



Synthesis, NoSpherA2 refinement, and noncovalent bonding of abiraterone bromide monohydrate

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Abstract

Abiraterone (AbirOH) is an active metabolite of abiraterone acetate (AbirAc) used as a therapeutic agent against prostate cancer. A variety of its solid forms includes polymorphs of pure abiraterone, abiraterone acetate, and cocrystals of abiraterone acetate with carboxylic acids, and a pyridinium-containing (HAbirOH)⁺ salt as well. Herein, we present the crystal structure of (HAbirOH)Br · H₂O as obtained by means of NoSpherA2 refinement applied to reveal charge density distribution for abiraterone from routine single-crystal X-ray diffraction experiment. Peculiarities of noncovalent interactions in this salt are discussed in terms of molecular Voronoi and Hirshfeld surfaces. The AbirOH packing in this salt and in previously reported solids is analyzed by means of the “Crystal Packing Similarity” tool.

Keywords Abiraterone · Active pharmaceutical ingredient · Crystal Packing Similarity · Hirshfeld molecular surface · NoSphereA2 refinement · Single-crystal X-ray diffraction · Voronoi molecular surface

Introduction

Abiraterone (AbirOH) is an active metabolite of the commercially available drug *Zytiga* (abiraterone acetate, AbirAc) used to treat prostate cancer. According to the results provided by macromolecular crystallography, nitrogen atom can coordinate heme fragment of androgen receptors. A detailed study of the ligand-receptor binding that was carried out on the base of combined study utilizing experimental charge

density and Voronoi tessellation [1] as main methods has shown that weak hydrophobic H...H interactions between steroid fragment and amino acid residues are also very important for retention of abiraterone in cavity of receptor formed by peptide chains and heme fragment. Mainly the hydrophobic molecular surface of AbirAc results in its poor solubility; thus, by cocrystallization of AbirAc with some cofomers, novel solid forms of AbirAc with enhanced solubility were obtained [2, 3]. Acidic media hydrolysis of AbirAc [4] and protonation of the pyridine moiety can easily occur, which also allows to afford new solid forms of abiraterone for pharmaceuticals [5]. Herein, we present the structure of (HAbirOH)Br · H₂O (17-(pyridinium-3-yl)androsta-5,16-dien-3-ol bromide monohydrate, **1**; Scheme 1) that was found to be isostructural with previously studied chloride-containing analog [5].

Solid (HAbirOH)Cl · H₂O and (HAbirOH)Br · H₂O give an opportunity to obtain charge distribution in the region of abiraterone that can be of interest for biochemists. However, the chloride-containing salt was investigated at 298 and 100 K [5, 6] using a standard independent atom model only. We also failed to obtain a single crystal of the bromide-containing analog suitable for high-resolution X-ray diffraction experiment. Nevertheless, data collected by us allow application of aspherical structure factors [7] to obtain wavefunction related to unique parts of the unit cell. This

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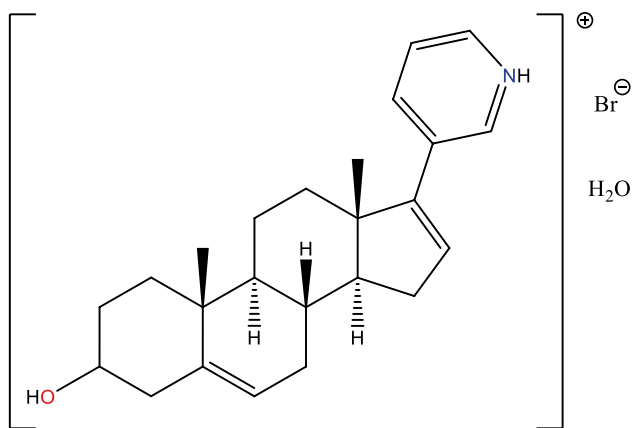
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Scheme 1 Schematic representation of 17-(pyridinium-3-yl)androsta-5,16-dien-3-ol bromide monohydrate

approach is suitable for the evaluation of many properties of small molecules and their solids, for instance, low-frequency lattice vibrations [8], charge density and NCI analysis [9, 10], and proton migration study of strong H-bonds [11]. To our knowledge, data about charge distribution in solids of active pharmaceutical ingredients obtained by means of the NoSphereA2 refinement are limited by recent work concerning crystal structure of pure imatinib [12].

