

Synthesis, spectroscopic characterization, X-Ray analysis, and DFT-HF calculations of 5-ethoxymethyl-8-hydroxyquinoline

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Abstract 5-ethoxymethyl-8-hydroxyquinoline was synthesized and characterized using spectroscopic methods (1 H, 13 C NMR, IR). The crystal structure determined at room temperature (295 K) by means of X-ray powder diffraction is orthorhombic, with space group Pbca and eight molecules per unit cell (Z = 8, Z' = 1). The lattice parameters are: a = 7.9551(12) Å, b = 17.981(3) Å, c = 15.125(2) Å and c = 163.5(6) Å 3 . Geometric parameters and properties depending on the charge distribution around the different types of donors and acceptors bonds within the molecule are calculated by density functional theory (DFT/B3LYP) and Hartree–Fock methods. Atomic charges and dipole moment value permit qualitative predictions about high reactivity of this molecule. The 5-ethoxymethyl-8-hydroxyquinoline adopts a non-planar structure in the solid state and the molecule is stabilized by contact system as π – π stacking interactions, weak intra and intermolecular H-Bonding O–H···N and C–H···O types, this latter involving the rings of both adjacent molecules in plans with a gap from 0.638 Å.

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1 Introduction

8-hydroxyquinoline derivatives are very attractive bioactive molecules especially in the field of drugs and pharmaceuticals (Di Varia et al. 2004; Salkovic-Petrisic et al. 2015; Cacciatore et al. 2013) and as chelating agents for a large number of metal ions (Pollak et al. 2013; La Deda et al. 2004). Derivatives such as zinc(II) bis (8-hydroxyquinoline) complex have been investigated as electroluminescent (EL) materials over the past decades (Yang et al. 2013; Yuan et al. 2012; Sokołowski et al. 2012; Yuan et al. 2013; Zawadzka et al. 2013; Yuan et al. 2015; Wang et al. 2013; D'Souza et al. 2008) including electrotransporting and emitting materials, which are applied in molecular-based OLEDs for their thermal stability, high fluorescence and excellent electron-transporting mobility (Shan et al. 2014; Huo et al. 2010; Zawadzka et al. 2002; Zawadzka et al. 2016; Zawadzka et al. 2013; Zhang et al. 2015; Yuan et al. 2012).

In this way, many studies have been published to understand the reactivity of this derivatives in different solvents (Himmi et al. 2008) and to determine the most stable conformation in acidic and alkaline aqueous solutions (Saylam et al. 2014). But up today only few structural data on 8-hydroxyquinoline derivatives have been published, showing, for example dependence of the molecule structure on the synthesis procedures of 8-hydroxyquinoline derivatives (Pearson et al. 1967; Gershon et al. 1969; Collis et al. 2003). In this study, we focus on a new one of 8-hydroxyquinoline derivatives named 5-(ethoxymethyl)quinolin-8-ol (5-EMQO) (Scheme 1). This molecule was synthesized and analyzed by different spectroscopic methods. The X-ray analysis, DFT and HF calculations permit us to study the geometric and charge transfer properties in the ground state of the compound.

2 Experimental section

2.1 Synthesis method

5-EMQ was synthesized according to the method described by Mishra et al. (2005). The obtained product (Scheme 1) was characterized by IR and ¹H, ¹³C NMR methods, the spectral data are shown in Table 1.

Scheme 1 Synthesis of 5-ethoxymethyl-8-hydroxyquinoline

$$C1$$
 C_2H_5OH, K_2CO_3
 $Reflux, 1 h$
 $O+C_2H_5$
 $O+C_2H_5$
 $O+C_2H_5$
 $O+C_2H_5$
 $O+C_2H_5$
 $O+C_2H_5$



 Table 1
 Spectral data of 5-ethoxymethyl-8-hydroxyquinoline

| ¹ H NMR (300 MHz, DMSO-d ₆) | IR (KBr) (cm ⁻¹) | ¹³ C NMR (75 MHz, DMSO-d6) |
|---|---|---|
| $\begin{split} \overline{\delta_{ppm}} &= 7.049.07 \text{ (m, 4H, Quinoline),} \\ 4.91 \text{ (s, 1H, OH), 4.83 (s, 2H, aromatic-CH2-O), 3.55 (m, 2H, O-CH2-), 1.23 (m, 3H, -CH3)} \end{split}$ | v = 3800–2700 (-OH Stretching), 1599 and 3026 (Ar C=C and C-H stretching), 1275–1298 (C-N stretching), 2850, 2940, 1450 (-CH ₂ - stretching) | $\delta_{\text{ppm}} = 15.23, 65.91,$ 77.02, 109.90, 120.30, 126.48, 146.80, 150.37 |

