# organic compounds

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# 3,3'-Bis(3-methoxybenzyl)-1,1'-ethylenediimidazolium dibromide

#### Hon Man Lee\* and Chi-Ying Lu

National Changhua University of Education, Department of Chemistry, Changhua 50058, Taiwan Correspondence e-mail: leehm@cc.ncue.edu.tw

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Key indicators: single-crystal X-ray study; T = 298 K; mean  $\sigma$ (C–C) = 0.006 Å; R factor = 0.052; wR factor = 0.143; data-to-parameter ratio = 18.0.

In the title compound,  $C_{24}H_{28}N_4O_2^{2+}\cdot 2Br^-$ , the imidazolium cation is located on an inversion centre. The two imidazole rings are parallel to each other, whereas the imidazole and benzene rings make a dihedral angle of 77.25 (16)°. Non-classical intermolecular C-H···Br hydrogen bonds link the imidazolium cations and the bromide anions into a three-dimensional network.

#### **Related literature**

For the structure of 1,1'-bis(3-methoxybenzyl)-3,3'-methylenediimidazolium dibromide, see: Lee & Chiu (2004). For the structures of other related bis(imidazolium) salts, see: Cheng *et al.* (2006); Lee *et al.* (2004, 2007). For a review of *N*heterocyclic carbenes, see: Hillier *et al.* (2002).



#### **Experimental**

Crystal data  $C_{24}H_{28}N_4O_2^{2+}\cdot 2Br^{-1}M_r = 564.32$ 

Monoclinic,  $P2_1/c$ *a* = 18.340 (6) Å

b = 5.3566 (17) Å	
c = 12.340 (4) Å	
$\beta = 91.491 \ (9)^{\circ}$	
V = 1211.9 (7) Å <sup>3</sup>	
7 - 2	

### Data collection

Bruker SMART APEXII	6855 measured reflections
diffractometer	2609 independent reflections
Absorption correction: multi-scan	1838 reflections with $I > 2\sigma$
(SADABS; Sheldrick, 2003)	$R_{\rm int} = 0.050$
$T_{\min} = 0.366, T_{\max} = 0.600$	
Refinement	

 $R[F^{2} > 2\sigma(F^{2})] = 0.052$   $WR(F^{2}) = 0.143$  S = 1.00  $\Delta\rho_{max} = 0.71 \text{ e } \text{\AA}^{-3}$   $\Delta\rho_{min} = -1.24 \text{ e } \text{\AA}^{-3}$ 

Mo  $K\alpha$  radiation  $\mu = 3.37 \text{ mm}^{-1}$ 

 $0.35 \times 0.20 \times 0.15$  mm

T = 298 (2) K

# Table 1

Hydrogen-bond geometry (Å, °).  $\overline{D_{-}H_{\cdots}4}$   $D_{-}H_{-}H_{\cdots}4$ 

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$C2-H2A\cdots Br1^i$	0.93	2.77	3.657 (4)	161
C3−H3A···Br1 <sup>ii</sup>	0.93	2.91	3.729 (4)	148
$C4 - H4B \cdots Br1^{iii}$	0.97	2.85	3.669 (4)	143
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Symmetry codes: (i) x, y - 1, z; (ii)  $-x + 1, y - \frac{1}{2}, -z + \frac{1}{2}$ ; (iii)  $x, -y + \frac{3}{2}, z - \frac{1}{2}$ .

Data collection: *APEX2* (Bruker, 2004); cell refinement: *APEX2*; data reduction: *SAINT* (Bruker, 2004); program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: WN2283).

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supplementary materials

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## 3,3'-Bis(3-methoxybenzyl)-1,1'-ethylenediimidazolium dibromide

### H. M. Lee and C.-Y. Lu

#### Comment

In the past decade, N-heterocyclic carbenes (NHCs) and their palladium complexes have attracted much interest due to their catalytic activities in C—C coupling reactions (Hillier *et al.*, 2002). The structure of 1,1'-bis(3-methoxybenzyl)-3,3'-methylenediimidazolium dibromide has already been reported (Lee & Chiu, 2004). The structures of other related bis(imidazolium) salts have also been reported (Cheng *et al.*, 2006; Lee *et al.*, 2007).

One of the common methods for the preparation of palladium NHC complexes is a one-pot reaction between an imidazolium salt and a palladium precursor in the presence of base (Lee *et al.*, 2004). By this method, we prepared a palladium bis(NHC) complex from the title compound. Here, we report the crystal structure of the title compound.

