organic compounds

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

N-(3,4-Difluorophenyl)-*N*'-(2,5-dimethoxyphenyl)urea

Hyeong Choi,^a Byung Hee Han,^a Taewoo Lee,^a Sung Kwon Kang^a* and Chang Keun Sung^b

^aDepartment of Chemistry, Chungnam National University, Daejeon 305-764, Republic of Korea, and ^bDepartment of Food Science and Technology, Chungnam National University, Daejeon 305-764, Republic of Korea Correspondence e-mail: skkang@cnu.ac.kr

Received 8 July 2010; accepted 10 August 2010

Key indicators: single-crystal X-ray study; T = 174 K; mean σ (C–C) = 0.006 Å; disorder in main residue; R factor = 0.061; wR factor = 0.145; data-to-parameter ratio = 11.2.

In the title compound, $C_{15}H_{14}F_2N_2O_3$, the dihedral angle between the benzene rings is 64.5 (1)°. One F atom is disordered over two *meta* positions, with occupancy factors of 0.72 and 0.28. In the crystal, molecules are linked by N– H···O hydrogen bonds involving two N–H and one C=O groups of the urea central fragment, leading to a supramolecular chain along [011].

Related literature

For general background to the development of potent inhibitory agents of tyrosinase and melanin formation used as whitening agents, see: Cabanes *et al.* (1994); Choi *et al.* (2010); Criton & Le Mellay-Hamon (2008); Germanas *et al.* (2007); Dawley & Flurkey (1993); Ha *et al.* (2007); Hong *et al.* (2008); Kwak *et al.* (2010); Lee *et al.* (2007); Nerya *et al.* (2003); Yi *et al.* (2009, 2010).



Experimental

Crystal data

 $\begin{array}{l} C_{15}H_{14}F_2N_2O_3\\ M_r = 308.28\\ \text{Monoclinic, } P2_1/c\\ a = 13.209 \ (2) \ \text{\AA}\\ b = 12.0887 \ (18) \ \text{\AA} \end{array}$

c = 9.0740 (12) Å $\beta = 104.990 (4)^{\circ}$ $V = 1399.6 (4) \text{ Å}^{3}$ Z = 4Mo $K\alpha$ radiation



 $\mu = 0.12 \text{ mm}^{-1}$ T = 174 K

Data collection

Bruker SMART CCD area-detector diffractometer 10104 measured reflections	2433 independent reflections 1211 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.120$
Refinement	
$R[F^2 > 2\sigma(F^2)] = 0.061$	H atoms treated by a mixture of
$wR(F^2) = 0.145$	independent and constrained
S = 0.96	refinement
2433 reflections	$\Delta \rho_{\rm max} = 0.25 \text{ e} \text{ Å}^{-3}$
218 parameters	$\Delta \rho_{\rm min} = -0.26 \text{ e} \text{ Å}^{-3}$

 $0.09 \times 0.04 \times 0.02 \ \mathrm{mm}$

Table 1 Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N9-H9\cdotsO11^{i}$ $N12-H12\cdotsO11^{i}$	0.84(4)	2.09 (4)	2.907 (4)	163 (4)
	0.80(4)	2.30 (4)	3.002 (4)	147 (4)

Symmetry code: (i) $x, -y + \frac{1}{2}, z + \frac{1}{2}$.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

We wish to thank the DBIO Company for partial support of this work.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BH2300).

References

- Bruker (2002). SADABS, SAINT and SMART. Bruker AXS Inc., Madison, Wisconsin, USA.
- Cabanes, J., Chazarra, S. & Garcia-Carmona, F. (1994). J. Pharm. Pharmacol. 46, 982–985.
- Choi, H., Han, B. H., Lee, T., Kang, S. K. & Sung, C. K. (2010). Acta Cryst. E66, 01142.
- Criton, M. & Le Mellay-Hamon, V. (2008). Bioorg. Med. Chem. Lett. 18, 3607–3610.

Dawley, R. M. & Flurkey, W. H. (1993). J. Food Sci. 58, 609-610.

