

## Supporting Information

# A Crystal Engineering Rationale in Designing A Cd<sup>II</sup> Coordination Polymer Based Metallogel Derived from a C<sub>3</sub> Symmetric Tris-amide-tris-carboxylate Ligand

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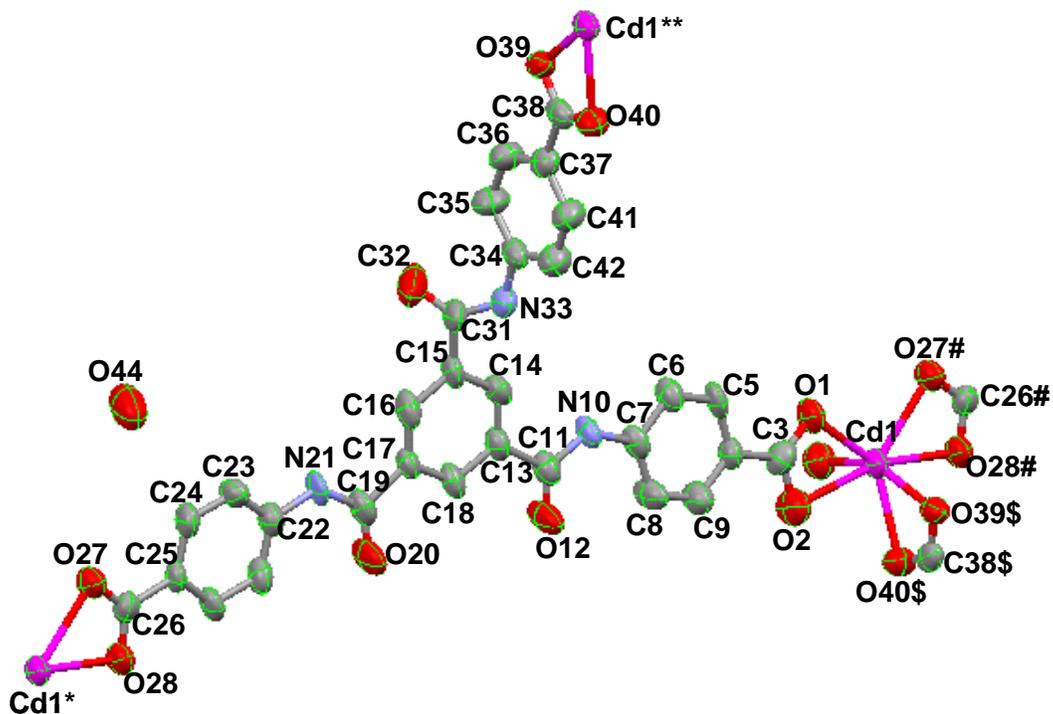
[ocpd@iacs.res.in](mailto:ocpd@iacs.res.in); [parthod123@rediffmail.com](mailto:parthod123@rediffmail.com)

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**Molecular plot and Hydrogen Bonding Parameters for CP1 (Figure S1)**

