

Molecular Self Assembly: Solvent Guests Tune the  
Conformation of a Series of 2,6-Bis(2-  
anilinoethynyl)pyridine-Based Ureas

*Jeffrey M. Engle, P. S. Lakshminarayanan, Calden N. Carroll, Lev N. Zakharov,\**

*Michael M. Haley\* and Darren W. Johnson\**

Department of Chemistry and Materials Science Institute,

University of Oregon,

Eugene, Oregon 97403-1253 (USA)

**Table S1.** Crystallographic Data and Data Collections and Refinements Parameters for the investigated structures:

	<b>L<sup>1</sup>•2DMSO</b>	<b>L<sup>1</sup>•2DMSO•CH<sub>3</sub>OH</b>	<b>L<sup>1</sup>•3DMF</b>
Empirical formula	C <sub>47</sub> H <sub>51</sub> N <sub>7</sub> O <sub>8</sub> S <sub>2</sub>	C <sub>48</sub> H <sub>55</sub> N <sub>7</sub> O <sub>9</sub> S <sub>2</sub>	C <sub>52</sub> H <sub>60</sub> N <sub>10</sub> O <sub>9</sub>
Formula weight	906.07	938.11	969.10
Temperature	173(2) K	173(2) K	173(2) K
Wavelength	0.71073 Å	0.71073 Å	0.71073 Å
Crystal system	Monoclinic	Triclinic	Triclinic
Space group	<i>P2<sub>1</sub>/c</i>	<i>P</i> -1	<i>P</i> -1
Unit cell dimensions	a = 14.7007(13) Å b = 25.956(2) Å c = 12.3794(11) Å α = 90° β = 103.568(1)° γ = 90°.	a = 13.037(8) Å b = 14.106(9) Å c = 14.707(9) Å α = 112.755(10) β = 103.142(10)° γ = 92.983(11)°	a = 9.6860(14) Å b = 12.7435(18) Å c = 20.683(3) Å α = 93.436(3)° β = 98.492(3)° γ = 95.516(3)°
Volume	4591.8(7) Å <sup>3</sup>	2399(3) Å <sup>3</sup>	2506.1(6) Å <sup>3</sup>
Z	4	2	2
Density (calculated)	1.311 Mg/m <sup>3</sup>	1.299 Mg/m <sup>3</sup>	1.284 Mg/m <sup>3</sup>
Absorption coefficient	0.177 mm <sup>-1</sup>	0.173 mm <sup>-1</sup>	0.090 mm <sup>-1</sup>
F(000)	1912	992	1028
Crystal size	0.33 x 0.23 x 0.12 mm <sup>3</sup>	0.38 x 0.22 x 0.05 mm <sup>3</sup>	0.19 x 0.04 x 0.02 mm <sup>3</sup>
Theta range for data collection	1.57 to 25.00°	1.58 to 24.00°.	1.00 to 25.00°.
Index ranges	-17<=h<=17, -30<=k<=30, -14<=l<=14	-14<=h<=14, -16<=k<=16, -16<=l<=16	-11<=h<=11, -15<=k<=15, -24<=l<=24

Reflections collected	44074	20819	24347
Independent reflections	8098 [R(int) = 0.0233]	7533 [R(int) = 0.0580]	8816 [R(int) = 0.0638]
Completeness to theta = 25.00°	100.0 %	99.9 %	99.8 %
Absorption correction	Semi-empirical equivalents	from Semi-empirical equivalents	from Semi-empirical equivalents
Max. and min. transmission	0.9791 and 0.9439	0.9914 and 0.9370	0.9982 and 0.9832
Refinement method	Full-matrix least-squares on F <sup>2</sup>	Full-matrix least-squares on F <sup>2</sup>	Full-matrix least-squares on F <sup>2</sup>
Data / restraints / parameters	8098 / 0 / 593	7533 / 6 / 666	8816 / 6 / 654
Goodness-of-fit on F <sup>2</sup>	1.014	1.032	1.005
Final R indices [I > 2sigma(I)]	R1 = 0.0473, wR2 = 0.1226	R1 = 0.0867, wR2 = 0.2233	R1 = 0.0698, wR2 = 0.1632
R indices (all data)	R1 = 0.0538, wR2 = 0.1290	R1 = 0.1382, wR2 = 0.2502	R1 = 0.1309, wR2 = 0.2028
Largest diff. peak and hole	0.748 and -0.613 e.Å <sup>-3</sup>	0.609 and -0.759 e.Å <sup>-3</sup>	0.499 and -0.270 e.Å <sup>-3</sup>

