



# Crystal structure of 2-amino-4-phenyl-4*H*-benzo[*h*]chromene-3-carbonitrile

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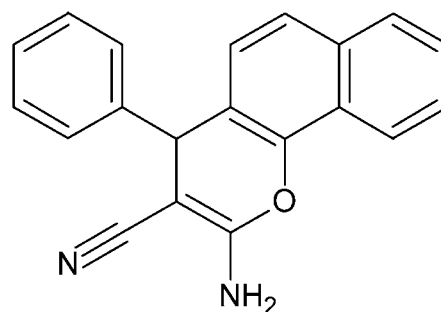
In the title compound, C<sub>20</sub>H<sub>14</sub>N<sub>2</sub>O, the plane of the phenyl ring is almost normal to that of the naphthalene ring system, forming a dihedral angle of 83.15 (8)°. The 4*H*-pyran ring fused with the naphthalene ring system has a flattened boat conformation. In the crystal, molecules are linked by pairs of N—H...N hydrogen bonds, forming inversion dimers with an *R*<sub>2</sub><sup>2</sup>(12) ring motif. The dimers are connected by C—H... $\pi$  interactions, forming supramolecular chains along [010].

**Keywords:** crystal structure; aminochromene; fused chromene; hydrogen bonding; C—H... $\pi$  interactions.

**CCDC reference:** 1406770

## 1. Related literature

For synthesis of chromene-containing compounds, see: Elagamey *et al.* (1988); El-Maghraby (2014). For industrial applications of aminochromenes, see: Ellis (1977); Hafez *et al.* (1987). For various biological activities of fused chromenes, see: Hiramoto *et al.* (1997); Bianchi & Tava (1987); Eiden & Denk (1991); Smith *et al.* (1998); Taylor *et al.* (1998). For the crystal structure of the isomer of the title compound, 3-amino-1-phenyl-1*H*-benzo[*f*]chromene-2-carbonitrile, see: Akkurt *et al.* (2013).



## 2. Experimental

### 2.1. Crystal data

C<sub>20</sub>H<sub>14</sub>N<sub>2</sub>O  
*M<sub>r</sub>* = 298.33  
 Monoclinic, *P*2<sub>1</sub>  
*a* = 9.1662 (1) Å  
*b* = 5.7246 (1) Å  
*c* = 13.9177 (2) Å  
 $\beta$  = 90.153 (1)°  
*V* = 730.30 (2) Å<sup>3</sup>  
*Z* = 2  
 Cu *K* $\alpha$  radiation  
 $\mu$  = 0.67 mm<sup>−1</sup>  
*T* = 100 K  
 0.28 × 0.13 × 0.10 mm

### 2.2. Data collection

Rigaku AFC11 diffractometer  
 Absorption correction: multi-scan  
 (*CrystalClear-SM Expert*; Rigaku, 2012)  
*T*<sub>min</sub> = 0.883, *T*<sub>max</sub> = 1.000  
 5778 measured reflections  
 2201 independent reflections  
 2184 reflections with *I* > 2 $\sigma$ (*I*)  
*R*<sub>int</sub> = 0.029

### 2.3. Refinement

*R* [*F*<sup>2</sup> > 2 $\sigma$ (*F*<sup>2</sup>)] = 0.031  
*wR* (*F*<sup>2</sup>) = 0.093  
*S* = 1.09  
 2201 reflections  
 216 parameters  
 1 restraint  
 H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\text{max}}$  = 0.14 e Å<sup>−3</sup>  
 $\Delta\rho_{\text{min}}$  = −0.15 e Å<sup>−3</sup>  
 Absolute structure: Flack *x* determined using 775 quotients [(*I*<sup>+</sup>) − (*I*<sup>−</sup>)] / [(*I*<sup>+</sup>) + (*I*<sup>−</sup>)] (Parsons *et al.*, 2013)  
 Absolute structure parameter: 0.2 (3)

**Table 1**

Hydrogen-bond geometry (Å, °).

Cg1 is the centroid of the C15–C20 phenyl ring.

