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1	A combined experimental and theoretical insight into				
2	the drug delivery of nanoporous metal-organic				
3	frameworks				
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15	Abstract				
16					
17	Two isostructural nanoporous MOFs with $[Zn_3(\mu_3-O)(BTC)_2(H_3O)]_n$ (NTU-Z11)				
18	and $\{[Zn_3(\mu_3-O)(BTC)_2(DMF)] \cdot 2NH_2(CH_3)_2 \cdot 4H_2O\}_n$ (GDMU) (BTC)				
19	=1,3,5-benzenetricarboxylate) have been used as drug carriers of 5-fluorouracil				
20	(5-FU). The incorporation of the 5-FU into the desolvated NTU-Z11 and GDMU was				
21	around 0.38 g/g and 0.22 g/g, respectively. NTU-Z11 presents a pH-triggered				
22	controlled drug release property in 6.0, 7.4, 9.18 and water media. In addition, we				
23	performed GCMC simulations to investigate the loading of 5-FU to NTU-Z11 and				
24	GDMU at the molecular level. The results from simulations reproduce the				
25	experimental trend with respect to drug loading capacity of each material.				
26	Comparison between calculated drug loading values and some molecular level				
27	properties indicates the existence of a relationship between the void space of material				
28	and drug loading capacity.				
29	Introduction				

30 Porous metal-organic frameworks (MOFs) have particularly highlighted for their

excellent gas-storage and catalysis properties [1-3]. Recently, tremendous efforts on MOF carriers have been made to boost their way toward medical applications [5-8]. Férey's group first described the potential loading and release properties of some drugs on MOFs[9], whereas Lin *et al.* have constructed a Pt-based drug at the nanoscale by using it as one building block to create a new coordination polymer[10]. Horcajada and his co-workers had also reported that porous MOFs can load and release drugs, acting as a promising non-toxic drug carrier [11].

8 Zhang and his co-worker reported a facile route to synthesize a series of 9 NTU-based MOFs [12]. NTU-Z11 is the isostructure of MOF-38 and can be repeatedly synthesized with high yield [13]; moreover, its channels are empty and 10 have a dimension of about 11.5×11.5 Å. Inspired by these works, our strategy is to 11 12 explore a neutral MOF that its structural feature is similar with NTU-Z11. 13 Unfortunately, only a negative **GDMU** was obtained. But we are still interested to 14 develop the loading and release properties of 5-FU on the two MOFs because the 15 efficiency of drug delivery is related to the pore characteristics and the nature of 16 host-guest interactions. GCMC simulation is a powerful technique to explain and 17 predicate the gas adsorption to porous materials. However, there is still a challenging 18 work to use the GCMC simulations to investigate the loading of large molecules to 19 porous materials due to the requirement of the conformational sampling and fitting of 20 such molecules inside tight pores [14-16].

21 Herein, we demonstrated two Zn(II)-based frameworks with additional negative 22 charges that have been used as drug carriers of 5-FU. The incorporation of the 5-FU 23 into the desolvated NTU-Z11 and GDMU was around 0.38 g/g and 0.22 g/g, 24 respectively. **NTU-Z11** presents a pH-triggered controlled drug release property in pH 25 6.0, 7.4, 9.18 and water media. In addition, we performed GCMC simulations to 26 investigate the loading of 5-FU in NTU-Z11 and GDMU at the molecular level. 27 Comparison between calculated drug loading values and some molecular level 28 properties indicates the existence of an important relationship between the void space 29 of material and drug loading capacity.

30 Materials and Method

All reagents were purchased from commercial sources and used as received. IR spectra were recorded with a Perkin–Elmer Spectrum One spectrometer in the region 4000-400cm⁻¹ using KBr pellets. TGA were carried out with a Metter–Toledo TA 50 under dry dinitrogen flux (60mL.min⁻¹) at a heating rate of 5°C min⁻¹. X-ray powder diffraction (PXRD) data were recorded on a Rigaku RU200 diffractometer at 60KV, 300mA for *Cu K_a* radiation ($\lambda = 1.5406$ Å), with a scan speed of 2 °C/min and a step size of 0.02° in 20.