	<b>L<sup>2</sup>•H<sub>2</sub>O •0.5DMSO</b>	<b>L<sup>3</sup>•H<sub>2</sub>O•DMSO (I)</b>	<b>L<sup>3</sup>•H<sub>2</sub>O•DMSO (II)</b>
Empirical formula	C <sub>38</sub> H <sub>26</sub> F <sub>6</sub> N <sub>7</sub> O <sub>7.50</sub> S <sub>0.50</sub>	C <sub>41</sub> H <sub>35</sub> F <sub>6</sub> N <sub>5</sub> O <sub>6</sub> S	C <sub>41</sub> H <sub>35</sub> F <sub>6</sub> N <sub>5</sub> O <sub>6</sub> S
Formula weight	830.69	839.80	839.80
Temperature	173(2) K	173(2) K	173(2) K
Wavelength	0.71073 Å	0.71073 Å	0.71073 Å
Crystal system	Monoclinic	Triclinic	Triclinic
Space group	<i>P2/n</i>	<i>P</i> -1	<i>P</i> -1
Unit cell dimensions	<i>a</i> = 14.8341(13) Å <i>b</i> = 9.5759(9) Å <i>c</i> = 29.903(3) Å $\alpha$ = 90° $\beta$ = 90.690(2)° $\gamma$ = 90°	<i>a</i> = 8.7292(10) Å <i>b</i> = 13.3810(15) Å <i>c</i> = 17.3731(19) Å $\alpha$ = 84.395(2)° $\beta$ = 78.492(2)° $\gamma$ = 80.087(2)°	<i>a</i> = 9.533(2) Å <i>b</i> = 14.185(3) Å <i>c</i> = 16.112(4) Å $\alpha$ = 73.871(4)° $\beta$ = 78.002(4)° $\gamma$ = 71.375(4)°
Volume	4247.3(7) Å <sup>3</sup>	1954.6(4) Å <sup>3</sup>	1966.3(7) Å <sup>3</sup>
Z	4	2	2
Density (calculated)	1.299 Mg/m <sup>3</sup>	1.427 Mg/m <sup>3</sup>	1.418 Mg/m <sup>3</sup>
Absorption coefficient	0.133 mm <sup>-1</sup>	0.167 mm <sup>-1</sup>	0.166 mm <sup>-1</sup>
F(000)	1700	868	868
Crystal size	0.43 x 0.12 x 0.03 mm <sup>3</sup>	0.33 x 0.12 x 0.03 mm <sup>3</sup>	0.45 x 0.06 x 0.04 mm <sup>3</sup>
Theta range for data collection	1.36 to 25.00°.	1.20 to 25.00°.	1.56 to 25.00°
Index ranges	-17<= <i>h</i> <=17, -11<= <i>k</i> <=11, - 35<= <i>l</i> <=35	10<= <i>h</i> <=10, - 15<= <i>k</i> <=15, -20<= <i>l</i> <=20	- 11<= <i>h</i> <=11, -16<= <i>k</i> <=16, - 19<= <i>l</i> <=19
Reflections collected	39732	19048	18895
Independent reflections	7464 [R(int) = 0.0439]	6884 [R(int) = 0.0204]	6893 [R(int) = 0.0428]

Completeness to theta = 25.00°	99.7 %	99.8 %	99.7 %
Absorption correction	Semi-empirical from equivalents	Semi-empirical from equivalents	Semi-empirical from equivalents
Max. and min. transmission	0.9935 and 0.9128	0.9950 and 0.9471	0.9934 and 0.9292
Refinement method	Full-matrix least-squares on F <sup>2</sup>	Full-matrix least-squares on F <sup>2</sup>	Full-matrix least-squares on F <sup>2</sup>
Data / restraints / parameters	7464 / 0 / 561	6884 / 0 / 648	6893 / 0 / 556
Goodness-of-fit on F <sup>2</sup>	1.049	1.002	1.024
Final R indices [I>2sigma(I)]	R1 = 0.0722, wR2 = 0.1943	R1 = 0.0569, wR2 = 0.1643	R1 = 0.0671, wR2 = 0.1623
R indices (all data)	R1 = 0.0963, wR2 = 0.2080	R1 = 0.0711, wR2 = 0.1820	R1 = 0.1144, wR2 = 0.1919
Largest diff. peak and hole	0.574 and -0.342 e.Å <sup>-3</sup>	0.882 and -0.518 e.Å <sup>-3</sup>	0.818 and -0.361 e.Å <sup>-3</sup>

**Table S2.** Hydrogen bonding interactions in **L<sup>1</sup>•2DMSO** [Å and °].

D-H...A	d(D-H)	d(H...A)	d(D...A)	<(DHA)
N(5)-H(5N)...O(8)	0.81(3)	2.20(3)	2.935(3)	151(2)
N(6)-H(6N)...O(8)	0.85(3)	1.99(3)	2.790(3)	156(2)
N(3)-H(3N)...O(7)	0.83(3)	1.93(3)	2.757(2)	169(2)
N(2)-H(2N)...O(7)	0.80(3)	2.42(3)	3.122(2)	148(2)

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**Table S3.** Hydrogen bonding interactions in  $L^1 \cdot 2DMSO \cdot CH_3OH$  [ $\text{\AA}$  and  $^\circ$ ].

