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## Di-tert-butyl 3,3'-(2,2'-bi-1H-imidazole-1,1'-diyl)dipropanoate

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Received 3 April 2009; accepted 5 June 2009
Key indicators: single-crystal X-ray study; $T=295 \mathrm{~K}$; mean $\sigma(\mathrm{C}-\mathrm{C})=0.005 \AA$; $R$ factor $=0.077 ; w R$ factor $=0.190 ;$ data-to-parameter ratio $=19.2$.

In the title compound, $\mathrm{C}_{20} \mathrm{H}_{20} \mathrm{~N}_{4} \mathrm{O}_{4}$, the complete molecule is generated by a crystallographic centre of symmetry. The conformation is stabilized by two intramolecular $\mathrm{C}-\mathrm{H} \cdots \mathrm{N}$ links.

## Related literature

For the background to 2, 2'-biimidazole derivatives, see: Barnett et al. (1999, 2002); Liang et al. (2009); Zhang \& Liang (2009); Zhang, Zhang, Ren et al. (2009); Zhang, Zhang, Xu et al. (2009). For the synthesis, see: Barnett et al. (1999).


## Experimental

## Crystal data

$$
\begin{aligned}
& \mathrm{C}_{20} \mathrm{H}_{30} \mathrm{~N}_{4} \mathrm{O}_{4} \\
& M_{r}=390.48 \\
& \text { Monoclinic, } P 2_{2} / c \\
& a=7.0321(14) \AA \\
& b=17.484(4) \AA \\
& c=8.9681(18) \AA \\
& \beta=100.80(3)^{\circ}
\end{aligned}
$$

## Data collection

Rigaku R-AXIS RAPID diffractometer
Absorption correction: multi-scan (ABSCOR; Higashi, 1995) $T_{\text {min }}=0.960, T_{\text {max }}=0.988$

## Refinement

$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.077$

## 128 parameters

$w R\left(F^{2}\right)=0.190$
H -atom parameters constrained
$S=1.02$
2453 reflections

9620 measured reflections 2453 independent reflections
1474 reflections with $I>2 \sigma(I)$ $R_{\text {int }}=0.077$

Table 1
Hydrogen-bond geometry ( $\AA,{ }^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{C} 10-\mathrm{H} 10 B \cdots \mathrm{~N} 2^{\mathrm{i}}$ | 0.97 | 2.46 | $2.960(3)$ | 111 |
| Symmetry code: (i) $-x+1,-y+1,-z+1$. |  |  |  |  |

Data collection: RAPID-AUTO (Rigaku, 1998); cell refinement: RAPID-AUTO; data reduction: CrystalStructure (Rigaku/MSC, 2004); program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXL97.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FB2148).

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## supporting information

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## Di-tert-butyl 3,3'-(2,2'-bi-1H-imidazole-1,1'-diyl)dipropanoate

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## S1. Comment

Biimidazole is a potentially polydentate ligand, but its chemistry is less developed in comparison to the imidazole chemistry. The reason may be a limited solubility of biimidazole in common organic solvents. Although several new disubstituted 2,2'-biimidazoles have been recently synthesized (Barnett et al., 1999; Barnett et al., 2002), a few metal complexes based on biimidazole derivatives are reported (Zhang, Zhang, Ren et al., 2009). In the course of our ongoing study, we have successfully synthesized a series of biimidazole derivatives with terminal carboxylic, hydroxyl, phosphino, imino groups which can be used as ligands in the coordination chemistry (Zhang \& Liang, 2009; Zhang, Zhang, Xu et al., 2009) and cross-coupling reactions (Liang et al., 2009). These ligands exhibit rich coordination patterns and catalytic properties. Here we report the synthesis and the crystal structure of the title compound which is an intermediate of those above mentioned ligands. As shown in Fig. 1, the biimidazole ring atoms (C6, C7, C9, N1, N2 and their inversion-related partners) exhibit essentially coplanar mutual orientation [the dihedral angle is $0.00(1)^{\circ}$ ], and the value of the torsion angle $\mathrm{C} 9-\mathrm{N} 1-\mathrm{C} 10-\mathrm{C} 5$ is $-77.53(30)^{\circ}$. In the crystal structure, there are weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{N}$ interactions (Tab. 1, Fig. 2).