The structure of the title compound is shown in Fig. 1. The bis(imidazolium) dication is located on an inversion center, with the two imidazole rings parallel to each other. The imidazole and benzene rings make a dihedral angle of 77.25 (16)°. The bromide anions are involved in intermolecular hydrogen bonds of the type C—H…Br with the imidazolium cations, forming a three-dimensional hydrogen-bonded network (Fig. 2 and Table 1).

#### **Experimental**

The compound was prepared according to the literature procedure (Lee *et al.*, 2004). Suitable crystals were obtained by slow diffusion of diethyl ether into a DMF solution of the compound at room temperature. The average dimensions of the colorless, rod-like crystals are about  $0.35 \times 0.20 \times 0.20$  mm.

#### Refinement

All hydrogen atoms could have been located in the difference Fourier map; nevertheless, they were all positioned geometrically and refined as riding atoms, with  $C_{aryl}$ —H = 0.93,  $C_{methyl}$ —H = 0.96,  $C_{methylene}$ —H = 0.97 Å;  $U_{iso}(H) = 1.5U_{eq}(C)$  for the methyl H atoms and  $U_{iso}(H) = 1.2U_{eq}(C)$  for all the other H atoms.

#### **Figures**



Fig. 1. The structure of the title compound, showing 50% probability displacement ellipsoids for the non-hydrogen atoms. The H atoms are depicted as circles of arbitrary radius. The unlabelled atoms of the imidazolium cation are related to the labelled ones by 3/2 - x, 3/2 - y, -z; for the anion, the symmetry operation for Br1 is 1 - x + 1, y, 1/2 - z.



Fig. 2. A view of the crystal packing along the b axis. Hydrogen bonds are shown as dashed lines.

# 3,3'-Bis(3-methoxybenzyl)-1,1'-ethylenediimidazolium dibromide

$C_{24}H_{28}N_4O_2^{2+}\cdot 2Br^-$	$F_{000} = 572$
$M_r = 564.32$	$D_{\rm x} = 1.546 {\rm ~Mg~m}^{-3}$
Monoclinic, $P2_1/c$	Mo K $\alpha$ radiation $\lambda = 0.71073$ Å
Hall symbol: -P 2ybc	Cell parameters from 1952 reflections
a = 18.340 (6) Å	$\theta = 3.3 - 26.3^{\circ}$
b = 5.3566 (17)  Å	$\mu = 3.37 \text{ mm}^{-1}$
c = 12.340 (4)  Å	T = 298 (2)  K
$\beta = 91.491 \ (9)^{\circ}$	Rod, white
$V = 1211.9 (7) \text{ Å}^3$	$0.35\times0.20\times0.15~mm$
Z = 2	

### Data collection

Bruker SMART APEXII diffractometer	1838 reflections with $I > 2\sigma$
Monochromator: graphite	$R_{\rm int} = 0.050$
T = 298(2)  K	$\theta_{\text{max}} = 27.0^{\circ}$
ω scans	$\theta_{\min} = 3.3^{\circ}$
Absorption correction: multi-scan (SADABS; Sheldrick, 2003)	$h = -14 \rightarrow 23$
$T_{\min} = 0.366, \ T_{\max} = 0.600$	$k = -6 \rightarrow 6$
6855 measured reflections	$l = -14 \rightarrow 15$
2609 independent reflections	

### Refinement

Refinement on $F^2$	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: difference Fourier map
$R[F^2 > 2\sigma(F^2)] = 0.052$	H-atom parameters constrained
$wR(F^2) = 0.143$	$w = 1/[\sigma^2(F_o^2) + (0.0825P)^2 + 0.0975P]$ where $P = (F_o^2 + 2F_c^2)/3$
S = 1.00	$(\Delta/\sigma)_{\rm max} = 0.001$
2609 reflections	$\Delta \rho_{max} = 0.71 \text{ e } \text{\AA}^{-3}$
145 parameters	$\Delta \rho_{\rm min} = -1.24 \text{ e } \text{\AA}^{-3}$

Primary atom site location: structure-invariant direct Extinction correction: none

#### Special details

**Geometry**. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

**Refinement**. Refinement of  $F^2$  against ALL reflections. The weighted *R*-factor *wR* and goodness of fit *S* are based on  $F^2$ , conventional *R*-factors *R* are based on *F*, with *F* set to zero for negative  $F^2$ . The threshold expression of  $F^2 > \sigma(F^2)$  is used only for calculating *R*-factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. *R*-factors based on  $F^2$  are statistically about twice as large as those based on *F*, and *R*- factors based on ALL data will be even larger.