<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—H1 <i>B</i> ...N2 <sup>i</sup>	0.91 (3)	2.09 (3)	2.970 (2)	163 (3)
C9—H9...Cg1 <sup>ii</sup>	0.95	2.88	3.574 (2)	131

Symmetry codes: (i)  $-x + 1, y + \frac{1}{2}, -z$ ; (ii)  $-x + 2, y + \frac{1}{2}, -z + 1$ .

Data collection: *CrystalClearSM Expert* (Rigaku, 2012); cell refinement: *CrystalClearSM Expert*; data reduction: *CrystalClearSM Expert*; program(s) used to solve structure: *SUPERFLIP* (Palatinus & Chapuis, 2007); program(s) used to refine structure: *SHELXL2014* (Sheldrick, 2015); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012); software used to prepare material for publication: *WinGX* (Farrugia, 2012).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: TK5369).

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## supporting information

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## Crystal structure of 2-amino-4-phenyl-4*H*-benzo[*h*]chromene-3-carbonitrile

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### S1. Comment

Among synthetic heterocyclic compounds, aminochromenes represent an important class of organic compounds being the main components of many naturally occurring products (Elagamey *et al.*, 1988; El-Maghraby, 2014). They are used for the chemical synthesis of cosmetics, pigments (Ellis, 1977), and potentially biodegradable agrochemicals (Hafez, *et al.*, 1987). Fused chromene systems have displayed a broad spectrum of biological activities such as mutagenicity (Hiramoto, *et al.*, 1997), sex pheromonal (Bianchi & Tava, 1987), central nervous system (CNS) activities (Eiden & Denk, 1991) and inhibitors for influenza virus sialidases (Smith *et al.*, 1998; Taylor *et al.*, 1998). In this context and following our strategy for the synthesis of bio-active molecules, we herein report the synthesis and crystal structure of the title compound.

As seen in Fig. 1, the C4–C13 naphthalene ring system of the title compound is essentially planar [maximum deviations = -0.020 (2) Å for C4 and -0.016 (2) Å for C8]. The C15–C20 phenyl ring makes a dihedral angle of 83.15 (8)° with the mean plane of the naphthalene ring. The 4*H*-pyran ring (O1/C1–C4/C13) in the title compound is puckered [the puckering parameters (Cremer & Pople, 1975) are  $Q_T = 0.177$  (2) Å,  $\theta = 98.2$  (6)° and  $\varphi = 342.9$  (7)°]. The structural geometric parameters of the title compound are normal and are consistent with those of the isomer compound 3-amino-1-phenyl-1*H*-benzo[*f*]chromene-2-carbonitrile (Akkurt *et al.*, 2013). Both isomers crystallize in the same monoclinic space group  $P2_1$  and their unit-cell parameters are almost equal.

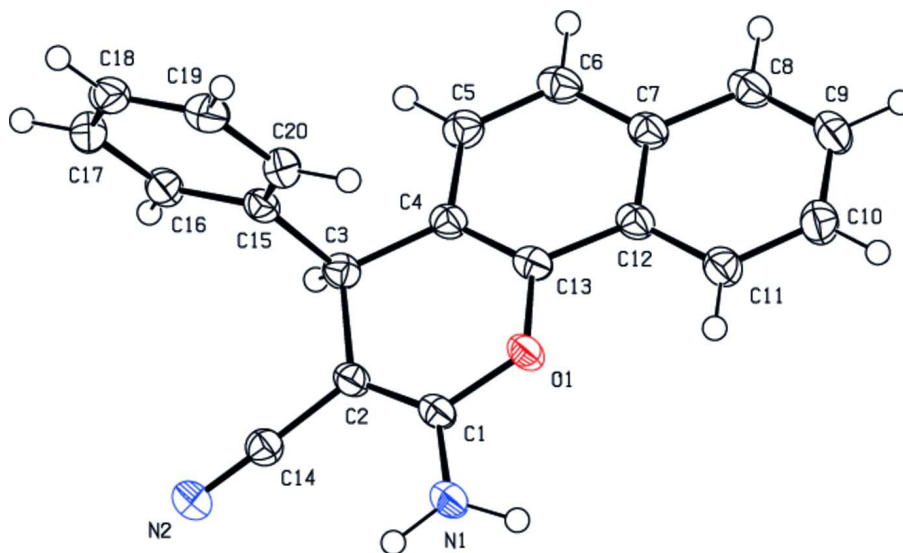
In the crystal, pairs of N—H⋯N hydrogen bonds form inversion dimers with an  $R_2^2(12)$  ring motif (Table 1 and Fig. 2). In addition, C—H⋯ $\pi$  interactions are observed.

