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4-(4-Chlorobenzyl)-5-methyl-2-phenyl-1H-pyrazol-3(2H)-one

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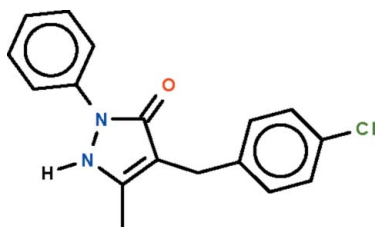
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 Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.036; wR factor = 0.097; data-to-parameter ratio = 13.5.

The five-membered ring of the title compound, $\text{C}_{17}\text{H}_{15}\text{ClN}_2\text{O}$, is almost planar (r.m.s. deviation = 0.008 Å), and its phenyl substituent is aligned at 34.9 (1)° with respect to this ring. The angle at the methylene C atom is opened to 116.4 (2)°. In the crystal, adjacent molecules are linked by an $\text{N}-\text{H}\cdots\text{O}$ hydrogen bond, generating a linear chain along the a axis.

Related literature

 For the synthesis, see: Pettinari *et al.* (1994).


Experimental

Crystal data

 $\text{C}_{17}\text{H}_{15}\text{ClN}_2\text{O}$
 $M_r = 298.76$

 Orthorhombic, $Fdd2$
 $a = 23.1540$ (3) Å

 $b = 43.8905$ (6) Å
 $c = 5.6239$ (1) Å
 $V = 5715.23$ (15) Å³
 $Z = 16$

 Cu $K\alpha$ radiation
 $\mu = 2.36$ mm⁻¹
 $T = 100$ K
 $0.30 \times 0.30 \times 0.03$ mm

Data collection

 Agilent SuperNova Dual
 diffractometer with an Atlas
 detector
 Absorption correction: multi-scan
 (*CrysAlis PRO*; Agilent, 2010)
 $T_{\min} = 0.538$, $T_{\max} = 0.933$

 10182 measured reflections
 2623 independent reflections
 2611 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.028$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.097$
 $S = 1.08$
 2623 reflections
 195 parameters
 1 restraint

 H atoms treated by a mixture of
 independent and constrained
 refinement
 $\Delta\rho_{\text{max}} = 0.21$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.41$ e Å⁻³
 Absolute structure: Flack (1983),
 1011 Friedel pairs
 Flack parameter: 0.000 (12)

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N2}-\text{H2}\cdots\text{O1}^i$	0.85 (3)	1.82 (3)	2.6516 (18)	165 (2)

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

Data collection: *CrysAlis PRO* (Agilent, 2010); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *publCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT5516).

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supporting information

Acta Cryst. (2011). E67, o1153 [doi:10.1107/S1600536811013791]

4-(4-Chlorobenzyl)-5-methyl-2-phenyl-1H-pyrazol-3(2H)-one

Shaaban K. Mohamed, Mahmoud A. A. El-Remaly, Ahmed M. Soliman, Hossam Abdel-Ghany and Seik Weng Ng

S1. Comment

3-Methyl-1-phenyl-4,5-dihydro-1H-5-pyrazolone possesses an active methylene linkage that undergoes condensation with aromatic aldehydes to yield compounds that react with metal salts (Pettinari *et al.*, 1994). In these organic compounds, the pyrazole ring is connected to the aromatic system (of the aldehyde precursor) by a methylene linkage. The five-membered ring of C₁₇H₁₅ClN₂O (Scheme I) is planar, and its phenyl substituent is aligned at 34.9 (1) ° with respect to this ring. The angle at the methylene C atom is opened to 116.4 (2) ° (Fig. 1). Adjacent molecules are linked by an N–H···O hydrogen bond to generate a linear chain along the *a*-axis of the orthorhombic unit cell (Fig. 2).

