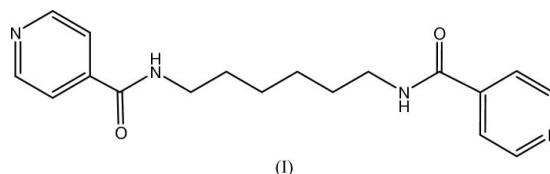


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zhangshush@public.qd.sd.cn**Key indicators**Single-crystal X-ray study  
 $T = 293$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å  
 $R$  factor = 0.041  
 $wR$  factor = 0.113  
Data-to-parameter ratio = 14.8For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.**1,6-Bis(pyridine-4-carboxamido)hexane**

In the title compound,  $\text{C}_{18}\text{H}_{22}\text{N}_4\text{O}_2$ , the molecule has a crystallographically imposed centre of symmetry. Intermolecular  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds link the molecules into ribbons along the  $a$  axis. The crystal packing is further stabilized by weak  $\text{C}-\text{H}\cdots\text{N}$  interactions.

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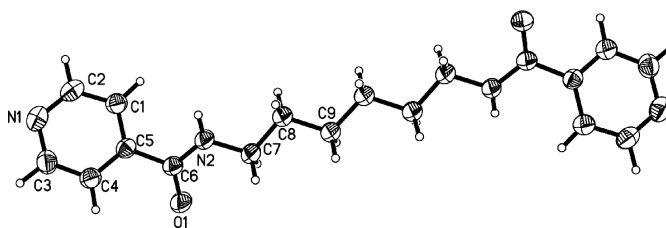
In recent years, the synthesis and characterization of coordination polymers have attracted increased interest, owing to their fascinating structural diversity and potential applications as functional materials (Tang & Guloy, 1999; Debajyoti *et al.*, 2004). As a good candidate for rigid rodlike spacers in the construction of metal-organic polymers, 4,4'-bipyridine has been relatively well known and has shown hundreds of interesting supramolecular architectures (Chen *et al.*, 2001). For this reason, and as a continuation of our search for new pyridyl-donor ligands, we have synthesized the title compound, (I) (Fig. 1).



The bond lengths and angles (Table 1) in (I) show normal values (Allen *et al.*, 1987). The molecule has a crystallographic inversion centre. Intermolecular  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds (Table 2) link the molecules into ribbons along the  $a$  axis. The packing is further stabilized into a three-dimensional framework by weak intermolecular  $\text{C}-\text{H}\cdots\text{N}$  hydrogen bonds (Table 2).

**Experimental**

Hexane-1,6-diamine (1 g, 17 mmol) was added slowly to a pyridine solution (50 ml) of pyridine-4-carboxylic acid (4.23 g, 34 mmol). The



**Figure 1**  
View of (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme. Unlabelled atoms are related to labelled atoms by  $(-1-x, -y, -z)$ .

mixture was stirred for 15 min and triphenyl phosphite (9 ml, 34 mmol) was added slowly through a dropping funnel over a period of 15 min. The mixture was refluxed for 6 h and the volume was then reduced to 10 ml by vacuum evaporation. A white precipitate was obtained from the solution after allowing it to stand at room temperature for 24 h. The solid was filtered off and washed with cold water. Yellow crystals were obtained by slow evaporation of an ethanol solution of the compound.

Crystal data

$C_{18}H_{22}N_4O_2$   $D_x = 1.308 \text{ Mg m}^{-3}$   
 $M_r = 326.40$  Mo  $K\alpha$  radiation  
 Monoclinic,  $P2_1/c$  Cell parameters from 1592 reflections  
 $a = 5.0940(9) \text{ \AA}$   $\theta = 2.6\text{--}25.9^\circ$   
 $b = 5.2911(10) \text{ \AA}$   $\mu = 0.09 \text{ mm}^{-1}$   
 $c = 30.759(6) \text{ \AA}$   $T = 293(2) \text{ K}$   
 $\beta = 91.067(3)^\circ$  Plate, yellow  
 $V = 828.9(3) \text{ \AA}^3$   $0.41 \times 0.22 \times 0.06 \text{ mm}$   
 $Z = 2$