- Farrugia, L. J. (1997). J. Appl. Cryst. 30, 565.
- Farrugia, L. J. (1999). J. Appl. Cryst. 32, 837-838.
- Germanas, J. P., Wang, S., Miner, A., Hao, W. & Ready, J. M. (2007). Bioorg. Med. Chem. Lett. 17, 6871–6875.
- Ha, Y. M., Chung, S. W., Song, S., Lee, H., Suh, H. & Chung, H. Y. (2007). Biol. Pharm. Bull. 30, 1711–1715.
- Hong, W. K., Heo, J. Y., Han, B. H., Sung, C. K. & Kang, S. K. (2008). Acta Cryst. E64, 049.
- Kwak, S.-Y., Noh, J.-M., Park, S.-H., Byun, J.-W., Choi, H.-R., Park, K.-C. & Lee, Y.-S. (2010). Bioorg. Med. Chem. Lett. 20, 738–741.
- Lee, C. W., Son, E.-M., Kim, H. S., Xu, P., Batmunkh, T., Lee, B.-J. & Koo, K. A. (2007). *Bioorg. Med. Chem. Lett.* 17, 5462–5464.
- Nerya, O., Vaya, J., Musa, R., Izrael, S., Ben-Arie, R. & Tamir, S. (2003). J. Agric. Food Chem. 51, 1201–1207.
- Sheldrick, G. M. (2008). Acta Cryst. A64, 112-122.
- Yi, W., Cao, R.-H., Chen, Z.-Y., Yu, L., Ma, L. & Song, H.-C. (2009). Chem. Pharm. Bull. 57, 1273–1277.
- Yi, W., Cao, R., Peng, W., Wen, H., Yan, Q., Zhou, B., Ma, L. & Song, H. (2010). *Eur. J. Med. Chem.* 45, 639–646.

supporting information

Acta Cryst. (2010). E66, o2320 [https://doi.org/10.1107/S1600536810032095] N-(3,4-Difluorophenyl)-N'-(2,5-dimethoxyphenyl)urea

Hyeong Choi, Byung Hee Han, Taewoo Lee, Sung Kwon Kang and Chang Keun Sung

S1. Comment

Melanin is the pigment responsible for the color of human skin and it is formed through a series of oxidative reaction in the presence of key enzyme tyrosinase (Ha et al., 2007) that converts tyrosine into melanin. It is secreted by melanocyte cells distributed in the basal layer of the dermis. Its role is to protect the skin from ultraviolet (UV) damage by absorbing the ultraviolet sunlight and removing reactive oxygen species. Therefore these inhibitors are target molecules for developing anti-pigmentation agents. Common tyrosinase inhibitors (Dawley & Flurkey, 1993; Nerya et al., 2003) are hydroquinone, ascorbic acid, kojic acid and arbutin (Cabanes et al., 1994). Recently, numerous reports have focused on the inhibition of tyrosinase. They are containing aromatic, methoxy, hydroxyl (Hong et al., 2008; Lee et al., 2007), aldehyde (Yi et al., 2010), amide (Kwak et al., 2010; Choi et al., 2010), thiosemicarbazone (Yi et al., 2009), thiazole (Germanas et al., 2007), thiourea (Criton & Le Mellay-Hamon, 2008) groups in their structures, and act as a specific functional group to make the skin white by inhibiting the production of melanin. However, most of them are not potent enough to put into practical use due to their weak individual activities, poor skin penetration, and low stability of formulations, as well as toxicity or safety concerns. Consequently, there is still need to search and develop novel tyrosinase inhibitors with better activities together with lower side effects. To complement the inadequacy of current whitening agent above mentioned and maximize the effect of inhabitation of melanin creation, we have synthesized the title compound, (I), from the reaction of 3.4-difluoroaniline with 2,5-dimethoxyphenyl isocvanate, under ambient conditions. Herein, the crystal sturucture of (I) is described (Fig. 1).

The 3,4-difluoroaniline group and 2,5-dimethoxyphenyl moiety are essentially planar, with a mean deviations of 0.007 Å and 0.016 Å, respectively, from the corresponding least-squares planes defined by each nine constituent atoms. The dihedral angle between the benzene rings is 64.5 (1) $^{\circ}$. The presence of intermolecular N—H…O hydrogen bonds lead to the formation a supramolecular chain along [011].

S2. Experimental

2,5-Dimethoxyphenyl isocyanate and 3,4-difluoroaniline were purchased from Sigma Chemical Co. Solvents used for synthesis were redistilled before use. All other chemicals and solvents were of analytical grade and used without further purification. The title compound was prepared from the reaction of 3,4-difluoroaniline (0.28 ml, 3 mmol) and 2,5-dimethoxyphenyl isocyanate (0.5 g, 3 mmol) in acetonitrile (6 ml). The mixture was refluxed for 8 h at 353 K, and then treated with water and extracted with methylene chloride (2×50 mL). The combined extracts were dried over anhydrous magnesium sulfate. Removal of solvent gave a white solid (90%, m.p. 454 K). Single crystals were obtained by slow evaporation of a methylene chloride solution at room temperature.