D-H...A	d(D-H)	d(H...A)	d(D...A)	<(DHA)
N(2)-H(2N)...O(1S)	1.01(6)	2.41(6)	3.202(5)	135(4)
N(6)-H(6N)...O(2S)	1.06(5)	1.71(5)	2.767(6)	172(4)
O(1S)-H(1S)...N(1)	1.07(7)	1.69(7)	2.731(5)	163(6)
N(5)-H(5N)...O(2S)	1.15(6)	2.25(6)	3.229(6)	141(4)
N(3)-H(3N)...O(1S)	0.95(5)	1.84(5)	2.786(5)	174(4)

**Table S4.** Hydrogen bonding interactions of  $L^1 \cdot 3DMF$  [ $\text{\AA}$  and  $^\circ$ ].

D-H...A	d(D-H)	d(H...A)	d(D...A)	<(DHA)
N(2)-H(2N)...O(1S)#1	0.86(4)	2.35(4)	3.137(4)	151(3)
N(3)-H(3N)...O(1S)#1	0.88(4)	2.02(4)	2.851(4)	159(4)
N(5)-H(5N)...O(2S)	0.84(3)	2.36(4)	3.052(4)	140(3)
N(6)-H(6N)...O(2S)	0.91(4)	1.93(4)	2.825(4)	168(3)

Symmetry transformations used to generate equivalent atoms: #1 x+1, y, z

**Table S5.** Hydrogen bonding interactions in  $L^2 \cdot H_2O \cdot 0.5DMSO$  [ $\text{\AA}$  and  $^\circ$ ].

D-H...A	d(D-H )	d(H...A)	d(D...A)	<(DHA)
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N(2)-H(2N)···O(1S)	0.91(4)	2.49(4)	3.229(4)	138(3)
N(3)-H(3N)···O(1S)	0.89(4)	1.92(4)	2.795(4)	167(3)
O(1S)-H(1S)···N(1)	0.75(5)	2.12(5)	2.867(4)	174(5)
O(1S)-H(2S)···O(4)#2	0.90(10)	1.92(10)	2.821(4)	177(9)

Symmetry transformations used to generate equivalent atoms:

#1 -x+1/2,y,-z+1/2 #2 -x+1,-y+1,-z

**Table S6.** Hydrogen bonding interactions of  $L^3 \cdot H_2O \cdot DMSO$  (I) [ $\text{\AA}$  and  $^\circ$ ]

D-H···A	d(D-H)	d(H···A)	d(D···A)	<(DHA)
N(5)-H(5N)···S(1S)#1	0.86(4)	2.92(4)	3.660(3)	145(3)
N(5)-H(5N)···O(1S)#1	0.86(4)	1.95(4)	2.783(3)	163(3)
O(5)-H(5B)···O(3)#1	0.71(5)	2.10(5)	2.804(3)	169(5)
O(5)-H(5A)···N(1)	0.77(4)	2.09(4)	2.809(3)	157(3)
N(3)-H(3N)···O(5)	0.89(3)	1.93(3)	2.799(3)	165(3)
N(4)-H(4N)···O(1S)#1	0.81(3)	2.33(3)	3.030(3)	147(2)
N(2)-H(2N)···O(5)	0.84(3)	2.28(3)	3.030(3)	148(3)

Symmetry transformations used to generate equivalent atoms:

