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SYNTHESIS, CRYSTAL STRUCTURE AND DFT STUDY OF 5-BROMO-1-TOSYL-1*H*-PYRROLO[2,3-*b*]PYRIDINE*

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The 7-azaindoles derivatives, as a major category of compounds, show a widespread of biological activities. The compound 5-bromo-1-tosyl-1*H*-pyrrolo[2,3-*b*]pyridine (1) was obtained by one-step substitution reaction. In the meantime, the structure of compound 1 was confirmed by ¹H NMR, ¹³C NMR, FTIR and X-ray diffraction. The optimized molecular crystal structure was preliminarily determined by using density functional theory (DFT) and compared with the X-ray diffraction values. In addition, more physical and chemical properties of compound 1 have been studied by further studying the molecular electrostatic potential and frontier molecular orbital of compound 1.

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INTRODUCTION

Indazole derivatives have been reported to have anticancer, antiviral, antibacterial, antiprotozoal, antipsychotic, antiinflammatory, analgesic and radio sensitizing properties. Nitrogen-containing heterocyclic compounds are widely found in nature and in living organisms, and are of great interest in the fields of medicine, agriculture, functional materials and chemicals because of their high efficiency, low toxicity, environmental friendliness, and unique biological activity and structural diversity [1-4]. Novel molecularly targeted drugs usually have relatively clear targets, mainly acting on key molecules and their signal transduction pathways that regulate cell growth and proliferation, which differ greatly between normal and tumor cells, and can inhibit the growth of tumor cells while reducing the effect on normal cells, thus increasing the selectivity to tumors and reducing the toxicity to normal tissues. Among them, 7-azaindoles can inhibit the activity of various proteases in pharmacological effects, and have anticancer, antiviral, and other effects [5-8]. As widely studied intermediates of C3 and C5-substituted 7-azaindoles derivatives, 5-bromo-1-tosyl-1*H*-pyrrolo[2,3-*b*]pyridine (1) is available at the C3 position via Negishi reaction, Suzuki cross-coupling reaction, and at the C5 position via Buchwald-Hartwig

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Fig. 1. Crystal structure (a) and DFT optimized structure of compound 1 (b) and their overlap figure (c) (green: crystal structure; pink: DFT optimized structure) (see the electronic version).

coupling, Suzuki-Miyura coupling, which can be further modified to develop novel 7-azaindole derivatives. In addition, the synthesis method of the compound reported so far is to react under strong bases such as potassium hydroxide, sodium hydroxide and sodium hydroxide, while the method used in this paper is based on triethylamine and carried out under the condition of catalyst 4-dimethylaminopyridine (DMAP). The advantages of this method are mild conditions, simple post-treatment and high yield of the target compound, so it is suitable for industrial production. It is understood that the melting point of the measured substance can qualitatively identify the purity of the measured substance. The closer the melting point of the measured substance is to its standard melting point, the purer the substance will be. So we determined the melting point of compound 1. The melting range of compound 1 reported in the present literature is between 140-141 °C, while the melting point measured in this experiment is 138-140 °C, which is consistent with the melting range reported in the literature [9].

In this paper, the target product 5-bromo-1-[(4-methylphenyl)sulfonyl]-1*H*-pyrrolo[2,3-*b*]pyridine was synthesized by a one-step substitution reaction (Fig. 1) using 5-bromo-7-azaindole as the starting material, based on the principles of collocation, local modification, bioelectronic equipartition and bioequivalence, while preserving the mode of action of the pharmacophore. The structure was confirmed by FTIR, hydrogen spectroscopy, carbon spectroscopy, NMR spectroscopy and mass spectroscopy. The dominant conformation was obtained by density functional theory (DFT) calculations using Gaussian09 software package in B3LYP/6-311+G(2*d*,*p*) mode [10, 11]. Crystallographic and conformational analysis of compound **1** and comparison of the structure of X-ray single-crystal diffraction with the DFT-optimized structure revealed that the structure of the title compound **1** is indeed identical to the single-crystal structure determined by experiment. In order to reveal the pharmaceutical aspects of 5-bromo-1-[(4-methylphenyl)sulfonyl]-1*H*-pyrrolo[2,3-*b*]pyridine, we also investigated the physicochemical characteristics, molecular electrostatic potential and frontier molecular orbitals of the title compound by DFT.