8 X-ray Crystallography: Single crystal X-ray diffraction analyses of the two 9 compounds were carried out on a Bruker SMART APEX II CCDdiffractometer equipped with a graphite monochromated MoK α radiation ($\lambda = 0.71073$ Å) by using 10 ϕ/ψ scan technique at room temperature. The intensities were corrected for Lorentz 11 12 and polarization effects as well as for empirical absorption based on multi-scan 13 techniques; all structures were solved by direct methods and refined by full-matrix least-squares fitting on F^2 by SHELX-97[17]. Absorption corrections were applied by 14 15 using multi-scan program SADABS[18]. Non-hydrogen atoms were refined 16 anisotropically. The structure contains large (37 % volume) regions of intensely 17 disordered cations and solvent. These were impossible to model at atomic resolution 18 and their presence in the structure is assumed on the basis of the elemental analysis, 19 TGA and the PLATON/SQUEEZE calculations [19]. The latter were used to calculate 20 the diffraction contribution of the solvent molecules and, thereby, to produce a set of 21 solvent-free diffraction intensities for the refinement of the MOF structure. Crystallographic data for complexes **GDMU** are given in Table 1. Selected bond 22 23 distances and bond angles are listed in Table 2. CCDC: 1405443 for GDMU.

24 Syntheses of these complexes

25 $[Zn_3(\mu_3-O)(BTC)_2(H_3O)]_n$ (NTU-Z11)

We only synthesized the **NTU-Z11** according to the reference. The sample purity was confirmed by the PXRD.

28 $\{[Zn_3(\mu_3-O)(BTC)_2(DMF)] \cdot 2NH_2(CH_3)_2 \cdot 4H_2O\}_n (GDMU)$

29 A mixture of $Zn(NO_3)_2 \cdot 6H_2O$ (0.450g, 0.1mmol), L(4,4'-bis(pyrid-4-yl)biphenyl) 30 (0.015g, 0.04mmol), and H₃BTC (0.450mg, 0.2mmol), DMF (4mL) in a

screw-capped vial. After five drops of HNO₃ was added into the mixture. The vial was capped and placed in an oven at 110 °C for 3 days. The resulting colorless single crystals were washed with absolute CH₃CH₂OH three times to give **1**. Anal. Calcd for $C_{25}H_{37}N_3O_{18}Zn_3$ (863.68), C, 34.77; H, 4.32; N, 4.87. Found C, 34.28.; H, 4.15; N,4.55. IR (KBr, cm⁻¹) : 3480(vs); 2940(m); 1632(vs); 1428(v); 1390(v); 1099(m); 938(m); 708(v); 547(m).

7 **Computational Details**

8 The 5-FU adsorption in NTU-11 and GDMU was studied using grand canonical 9 ensemble Monte Carlo (GCMC) simulations, employed with the RASPA code at 298 10 K[20]. The structures of 5-FU and MOFs are described with an all-atom model in this 11 work. The structures for 5-FU and MOFs can be found in Figures S1 and S2. For 12 MOFs structures, the framework atoms were kept rigid during the simulations. The 13 guest-guest and guest-host interactions were computed with a Lennard-Jones (LJ) and 14 Coulombic potential. The Antechamber program of AmberTools1.27 was used to 15 generate the force field for 5-FU with the general amber force field parameters [21]. 16 The atomic partial charges for 5-FU were computed with CHELPG method based on the Gaussian 03 suite with the 6-31++g* basis set. The Lennard-Jones parameters and 17 18 partial charges can be found in Table S1-S3.

19 The atomic positions of NTU-Z11 and GDMU structures were taken from the 20 PXRD data(The Rietveld refinement for the 5-FU@MOFs complexes was performed 21 with the software GSAS/EXPGUI, using the X-ray structure of the MOF as initial atomic coordinates.). The cations of H_3O^+ and $NH_2(CH_3)_2^+$ are included in NTU-Z11 22 and in **GDMU**, respectively. The cations of $NH_2(CH_3)_2^+$ were not removed in the 23 uptake of 5-FU. Thus we also did not remove these molecules in these simulated 24 25 structures. The Lennard-Jones parameters for the MOFs structure atoms were taken 26 from the UFF force field (listed in Table S2) [22]. The accurate prediction of 27 adsorption in various MOFs could be achieved by a number of simulation 28 investigations using the UFF force field [23-24]. The molecular geometries for cations 29 were optimized by DFT method. In canonical ensemble, the desired number of cation were inserted into the pore and attempted to accelerate the equilibrium with 30

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- reinsertion-move. The obtained configurations were used to simulate the adsorption of
 5-FU in MOFs. The solvent molecules of the MOFs in the simulations are allowed to
 move (including translation and rotation).
- 4 The heats of adsorption were computed using the equation:

$$Q_{st} = RT - \frac{\langle UN \rangle - \langle U \rangle \langle N \rangle}{\langle N^2 \rangle - \langle N \rangle^2}$$

5

where < > refers to the average over the simulation, and U is the energy, N is the
number of adsorbed molecules.