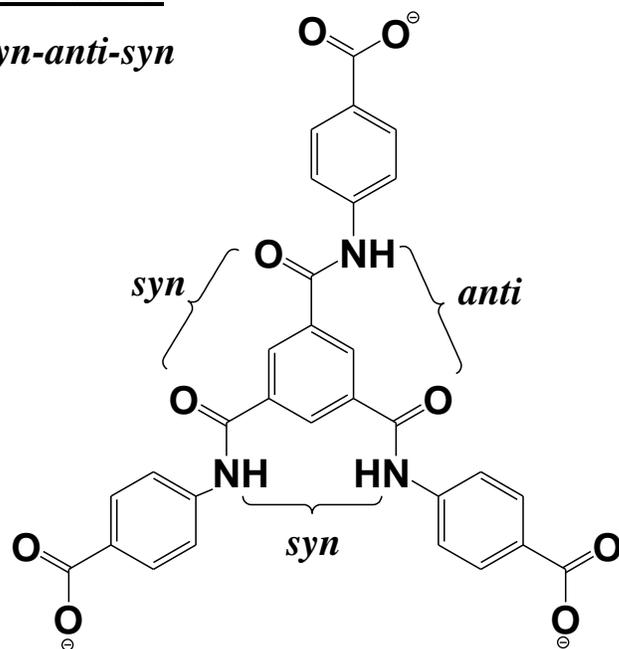


\* 1+x, y, 2+z; \*\* -x, -1/2+y, 1/2-z ; # -1+x, y, -2+z ; \$ -x, 1/2+y, 1/2-z

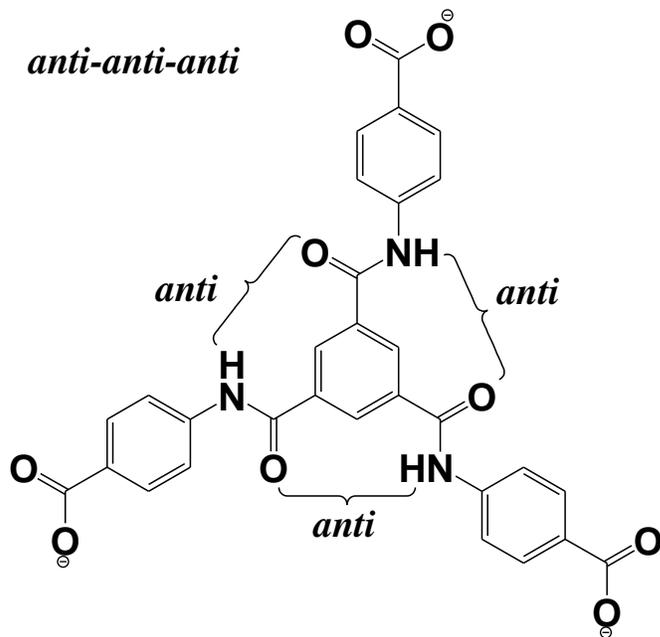
Hydrogen bonding parameters of CP1					
D-H...A	D-H (Å)	H...A (Å)	D...A (Å)	D-H...A (°)	Symmetry operation for A
N(10)-H(10)...O(39)	0.86	2.13	2.929(6)	154.1	x, -y+3/2, z-1/2
N(33)-H(33)...O(28)	0.86	2.07	2.881(6)	157.1	-x+1, -y+2, -z+2
O(44)...O(43)			2.745(7)		
O(44)...O(27)			2.744(7)		
O(43)...O(1)			2.618		

**Various conformations of L1**

*syn-anti-syn*



*anti-anti-anti*



## Scheme S1

S2

### TGA of CP1

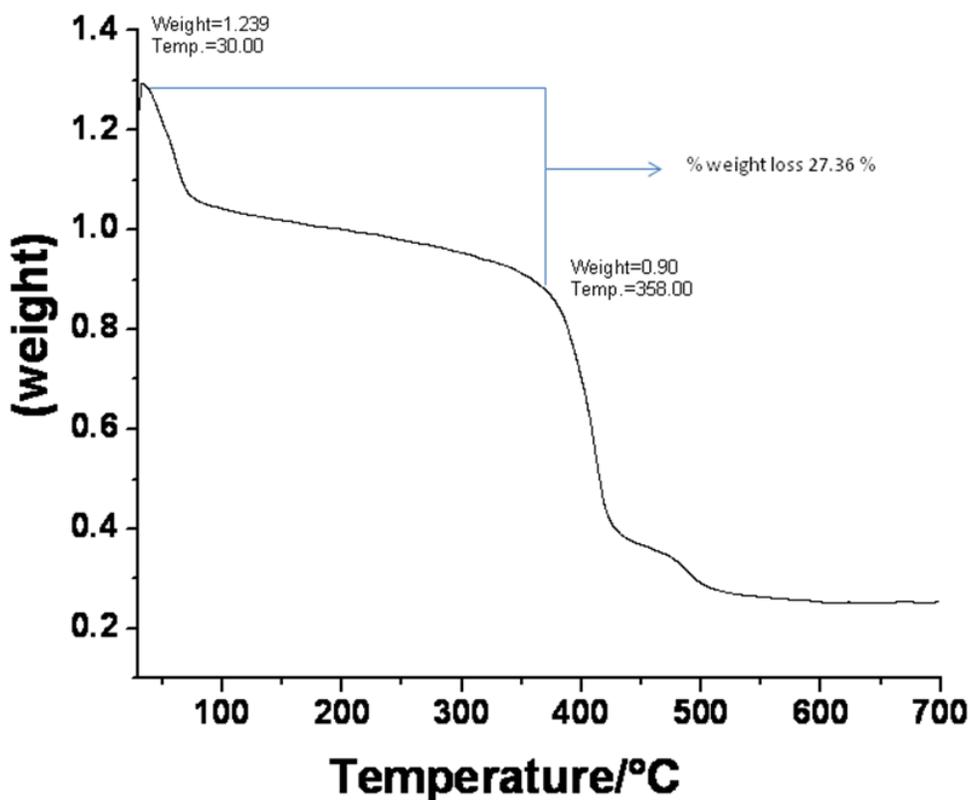


Figure. S2

Unit cell contents = 4 Ligand(L1) + 4 Cd<sup>2+</sup> + 4 H<sub>2</sub>O (occluded) + 4 H<sub>2</sub>O (coordinated) + 48 disordered H<sub>2</sub>O (488 electrons from SQUEEZE results approximately attributed to 12 H<sub>2</sub>O molecule)