EXPERIMENTAL

General remarks

All the experimental materials used, including chemicals and reagents, were commercial suppliers and were used without further purification. The default TMS was an internal standard, and ¹³C and ¹H NMR spectra (400 MHz) in DMSO- d_6 solvent were recorded on a JEOL-ECX NMR spectrometer. Mass spectrometry studies were performed on the Agilent Technologies 1100 organic mass spectrometer. The UV study was performed on a UV spectrophotometer (Shanghai Yuanxi Instrument Co., Ltd.). The IR spectra of compound **1** were recorded in the 4000-400 cm⁻¹ regions on the Bruker IFS-55V IR spectrometer (Bruker, Germany). X-ray diffraction data were recorded on a Bruker Apex II X-ray diffractometer and collected using graphite monochromatic K_{α} radiation ($\lambda = 0.71073$ Å). Compound **1** was synthesized by one step, and the reaction was monitored by thin layer chromatography.

Synthetic procedure

In a 500 mL single-mouth flask, 5-bromo-7-azaindole (20 g, 101.5 mmol) was dissolved in 200 mL tetrahydrofuran, and 4-dimethylaminopyridine (DMAP) (0.62 g, 5.1 mmol) and triethylamine (20.54 g, 202.6 mmol) were added sequentially. *p*-Toluenesulfonyl chloride was slowly added under stirring and the reaction was carried out at room temperature for 1 h. The reaction was monitored by TLC. After the reaction, the reaction solution was poured into 300 mL of water with slow stirring, a solid was precipitated, filtered, and the filter cake was washed with a little water to obtain a light yellow solid, the filter cake was dried to obtain 35.90 g of crude product, the crude product was pulped with petroleum ether to obtain 5-bromo-1-[(4-methylphenyl)sulfonyl]-1*H*-pyrrolo[2,3-*b*]pyridine as a white solid (34.30 g, yield: 96.2%). ¹H NMR (400 MHz, CDCl₃) 8.46(d, J = 4.0 Hz, 1H), 8.07 (d, J = 8.0 Hz, 2H), 7.98 (d, J = 4.0 Hz, 1H), 7.76 (d, J = 4.0 Hz, 1H), 7.31 (d, J = 12.0 Hz, 2H), 6.55 (d, J = 4.0 Hz, 1H), 2.39 (s, 3H); ¹³C NMR (101 MHz, CDCl₃) δ , ppm 145.51, 145.45, 135.05, 131.72, 129.74, 128.09, 127.94, 124.43, 115.18, 104.48, 21.67.



Scheme 1. Synthesis route of compound 1.

X-ray crystal structure determination

The crystals of compound **1** were successfully obtained by solvent evaporation method. 10 mg of compound **1** was weighed and dissolved in 5 mL of a mixture of acetonitrile and DMF, and the solution was filtered through an organonylon syringe filter ($\emptyset = 0.22 \,\mu$ m) and transferred to a 10 mL reagent bottle. The bottle was sealed with a film and three small holes of the same size were punched in the film with a clean needle; single crystals of 5-bromo-1-tosyl-1*H*-pyrrolo[2,3-*b*]pyridine were obtained after standing at room temperature and slow evaporation of the solvent. Crystals that were transparent, free of cracks and had a clean and shiny surface were selected for data collection under microscopic observation, and the size of the selected crystals was $0.200 \times 0.200 \times 0.200 \,\mathrm{mm^3}$. Multiple scans and absorption corrections were performed at 293 K using a Bruker APEX II X-diffractometer in scanning mode with rays (0.071073 Å). The collected diffraction data were corrected using the full matrix least squares method and then refined with the SHELXL-2017/1 program. The crystal data for compound **1** are shown in Table 1.