Experimental

Solution of abiraterone acetate (0.012 g, 0.05 mmol) in 1 ml of ethanol was added to a solution of FeBr_3 (0.087 g, 0.5 mmol) in 2 ml of ethanol. Mixture was heated and solution was cooled on air. After 2 days of standing at RT, white precipitate formed. Single crystals of $\text{C}_{24}\text{H}_{34}\text{BrNO}_2$ were obtained from this precipitate. ^1H and ^{13}C NMR spectra were recorded on 400 MHz and 101 spectrometers in DMSO-d_6 solution. The powder X-ray diffraction patterns were obtained in reflection mode. The measurements were performed with a Bruker D8 Advance diffractometer (Bragg–Brentano geometry) equipped with motorized slits and a LynxEye 1D position-sensitive detector (CuK α , Ni-filter) in the range $2\theta = 3\text{--}50^\circ$. FTIR spectra were recorded on a Perkin Elmer Spectrum 65 spectrometer in the range of 400–4000 cm^{-1} .

^1H NMR (400 MHz, DMSO-d_6) δ : 0.92–1.08 (m, 2H, CH), 0.99 (s, 3H, CH_3), 1.03 (s, 3H, CH_3), 1.27–2.34 (m, 18H, other CH and CH_2), 3.19–3.31 (m (seems like quintet, $J = 5.3$ Hz), 1H, CH(OH)), 4.30 (ws, OH, H $^+$, H_2O), 5.30 (d, $J = 5.0$ Hz, 1H, CH=C), 6.46 (s, 1H, CH=C), 7.93 (t, $J = 7.0$ Hz, 1H, Ar), 8.49 (d, $J = 8.2$ Hz, 1H, Ar), 8.73 (d, $J = 5.6$ Hz, 1H, Ar), 8.86 (s, 1H, Ar). ^{13}C NMR (101 MHz,

DMSO-d_6) δ : 16.39, 19.53, 20.80, 30.32, 31.33, 31.84, 32.03, 34.47, 36.71, 37.29, 42.68, 47.15, 50.17, 57.45, 70.42, 120.58, 127.05, 134.33, 135.36, 140.48, 141.37, 141.75, 142.10, 148.74. IR-spectrum (FTIR) (cm^{-1}): 3328 s, 2960 m, 2909 s, 2889 s, 1550 s, 1057 s, 804 s, 684 s.

The intensities of 10,124 reflections were collected at “Belok/XSA” beamline of the Kurchatov Synchrotron Radiation Source [13, 14]. Diffraction patterns were collected using a 1-axis MarDTB goniometer equipped with a Rayonix SX165 CCD 2D positional sensitive CCD detector ($\lambda = 0.745$ Å, φ -scanning in 1.0° steps) in the direct geometry with a detector plane perpendicular to its beam. Approximately 180 diffraction frames were collected for each data set. Thus, obtained data were indexed and integrated using the XDS software suite [15]. At 100 K, crystal system is monoclinic, space group $P2_1$; $a = 6.6520(13)$, $b = 11.136(2)$, $c = 14.513(3)$ Å, $\alpha = 90$, $\beta = 94.25(3)$, $\gamma = 90^\circ$, $V = 1072.1(4)$ Å 3 , $Z = 2$, $\mu = 2.178$ mm^{-1} , $D_{\text{calc}} = 1.389$ g cm^{-3} , $F(000) = 471.2$.