Monoclinic, P2<sub>1</sub>/c space group, Z = 4

Therefore FW = Unitcell contents/4

$$\begin{aligned} &= 1\text{Ligand(L1)} + 1\text{ Cd}^{2+} + 1\text{ H}_2\text{O (coordinated)} + 1\text{(occluded)} + 12\text{ H}_2\text{O (occluded)} \\ &= 1 \times 564.11 + 113.9 + 18.01 + 18.01 + 12 \times 18.01 = 929.11 \end{aligned}$$

Weight loss for 13H<sub>2</sub>O = 14 X 18.01/929.11

$$\begin{aligned} &= 252/912.11 \\ &= 27.12 \% \text{ (experimental 27.36 \%)} \end{aligned}$$

S3

### PXRD comparison between bulk and xerogel state of G2,G3, G4

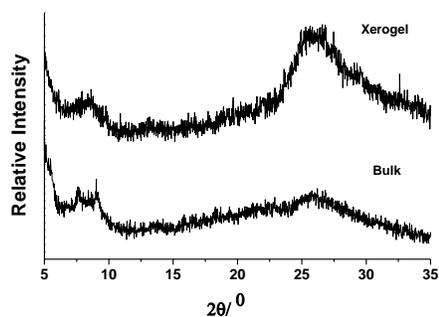


Figure S3 : PXRD comparison between bulk (CP2) and xerogel of G2

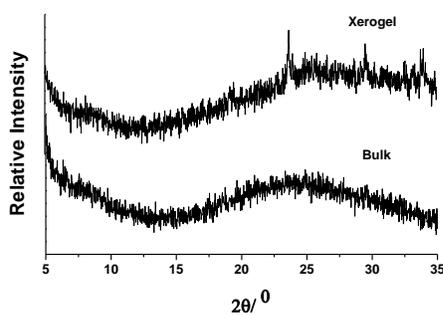
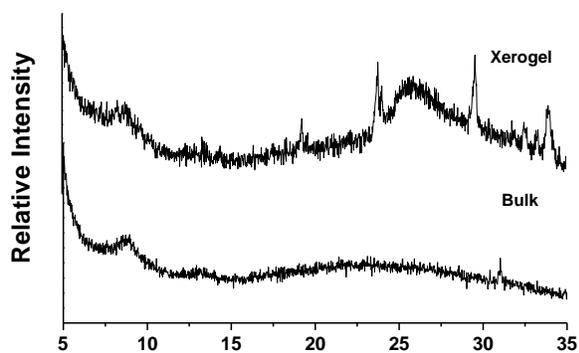


Figure S4 : PXRD comparison between bulk (CP3) and xerogel of G3



5

Figure **S5** : PXRD comparison between bulk (**CP4**) and xerogel of G4

**S4**

## checkCIF/PLATON report

You have not supplied any structure factors. As a result the full set of tests cannot be run.

No syntax errors found.    CIF dictionary    Interpreting this report

### Datablock: pdann

---

Bond precision:    C-C = 0.0096 Å                      Wavelength=0.71073

Cell:                      a=17.9224 (13)              b=26.390 (2)              c=9.0841 (7)  
                            alpha=90                      beta=99.700 (2)              gamma=90

Temperature:              298 K

	Calculated	Reported
Volume	4235.1 (6)	4235.1 (6)
Space group	P 21/c	P2 (1)/c
Hall group	-P 2ybc	?
Moiety formula	C30 H18 Cd N3 O10, O	?
Sum formula	C30 H18 Cd N3 O11	C30 H46 Cd N3 O23
Mr	708.88	929.10
Dx, g cm-3	1.112	1.457
Z	4	4
Mu (mm-1)	0.562	0.601
F000	1420.0	1916.0
F000'	1417.52	
h,k,lmax	17,25,8	16,25,8
Nref	3819	3803
Tmin,Tmax	0.829,0.919	0.831,0.921
Tmin'	0.825	

Correction method= MULTI-SCAN

Data completeness= 0.996                      Theta(max)= 19.750

R(reflections)= 0.0437 ( 3109)              wR2(reflections)= 0.1023 ( 3803)

S = 1.107                      Npar= 406

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The following ALERTS were generated. Each ALERT has the format  
**test-name\_ALERT\_alert-type\_alert-level**.  
Click on the hyperlinks for more details of the test.