### S2. Experimental

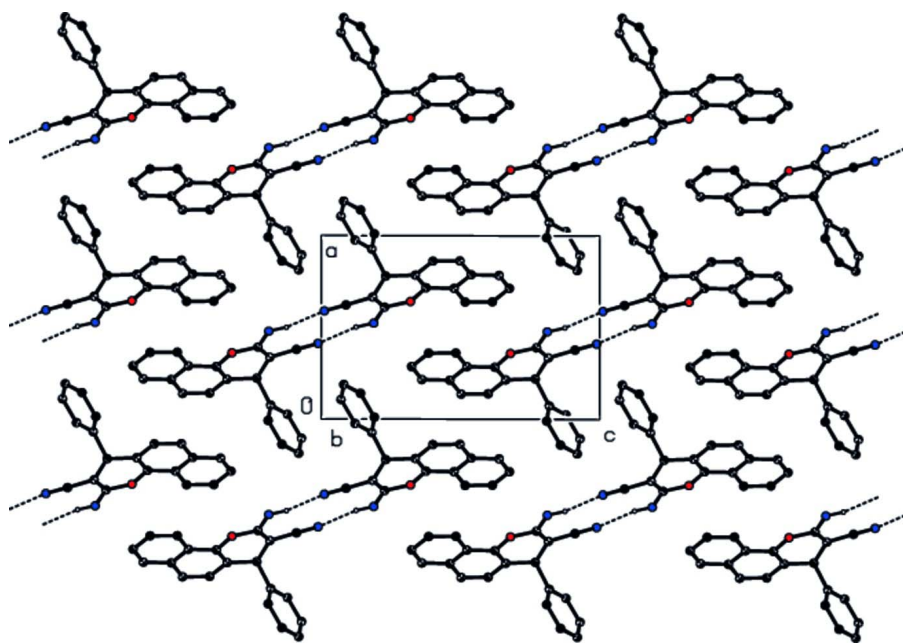
To a solution of 1-naphthol (144 mg; 1 mmol) in 10 ml absolute ethanol, an equimolar amount of benzylidene-malononitrile (154 mg; 1 mmol) was added with constant stirring. The reaction mixture was refluxed for 3 h in the presence of a catalytic amount of piperidine. The reaction progress was monitored by TLC and after cooling, the formed precipitate was filtered off, washed with cold ethanol and dried under vacuum in a desiccator for 24 h. The solid was recrystallized from ethanol. Crystals suitable for X-ray crystallography were obtained by slow evaporation of a solution of the title compound in ethanol (yield 92%; m.p. 483 K).

### S3. Refinement

The H atoms of the NH<sub>2</sub> group were refined. The H atoms attached to the C atoms were positioned geometrically, with C—H = 0.95 Å and C—H = 1.00 Å for aromatic and methine H, respectively, and with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ .

**Figure 1**

View of the title compound with the atom-numbering scheme. Displacement ellipsoids for non-H atoms are drawn at the 50% probability level.

**Figure 2**

View of the dimers formed by N—H...O hydrogen bonds.

