## Evidence for O-H--C and N-H--C Hydrogen Bonding in Crystalline Alkynes, Alkenes, and Aromatics

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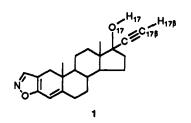
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Abstract: Attempts have been made to understand the nature and significance of hydrogen bonds of the type X-H--C (X = O, N). These unusual interactions have been discussed recently. Crystallographic studies on  $17\alpha$ -ethynylandrosta-2,4-dieno [2,3-d] dihydroxazol-17 $\beta$ -ol (donazole) provide direct evidence of such an O-H...C interaction. Ab initio computations, IR spectroscopy, and database studies show that these hydrogen bonds, while uncommon, are energetically and structurally significant.

Unusual hydrogen bonds of the type  $X-H\cdots C$  (X = O, N) have been discussed recently.<sup>1-7</sup> While C-H-O hydrogen bonds are widespread,<sup>8-10</sup> the complementary O-H…C interaction is rare because C is not as electronegative as O and also because C atoms are not often situated in sterically-unhindered positions (unlike carbonyl and ethereal O atoms which permit easy access by C-H groups to form C-H···O bonds). This report shows that, in spite of these limitations, a sufficiently electron-rich C atom (alkyne, alkene, aromatic) has a propensity to form a hydrogenbond-like interaction with X-H groups (X = O, N) thus contributing to crystal stability. This interaction has been referred to as a  $\pi$ -hydrogen bond.<sup>3</sup>

A striking example of such an O-H…C hydrogen bond is found in the crystal structure of  $17\alpha$ -ethynylandrosta-2,4-dieno[2,3d]dihydroxazol-17 $\beta$ -ol (donazole, 1) (Figure 1:  $P2_12_12_1$ , Z = 4;



a = 6.595, b = 10.519, c = 26.166 Å;  $R = 0.036, R_w = 0.034$ ; 1500 observed reflections; C, N, O anisotropic; all H-atoms located from difference maps and refined isotropically). The  $17\beta$ hydroxyl H atom is directed toward the  $C \equiv C$  bond of a

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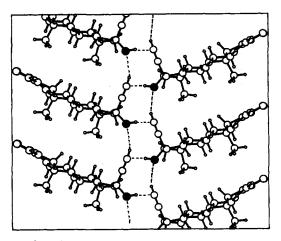


Figure 1. Crystal structure of donazole (1) down [010] to show the O-H···C and C-H···O bonds between the 178-hydroxyl and ethynyl groups. Notice the cooperative network of hydrogen bonds. The a-axis is vertical in the plane of the paper. O atoms are shaded.

neighboring 2<sub>1</sub>-related molecule  $(O(17) - C(17\beta) = 3.25$  Å;  $H(17)\cdots C(17\beta) = 2.45 \text{ Å}; O(17) - H(17)\cdots C(17\beta) = 144^{\circ}).$  This intermolecular interaction therefore has all the geometrical attributes of a hydrogen bond.<sup>8,11</sup> Additionally, the alkyne C-H forms a conventional C-H-O hydrogen bond to the 17-hydroxyl group of another 2<sub>1</sub>-screw related molecule (C(17 $\beta$ )...O(17) = 3.19 Å;  $H(17\beta) \cdots O(17) = 2.45$  Å;  $C(17\beta) - H(17\beta) \cdots O(17) =$ 145°). These O-H-C and C-H-O hydrogen bonds assist each other cooperatively and constitute the dominant motif in the crystal packing. Evidence for the O-H-C interaction in 1 is also available from IR data. While the O-H and C-H (alkyne) stretches appear at 3565 and 3290 cm<sup>-1</sup> (CH<sub>2</sub>Cl<sub>2</sub>), these values are red shifted by 75 and 50  $cm^{-1}$  in the solid state (KBr).

Ab initio calculations (GAUSSIAN80, 3-21G) on the model system, methylacetylene-methanol, show that the O-H…C interaction is attractive in nature. The computed total energies for CH<sub>3</sub>-C=C-H and CH<sub>3</sub>-OH respectively are -115.224 962 1 and -114.397 187 1 au. For an optimized H...C distance of 2.66 Å, the energy of the complex was found to be -229.625 004 3 au. The attractive force between the two species is therefore equivalent to -0.002 855 1 au or -1.79 kcal/mol. The calculation also shows that the H…C=C angle lies ideally between

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Table I. Results of Database Analysis of Intermolecular X-H--C Hydrogen Bonds

|                         | no. of<br>entries with | no. of                        | mean         |           |                  |
|-------------------------|------------------------|-------------------------------|--------------|-----------|------------------|
| X–H···C                 | X–H and $\pi$ -groups  | X–H•••C<br>bonds <sup>a</sup> | Х•••С<br>(Å) | HC<br>(Å) | X-H···C<br>(deg) |
| 0—H…C≡C                 | 154                    | 5°                            | 3.33         | 2.69      | 127              |
| N−H…C≡C                 | 68                     | 5 <sup>d</sup>                | 3.30         | 2.61      | 129              |
| 0—н…С=С                 | 4393                   | 20 <sup>e</sup>               | 3.26         | 2.77      | 113              |
| N—H…C=C                 | 3217                   | 51                            | 3.31         | 2.82      | 111              |
| O-H…phenyl <sup>b</sup> | 2823                   | 58                            | 3.29         | 2.43      | 150              |
| N—H…phenyl <sup>b</sup> | 3057                   | 12 <sup>h</sup>               | 3.31         | 2.47      | 148              |

