STRUCTURE ANALYSIS OF NEWLY SYNTHESIZED STEROIDAL TETRAZOLES

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Introduction

Bearing in mind that androstane derivatives with tethered heterocyclic groups at position C–17 show anticancer activity, the aim of this study was a synthesis of novel steroidal derivatives with a tetrazole ring as a substituent at C–17 position, as well as a detailed investigation of their structures. The crystal structures of the two novel androstane derivatives: 3β -acetoxy-17-(1-methyl-1*H*tetrazol-5-yl)androsta-5,16-diene (**1**) и 3β-acetoxy-17-(2methyl-2*H*-tetrazol-5-yl)androsta-5,16-diene (2) were determined at room temperature.

Methods

COMPOUNDS

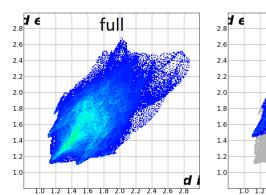
- 3β-acetoxy-17-(1-methyl-1*H*-tetrazol-5-yl)androsta-1. 5,16-diene (**1**)
- 2. 3β-acetoxy-17-(2-methyl-2*H*-tetrazol-5-yl)androsta-5,16-diene (2)

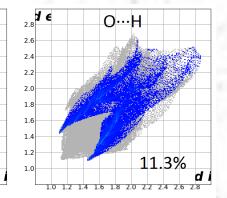
Hirshfeld surface analysis and 2D fingerprint plots

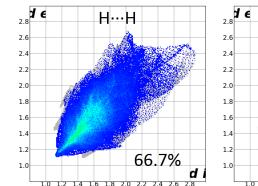
Fig. 1. Hirshfeld surface for conformer A of tetrazole 1 mapped with d_{norm} over the range of -0.0673 to 1.8356 and neighboring molecules associated with close contacts.

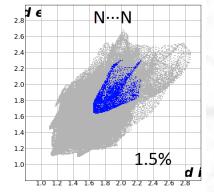
2.536 Å











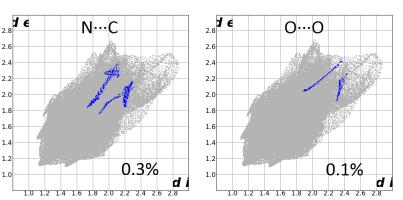
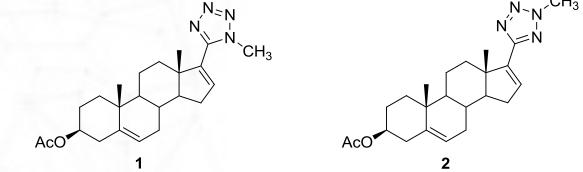


Fig 4. Two-dimensional fingerprint plots of conformer 1B

3.4%

 $N_N - N^CH_3$

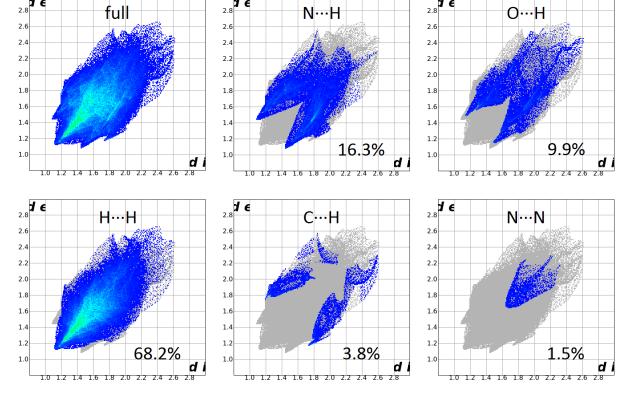


X-RAY CRYSTALLOGRAPHY

The diffraction data for **1** (MoK α , λ = 0.71071 Å) and **2** (CuK α , λ = 1.54184 Å) were collected at room temperature on Rigaku (Oxford Diffraction) Gemini S diffractometer using program package CrysAlis CCD. The data reduction was performed with program package CrysAlis RED. Collected data were corrected for absorption effects by using Multi-scan absorption correction. The structures were solved by direct methods using SHELXT. The structures were refined by full-matrix least-squares procedures on F^2 using SHELXL-2018/3 program.

| Crystallographic data for $C_{23}H_{32}N_4O_2$ (1) | |
|--|---|
| Crystal system | Monoclinic |
| Space group | P2 ₁ |
| Unit cell parameters | a = 9.7390 (5) Å b = 7.3412 (3) Å c = 30.5185 (15) Å $\beta = 96.970 (5)^{\circ}$ |
| Data/Restrains/Parameters | 7761/1/547 |
| Goodness-of-fit on F^2 | 1.036 |
| Final <i>R</i> indices $[l > 2\sigma(l)]$ | R1 = 0.0591 wR2 = 0.1148 |
| R indices [all data] | R1 = 0.1157 wR2 = 0.1387 |

| Crystallographic data for C_{23} H_{32} N_4 O_2 (2) | | |
|---|---|--|
| Crystal system | Orthorhombic | |
| Space group | P2 ₁ 2 ₁ 2 ₁ | |
| Unit cell parameters | a = 7.3721(2) Å b = 9.8807(2) Å c = 30.1493(9) Å | |
| Data/Restrains/Parameters | 4186/0/267 | |
| Goodness-of-fit on F^2 | 1.034 | |
| Final <i>R</i> indices $[I > 2\sigma(I)]$ | R1 = 0.0411 wR2 = 0.1084 | |
| <i>R</i> indices [all data] | R1 = 0.0519 wR2 = 0.1183 | |