8 For the interactions of unlike sites were computed with Lorentz-Berthelot mixing 9 rules. The Lennard-Jones interactions were cut and shifted at the 13 Å. The partial 10 charges of the NTU-Z11 and GDMU atoms were computed from density functional 11 theory (DFT) with the B3LYP functional. For the metal atoms, the LanL2DZ basis set 12 was applied. The 6-31++g* basis set was used to optimized for all other atoms. The 13 atomic partial charges can be obtained by fitting the electrostatic potentials after DFT 14 computation. The Coulombic interactions were computed using the Ewald sum technique. The details of simulated boxes are listed in Table S4. After the initial 10^6 15 Monte Carlo (MC) cycles, the production of 10^6 cycles was used to compute the 16 ensemble averages properties. For each cycle, the MC moves include the molecule of 17 18 insertion, deletion, translation, rotation or re-growth. We used the equal probability 19 for each MC moves.

20 Results and Discussion

21 $[Zn_3(\mu_3-O)(BTC)_2(H_3O)]_n$ (NTU-Z11) and $\{[Zn_3(\mu_3-O)(BTC)_2(DMF)] \cdot 2NH_2(CH_3)_2 \cdot 4H_2O\}_n$ 22 (GDMU)

The NTU-Z11 and GDMU are isostructural, which are composed of the [Zn₃(μ_3 -O)(COO)₆] subunits (Fig. 1a-1b). The subunits are connected by BTC ligands, which results in an infinite 3-D (3,6)- connected framework with 1-D channel of about 11.5 × 11.5 Å dimension along the *c*-axis (Figure 1c-1d). But we should state herein, if the L was absent in this reactive system, the final product of GDMU could not be obtained. Furthermore, the pores of GDMU were occupied by the NH₂(CH₃)₂

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and DMF molecules. This structural feature was also similar with MOF-38, which
holds some disordered HTEA molecules. However, the MOF-38 cannot be repeated
as mentioned in the literature[13].

4 Thermogravimetric Analyses

5 The thermogravimetric analyses (TGA) of complex **GDMU** was performed (Fig. S3). It shows three weight loss steps. The first weight loss begins at 25°C and is 6 7 completed at 80°C. The observed weight loss of 8.6% is corresponding to the loss of 8 the free water molecules (calcd 8.3%). The second weight loss occurs latterly, and can 9 be attributed to the elimination of $NH_2(CH_3)_2$ cations(obsd: 9.5%; calcd 10.4%). A gradual weight loss from 210 °C indicates that the complex decomposes continuously 10 when the temperature is raised. The mass remnant at ~700 °C of 25.4 % is roughly 11 consistent with the deposition of ZnO (calcd 28.3%) (a weight loss of 4.0% is larger 12 13 than the calculated value, probably resulting from the sensitivity to temperature and 14 humidity or a very slow absorbability of the guest molecules from the air at room 15 temperature).

16 Both of NTU-Z11 and GDMU were desolvated at 120 °C for 10 h prior to 17 insertion of the drug. As confirmed by PXRD and TGA, 5-FU containing sample 18 maintains its crystallinity (Fig. S3 and Fig. S4), thus, the drug encapsulation did not 19 alter the structure of these materials. Only a decrease in the intensity of the low angle reflections on the PXRD patterns (\sim 5-8° 2 θ) was observed after encapsulation, 20 following the change in pore content that is known to strongly affect the relative 21 intensities of the Bragg peaks^{2b}. This was confirmed by N₂ adsorption analyses 22 23 showing that the BET surface area significantly decease upon drug molecules loading 24 (see Supplementary Information Fig. S5).

Incorporation of the drug molecule during loading process has been recorded by Fourier transformed infrared spectroscopy (FTIR) (Figure S6). The absorption bands of C–F deformations were discovered in the 820–550 cm⁻¹ regions. The absorption band at about 1240 cm⁻¹ may be due to fluorine atom on the ring [25-26]. Based on the above structural analyses, these two compounds may be taken as a good drug carrier. The loading of anticancer 5-FU was carried out by impregnating **NTU-Z11** 2

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UV-vis absorption spectroscopy has been used to determine the effective storage

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1 and **GDMU** under stirring in 5-FU containing ethanol solutions.