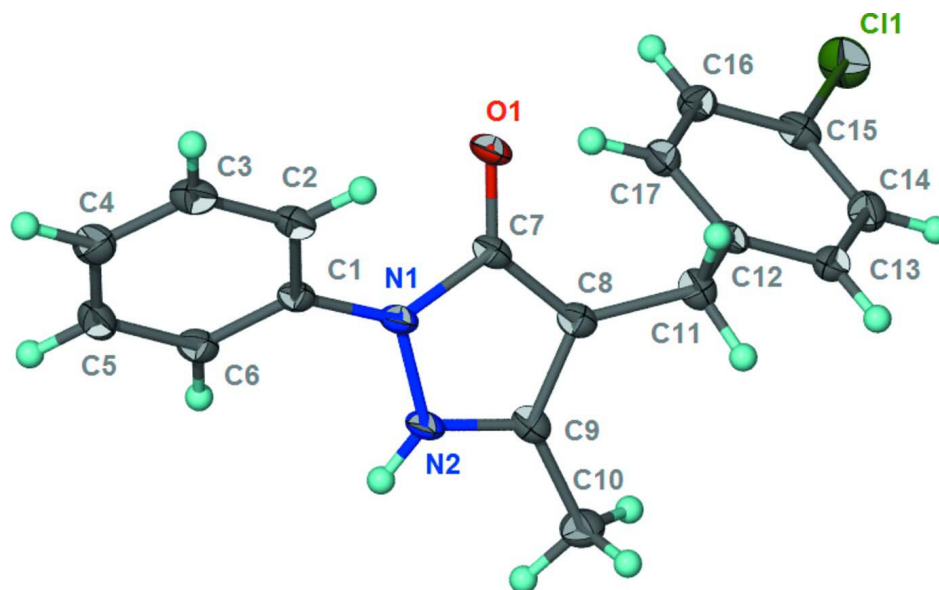
S2. Experimental

3-Methyl-1-phenyl-4,5-dihydro-1H-5-pyrazolone (10 mmol) and 4-chlorobenzaldehyde (10 mmol) along with few drops of concentrated hydrochloric acid were heated at 426 K in *N,N*-dimethylformamide (50 ml) for 8 h. The product was collected and recrystallized from ethanol; m.p. 471 K.

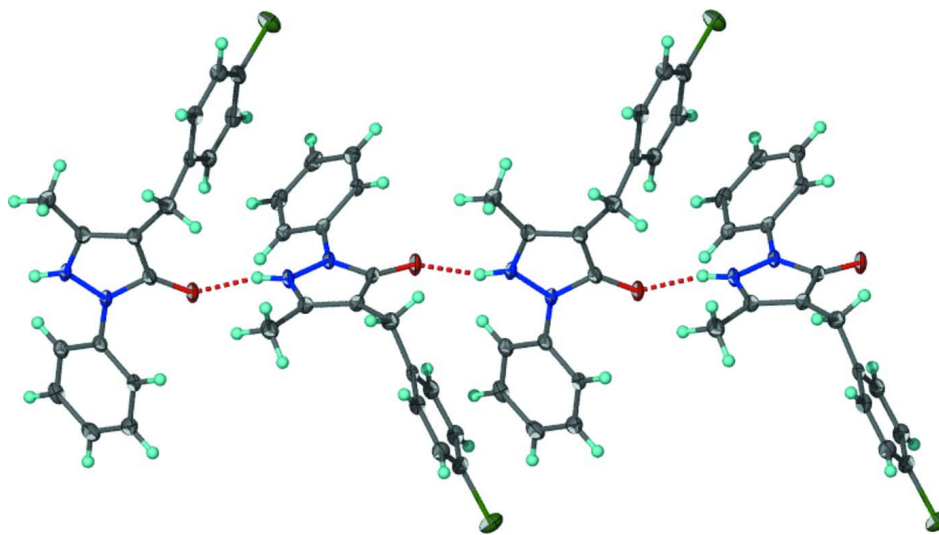
S3. Refinement

Carbon-bound H-atoms were placed in calculated positions [C—H 0.95 to 0.98 Å, $U_{\text{iso}}(\text{H})$ 1.2 to 1.5 $U_{\text{eq}}(\text{C})$] and were included in the refinement in the riding model approximation.

The amino H-atom was located in a difference Fourier map, and was freely refined.

**Figure 1**

Anisotropic displacement ellipsoid plot (Barbour, 2001) of $C_{17}H_{15}ClN_2O$ at the 70% probability level; hydrogen atoms are drawn as spheres of arbitrary radius.

**Figure 2**

Hydrogen-bonded chain structure.

4-(4-Chlorobenzyl)-5-methyl-2-phenyl-1*H*-pyrazol-3(2*H*)-one

Crystal data

$C_{17}H_{15}ClN_2O$

$M_r = 298.76$

Orthorhombic, *Fdd2*

Hall symbol: *F* 2 -2*d*

$a = 23.1540$ (3) Å

$b = 43.8905$ (6) Å

$c = 5.6239$ (1) Å

$V = 5715.23$ (15) Å³

$Z = 16$

$F(000) = 2496$

$D_x = 1.389$ Mg m⁻³

Cu $K\alpha$ radiation, $\lambda = 1.54184$ Å

Cell parameters from 8414 reflections

$\theta = 3.8$ – 74.2°

$\mu = 2.36 \text{ mm}^{-1}$
 $T = 100 \text{ K}$

Plate, colorless
 $0.30 \times 0.30 \times 0.03 \text{ mm}$

Data collection

Agilent SuperNova Dual
 diffractometer with an Atlas detector
 Radiation source: SuperNova (Cu) X-ray
 Source
 Mirror monochromator
 Detector resolution: $10.4041 \text{ pixels mm}^{-1}$
 ω scans
 Absorption correction: multi-scan
 (CrysAlis PRO; Agilent, 2010)