Quantum chemistry/DFT calculation

The DFT calculations, which for the purpose of verifying the dependability of the X-ray crystal data, were performed in the ground state (in vacuo) with Gaussian09 software package [12] by using the B3LYP method with the 6-311+G(2d,p) basis set [13]. Furthermore, based on the optimized structure [14], information such as geometrical, electronic and energy parameters, was extracted from the GuassView 5.0 program [12].

RESULTS AND DISCUSSION

Synthesis and characterization

The ¹³C NMR, ¹H NMR spectroscopy and MS of title compound **1**, as shown in Fig. S1-S3 of Supplementary Material, they all confirmed that the structure of the compound was correct.

TABLE 1. Crystal Data and Parameters for Structure Refinement of Compound 1

Compound	1		
CCDC	2216318		
Molecular formula	C ₁₄ H ₁₁ Br N ₂ O ₂ S		
Molecular weight	351.22		
Crystal system	Monoclinic		
Space group	$P2_{1}/c$		
<i>a</i> , <i>b</i> , <i>c</i> , Å	11.174(3), 17.298(4), 7.7006(17)		
α , β , γ , deg	90, 108.677(6), 90		
$V, Å^3$	1410.1(6)		
Ζ	4		
Density (calculated), mg/m ³	1.654		
μ , mm ⁻¹	3.065		
Radiation λ, Å	0.71073		
Ranges / indices (h, k, l)	$-13 \le h \le 13, -20 \le k \le 20, -9 \le l \le 9$		
N(hkl) _{measured} , N(hkl) _{unique}	2499, 1792		
$R_{ m int}$	0.0672		

Crystallographic analysis

Further study on the structural characteristics of the compound can better analyze its various physical and chemical properties. Therefore, X-ray diffraction is adopted to study its structure. A ORTEP diagram can clearly and intuitively display the structural characteristics of compound **1**, as shown in Fig. 1. Among them, we stack the structure of the obtained compound with the theoretical calculation results, as shown in Fig. 1, the green part is the obtained crystal, and the pink part is the theoretical calculation. The crystal packing in the compound **1** is stabilized by C4–H4…O1 and weak C8–H8… π interactions, the perpendicular distance of different molecules is 2.765 Å, as shown in Fig. 2. In general, the crystal structure shows that the intermolecular packing is stabilized by π – π interactions, C–H… π interactions and C–H…O hydrogen bond. The crystallographic and refinement data are shown in Table 1. The experimental and theoretical values of all bond lengths,



Fig. 2. The crystal packing of title compound (*a*); π – π stacking of compound 1 and centroid–centroid distances between two molecules (*b*); the distance of the hydrogen bond of the title compound (*c*).

bond angles and torsion angles as well as the differences between the two values are within reasonable range, as shown in Table 2 and Table S1 of Supplementary Materials.

Conformational determination

The biological activity and physicochemical properties of a molecule is dramatically affected by its conformation [15, 16]. Therefore, what takes a salient role in further study the structure of a compound is the reliable conformational analysis. The initial conformation is very important for the study of the compound **1**, and this initial conformation is searched by means of Spartan08 program [17] with an MMFF [18, 19] molecular mechanics force field. Whereafter, at the level of DFT/B3LYP/6-311+G** [20, 21], frequency calculation and geometric optimization of all possible conformations are