The structure was solved by the dual-space algorithm [16] and initially refined by full-matrix least squares against F^2 using the IAM model (SHELXL program [17]) to $R_1 = 0.0376$ and $wR_2 = 0.0949$, $\rho_{\text{min}}/\rho_{\text{max}} = -0.93/0.87$ e. Non-hydrogen atoms were refined in an anisotropic approximation; anharmonic refinement for bromine atom was applied. Hydrogen atoms were located on difference Fourier maps and refined with unfixed bond distances; their ADPs were calculated with SHADE3 [18] and fixed at the calculated values. The application of the NoSphereA2 algorithm [7] implemented within the Olex2 package [19] for subsequent refinement led to a noticeable decrease of residual factors. Refinement converged to $R_1 = 0.0295$ (for 5468 observed reflections and 382 parameters), $wR_2 = 0.0693$, and $\text{GOF} = 1.039$ (for 5852 independent reflections, $R_{\text{int}} = 0.0338$), $\rho_{\text{min}}/\rho_{\text{max}} = -0.31/0.51$ e.

Peculiarities of the Voronoi molecular polyhedra were calculated using the ToposPro package [20]. The Hirshfeld surface analysis was performed by the CrystalExplorer 17.5 package [21]. Analysis of electron density function and visualization of results were carried out with Multiwfn [22] and VMD software [23].

Results and discussion

In our study of novel solid forms of known active pharmaceutical ingredients [12, 24], the ability of abiraterone acetate to form salts with various metals was examined. Solution of abiraterone acetate in ethanol was added to a solution of FeBr_3 in equimolar ratio (1:1). After several days of standing on air at RT, red precipitate formed.

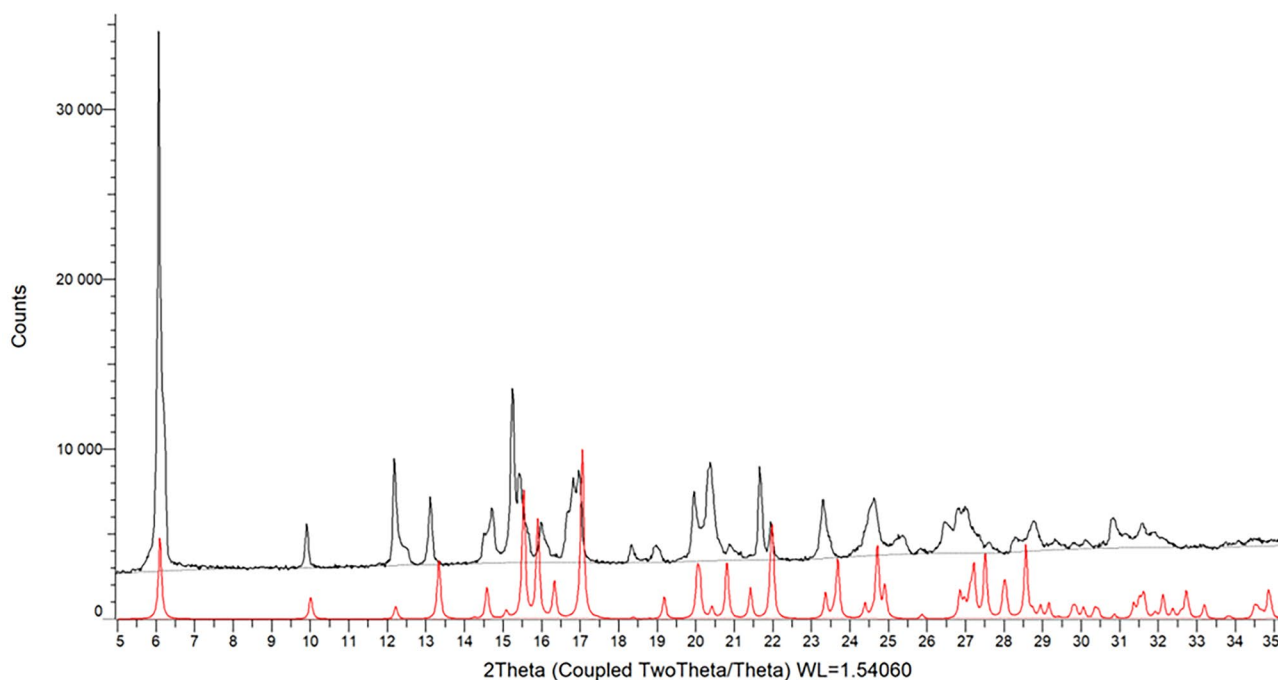


Fig. 1 PXRD patterns of (HAbirOH)Br · H₂O at RT (red line—calculated profile; black line—experimental profile) in 2 Ω range 5 – 35°