Melting points were determined on an automatic electrothermal IA 9200 digital melting point apparatus in capillary Mp: 78–80 °C. $^1\mathrm{H}$ NMR spectra were recorded on a Bruker 300 WB spectrometer at 300 MHz for solutions in Me₂SO-d₆. Chemical shifts are given as δ values with reference to tetramethylsilane (TMS) as internal standard. Infrared spectra were recorded from 400 to 4000 cm $^{-1}$ on a Bruker IFS 66v Fourier transform spectrometer using KBr pellets. Mass spectrum was recorded on THERMO Electron DSQ II.

2.2 X-ray diffraction

High quality powder X-ray diffraction measurements were performed with an Inel CPS120 powder diffractometer (Debye–Scherrer geometry, transmission mode). The sample was introduced into 0.5 mm diameter Lindemann glass capillary. The latter was rotated around its axis in order to minimize preferential orientations of the crystallites and to obtain reflections with exploitable intensities. The time of acquisition was set to 8 h approximately for a good signal-to-nose ratio. Monochromatic Cu-K α_1 radiation ($\lambda=1.54056$ Å) was selected with asymmetric focusing incident-beam curved quartz monochromator, the generator power was set to 40 kV and 25 mA. The diffracted lines were collected on a 4096 channel detector over an arc of 120° and centered on the sample. External calibration using the Na₂Ca₂Al₂F₁₄ (NAC) cubic phase mixed with silver behenate was performed using cubic spline fittings.

2.3 Theoretical calculation

Geometrical data of the molecule in ground state was optimized by density functional theory (DFT) and Hartree–Fock (HF) methods with 6-311G(d,p) basis set, using Gaussian 03 program package at room temperature (Frisch et al. 2004). These methods are more used, actually, to study chemical and biochemical phenomena within a large variety of molecules (Prasad et al. 2013; Li et al. 2014). Because they can reach exactitude similar to other methods as ab initio in less time and less expensive from the computational point of view, on the other hand, they offers similar accuracy for middle sized and large systems.

3 Results and discussion

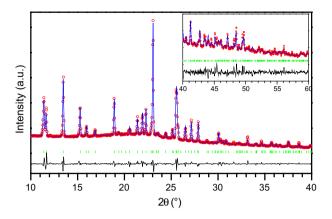
3.1 Crystal structure analysis

The chemical structure of the compound deduced from its IR, 1 H, 13 C and NMR spectra was confirmed by X-ray analysis. Thermal analysis was performed with Perkin-Elmer



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Fig. 1 Experimental (red) and calculated (blue) diffraction patterns along with the difference profile (black) and calculated Bragg reflections (green) of 5-ethoxymethyl-8-hydroxyquinoline at 295 K. The insert is enlarged 15×



DSC-7 apparatus using heating rates of $10~\rm K~min^{-1}$ revealed no solid–solid phase transition between room temperature and the melting point. The melting point temperature (Mp = 351.8 K) was in good agreement with that measured by using digital melting point apparatus (Electrothermal IA 9200) and enthalpy of melting was equal to $\Delta H = 21.136~\rm kJ/mol$.

The crystal structure was determined with materials studio modeling 5.5 (http://accelrys.com/products/materials-studio). First, the pattern was indexed by means of the peak picking option of the software package and using the X-cell algorithm (Neumann 2003). Then, a Pawley profile-fitting procedure (Pawley 1981) was applied including refined cell parameters, experimental profile fitting with pseudo-Voigt function, zero shift and asymmetry Finger-Cox-Jephcoat function. Distances, angles and torsions in the molecule were obtained via energy-minimization calculations using the Dreiding force field (Mayo et al. 1990). Then, a Monte-Carlo approach was carried out in the direct space to solve the structure (Serbezov et al. 2013) moving the molecule as a rigid-body and allowing the change of the chain torsion angles. Rwp agreement factor of 4.78 % is reached after Rietveld refinement (Rietveld 1969). Describing a good agreement between the measured and calculated spectra for 5-EMQ molecule at 295 K (Fig. 1).