$T_{\min} = 0.538$, $T_{\max} = 0.933$
 10182 measured reflections
 2623 independent reflections
 2611 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.028$
 $\theta_{\max} = 74.3^\circ$, $\theta_{\min} = 4.0^\circ$
 $h = -24 \rightarrow 28$
 $k = -53 \rightarrow 54$
 $l = -6 \rightarrow 6$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.097$
 $S = 1.08$
 2623 reflections
 195 parameters
 1 restraint
 Primary atom site location: structure-invariant
 direct methods
 Secondary atom site location: difference Fourier
 map

Hydrogen site location: inferred from
 neighbouring sites
 H atoms treated by a mixture of independent
 and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0771P)^2 + 3.0966P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.21 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.41 \text{ e } \text{\AA}^{-3}$
 Absolute structure: Flack (1983), 1011 Friedel
 pairs
 Absolute structure parameter: 0.000 (12)

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Cl1	0.122967 (18)	0.029555 (10)	0.59020 (10)	0.02795 (14)
O1	0.20050 (5)	-0.13339 (3)	0.3426 (3)	0.0201 (3)
N1	0.29664 (5)	-0.13658 (3)	0.2283 (3)	0.0152 (3)
N2	0.34778 (6)	-0.12275 (3)	0.2955 (3)	0.0161 (3)
H2	0.3806 (10)	-0.1243 (5)	0.228 (5)	0.019 (5)*
C1	0.29551 (6)	-0.16087 (4)	0.0632 (3)	0.0143 (3)
C2	0.25718 (7)	-0.18504 (4)	0.1001 (3)	0.0168 (3)
H2A	0.2333	-0.1854	0.2373	0.020*
C3	0.25436 (7)	-0.20838 (4)	-0.0648 (4)	0.0200 (4)
H3	0.2279	-0.2247	-0.0416	0.024*
C4	0.28990 (7)	-0.20825 (4)	-0.2645 (4)	0.0223 (4)
H4	0.2875	-0.2243	-0.3776	0.027*
C5	0.32888 (7)	-0.18450 (4)	-0.2971 (4)	0.0205 (4)
H5	0.3537	-0.1845	-0.4316	0.025*
C6	0.33170 (7)	-0.16074 (4)	-0.1343 (3)	0.0169 (3)
H6	0.3582	-0.1445	-0.1577	0.020*
C7	0.25237 (6)	-0.12512 (4)	0.3689 (3)	0.0150 (3)
C8	0.27831 (7)	-0.10413 (4)	0.5272 (3)	0.0156 (3)
C9	0.33640 (7)	-0.10305 (4)	0.4713 (3)	0.0162 (3)
C10	0.38394 (7)	-0.08494 (4)	0.5800 (4)	0.0210 (4)
H10A	0.4167	-0.0841	0.4700	0.031*