## 2-Amino-4-phenyl-4H-benzo[h]chromene-3-carbonitrile

### Crystal data

$C_{20}H_{14}N_2O$

$M_r = 298.33$

Monoclinic,  $P2_1$

Hall symbol: P 2yb

$a = 9.1662 (1) \text{ \AA}$

$b = 5.7246 (1) \text{ \AA}$

$c = 13.9177 (2) \text{ \AA}$

$\beta = 90.153 (1)^\circ$

$V = 730.30 (2) \text{ \AA}^3$   
 $Z = 2$   
 $F(000) = 312$   
 $D_x = 1.357 \text{ Mg m}^{-3}$   
 Cu  $K\alpha$  radiation,  $\lambda = 1.54178 \text{ \AA}$   
 Cell parameters from 5573 reflections

$\theta = 6.3\text{--}66.6^\circ$   
 $\mu = 0.67 \text{ mm}^{-1}$   
 $T = 100 \text{ K}$   
 Block, brown  
 $0.28 \times 0.13 \times 0.10 \text{ mm}$

#### Data collection

Rigaku AFC11  
 diffractometer  
 Radiation source: Rotating Anode  
 Detector resolution:  $22.2222 \text{ pixels mm}^{-1}$   
 profile data from  $\omega$ -scans  
 Absorption correction: multi-scan  
 (*CrystalClear-SM Expert*; Rigaku, 2012)  
 $T_{\min} = 0.883$ ,  $T_{\max} = 1.000$

5778 measured reflections  
 2201 independent reflections  
 2184 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.029$   
 $\theta_{\max} = 66.7^\circ$ ,  $\theta_{\min} = 4.8^\circ$   
 $h = -10 \rightarrow 10$   
 $k = -6 \rightarrow 6$   
 $l = -16 \rightarrow 16$

#### Refinement

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.031$   
 $wR(F^2) = 0.093$   
 $S = 1.09$   
 2201 reflections  
 216 parameters  
 1 restraint  
 Hydrogen site location: mixed

H atoms treated by a mixture of independent  
 and constrained refinement  
 $w = 1/[\sigma^2(FO^2) + (0.0642P)^2 + 0.1186P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.14 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.15 \text{ e \AA}^{-3}$   
 Absolute structure: Flack  $x$  determined using  
 775 quotients  $[(I^+)-(I^-)]/[(I^+)+(I^-)]$  (Parsons *et al.*,  
 2013)  
 Absolute structure parameter: 0.2 (3)

#### Special details

**Geometry.** Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

**Refinement.** Refinement on  $F^2$  for ALL reflections except those flagged by the user for potential systematic errors. Weighted  $R$ -factors  $wR$  and all goodnesses 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 observed criterion of  $F^2 > \sigma(F^2)$  is used only for calculating  $-R$ -factor-obs *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.

#### Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.64120 (15)	0.8755 (3)	0.32053 (9)	0.0250 (4)
N1	0.5228 (2)	0.9831 (4)	0.18837 (13)	0.0279 (6)
N2	0.58898 (19)	0.5238 (4)	0.01142 (12)	0.0305 (6)
C1	0.6097 (2)	0.8195 (4)	0.22720 (13)	0.0223 (5)
C2	0.6647 (2)	0.6252 (4)	0.18462 (14)	0.0228 (6)
C3	0.7792 (2)	0.4700 (4)	0.23142 (13)	0.0222 (6)
C4	0.7851 (2)	0.5235 (4)	0.33777 (13)	0.0225 (6)
C5	0.8612 (2)	0.3703 (4)	0.40074 (14)	0.0264 (6)
C6	0.8631 (2)	0.4068 (4)	0.49782 (15)	0.0286 (6)
C7	0.7877 (2)	0.5997 (4)	0.53815 (14)	0.0255 (6)

