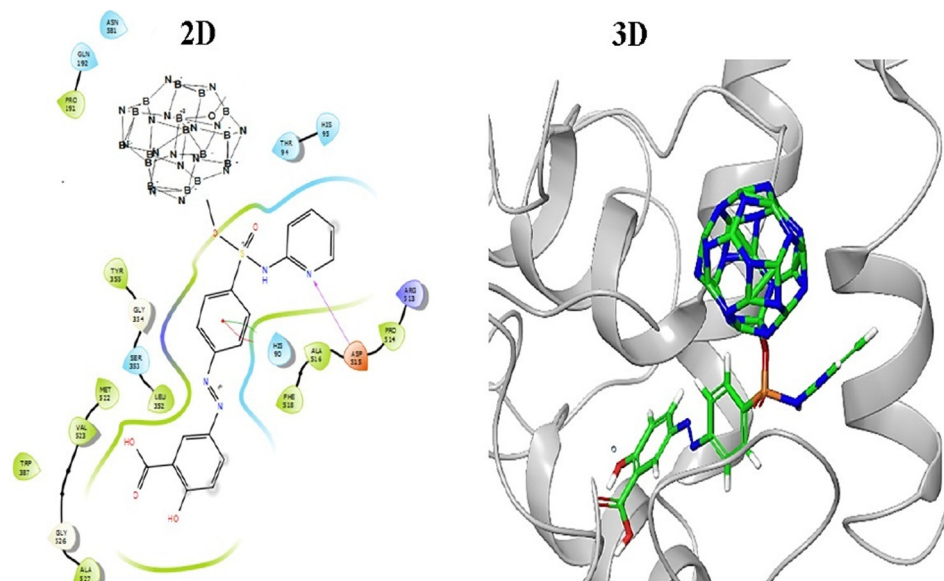


Fig. 7. Presentation of 2D and 3D models of interactions between complex B and COX-2 (PDB ID: 1CX2).

Table 4

Molecular docking simulations results for the complexes and TNF- α receptor (PDB ID: 2AZ5), IL-1 receptor (PDB ID: 2L5X), COX-2 receptor (PDB ID:1CX2), HER2 receptor (PDB ID:3RCD and 5O4G).

Compound	PDB ID: 2AZ5		PDB ID:2L5X		PDB ID:1CX2		PDB ID:3RCD		PDB ID:2IOK	
	BE (kcal/mol)	Ki (μ M)	BE (kcal/mol)	Ki (μ M)	BE (kcal/mol)	Ki (μ M)	BE (kcal/mol)	Ki (μ M)	BE (kcal/mol)	Ki (μ M)
Complex B	-9.8	5.5	-9.7	5.5	-10.8	3.4	-10.1	4.7	-9.9	4.7
Complex C	-9.9	5.4	-8.7	7.2	-9.7	5.6	-10.2	4.4	-9.2	6.2
Complex E	-10.6	3.7	-8.5	7.4	-9.1	6.4	-10.7	3.6	-9.6	5.1
Complex F	-10.1	4.6	-9.6	6.3	-9.5	6.1	-10.1	4.8	-8.8	7.4
Complex H	-10.3	4.3	-9.7	5.6	-9.2	6.3	-9.8	5.2	-9.4	5.4
Complex G	-10.5	3.9	-9.2	6.7	-9.1	6.4	-10.4	4.2	-9.1	6.5
SSZ	-7.1	9.4	-7.4	8.7	-7.8	8.1	-7.9	8.1	-7.8	7.7
B ₁₅ GeN ₁₆	-6.4	10.5	-6.4	11.4	-6.7	10.2	-6.5	10.4	-6.3	10.7
B ₁₅ SiN ₁₆	-6.2	10.8	-6.1	11.6	-6.5	10.4	-6.3	10.7	-6.1	10.9
B ₁₆ N ₁₆	-6.1	11.1	-5.9	12.1	-6.3	10.5	-6.1	10.8	-6.0	11.3

tain dose have the capability to decrease the risk of cardiovascular diseases in patients because of potential NF- κ B and TNF- α inhibitory activity [80,81]. The calculated results from molecular docking indicate that the SSZ functionalized B₁₆N₁₆ and B₁₅GeN₁₆ nanocages can be utilized to reduce the risk of cardiovascular disease via decreased levels of pro-inflammatory cytokines like IL-1, TNF- α , and COX-2 in patients with rheumatoid arthritis [82-85].

