

## Ethyl 5-ethoxymethylene-2-methyl-6-oxo-4-phenyl-1,4,5,6-tetrahydropyridine-3-carboxylate

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In the title compound,  $C_{18}H_{21}NO_4$ , the molecules form dimers by means of a pair of  $N-H \cdots O$  hydrogen bonds. The  $2(1H)$ -pyridone ring displays a screw-boat conformation.

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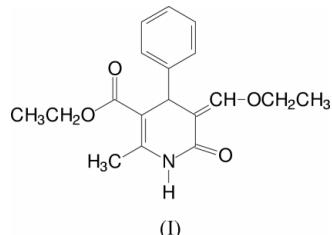
### Comment

A wide variety of compounds containing the  $2(1H)$ -pyridone ring are found in nature, and in some cases they display useful biological properties (Overman *et al.*, 1980).  $2(1H)$ -Pyridones are very close in structure to the 1,4-dihydropyridines, which have been used as effective drugs in the treatment of cardiovascular diseases. In a previous study, we have reported a general procedure (Verdecia *et al.*, 1996) for the synthesis of novel *o*-chloroformyl-substituted ethyl 1,4-dihydropyridine-5-carboxylates from the corresponding 3,4-dihydropyridones (Goldmann & Stoltfuss, 1991). The reaction proceeds through a mechanism involving several steps and one of the intermediates formed in the reaction is the title compound, (I).

### Key indicators

Single-crystal X-ray study  
 $T = 293\text{ K}$   
 $\text{Mean } \sigma(\text{C-C}) = 0.004\text{ \AA}$   
 $R \text{ factor} = 0.067$   
 $wR \text{ factor} = 0.219$   
Data-to-parameter ratio = 14.0

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.



The  $2(1H)$ -pyridone ring of (I) has a screw-boat conformation, with puckering parameters (Cremer & Pople, 1975)  $Q = 0.280(2)\text{ \AA}$ ,  $\theta = 109.5(4)^\circ$  and  $\varphi = 339.1(5)^\circ$ , with the axis through C2–C3, for N1–C1–C2–C3–C4–C5–C6. The mean  $Csp^2$ – $Csp^2$  bond length within this ring is  $1.434(1)\text{ \AA}$ . A centrosymmetric pair of  $N-H \cdots O$  hydrogen bonds holds the molecules together in a dimeric association (Table 2).

### Experimental

The title compound was prepared by stirring a solution of 5-ethoxycarbonyl-6-methyl-4-phenyl-3,4-dihydro-2( $1H$ )-pyridone (10 mmol) in anhydrous *N,N*-dimethylformamide (40 mmol), and phosphorus oxychloride (40 mmol) in dry chloroform (50 ml). After stirring at room temperature for 18 h, 20 ml of anhydrous ethanol was added. After 0.5 h, a white solid had precipitated and was filtered off. Further purification was accomplished by recrystallization from methanol (yield 68%; m.p. 746–747 K). IR ( $\text{KBr}, \text{cm}^{-1}$ ): 3202 (NH), 1697 (C=O), 1674 (C=O), 1637 (C=C);  $^1\text{H}$  NMR ( $\text{DMSO}-d_6$ , p.p.m.): 9.68 (*s*, 1H, NH), 7.36 (*s*, 1H, =CH), 7.31–7.11 (*m*, 5H, Ph), 4.89 (*s*, 1H, H-4), 4.13 (*q*, 2H, CH<sub>2</sub>), 3.96 (*q*, 2H, CH<sub>2</sub>), 2.29 (*s*, 3H, CH<sub>3</sub>), 1.22 (*t*, 3H, CH<sub>3</sub>), 1.06 (*t*, 3H, CH<sub>3</sub>);  $^{13}\text{C}$  NMR ( $\text{DMSO}-d_6$ ,

p.p.m.): 171.4 (C2), 169.7 (COO), 161.2 (=CH), 152.2 (C6), 149.5 (C1'), 133.4 (C2', C6'), 131.9 (C3', C5'), 131.4 (C4'), 114.5 (C3), 109.2 (C5), 75.3 (CH<sub>2</sub>), 64.5 (CH<sub>2</sub>), 43.4 (C4), 23.5 (CH<sub>3</sub>), 20.5 (CH<sub>3</sub>), 19.3 (CH<sub>3</sub>); MS: *m/z* (intensity%): 315 (*M*<sup>+</sup>, 64), 286 (60), 270 (31), 238 (65), 210 (100), 182 (44). Crystals suitable for X-ray analysis were obtained by slow evaporation from methanol.

