

2,2'-Bis(1-ethylbenzimidazol-2-yl)biphenyl

Robert T. Stibrany, Harvey J. Schugar and Joseph A. Potenza*

Department of Chemistry and Chemical Biology,
Rutgers, The State University of New Jersey,
610 Taylor Road, Piscataway, New Jersey
08854-8087, USACorrespondence e-mail:
potenza@rutchem.rutgers.edu

Key indicators

Single-crystal X-ray study
 $T = 294$ K
Mean $\sigma(\text{C}-\text{C}) = 0.004$ Å
 R factor = 0.060
 wR factor = 0.174
Data-to-parameter ratio = 10.2For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

The title compound, $\text{C}_{30}\text{H}_{26}\text{N}_4$, adopts a folded, twisted clamshell geometry in the solid state in which the ethyl groups are *trans*, and in which the imidazole fragments of the benzimidazole units are nearly eclipsed. Distances between imidazole fragments indicate extensive intramolecular $\pi-\pi$ interactions. The molecules pack in columns along b with little, if any, intermolecular $\pi-\pi$ interactions. In contrast, the *n*-propyl analog, 2,2'-bis(1-propylbenzimidazol-2-yl)biphenyl, adopts an extended conformation with little, if any, $\pi-\pi$ interactions of any kind.

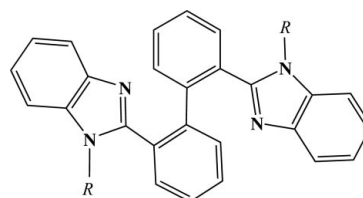
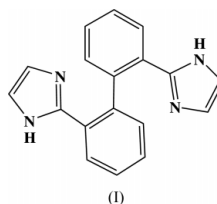
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Comment

Bis(imidazole)biphenyl and bis(benzimidazole)biphenyl species such as (I) and (II), with three torsional degrees of freedom, have been used as ligands to study electron self-exchange (Knapp *et al.*, 1990; Xie *et al.*, 1999), and to enforce near-tetrahedral coordination in transition metal complexes (Stibrany *et al.*, 2003). They have also found use in catalysis (Stibrany, Matturo *et al.*, 2002) and have been shown to act as proton sponges (Stibrany, Schugar & Potenza, 2002).



The title compound, (IIa) (Fig. 1), adopts a compact twisted clamshell conformation with the ethyl groups *trans*. Both the substituted phenyl (ph) and benzimidazole (bz) fragments are planar. Fragment planarity, coupled with the ph/ph [63.10 (8)°] and ph/bz [54.47 (6) and 48.29 (6)°] dihedral angles, give the molecule a pseudo-twofold axis passing through the midpoint of the C21–C31 bond and between the bz fragments. The shortest distances between C and N atoms of the nearly eclipsed imidazole fragments range from 3.167 (3) Å for C12...C42 to 3.673 (3) Å for N11...C43, indicative of intramolecular $\pi-\pi$ interactions. In type (II) species, the molecular shape correlates with the ph/ph dihedral angle. Thus, the

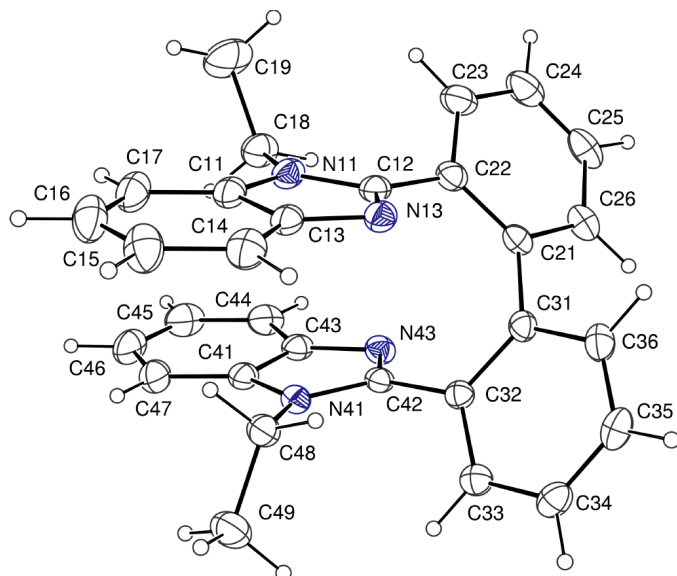
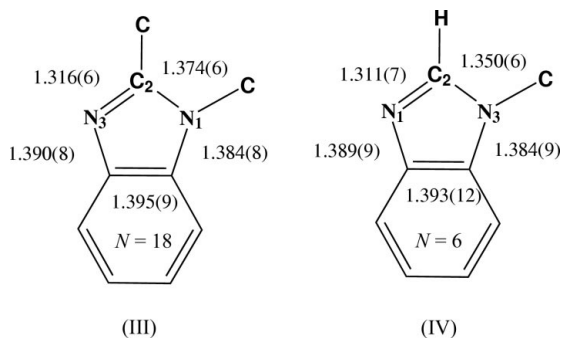


Figure 1
The molecular structure of the title compound, showing 25% probability displacement ellipsoids.

proton sponge IbH^+ (Stibrany, Schugar & Potenza, 2002) adopts a more open structure than (IIa), consistent with its larger ph/ph dihedral angle [$88.6(6)^\circ$], while molecules of (IIb), with ph/ph dihedral angles of $126.71(1)$ and $129.93(1)^\circ$ for two unique molecules, exhibit an open *trans* structure with little evidence of intra- or intermolecular π - π interactions (Stibrany *et al.*, 2003).