<sup>a</sup> Intramolecular bonds are excluded. <sup>b</sup> For the X-H---phenyl bonds, the shorter of the distances between X/H to the centroid of the sixmembered ring and X/H to the nearest edge C atom was chosen to compute the mean. <sup>c</sup> Refcodes BETXAZ, CERYED, DAXWOO, DOJHEP, MNPREY. <sup>d</sup> Refcodes GERYEH (2), GICPIR, TEYNBH (2). <sup>c</sup> Refcodes BACXAE10 (2), BIWDEQ, BIWSOP (2), CAFVEK, CIMTEX, CYTOCH10, DULHUN, FADWOW, FENZED, FUPXUJ, GEZPUW, GIMZIL, LUTOSK, MOXCTS10, OHDLST, PROSTE, RTRSIN, RUGULA. <sup>f</sup> Refcodes ANTMYCO1, BPURMN, DEFMUW, FIWRAE, URPOAD10. <sup>s</sup> Refcodes APALTY, BHPHOL, DOZMIO, GEYZAL, MTBCHO. <sup>h</sup> Refcodes ABHYTZ, AMPHOM02, BAGFIY, BARKIO10, BIRKUI, FAJVIV, GEYXUD, KARHUG, MAMPOL, MXTRYP, TRYPTB, VUXYUI.

60 and 120°, presumably due to steric hindrance in other alternative positions. In steroid 1, this angle is 93°.

The generality of this novel X-H-C interaction was probed with the 1990 version of the Cambridge Structural Database<sup>12</sup> (82 129 entries). Error-free, nondisordered structures with R <0.100 and refined H-atom positions were accepted. The intermolecular contacts were stringently evaluated with the program GSTAT90: the X...C distance was specified to lie between 2.9 and 3.4 Å, even though C...O distances up to 3.75 Å have been considered viable for C-H-O hydrogen bonds;9 the X-H-C angle was constrained to lie between 100 and 180°; the hits were always scrutinized manually; and in several cases we referred to the original literature, and obviously inappropriate compounds were eliminated. With these stipulations, five unambiguous O-H-C=C contacts were found. We obtained similar results for N-H-C=C (5 bonds), O-H-C=C (20 bonds), N-H-C=C (5 bonds), O-H...phenyl (5 bonds), and N-H...phenyl (12 bonds). The results are summarized in Table I and, when considered along with Figure 1 and the details of the individual structures, provide compelling evidence that X-H-C hydrogen bonds, though rare and weak, are structurally significant when they do occur.

Table I shows that the more acidic hydrogens and the more basic  $\pi$ -systems are more likely to form better X-H…C hydrogen bonds. Further, contacts involving weaker acids (N-H) and weaker  $\pi$ -bases (alkene, phenyl) are more likely to be intramolecular and/or accompanied by bifurcation to stronger hydrogen bond acceptors (O, N). The O-H…C=C contacts are distinctive and seem able to control crystal packing in a decisive manner as in steroid 1.

The database analysis also reveals that, for X-H--phenyl contacts, the hydrogen atom is directed into the middle of the aromatic ring system (centroid type).<sup>3</sup> There is also the occurrence of a second type of interaction, where the X-H group makes a close approach to only two adjacent carbons of a particular phenyl group (edge type). This is observed in crystalline 2,6-diphenylphenol<sup>13</sup> and 4-nitro-2,6-diphenylphenol<sup>1</sup> though the contacts are intramolecular and as such not readily accessible via GSTAT90. Interestingly this edge-type X-H--phenyl contact has not been studied theoretically. The distinction, if any, between the centroid type and the edge type of X-H...phenyl contact is still elusive and warrants further study. Very little is known about N-H-C interactions, barring an ab initio calculation on the ammonia-benzene system<sup>2</sup> and a recent structural report,<sup>7</sup> and we believe that the present work may be the first to systematically study these interactions. To continue this line of reasoning, can a sufficiently acidic C-H group form a hydrogenbond-like C-H-C interaction? Or is such a geometry merely an extreme manifestation of the C-H( $\delta$ +)...C( $\delta$ -) herringbone interaction?<sup>14</sup> The crystal structures of propiolic and butynoic acids<sup>15</sup> are of some relevance in this context.

Table I includes biologically significant substances such as nucleic acid model compounds, steroids, and amino acids. Additionally, it is expected that this X-H···C hydrogen bond occurs not only in crystals but also in other environments such as aqueous media, either alone or in conjunction with other interactions.

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Supplementary Material Available: Tables of crystal data, positional parameters, general displacement parameter expressions, and distances and angles and a list of bibliographic references for the compounds in this study (15 pages); tables of structure factors (4 pages). Ordering information is given on any current masthead page.

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