Bond distance, Å	Experimental geometry parameters for molecule 1	Calculated geometry parameters for conformer 1-2	Difference
Br1–C13	1.896(5)	1.91571	0.01971
C1–C2	1.501(7)	1.50602	0.00502
C2–C3	1.399(7)	1.39860	-0.0004
C3–C4	C3–C4 1.373(7) 1.		0.01376
C4–C5	1.385(6)	1.385(6) 1.39192	
C5–C6	1.393(6)	1.39022	-0.00278
C5–S1	1.749(5)	1.77431	0.02531
C6–C7	1.380(7)	1.38821	0.00821
C8–C9	1.350(7)	1.35973	0.00973
C8-N1	1.392(5)	1.39501	0.00301
C9–C10	1.436(6)	1.43396	-0.00204
C10-C14	1.378(6)	1.39470	0.0167
C10-C11	1.397(6)	1.41663	0.01963
C11-N2	1.328(6)	1.32173	-0.00627
C11–N1	1.392(5)	1.39430	0.0023
C12-N2	1.336(6)	1.33217	-0.00383
C12-C13	1.389(6)	1.39983	0.01083
C13–C14	1.375(6)	1.38355	0.00855
N1-S1	1.677(4)	1.72253	0.04553
O1–S1	1.426(3)	1.44419	0.01819
O2–S1	1.423(3)	1.43679	0.01379

TABLE 2. Selected Experimental and Calculated Geometry Parameters for 1



Fig. 3. Stable conformers of compound 1.

TABLE 3.	Gibbs Free	Energy ((G), Rela	ative Gibbs	s Free E	nergy $(\Delta G)^{\#}$	1 , and 1	Boltzmann
	Weighting	g Factor ($Pi, \%)^{#2}$	of the Con	formers	s of Compou	ınd 1	

Conformer	G , kcal/mol ΔG , kcal/m		Pi, %
1-1	-3772.436248	0	56.12
1-2	-3772.436018	0.144326357	43.88

^{#1} Related to the most stable conformer.

^{#2} Boltzmann weighting factor (*Pi*, %) based on ΔG .

realized using Gaussian09 software package. On the basis of the relative free energies, it was feasible for us to predicted the overall percentage of each conformation in the room-temperature equilibrium mixture. The Gibbs free energy (*G*), relative Gibbs free energy ($\Delta G = \exp(-Gi/RT)$) and Boltzmann distribution (Boltzmann weighting factor $Pi = \frac{\exp(-Gi/RT)}{\sum_{i} \exp(-Gi/RT)} \cdot 100\%$) for different conformers of compound **1** are shown in Table 3.

There are two relatively stable conformers of the title compound 1 in Fig. 3, which are conformers 1-1 (56.1%), 1-2 (43.88%), and both of conformers are dramatically existed at room temperature. What the distinctions between the two conformers were principally classified into the rotational orientation of the S1–N1.

A comparison was drawn between the DFT-optimized structure and the crystal structure of the compound **1**. The crystal conformation obtained using X-ray diffraction, as shown in Table 3 and Table S1, the experimental value of the crystal is nearly as same as the calculated value of the conformer **1-2**. For example, bond angle C7–C2–C3, C7–C2–C1, C3–C2–C1, what the difference between the experimental values of crystal structure and the calculated values of conformer **1-2** are 0.6, -0.8, -0.03. Thus, as expected, the crystal conformation offered by X-ray diffraction is as consistent as the result of the conformer **1-2** which was calculated by DFT.

Not only the hydrogen bonds but the π - π stacking interactions were formed the molecular packing of the crystal structure of the compound **1**, as clearly shown in Fig. 3. Hydrogen bonds, similar to the X–H···Y, can be formed between a hydrogen atom and two other atoms that are more electronegative but have smaller atomic radius. Hence, there is an intermolecular hydrogen bonding: C4–H4…O1, which was found in the title compound **1** and played an extremely crucial part in stabilizing the molecule (Table 4). Except for the aforementioned interactions, what worth noting is that weak π - π stacking interactions have better ability to stabilize structures, including the distance between the ring formed by *Cg*1 and C1, between the ring formed by *Cg*1 and C1, between the ring formed by *Cg*1 and C1, between the ring formed by *Cg*2 and C1, as a result, the perpendicular distance of two rings of two molecules respectively are $d_{Cg-Cg} = 3.851$ Å, $d_{Cg-Cg} = 3.660$ Å (Fig. 3). The detailed information of the title compound hydrogen bonds is listed in Table 4.