To investigate for plausible mechanism of anti-cancer activity, in silico molecular docking were performed towards two targets (PDB IDs: 3RCD and 2IOK). Based on obtained results, complex E revealed the lowest dock score (binding energy) and best orientations toward HER2 receptor [86-88]. Structural analysis exhibited that complex E have sulfasalazine and B₁₆N₁₆ nanocage possessing germanium that joined together through pyridine moiety. As shown in Fig. 8 complex E interact with amino acid residues such as Met 774, Val707, Ile752, Leu785, Leu796, Ala751, Val734, Leu726, Phe1004, Met801, Leu800, Leu852 and Phe864 by hydrophobic interaction. Additionally, complex E indicated polar interactions with amino acid residues like Lys753, Arg734, Ser783, Asp863, Thr862, Thr798, His878, Asp808, and Cys805. Furthermore, Hydroxyl and carbonyl groups of 5-aminoSalicylic acid moiety of sulfasalazine interacted via four hydrogen bonds with Ser783, Leu785 and Asp863 residues at the distances of 1.90, 2.82, 3.09 and 1.90 Å, respectively. The noncovalent interactions also have a significant role in occupation of active sites of the target

[89,90]. Also, complex E had highest binding affinity towards active site of HER2 receptors. Therefore, complex E can be considered as HER2 potential inhibitor. A comparison with the previous literature reveals that complex E binds with similar key amino acid residues in the active site of the HER2 target [91]. Based on obtained results, it is deduced that presence of germanium element (B₁₅GeN₁₆) is effectively crucial in binding interaction between the complex and binding pocket of HER2 receptor and it can lead to the possibility of improving the antitumor activity [92-99]. As a result, noncovalent interactions including hydrogen bond and hydrophobic interaction play significant role in occupation of active site of the target [100]. Accordingly, complex E could be considered as high potency inhibitor of HER2 target.

3.6. Drug likeness and ADMET prediction

Further, pharmacokinetic and toxicity properties of selected complexes were determined. The results of predicted ADMET properties of selected complexes including solubility, Number of H-bond acceptor (HBA), Number of H-bond donors (HBD), Molecular refractivity (MR), Topological polar surface area (TPSA), Drug likeness (DL), and toxicity profiles are presented in Table 5. To get an insight about the compliance of selected complexes with Lipinski's Rule of Five, the 8 complexes were screened further for more analysis. Notably, most complexes passed the Lipinski rule and did not

