

2,4-Dimethyl-1,5-diphenylpenta-1,4-dien-3-one

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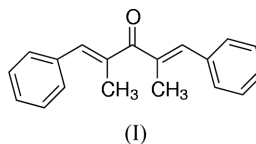
Key indicators

Single-crystal X-ray study
 $T = 296$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.034
 wR factor = 0.099
Data-to-parameter ratio = 7.0For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

In the crystal structure of the title compound, $\text{C}_{19}\text{H}_{18}\text{O}$, there is one-half molecule in the asymmetric unit and the carbonyl group is located on a crystallographic twofold rotation axis. The olefinic bond is in an *anti* configuration.

Comment

α,β -Unsaturated ketones have attracted considerable interest because they are versatile intermediates in organic synthesis, especially in the synthesis of natural and bioactive products (Fuchs & Paquette, 1994).



The molecule of the title compound, (I), lies on a crystallographic twofold rotation axis (Fig. 1). The olefinic bond is in an *anti* configuration. Some unexpected bond angles (Table 1) may be the result of steric repulsions in the molecule. The molecules are stretched into an undulating ribbon structure along the c axis and pack into infinite chains parallel to each other along the b axis (Fig. 2).

Experimental

Pentan-3-one (0.86 g, 10 mmol) and benzaldehyde (1.06 g, 10 mmol) were dissolved in ethanol (25 ml). To the mixture was added 10% aqueous NaOH (10 ml, 25 mmol). The mixture was stirred at room temperature for 2 h. The reaction mixture was extracted with CH_2Cl_2 , dried with anhydrous MgSO_4 and recrystallized from ethanol to produce the title compound, (I) (Concellon & Huerta, 2003).

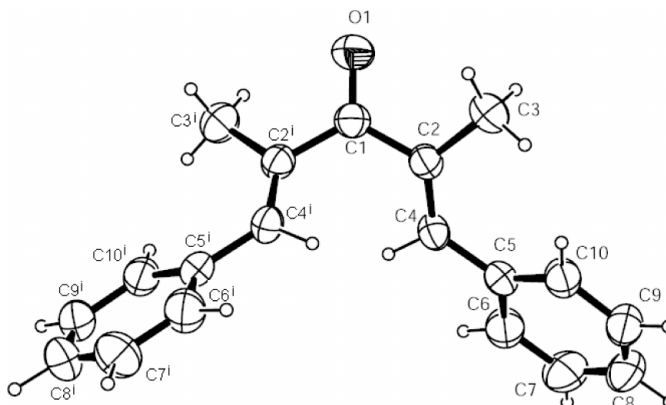


Figure 1
The molecular structure of (I), shown with 50% probability displacement ellipsoids. [Symmetry code: (i) $1 - x, y, -z$.]

Crystal data

C₁₉H₁₈O
M_r = 262.35
 Monoclinic, C2
a = 10.5362 (6) Å
b = 6.5254 (2) Å
c = 10.6656 (4) Å
 β = 97.790 (1)°
V = 726.52 (5) Å³
Z = 2

D_x = 1.199 Mg m⁻³
 Mo Kα radiation
 Cell parameters from 2926 reflections
 θ = 3.7–27.3°
 μ = 0.07 mm⁻¹
T = 296 (1) K
 Chunk, colorless
 0.33 × 0.30 × 0.17 mm

Data collection

Rigaku R-AXIS RAPID diffractometer
 ω scans
 Absorption correction: multi-scan (ABSCOR; Higashi, 1995)
T_{min} = 0.953, *T_{max}* = 0.988
 3380 measured reflections

904 independent reflections
 715 reflections with *F*² > 2σ(*F*²)
R_{int} = 0.027
 θ_{max} = 27.4°
h = -13 → 13
k = -8 → 8
l = -13 → 13

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.034
wR(*F*²) = 0.099
S = 1.01
 715 reflections
 93 parameters
 H-atom parameters constrained

w = 1/[0.0019*F_o*² + σ(*F_o*²)]/(4*F_o*²)
 (Δ/σ)_{max} < 0.001
 Δρ_{max} = 0.13 e Å⁻³
 Δρ_{min} = -0.10 e Å⁻³
 Extinction correction: (Larson, 1970)
 Extinction coefficient: 3.0 (5) × 10²

Table 1

Selected geometric parameters (°).

O1—C1—C2	119.1 (1)	C2—C4—C5	128.8 (2)
C1—C2—C3	114.4 (2)	C4—C5—C10	124.2 (2)
C1—C2—C4	119.7 (2)	C4—C5—C6	117.9 (2)
C4—C2—C3	125.5 (2)		
C3—C2—C4—C5	-0.8 (2)	C2—C4—C5—C10	-31.5 (2)

In the absence of significant anomalous dispersion effects, Friedel-pair reflections were merged before the final refinement. The H

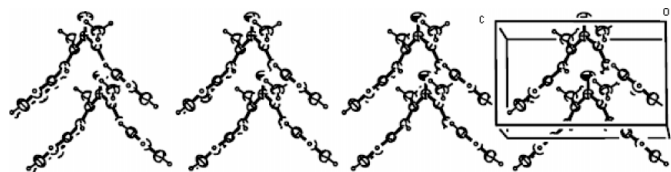


Figure 2
 The crystal structure of (I).

atoms were positioned geometrically and treated as riding, with C—H distances of 0.97 Å and *U_{iso}*(H) = 1.2*U_{eq}*(parent atom).

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/MSK, 2004); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *CRYSTALS* (Watkin *et al.*, 1996); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *CrystalStructure*.

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