S3. Refinement

The amide H atoms were located in a difference map and refined freely. The remaining H atoms were positioned geometrically and refined using a riding model with C—H = 0.93–0.96 Å, and with $U_{iso}(H) = 1.2U_{eq}(C)$ for aromatic and $1.5U_{eq}(C)$ for methyl H atoms. Atom F7 is disordered over two positions and the two split atoms are designated by having the suffix A after the atom number. The final occupancy factors are F7 0.72 and F7A 0.28. The measured diffraction fraction is relatively low of 95.5% due to a tiny single-crystal for data collection. This single-crystal was the largest one we could produce as described in the experimental section.



Figure 1

Molecular structure of (l), showing the atom-numbering scheme and 50% probability ellipsoids.

N-(3,4-Difluorophenyl)-N'-(2,5-dimethoxyphenyl)urea

Crystal data

 $C_{15}H_{14}F_2N_2O_3$ $M_r = 308.28$ Monoclinic, $P2_1/c$ Hall symbol: -P 2ybc *a* = 13.209 (2) Å b = 12.0887 (18) Åc = 9.0740 (12) Å $\beta = 104.990 \ (4)^{\circ}$ V = 1399.6 (4) Å³ Z = 4

Data collection

 $wR(F^2) = 0.145$

Bruker SMART CCD area-detector diffractometer	1211 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.120$
Radiation source: fine-focus sealed tube	$\theta_{\rm max} = 25.3^{\circ}, \theta_{\rm min} = 1.6^{\circ}$
φ and ω scans	$h = -15 \rightarrow 14$
10104 measured reflections	$k = -12 \rightarrow 14$
2433 independent reflections	$l = -4 \rightarrow 10$
Refinement	
Refinement on F^2	S = 0.96
Least-squares matrix: full	2433 reflections
$R[F^2 > 2\sigma(F^2)] = 0.061$	218 parameters

F(000) = 640 $D_{\rm x} = 1.463 {\rm Mg} {\rm m}^{-3}$ Melting point: 454 K Mo *K* α radiation, $\lambda = 0.71073$ Å Cell parameters from 398 reflections $\theta = 3.0 - 18.6^{\circ}$ $\mu = 0.12 \text{ mm}^{-1}$ T = 174 KNeedle, colourless $0.09 \times 0.04 \times 0.02 \text{ mm}$

218 parameters 0 restraints

0 constraints

H atoms treated by a mixture of independent

and constrained refinement

$$\begin{split} &w = 1/[\sigma^2(F_o{}^2) + (0.0573P)^2] \\ & \text{where } P = (F_o{}^2 + 2F_c{}^2)/3 \\ & (\Delta/\sigma)_{\text{max}} < 0.001 \\ & \Delta\rho_{\text{max}} = 0.25 \text{ e } \text{ Å}{}^{-3} \\ & \Delta\rho_{\text{min}} = -0.26 \text{ e } \text{ Å}{}^{-3} \end{split}$$