#### Crystal data

C<sub>18</sub>H<sub>21</sub>NO<sub>4</sub>  
M<sub>r</sub> = 315.36  
Monoclinic, C2/c  
*a* = 22.652 (1) Å  
*b* = 10.5508 (3) Å  
*c* = 15.7162 (7) Å  
β = 114.921 (4)<sup>o</sup>  
V = 3406.4 (3) Å<sup>3</sup>  
Z = 8

D<sub>x</sub> = 1.230 Mg m<sup>-3</sup>  
Cu Kα radiation  
Cell parameters from 40 reflections  
θ = 10.8–27.9<sup>o</sup>  
μ = 0.71 mm<sup>-1</sup>  
T = 293 K  
Prism, colourless  
0.60 × 0.40 × 0.40 mm

#### Data collection

Siemens P4 four-circle diffractometer  
ω/2θ scans  
Absorption correction: ψ scan (North *et al.*, 1968)  
T<sub>min</sub> = 0.565, T<sub>max</sub> = 0.753  
7611 measured reflections  
2962 independent reflections  
2690 reflections with F<sup>2</sup> > 2σ(F<sup>2</sup>)

R<sub>int</sub> = 0.030  
θ<sub>max</sub> = 69.1<sup>o</sup>  
h = -1 → 27  
k = -1 → 12  
l = -19 → 17  
3 standard reflections every 100 reflections  
intensity decay: none

#### Refinement

Refinement on F<sup>2</sup>  
R[F<sup>2</sup> > 2σ(F<sup>2</sup>)] = 0.067  
wR(F<sup>2</sup>) = 0.219  
S = 0.98  
2962 reflections  
212 parameters  
H-atom parameters constrained

w = 1/[σ<sup>2</sup>(F<sub>o</sub><sup>2</sup>) + (0.1525P)<sup>2</sup> + 3.0912P]  
where P = (F<sub>o</sub><sup>2</sup> + 2F<sub>c</sub><sup>2</sup>)/3  
(Δ/σ)<sub>max</sub> = 0.110  
Δρ<sub>max</sub> = 0.62 e Å<sup>-3</sup>  
Δρ<sub>min</sub> = -0.39 e Å<sup>-3</sup>  
Extinction correction: SHELXL97  
Extinction coefficient: 0.0018 (3)

**Table 1**

Selected geometric parameters (Å, °).

O21—C2	1.234 (3)	O53—C51	1.333 (3)
O32—C31	1.342 (3)	O53—C54	1.453 (3)
O32—C33	1.445 (4)	N1—C2	1.368 (3)
O52—C51	1.204 (3)	N1—C6	1.394 (3)
C31—O32—C33	114.7 (2)	N1—C6—C61	111.3 (2)
C51—O53—C54	116.3 (2)	O32—C31—C3	121.9 (2)
C2—N1—C6	125.4 (2)	O32—C33—C34	109.8 (3)
O21—C2—N1	119.6 (2)	O52—C51—O53	122.0 (2)
O21—C2—C3	125.0 (2)	O52—C51—C5	122.3 (2)
N1—C2—C3	115.4 (2)	O53—C51—C5	115.8 (2)
N1—C6—C5	120.2 (2)	O53—C54—C55	107.7 (2)

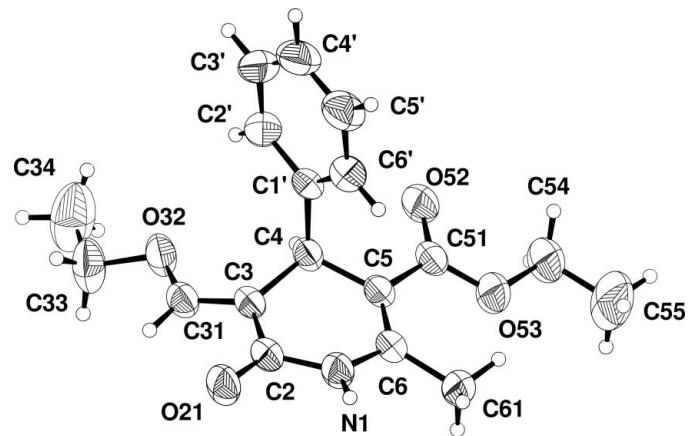
**Table 2**

Hydrogen-bonding geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1···O21 <sup>i</sup>	0.86	2.03	2.885 (3)	171
C31—H31···O21	0.93	2.46	2.785 (3)	100
C61—H61B···O53	0.96	2.20	2.803 (3)	119

Symmetry code: (i) -*x*, 1 - *y*, -*z*.

The value of the max shift/s.u. parameter corresponds to the torsion angle variation of atom H34 attached to C34 of a terminal



**Figure 1**

Plot showing the atomic numbering scheme in (I). Displacement ellipsoids for non-H atoms are shown at the 50% probability level.

methyl group. H atoms were positioned geometrically and included in the refinement, but were constrained to ride on their parent atoms. The isotropic displacement parameters of the H atoms were fixed at 1.3U<sub>eq</sub> of their parent atoms.

Data collection: *XSCANS* (Siemens, 1996); cell refinement: *XSCANS*; data reduction: *XSCANS*; program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *DIAMOND* (Bergerhoff, 1996); software used to prepare material for publication: *PLATON* (Spek, 1990), *PARST* (Nardelli, 1995) and *PARSTCIF* (Nardelli, 1991).

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