	x	У	Ζ	$U_{\rm iso}*/U_{\rm eq}$
N1	0.43660 (17)	0.3916 (6)	0.1046 (3)	0.0292 (7)
N2	0.33797 (15)	0.2134 (6)	0.1563 (2)	0.0272 (7)
01	0.09061 (18)	0.8336 (7)	0.1978 (3)	0.0551 (9)
Br1	0.34564 (2)	0.59558 (9)	0.39315 (4)	0.0443 (2)
C1	0.3639 (2)	0.4034 (7)	0.1010 (3)	0.0318 (9)
H1A	0.3361	0.5252	0.0655	0.038*
C2	0.3950 (2)	0.0735 (7)	0.1966 (3)	0.0305 (9)
H2A	0.3917	-0.0702	0.2385	0.037*
C3	0.4573 (2)	0.1845 (8)	0.1637 (3)	0.0331 (9)
H3A	0.5048	0.1309	0.1783	0.040*
C4	0.4858 (2)	0.5616 (7)	0.0499 (3)	0.0301 (9)
H4A	0.5262	0.6056	0.0984	0.036*
H4B	0.4600	0.7136	0.0299	0.036*
C5	0.2612 (2)	0.1516 (9)	0.1779 (4)	0.0478 (13)
H5A	0.2538	0.1573	0.2554	0.057*
H5B	0.2511	-0.0173	0.1534	0.057*
C6	0.2085 (2)	0.3286 (8)	0.1218 (4)	0.0353 (10)
C7	0.1942 (2)	0.3068 (10)	0.0106 (4)	0.0504 (12)
H7A	0.2172	0.1854	-0.0300	0.060*
C8	0.1452 (3)	0.4690 (11)	-0.0375 (4)	0.0549 (13)
H8A	0.1360	0.4578	-0.1119	0.066*
C9	0.1094 (2)	0.6460 (9)	0.0203 (4)	0.0470 (12)
H9A	0.0763	0.7530	-0.0142	0.056*
C10	0.1230 (2)	0.6648 (8)	0.1315 (4)	0.0375 (10)
C11	0.1735 (2)	0.5067 (9)	0.1817 (4)	0.0361 (9)
H11A	0.1835	0.5211	0.2557	0.043*
C12	0.0360 (3)	0.9930 (12)	0.1522 (5)	0.0641 (15)
H12A	0.0182	1.1019	0.2073	0.096*
H12B	-0.0035	0.8940	0.1231	0.096*
H12C	0.0564	1.0909	0.0953	0.096*

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

# Atomic displacement parameters $(Å^2)$

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
N1	0.0268 (16)	0.0313 (18)	0.0295 (17)	-0.0003 (13)	0.0001 (13)	0.0021 (14)
N2	0.0241 (16)	0.0276 (18)	0.0298 (17)	0.0021 (13)	-0.0011 (13)	0.0040 (14)
01	0.053 (2)	0.053 (2)	0.060 (2)	0.0171 (16)	-0.0040 (16)	-0.0099 (17)
Br1	0.0467 (3)	0.0480 (3)	0.0380 (3)	-0.0161 (2)	-0.00102 (19)	0.0012 (2)
C1	0.030 (2)	0.030 (2)	0.036 (2)	0.0073 (16)	-0.0012 (16)	0.0039 (17)
C2	0.029 (2)	0.030 (2)	0.032 (2)	-0.0002 (16)	-0.0018 (16)	0.0090 (17)
C3	0.029 (2)	0.036 (2)	0.034 (2)	0.0047 (17)	-0.0043 (16)	0.0071 (18)
C4	0.031 (2)	0.031 (2)	0.028 (2)	-0.0029 (16)	-0.0002 (16)	0.0006 (16)
C5	0.021 (2)	0.051 (3)	0.071 (3)	-0.0005 (19)	0.004 (2)	0.024 (2)
C6	0.0225 (19)	0.037 (2)	0.047 (2)	-0.0039 (17)	-0.0012 (17)	0.0074 (19)
C7	0.038 (3)	0.059 (3)	0.055 (3)	0.008 (2)	0.007 (2)	-0.008 (3)
C8	0.054 (3)	0.075 (4)	0.036 (3)	0.009 (3)	-0.006 (2)	-0.002 (2)
C9	0.036 (2)	0.055 (3)	0.049 (3)	0.011 (2)	-0.006 (2)	0.010 (2)
C10	0.029 (2)	0.037 (2)	0.046 (3)	-0.0011 (18)	0.0003 (18)	-0.0017 (19)
C11	0.028 (2)	0.043 (2)	0.038 (2)	-0.0042 (19)	-0.0045 (17)	0.003 (2)
C12	0.052 (3)	0.056 (3)	0.084 (4)	0.020 (3)	0.003 (3)	-0.006 (3)