MEP

The B3LYP/6-311+G(2d,p) method was selected to investigate the molecular electrostatic potential of the conformer 1-2 (same as the crystal structure). Because it indicated the information about the intermolecular interaction region of the title compound 1, which was great help to further understand the compound 1. In the MEP map, as shown at the surface of the

<i>D</i> –Н…А	<i>d</i> (<i>D</i> –H), Å	<i>d</i> (H <i>A</i>), Å	<i>d</i> (<i>DA</i>), Å	\angle (<i>D</i> -H <i>A</i>), deg
C4–H4O1	0.93	2.56	2.922(6)	103

TABLE 4. Hydrogen-Bond Geometry of Compound 1



Fig. 4. Molecular electrostatic potential map of conformer **1-2**.

map, different colors suggest the different electrostatic potentials. And the underlying increases in the order of red < orange < yellow << green < blue. The color code of the maps in the range $-5.0 e^{-2}$ (deepest red) to $5.0 e^{-2}$ (deepest blue) in title molecule surface, where red color denotes the electron rich area and blue region indicates electron deficient region. As shown in Fig. 4, The O1 and O2 atom in the 5-bromo-1-tosyl-1*H*-pyrrolo[2,3-*b*]pyridine as well as the N1 and N2 atoms in the conformer **1-2** are surrounded by negative charges, which indicate some possible nucleophilic attack sites. Besides, the H atom of the C4 atom, C5 atom and the C6 are located on the positive charge regions.

FMOs

Frontier molecular orbitals investigation invariably plays an extremely important role in reactive prediction [22, 23]. As an electrons donor, on the one hand, the highest occupied molecular orbital (HOMO) is always related to potential electron delocalization directly. Indeed, the ability of charges transfer revealed from the energy of HOMO. On the other hand, the lowest unoccupied molecular orbital (LUMO) manifests an area where electrons can be accepted, and the energy of LUMO is regarded as electron affinity. What the energy of band gap point to the energy difference between this two important FMOs which implies the stability of molecular structure.

As a high-efficiency method to further investigate the chemical stability of conformer 1-2, the B3LYP/6-311+G(2*d*,*p*) was selected to calculate the energies of the highest occupied molecular orbital (HOMO), the lowest unoccupied molecular orbital (LUMO), and their orbital energy gap were calculated. The pictorial illustration of the frontier molecular orbitals (FMOs) and their respective positive and negative regions represented by red and green colors are shown in Fig. 5. The values of LUMO and HOMO were -1.7374 eV and -6.7378 eV, respectively. The value of the energy separation between the HOMO and LUMO was 5.004 eV for conformer 1-2. The large HOMO–LUMO energy gap means high excitation energy of the excited state, good chemical stability and large hardness for the calculated conformer. In addition, the ionization energy and electron affinity can be expressed as: $I = -E_{HOMO} = 6.7378$ eV, $A = -E_{LUMO} = 1.7374$ eV. The hardness, which can be denoted as: $\eta = (I - A)/2$, indicates the resistance toward the deformation of the electron cloud of chemical systems under small perturbation encountered during chemical process [24-26]. Therefore, the hardness of title compound **1** is 2.5002.

Ultraviolet-visible spectral analysis

The experimental procedure of a 5 ppm concentration in the methanol was recorded. The experimental and theoretical UV-Vis spectra of compound 1 were relatively analyzed, as shown in Fig. 6, and the theoretical spectrum of compound 1 shifted by 34 nm to a higher wavelength agrees well with the experimental spectrum. For compound 1, the experimental spectra showed a strong absorption at 222 nm and a weak absorption at 241 nm and 291 nm. The strong band is



Fig. 5. The highest occupied and lowest unoccupied molecular orbitals of conformer 1-2 obtained by the DFT/6-311+G(2d,p) method.