H10B	0.3961	-0.0946	0.7290	0.031*
H10C	0.3703	-0.0642	0.6126	0.031*
C11	0.24961 (7)	-0.08915 (4)	0.7360 (3)	0.0178 (3)
H11A	0.2789	-0.0863	0.8621	0.021*
H11B	0.2200	-0.1033	0.7989	0.021*
C12	0.22098 (6)	-0.05861 (4)	0.6893 (3)	0.0154 (3)
C13	0.22392 (7)	-0.03570 (4)	0.8595 (3)	0.0172 (3)
H13	0.2465	-0.0388	0.9985	0.021*
C14	0.19446 (7)	-0.00837 (4)	0.8301 (4)	0.0193 (3)
H14	0.1968	0.0072	0.9467	0.023*
C15	0.16151 (7)	-0.00439 (4)	0.6266 (3)	0.0188 (4)
C16	0.15824 (7)	-0.02657 (4)	0.4517 (3)	0.0184 (3)
H16	0.1357	-0.0234	0.3128	0.022*
C17	0.18871 (7)	-0.05357 (4)	0.4839 (3)	0.0175 (3)
H17	0.1875	-0.0688	0.3640	0.021*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.0289 (2)	0.0233 (2)	0.0317 (3)	0.01125 (15)	-0.00111 (18)	0.00169 (18)
O1	0.0092 (5)	0.0249 (6)	0.0261 (7)	-0.0009 (4)	0.0031 (5)	-0.0062 (6)
N1	0.0089 (6)	0.0176 (6)	0.0191 (7)	-0.0009 (5)	0.0009 (5)	-0.0015 (6)
N2	0.0086 (6)	0.0191 (6)	0.0206 (8)	-0.0025 (5)	0.0023 (6)	-0.0025 (6)
C1	0.0110 (6)	0.0154 (7)	0.0164 (8)	0.0031 (5)	-0.0018 (6)	0.0000 (6)
C2	0.0117 (6)	0.0168 (7)	0.0220 (9)	0.0010 (5)	0.0014 (7)	0.0001 (7)
C3	0.0148 (7)	0.0174 (8)	0.0277 (10)	0.0010 (6)	-0.0018 (7)	-0.0003 (7)
C4	0.0190 (8)	0.0216 (8)	0.0262 (10)	0.0053 (6)	-0.0021 (7)	-0.0062 (8)
C5	0.0167 (7)	0.0263 (8)	0.0185 (9)	0.0057 (6)	0.0021 (7)	-0.0009 (8)
C6	0.0128 (7)	0.0187 (7)	0.0191 (9)	0.0026 (5)	-0.0007 (6)	0.0021 (7)
C7	0.0118 (7)	0.0157 (7)	0.0176 (9)	0.0027 (5)	0.0021 (6)	0.0020 (6)
C8	0.0133 (7)	0.0159 (7)	0.0177 (9)	0.0015 (6)	0.0006 (6)	0.0020 (6)
C9	0.0147 (7)	0.0162 (7)	0.0177 (9)	0.0010 (6)	0.0024 (7)	0.0018 (7)
C10	0.0174 (7)	0.0213 (8)	0.0242 (10)	-0.0036 (6)	-0.0005 (7)	-0.0023 (8)
C11	0.0185 (7)	0.0174 (7)	0.0176 (9)	0.0018 (6)	0.0024 (7)	0.0005 (7)
C12	0.0104 (6)	0.0178 (8)	0.0178 (9)	-0.0013 (5)	0.0042 (6)	0.0001 (6)
C13	0.0145 (7)	0.0212 (7)	0.0157 (9)	-0.0005 (6)	0.0003 (6)	-0.0006 (7)
C14	0.0188 (7)	0.0190 (7)	0.0203 (9)	-0.0009 (6)	0.0023 (7)	-0.0030 (7)
C15	0.0150 (7)	0.0185 (8)	0.0229 (10)	0.0032 (6)	0.0029 (6)	0.0023 (7)
C16	0.0154 (7)	0.0250 (9)	0.0149 (8)	0.0001 (6)	-0.0009 (6)	0.0019 (7)
C17	0.0149 (7)	0.0200 (8)	0.0177 (9)	-0.0015 (6)	0.0028 (6)	-0.0030 (7)

Geometric parameters (Å, °)

Cl1—C15	1.7488 (16)	C8—C9	1.382 (2)
O1—C7	1.263 (2)	C8—C11	1.501 (2)
N1—N2	1.3833 (18)	C9—C10	1.489 (2)
N1—C7	1.389 (2)	C10—H10A	0.9800
N1—C1	1.414 (2)	C10—H10B	0.9800