The molecular structure is represented in Fig. 2a. Relevant crystallographic data are summarized in Table 2. The parameters in the CIF form are available as Electronic Supplementary Information from the Cambridge Crystallographic Database Centre (1029560).

The molecule exibited non-planar structure, the dihedral angle calculated between the methylquinolin-8-ol group and other part of the molecule is 77.08° (Fig. 2b).

The stabilized crystal packing is mainly governed by hydrogen bonds (Fig. 3) and π - π stacking interactions (Fig. 4). The intramolecular hydrogen bonds observed in the structure involves the OH functional group, this latter presents strong interactions in many solvents, modify physical properties at room temperature (Kadiri et al. 1994; Cazeau-Dubroca 1991) and enhancing the biological activity of the molecule. Weak intra and intermolecular hydrogen bonds O-H···N and C-H···O types are detected in the structure involving



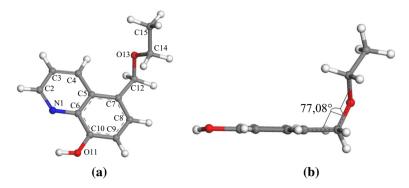


Fig. 2 a Crystal molecular conformation of 5-ethoxymethyl-8-hydroxyquinoline, b geometric display showing the non planar conformation of the molecule

Table 2 Crystallographic data of 5-ethoxymethyl-8hydroxyquinoline

| Empirical formula | C ₁₂ H ₁₃ N O ₂ |
|----------------------------------|--|
| Formula weight | 203.23 |
| Temperature | 295 K |
| Wavelength | 1.54056 Å |
| Structure | Orthorhombic |
| Space group | Pbca |
| a (Å) | 7.9551(12) |
| b (Å) | 17.981(3) |
| c (Å) | 15.125(2) |
| $(\alpha = \beta = \nu)^{\circ}$ | 90.0000 |
| Volume (ų) | 2163.5(6) |
| Z(Z') | 8(1) |
| Density (calculated) | 1.248 g/cm^3 |
| | |

adjacent molecules in different plans, with a gap from 0.638 Å (Fig. 3b). Selected distances (Å) and angles (°) describing the hydrogen bonds are reported in Table 3.

In addition to the hydrogen bond, the crystal packing structure of 5-EMOO shows $\pi-\pi$ stacking interactions between 8-quinolinol rings on adjacent molecules (Fig. 4a). Those contacts may stabilize the structure, with ring centroid to ring centroid distance of 3.473 Å (Fig. 4b). This value is similar to the values reported for other π - π interactions (Shi et al. 2007; Zawadzka et al. 2012).

3.2 Theoretical calculation

Figure 5a shows an optimized calculated structure of the compound. In Table 4 we report the most optimized structural parameters (bond distances, angles and torsions) obtained from crystal structure data (1), optimized by DFT (B3LYP) (2) and by HF (3), in accordance with the atom numbering scheme given in Fig. 2a. The global minimum energy obtained by DFT and HF structure optimization is calculated as -18244.67 and -18131.48 eV respectively. The dipole moment is equal to 3.0245 Debye.



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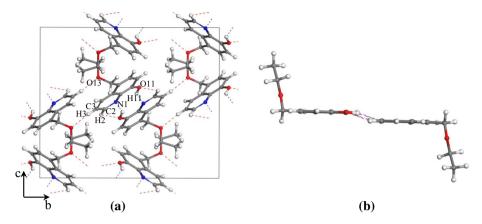


Fig. 3 Hydrogen Bonding contacts (*dashed lines*) in crystal lattice of molecule: **a** along the (bc) plan, **b** geometric display of the hydrogen bonds