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#### Alert level A

CHEMW03\_ALERT\_2\_A ALERT: The ratio of given/expected molecular weight as  
calculated from the \_atom\_site\* data lies outside  
the range 0.90 <> 1.10  
From the CIF: \_cell\_formula\_units\_Z                      4

```
From the CIF: _chemical_formula_weight          929.10
TEST: Calculate formula_weight from _atom_site_*
atom      mass      num      sum
C         12.01     30.00    360.33
H          1.01     18.00     18.14
N         14.01      3.00     42.02
O         16.00     11.00    175.99
Cd        112.41      1.00    112.41
Calculated formula weight                      708.89
```

**Author Response:** We have used SQUEEZE to remove the contributions of disordered solvent molecules (which we could not be modeled) within the crystal lattice. Therefore, the contribution of disordered solvent molecules removed by the SQUEEZE process are included in the overall formula, formula weight, density, F(000) etc. The output of the PLATON (generated as .sqf file) is also appended to the bottom of the CIF.

```
THETM01_ALERT_3_A The value of sine(theta_max)/wavelength is less than 0.550
Calculated sin(theta_max)/wavelength =      0.4755
```

**Author Response:** The crystal was poorly diffracting at higher Bragg angles. The situation did not improve even after repeating crystallization and fresh data collection.

```
PLAT043_ALERT_1_A Check Reported Molecular Weight .....          929.10
```

**Author Response:** We have used SQUEEZE to remove the contributions of disordered solvent molecules (which we could not be modeled) within the crystal lattice. Therefore, the contribution of disordered solvent molecules removed by the SQUEEZE process are included in the overall formula, formula weight, density, F(000) etc. The output of the PLATON (generated as \ .sqf file) is also appended to the bottom of the CIF.

```
PLAT306_ALERT_2_A Isolated Oxygen Atom (H-atoms Missing ?) .....          044
```

**Author Response:** The hydrogen atoms of the lattice included water molecules could not be located from difference Fourier map.

---

**Alert level B**

```
PLAT051_ALERT_1_B Mu(calc) and Mu(CIF) Ratio Differs from 1.0 by .          6.45 Perc.
```

**Author Response:** We have used SQUEEZE to remove the contributions of disordered solvent molecules (which we could not be modeled) within the crystal lattice. Therefore, the contribution of disordered solvent molecules removed by the SQUEEZE process are included in the overall formula, formula weight, density, F(000) etc. The output of the PLATON (generated as .sqf file) is also appended to the bottom of the CIF.

PLAT430\_ALERT\_2\_B Short Inter D...A Contact O27 .. O44 .. 2.74 Ang.

**Author Response:** The short contact is due to the hydrogen bonding between the lattice included water and carboxylate. Since the hydrogen atom associated with the lattice included water could not located in the difference fourier map, it shows as short contact alert.

PLAT430\_ALERT\_2\_B Short Inter D...A Contact O43 .. O44 .. 2.75 Ang.

**Author Response:** The short contact is due to the hydrogen bonding between the lattice included water and carboxylate. Since the hydrogen atom associated with the lattice included water could not located in the difference fourier map, it shows as short contact alert.

● Alert level C

REFNR01\_ALERT\_3\_C Ratio of reflections to parameters is < 10 for a centrosymmetric structure  
sine(theta)/lambda 0.4755  
Proportion of unique data used 1.0000  
Ratio reflections to parameters 9.3670  
PLAT041\_ALERT\_1\_C Calc. and Reported SumFormula Strings Differ ?  
PLAT068\_ALERT\_1\_C Reported F000 Differs from Calcd (or Missing)... ?  
PLAT088\_ALERT\_3\_C Poor Data / Parameter Ratio ..... 9.37  
PLAT230\_ALERT\_2\_C Hirshfeld Test Diff for O1 -- C3 .. 5.3 su  
PLAT230\_ALERT\_2\_C Hirshfeld Test Diff for C4 -- C5 .. 5.5 su  
PLAT230\_ALERT\_2\_C Hirshfeld Test Diff for C16 -- C17 .. 5.2 su  
PLAT241\_ALERT\_2\_C Check High Ueq as Compared to Neighbors for O2  
PLAT241\_ALERT\_2\_C Check High Ueq as Compared to Neighbors for C35  
PLAT242\_ALERT\_2\_C Check Low Ueq as Compared to Neighbors for Cd1  
PLAT250\_ALERT\_2\_C Large U3/U1 Ratio for Average U(i,j) Tensor .... 2.2  
PLAT334\_ALERT\_2\_C Small Average Benzene C-C Dist. C34 -C42 1.37 Ang.  
PLAT342\_ALERT\_3\_C Low Bond Precision on C-C Bonds ..... 0.0096 Ang  
PLAT369\_ALERT\_2\_C Long C(sp2)-C(sp2) Bond C11 - C13 ... 1.53 Ang.  
PLAT414\_ALERT\_2\_C Short Intra D-H...H-X H10 .. H14 .. 1.96 Ang.  
PLAT414\_ALERT\_2\_C Short Intra D-H...H-X H16 .. H21 .. 1.93 Ang.  
PLAT420\_ALERT\_2\_C D-H Without Acceptor N21 - H21 ... ?