The red color of precipitate can be easily explained by the formation of colloid Fe(OH)₃. At the same time, small colorless crystals were also found in a microscopic study. The phase purity of the colorless phase was checked using XRPD. No significant differences in peak positions and intensities between experimental and calculated patterns were found (Fig. 1). Subsequently, the synchrotron X-ray diffraction study has revealed that colorless crystals correspond to salt **1**. Thus, it was proved that abiraterone acetate can be hydrolyzed in acidic media at presence of iron(III) salt, although no complexation with metal atom occurred. Note also that recrystallization of AbirAc from 2 M and concentrated 48% HBr affords, respectively, starting AbirAc or an amorphous product as obtained from powder X-ray diffraction.

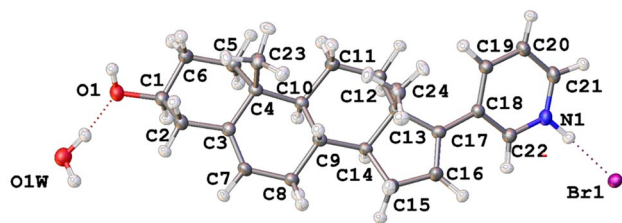


Fig. 2 Molecular view of **1** in representation of atoms with thermal ellipsoids ($p=50\%$)

Molecular and crystal structure of **1**

The unit cell of (HAbirOH)Br · H₂O is represented in Fig. 2. The positions of all hydrogen atoms can be easily revealed from difference Fourier maps and the most expected H-bonds. These should be present for O1...Br1, O1W...Br, and O1W...O1 distances as short as 3.245(2), 3.356(2), and 2.778(3) Å. In addition, the protonation of pyridine fragment is expected from N1...Br1 distance as short as 3.211(2) Å (Table 1). Thus, the positions of all hydrogen atoms were revealed on difference Fourier maps and refined with unfixed bond distances, while ADPs of hydrogen atoms were calculated with SHADE3 web application [18] and fixed at the calculated values as it was previously suggested by Woińska et al. [25]. N1-H1A, O1-H1, and O1W-H

Table 1 H-bond lengths (Å, °) for salt **1**

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1A...Br1	1.04(4)	2.16(4)	3.211(2)	178(4)
O1—H1...Br1 ⁱ	0.93(4)	2.34(4)	3.245(2)	165(3)
O1W—H1WA...O1	1.02(4)	1.77(4)	2.778(3)	172(4)
O1W—H1WB...Br1 ⁱⁱ	0.97(3)	2.39(3)	3.356(2)	174(4)
C16—H16...O1W ⁱⁱⁱ	1.09(3)	2.27(3)	3.325(3)	167(3)

Symmetry transformations used: (i) $1-x, -1/2+y, 1-z$; (ii) $1+x, y, -1+z$; (iii) $x, y, 1+z$

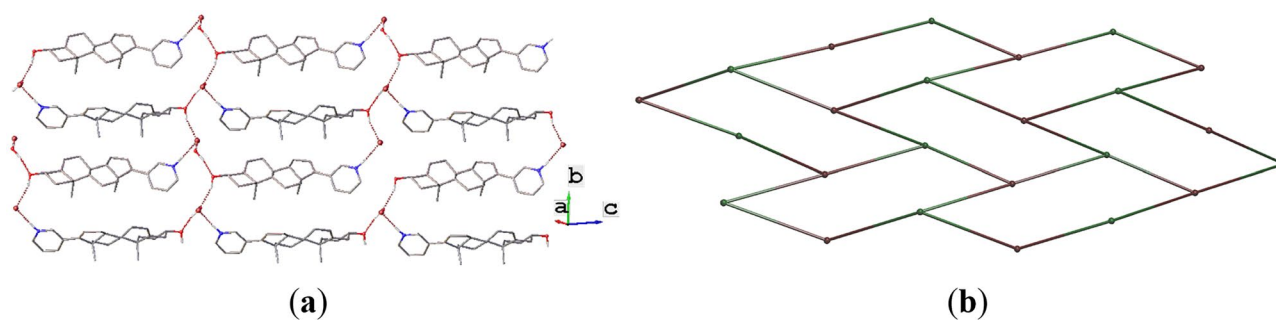


Fig. 3 **a** Fragment of H-bonded layers in (HAbirOH)Br · H₂O, and **b** underlying H-bonded net of salt **1** (centers of cations and anions are depicted as brown and green spheres, respectively, keeping connectivity of the network)

