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# 5-(4-Hexyl-1H-1,2,3-triazol-1-yl)-2,1,3benzoxadiazole

## Jessie A. Key,<sup>a</sup> Christopher W. Cairo<sup>a</sup> and Michael J. Ferguson<sup>b</sup>\*

<sup>a</sup>Alberta Glycomics Centre, Department of Chemistry, University of Alberta, Edmonton, AB T6G 2G2, Canada, and <sup>b</sup>X-ray Crystallography Laboratory. Department of Chemistry, University of Alberta, Edmonton, AB T6G 2G2, Canada Correspondence e-mail: michael.ferguson@ualberta.ca

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Key indicators: single-crystal X-ray study; T = 173 K; mean  $\sigma$ (C–C) = 0.002 Å; R factor = 0.036; wR factor = 0.099; data-to-parameter ratio = 17.2.

The title compound,  $C_{14}H_{17}N_5O$ , a 1,2,3-triazole derivative of benzoxadiazole (C<sub>14</sub>H<sub>17</sub>N<sub>5</sub>O), was synthesized via Cu-catalvsed azide-alkyne cycloaddition (CuAAC) from the corres*n*-octyne 4-azidobenzoxadiazole. ponding and The benzoxadiazole and triazole rings show a roughly planar orientation [dihedral angle between the ring planes =  $12.18(5)^{\circ}$ ]. The alkane chain adopts a zigzag conformation, which deviates from the central triazole ring by  $20.89 (6)^{\circ}$ . These two torsion angles result in an overall twist to the structure, with a dihedral angle of  $32.86(7)^{\circ}$  between the benzoxadiazole group and the hexyl chain. The crystal structure features  $C-H \cdots N$  hydrogen bonds leading to chains propagating along  $[2\overline{1}0]$  and offset parallel stacking interactions of the triazole and benzoxadiazole rings. The centroid of the extended  $\pi$ -system formed by the benzoxadiazole and triazole rings (14 atoms total) was calculated; the centroid-centroid distance was 4.179 Å, interplanar separation was 3.243 Å, and the resulting offset was 2.636 Å.

## **Related literature**

For the synthesis of the title compound and related benzoxadiazole analogs, see: Key & Cairo (2011). For computational studies of the absorption and fluorescence properties of this series of compounds, see: Brown et al. (2012). For structures with 1-aryl-substituted 1,2,3-triazole rings, see: Costa et al. (2006). For the use of fluorophores as chemical or biological probes, see: Cairo et al. (2010); Lavis & Raines (2008). For related benzoxadiazole structures, see: Key et al. (2012*a*,*b*). For triazole-substituted coumarin derivatives, see: Key et al. (2009).



 $\gamma = 85.6240 \ (17)^{\circ}$ 

Mo  $K\alpha$  radiation

 $1.02 \times 0.35 \times 0.03 \text{ mm}$ 

 $\mu = 0.09 \text{ mm}^-$ 

T = 173 K

Z = 2

 $V = 685.04 (17) \text{ Å}^3$ 

## **Experimental**

Crystal data C14H17N5O  $M_r = 271.33$ Triclinic,  $P\overline{1}$ a = 5.3604 (8) Å b = 7.8585 (11) Å c = 16.357 (2) Å  $\alpha = 87.4656(17)^{\circ}$  $\beta = 86.2519 \ (16)^{\circ}$ 

#### Data collection

Bruker APEXII CCD	6114 measured reflections
diffractometer	3120 independent reflections
Absorption correction: multi-scan	2568 reflections with $I > 2\sigma(I)$
(SADABS; Bruker, 2008)	$R_{\rm int} = 0.013$
$T_{\min} = 0.915, \ T_{\max} = 0.997$	

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.036$	181 parameters
$wR(F^2) = 0.099$	H-atom parameters constrained
S = 1.04	$\Delta \rho_{\rm max} = 0.20 \text{ e } \text{\AA}^{-3}$
3120 reflections	$\Delta \rho_{\rm min} = -0.22 \text{ e } \text{\AA}^{-3}$

### Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C3 - H3 \cdot \cdot \cdot N2^{i}$	0.95	2.52	3.4674 (15)	177
C3-H3···N4	0.93	2.40	5.5445 (15)	134

Symmetry codes: (i) -x + 1, -y, -z + 1; (ii) -x - 1, -y + 1, -z + 1.

Data collection: APEX2 (Bruker, 2008); cell refinement: SAINT (Bruker, 2008); data reduction: SAINT; program(s) used to solve structure: SHELXD (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: MW2081).

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