● Alert level G

FORMU01\_ALERT\_2\_G There is a discrepancy between the atom counts in the chemical formula sum and the formula from the atom\_site\* data.  
Atom count from \_chemical\_formula\_sum: C30 H46 Cd1 N3 O23  
Atom count from the \_atom\_site data: C30 H18 Cd1 N3 O11  
CELLZ01\_ALERT\_1\_G Difference between formula and atom\_site contents detected.  
CELLZ01\_ALERT\_1\_G ALERT: Large difference may be due to a symmetry error - see SYMMG tests  
From the CIF: \_cell\_formula\_units\_Z 4  
From the CIF: \_chemical\_formula\_sum C30 H46 Cd N3 O23  
TEST: Compare cell contents of formula and atom\_site data

atom	Z*formula	cif sites	diff
C	120.00	120.00	0.00
H	184.00	72.00	112.00
Cd	4.00	4.00	0.00
N	12.00	12.00	0.00

```
          O          92.00    44.00    48.00
PLAT004_ALERT_5_G Info: Polymeric Structure Found with Dimension .      2
PLAT005_ALERT_5_G No _iucr_refine_instructions_details in CIF ....      ?
PLAT007_ALERT_5_G Note: Number of Unrefined D-H Atoms .....            3
PLAT194_ALERT_1_G Missing _cell_measurement_reflms used datum ....      ?
PLAT195_ALERT_1_G Missing _cell_measurement_theta_max datum ....        ?
PLAT196_ALERT_1_G Missing _cell_measurement_theta_min datum ....        ?
PLAT606_ALERT_4_G VERY LARGE Solvent Accessible VOID(S) in Structure    !
PLAT764_ALERT_4_G Overcomplete CIF Bond List Detected (Rep/Expd) .      1.13 Ratio
PLAT804_ALERT_5_G ARU-Pack Problem in PLATON Analysis .....            6 Times
PLAT869_ALERT_4_G ALERTS Related to the use of SQUEEZE Suppressed      !
```

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4 **ALERT level A** = Most likely a serious problem - resolve or explain  
3 **ALERT level B** = A potentially serious problem, consider carefully  
17 **ALERT level C** = Check. Ensure it is not caused by an omission or oversight  
13 **ALERT level G** = General information/check it is not something unexpected

9 ALERT type 1 CIF construction/syntax error, inconsistent or missing data  
17 ALERT type 2 Indicator that the structure model may be wrong or deficient  
4 ALERT type 3 Indicator that the structure quality may be low  
3 ALERT type 4 Improvement, methodology, query or suggestion  
4 ALERT type 5 Informative message, check

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It is advisable to attempt to resolve as many as possible of the alerts in all categories. Often the minor alerts point to easily fixed oversights, errors and omissions in your CIF or refinement strategy, so attention to these fine details can be worthwhile. In order to resolve some of the more serious problems it may be necessary to carry out additional measurements or structure refinements. However, the purpose of your study may justify the reported deviations and the more serious of these should normally be commented upon in the discussion or experimental section of a paper or in the "special\_details" fields of the CIF. checkCIF was carefully designed to identify outliers and unusual parameters, but every test has its limitations and alerts that are not important in a particular case may appear. Conversely, the absence of alerts does not guarantee there are no aspects of the results needing attention. It is up to the individual to critically assess their own results and, if necessary, seek expert advice.

#### Publication of your CIF in IUCr journals

A basic structural check has been run on your CIF. These basic checks will be run on all CIFs submitted for publication in IUCr journals (*Acta Crystallographica*, *Journal of Applied Crystallography*, *Journal of Synchrotron Radiation*); however, if you intend to submit to *Acta Crystallographica Section C* or *E*, you should make sure that full publication checks are run on the final version of your CIF prior to submission.

#### Publication of your CIF in other journals

Please refer to the *Notes for Authors* of the relevant journal for any special instructions relating to CIF submission.

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PLATON version of 19/04/2012; check.def file version of 14/04/2012

Datablock pdann - ellipsoid plot