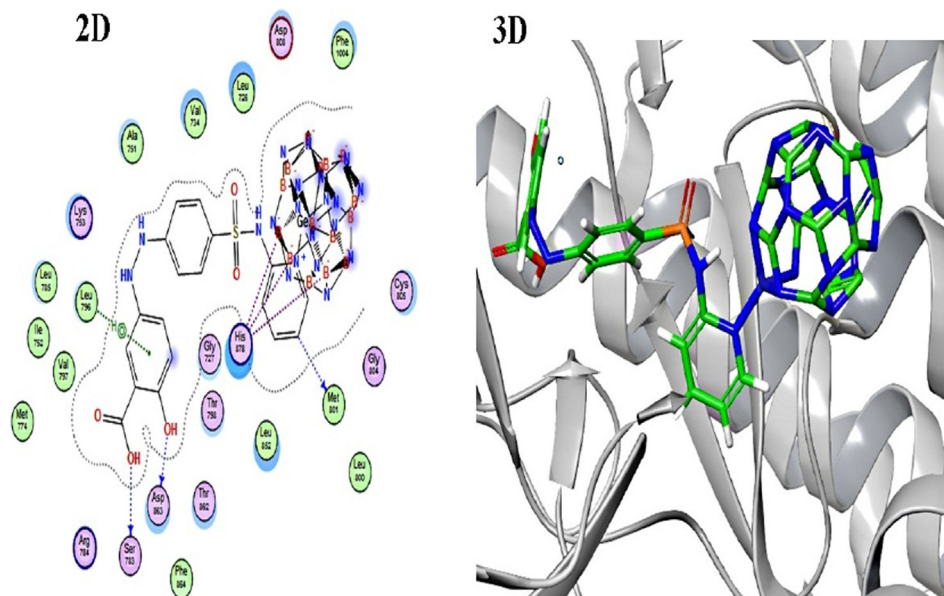


Fig. 8. Presentation of 2D and 3D models of interactions between complex E and HER-2 (PDB ID: 3RCD).

Table 5
Drug-likeness properties of compounds.

Compound	Log S	MW	Clog P	HBA	HBD	NRB	MR	TPSA	DL
ref	>-4	-	<=5	<=10	<=5	<=10	40-130	less than140	-
Complex B	-3.3	795	-7.8	24	3	8	131	187	0.75
Complex C	-3.4	796	-8.2	25	3	7	135	181	0.5
Complex E	-0.34	857	-8.2	24	3	7	131	181	0.65
Complex F	-3.5	812	-7.9	22	3	7	137	187	0.52
Complex H	-0.35	814	-8.2	24	5	9	139	182	0.67
Complex G	-0.34	812	-8.2	22	3	7	134	181	0.51
SSZ	-5.1	398	3.7	7	3	6	100	141	0.7
B ₁₅ GeN ₁₆	-3.0	414	-11.1	16	0	0	95	50.4	0.61
B ₁₅ SiN ₁₆	-3.1	458	-11.1	16	0	0	92	50.4	0.53
B16N16	-3.1	397	-11.0	16	0	0	98	49	0.52

Abbreviations: Logs: Logarithm of water solubility; MW: molecular weight; logP: Logarithm of compound partition coefficient between *n*-octanol and water; HBA: Number of hydrogen bonds acceptors; HBD: Number of hydrogen bond donors; TPSA: Topological polar surface area; NRB: Number of rotatable bonds, MR: Molecular refractivity, DL: Drug likeness.

Table 6
ADMET profile of compounds.

Compound	BBB	HIA	Caco2	P-GI	CYP450-2C9	CYP450-2D6	CYP450-3A4	AMES	CIG	HPT	AOC
ref	-	-		No	No	No	No	No	No	No	-
Complex B	Yes	Yes	No	No	No	No	No	No	No	No	2.4
Complex C	Yes	Yes	No	No	Yes	No	No	No	No	No	2.5
Complex E	Yes	Yes	No	No	No	No	No	No	No	No	2.44
Complex F	Yes	Yes	No	No	No	No	No	No	No	No	2.47
Complex H	Yes	Yes	No	No	No	No	No	No	No	No	2.4
Complex G	Yes	Yes	No	No	No	No	No	No	No	No	2.5
SSZ	Yes	Yes	No	No	Yes	No	No	No	No	No	2.63
B ₁₅ GeN ₁₆	Yes	Yes	No	No	No	No	No	No	No	No	1.56
B ₁₅ SiN ₁₆	Yes	Yes	No	No	No	No	Yes	No	No	No	1.5
B16N16	Yes	Yes	No	No	No	No	No	No	No	No	1.7

Abbreviations: BBB: Blood Brain Barrier; HIA: Human Intestinal Absorption; P-GI: P-glycoprotein inhibitor; CIG: Carcinogens; HPT: hepatotoxicity; AOC: Acute oral Toxicity.