## S2. Experimental

The title compound was prepared according to a published procedure (Barnett et al., 1999). $0.2 \mathrm{~g}(5 \mathrm{mmol})$ of NaOH was added to a suspension of $3 \mathrm{~g}(22.4 \mathrm{mmol})$ of $2,2^{\prime}$-biimidazole in 100 ml of DMF (dimethylformamide) at $80^{\circ} \mathrm{C}$. The resulting mixture was stirrred for 30 min . In the course of this time the mixture gradually turned into a clear pale yellow solution. $7.12 \mathrm{~g}(55.6 \mathrm{mmol})$ butyl acrylate in 10 ml of DMF was added dropwise in several minutes to the solution and the reaction was stirred at $80^{\circ} \mathrm{C}$ for 8 h until the heating was stopped. The DMF was removed via vacuum distillation in a hot oil bath at $100^{\circ} \mathrm{C}$. The resulting black brown oil was dissolved in water $(30 \mathrm{ml})$ and extracted with $\mathrm{CH}_{2} \mathrm{Cl}_{2}$. The organic layer was washed with water, and then evaporated under reduced pressure to yield a white product ( $7.4 \mathrm{~g}, 85 \%$ ). The product was dissolved in $95 \%$ ethanol $(30 \mathrm{ml})$ and cooled slowly in a refrigerator to afford colourless block crystals of average size $1.5 \mathrm{~mm} \times 1.2 \mathrm{~mm} \times 0.5 \mathrm{~mm}$ that were suitable for the X-ray analysis.

## S3. Refinement

All the hydrogens were discernible in the difference electron density map. Nevertheless, the hydrogens were situated into the idealized positions. The C-H distances were constrained to $0.93,0.96$ and $0.97 \AA$ for aryl, methylene and methyl hydrogens, respectively. $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}\left(\mathrm{C}_{\text {aryl }}\right) ; U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}\left(\mathrm{C}_{\text {methylene }}\right) ; U_{\text {iso }}(\mathrm{H})=1.5 U_{\text {eq }}\left(\mathrm{C}_{\text {methyl }}\right)$.


## Figure 1

View of the title molecule with the displacement ellipsoids at the $45 \%$ probability level. The labelling of the non-H atoms is also given. The symmetry code (i):-x+1, $-y+1,-z+1$.


Figure 2
A section of the title structure. The C-H $\cdots \mathrm{N}$ hydrogen bonds (Tab. 1) are shown as dashed lines.

## Di-tert-butyl 3,3'-(2,2'-bi-1 H-imidazole-1,1'-diyl)dipropanoate

## Crystal data

$\mathrm{C}_{20} \mathrm{H}_{30} \mathrm{~N}_{4} \mathrm{O}_{4}$
$M_{r}=390.48$
Monoclinic, $P 2_{1} / c$
Hall symbol: -P 2ybc
$a=7.0321$ (14) $\AA$
$b=17.484$ (4) $\AA$
$c=8.9681$ (18) $\AA$
$\beta=100.80(3)^{\circ}$
$V=1083.1$ (4) $\AA^{3}$
$Z=2$

$$
\begin{aligned}
& F(000)=420 \\
& D_{\mathrm{x}}=1.197 \mathrm{Mg} \mathrm{~m}^{-3} \\
& \text { Mo Ka radiation, } \lambda=0.71073 \AA \\
& \text { Cell parameters from } 1478 \text { reflections } \\
& \theta=3.0-27.5^{\circ} \\
& \mu=0.08 \mathrm{~mm}^{-1} \\
& T=295 \mathrm{~K} \\
& \text { Plate, colourless } \\
& 0.48 \times 0.42 \times 0.14 \mathrm{~mm}
\end{aligned}
$$

## Data collection

Rigaku R-AXIS RAPID
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
Detector resolution: 0 pixels $\mathrm{mm}^{-1}$

