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**(±)-1-[[4-Methoxybenzylidene)-amino](4-methoxyphenyl)methyl]-2-naphthol**

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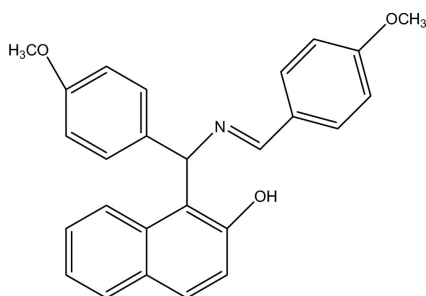
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Key indicators: single-crystal X-ray study;  $T = 298$  K; mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å;  $R$  factor = 0.071;  $wR$  factor = 0.156; data-to-parameter ratio = 13.1.

The title compound,  $\text{C}_{26}\text{H}_{23}\text{NO}_3$ , is a key intermediate used in the synthesis of 1-[amino(4-methoxyphenyl)methyl]-2-naphthol, which has been found to be a potent chiral catalyst in the addition reactions of dialkylzinc compounds with aldehydes. The  $\pi$ -conjugated system formed by the methoxybenzylidene-imine tautomer is essentially planar. An intramolecular  $\text{O}-\text{H}\cdots\text{N}$  hydrogen bond is formed between the hydroxyl OH group attached to the naphthalene group and the Schiff base N atom. The crystal packing is stabilized by van der Waals forces.

## Related literature

For related literature, see: Sztamari *et al.* (2003).

## Experimental

## Crystal data

 $\text{C}_{26}\text{H}_{23}\text{NO}_3$   
 $M_r = 397.45$ Monoclinic,  $P2_1/c$   
 $a = 20.790$  (5) Å $b = 10.717$  (3) Å  
 $c = 9.470$  (2) Å  
 $\beta = 102.952$  (4)°  
 $V = 2056.4$  (9) Å<sup>3</sup>  
 $Z = 4$ Mo  $K\alpha$  radiation  
 $\mu = 0.08$  mm<sup>-1</sup>  
 $T = 298$  (2) K  
 $0.46 \times 0.45 \times 0.16$  mm

## Data collection

Bruker APEX area-detector diffractometer  
Absorption correction: multi-scan (SADABS; Bruker, 2001)  
 $T_{\min} = 0.963$ ,  $T_{\max} = 0.987$ 10108 measured reflections  
3595 independent reflections  
3150 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.022$ 

## Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.071$   
 $wR(F^2) = 0.156$   
 $S = 1.22$   
3594 reflections274 parameters  
H-atom parameters constrained  
 $\Delta\rho_{\max} = 0.19$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.22$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O3}-\text{H3}\cdots\text{N1}$	0.82	1.82	2.544 (3)	146

Data collection: *SMART* (Bruker, 2001); cell refinement: *S SAINT* (Bruker, 2001); data reduction: *S SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP III* (Burnett & Johnson, 1996) and *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: DN2180).

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