

Acta Crystallographica Section E

## Structure Reports

Online

ISSN 1600-5368

## 4-(3,5-Dimethyl-1H-pyrazol-1-yl)-benzenesulfonamide

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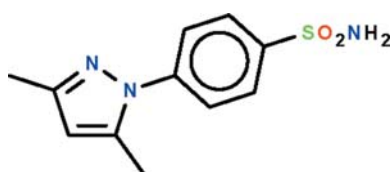
Received 11 August 2011; accepted 13 August 2011

Key indicators: single-crystal X-ray study;  $T = 100$  K; mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å;  
 $R$  factor = 0.031;  $wR$  factor = 0.083; data-to-parameter ratio = 14.1.

The two aromatic rings of the title compound,  $\text{C}_{11}\text{H}_{13}\text{N}_3\text{O}_2\text{S}$ , are inclined at an angle of  $47.81$  (4)°. The N atom of the amino unit is pyramidally coordinated; one H atom interacts with the sulfamyl O atom of an adjacent molecule, forming a centrosymmetric hydrogen-bonded dimer. The dimers are linked by  $\text{N}-\text{H}\cdots\text{N}$  hydrogen bonds, generating a three-dimensional network.

## Related literature

For the synthesis and medicinal properties of the title compound, see: Grueneberg *et al.* (2002); Wright *et al.* (1964).



## Experimental

## Crystal data

 $\text{C}_{11}\text{H}_{13}\text{N}_3\text{O}_2\text{S}$  $M_r = 251.30$ Monoclinic,  $P2_1/n$  $a = 7.9649$  (1) Å $b = 11.7827$  (2) Å $c = 12.2720$  (2) Å $\beta = 91.720$  (1)°  
 $V = 1151.18$  (3) Å<sup>3</sup>  
 $Z = 4$   
Cu  $K\alpha$  radiation $\mu = 2.47$  mm<sup>-1</sup>  
 $T = 100$  K  
 $0.30 \times 0.20 \times 0.02$  mm

## Data collection

Agilent SuperNova Dual  
diffractometer with an Atlas  
detector  
Absorption correction: multi-scan  
(*CrysAlis PRO*; Agilent, 2010)  
 $T_{\min} = 0.525$ ,  $T_{\max} = 0.952$ 8510 measured reflections  
2312 independent reflections  
2215 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.018$ 

## Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.031$   
 $wR(F^2) = 0.083$   
 $S = 1.07$   
2312 reflections  
164 parameters  
2 restraintsH atoms treated by a mixture of  
independent and constrained  
refinement  
 $\Delta\rho_{\max} = 0.35$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.41$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N3}-\text{H1}\cdots\text{O1}^i$	0.87 (1)	2.13 (1)	2.966 (2)	160 (2)
$\text{N3}-\text{H1}\cdots\text{N2}^{ii}$	0.87 (1)	2.94 (2)	3.501 (2)	124 (2)

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

Data collection: *CrysAlis PRO* (Agilent, 2010); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; 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).

We thank King Abdulaziz University and the University of Malaya for supporting this study.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT5610).

## References

- Agilent (2010). *CrysAlis PRO*. Agilent Technologies, Yarnton, Oxfordshire, England.
- Barbour, L. J. (2001). *J. Supramol. Chem.* **1**, 189–191.
- Grueneberg, S., Stubbs, M. T. & Klebe, G. (2002). *J. Med. Chem.* **45**, 3588–3602.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Westrip, S. P. (2010). *J. Appl. Cryst.* **43**, 920–925.
- Wright, J. B., Dulin, W. E. & Markillie, J. H. (1964). *J. Med. Chem.* **7**, 102–105.