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Ethyl 4-(tert-butylamino)-3-nitrobenzoate

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Key indicators: single-crystal X-ray study; T = 100 K; mean σ (C–C) = 0.002 Å; R factor = 0.047; wR factor = 0.134; data-to-parameter ratio = 23.2.

In the title compound, C13H18N2O4, intramolecular N- $H \cdots O$, $N - H \cdots N$ and $C - H \cdots O$ (× 3) hydrogen bonds generate S(6) and S(5) ring motifs. There are two crystallographically independent molecules (A and B) in the asymmetric unit. The nitro group is coplanar with the benzene ring, with O-N-C-C torsion angles of -0.33 (13) and $0.93 (14)^{\circ}$ in molecules A and B, respectively. In the crystal structure, neighbouring molecules are linked together by intermolecular C-H···O hydrogen bonds. In addition, the crystal structure is stabilized by π - π interactions with centroid-centroid distances ranging from 3.7853 (6) to 3.8625 (6) Å.

Related literature

For literature on hydrogen-bond motifs, see: Bernstein et al. (1995). For values of bond lengths, see: Allen et al. (1987). For related literature, see, for example: Göker et al. (1998); Anderson (2005); Kakei et al. (1993).



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Experimental

Crystal data

β

$C_{13}H_{18}N_2O_4$	$V = 2722.37 (14) \text{ Å}^3$
$M_r = 266.29$	Z = 8
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
a = 16.0471 (5) Å	$\mu = 0.10 \text{ mm}^{-1}$
b = 6.6417 (2) Å	T = 100.0 (1) K
c = 30.0180 (9) Å	$0.51 \times 0.43 \times 0.17 \text{ mm}$
$\beta = 121.688 \ (2)^{\circ}$	

Data collection

Bruker SMART APEXII CCD area-detector diffractometer Absorption correction: multi-scan (SADABS; Bruker, 2005) $T_{\min} = 0.879, \ T_{\max} = 0.984$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.046$	351 parameters
$wR(F^2) = 0.133$	H-atom parameters constrained
S = 1.04	$\Delta \rho_{\rm max} = 0.53 \ {\rm e} \ {\rm \AA}^{-3}$
8141 reflections	$\Delta \rho_{\rm min} = -0.24 \text{ e } \text{\AA}^{-3}$

63326 measured reflections

 $R_{\rm int} = 0.033$

8141 independent reflections

6368 reflections with $I > 2\sigma(I)$

Table 1

Selected centroid-centroid distances (Å).

Cg1 and Cg2 are the centroids of the C1A-C6A and C1B-C6B rings, respectively.

$Cg1 \cdots Cg2^{i}$	3.7853 (6)	$Cg1 \cdots Cg2^{ii}$	3.8625 (6)
Symmetry codes: (i) x	$, -y - \frac{1}{2}, z - \frac{1}{2};$ (ii) $x, z = \frac{1}{2}$	$-y + \frac{1}{2}, z - \frac{1}{2}.$	

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N2A - H2NA \cdots O4A$	0.86	1.93	2.6299 (15)	138
$N2A - H2NA \cdots N1A$	0.86	2.54	2.9361 (15)	109
$N2B - H2NB \cdots O4B$	0.87	1.95	2.6355 (15)	134
$N2B - H2NB \cdot \cdot \cdot N1B$	0.87	2.58	2.9419 (15)	106
$C1A - H1A \cdots O3A$	0.95	2.31	2.6522 (16)	100
$C1B - H1B \cdots O3B$	0.95	2.31	2.6498 (16)	100
$C4A - H4A \cdots O3B^{iii}$	0.95	2.50	3.4124 (17)	160
$C4B - H4B \cdots O3A^{iv}$	0.95	2.42	3.2566 (18)	147
$C5A - H5A \cdots O1A$	0.95	2.41	2.7326 (13)	100
$C11A - H11A \cdots O2B^{v}$	0.98	2.55	3.4714 (17)	157
$C11B - H11F \cdot \cdot \cdot O2A^{ii}$	0.98	2.52	3.4446 (17)	158
$C13B - H13D \cdots O2A^{vi}$	0.98	2.60	3.5071 (17)	154

 $-x + 1, -y + 1, -z + 1; (v) x, -y + \frac{1}{2}, z + \frac{1}{2}; (vi) x, -y + \frac{3}{2}, z - \frac{1}{2}.$

Data collection: APEX2 (Bruker, 2005); cell refinement: APEX2; data reduction: SAINT (Bruker, 2005); program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL and PLATON (Spek, 2003).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: AT2594).