capacity, To reach a maximal drug loading, 5-FU to porous solid relative ratio and 3 contact time were evaluated (Table S5)[6]. The loading amount of 5-FU increased 4 5 with initial 5-FU/material ratio repressed in weight and optimal value 1:1 and 1:3 for NTU-Z11 and GDMU in ethanol, respectively. The contact time was also important, 6 7 the maximum adsorption was obtained after 2 days and 3 days for NTU-Z11 and GDMU, respectively. Thus, the best results were obtained when NTU-Z11 was 8 9 soaked for 2 days within a 5-FU to material weight ratio of 1:1, while GDMU was 10 soaked for 3 days within a 5-FU to material weight ratio of 1:3. 5-FU was incorporated into desolvated NTU-Z11 and GDMU with loadings of 0.382 and 0.206 11 12 g/g, respectively. The difference between NTU-Z11 and GDMU shows that the 13 $NH_2(CH_3)_2$ takes as gate and blocks the drug molecules access to inner pores [27].

14 Fig. 2 shows the release profile of the drug delivery system of NTU-Z11 and 15 GDMU in PBS solution at 37 °C. At the first stage (24 hours), the NTU-Z11 and 16 **GDMU** have the similar releasing behavior and approximately 65 % of the drug was 17 released. However, the other part released gradually in GDMU, implying a strong 18 host-guest interaction involved in this process. Compared with the NTU-Z11 carrier, 19 there is a big cation in the host channels in **GDMU**, which can take as donor/acceptor 20 and bind to drug molecules resulting in the dramatically releasing behavior. Thus, 21 5-FU with flat molecular shape diffuses along the hexagonal channels. Similar results were also found in MIL-53 with a pore size of 8.6 Å exhibited a drug loading capacity 22 23 of 0.22 g/g for drug IBU (IBU = ibuprofen) [3].

24 To further explore the pH-responsive drug release feature of NTU-Z11, release 25 profile were performed in pH 6.0, 9.18 and water medium. Around 62.5% of the 26 loaded 5-FU was released fast within 24 h, and 63.1% within 30 h. More than 40% of 27 5-FU released around one hour, which consistent with dissolution of NTU-Z11 in 28 acidic environment. Compared with other MOF carriers[28], NTU-Z11 shows a fast 29 release rate for 5-FU. In the water medium, the released profile of 5-FU exhibits a flat shape and occurs no burst effect. The delivery of 5-FU occurred within 96 h and 47 % 30

7

of the loaded drug was released. However, three stages related to the drug release could be distinguished in pH 9.18, around 48% of the loaded drug was released in the first stage (33 h) and only almost not more than 10% of the loaded drug was released. Thus, a rapid releasing process was observed during the first stage followed by a slower in the high pH. These results imply that the loaded drug can be decreased during blood circulation and the drug release rate is suddenly accelerated after release into cancer cells [25, 29].

8 Computational Simulations of 5-FU Adsorption

9 The amount of drug per porous material or drug loading is one of the main 10 quantities of interest in the use of MOFs for controlled drug release [30]. We have 11 used GCMC simulations to investigate the loading of 5-FU to two compounds at the 12 molecular level. These simulations were used to determine the preferential binding 13 sites of the 5-FU in the porous materials, to estimate the maximum drug loading 14 capacity of each material, and propose a molecular mechanism for drug loading and 15 release.

16 Adsorption isotherm of 5-FU in MOFs

We calculated the adsorption isotherms of 5-FU in NTU-Z11 and GDMU 298 K. 17 18 As observed in Figure 3, there are some differences between NTU-Z11 and GDMU. 19 The NTU-Z11 has much higher saturation capacity for 5-FU, which is about 0.4 g/g. 20 The saturation capacity is around 0.22 g/g for **GDMU**. The bigger molecules of 21 **GDMU** result in a low saturation capacity compared to that of **NTU-Z11**. Also, 22 **GDMU** show a saturation uptake at the low fugacity range due to the stronger 23 5-FU-MOF interactions. The presence of stronger interaction due to the existence of 24 cations in **GDMU** strengthens the host-guest interactions and results in the steep 25 adsorption of 5-FU at lower fugacity than in NTU-Z11.

