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2-[4-(4,5-Dihydro-1*H*-pyrrol-2-yl)-phenyl]-4,5-dihydro-1*H*-imidazole. Corrigendum

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The chemical name of the title compound in the paper by Kia, Fun & Kargar [*Acta Cryst.* (2009), **E64**, o2406] is corrected.

In the paper by Kia, Fun & Kargar [*Acta Cryst.* (2009), **E64**, o2406], the chemical name given in the *Title* should be '2,2'-(*p*-phenylene)bis(4,5-dihydro-1*H*-imidazole)'.
C1CN(C1)c2ccc(cc2)N3CCCN3

2-[4-(4,5-Dihydro-1H-pyrrol-2-yl)-phenyl]-4,5-dihydro-1H-imidazole

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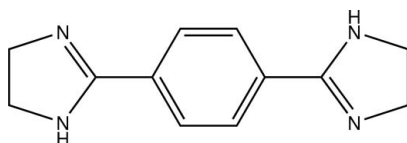
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.001$ Å; R factor = 0.041; wR factor = 0.116; data-to-parameter ratio = 12.8.

The molecule of the title compound, $\text{C}_{12}\text{H}_{14}\text{N}_4$, lies about a crystallographic inversion centre. The five- and six-membered rings are twisted from each other, forming a dihedral angle of 18.06 (7)°. In the crystal structure, neighbouring molecules are linked by intermolecular $\text{N}-\text{H}\cdots\text{N}$ hydrogen bonds into one-dimensional infinite chains forming 18-membered rings with $R_2^2(18)$ motifs. The crystal structure is further stabilized by weak intermolecular $\pi-\pi$ stacking [centroid-centroid distance = 3.8254 (6) Å] and $\text{C}-\text{H}\cdots\pi$ interactions.

Related literature

For details of hydrogen-bond motifs, see: Bernstein *et al.* (1995). For a related structure and synthesis, see: Stibrany *et al.* (2004). For applications, see: Blancafart (1978); Chan (1993); Vizi (1986); Li *et al.* (1996); Ueno *et al.* (1995); Corey & Grogan (1999).



Experimental

Crystal data

$\text{C}_{12}\text{H}_{14}\text{N}_4$	$\gamma = 94.207$ (2)°
$M_r = 214.27$	$V = 247.52$ (2) Å ³
Triclinic, $P\bar{1}$	$Z = 1$
$a = 4.8863$ (2) Å	Mo $K\alpha$ radiation
$b = 5.1472$ (2) Å	$\mu = 0.09$ mm ⁻¹
$c = 10.2295$ (4) Å	$T = 100.0$ (1) K
$\alpha = 104.414$ (2)°	$0.56 \times 0.17 \times 0.15$ mm
$\beta = 93.885$ (2)°	

Data collection

Bruker SMART APEXII CCD area-detector diffractometer	4616 measured reflections 1296 independent reflections 1208 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Bruker, 2005)	$R_{\text{int}} = 0.026$
$T_{\text{min}} = 0.950$, $T_{\text{max}} = 0.986$	

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.041$	101 parameters
$wR(F^2) = 0.116$	All H-atom parameters refined
$S = 1.07$	$\Delta\rho_{\text{max}} = 0.42$ e Å ⁻³
1296 reflections	$\Delta\rho_{\text{min}} = -0.24$ e Å ⁻³

Table 1

Hydrogen-bond geometry (Å, °).

Cg1 is the centroid of the N1/C1/C2/N2/C3 imidazoline ring.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N1}-\text{H1N1}\cdots\text{N2}^{\text{i}}$	0.87 (2)	2.18 (2)	3.0060 (13)	158.1 (15)
$\text{C2}-\text{H2B}\cdots\text{Cg1}^{\text{ii}}$	1.015 (15)	2.980 (15)	3.8882 (11)	149.6 (11)

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

Data collection: APEX2 (Bruker, 2005); cell refinement: APEX2; data reduction: SAINT (Bruker, 2005); 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 and PLATON (Spek, 2003).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: TK2331).

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supporting information

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2-[4-(4,5-Dihydro-1H-pyrrol-2-yl)phenyl]-4,5-dihydro-1H-imidazole**Reza Kia, Hoong-Kun Fun and Hadi Kargar****S1. Comment**

Imidazoline derivatives are of great importance because they exhibit significant biological and pharmacological activities including anti-hypertensive (Blancafort 1978), anti-hyperglycemic (Chan 1993), anti-depressive (Vizi 1986), anti-hypercholesterolemic (Li *et al.*, 1996) and anti-inflammatory (Ueno *et al.*, 1995) activities. These compounds are also used as catalysts and synthetic intermediates in some organic reactions (Corey & Grogan 1999). In consideration of the important applications of imidazolines, herein the crystal structure of the title compound, (I), is reported.