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Ethyl 4-(tert-butylamino)-3-nitrobenzoate

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Comment

As a part of our ongoing studies on new nitro benzoic acid derivatives, we have synthesized the title compound (I) employing a modified protocol of previous procedure (Göker *et al.*, 1998). The nitro benzoic acid intermediates are a convenient starting material for the synthesis of heterocycles targeting important biological processes, *e.g.* antifungal (Fluconazole) (Anderson, 2005) and proton pump inhibitor (Omeprazole) (Kakei *et al.*, 1993). The crystal structure of the *tert*-butylamino functionalized nitro benzoic acid (I) has been determined, and herein, we present a full report on its crystal structure.

In the title compound (I) (Fig. 1), intramolecular N—H···O (x 2), N—H···N (x 2), and C—H···O (x 3) hydrogen bonds (Table 2) generate *S*(6) and *S*(5) ring motifs, respectively (Bernstein *et al.*, 1995). There are two crystallographically independent molecules, A and B in the asymmetric unit of the title compound. The nitro group is coplanar with the benzene ring with torsion angle of -0.33 (13) and 0.93 (14)° in the molecule A and B, respectively. In the crystal structure neighbouring molecules are linked together by intermolecular C—H···O hydrogen bonds (Table 1). In the crystal packing (Table 2 & Fig. 2), molecules are stacked down the *b* axis, being consolidated by π – π interactions with centroid to centroid distances ranging from 3.7853 (6) – 3.8625 (6) Å.

The crystal structure is stabilized by intramolecular N—H···O (x 2), N—H···N (x 2), C—H···O (x 3), and intermolecular C—H···O (x 5) hydrogen bonds and π - π interactions.

Experimental

Ethyl 4-fluoro-3-nitrobenzoate (200 mg, 0.93 mmol) was dissolved in dry dichloromethane (10 ml). *N*, *N*-diisopropylethylamine (DIPEA) (0.20 ml, 1.12 mmol) was added to the stirred mixture. Then, *tert*-butylamine (0.11 ml, 1.03 mmol) was added dropwise using syringe and stirred at room temperature under N₂ overnight. After completion of the reaction, the mixture was washed with 10% NaCO₃ (10 ml). The aqueous layer was washed with dichloromethane (3 x 15 ml). The organic layers were collected and dried over MgSO₄ (anhydrous). The solvent was removed under reduced pressure to yield the crude product. Recrystallisation with hot hexane revealed the title compound (I) as bright yellow crystals.

Refinement

The H-atoms attached to N2A and N2B were located from the difference Fourier map and refined as riding with the parent atom with an isotropic thermal parameter 1.2 times that of the parent atom. The rest of the hydrogen atoms were positioned geometrically [C-H = 0.95-98 Å] and refined using a riding model. A rotating-group model was used for the methyl groups. The highest peak is located 0.63 Å from C6B and the deepest hole is located 0.59 Å from N1A.

Figures



Fig. 1. The molecular structure of the title compound, showing 50% probability displacement ellipsoids and the atomic numbering. Intramolecular hydrogen bonds are drawn as dashed lines.



Fig. 2. The crystal packing of (I), showing stacking arrangement viewed down the *b*-axis. Intramolecular and intermolecular interactions are drawn as dashed lines.