distances (1.04(4), 0.93(4), and 0.97(3)—1.02(4) Å) refined with NoSpherA2 [7] are close to the values obtained from neutron diffraction studies (1.05, 0.98, and 0.97 [26, 27]).

Strong hydrogen bonding in solid **1** results in the formation of H-bonded layers. Parameters of H-bonds are listed in Table 1, and the fragment of the layer is depicted in Fig. 3. The bromide anions and the cations both act as three-connected nodes of the layer (these ions take part in three hydrogen bonds), and water molecules are two-connected nodes or simple linkers (weak C–H...O bonding is not considered). The resulting H-bonded network obtained by simplification of cations, anions, and water molecules to their centers followed by removal of simple linkers as described in Ref. [28] is given in Fig. 3b. It has the honeycomb topology which is one of the most widespread among H-bonded nets of molecular crystals [29].

Packing similarity of **1** and previously reported forms of abiraterone

In several previously reported X-ray studies of solid forms of AbirOH, AbirAc, and (HAbirOH)⁺, it was shown that the

character of intermolecular pairwise interactions is almost independent from the presence of the acetate group or protonation of the pyridine fragment. Particularly, high (up to 82%) contribution of H...H interactions to the Voronoi molecular surface was found [1]. This quantity and the steric repulsion of methyl groups from one another are favorable for retaining the conformation of steroid core unchanged. The crystal packing in salt **1** is very similar to its isostructural chloride-containing analog; both compounds form H-bonded layers (Fig. 4a); moreover, the “Crystal Packing Similarity” tool described in Refs. [30, 31] demonstrates that clusters composed of 20 HAbirOH⁺ cations in these two salts are also close to each other (Fig. 4a) demonstrating similarity of layers packing through hydrophobic interactions. The role of hydrophobic interactions in packing of abiraterone and its derivatives can be highlighted also by the fact that the “Crystal Packing Similarity” tool also revealed similarly packed chains through H...H interaction molecules in (HAbirOH)Br · H₂O **1** and AbirAc · *p*-HOC₆H₄COOH [2], AbirAc · HOOC(CH₂)₃COOH [2], AbirAc · TAA (TAA = *trans*-aconitic acid [3]), and pure AbirOH (polymorph II [5]; Fig. 4b).