# Geometric parameters (Å, °)

N1—C1	1.334 (5)	С5—Н5А	0.9700
N1—C3	1.375 (5)	С5—Н5В	0.9700
N1—C4	1.460 (5)	C6—C11	1.376 (6)
N2—C1	1.321 (5)	C6—C7	1.396 (6)
N2—C2	1.369 (5)	С7—С8	1.374 (7)
N2—C5	1.477 (5)	С7—Н7А	0.9300
O1—C10	1.366 (5)	C8—C9	1.365 (7)
O1—C12	1.422 (6)	C8—H8A	0.9300
C1—H1A	0.9300	C9—C10	1.391 (6)
C2—C3	1.360 (6)	С9—Н9А	0.9300
C2—H2A	0.9300	C10—C11	1.389 (6)
С3—НЗА	0.9300	C11—H11A	0.9300
C4—C4 <sup>i</sup>	1.502 (7)	C12—H12A	0.9600
C4—H4A	0.9700	C12—H12B	0.9600
C4—H4B	0.9700	C12—H12C	0.9600
C5—C6	1.509 (6)		
C1—N1—C3	108.6 (3)	С6—С5—Н5В	109.2
C1—N1—C4	125.7 (3)	H5A—C5—H5B	107.9
C3—N1—C4	125.6 (3)	C11—C6—C7	120.5 (4)
C1—N2—C2	109.1 (3)	C11—C6—C5	119.5 (4)
C1—N2—C5	128.6 (3)	C7—C6—C5	120.0 (4)
C2—N2—C5	122.3 (3)	C8—C7—C6	118.5 (5)
C10—O1—C12	118.2 (4)	С8—С7—Н7А	120.8
N2—C1—N1	108.6 (3)	С6—С7—Н7А	120.8
N2—C1—H1A	125.7	C9—C8—C7	122.1 (5)

N1—C1—H1A	125.7	С9—С8—Н8А	119.0
C3—C2—N2	107.0 (3)	С7—С8—Н8А	119.0
C3—C2—H2A	126.5	C8—C9—C10	119.4 (4)
N2—C2—H2A	126.5	С8—С9—Н9А	120.3
C2—C3—N1	106.7 (3)	С10—С9—Н9А	120.3
С2—С3—НЗА	126.6	O1-C10-C11	115.5 (4)
N1—C3—H3A	126.6	O1—C10—C9	124.7 (4)
N1—C4—C4 <sup>i</sup>	109.6 (4)	C11—C10—C9	119.7 (4)
N1—C4—H4A	109.7	C6—C11—C10	119.9 (4)
C4 <sup>i</sup> —C4—H4A	109.7	C6—C11—H11A	120.1
N1—C4—H4B	109.7	C10-C11-H11A	120.1
C4 <sup>i</sup> —C4—H4B	109.7	O1—C12—H12A	109.5
H4A—C4—H4B	108.2	O1-C12-H12B	109.5
N2—C5—C6	112.3 (3)	H12A—C12—H12B	109.5
N2—C5—H5A	109.2	O1-C12-H12C	109.5
С6—С5—Н5А	109.2	H12A—C12—H12C	109.5
N2—C5—H5B	109.2	H12B—C12—H12C	109.5
Symmetry address (i) $w + 1 = w + 1 = w$			

Symmetry codes: (i) -x+1, -y+1, -z.

*Hydrogen-bond geometry (Å, °)* 

D—H···A	<i>D</i> —Н	$H \cdots A$	$D \cdots A$	D—H··· $A$
C2—H2A…Br1 <sup>ii</sup>	0.93	2.77	3.657 (4)	161
C3—H3A…Br1 <sup>iii</sup>	0.93	2.91	3.729 (4)	148
C4—H4B…Br1 <sup>iv</sup>	0.97	2.85	3.669 (4)	143

Symmetry codes: (ii) x, y-1, z; (iii) -x+1, y-1/2, -z+1/2; (iv) x, -y+3/2, z-1/2.

Fig. 1





Fig. 2