Ethyl 4-(tert-butylamino)-3-nitrobenzoate

Crystal data	
$C_{13}H_{18}N_2O_4$	$F_{000} = 1136$
$M_r = 266.29$	$D_{\rm x} = 1.299 {\rm ~Mg~m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation $\lambda = 0.71073$ Å
Hall symbol: -P 2ybc	Cell parameters from 9969 reflections
<i>a</i> = 16.0471 (5) Å	$\theta = 2.5 - 30.1^{\circ}$
b = 6.6417 (2) Å	$\mu = 0.10 \text{ mm}^{-1}$
c = 30.0180 (9) Å	T = 100.0 (1) K
$\beta = 121.688 \ (2)^{\circ}$	Plate, yellow
$V = 2722.37 (14) \text{ Å}^3$	$0.51 \times 0.43 \times 0.17 \text{ mm}$
Z = 8	

Data collection

Bruker SMART APEXII CCD area-detector diffractometer	8141 independent reflections
Radiation source: fine-focus sealed tube	6368 reflections with $I > 2\sigma(I)$
Monochromator: graphite	$R_{\rm int} = 0.033$
T = 100.0(1) K	$\theta_{\text{max}} = 30.3^{\circ}$
ϕ and ω scans	$\theta_{\min} = 1.5^{\circ}$
Absorption correction: multi-scan (SADABS; Bruker, 2005)	$h = -21 \rightarrow 22$
$T_{\min} = 0.879, T_{\max} = 0.984$	$k = -9 \rightarrow 9$
63326 measured reflections	$l = -41 \rightarrow 42$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.047$	H-atom parameters constrained

$P(F^2) = 0.122$	$w = 1/[\sigma^2(F_0^2) + (0.0652P)^2 + 0.787P]$
$WR(F^{-}) = 0.133$	where $P = (F_0^2 + 2F_c^2)/3$
<i>S</i> = 1.04	$(\Delta/\sigma)_{\text{max}} = 0.001$
8141 reflections	$\Delta \rho_{max} = 0.53 \text{ e } \text{\AA}^{-3}$
351 parameters	$\Delta \rho_{min} = -0.24 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	Extinction correction: none

Special details

Experimental. The low-temperature data was collected with the Oxford Cyrosystem Cobra low-temperature attachment.

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted *R*-factor *wR* and goodness of fit *S* are based on F^2 , conventional *R*-factors *R* are based on *F*, with *F* set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating *R*-factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. *R*-factors based on F^2 are statistically about twice as large as those based on *F*, and *R*- factors based on ALL data will be even larger.

	x	У	z	$U_{\rm iso}*/U_{\rm eq}$
O1A	1.01188 (6)	0.50707 (13)	0.90700 (3)	0.02027 (18)
O2A	0.87073 (6)	0.53054 (14)	0.90534 (3)	0.02371 (19)
O3A	0.57249 (6)	0.54152 (13)	0.73478 (3)	0.02243 (18)
O4A	0.56803 (6)	0.54680 (13)	0.66152 (3)	0.02250 (18)
N1A	0.61441 (6)	0.54228 (13)	0.71013 (4)	0.01589 (18)
N2A	0.73120 (6)	0.54415 (13)	0.66080(3)	0.01583 (18)
H2NA	0.6682	0.5497	0.6444	0.019*
C1A	0.76647 (7)	0.53258 (15)	0.79329 (4)	0.0143 (2)
H1A	0.7281	0.5335	0.8089	0.017*
C2A	0.72036 (7)	0.53705 (15)	0.73898 (4)	0.01349 (19)
C3A	0.77441 (7)	0.53716 (14)	0.71324 (4)	0.01347 (19)
C4A	0.87825 (7)	0.52932 (15)	0.74731 (4)	0.0151 (2)
H4A	0.9180	0.5271	0.7325	0.018*
C5A	0.92253 (7)	0.52490 (15)	0.80055 (4)	0.0149 (2)
H5A	0.9920	0.5204	0.8217	0.018*
C6A	0.86747 (7)	0.52682 (15)	0.82473 (4)	0.0142 (2)
C7A	0.91408 (8)	0.52224 (16)	0.88226 (4)	0.0159 (2)
C8A	1.06291 (8)	0.5008 (2)	0.96363 (4)	0.0252 (3)
H8A	1.0412	0.3828	0.9751	0.030*
H8B	1.0489	0.6242	0.9771	0.030*
C9A	1.17025 (9)	0.4861 (2)	0.98412 (5)	0.0347 (3)
H9A	1.2069	0.4877	1.0225	0.052*
H9B	1.1904	0.6006	0.9713	0.052*
H9C	1.1837	0.3603	0.9720	0.052*