## $\omega$ scans

Absorption correction: multi-scan
(ABSCOR; Higashi, 1995)
$T_{\text {min }}=0.960, T_{\text {max }}=0.988$

## Refinement

Refinement on $F^{2}$
Least-squares matrix: full
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.077$
$w R\left(F^{2}\right)=0.190$
$S=1.02$
2453 reflections
128 parameters
0 restraints
59 constraints

> 9620 measured reflections
> 2453 independent reflections
> 1474 reflections with $I>2 \sigma(I)$
> $R_{\text {int }}=0.077$
> $\theta_{\max }=27.5^{\circ}, \theta_{\min }=3.2^{\circ}$
> $h=-9 \rightarrow 9$
> $k=-22 \rightarrow 22$
> $l=-11 \rightarrow 10$

Primary atom site location: structure-invariant direct methods
Secondary atom site location: difference Fourier map
Hydrogen site location: difference Fourier map
H -atom parameters constrained
$w=1 /\left[\sigma^{2}\left(F_{0}^{2}\right)+(0.0718 P)^{2}+0.3943 P\right]$
where $P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3$
$(\Delta / \sigma)_{\text {max }}<0.001$
$\Delta \rho_{\text {max }}=0.32 \mathrm{e} \AA^{-3}$
$\Delta \rho_{\text {min }}=-0.24 \mathrm{e} \AA^{-3}$

## 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 1.s. planes.
Refinement. Refinement of $F^{2}$ against ALL reflections. The weighted $R$-factor $w R$ 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 $F^{2}>\sigma\left(F^{2}\right)$ 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.

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

|  | $x$ | $y$ | $z$ | $U_{\text {iso }} * / U_{\text {eq }}$ |
| :--- | :--- | :--- | :--- | :--- |
| N1 | $0.3460(3)$ | $0.51573(11)$ | $0.30746(19)$ | $0.0410(5)$ |
| N2 | $0.6689(3)$ | $0.51901(13)$ | $0.3698(2)$ | $0.0522(6)$ |
| C10 | $0.1400(3)$ | $0.50398(14)$ | $0.3101(3)$ | $0.0454(6)$ |


|  |  |  |  |  |
| :--- | :--- | :--- | :--- | :--- |
| H10A | 0.0628 | 0.5297 | 0.2237 | $0.054^{*}$ |
| H10B | 0.1099 | 0.5268 | 0.4016 | $0.054^{*}$ |
| O2 | $0.2468(3)$ | $0.32667(12)$ | $0.1854(2)$ | $0.0717(6)$ |
| O1 | $0.0366(3)$ | $0.40123(12)$ | $0.0365(2)$ | $0.0774(7)$ |
| C9 | $0.5051(3)$ | $0.50815(13)$ | $0.4210(2)$ | $0.0415(6)$ |
| C8 | $0.1171(4)$ | $0.38275(15)$ | $0.1612(3)$ | $0.0525(6)$ |
| C7 | $0.6104(4)$ | $0.53329(17)$ | $0.2169(3)$ | $0.0568(7)$ |
| H7A | 0.6939 | 0.5428 | 0.1501 | $0.068^{*}$ |
| C6 | $0.4159(4)$ | $0.53160(15)$ | $0.1774(2)$ | $0.0504(6)$ |
| H6A | 0.3431 | 0.5396 | 0.0809 | $0.061^{*}$ |
| C5 | $0.0864(4)$ | $0.41992(15)$ | $0.3056(3)$ | $0.0520(6)$ |
| H5A | 0.1644 | 0.3940 | 0.3914 | $0.062^{*}$ |
| H5B | -0.0484 | 0.4147 | 0.3144 | $0.062^{*}$ |
| C4 | $0.2854(6)$ | $0.2868(2)$ | $0.0492(4)$ | $0.0905(11)$ |
| H4A | 0.3139 | 0.3238 | -0.0242 | $0.109^{*}$ |
| H4B | 0.1724 | 0.2577 | 0.0027 | $0.109^{*}$ |
| C3 | $0.4524(6)$ | $0.2349(2)$ | $0.0943(5)$ | $0.0958(12)$ |
| H3A | 0.4239 | 0.2005 | 0.1722 | $0.115^{*}$ |
| H3B | 0.4664 | 0.2039 | 0.0072 | $0.115^{*}$ |
| C2 | $0.6387(6)$ | $0.2728(2)$ | $0.1520(5)$ | $0.1045(13)$ |
| H2B | 0.6290 | 0.3010 | 0.2434 | $0.125^{*}$ |
| H2C | 0.6653 | 0.3092 | 0.0770 | $0.125^{*}$ |
| C1 | $0.8074(6)$ | $0.2163(3)$ | $0.1876(7)$ | $0.1289(18)$ |
| H1A | 0.9241 | 0.2435 | 0.2285 | $0.193^{*}$ |
| H1B | 0.8229 | 0.1904 | 0.0962 | $0.193^{*}$ |
| H1C | 0.7809 | 0.1795 | 0.2605 |  |

