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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=298 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.005 \AA$
$R$ factor $=0.038$
$w R$ factor $=0.104$
Data-to-parameter ratio $=7.0$

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

In the paper by Hansen, Perlovich \& Bauer-Brandl [Acta Cryst. (2003), E59, o1357-o1358], the coordinates of the $R$ enantiomer of the title compound, $\mathrm{C}_{13} \mathrm{H}_{18} \mathrm{O}_{2}$, were incorrectly given instead of those of the $S$ enantiomer. The correct coordinates of the $S$ enantiomer are given here.

## Comment

In the paper by Hansen et al. (2003), the coordinates of the $R$ enantiomer were incorrectly given instead of those of the $S$ enantiomer, (I). The correct coordinates of the $S$ enantiomer are given in the deposited replacement CIF. Molecular geometry parameters are not affected, except for the signs of torsion angles; the correct values are given in Table 1 below for the torsion angles in Table 2 of the previous report (where there was also an error in the atom numbering). Fig. 1 shows the correct structure of the two independent molecules, which form a hydrogen-bonded dimer without crystallographic symmetry.

(I)

## Experimental

Table 1
Selected torsion angles ( ${ }^{\circ}$ ).

| $\mathrm{C} 5 B-\mathrm{C} 4 B-\mathrm{C} 2 B-\mathrm{C} 3 B$ | $29.1(4)$ | $\mathrm{O} 1 A-\mathrm{C} 1 A-\mathrm{C} 2 A-\mathrm{C} 4 A$ | -81.7 (4) |
| :--- | ---: | :--- | ---: |
| $\mathrm{C} 7 B-\mathrm{C} 10 B-\mathrm{C} 11 B-\mathrm{C} 12 B$ | $-68.0(5)$ | $\mathrm{C} 3 A-\mathrm{C} 2 A-\mathrm{C} 4 A-\mathrm{C} 5 A$ | -144.4 (4) |
| $\mathrm{C} 4 B-\mathrm{C} 2 B-\mathrm{C} 1 B-\mathrm{O} 1 B$ | $83.5(3)$ | $\mathrm{C} 7 A-\mathrm{C} 10 A-\mathrm{C} 11 A-\mathrm{C} 13 A$ | 67.9 (5) |

All H atoms were refined freely $[\mathrm{C}-\mathrm{H}=0.85$ (3)-114 (5) $\AA$ ].
Data collection: CAD-4-PC Software (Enraf-Nonius, 1992); cell refinement: CELDIM in CAD-4-PC Software; data reduction: $X C A D$ (McArdle \& Higgins, 1995); program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEX (McArdle, 1995); software used to prepare material for publication: OSCAIL (McArdle, 1993).

## References

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Figure 1
The structure of the two independent molecules of (I). Displacement ellipsoids are shown at the $30 \%$ probability level. Hydrogen bonds are shown as dashed lines.

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