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (A^2)

C10A	0.77667 (8)	0.54745 (16)	0.62878 (4)	0.0163 (2)
C11A	0.83246 (8)	0.35165 (17)	0.63510 (4)	0.0203 (2)
H11A	0.7877	0.2370	0.6254	0.030*
H11B	0.8861	0.3373	0.6716	0.030*
H11C	0.8593	0.3555	0.6124	0.030*
C12A	0.68959 (9)	0.56150 (19)	0.57218 (4)	0.0233 (2)
H12A	0.6458	0.4465	0.5647	0.035*
H12B	0.7136	0.5600	0.5481	0.035*
H12C	0.6537	0.6869	0.5676	0.035*
C13A	0.84047 (8)	0.73502 (17)	0.64034 (4)	0.0195 (2)
H13A	0.8927	0.7359	0.6774	0.029*
H13B	0.8000	0.8560	0.6324	0.029*
H13C	0.8695	0.7331	0.6186	0.029*
O1B	0.49200 (5)	0.43211 (13)	0.10402 (3)	0.02139 (18)
O2B	0.63293 (6)	0.45058 (13)	0.10484 (3)	0.02286 (18)
O3B	0.93204 (6)	0.47400 (13)	0.27341 (3)	0.02111 (18)
O4B	0.93898 (6)	0.47557 (13)	0.34743 (3)	0.02151 (18)
N1B	0.89147 (6)	0.47252 (13)	0.29872 (4)	0.01529 (18)
N2B	0.77682 (6)	0.47199 (14)	0.34950 (3)	0.01565 (18)
H2NB	0.8407	0.4767	0.3671	0.019*
C1B	0.73817 (7)	0.46115 (15)	0.21633 (4)	0.01401 (19)
H1B	0.7757	0.4608	0.2002	0.017*
C2B	0.78566 (7)	0.46722 (15)	0.27069 (4)	0.01293 (19)
C3B	0.73273 (7)	0.46737 (15)	0.29702 (4)	0.01340 (19)
C4B	0.62889 (8)	0.46322 (16)	0.26366 (4)	0.0157 (2)
H4B	0.5900	0.4645	0.2790	0.019*
C5B	0.58317 (7)	0.45740 (15)	0.21029 (4)	0.0154 (2)
H5B	0.5137	0.4545	0.1895	0.019*
C6B	0.63721 (7)	0.45564 (15)	0.18553 (4)	0.01401 (19)
C7B	0.58976 (8)	0.44669 (16)	0.12800 (4)	0.0168 (2)
C8B	0.43816 (8)	0.4179 (2)	0.04737 (4)	0.0260 (3)
H8C	0.3751	0.3484	0.0350	0.031*
H8D	0.4762	0.3370	0.0365	0.031*
C9B	0.41883 (9)	0.6226 (2)	0.02271 (5)	0.0302 (3)
H9D	0.3804	0.6089	-0.0155	0.045*
H9E	0.4812	0.6888	0.0335	0.045*
H9F	0.3822	0.7038	0.0340	0.045*
C10B	0.73206 (8)	0.46992 (16)	0.38189 (4)	0.0172 (2)
C11B	0.67381 (8)	0.27595 (17)	0.37340 (4)	0.0217 (2)
H11D	0.6189	0.2698	0.3370	0.032*
H11E	0.6487	0.2747	0.3969	0.032*
H11F	0.7166	0.1593	0.3809	0.032*
C12B	0.81999 (9)	0.47351 (19)	0.43837 (4)	0.0246 (2)
H12D	0.8582	0.5964	0.4441	0.037*
H12E	0.8613	0.3555	0.4445	0.037*
H12F	0.7966	0.4710	0.4627	0.037*
C13B	0.66997 (8)	0.65807 (18)	0.37269 (5)	0.0222 (2)
H13D	0.7108	0.7784	0.3806	0.033*
H13E	0.6433	0.6541	0.3955	0.033*