#1 -x+1,-y+2,-z+1

**Table S7.** Hydrogen bonding interactions in  $L^3 \cdot H_2O \cdot DMSO$  (II) [ $\text{\AA}$ ,  $^\circ$ ]

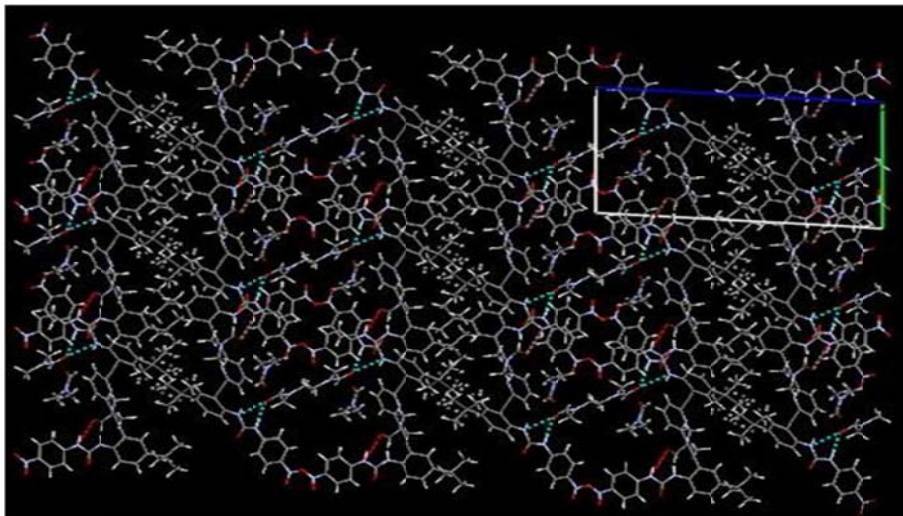
D-H···A	d(D-H)	d(H···A)	d(D···A)	<(DHA)
N(2)-H(2N)···O(5)	0.76(3)	2.35(4)	3.039(5)	150(3)
N(3)-H(3N)···O(5)	0.81(3)	2.04(4)	2.828(5)	165(3)

N(4)-H(4N)···O(1S)#1	0.74(4)	2.32(4)	2.950(4)	145(4)
N(5)-H(5N)···O(1S)#1	0.80(4)	2.02(4)	2.795(5)	163(4)
O(5)-H(5A)···N(1)	0.84(5)	2.04(5)	2.871(5)	172(5)
O(5)-H(5B)···O(3)#2	0.73(5)	2.07(6)	2.774(5)	163(6)

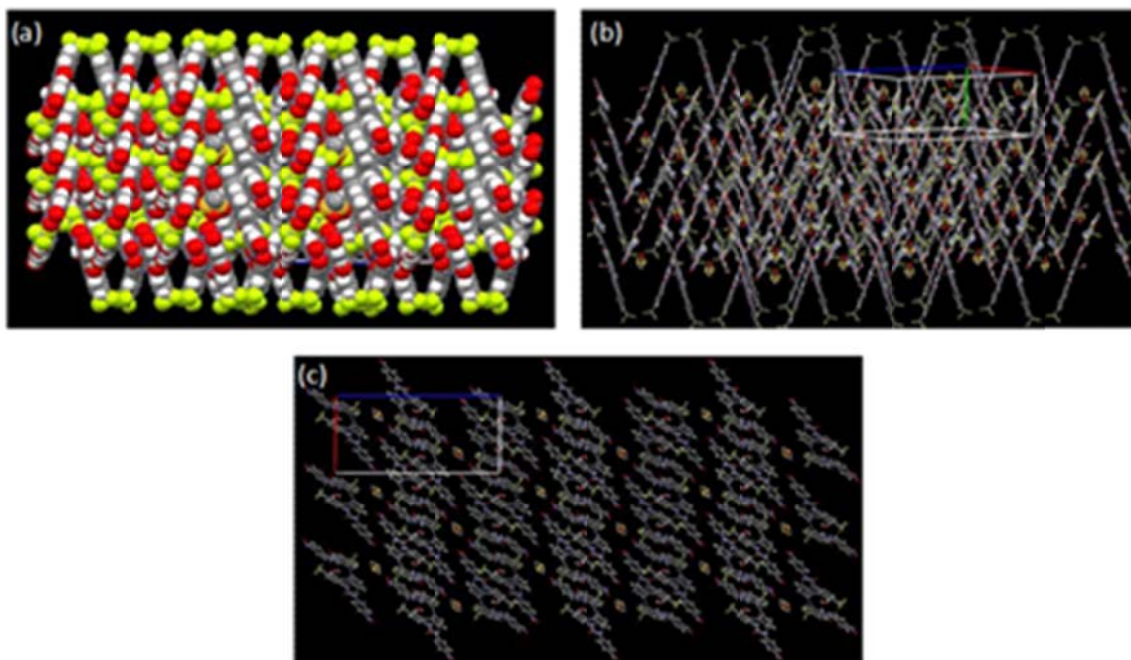
Symmetry transformations used to generate equivalent atoms:

#1 -x+1,-y+1,-z+1 #2 -x+1,-y,-z+1





**Figure S1.** General packing diagram of  $L^1 \cdot 3DMF$  viewed down the crystallographic  $a$ -axis showing the various hydrogen-bonding interactions.



**Figure S2.** (a) Packing model of  $L^2 \cdot H_2O \cdot 0.5DMSO$  along the pseudo  $c$ -axis as a space-filling model (solvent water and DMSO molecules are omitted for clarity). (b) Packing model of  $L^2 \cdot H_2O \cdot 0.5DMSO$  and solvents along the pseudo  $c$ -axis and (c)  $b$ -axis.