C8	0.7844 (2)	0.6408 (5)	0.63859 (14)	0.0297 (6)
C9	0.7137 (2)	0.8309 (5)	0.67554 (14)	0.0301 (6)
C10	0.6403 (2)	0.9868 (5)	0.61433 (15)	0.0307 (6)
C11	0.6388 (2)	0.9514 (4)	0.51680 (14)	0.0275 (6)
C12	0.7134 (2)	0.7592 (4)	0.47686 (14)	0.0239 (6)
C13	0.7165 (2)	0.7126 (4)	0.37642 (13)	0.0226 (6)
C14	0.6217 (2)	0.5720 (4)	0.08901 (13)	0.0241 (6)
C15	0.9263 (2)	0.4943 (4)	0.18151 (13)	0.0222 (6)
C16	0.9754 (2)	0.3182 (4)	0.12136 (13)	0.0261 (6)
C17	1.1057 (2)	0.3423 (5)	0.07107 (14)	0.0301 (6)
C18	1.1876 (2)	0.5433 (5)	0.08102 (14)	0.0301 (6)
C19	1.1396 (2)	0.7208 (5)	0.14112 (14)	0.0293 (6)
C20	1.0097 (2)	0.6961 (4)	0.19070 (14)	0.0260 (6)
H1A	0.474 (3)	1.088 (5)	0.2297 (18)	0.034 (7)*
H1B	0.501 (3)	0.971 (6)	0.125 (2)	0.040 (7)*
H3	0.74660	0.30430	0.22390	0.0270*
H5	0.91180	0.24000	0.37480	0.0320*
H6	0.91520	0.30260	0.53850	0.0340*
H8	0.83200	0.53470	0.68080	0.0360*
H9	0.71420	0.85760	0.74290	0.0360*
H10	0.59130	1.11800	0.64070	0.0370*
H11	0.58740	1.05650	0.47620	0.0330*
H16	0.91940	0.17950	0.11430	0.0310*
H17	1.13830	0.22050	0.03000	0.0360*
H18	1.27650	0.56020	0.04680	0.0360*
H19	1.19580	0.85910	0.14820	0.0350*
H20	0.97710	0.81850	0.23150	0.0310*

*Atomic displacement parameters ( $\text{\AA}^2$ )*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O1	0.0306 (8)	0.0258 (8)	0.0186 (6)	0.0031 (7)	−0.0059 (5)	−0.0008 (6)
N1	0.0309 (9)	0.0322 (11)	0.0205 (9)	0.0045 (9)	−0.0056 (7)	−0.0012 (8)
N2	0.0297 (9)	0.0375 (12)	0.0243 (9)	0.0007 (9)	−0.0046 (7)	−0.0041 (8)
C1	0.0219 (9)	0.0273 (11)	0.0178 (8)	−0.0041 (9)	−0.0026 (7)	0.0026 (8)
C2	0.0202 (9)	0.0293 (12)	0.0190 (9)	−0.0025 (9)	−0.0022 (7)	0.0002 (8)
C3	0.0226 (9)	0.0205 (10)	0.0236 (10)	−0.0015 (9)	−0.0021 (7)	−0.0009 (8)
C4	0.0191 (9)	0.0272 (12)	0.0213 (9)	−0.0039 (8)	−0.0007 (7)	0.0010 (8)
C5	0.0249 (9)	0.0270 (12)	0.0272 (10)	0.0003 (10)	0.0001 (7)	0.0025 (9)
C6	0.0276 (10)	0.0325 (13)	0.0256 (10)	−0.0010 (10)	−0.0044 (8)	0.0074 (9)
C7	0.0220 (9)	0.0321 (12)	0.0223 (9)	−0.0050 (9)	−0.0013 (7)	0.0029 (9)
C8	0.0259 (10)	0.0414 (14)	0.0219 (9)	−0.0058 (10)	−0.0031 (8)	0.0058 (9)
C9	0.0275 (10)	0.0427 (14)	0.0200 (9)	−0.0106 (11)	0.0002 (7)	−0.0028 (10)
C10	0.0296 (10)	0.0356 (13)	0.0268 (10)	−0.0044 (11)	0.0024 (8)	−0.0060 (9)
C11	0.0280 (10)	0.0298 (12)	0.0246 (10)	−0.0018 (10)	−0.0018 (8)	−0.0021 (9)
C12	0.0202 (9)	0.0301 (12)	0.0214 (9)	−0.0064 (9)	−0.0012 (7)	0.0006 (8)
C13	0.0214 (9)	0.0251 (11)	0.0213 (9)	−0.0029 (9)	−0.0031 (7)	0.0036 (8)
C14	0.0226 (9)	0.0263 (11)	0.0235 (10)	−0.0010 (9)	0.0001 (7)	0.0003 (9)