H13F	0.6160	0.6623	0.3360	0.	033*	
Atomic displacer	nent parameters ((A^2)				
	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
01A	0.0132 (3)	0.0327 (5)	0.0134 (4)	0.0004 (3)	0.0059 (3)	-0.0014 (3)
O2A	0.0181 (4)	0.0365 (5)	0.0190 (4)	0.0000 (3)	0.0114 (3)	-0.0013 (3)
O3A	0.0154 (4)	0.0298 (5)	0.0260 (4)	0.0006 (3)	0.0135 (3)	0.0003 (3)
O4A	0.0136 (4)	0.0309 (5)	0.0186 (4)	0.0002 (3)	0.0055 (3)	-0.0002 (3)
N1A	0.0126 (4)	0.0142 (4)	0.0205 (4)	-0.0004 (3)	0.0084 (3)	-0.0004 (3)
N2A	0.0120 (4)	0.0192 (4)	0.0151 (4)	0.0004 (3)	0.0062 (3)	0.0003 (3)
C1A	0.0149 (5)	0.0111 (4)	0.0189 (5)	-0.0006 (3)	0.0104 (4)	-0.0004 (4)
C2A	0.0102 (4)	0.0119 (5)	0.0175 (5)	0.0002 (3)	0.0068 (4)	0.0002 (4)
C3A	0.0135 (4)	0.0099 (4)	0.0168 (5)	-0.0006 (3)	0.0078 (4)	-0.0003 (3)
C4A	0.0127 (4)	0.0155 (5)	0.0182 (5)	0.0003 (4)	0.0088 (4)	0.0005 (4)
C5A	0.0124 (4)	0.0145 (5)	0.0175 (5)	0.0003 (4)	0.0076 (4)	0.0006 (4)
C6A	0.0147 (5)	0.0131 (5)	0.0148 (5)	-0.0004 (4)	0.0077 (4)	-0.0005 (4)
C7A	0.0140 (5)	0.0160 (5)	0.0170 (5)	-0.0009 (4)	0.0075 (4)	-0.0013 (4)
C8A	0.0188 (5)	0.0421 (7)	0.0122 (5)	0.0004 (5)	0.0065 (4)	-0.0021 (5)
C9A	0.0184 (6)	0.0630 (10)	0.0180 (6)	0.0030 (6)	0.0063 (5)	-0.0027 (6)
C10A	0.0161 (5)	0.0183 (5)	0.0154 (5)	-0.0005 (4)	0.0088 (4)	0.0001 (4)
C11A	0.0208 (5)	0.0181 (5)	0.0240 (5)	-0.0002 (4)	0.0131 (4)	-0.0028 (4)
C12A	0.0212 (5)	0.0303 (6)	0.0157 (5)	-0.0009 (5)	0.0078 (4)	-0.0005 (4)
C13A	0.0205 (5)	0.0190 (5)	0.0203 (5)	-0.0011 (4)	0.0116 (4)	0.0018 (4)
O1B	0.0133 (4)	0.0343 (5)	0.0138 (4)	-0.0022 (3)	0.0053 (3)	0.0013 (3)
O2B	0.0181 (4)	0.0339 (5)	0.0178 (4)	0.0012 (3)	0.0102 (3)	-0.0004 (3)
O3B	0.0150 (4)	0.0280 (4)	0.0238 (4)	0.0003 (3)	0.0127 (3)	0.0009 (3)
O4B	0.0136 (4)	0.0302 (5)	0.0164 (4)	0.0002 (3)	0.0049 (3)	0.0004 (3)
N1B	0.0135 (4)	0.0134 (4)	0.0186 (4)	0.0001 (3)	0.0081 (3)	0.0002 (3)
N2B	0.0129 (4)	0.0195 (4)	0.0143 (4)	0.0001 (3)	0.0070 (3)	0.0001 (3)
C1B	0.0144 (4)	0.0125 (5)	0.0168 (5)	0.0004 (3)	0.0093 (4)	0.0006 (4)
C2B	0.0108 (4)	0.0113 (4)	0.0166 (5)	0.0003 (3)	0.0072 (4)	0.0007 (3)
C3B	0.0147 (5)	0.0101 (4)	0.0151 (5)	0.0004 (3)	0.0077 (4)	0.0010 (3)
C4B	0.0137 (4)	0.0173 (5)	0.0185 (5)	0.0004 (4)	0.0100 (4)	0.0005 (4)
C5B	0.0121 (4)	0.0151 (5)	0.0185 (5)	0.0000 (4)	0.0076 (4)	0.0007 (4)
C6B	0.0135 (4)	0.0133 (5)	0.0148 (5)	0.0000 (3)	0.0071 (4)	0.0009 (3)
C7B	0.0152 (5)	0.0170 (5)	0.0163 (5)	0.0001 (4)	0.0070 (4)	0.0007 (4)
C8B	0.0176 (5)	0.0410 (7)	0.0148 (5)	-0.0037 (5)	0.0055 (4)	-0.0017 (5)
C9B	0.0236 (6)	0.0469 (8)	0.0197 (6)	0.0022 (5)	0.0111 (5)	0.0074 (5)
C10B	0.0194 (5)	0.0187 (5)	0.0158 (5)	0.0009 (4)	0.0107 (4)	0.0007 (4)
C11B	0.0255 (6)	0.0209 (5)	0.0226 (5)	-0.0013 (4)	0.0154 (5)	0.0024 (4)
C12B	0.0265 (6)	0.0306 (6)	0.0152 (5)	0.0009 (5)	0.0099 (5)	0.0008 (4)
C13B	0.0242 (5)	0.0205 (5)	0.0251 (6)	0.0019 (4)	0.0151 (5)	-0.0022 (4)

