

A phthaloyl-linked benzotriazole corrosion inhibitor:
1,1'-(phthaloyl)bis(benzotriazole)Graham Smith,* Steven E. Bottle,
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Key indicators

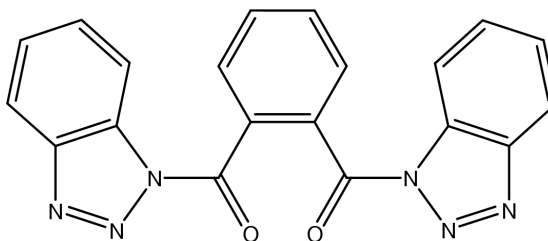
Single-crystal X-ray study
T = 295 K
Mean $\sigma(\text{C}-\text{C}) = 0.007 \text{ \AA}$
R factor = 0.037
wR factor = 0.138
Data-to-parameter ratio = 8.8For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.The crystal structure of the phthaloyl-linked benzotriazole 1,1'-(phthaloyl)bis(benzotriazole), $\text{C}_{20}\text{H}_{12}\text{N}_6\text{O}_2$, having potential as a corrosion inhibitor for copper, confirms that the *ortho*-related carboxamide substituents in the benzene parent ring adopt a *syn-syn* conformation with respect to one another.

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Comment

Benzotriazole (BTAH) has long been used and remains one of the most effective inhibitors of copper corrosion in neutral and alkaline conditions (Cotton, 1963; Cotton & Scholes, 1967; Poling, 1970; Mansfeld *et al.*, 1971). Since the mechanism of its inhibition effect is largely attributed to the formation of a polymeric BTA-Cu-BTA-Cu film (Rubim *et al.*, 1993; Ashour *et al.*, 1996), the BTAH derivatives, therefore, may offer higher inhibition efficiencies than the parent molecule, with the extended molecule giving better interaction with the metal surface. With this goal in mind and in an attempt to further understand the mechanism of corrosion inhibition of the carboxybenzotriazoles, a new dicarboxamide, 1,1'-(phthaloyl)bis(benzotriazole), (I), was synthesized (Reid, 1996) and its structure determined using single-crystal X-ray methods.

(I)

The analysis shows (Fig. 1) that the *ortho*-substituted carboxybenzotriazole side chains adopt a *syn-syn* conformation with respect to one another [torsion angles: $\text{C}1-\text{C}2-\text{C}21-\text{O}21 = -127.0 (5)^\circ$ and $\text{C}6-\text{C}1-\text{C}11-\text{O}11 = -135.2 (5)^\circ$]. These chains are slightly distorted from planarity [torsion angles: $\text{C}1-\text{C}11-\text{N}11-\text{N}12 = 16.3 (6)^\circ$ and $\text{C}2-\text{C}21-\text{N}21-\text{N}22 = -9.5 (6)^\circ$]. There are no intramolecular associations between the chains and, in addition, the packing in the unit cell shows no significant intermolecular interactions.

Experimental

The title compound was synthesized by the slow addition of 20 ml of a solution of phthaloyl chloride (4.3 ml, 60 mmol) in tetrahydrofuran to

100 ml of a solution containing benzotriazole (7.14 g, 60 mmol) and triethylamine (8.4 ml, 60 mmol) (also in tetrahydrofuran), maintained at 273–278 K. After stirring for *ca* 12 h, the mixture was poured on to ice and the resulting precipitate washed with water and recrystallized from chloroform from which data-quality crystals were obtained.

Crystal data

$C_{20}H_{12}N_6O_2$
 $M_r = 368.36$
 Orthorhombic, $P2_12_12_1$
 $a = 10.744$ (5) Å
 $b = 22.193$ (4) Å
 $c = 7.079$ (5) Å
 $V = 1687.9$ (14) Å³
 $Z = 4$
 $D_x = 1.450$ Mg m⁻³

Mo $K\alpha$ radiation
 Cell parameters from 25 reflections
 $\theta = 8.4$ – 14.7°
 $\mu = 0.10$ mm⁻¹
 $T = 295$ (2) K
 Plate, colourless
 $0.38 \times 0.25 \times 0.20$ mm

Data collection

Rigaku AFC-7R diffractometer
 ω - 2θ scans
 2231 measured reflections
 2231 independent reflections
 992 reflections with $I > 2\sigma(I)$
 $\theta_{\max} = 27.5^\circ$

$h = 0 \rightarrow 13$
 $k = 0 \rightarrow 28$
 $l = 0 \rightarrow 9$
 3 standard reflections
 every 150 reflections
 intensity decay: none

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.138$
 $S = 0.89$
 2231 reflections
 254 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0638P)^2 + 0.2455P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.18$ e Å⁻³
 $\Delta\rho_{\min} = -0.18$ e Å⁻³
 Extinction correction: *SHELXL97* (Sheldrick, 1997)
 Extinction coefficient: 0.014 (2)

The positional and isotropic displacement parameters of the H atoms were fixed in the refinement.

Data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1999a); cell refinement: *MSC/AFC Diffractometer Control Software*; data reduction: *TEXSAN for Windows* (Molecular Structure Corporation, 1999b); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON for Windows* (Spek, 1999); software used to prepare material for publication: *TEXSAN for Windows*.

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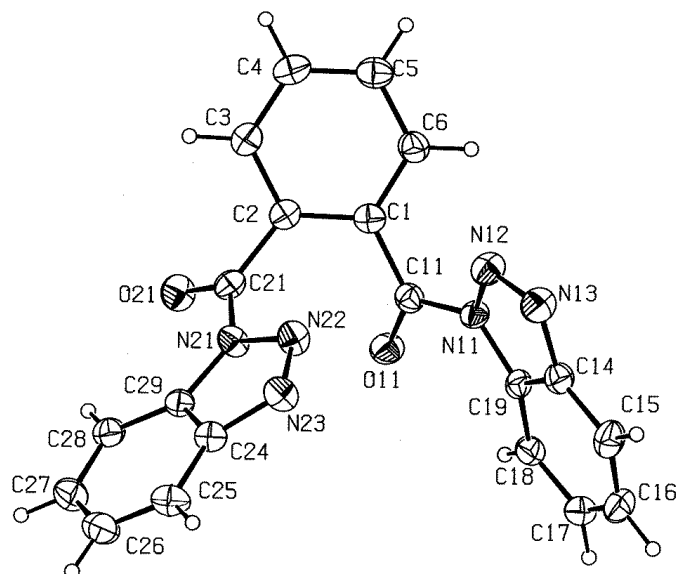


Figure 1

The molecular configuration and atom-numbering scheme for (I). Displacement ellipsoids are shown as the 30% probability level.

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