Data collection

Bruker SMART 1000 CCD area-detector diffractometer 1616 independent reflections  
 1363 reflections with  $I > 2\sigma(I)$   
 $\omega$  scans  $R_{int} = 0.016$   
 Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)  $\theta_{max} = 26.0^\circ$   
 $T_{min} = 0.965$ ,  $T_{max} = 0.995$   $h = -6 \rightarrow 5$   
 4366 measured reflections  $k = -6 \rightarrow 6$   
 $l = -37 \rightarrow 31$

Refinement

Refinement on  $F^2$   $w = 1/[\sigma^2(F_o^2) + (0.0567P)^2 + 0.1501P]$   
 $R[F^2 > 2\sigma(F^2)] = 0.041$  where  $P = (F_o^2 + 2F_c^2)/3$   
 $wR(F^2) = 0.113$   $(\Delta/\sigma)_{max} < 0.001$   
 $S = 1.07$   $\Delta\rho_{max} = 0.16 \text{ e \AA}^{-3}$   
 1616 reflections  $\Delta\rho_{min} = -0.14 \text{ e \AA}^{-3}$   
 109 parameters  
 H-atom parameters constrained

Table 1

Selected geometric parameters ( $\text{\AA}$ ,  $^\circ$ ).

O1—C6	1.2315 (16)	N2—C6	1.3325 (18)
N1—C2	1.330 (2)	N2—C7	1.4584 (18)
N1—C3	1.334 (2)		
C6—N2—C7	120.91 (11)	O1—C6—C5	120.73 (13)
O1—C6—N2	122.12 (13)	N2—C6—C5	117.14 (12)

Table 2

Hydrogen-bonding geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N2—H2A $\cdots$ O1 <sup>i</sup>	0.85	2.16	2.985 (1)	162
C2—H2B $\cdots$ N1 <sup>ii</sup>	0.93	2.60	3.466 (2)	156

Symmetry codes: (i)  $x - 1, y, z$ ; (ii)  $-1 - x, y - \frac{1}{2}, \frac{1}{2} - z$ .

All H atoms were located in difference Fourier maps and refined isotropically. All H atoms were placed at idealized positions and

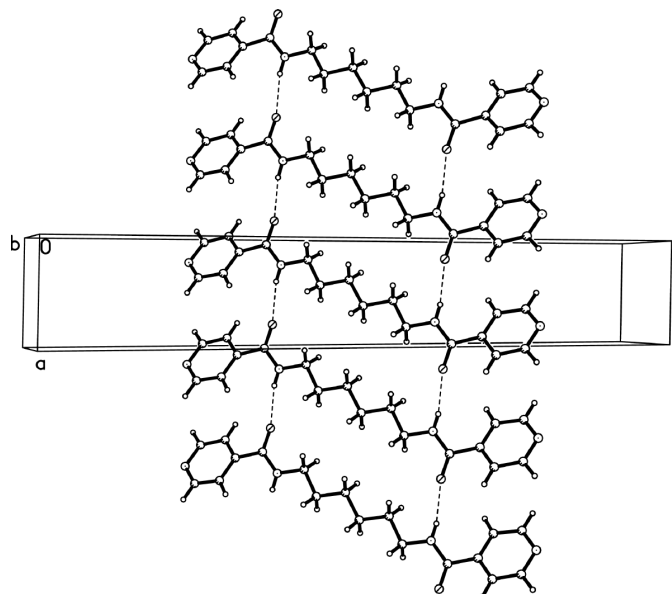


Figure 2

Packing diagram of (I), showing the hydrogen-bonded (dashed lines) ribbons.

allowed to ride on their parent C atom, with C—H = 0.93–0.97  $\text{\AA}$  and  $U_{iso}(H) = 1.2$  and  $1.5U_{eq}(C)$ .

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINTE* (Siemens, 1996); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 1997); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*, *PARST* (Nardelli, 1995) and *PLATON* (Spek, 2003).

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