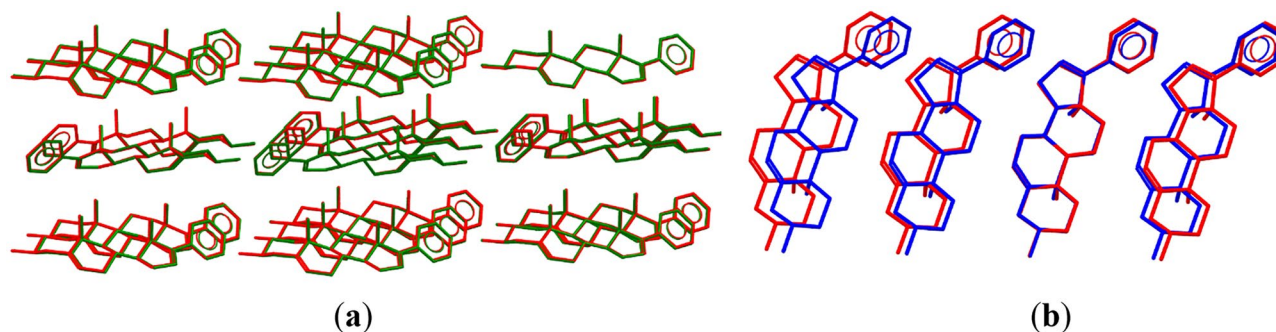
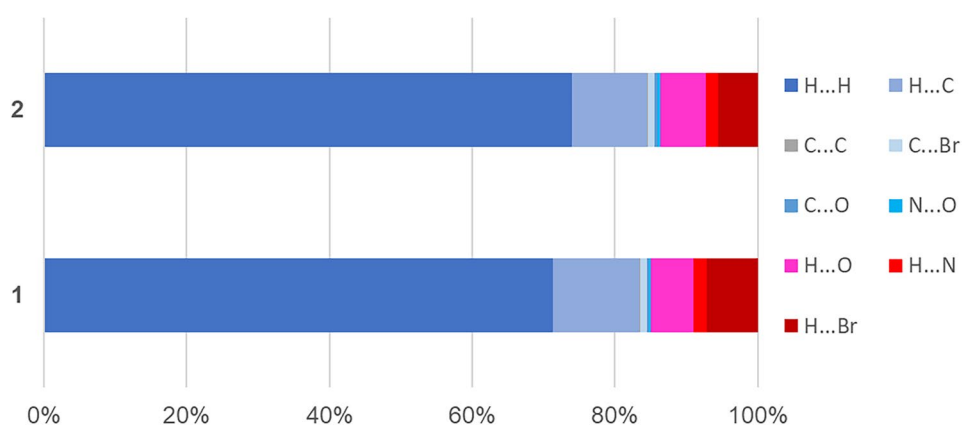


Fig. 4 **a** Comparison of molecular clusters in (HAbirOH)Br · H₂O (red) and (HAbirOH)Cl · H₂O (green); and **b** H...H connected chains in (HAbirOH)Br · H₂O (red) and form II of pure AbirOH (blue)

Fig. 5 Contribution of various noncovalent interactions to the molecular Voronoi (1) and Hirshfeld (2) surfaces



Contribution of various types of interactions to the molecular surface of abiraterone

The molecular Hirshfeld [32] and Voronoi [33] surfaces are well-known approaches for analysis of noncovalent interactions in crystals of molecular compounds. Although these are mainly applied to monomolecular compounds, examples of their application to salts and solvates are also known [1, 34–37]. We performed the analysis of partial contributions of various types of noncovalent interactions to the surrounding of the HAbirOH⁺ in solid **1** by means of the Voronoi and Hirshfeld molecular surfaces. These approaches give the area of the molecular surface for the cation equal to, respectively, 434.5 and 370.3 Å². Solid **1** contains atoms of five element; hence, 15 different types of noncovalent interactions can theoretically be found;

however, both methods detect nine types only (Fig. 5). Six of them (H...H, H...C, C...C, C...Br, C...O, and N...O) belong to the hydrophobic interactions, and three (H...O, H...N, and H...Br) to hydrophilic, and both approaches indicate that 85% of the molecular surface goes to hydrophobic interactions. This fact is in accord with previously reported data for abiraterone-containing solids [1]. Good correlation between partial contributions of noncovalent interactions to the molecular Voronoi and Hirshfeld surfaces is also typical for molecular solids [38–40]. Difference of molecular volume for two approaches is less pronounced; total volume of the cation is equal to 488.3 and 464.4 Å³ for the Voronoi and Hirshfeld surfaces.

Both surfaces are depicted in Fig. 6. Note that color schemes for the surfaces have different advantages and disadvantages for analysis of noncovalent interactions.

Fig. 6 The molecular Voronoi surface of HAbirOH⁺ colored in accord with the nature of **a** internal and **b** external atoms (Br, green; C, dark gray; H, gray; N, blue; O, red). The molecular Hirshfeld surface of HAbirOH⁺ colored with the **c** d_{norm} and **d** curvedness