Fig. 6. Theoretical and experimental UV spectra of compound 1 (a), the excited state energy (b).

obviously due to the conjugated double bond $\pi \rightarrow \pi^*$ transition of the C=C double bond. The weak band is due to heteroatom N and $n \rightarrow \pi^*$ transitions of the C-Br, C-S.

The data in this paper are based on the analysis of DFT calculation of UV-Vis spectrum, we used the Gauss08 software package to perform the DFT calculation in the ground state. According to the optimized structure, the electronic, geometric and energy parameters were extracted from the GuassView 6.0 program. At the same time, we calculated the excited state of 40.

From TD-DFT calculations, the excited state energy and the corresponding absorption wavelength are shown together with the approximate curve in Fig. 6.

Vibration Analysis

It has been proved that the vibration of functional groups is bound to have different effects due to the different amplitudes of each functional group and the environment in which the molecule is located. Therefore, infrared spectroscopy



Fig. 7. Theoretical and experimental FTIR spectra of compound 1.

is widely used to analyze and determine the structure of compounds. As shown in Fig. 7, the theoretical and experimental infrared spectra of compound **1** are clearly listed, and the measured, calculated.

C–Br vibrations. Compound 1 has a bromine substituent with an absorption frequency of 690-515 cm⁻¹ [27]. Therefore, 701.53 cm⁻¹ in the IR spectrum can be considered as C–Br stretching vibration.

Ring vibrations. In general, the carbonyl vibrations of aromatic heterocycles such as pyridine ring, pyrrole ring, and benzene ring have about characteristic peaks around 1600 cm^{-1} , 1580 cm^{-1} , 1500 cm^{-1} , and 1450 cm^{-1} [28]. Thus, the characteristic peaks at 1592.72 cm^{-1} , 1513.88 cm^{-1} , and 1439.48 cm^{-1} in the IR spectra can be attributed to the stretching vibrations of the aromatic rings.

For aromatic compounds containing benzene rings, absorption bands were observed between 1250 cm^{-1} and 950 cm^{-1} for C–H in-plane bending vibrations and between 900 cm^{-1} and 600 cm^{-1} for out-of-plane bending vibrations [28]. In the chromatogram of the title compound **1**, the absorption band at the in-plane bending vibration band with the aromatic ring is clearly observed at 912.40 cm^{-1} .

Ar–N vibrations. Amines with C–N bonds, such as aromatic or unsaturated amines, stretch at 1360-1250 cm⁻¹. In secondary amines, when the aryl group is attached to the nitrogen atom, two peaks can be seen in the IR spectrum of compound **1** with C–N stretching vibrations at 1243.03 cm⁻¹ and 1355.53 cm⁻¹.

Sulfone (S=O=S) vibrations. The frequency of the absorption band where the sulfonyl group is located is 1340-1290 cm⁻¹. In the IR spectrogram of compound 1, the position where the absorption band is 1335.53 cm⁻¹ can be considered as the stretching vibration.

CONCLUSIONS

In this study, 5-bromo-1-[(4-methylphenyl)sulfonyl]-1*H*-pyrrolo[2,3-*b*]pyridine and was synthesized using a onestep reaction route. The structure of the compound was confirmed by MS and analyzed by ¹H NMR spectra, IR spectra and ¹³C NMR spectra, in full agreement with the validation results. In addition, the conformation **1-2** obtained by DFT optimization are consistent with those determined by X-ray single crystal diffraction. The physical properties of compound **1** were further analyzed using the values of molecular electrostatic potential and frontier molecular orbitals.

CONFLICT OF INTERESTS

The authors declare that they have no conflicts of interests.

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