C15	0.0227 (10)	0.0265 (11)	0.0173 (8)	0.0004 (9)	−0.0025 (7)	0.0029 (8)
C16	0.0301 (10)	0.0250 (12)	0.0231 (9)	0.0024 (10)	−0.0032 (7)	−0.0016 (8)
C17	0.0319 (11)	0.0346 (13)	0.0238 (9)	0.0069 (10)	0.0008 (7)	−0.0020 (10)
C18	0.0245 (10)	0.0429 (14)	0.0230 (9)	0.0041 (10)	0.0002 (7)	0.0067 (9)
C19	0.0264 (10)	0.0328 (12)	0.0286 (10)	−0.0040 (10)	−0.0030 (8)	0.0056 (9)
C20	0.0274 (10)	0.0263 (12)	0.0244 (10)	0.0001 (10)	−0.0001 (8)	−0.0024 (9)

*Geometric parameters (Å, °)*

O1—C1	1.368 (2)	C11—C12	1.410 (3)
O1—C13	1.396 (3)	C12—C13	1.424 (3)
N1—C1	1.342 (3)	C15—C20	1.391 (3)
N2—C14	1.154 (3)	C15—C16	1.386 (3)
C1—C2	1.358 (3)	C16—C17	1.393 (3)
N1—H1B	0.91 (3)	C17—C18	1.381 (4)
N1—H1A	0.95 (3)	C18—C19	1.388 (3)
C2—C14	1.420 (3)	C19—C20	1.385 (3)
C2—C3	1.520 (3)	C3—H3	1.0000
C3—C4	1.512 (3)	C5—H5	0.9500
C3—C15	1.525 (3)	C6—H6	0.9500
C4—C13	1.363 (3)	C8—H8	0.9500
C4—C5	1.422 (3)	C9—H9	0.9500
C5—C6	1.367 (3)	C10—H10	0.9500
C6—C7	1.419 (3)	C11—H11	0.9500
C7—C12	1.422 (3)	C16—H16	0.9500
C7—C8	1.418 (3)	C17—H17	0.9500
C8—C9	1.368 (4)	C18—H18	0.9500
C9—C10	1.404 (3)	C19—H19	0.9500
C10—C11	1.373 (3)	C20—H20	0.9500
C1—O1—C13	118.39 (17)	C3—C15—C20	121.36 (18)
O1—C1—N1	110.02 (18)	C16—C15—C20	118.70 (17)
O1—C1—C2	121.91 (18)	C15—C16—C17	120.8 (2)
N1—C1—C2	128.06 (18)	C16—C17—C18	119.9 (2)
C1—N1—H1B	118 (2)	C17—C18—C19	119.82 (18)
H1A—N1—H1B	123 (3)	C18—C19—C20	120.0 (2)
C1—N1—H1A	118.7 (16)	C15—C20—C19	120.8 (2)
C1—C2—C3	123.25 (17)	C2—C3—H3	108.00
C1—C2—C14	118.83 (19)	C4—C3—H3	108.00
C3—C2—C14	117.76 (18)	C15—C3—H3	108.00
C2—C3—C15	111.21 (16)	C4—C5—H5	119.00
C4—C3—C15	113.46 (15)	C6—C5—H5	119.00
C2—C3—C4	108.89 (17)	C5—C6—H6	120.00
C3—C4—C13	122.12 (18)	C7—C6—H6	120.00
C3—C4—C5	119.65 (19)	C7—C8—H8	120.00
C5—C4—C13	118.21 (17)	C9—C8—H8	120.00
C4—C5—C6	121.3 (2)	C8—C9—H9	120.00
C5—C6—C7	120.32 (19)	C10—C9—H9	120.00