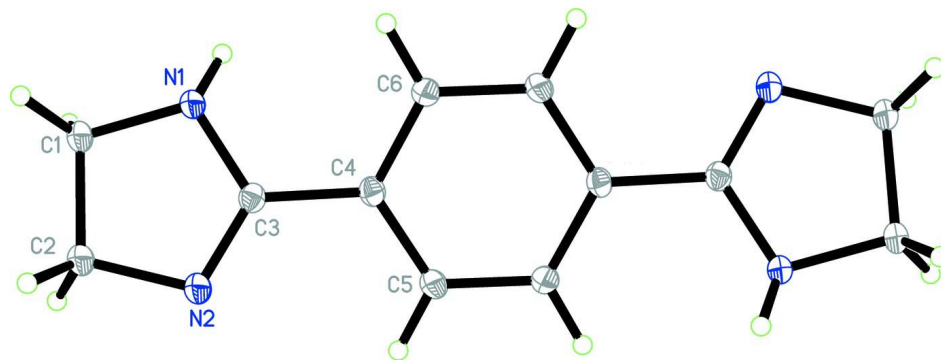
In compound (I), Fig. 1, bond lengths and angles are within the normal ranges and are comparable with a related structure (Stibrany *et al.*, 2004). The molecule lies about a crystallographic inversion centre. The five- and six-membered rings are twisted from each other, forming a dihedral angle of 18.06 (7)°. Intermolecular N—H...N hydrogen bonds form 18-membered rings producing $R_2^2(18)$ ring motifs to link molecules into one-dimensional infinite chains along the *b*-axis, Table 1 and Fig. 2. The crystal structure is further stabilized by weak intermolecular π – π stacking [$Cg1 \cdots Cg2^i = 3.8254(6)$ Å; (i) $1 + x, y, z$] and C—H... π ($Cg1$ and $Cg2$ are the centroids of the N1/C1/C2/N2/C3 imidazoline ring and the benzene ring, respectively) interactions, Table 1.

S2. Experimental

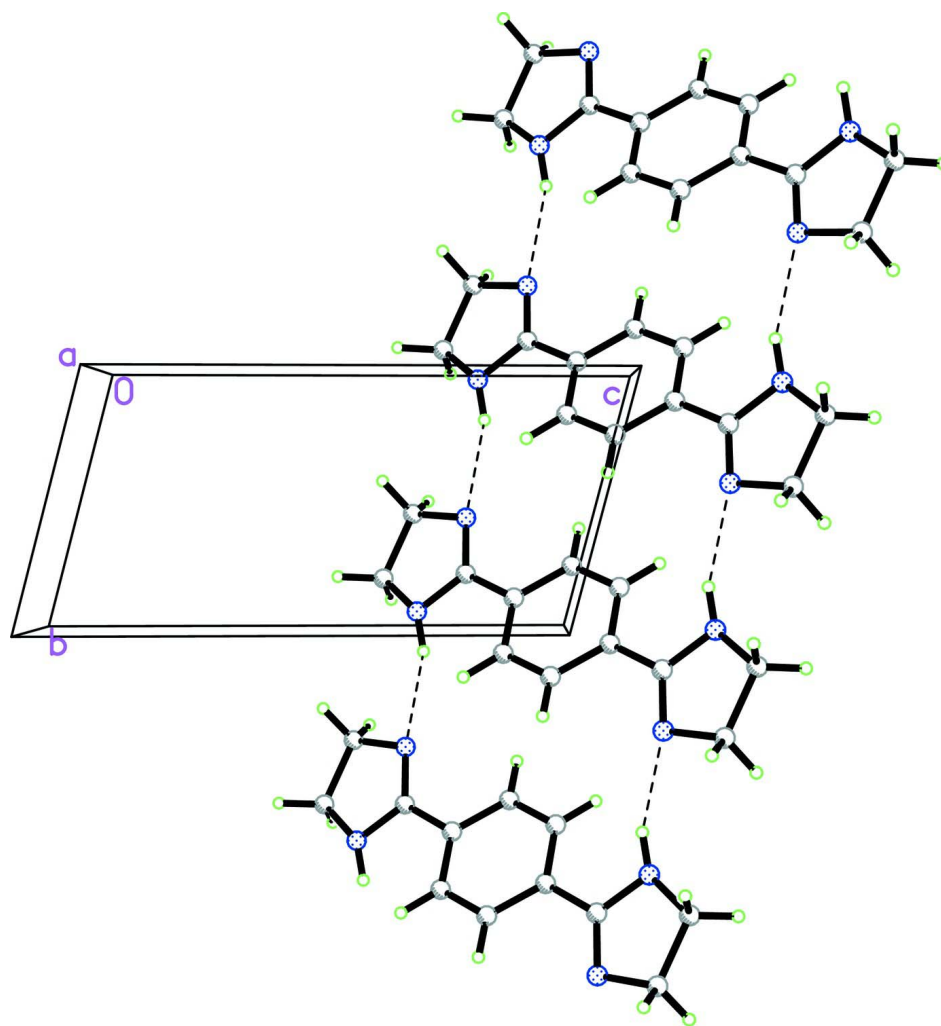
The synthetic method used for the preparation of (I) was based on previous work (Stibrany *et al.* 2004), except that 1,4-dicyanobenzene (10 mmol) and ethylenediamine (40 mmol) were used. Single crystals suitable for X-ray diffraction were obtained by evaporation of a methanol solution of (I) held at room temperature.