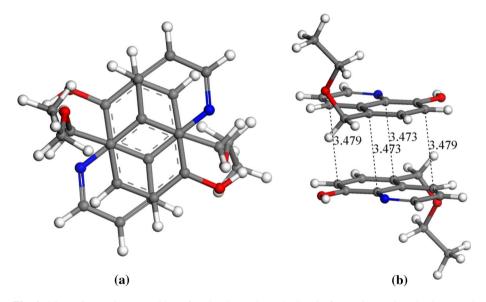


Fig. 4 View of crystal π - π stacking of molecule: **a** along the b axis, **b** π - π interactions in the crystal packing of molecule

The C–C bonds lengths optimized with both methods are in a fairly good agreement with the corresponding bond lengths obtained from X-ray analysis, however slight differences on C–C bond lengths may be noticed between DFT(B3LYP) and the HF calculations. The biggest gap between calculation and experience is observed for C(3)–C(4), C(7)–C(8) and C(9)–C(10) bonds at HF method, values are 0.045, 0.056 and 0.050 Å respectively. For the rest of bond lengths type, the highest difference between crystal structure and calculation is observed for the N(1)–C(2) bond with a value of 0.04 and 0.06 Å for DFT and HF methods respectively. Generally the HF method underestimate



| D | Н | A | D-H | Н…А | D···A | D–H···A | symmetry (A) |
|-----|-----|-----|------------|------------|------------|-----------|--------------|
| O11 | H11 | N1 | 0.977 (11) | 2.508 (10) | 2.794 (9) | 96.6 (7) | (1) |
| O11 | H11 | N1 | 0.977 (11) | 2.053 (9) | 2.905 (9) | 144.6 (7) | (2) |
| C3 | Н3 | O13 | 1.015 (9) | 2.403 (10) | 3.329 (9) | 151.2 (9) | (3) |
| C2 | H2 | 011 | 1.024 (10) | 2.547 (11) | 3.061 (10) | 110.6 (9) | (2) |

Table 3 Intermolecular hydrogen bonding geometries of 5-EMQO (Å, °)

Symmetry applied to the atom A: (1) x, y, z; (2) -x, -y, -z; (3) x-1/2, 1/2-y, -z

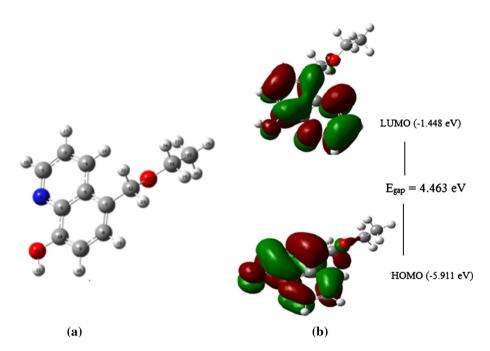


Fig. 5 a Optimized structure (B3LYP/6-311) of molecule; b Molecular orbital surfaces and energy levels for the HOMO, LUMO

some bond lengths by Inclusion of the electron correlation which makes this bond shorter because of the increased resonance in the correlated wave function (Arici et al. 2005; Lakshmi et al. 2011; Zawadzka et al. 2015; Kosar and Albayrak 2011). As a result, the DFT method for bond lengths is much closer to experimental data.

The calculated bond angles values obtained with DFT and HF are very similar, but compared to crystal data some discrepancies appears for C(2)–N(1)–C(6) and C(9)C(10)–O(11) with a difference of 5.28° for DFT method and 5.34° for HF method, respectively. Concerning torsion angles, we presented in Table 3 only those with notable discrepancies between X-ray data and the two calculations (DFT and HF) methods. The Azine ring appears a little distorted with smaller C(6)–N(1)–C(2)–C(3), C(2)–N(1)–C(6)–C(5), N(1)–C(2)–C(3)–C(4) torsion angles and bigger C(2)–C(3)–C(4)–C(5), C(3)–C(4)–C(5)–C(6),



