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Key indicators

Single-crystal X-ray study T = 150 KMean $\sigma(C-C) = 0.004 \text{ Å}$ Disorder in main residue R factor = 0.064 wR factor = 0.179 Data-to-parameter ratio = 18.6

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. Ethyl 4-cyclohexyl-7-(dicyanomethylene)tricyclo[4.2.1.0^{2,5}]non-3-ene-3-carboxylate

The regio- and stereochemistry of the title compound, $C_{21}H_{24}N_2O_2$, has been established by X-ray analysis. There are two independent molecules in the asymmetric unit, one of which is disordered.

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Comment

Recently, we studied the remote substituent effects on ruthenium-catalysed [2 + 2]-cycloaddition reactions between 2-substituted 5-norbornenes and unsymmetrically substituted alkynes. Two different regio- and stereoisomers could be formed in the cycloadditions. When the substituent of the norbornene is a dicyanomethylene group, two regioisomers were obtained in a ratio of 15:1. These regioisomers were separated by fractional recystallization. The regio- and stereochemistry of the major isomer, (I), has been established by our single-crystal X-ray diffraction analysis.



In the structure of (I), there are two molecules in the asymmetric unit, namely A and B (Figs. 1 and 2). In molecule B, atom C15 of the ethoxy group is disordered over three sites. The occupancies of the three disorder sites, named C15B, C15* and C15\$, are 0.50, 0.25 and 0.25, respectively (see Fig. 2). The three sites are the result of rotational disorder about the O2B-C14B bond. Torsion angles for C13-O2-C14-C15, which describes the disorder and contrasts the geometry of the ethoxy group in molecules A and B, are given in Table 1. In addition, there is a slight difference in the orientation of the cyclohexyl group in each of the independent molecules, which is described by the torsion angle C7-C6-C16-C17 (see Table 1).

Experimental

Addition of 2-dicyanomethylene-5-norbornene and ethyl 3-cyclohexylpropiolate to Cp*RuCl(COD) in THF at 298 K provided two regioisomers in a ratio of 15:1. Fractional recystallization from an ethyl acetate/hexanes (1:4) mixture provided the major regioisomer, (I). Suitable crystals were grown from an ethyl acetate/hexanes (1:4) mixture.

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Figure 1

View of molecule A of (I), with the crystallographic labeling scheme. Displacement ellipsoids are drawn at the 30% probability level.



Figure 2

View of molecule B of (I), with the crystallographic labeling scheme. Displacement ellipsoids are drawn at the 30% probability level. The minor disorder components are shown with open and dashed bonds.

Crystal data

$C_{21}H_{24}N_2O_2$	Z = 4
$M_r = 336.42$	$D_x = 1.198 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 11.6420 (10) Å	Cell parameters from 16348
b = 12.2348(11) Å	reflections
c = 13.5739(13) Å	$\theta = 2.6-27.5^{\circ}$
$\alpha = 104.761 \ (5)^{\circ}$	$\mu = 0.08 \text{ mm}^{-1}$
$\beta = 90.427 \ (4)^{\circ}$	T = 150 (1) K
$\gamma = 93.727 \ (5)^{\circ}$	Plate, colourless
$V = 1865.1 (3) \text{ Å}^3$	$0.35 \times 0.30 \times 0.05 \text{ mm}$

Data collection

Nonius KappaCCD diffractometer	$R_{\rm int} = 0.100$
φ scans and ω scans with κ offsets	$\theta_{\rm max} = 27.6^{\circ}$
Absorption correction: none	$h = -15 \rightarrow 15$
20141 measured reflections	$k = -15 \rightarrow 15$
8463 independent reflections	$l = -17 \rightarrow 17$
3937 reflections with $I > 2\sigma(I)$	

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0702P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.064$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.180$	$(\Delta/\sigma)_{\rm max} = 0.001$
S = 0.98	$\Delta \rho_{\rm max} = 0.26 \ {\rm e} \ {\rm \AA}^{-3}$
8463 reflections	$\Delta \rho_{\rm min} = -0.30 \text{ e} \text{ \AA}^{-3}$
456 parameters	Extinction correction: SHELXTL
H-atom parameters constrained	Extinction coefficient: 0.0087 (18)

Table 1 Selected torsion angles (°).

C13A - O2A - C14A - C15A - 87.4(3)	C13B-O2B-C14B-C15\$	-114.0(8)
C13B-O2B-C14B-C15B 169.6 (4)	C7A-C6A-C16A-C17A	-85.1(3)
$C13B - O2B - C14B - C15^* - 159.9(6)$	C7B-C6B-C16B-C17B	-66.3 (4)

In the refinement, the bond lengths of the three disorder components (C14B-C15B, C14B-C15* and C14B-C15\$) were constrained to be equal, and the partial occupancy atoms were refined with isotropic displacement parameters. All H atoms were placed in calculated positions, with C-H distances ranging from 0.98 to 1.00 Å and included in the refinement in riding-motion approximation, with $U_{iso} = 1.2U_{eq}$ (1.5 U_{eq} for methyl) of the carrier atom.

Data collection: COLLECT (Nonius, 1997-2002); cell refinement: DENZO-SMN (Otwinowski & Minor, 1997); data reduction: DENZO-SMN; program(s) used to solve structure: SIR92 (Altomare et al., 1993); program(s) used to refine structure: SHELXTL (Sheldrick, 2001); molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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