S3. Refinement

All hydrogen atoms were located from a difference Fourier map and refined freely: C—H ranged from 0.961 (16) to 1.015 (15) Å and N—H was 0.874 (18) Å.

**Figure 1**

The molecular structure of (I) with atom labels and 50% probability ellipsoids for non-H atoms. Unlabelled atoms are related by $-x + 1, -y, -z$.

**Figure 2**

Partial crystal packing in (I), viewed down the a -axis showing one-dimensional infinite chains along the b -axis mediated by intermolecular N—H...N interactions (dashed lines).

2-[4-(4,5-Dihydro-1H-pyrrol-2-yl)-phenyl]-4,5-dihydro-1H-imidazole

Crystal data

$C_{12}H_{14}N_4$	$Z = 1$
$M_r = 214.27$	$F(000) = 114$
Triclinic, $P\bar{1}$	$D_x = 1.437 \text{ Mg m}^{-3}$
Hall symbol: -P 1	Melting point: 312 K
$a = 4.8863 (2) \text{ \AA}$	Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
$b = 5.1472 (2) \text{ \AA}$	Cell parameters from 3789 reflections
$c = 10.2295 (4) \text{ \AA}$	$\theta = 2.5\text{--}30.3^\circ$
$\alpha = 104.414 (2)^\circ$	$\mu = 0.09 \text{ mm}^{-1}$
$\beta = 93.885 (2)^\circ$	$T = 100 \text{ K}$
$\gamma = 94.207 (2)^\circ$	Block, colourless
$V = 247.52 (2) \text{ \AA}^3$	$0.56 \times 0.17 \times 0.15 \text{ mm}$

Data collection

Bruker SMART APEXII CCD area-detector diffractometer	4616 measured reflections
Radiation source: fine-focus sealed tube	1296 independent reflections
Graphite monochromator	1208 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\text{int}} = 0.026$
Absorption correction: multi-scan (SADABS; Bruker, 2005)	$\theta_{\text{max}} = 29.0^\circ$, $\theta_{\text{min}} = 4.1^\circ$
$T_{\text{min}} = 0.950$, $T_{\text{max}} = 0.986$	$h = -6 \rightarrow 6$
	$k = -6 \rightarrow 6$
	$l = -13 \rightarrow 13$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.041$	All H-atom parameters refined
$wR(F^2) = 0.116$	$w = 1/[\sigma^2(F_o^2) + (0.0717P)^2 + 0.0714P]$
$S = 1.07$	where $P = (F_o^2 + 2F_c^2)/3$
1296 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
101 parameters	$\Delta\rho_{\text{max}} = 0.42 \text{ e \AA}^{-3}$
0 restraints	$\Delta\rho_{\text{min}} = -0.24 \text{ e \AA}^{-3}$
Primary atom site location: structure-invariant direct methods	

Special details

Experimental. The low-temperature data was collected with the Oxford Cyrosystem Cobra low-temperature attachment.

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.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	0.99649 (17)	0.02202 (18)	0.30177 (9)	0.0140 (2)
N2	1.05027 (18)	0.42785 (18)	0.25064 (9)	0.0161 (2)