Table 4 Principal geometry parameters for the 5-ethoxymethyl-8-hydroxyquinoline, (1) from X-ray data analysis, (2) calculated by the DFT/B3LYP/6-311G(d,p), (3) calculated by HF/6-311G(d,p)

| Parameters | (1) | (2) | (3) |
|---------------------|------------|--------|--------|
| Bonds (Å) | | | |
| N(1)–C(2) | 1.355 (10) | 1.314 | 1.290 |
| N(1)-C(6) | 1.360 (7) | 1.357 | 1.349 |
| C(2)–C(3) | 1.403 (10) | 1.412 | 1.412 |
| C(3)–C(4) | 1.401 (9) | 1.372 | 1.356 |
| C(4)–C(5) | 1.416 (6) | 1.418 | 1.418 |
| C(5)–C(6) | 1.425 (6) | 1.429 | 1.404 |
| C(5)–C(7) | 1.428 (7) | 1.429 | 1.432 |
| C(6)-C(10) | 1.403 (9) | 1.433 | 1.432 |
| C(7)–C(8) | 1.410 (9) | 1.375 | 1.354 |
| C(7)–C(12) | 1.487 (8) | 1.505 | 1.506 |
| C(8)–C(9) | 1.398 (10) | 1.410 | 1.415 |
| C(9)-C(10) | 1.406 (5) | 1.377 | 1.356 |
| C(10)-O(11) | 1.354 (9) | 1.355 | 1.340 |
| C(12)-O(13) | 1.439 (11) | 1.424 | 1.397 |
| O(13)-C(14) | 1.434 (10) | 1.420 | 1.397 |
| C(14)–C(15) | 1.535 (14) | 1.517 | 1.514 |
| Angles (°) | | | |
| C(2)-N(1)-C(6) | 123.3 (5) | 118.02 | 118.58 |
| N(1)-C(2)-C(3) | 119.0 (5) | 123.74 | 123.54 |
| C(2)-C(3)-C(4) | 119.7 (5) | 118.84 | 118.60 |
| C(3)–C(4)–C(5) | 120.7 (5) | 119.79 | 119.69 |
| C(4)-C(5)-C(6) | 117.2 (4) | 116.38 | 116.43 |
| C(4)-C(5)-C(7) | 123.6 (5) | 123.17 | 123.06 |
| C(6)–C(5)–C(7) | 119.3 (4) | 120.43 | 120.50 |
| N(1)-C(6)-C(5) | 120.1 (4) | 123.19 | 123.13 |
| N(1)-C(6)-C(10) | 119.6 (5) | 117.98 | 117.84 |
| C(5)-C(6)-C(10) | 120.3 (4) | 118.81 | 119.01 |
| C(5)–C(7)–C(8) | 119.1 (5) | 118.28 | 118.16 |
| C(5)–C(7)–C(12) | 122.5 (6) | 121.44 | 121.45 |
| C(8)–C(7)–C(12) | 118.4 (5) | 120.25 | 120.36 |
| C (7)–C(8)–C(9) | 121.0(5) | 122.14 | 122.10 |
| C (8)–C(9)–C(10) | 120.3 (5) | 120.69 | 120.62 |
| C (6)–C(10)–C(9) | 120.0 (5) | 119.62 | 119.58 |
| C (6)–C(10)–O(11) | 121.9 (5) | 117.04 | 116.97 |
| C (9)–C(10)–O(11) | 118.1(5) | 123.33 | 123.44 |
| C (7)–C(12)–O(13) | 112.2(5) | 110.48 | 110.56 |
| C (12)–O(13)–C(14) | 116.9 (7) | 112.89 | 114.35 |
| O(13)-C(14)-C(15) | 108.9 (8) | 108.48 | 108.77 |
| Torsions (°) | | | |
| C(6)-N(1)-C(2)-C(3) | 0.7 (9) | 0.33 | 0.211 |
| C(2)-N(1)-C(6)-C(5) | -0.4(7) | 0.07 | 0.02 |



| Table 4 | continued |
|---------|-----------|
|---------|-----------|

| Parameters | (1) | (2) | (3) | |
|------------------------|-----------|----------|---------|--|
| N(1)-C(2)-C(3)-C(4) | -0.5 (9) | -0.15 | 0.01 | |
| C(2)-C(3)-C(4)-C(5) | -0.1(9) | -0.44 | -0.45 | |
| C(3)-C(4)-C(5)-C(6) | 0.3 (7) | 0.78 | 0.64 | |
| C(4)-C(5)-C(6)-N(1) | 0.3 (7) | -0.62 | -0.44 | |
| C(5)-C(7)-C(12)-O(13) | 62.7 (8) | -65.24 | -65.28 | |
| C(7)-C(12)-O(13)-C(14) | 77.1 (13) | -178. 63 | -178.44 | |