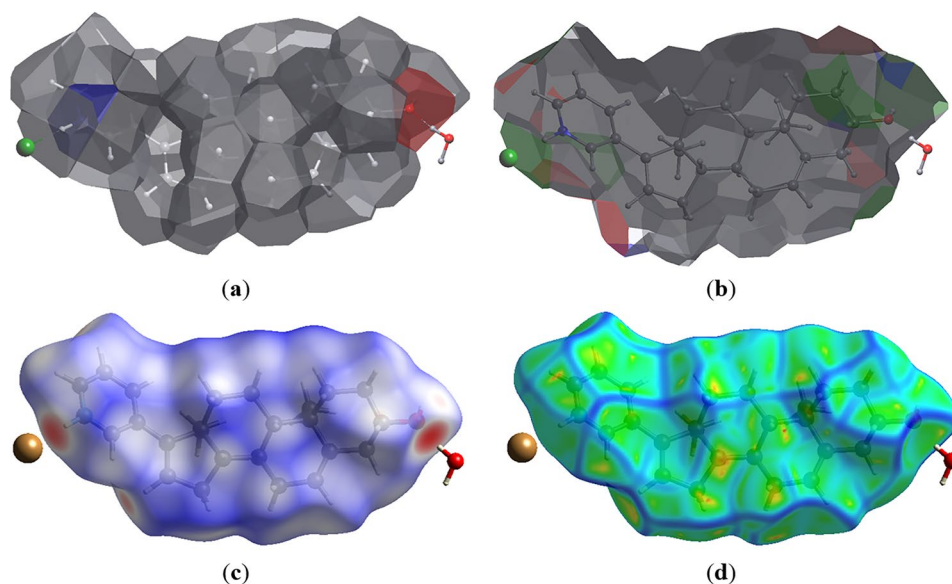
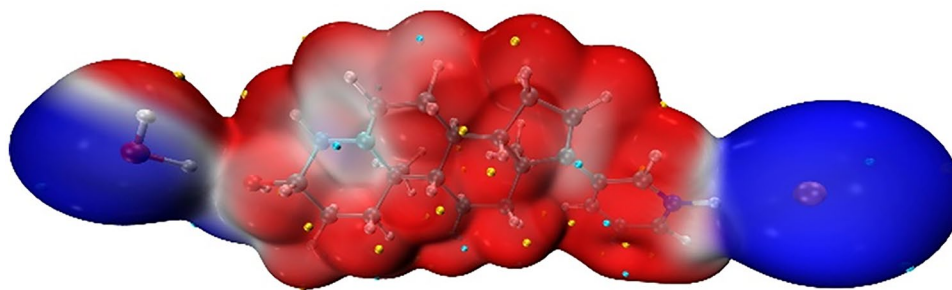


Fig. 7 Electrostatic potential function ranged from -0.03 a.u. (blue) to 0.03 a.u. (red) mapped on 0.001 isosurface of electron density function



The Voronoi surface allows differentiation between atom types external and internal towards the surface, and gives possibility to extract the surface corresponding towards any pairwise interactions. The Hirshfeld surfaces allow revealing the closest interactions, molecular curvedness, and shape index. Particularly, the N–H...Br bond manifests itself as a red spot on the Hirshfeld surface colored with d_{norm} , and as a region colored with gray and green on the Voronoi surface colored with the nature of an internal and external atoms (Fig. 6a, b, respectively).

Charge density distribution in **1**

The wavefunction obtained in NoSpherA2 refinement was utilized to rationalize the information obtained from the analysis of geometry and short contacts. The function of electrostatic potential mapped on 0.001 a.u. demonstrates the character of charge distribution in the unique part of the unit cell of **1**. Bromine atoms, water molecule, hydroxyl group, and double bond C3–C7 appeared to be the regions of negative charge concentration as compared to steroid core that is mostly positively charged (Fig. 7). The comparative analysis of electron density distribution in terms of the QTAIM theory [41] and NCI method [42, 43] has

revealed that hydrogen atoms in the steroid core are involved in intramolecular interactions that can be classified as steric repulsion. In most cases, the presence of isosurfaces of RDG function [42, 43] colored by brown and green colors is indicative for steric interactions between hydrogen atoms. Besides, the occurrence of bond critical points (bcps) is also observed (Fig. 8a) for three H...H interactions. Recently, we proposed that bonding H...H interactions can be extracted from all H...H contacts based on atomic Voronoi polyhedra. Particularly, such interactions should be characterized by solid angles Ω of corresponding faces above 10% of full solid angle (4π steradian) [44]. Among 72 faces of atomic Voronoi polyhedra which correspond to intramolecular H...H interactions, only 32 and 5 faces are characterized by $\Omega > 10$ and 15%, respectively. Particularly, the interactions connected by bond paths in Fig. 8a are characterized by $\Omega = 17, 16,$ and 16% (Fig. 8b).