C1	1.2376 (2)	0.1595 (2)	0.39310 (10)	0.0148 (2)
C2	1.2547 (2)	0.4433 (2)	0.36664 (10)	0.0158 (2)
C3	0.92585 (19)	0.1869 (2)	0.21989 (10)	0.0124 (2)
C4	0.70754 (19)	0.08912 (19)	0.10716 (10)	0.0124 (2)
C5	0.6787 (2)	0.2263 (2)	0.00581 (10)	0.0138 (2)
C6	0.5270 (2)	-0.1391 (2)	0.09997 (10)	0.0135 (2)
H1A	1.402 (3)	0.062 (3)	0.3667 (15)	0.020 (3)*
H1B	1.212 (3)	0.169 (3)	0.4906 (16)	0.024 (4)*
H2A	1.443 (3)	0.498 (3)	0.3429 (16)	0.026 (4)*
H2B	1.214 (3)	0.587 (3)	0.4488 (15)	0.020 (3)*
H5	0.802 (3)	0.383 (3)	0.0096 (16)	0.025 (4)*
H6	0.537 (3)	-0.239 (3)	0.1687 (15)	0.020 (3)*
H1N1	0.989 (3)	-0.151 (4)	0.2656 (17)	0.029 (4)*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0135 (4)	0.0131 (4)	0.0147 (4)	-0.0009 (3)	-0.0033 (3)	0.0041 (3)
N2	0.0147 (4)	0.0150 (4)	0.0177 (5)	-0.0009 (3)	-0.0044 (3)	0.0047 (3)
C1	0.0125 (4)	0.0157 (5)	0.0153 (5)	-0.0008 (3)	-0.0030 (3)	0.0037 (4)
C2	0.0144 (5)	0.0153 (5)	0.0167 (5)	-0.0011 (4)	-0.0038 (3)	0.0041 (4)
C3	0.0102 (4)	0.0142 (5)	0.0130 (4)	0.0015 (3)	0.0006 (3)	0.0036 (3)
C4	0.0098 (4)	0.0134 (5)	0.0133 (4)	0.0009 (3)	0.0000 (3)	0.0025 (3)
C5	0.0109 (4)	0.0143 (5)	0.0159 (5)	-0.0009 (3)	0.0001 (3)	0.0041 (3)
C6	0.0128 (4)	0.0140 (5)	0.0142 (5)	0.0002 (3)	0.0003 (3)	0.0049 (3)

Geometric parameters (Å, °)

N1—C3	1.3780 (13)	C2—H2B	1.015 (15)
N1—C1	1.4700 (12)	C3—C4	1.4787 (13)
N1—H1N1	0.874 (18)	C4—C5	1.3973 (14)
N2—C3	1.2944 (13)	C4—C6	1.4000 (14)
N2—C2	1.4808 (12)	C5—C6 ⁱ	1.3881 (13)
C1—C2	1.5479 (14)	C5—H5	0.961 (16)
C1—H1A	0.997 (14)	C6—C5 ⁱ	1.3881 (13)
C1—H1B	1.004 (16)	C6—H6	0.970 (15)
C2—H2A	1.006 (16)		
C3—N1—C1	107.38 (8)	C1—C2—H2B	112.1 (9)
C3—N1—H1N1	118.4 (11)	H2A—C2—H2B	106.8 (12)
C1—N1—H1N1	119.9 (10)	N2—C3—N1	116.89 (9)
C3—N2—C2	106.43 (8)	N2—C3—C4	123.28 (9)
N1—C1—C2	102.00 (8)	N1—C3—C4	119.77 (9)
N1—C1—H1A	108.9 (8)	C5—C4—C6	118.99 (9)
C2—C1—H1A	112.8 (9)	C5—C4—C3	119.82 (9)
N1—C1—H1B	112.3 (9)	C6—C4—C3	121.19 (9)
C2—C1—H1B	111.4 (9)	C6 ⁱ —C5—C4	120.61 (9)
H1A—C1—H1B	109.3 (12)	C6 ⁱ —C5—H5	119.7 (9)

N2—C2—C1	106.30 (8)	C4—C5—H5	119.6 (9)
N2—C2—H2A	109.2 (9)	C5 ⁱ —C6—C4	120.40 (9)
C1—C2—H2A	111.9 (9)	C5 ⁱ —C6—H6	117.6 (9)
N2—C2—H2B	110.6 (8)	C4—C6—H6	122.0 (9)
C3—N1—C1—C2	9.79 (10)	N1—C3—C4—C5	-165.14 (9)
C3—N2—C2—C1	3.06 (11)	N2—C3—C4—C6	-161.93 (10)
N1—C1—C2—N2	-7.84 (10)	N1—C3—C4—C6	15.11 (14)
C2—N2—C3—N1	3.71 (12)	C6—C4—C5—C6 ⁱ	0.46 (16)
C2—N2—C3—C4	-179.17 (8)	C3—C4—C5—C6 ⁱ	-179.29 (8)
C1—N1—C3—N2	-9.24 (12)	C5—C4—C6—C5 ⁱ	-0.46 (16)
C1—N1—C3—C4	173.53 (8)	C3—C4—C6—C5 ⁱ	179.29 (8)
N2—C3—C4—C5	17.82 (14)		

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

Hydrogen-bond geometry (Å, °)

<i>D—H...A</i>	<i>D—H</i>	<i>H...A</i>	<i>D...A</i>	<i>D—H...A</i>
N1—H1N1...N2 ⁱⁱ	0.87 (2)	2.18 (2)	3.0060 (13)	158.1 (15)
C2—H2B...Cg1 ⁱⁱⁱ	1.015 (15)	2.980 (15)	3.8882 (11)	149.6 (11)

Symmetry codes: (ii) $x, y-1, z$; (iii) $-x, -y+1, -z+1$.