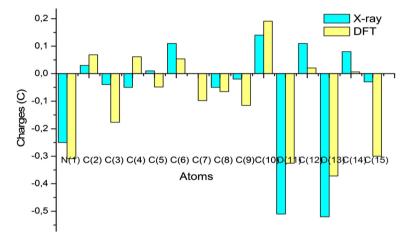


Fig. 6 The atomic charges for 5-ethoxymethyl-8-hydroxyquinoline

C(4)–C(5)–C(6)–N(1) torsion angles (Table 4), comparing values from DFT and HF methods to the crystal data. The highest difference corresponds to the C(7)–C(12)–O(13)–C(14) torsion angle with the value of 77.1(13)° in the crystal structure and −178.6° and −178.4° respectively for DFT and HF calculations. The discrepancies between calculated and experimental structural parameters are related to the fact, that calculations refer to the isolated molecule, contrarily experimental results which correspond to interacting molecules in the crystal lattice where intra and intermolecular interactions with the neighboring molecules are present (Kavitha et al. 2010; Ghiasi et al. 2014; Pournamdari et al. 2012).

Highest occupied molecular orbital (HOMO) and lowest unoccupied molecular orbital (LUMO) are been computed using B3LYP level 6-311(d,p) of DFT. Figure 5b shows the patterns of the HOMO and LUMO of 5-EMQO. The HOMO–LUMO gap is 4.463 eV. The HOMO energy shows the ability of giving an electron, whereas, the LUMO energy represents the ability of gaining an electron (Pournamdari et al. 2012). In many recent works, the energy gap between HOMO and LUMO has been used to prove the bioactivity from intra molecular charge transfer and to explain the chemical reactivity of the molecule (Fukui 1982; Gunasekaran et al. 2008).

The charge distribution on different atoms (C, N and O) for molecule obtained by using X-ray analysis data B3LYP methods and HF methodes is plotted in Fig. 6. Results



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obtained by these two methods are similar. This distribution depending on the electronegativity of atoms; the most negative are those present at nitrogen and oxygen atoms. The values obtained from DFT calculations are in fairly agreement with crystal structure. The most negative values are those that are present at nitrogen and oxygen: N1, O11 and O13. It has to be noticed that the carbon atom attached to oxygen and nitrogen atoms: C2, C6, C10, C12 and C14 in both X-ray and B3LYP are electron deficient atoms possessing positive electronic charge due to the oxygen and nitrogen electronegativity. Remaining carbon atoms possess negative electronic charge. Except some discrepancies about C4 and C5 which possess a different electronic charge in DFT calculation, this deviation is probably due to the fact that the intermolecular hydrogen bonding interaction C3-H3-O13 in the crystalline state are dominant influencing the charge distribution in C4 and C5. Furthermore, the DFT calculation refers to the isolated molecule where the intermolecular interactions with the neighboring molecules are absent. This distribution within the molecule and the high dipole moment explain the notable polarization given by the small HOMO-LUMO energy calculated. The molecule will exhibit a significant degree of intramolecular charge transfer from the electron donor section to the electron acceptor section.

4 Conclusion

In this work, we have synthesized 5-ethoxymethyl-8-hydroxyquinoline and determined its crystal structure by X-ray powder diffraction pattern. The crystal system is orthorhombic with Pbca space group and eight molecules per unit cell. The lattice parameters are: a=7.9551(12) Å, b=17.981(3) Å, c=15.125(2) Å. The crystal structure is mainly governed by intra and intermolecular O–H···N and C–H···O hydrogen-bonding type and π – π stacking interactions. Density functional theory DFT(B3LYP) and Hartree–Fock (HF) calculations were compared with the measured data, some discrepancies appears, the big one occurs in the C(7)–C(12)–O(13)–C(14) torsion angle, showing the orientation of the O–C₂H₅ group in the opposite direction to that given by X-ray analysis. Furthermore, HOMO–LUMO patterns and Mulliken atomic charges have been analyzed by DFT to explain the possibility of charge transfer within the molecule. The small obtained gap and atomic charges distribution reveled possibility of the charge transfer within the molecule and a high reactivity of 5-ethoxymethyl-8-hydroxyquinoline, who permits to this type of molecule to be used as bioactive molecule.

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