show any violation of standardized Lipinski rule of five. The values of calculated solubility of complexes range from -3.1 to -5.1, indicating moderate solubility of complexes. The calculated Log P values lie between 3.7 and -11.1 (unit). Moreover, molecular weight was more than 500 Dalton, number of Hydrogen bond donors were

less than 5, while number of H-acceptors of all complexes except sulfasalazine exhibited violation of Lipinski rule. Molar refractivity of most complexes were more than 130 while sulfasalazine and other nanocages have low values. According to the data presented in Table 5 and 6, pharmacokinetic parameters and ADMET profiles

(absorption, distribution, metabolism, excretion, and toxicity) were evaluated to predict the toxicity parameters of the complexes [101]. All the complexes are following the criteria of drug-likeness rules and indicated good absorption and permeability demonstrating moderate absorption [102–104]. The evaluation of the two key ADMET descriptors i.e., Human Intestinal Absorption (HIA) and blood brain barrier (BBB) exhibited appropriate profiles. In addition, all the selected complexes were indicated as non-inhibitors of the P-glycoprotein inhibitor (P-GI). Furthermore, the obtained results revealed that all the complexes were non-inhibitors of CYP450 enzymes (3A4, 2D6, and 2C9) acting as an effective factor in drug metabolism. Additionally, most complexes were non-carcinogenic, non-hepatotoxic and non-AMES toxic. Overall, regarding drug likeness properties and ADMET parameters, it could be concluded that most complexes had low risk, appropriate drug likeness behaviour and ADMET predictions; hence the selected complexes could be considered carefully as high potency inhibitors of anti-inflammatory and anti-cancer activities.

4. Conclusions

In this study, the therapeutic potential of $B_{16}N_{16}$, $B_{15}GeN_{16}$ and $B_{15}SiN_{16}$ nanocages as a novel carrier for sulfasalazine to treat inflammatory cardiovascular disease and breast cancer was compared with the pure sulfasalazine (SSZ). The structural and electronic features of SSZ, $B_{16}N_{16}$, $B_{15}GeN_{16}$ and $B_{15}SiN_{16}$ nanocages and their complexes were determined to show the efficiency of carriers as drug-delivery systems. Adsorption of SSZ via its pyridine ring over $B_{16}N_{16}$, $B_{15}GeN_{16}$ and $B_{15}SiN_{16}$ nanocages was established through electrostatic interaction, leading to improved dipole moment, which was remarkably higher than $B_{15}GeN_{16}$ and $B_{15}SiN_{16}$ nanocages. The *in silico* molecular docking simulation exhibited that complex B consisting of $B_{16}N_{16}$ nanocage and sulfasalazine had a strong interaction with targets of TNF- α , IL-1, and COX-2 in contrast to other complexes, while the complex E containing $B_{15}GeN_{16}$ nanocage and SSZ interacted strongly with HER2 receptor. Furthermore, the complexes revealed appropriate ADMET properties and hence the selected complexes could be considered as promising candidate for inhibition of binding pocket of COX-2 and HER-2 receptors. Thus, it is deduced that Ge decorated $B_{16}N_{16}$ is an excellent material to be utilized as carrier for delivery of sulfasalazine drug for treatment of cardiovascular diseases and breast cancer with minimum side effects.

Authorship contribution statement

Kadda Hachem: Editing, review and rewrite the paper, supervision, software, supervision.

Maria Jade Catalan Opulencia: Editing, revise the paper, rewrite.

Walid Kamal Abdelbasset: Editing, review and rewrite the paper, rewrite.

Andrey Sevbitov: Writing, review and editing.

Oleg R. Kuzichkin: Review, writing and editing.

Abdullah Mohamed: conceptualization, data curation, formal analysis, investigation.

Sahar Moazen Rad: methodology, writing the original draft, formal analysis, investigation.

Aref Salehi: methodology, writing the original draft, supervision, formal analysis, investigation.

Jupinder Kaur: formal analysis, software, formal analysis, investigation.

Ravinder Kumar: Methodology, validation, conceptualization, data curation.

Andrew Ng Kay Lup: writing, review and editing, validation, supervision.

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Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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