Atomic displacement parameters $\left(\AA^{2}\right)$

|  | $U^{11}$ | $U^{22}$ | $U^{33}$ | $U^{12}$ | $U^{13}$ | $U^{23}$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| N1 | $0.0380(10)$ | $0.0516(11)$ | $0.0319(9)$ | $0.0030(9)$ | $0.0030(7)$ | $0.0023(8)$ |
| N2 | $0.0407(11)$ | $0.0834(16)$ | $0.0336(10)$ | $0.0013(10)$ | $0.0097(8)$ | $0.0031(10)$ |
| C10 | $0.0364(11)$ | $0.0564(15)$ | $0.0413(12)$ | $0.0045(11)$ | $0.0017(9)$ | $0.0019(10)$ |
| O2 | $0.0728(13)$ | $0.0805(15)$ | $0.0564(12)$ | $0.0151(11)$ | $-0.0018(9)$ | $-0.0074(10)$ |
| O1 | $0.0988(16)$ | $0.0791(15)$ | $0.0457(11)$ | $0.0097(12)$ | $-0.0087(10)$ | $0.0001(9)$ |
| C9 | $0.0377(11)$ | $0.0553(14)$ | $0.0304(11)$ | $0.0037(11)$ | $0.0033(8)$ | $-0.0010(9)$ |
| C8 | $0.0518(14)$ | $0.0532(15)$ | $0.0491(15)$ | $-0.0082(12)$ | $0.0006(11)$ | $0.0019(11)$ |
| C7 | $0.0522(15)$ | $0.086(2)$ | $0.0343(12)$ | $-0.0015(13)$ | $0.0134(10)$ | $0.0047(12)$ |
| C6 | $0.0539(14)$ | $0.0675(17)$ | $0.0289(11)$ | $0.0016(12)$ | $0.0051(10)$ | $0.0033(11)$ |
| C5 | $0.0474(13)$ | $0.0629(16)$ | $0.0444(13)$ | $-0.0046(12)$ | $0.0054(10)$ | $0.0044(12)$ |
| C4 | $0.100(3)$ | $0.098(3)$ | $0.069(2)$ | $0.020(2)$ | $0.0058(18)$ | $-0.0199(19)$ |
| C3 | $0.097(3)$ | $0.095(3)$ | $0.098(3)$ | $-0.005(2)$ | $0.026(2)$ | $-0.021(2)$ |
| C2 | $0.101(3)$ | $0.095(3)$ | $0.115(3)$ | $-0.006(2)$ | $0.013(2)$ | $-0.001(2)$ |
| C1 | $0.087(3)$ | $0.101(3)$ | $0.198(5)$ | $0.000(2)$ | $0.025(3)$ | $0.031(3)$ |