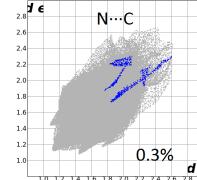
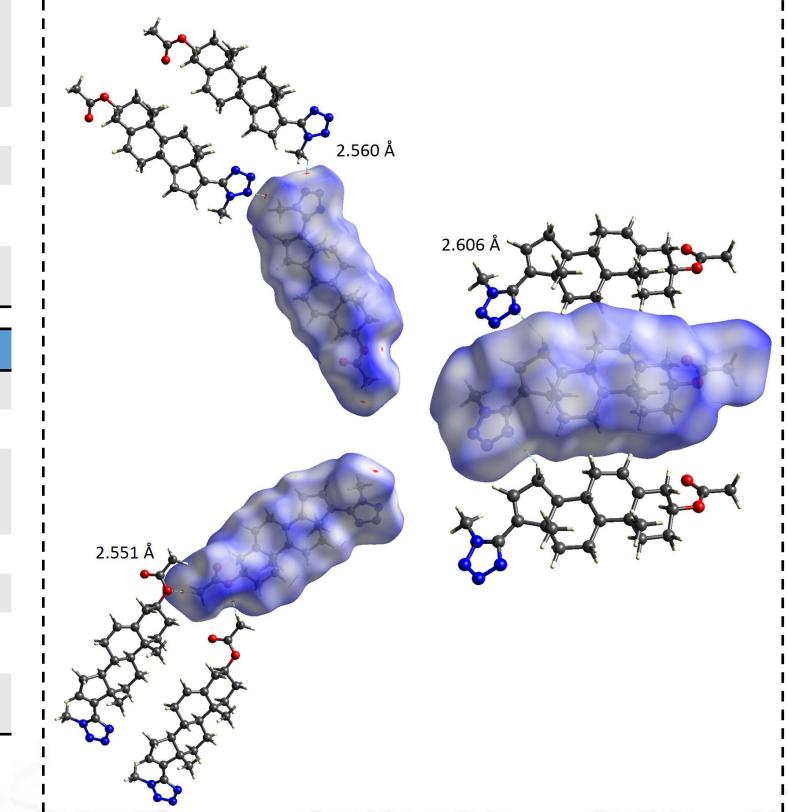


Fig 2. Two-dimensional fingerprint plots of conformer 1A



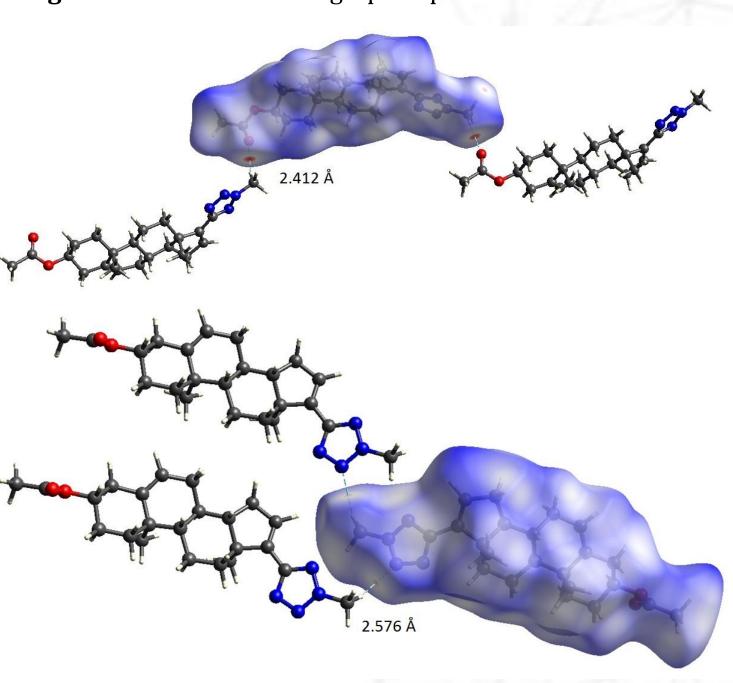
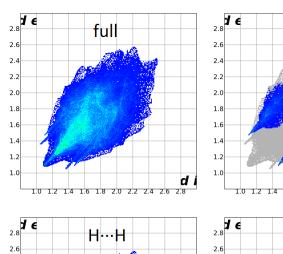
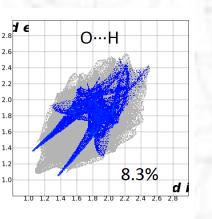
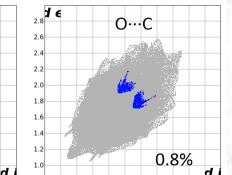


Fig. 5. Hirshfeld surface for tetrazole **2** mapped with d_{norm} over the range of -0.1413 to 1.7329 and neighboring molecules associated with close contacts.

16.8%

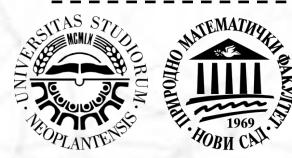






HIRSHFELD SURFACE ANALYSIS AND 2D FINGERPRINT PLOTS

CrystalExplorer 17.5 was used to analyze Hirshfeld surface and associated two-dimensional fingerprint plots for compounds 1 and 2.



Faculty of Sciences University of Novi Sad Fig. 3. Hirshfeld surface for conformer B of tetrazole 1 mapped with d_{norm} over the range of -0.0462 to 1.7465 and neighboring molecules associated with close contacts.

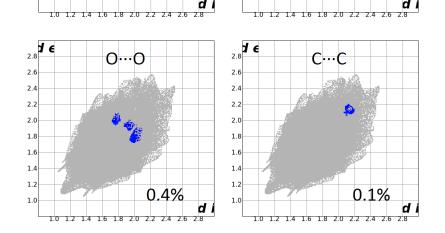


Fig 6. Two-dimensional fingerprint plots of conformer 1A

Acknowledgement

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