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N-(1,5-Dimethyl-3-oxo-2-phenyl-2,3-dihydro-1H-pyrazol-4-yl)acetamide-naphthalene-2,3-diol (1/1)

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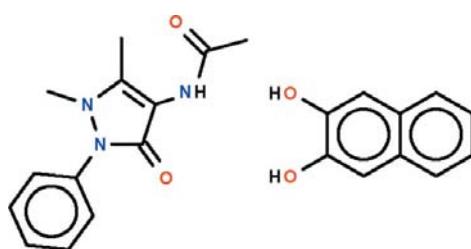
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Key indicators: single-crystal X-ray study; $T = 100\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.052; wR factor = 0.131; data-to-parameter ratio = 16.4.

In the reaction of naphthalene-2,3-diol and 4-aminoantipyrine in the presence of acetic acid, the amine function is acetylated and the resulting acetamide co-crystallizes with the diol in the title compound, $C_{13}H_{15}N_3O_2 \cdot C_{10}H_8O_2$, with 1:1 molar stoichiometry. The two components are linked by two $O\cdots H\cdots O=C$ hydrogen bonds. One of the hydroxy groups interacts with the pyrazolone carbonyl O atom and the other hydroxy group interacts with the amide O atom of another component, generating a chain motif. Adjacent chains are linked into a layer motif via $N\cdots H\cdots O$ interactions involving only the heterocyclic acetamide component.

Related literature

For the crystal structure of 4-acetamido-2,3-dimethyl-1-phenyl-5-pyrazol-3-one, see: Kuznetsov *et al.* (1999). For co-crystals of naphthalene-2,3-diol, see: Fritchie & Johnston (1975); Herbert & Truter (1980); Kuo *et al.* (1974); Nakamatsu *et al.* (2003); Wang *et al.* (2008); Wells *et al.* (1974).



Experimental

Crystal data

$C_{13}H_{15}N_3O_2 \cdot C_{10}H_8O_2$

$M_r = 405.44$

Monoclinic, $P2_1/c$
 $a = 12.426 (1)\text{ \AA}$
 $b = 14.304 (2)\text{ \AA}$
 $c = 12.959 (1)\text{ \AA}$
 $\beta = 117.845 (1)^\circ$
 $V = 2036.7 (4)\text{ \AA}^3$

$Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.09\text{ mm}^{-1}$
 $T = 100\text{ K}$
 $0.25 \times 0.25 \times 0.10\text{ mm}$

Data collection

Bruker SMART APEX
diffractometer
19263 measured reflections

4683 independent reflections
3189 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.061$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.052$
 $wR(F^2) = 0.131$
 $S = 1.02$
4683 reflections
286 parameters
27 restraints

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.31\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.36\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N1-H1\cdots O2^i$	0.87 (1)	2.07 (1)	2.924 (2)	169 (2)
$O3-H3\cdots O2$	0.85 (3)	1.81 (3)	2.639 (2)	163 (3)
$O4-H4\cdots O1^{ii}$	0.85 (3)	1.81 (3)	2.646 (2)	168 (3)

Symmetry codes: (i) $-x + 1, -y + 1, -z + 1$; (ii) $x, -y + \frac{3}{2}, z + \frac{1}{2}$.

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *publCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: IM2213).

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