Geometric parameters (Å, °)

O1A—C7A	1.3418 (13)	O1B—C7B	1.3424 (13)
O1A—C8A	1.4489 (13)	O1B—C8B	1.4503 (13)
O2A—C7A	1.2136 (13)	O2B—C7B	1.2125 (13)

O3A—N1A	1.2340 (12)	O3B—N1B	1.2332 (12)
O4A—N1A	1.2424 (12)	O4B—N1B	1.2443 (12)
N1A—C2A	1.4474 (13)	N1B—C2B	1.4462 (13)
N2A—C3A	1.3470 (13)	N2B—C3B	1.3463 (13)
N2A-C10A	1.4810 (13)	N2B—C10B	1.4816 (13)
N2A—H2NA	0.8617	N2B—H2NB	0.8733
C1A—C6A	1.3827 (14)	C1B—C6B	1.3809 (14)
C1A—C2A	1.3925 (14)	C1B—C2B	1.3924 (14)
C1A—H1A	0.9500	C1B—H1B	0.9500
C2A—C3A	1.4328 (14)	C2B—C3B	1.4321 (14)
C3A—C4A	1.4265 (14)	C3B—C4B	1.4238 (14)
C4A—C5A	1.3663 (14)	C4B—C5B	1.3683 (14)
C4A—H4A	0.9500	C4B—H4B	0.9500
C5A—C6A	1.4076 (14)	C5B—C6B	1.4076 (14)
С5А—Н5А	0.9500	C5B—H5B	0.9500
С6А—С7А	1.4784 (14)	C6B—C7B	1.4778 (14)
C8A—C9A	1.4973 (17)	C8B—C9B	1.5003 (19)
C8A—H8A	0.9900	C8B—H8C	0.9900
C8A—H8B	0 9900	C8B—H8D	0 9900
C9A—H9A	0.9800	C9B—H9D	0.9800
C9A—H9B	0.9800	C9B—H9E	0.9800
C9A—H9C	0.9800	C9B—H9F	0.9800
C10A - C13A	1 5320 (15)	C10B-C13B	1,5302(15)
C10A - C11A	1.5320(15) 1.5331(15)	C10B-C11B	1.5302(15) 1.5323(15)
	1.5351(15) 1.5343(15)	C10B C12B	1.5323(15) 1.5338(15)
C_{10A} $-C_{12A}$	0.0800	C11B H11D	0.0800
	0.9800		0.9800
	0.9800		0.9800
Clia Hia	0.9800	CIIB—HIIF	0.9800
CI2A—HI2A	0.9800	CI2B—HI2D	0.9800
CI2A—HI2B	0.9800	CI2B—HI2E	0.9800
CI2A—HI2C	0.9800	CI2B—HI2F	0.9800
CI3A—HI3A	0.9800	CI3B—HI3D	0.9800
С13А—Н13В	0.9800	C13B—H13E	0.9800
С13А—Н13С	0.9800	C13B—H13F	0.9800
Cg1…Cg2 ⁱ	3.7853 (6)	Cg1…Cg2 ⁱⁱ	3.8625 (6)
C7A—O1A—C8A	115.30 (9)	C7B—O1B—C8B	116.24 (9)
O3A—N1A—O4A	121.71 (9)	O3B—N1B—O4B	121.87 (9)
O3A—N1A—C2A	118.70 (9)	O3B—N1B—C2B	118.70 (9)
O4A—N1A—C2A	119.59 (9)	O4B—N1B—C2B	119.43 (9)
C3A—N2A—C10A	129.22 (9)	C3B—N2B—C10B	129.04 (9)
C3A—N2A—H2NA	113.4	C3B—N2B—H2NB	115.9
C10A—N2A—H2NA	117.3	C10B—N2B—H2NB	115.0
C6A—C1A—C2A	120.74 (9)	C6B—C1B—C2B	120.85 (9)
C6A—C1A—H1A	119.6	C6B—C1B—H1B	119.6
C2A—C1A—H1A	119.6	C2B—C1B—H1B	119.6
C1A—C2A—C3A	122.10 (9)	C1B—C2B—C3B	121.91 (9)
C1A—C2A—N1A	115.83 (9)	C1B—C2B—N1B	115.79 (9)
C3A—C2A—N1A	122.07 (9)	C3B—C2B—N1B	122.30 (9)