Unlike H...H interactions, the N1–H1A...Br1 and O1W–H1WA...O1 have mostly attractive nature. The energies of H1A...Br1 and H1WA...O1 bonds calculated using Espinosa, Mollins, and Lecomte correlation [45] are equal to -30.2 and -45.2 kJ/mol that correspond to weak hydrogen bonds. The energies of three H...H interactions are much weaker than hydrogen bonds (5.4 – 10.1 kJ/mol).

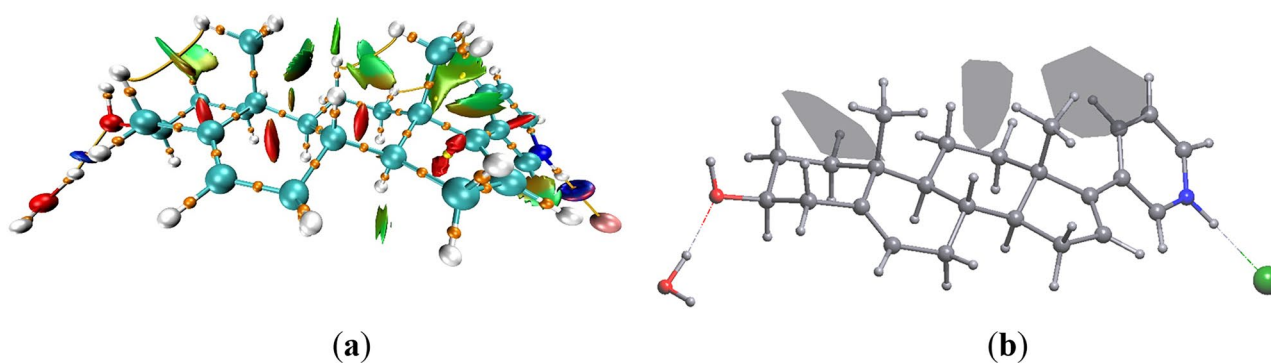


Fig. 8 **a** Molecular graph and RDG isosurface (0.4 a.u.) colored with $\text{sign}\lambda_2$ function and **b** faces of atomic Voronoi polyhedra which correspond to intramolecular H...H interactions (gray). The bcps and bond paths are shown by brown spheres and brown lines, respectively

Conclusions

To sum up, the complexes of abiraterone or abiraterone acetate with Fe(III) salts cannot be easily synthesized in polar media. Hydrolysis of abiraterone acetate to abiraterone and its further protonation was observed instead. The character of crystal packing is governed by combination of O-H...O, O-H...Br, and N-H...Br hydrogen bonds and hydrophobic H...H interactions. The hydrogen bonding between HAbirOH⁺ and anion or water molecule is too weak to affect molecular conformation. Large molecular surface formed by protons and rigid molecular conformation allow crystallization of H...H connected chains in solids, containing AbirAc, AbirOH, and HAbirOH⁺. Good correlation of partial contributions of noncovalent interactions to the molecular Voronoi and Hirshfeld surfaces of the cation was observed.

Supplementary Information The online version contains supplementary material available at <https://doi.org/10.1007/s11224-023-02210-3>.

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Author contribution A.A.K. and A.V.V. wrote the main manuscript text, P.A.B. prepared the sample and obtained spectral data, P.V.D. collected XRD data and performed an ordinary refinement, A.A.K. performed the NoSphereA2 refinement, and A.V.V. prepared all figures and collected powder XRD data. All authors read and approved the final manuscript.

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Availability of data and materials CCDC 2222682 contains crystallographic information for this salt and can be obtained from ccdc.cam.ac.uk/structures.

Declarations

Ethical approval Not applicable.

Competing interests A.V.V. declares that she is an Editorial Board Member of Structural Chemistry. The other co-authors declare that they have no competing interests.

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