Metric parameters in the benzimidazole fragments of (IIa) agree well with those reported for 18 neutral benzimidazole structures (III) having carbon substituents at the N1 and C2 positions [Cambridge Structural Database, Version 5.24 (Allen, 2002); refcodes: DIKFEI, HIXXAX, HOHHUH, JELROH, LEBGOO, MEVTAI, MEVTEM, NUBCOC, QAXKAB, QIBPEW, QIFPOK, SAGDIN, UBIPUQ, XITZOP, YAXDAC and ZENSOA]. Benzimidazoles unsubstituted at the C2 position, (IV), show equivalent distances for the imidazole fragments, except for the the C2-N(amine) distances which are significantly shorter in the unsubstituted compounds (Allen, 2002; refcodes: CAXKUH, MIFVUS, NEBFUV, SEMWAI, SIGVIG and VENSAT). We have not explored the extent to which this difference arises from steric and/or electronic effects.

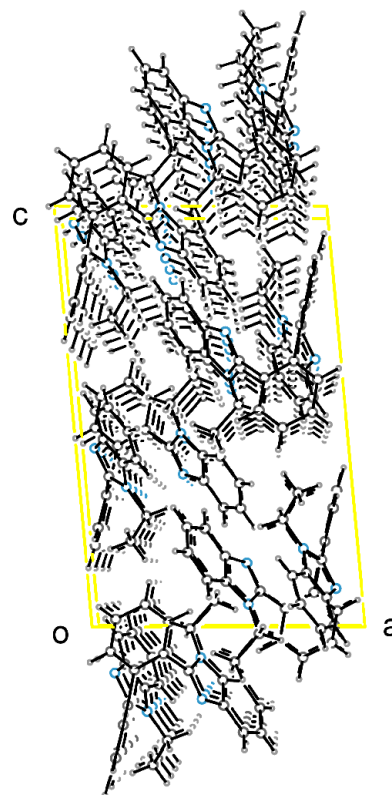


Figure 2
View of the unit cell along *b*. H atoms are omitted for clarity.

Molecules of (IIa) stack along *b* in columns (Fig. 2) related to each other by the centers of symmetry, glide planes and screw axes of the space group.

Experimental

The title compound, (IIa), was obtained by condensation of phenylenediamine with diphenic acid, followed by alkylation with ethyl iodide. Details of the preparation will be published elsewhere (Stibrany *et al.*, 2003). Crystals were obtained by slow evaporation of a solution of (IIa) in a mixture of ethanol and triethyl orthoformate. ^1H NMR (400 MHz, CDCl_3): δ 7.60 (*d*, $J = 7.0$ Hz, 2H), 7.41 (*d*, $J = 6.5$ Hz, 2H), 7.30 (*m*, 6H), 7.23 (*m*, 6H), 3.67 (*brd s*, 4H), 1.19 (*t*, $J = 7.0$ Hz, 6H). ^{13}C NMR (400 MHz, CDCl_3): δ 152.5 143.4, 141.1, 134.5, 131.5, 130.7, 130.0, 129.4, 127.41, 122.2, 121.8, 120.0, 110.0, 38.9, 14.8.

Crystal data

$\text{C}_{30}\text{H}_{26}\text{N}_4$
 $M_r = 442.55$
Monoclinic, $P2_1/c$
 $a = 12.0156(6)$ Å
 $b = 10.7001(6)$ Å
 $c = 18.6280(9)$ Å
 $\beta = 95.177(4)^\circ$
 $V = 2385.2(2)$ Å³
 $Z = 4$
 $D_x = 1.232$ Mg m⁻³
 $D_m = 1.19(1)$ Mg m⁻³

D_m , measured by flotation in carbon tetrachloride/cyclohexane
Mo $K\alpha$ radiation
Cell parameters from 821 reflections
 $\theta = 2.2$ – 23.1°
 $\mu = 0.07$ mm⁻¹
 $T = 294(1)$ K
Block, colorless
 $0.38 \times 0.36 \times 0.26$ mm

Data collection

Bruker SMART CCD area-detector diffractometer	4179 independent reflections 3375 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\text{int}} = 0.036$
Absorption correction: multi-scan (SADABS; Blessing, 1995)	$\theta_{\text{max}} = 25.0^\circ$
$T_{\text{min}} = 0.869$, $T_{\text{max}} = 1.000$	$h = -13 \rightarrow 14$
16281 measured reflections	$k = -12 \rightarrow 12$ $l = -22 \rightarrow 22$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.1086P)^2 + 0.4514P]$
$R[F^2 > 2\sigma(F^2)] = 0.060$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.174$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.00$	$\Delta\rho_{\text{max}} = 0.16 \text{ e } \text{\AA}^{-3}$
4179 reflections	$\Delta\rho_{\text{min}} = -0.13 \text{ e } \text{\AA}^{-3}$
411 parameters	
All H-atom parameters refined	

Table 1

Selected geometric parameters (Å).

N11—C12	1.383 (3)	N43—C43	1.387 (3)
N41—C42	1.385 (3)	N11—C11	1.387 (3)
N13—C12	1.314 (3)	N41—C41	1.385 (3)
N43—C42	1.312 (3)	C11—C13	1.395 (3)
N13—C13	1.387 (3)	C41—C43	1.393 (3)

Data collection: *SMART* (Bruker, 2000); cell refinement: *SMART*; data reduction: *SAINT* (Bruker, 2000); program(s) used to solve

structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPIII* (Burnett & Johnson, 1996) and *ORTEP32* (Farrugia, 1997); software used to prepare material for publication: *SHELXTL* (Bruker, 2000).

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