N2A—C3A—C4A	121.97 (9)	N2B—C3B—C4B	121.65 (9)
N2A—C3A—C2A	122.98 (9)	N2B—C3B—C2B	123.11 (9)
C4A—C3A—C2A	115.05 (9)	C4B—C3B—C2B	115.23 (9)
C5A—C4A—C3A	122.23 (10)	C5B—C4B—C3B	122.25 (9)
С5А—С4А—Н4А	118.9	C5B—C4B—H4B	118.9
C3A—C4A—H4A	118.9	C3B—C4B—H4B	118.9
C4A—C5A—C6A	121.42 (9)	C4B—C5B—C6B	121.22 (9)
C4A—C5A—H5A	119.3	C4B—C5B—H5B	119.4
С6А—С5А—Н5А	119.3	C6B—C5B—H5B	119.4
C1A—C6A—C5A	118.45 (9)	C1B—C6B—C5B	118.54 (9)
C1A—C6A—C7A	119.36 (9)	C1B—C6B—C7B	119.12 (9)
C5A—C6A—C7A	122.19 (9)	C5B—C6B—C7B	122.34 (9)
O2A—C7A—O1A	122.82 (10)	O2B—C7B—O1B	123.56 (10)
O2A—C7A—C6A	125.16 (10)	O2B—C7B—C6B	124.77 (10)
O1A—C7A—C6A	112.03 (9)	O1B—C7B—C6B	111.67 (9)
O1A—C8A—C9A	107.65 (9)	O1B—C8B—C9B	111.18 (11)
O1A—C8A—H8A	110.2	01B—C8B—H8C	109.4
C9A—C8A—H8A	110.2	C9B—C8B—H8C	109.4
O1A - C8A - H8B	110.2	O1B - C8B - H8D	109.4
C9A—C8A—H8B	110.2	C9B—C8B—H8D	109.4
H8A—C8A—H8B	108.5	H8C - C8B - H8D	108.0
C8A - C9A - H9A	109.5	C8B - C9B - H9D	109.5
C8A—C9A—H9B	109.5	C8B—C9B—H9E	109.5
H9A_C9A_H9B	109.5	H9D - C9B - H9E	109.5
C8A - C9A - H9C	109.5	C8B—C9B—H9F	109.5
H9A - C9A - H9C	109.5	H9D - C9B - H9F	109.5
H9B_C9A_H9C	109.5	H9F_C9B_H9F	109.5
N2A - C10A - C13A	111 31 (9)	N2B— $C10B$ — $C13B$	111 59 (9)
N2A $C10A$ $C11A$	111.31 (9)	N2B-C10B-C11B	111.37 (9)
$C_{13}A - C_{10}A - C_{11}A$	112.61 (9)	C_{13B} C_{10B} C_{11B}	111.27 (9)
N2A - C10A - C12A	104.30 (8)	N2B_C10B_C12B	104 10 (9)
$C_{13}A - C_{10}A - C_{12}A$	104.50(0) 108.24(9)	C13B - C10B - C12B	104.10 (9)
$C_{11}A - C_{10}A - C_{12}A$	108.24(9) 108.76(9)	C11B - C10B - C12B	108.88 (9)
C10A - C11A - H11A	108.70 ())	C10B-C11B-H11D	100.00 ())
C10A - C11A - H11B	109.5	C10B-C11B-H11F	109.5
	109.5	H11D C11B H11E	109.5
C10A - C11A - H11C	109.5	C10B-C11B-H11F	109.5
	109.5		109.5
H11B C11A H11C	109.5	H11E C11B H11E	109.5
C10A - C12A - H12A	109.5	C10B-C12B-H12D	109.5
C10A = C12A = H12R	109.5	C10B = C12B = H12E	109.5
H12A C12A H12B	109.5	H12D C12B H12E	109.5
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	109.5	$\begin{array}{cccc} \text{III2D} & \text{III2D} \\ \text{III2D} & \text{III2D} \\ \text{III2D} & \text{III2E} \\ \end{array}$	109.5
H12A C12A H12C	109.5	H12D C12B H12F	109.5
H12B_C12A_H12C	109.5	H12D - C12D - H12F H12F - C12B - H12F	109.5
	109.5	C10R C12R H12D	109.5
C10A C12A H12D	109.5	C10B - C13B - H13D	109.5
H12A C12A H12D	109.5	$\begin{array}{cccc} C10D & C13D & M13E \\ H13D & C13B & H13E \\ \end{array}$	109.5
ПІЗА—СІЗА—ПІЗВ	109.3	$\begin{array}{cccc} \Pi 1 \mathcal{D} & \Pi \mathcal{D} \\ \Pi \mathcal{D} & \Pi \mathcal{D} \\ $	109.5
CIUA-CISA-HISC	109.5	CINE-CISE-HISE	109.3