	x	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	Occ. (<1)
C1	0.6480 (3)	0.4327 (3)	0.4129 (4)	0.0243 (9)	
C2	0.5993 (3)	0.4687 (3)	0.2657 (4)	0.0278 (10)	
H2	0.5991	0.4245	0.1818	0.033*	
C3	0.5517 (3)	0.5705 (4)	0.2464 (4)	0.0351 (11)	
Н3	0.5189	0.5949	0.1484	0.042*	0.28
C4	0.5517 (3)	0.6365 (3)	0.3683 (5)	0.0354 (11)	
C5	0.5984 (3)	0.6018 (4)	0.5137 (4)	0.0336 (11)	
Н5	0.5973	0.6468	0.5964	0.04*	0.72
C6	0.6469 (3)	0.5008 (3)	0.5371 (4)	0.0278 (10)	
H6	0.6791	0.4775	0.6358	0.033*	
F7	0.5055 (3)	0.6079 (3)	0.1088 (3)	0.0597 (11)	0.72
F7A	0.6007 (6)	0.6720 (7)	0.6227 (8)	0.041 (2)	0.28
F8	0.5028 (2)	0.7365 (2)	0.3447 (3)	0.0586 (8)	
N9	0.6969 (2)	0.3289 (3)	0.4443 (3)	0.0274 (9)	
Н9	0.706 (3)	0.302 (3)	0.533 (4)	0.038 (12)*	
C10	0.7429 (3)	0.2699 (3)	0.3512 (4)	0.0271 (10)	
O11	0.74653 (19)	0.3012 (2)	0.2231 (2)	0.0301 (7)	
N12	0.7839 (3)	0.1710 (3)	0.4127 (4)	0.0315 (9)	
H12	0.775 (3)	0.151 (3)	0.492 (4)	0.036 (13)*	
C13	0.8518 (3)	0.1019 (3)	0.3556 (4)	0.0259 (10)	
C14	0.8499 (3)	-0.0108 (4)	0.3854 (4)	0.0295 (10)	
C15	0.9184 (3)	-0.0817 (4)	0.3393 (4)	0.0347 (11)	
H15	0.9177	-0.1569	0.3604	0.042*	
C16	0.9883 (3)	-0.0401 (4)	0.2614 (4)	0.0346 (11)	
H16	1.0341	-0.0875	0.2299	0.042*	
C17	0.9893 (3)	0.0715 (4)	0.2313 (4)	0.0317 (10)	
C18	0.9216 (3)	0.1432 (3)	0.2788 (4)	0.0295 (10)	
H18	0.9233	0.2186	0.259	0.035*	
O19	0.7767 (2)	-0.0423 (2)	0.4625 (3)	0.0381 (7)	
C20	0.7657 (4)	-0.1587 (4)	0.4843 (5)	0.0496 (13)	
H20A	0.7476	-0.1951	0.387	0.074*	
H20B	0.7114	-0.1709	0.5353	0.074*	
H20C	0.8307	-0.1879	0.5453	0.074*	
O21	1.0554 (2)	0.1220 (2)	0.1555 (3)	0.0414 (8)	
C22	1.1393 (3)	0.0546 (4)	0.1310 (4)	0.0423 (12)	
H22A	1.18	0.0265	0.2271	0.063*	
H22B	1.1833	0.0984	0.0846	0.063*	
H22C	1.1107	-0.006	0.0651	0.063*	

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters $(Å^2)$

supporting information

	$\overline{U^{11}}$	U^{22}	U^{33}	$\overline{U^{12}}$	U^{13}	U^{23}
C1	0.029 (2)	0.018 (2)	0.0284 (19)	-0.0011 (19)	0.0118 (16)	0.0004 (16)
C2	0.030 (2)	0.024 (3)	0.0296 (19)	-0.001 (2)	0.0088 (17)	-0.0009 (17)
C3	0.034 (3)	0.034 (3)	0.036 (2)	0.004 (2)	0.0063 (19)	0.007 (2)
C4	0.034 (3)	0.016 (3)	0.055 (3)	0.010 (2)	0.010(2)	0.004 (2)
C5	0.030 (3)	0.028 (3)	0.045 (2)	0.000 (2)	0.0121 (19)	-0.006 (2)
C6	0.030(2)	0.027 (3)	0.0278 (19)	-0.003 (2)	0.0082 (16)	-0.0031 (17)
F7	0.077 (3)	0.052 (3)	0.0428 (18)	0.022 (2)	0.0015 (17)	0.0155 (18)
F7A	0.050 (6)	0.031 (5)	0.050 (4)	0.006 (5)	0.025 (4)	-0.016 (4)
F8	0.066 (2)	0.0366 (18)	0.0715 (16)	0.0211 (15)	0.0138 (14)	0.0024 (13)
N9	0.039 (2)	0.025 (2)	0.0191 (16)	0.0072 (18)	0.0103 (14)	0.0042 (15)
C10	0.025 (3)	0.031 (3)	0.0251 (19)	-0.001 (2)	0.0065 (16)	-0.0030 (18)
O11	0.0381 (17)	0.0301 (18)	0.0248 (12)	0.0040 (14)	0.0133 (11)	0.0004 (11)
N12	0.043 (2)	0.025 (2)	0.0300 (18)	0.0083 (18)	0.0158 (16)	0.0063 (16)
C13	0.029 (3)	0.021 (3)	0.0270 (19)	0.004 (2)	0.0059 (17)	0.0016 (17)
C14	0.035 (3)	0.026 (3)	0.0265 (19)	-0.003 (2)	0.0066 (18)	0.0019 (18)
C15	0.046 (3)	0.020 (3)	0.037 (2)	0.003 (2)	0.008 (2)	-0.0021 (18)
C16	0.042 (3)	0.027 (3)	0.036 (2)	0.011 (2)	0.0121 (19)	-0.0048 (19)
C17	0.033 (3)	0.024 (3)	0.039 (2)	0.007 (2)	0.0094 (19)	0.0010 (19)
C18	0.037 (3)	0.018 (3)	0.034 (2)	0.001 (2)	0.0105 (18)	0.0033 (17)
O19	0.0450 (19)	0.0241 (19)	0.0492 (15)	-0.0003 (15)	0.0193 (14)	0.0043 (13)
C20	0.066 (3)	0.028 (3)	0.057 (3)	-0.012 (3)	0.019 (2)	0.003 (2)
O21	0.0436 (19)	0.035 (2)	0.0553 (16)	0.0122 (15)	0.0299 (14)	0.0078 (14)
C22	0.043 (3)	0.041 (3)	0.050 (2)	0.012 (2)	0.024 (2)	0.003 (2)