C8—C7—C12	118.3 (2)	C9—C10—H10	120.00
C6—C7—C8	122.1 (2)	C11—C10—H10	120.00
C6—C7—C12	119.69 (18)	C10—C11—H11	120.00
C7—C8—C9	120.9 (2)	C12—C11—H11	120.00
C8—C9—C10	120.29 (19)	C15—C16—H16	120.00
C9—C10—C11	120.6 (2)	C17—C16—H16	120.00
C10—C11—C12	120.1 (2)	C16—C17—H17	120.00
C11—C12—C13	122.97 (19)	C18—C17—H17	120.00
C7—C12—C11	119.76 (18)	C17—C18—H18	120.00
C7—C12—C13	117.26 (19)	C19—C18—H18	120.00
O1—C13—C12	114.26 (18)	C18—C19—H19	120.00
O1—C13—C4	122.59 (16)	C20—C19—H19	120.00
C4—C13—C12	123.14 (19)	C15—C20—H20	120.00
N2—C14—C2	178.3 (2)	C19—C20—H20	120.00
C3—C15—C16	119.85 (19)		
C13—O1—C1—N1	172.70 (17)	C4—C5—C6—C7	0.3 (3)
C13—O1—C1—C2	−8.3 (3)	C5—C6—C7—C8	178.6 (2)
C1—O1—C13—C4	13.6 (3)	C5—C6—C7—C12	−1.8 (3)
C1—O1—C13—C12	−165.44 (17)	C6—C7—C8—C9	178.5 (2)
O1—C1—C2—C3	−7.7 (3)	C12—C7—C8—C9	−1.1 (3)
O1—C1—C2—C14	176.97 (18)	C6—C7—C12—C11	−179.80 (18)
N1—C1—C2—C3	171.1 (2)	C6—C7—C12—C13	1.3 (3)
N1—C1—C2—C14	−4.2 (3)	C8—C7—C12—C11	−0.2 (3)
C1—C2—C3—C4	16.6 (3)	C8—C7—C12—C13	−179.15 (19)
C1—C2—C3—C15	−109.1 (2)	C7—C8—C9—C10	1.3 (3)
C14—C2—C3—C4	−168.01 (18)	C8—C9—C10—C11	−0.2 (3)
C14—C2—C3—C15	66.3 (2)	C9—C10—C11—C12	−1.1 (3)
C2—C3—C4—C5	167.12 (18)	C10—C11—C12—C7	1.3 (3)
C2—C3—C4—C13	−11.3 (3)	C10—C11—C12—C13	−179.9 (2)
C15—C3—C4—C5	−68.5 (3)	C7—C12—C13—O1	179.84 (17)
C15—C3—C4—C13	113.1 (2)	C7—C12—C13—C4	0.8 (3)
C2—C3—C15—C16	−105.1 (2)	C11—C12—C13—O1	0.9 (3)
C2—C3—C15—C20	71.3 (2)	C11—C12—C13—C4	−178.1 (2)
C4—C3—C15—C16	131.7 (2)	C3—C15—C16—C17	176.54 (18)
C4—C3—C15—C20	−51.9 (3)	C20—C15—C16—C17	0.1 (3)
C3—C4—C5—C6	−176.86 (18)	C3—C15—C20—C19	−176.69 (18)
C13—C4—C5—C6	1.6 (3)	C16—C15—C20—C19	−0.3 (3)
C3—C4—C13—O1	−2.7 (3)	C15—C16—C17—C18	0.1 (3)
C3—C4—C13—C12	176.26 (18)	C16—C17—C18—C19	−0.1 (3)
C5—C4—C13—O1	178.81 (18)	C17—C18—C19—C20	−0.2 (3)
C5—C4—C13—C12	−2.2 (3)	C18—C19—C20—C15	0.3 (3)

### Hydrogen-bond geometry ( $\text{\AA}$ , $^\circ$ )

Cg1 is the centroid of the C15–C20 phenyl ring.

$D\cdots H\cdots A$	$D\cdots H$	$H\cdots A$	$D\cdots A$	$D\cdots H\cdots A$
N1—H1B $\cdots$ N2 <sup>i</sup>	0.91 (3)	2.09 (3)	2.970 (2)	163 (3)



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C9—H9...Cg1 <sup>ii</sup>	0.95	2.88	3.574 (2)	131
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Symmetry codes: (i)  $-x+1, y+1/2, -z$ ; (ii)  $-x+2, y+1/2, -z+1$ .