H13A—C13A—H13C	109.5	H13D—C13B—H13F		109.5
H13B—C13A—H13C	109.5	H13E—C13B—H13F		109.5
Symmetry codes: (i) x , $-y-1/2$, $z-1/2$; (ii) <i>x</i> , − <i>y</i> +1/2, <i>z</i> −1/2.			
Hydrogen-bond geometry (Å, °)				
D—H···A	<i>D</i> —Н	H···A	$D \cdots A$	D—H··· A
N2A—H2NA···O4A	0.86	1.93	2.6299 (15)	138
N2A—H2NA…N1A	0.86	2.54	2.9361 (15)	109
N2B—H2NB····O4B	0.87	1.95	2.6355 (15)	134
N2B—H2NB…N1B	0.87	2.58	2.9419 (15)	106
C1A—H1A···O3A	0.95	2.31	2.6522 (16)	100
C1B—H1B···O3B	0.95	2.31	2.6498 (16)	100
C4A—H4A···O3B ⁱⁱⁱ	0.95	2.50	3.4124 (17)	160
C4B—H4B····O3A ^{iv}	0.95	2.42	3.2566 (18)	147
C5A—H5A···O1A	0.95	2.41	2.7326 (13)	100
C11A—H11A····O2B ^v	0.98	2.55	3.4714 (17)	157
C11B—H11F···O2A ⁱⁱ	0.98	2.52	3.4446 (17)	158
C13B—H13D····O2A ^{vi}	0.98	2.60	3.5071 (17)	154

Symmetry codes: (iii) -x+2, -y+1, -z+1; (iv) -x+1, -y+1, -z+1; (v) x, -y+1/2, z+1/2; (ii) x, -y+1/2, z-1/2; (vi) x, -y+3/2, z-1/2.



Fig. 1

Fig. 2

