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## Structure Reports

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## N,N-Dimethyl-5-[2-(hydroxymethyl)-4-nitro-phenyl]pent-4-ynamide

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## Key indicators

Single-crystal X-ray study
$T=161 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.054$
$w R$ factor $=0.149$
Data-to-parameter ratio $=19.6$

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

[^0]Molecules of title title compound, $\mathrm{C}_{14} \mathrm{H}_{16} \mathrm{~N}_{2} \mathrm{O}_{4}$, crystallize as centrosymmetric dimers, connected by intermolecular $\mathrm{O}-$ $\mathrm{H} \cdots \mathrm{O}$ hydrogen bonds.

## Comment

The title compound, (I), was prepared for an investigation of the addition of nucleophiles to $\mathrm{C} \equiv \mathrm{C}$ triple bonds and the $\mathrm{C}-$ H activation in the benzylic position by homogeneous gold catalysts (Hashmi, 2004).

(I)

The angle between the planes of the benzene ring and the nitro group is $3.9(1)^{\circ}$. The $\mathrm{C} 7-\mathrm{O} 3$ bond is almost coplanar with the benzene ring. This conformation results in a contact distance of $2.33 \AA$ between atoms O3 and H6. The amide group is approximately planar. The amide N atom shows no deviation from planarity; the sum of the three valence angles about atom N 2 is $360.0^{\circ}$. The molecules form centrosymmetric dimers connected by intermolecular $\mathrm{O}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonds. The structure of the dimer is shown in Fig. 1 and details


Figure 1
The structure of a centrosymmetric dimer of compound (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the $50 \%$ probability level. H atoms are drawn as small spheres of arbitrary radius and hydrogen bonds are shown as dotted lines.

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of the hydrogen bonding are given in Table 2. The hydrogenbond formation results in an approximate coplanarity of the benzene group and the amide group; the angle between the benzene ring and the plane of the $\mathrm{N}, \mathrm{N}$-dimethylamide group is $8.0(1)^{\circ}$. A very different conformation has been observed for the corresponding molecule without a nitro group attached to the benzene ring (Bats et al., 2006). Those molecules are arranged in hydrogen-bonded chains, and the angles between the benzene plane and the plane of the $\mathrm{N}, \mathrm{N}$-dimethylamide group range from 74.6 (1) to 82.2 (1) ${ }^{\circ}$.

The crystal packing is shown in Fig. 2. Neighboring dimers are connected by three different, very weak intermolecular $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ interactions, with $\mathrm{H} \cdots \mathrm{O}$ distances of $2.59,2.66$ and $2.66 \AA$ and $\mathrm{O}-\mathrm{H} \cdots \mathrm{O}$ angles of 144,160 and $109^{\circ}$. There is also a weak intermolecular C (methyl) $-\mathrm{H} \cdots \pi$ (alkyne) contact, with an $\mathrm{H} \cdots C g$ distance of $2.83 \AA$ and a $\mathrm{C}-\mathrm{H} \cdots C g^{\mathrm{ii}}$ angle of $142^{\circ}$ [ $C g$ is the mid-point of the $\mathrm{C} \equiv \mathrm{C}$ triple bond; symmetry code: (ii) $\left.x,-y+\frac{1}{2}, z+\frac{1}{2}\right]$.

## Experimental

The title compound was prepared using the Sonogashira coupling of 2-bromo-(5-nitrophenyl)methanol and $\mathrm{N}, \mathrm{N}$-dimethylpent-4-ynamide (Thorand \& Krause, 1998). Single crystals were obtained by evaporation of a solution of (I) in methanol/ethyl acetate ( $1 / 1 \mathrm{v} / \mathrm{v}$ ).

## Crystal data

$\mathrm{C}_{14} \mathrm{H}_{16} \mathrm{~N}_{2} \mathrm{O}_{4}$
$M_{r}=276.29$
Monoclinic, $P 2_{6} / c$
$a=12.877(3) \AA \AA$
$b=14.3165(18) \AA$
$c=7.5960(15) \AA$
$\beta=102.934(8){ }^{\circ}$
$V=1364.8(5) \AA^{3}$
$Z=4$

## Data collection

Siemens SMART 1K CCD
diffractometer
$\omega$ scans
Absorption correction: none
21386 measured reflections
3663 independent reflections

## Refinement

## Refinement on $F^{2}$

$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.054$
$w R\left(F^{2}\right)=0.149$
$S=1.03$
3663 reflections
187 parameters
H atoms treated by a mixture of independent and constrained refinement
$D_{x}=1.345 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
Cell parameters from 119 reflections
$\theta=3-23^{\circ}$
$\mu=0.10 \mathrm{~mm}^{-1}$
$T=161$ (2) K
Block, yellow
$0.26 \times 0.24 \times 0.18 \mathrm{~mm}$

2205 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.055$
$\theta_{\text {max }}=29.5^{\circ}$
$h=-17 \rightarrow 17$
$k=-19 \rightarrow 19$
$l=-10 \rightarrow 10$

$$
\begin{aligned}
& w= 1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.06 P)^{2}\right. \\
&+0.6 P] \\
& \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
&(\Delta / \sigma)_{\max }=0.003 \\
& \Delta \rho_{\max }=0.29 \mathrm{e}^{-3} \\
& \Delta \rho_{\min }=-0.25 \mathrm{e}^{-3}
\end{aligned}
$$

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

| $\mathrm{C} 6-\mathrm{C} 5-\mathrm{C} 7-\mathrm{O} 3$ | $6.3(3)$ | $\mathrm{C} 13-\mathrm{N} 2-\mathrm{C} 12-\mathrm{O} 4$ | $-176.00(18)$ |
| :--- | :--- | :--- | :--- |



Figure 2
The crystal packing of (I), viewed down c. Displacement ellipsoids are drawn at the $50 \%$ probability level and hydrogen bonds are shown as broken lines.

Table 2
Hydrogen-bond geometry ( $\AA{ }^{\circ},{ }^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| O3-H3B $\cdots \mathrm{O}^{\mathrm{i}}$ | $0.88(3)$ | $1.85(3)$ | $2.712(2)$ | $164(2)$ |

Symmetry code: (i) $-x+2,-y+1,-z$.
H atoms attached to C atoms were positioned geometrically and refined as riding atoms $\left[\mathrm{Csp} p^{2}-\mathrm{H}=0.95 \AA\right.$ and secondary $\mathrm{C}-\mathrm{H}=$ $0.99 \AA$, with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$, and methyl $\mathrm{C}-\mathrm{H}=0.98 \AA$ with $\left.U_{\text {iso }}(\mathrm{H})=1.5 U_{\text {eq }}(\mathrm{C})\right]$. The hydroxyl H atom was located in a difference Fourier map and was refined isotropically.

Data collection: SMART (Siemens, 1995); cell refinement: SMART; data reduction: SAINT (Siemens, 1995); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Sheldrick, 1996); software used to prepare material for publication: SHELXL97.

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