Geometric parameters (A, ${ }^{\circ}$ )

| N1-C9 | 1.371 (3) | C6-H6A | 0.9300 |
| :---: | :---: | :---: | :---: |
| N1-C6 | 1.376 (3) | C5-H5A | 0.9700 |
| N1-C10 | 1.468 (3) | C5-H5B | 0.9700 |
| N2-C9 | 1.331 (3) | C4-C3 | 1.480 (5) |
| N2-C7 | 1.378 (3) | C4-H4A | 0.9700 |
| C10-C5 | 1.516 (3) | C4-H4B | 0.9700 |
| C10-H10A | 0.9700 | C3-C2 | 1.473 (5) |
| C10-H10B | 0.9700 | C3-H3A | 0.9700 |
| O2-C8 | 1.329 (3) | C3-H3B | 0.9700 |
| O2-C4 | 1.475 (4) | C2-C1 | 1.531 (5) |
| O1-C8 | 1.199 (3) | C2-H2B | 0.9700 |
| C9- $\mathrm{C}^{\text {i }}$ | 1.460 (4) | C2-H2C | 0.9700 |
| C8-C5 | 1.500 (4) | C1-H1A | 0.9600 |
| C7-C6 | 1.347 (3) | C1-H1B | 0.9600 |
| C7-H7A | 0.9300 | C1-H1C | 0.9600 |
| C9-N1-C6 | 106.13 (18) | C10-C5-H5B | 109.3 |
| C9-N1-C10 | 130.15 (19) | H5A-C5-H5B | 108.0 |
| C6-N1-C10 | 123.53 (18) | $\mathrm{O} 2-\mathrm{C} 4-\mathrm{C} 3$ | 108.9 (3) |
| C9-N2-C7 | 104.6 (2) | $\mathrm{O} 2-\mathrm{C} 4-\mathrm{H} 4 \mathrm{~A}$ | 109.9 |
| N1-C10-C5 | 112.1 (2) | C3-C4-H4A | 109.9 |
| N1-C10-H10A | 109.2 | O2-C4-H4B | 109.9 |
| C5-C10-H10A | 109.2 | C3-C4-H4B | 109.9 |
| N1-C10-H10B | 109.2 | H4A-C4-H4B | 108.3 |
| C5-C10-H10B | 109.2 | C2-C3-C4 | 115.3 (4) |
| H10A-C10-H10B | 107.9 | C2-C3-H3A | 108.4 |
| C8-O2-C4 | 116.1 (2) | C4-C3-H3A | 108.4 |
| N2-C9-N1 | 111.59 (19) | C2-C3-H3B | 108.4 |
| N2-C9-C9 ${ }^{\text {i }}$ | 124.5 (2) | C4-C3- 33 B | 108.4 |
| N1-C9-C9 ${ }^{\text {i }}$ | 123.9 (2) | H3A-C3-H3B | 107.5 |
| O1-C8-O2 | 122.7 (3) | C3-C2-C1 | 112.7 (4) |
| O1-C8-C5 | 124.7 (3) | C3-C2-H2B | 109.0 |
| O2-C8-C5 | 112.6 (2) | C1-C2-H2B | 109.0 |
| C6-C7-N2 | 110.9 (2) | C3-C2-H2C | 109.0 |
| C6-C7-H7A | 124.5 | $\mathrm{C} 1-\mathrm{C} 2-\mathrm{H} 2 \mathrm{C}$ | 109.0 |
| N2-C7-H7A | 124.5 | H2B-C2-H2C | 107.8 |
| C7-C6-N1 | 106.7 (2) | C2-C1-H1A | 109.5 |
| C7-C6-H6A | 126.6 | C2-C1-H1B | 109.5 |
| N1-C6-H6A | 126.6 | H1A-C1-H1B | 109.5 |
| C8-C5-C10 | 111.6 (2) | C2-C1-H1C | 109.5 |
| C8-C5-H5A | 109.3 | H1A-C1-H1C | 109.5 |
| C10-C5-H5A | 109.3 | H1B-C1-H1C | 109.5 |
| C8-C5-H5B | 109.3 |  |  |

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

## supporting information

Hydrogen-bond geometry (A, o)

| $D — \mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{C} 10 — \mathrm{H} 10 B \cdots \mathrm{~N} 2^{\mathrm{i}}$ | 0.97 | 2.46 | $2.960(3)$ | 111 |

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