N2—C9	1.340 (2)	C10—H10C	0.9800
N2—H2	0.85 (3)	C11—C12	1.519 (2)
C1—C6	1.391 (2)	C11—H11A	0.9900
C1—C2	1.398 (2)	C11—H11B	0.9900
C2—C3	1.383 (3)	C12—C13	1.390 (2)
C2—H2A	0.9500	C12—C17	1.394 (3)
C3—C4	1.392 (3)	C13—C14	1.390 (2)
C3—H3	0.9500	C13—H13	0.9500
C4—C5	1.391 (3)	C14—C15	1.386 (3)
C4—H4	0.9500	C14—H14	0.9500
C5—C6	1.389 (3)	C15—C16	1.386 (3)
C5—H5	0.9500	C16—C17	1.391 (2)
C6—H6	0.9500	C16—H16	0.9500
C7—C8	1.415 (2)	C17—H17	0.9500
N2—N1—C7	108.49 (14)	C8—C9—C10	130.13 (17)
N2—N1—C1	121.74 (13)	C9—C10—H10A	109.5
C7—N1—C1	129.29 (13)	C9—C10—H10B	109.5
C9—N2—N1	108.46 (13)	H10A—C10—H10B	109.5
C9—N2—H2	123.9 (16)	C9—C10—H10C	109.5
N1—N2—H2	127.2 (16)	H10A—C10—H10C	109.5
C6—C1—C2	120.26 (16)	H10B—C10—H10C	109.5
C6—C1—N1	120.65 (14)	C8—C11—C12	116.41 (15)
C2—C1—N1	119.09 (15)	C8—C11—H11A	108.2
C3—C2—C1	119.48 (16)	C12—C11—H11A	108.2
C3—C2—H2A	120.3	C8—C11—H11B	108.2
C1—C2—H2A	120.3	C12—C11—H11B	108.2
C2—C3—C4	120.66 (16)	H11A—C11—H11B	107.3
C2—C3—H3	119.7	C13—C12—C17	118.84 (15)
C4—C3—H3	119.7	C13—C12—C11	119.89 (15)
C5—C4—C3	119.54 (17)	C17—C12—C11	121.16 (16)
C5—C4—H4	120.2	C12—C13—C14	121.22 (16)
C3—C4—H4	120.2	C12—C13—H13	119.4
C6—C5—C4	120.39 (17)	C14—C13—H13	119.4
C6—C5—H5	119.8	C15—C14—C13	118.51 (16)
C4—C5—H5	119.8	C15—C14—H14	120.7
C5—C6—C1	119.65 (15)	C13—C14—H14	120.7
C5—C6—H6	120.2	C14—C15—C16	121.82 (15)
C1—C6—H6	120.2	C14—C15—C11	119.01 (13)
O1—C7—N1	122.06 (16)	C16—C15—C11	119.16 (14)
O1—C7—C8	131.63 (15)	C15—C16—C17	118.58 (17)
N1—C7—C8	106.30 (13)	C15—C16—H16	120.7
C9—C8—C7	107.01 (15)	C17—C16—H16	120.7
C9—C8—C11	126.41 (16)	C16—C17—C12	121.00 (17)
C7—C8—C11	126.12 (15)	C16—C17—H17	119.5
N2—C9—C8	109.69 (14)	C12—C17—H17	119.5
N2—C9—C10	120.14 (14)		

C7—N1—N2—C9	-0.45 (19)	N1—C7—C8—C11	-170.82 (16)
C1—N1—N2—C9	-173.26 (14)	N1—N2—C9—C8	1.6 (2)
N2—N1—C1—C6	-39.1 (2)	N1—N2—C9—C10	179.61 (16)
C7—N1—C1—C6	149.68 (17)	C7—C8—C9—N2	-2.2 (2)
N2—N1—C1—C2	141.26 (16)	C11—C8—C9—N2	170.45 (15)
C7—N1—C1—C2	-29.9 (2)	C7—C8—C9—C10	-179.87 (18)
C6—C1—C2—C3	-1.8 (2)	C11—C8—C9—C10	-7.3 (3)
N1—C1—C2—C3	177.77 (15)	C9—C8—C11—C12	97.3 (2)
C1—C2—C3—C4	1.0 (2)	C7—C8—C11—C12	-91.5 (2)
C2—C3—C4—C5	0.5 (3)	C8—C11—C12—C13	-142.06 (16)
C3—C4—C5—C6	-1.2 (3)	C8—C11—C12—C17	41.7 (2)
C4—C5—C6—C1	0.4 (2)	C17—C12—C13—C14	1.3 (2)
C2—C1—C6—C5	1.1 (2)	C11—C12—C13—C14	-174.99 (15)
N1—C1—C6—C5	-178.48 (15)	C12—C13—C14—C15	0.4 (2)
N2—N1—C7—O1	178.60 (15)	C13—C14—C15—C16	-1.3 (2)
C1—N1—C7—O1	-9.3 (3)	C13—C14—C15—C11	178.64 (13)
N2—N1—C7—C8	-0.87 (19)	C14—C15—C16—C17	0.5 (3)
C1—N1—C7—C8	171.23 (16)	C11—C15—C16—C17	-179.46 (13)
O1—C7—C8—C9	-177.57 (18)	C15—C16—C17—C12	1.3 (2)
N1—C7—C8—C9	1.82 (19)	C13—C12—C17—C16	-2.2 (2)
O1—C7—C8—C11	9.8 (3)	C11—C12—C17—C16	174.10 (15)

Hydrogen-bond geometry (Å, °)

<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>
N2—H2...O1 ⁱ	0.85 (3)	1.82 (3)	2.6516 (18)	165 (2)

Symmetry code: (i) $x+1/4, -y-1/4, z-1/4$.