Atomic displacement parameters $(Å^2)$

Geometric parameters (Å, °)

C1—C2	1.394 (5)	C13—C14	1.390 (5)
C1—C6	1.400 (5)	C14—O19	1.385 (4)
C1—N9	1.406 (5)	C14—C15	1.387 (5)
С2—С3	1.371 (5)	C15—C16	1.394 (5)
С2—Н2	0.93	C15—H15	0.93
C3—C4	1.365 (5)	C16—C17	1.377 (5)
C3—F7	1.320 (5)	C16—H16	0.93
C4—F8	1.361 (4)	C17—O21	1.386 (5)
C4—C5	1.369 (5)	C17—C18	1.391 (5)
C5—C6	1.369 (5)	C18—H18	0.93
С5—Н5	0.93	O19—C20	1.434 (5)
С6—Н6	0.93	C20—H20A	0.96
N9—C10	1.362 (4)	C20—H20B	0.96
N9—H9	0.84 (4)	C20—H20C	0.96
C10-011	1.235 (4)	O21—C22	1.439 (4)
C10—N12	1.370 (5)	C22—H22A	0.96
N12—C13	1.418 (5)	C22—H22B	0.96
N12—H12	0.80 (4)	C22—H22C	0.96
C13—C18	1.385 (5)		

C2C1C6	119.3 (4)	C14—C13—N12	117.5 (4)
C2C1N9	123.2 (3)	O19—C14—C15	125.3 (4)
C6C1N9	117.5 (3)	O19—C14—C13	114.7 (4)
C3—C2—C1	119.0 (3)	C15—C14—C13	120.1 (4)
С3—С2—Н2	120.5	C14—C15—C16	120.0 (4)
C1—C2—H2	120.5	C14—C15—H15	120
F7—C3—C4	118.0 (4)	C16—C15—H15	120
F7—C3—C2	120.9 (4)	C17—C16—C15	119.7 (4)
C4—C3—C2	121.1 (4)	C17—C16—H16	120.2
С4—С3—Н3	119.4	C15—C16—H16	120.2
С2—С3—Н3	119.4	C16—C17—O21	124.7 (4)
F8—C4—C3	119.4 (4)	C16—C17—C18	120.7 (4)
F8—C4—C5	120.1 (4)	O21—C17—C18	114.6 (4)
C3—C4—C5	120.5 (4)	C13—C18—C17	119.7 (4)
C6—C5—C4	119.9 (4)	C13—C18—H18	120.1
С6—С5—Н5	120	C17—C18—H18	120.1
С4—С5—Н5	120	C14—O19—C20	116.7 (3)
C5—C6—C1	120.1 (4)	O19—C20—H20A	109.5
С5—С6—Н6	119.9	O19—C20—H20B	109.5
С1—С6—Н6	119.9	H20A—C20—H20B	109.5
C10—N9—C1	126.7 (3)	O19—C20—H20C	109.5
С10—N9—Н9	115 (3)	H20A—C20—H20C	109.5
С1—N9—H9	118 (3)	H20B—C20—H20C	109.5
O11—C10—N9	123.7 (4)	C17—O21—C22	115.9 (3)
O11—C10—N12	122.9 (4)	O21—C22—H22A	109.5
N9-C10-N12	113.4 (3)	O21—C22—H22B	109.5
C10—N12—C13	125.9 (3)	H22A—C22—H22B	109.5
C10—N12—H12	120 (3)	O21—C22—H22C	109.5
C13—N12—H12	114 (3)	H22A—C22—H22C	109.5
C18—C13—C14	119.9 (4)	H22B—C22—H22C	109.5
C18—C13—N12	122.6 (4)		

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D····A	<i>D</i> —H··· <i>A</i>
N9—H9…O11 ⁱ N12_H12…O11 ⁱ	0.84(4) 0.80(4)	2.09 (4) 2 30 (4)	2.907 (4) 3.002 (4)	163 (4) 147 (4)
1112-1112 011	0.00 (+)	2.30 (+)	5.002 (+)	14/(4)

Symmetry code: (i) x, -y+1/2, z+1/2.