26 Heat of adsorption

The heats of adsorption (Q_{st}) for 5-FU in **NTU-Z11** and **GDMU** studied are shown in Fig. 4. The heat of adsorption is closely related with the pore structure, in which could be taken as an index of the adsorption materials heterogeneity [30]. Fig. 4

shows the Qst for NTU-Z11 and GDMU as the function of uptake. As observed in 1 2 Fig.4, the NTU-Z11 shows the low Q_{st} (about 120-150 kJ/mol) at the loadings process. 3 This observation shows that the 5-FU molecules can load into MOFs pores with 4 strong interaction. The stronger interaction results in the higher adsorption heat of 5 5-FU than in NTU-Z11 because of the presence of DMF in GDMU. The GDMU shows the higher Qst (160-228 kJ/mol) at the range of loadings. The medicine 6 molecules can be strongly retained in the MOF structures due to the high Qst at the 7 loadings and it is very favorable for the long release process. **GDMU** shows higher 8 Q_{st} values than NTU-Z11 at high loadings, with the important contributions of the 9 10 solvent molecules. The results are consistent with experimental released process.

11 **Density plots**

12 NTU-Z11 consists of the trimetric SBU and BTC ligand. The 3-D framework has two channel systems with dimension of 7.5 \times 7.5 Å (refer to as A) and 11.5 \times 11.5 Å 13 14 (refer to as B) along the *c*-axis. As observed in Fig. 5, the 5-FU molecules are 15 primarily distributed in the two favorable regions. The 5-FU loading is closely packed 16 in the pores because the solvent molecules have a smaller size in A region. The bigger 17 cations present in GDMU hinder the adsorption of 5-FU in A region. Then the 18 adsorption of 5-FU in NTU-Z11 increases with the increase of fugacity. However, the 19 adsorption of 5-FU in **GDMU** rapidly approaches to a platform at same range of 20 pressure.

21 Conclusion

22 In summary, two isostructural nanoporous MOFs were used to load anti-cancer 23 chemotherapy drug 5-FU and demonstrated a remarkable different capacity due to 24 their various pore spaces. Owing to pH-sensitive property of NTU-Z11, it was 25 observed that it released much faster in mild acidic buffer solution than at a neutral 26 medium, suggesting that this pH-triggered feature may be useful property for drug 27 delivery to tumors. GCMC simulations suggested that the anti-cancer drug 5-FU 28 could load to the NTU-Z11 in high loading capacity. Our findings indicate that the 29 combined experimental-computational approach is a powerful strategy for the 30 efficient identification and incorporation of bioactive compounds in porous materials.

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43 density of 5-FU in **NTU-Z11** at 298 K (a) 10⁻⁹ mPa (b) 10⁻⁷ mPa (below).

2	F				-	
3	Crystal system		tetragonal			
4			I4cm			
5 6	Crystal color		Colorless			
7			20,5128(10)			
8	<i>u</i> , A		17.0100(0)			
9 10	с, А		17.8100(8)			
11	γ		90			
12	<i>V</i> , Å ³		7494.7(8)			
13 14	Z		8			
15	-	ρ_{calcd} , g/cm ³		1.531		
16	-	F(0.00)			526	
17	-	F(000)		3536		
18		θ Range, deg		2.49-27.92		
19 20	Reflns collected/unique(Rint)		21512/ 4425 (0.0299)			
21		GOF		1.092		
22				0.0310.0.0000		
23 24	-	$R_1, wR_2 (I > 2\sigma(I))^*$		0.0319, 0.0900		
24 25		R_1 , wR_2 (all data)**		0.0357, 0.0924		
26	-					
27	$R = \sum (F_{o} - F_{o})$	$F_{c})/\sum(F_{o}), ** wR_{2} =$	$= \{\sum [w(F_0^2 - F_c^2)^2]\}$	$\sum (F_0^2)^2 \}^{1/2}$		
28						
29	Table 2.	Selected bond di	stances (Å) and ar	ngles (deg)	of structure GE	OMU
30						
31	Zn1-01	1.945(4)	Zn1-	08	1.9458(17)	
32	Zn1- 04	1.969(3)	Zn1- (05	1.972(3)	
33	Zn2- 07	2.059(5)	Zn2- (03	2.091(3)	
34	Zn2- 03	2.091(3)	Zn2- (28	2.103(4)	
35	Zn2- 06	2.106(3)	Zn2- (06	2.106(3)	
36	01- Zn1 -08	113.76(18)	01- Zr	n1- 04	123.14(16)	
37	08- Zn1- 04	105.27(16)	01- Zr	n1- 05	103.53(16)
38	08 -Zn1- 05	103.27(17)	04- 2	Zn1- 05	105.86(18	5)

1	Table 1. Crystal	data and structure	refinement inform	nation for	compound	GDMU
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1	07 -Zn2- 03	86.24(14)	03- Zn2- 03	97.10(17)
2	07- Zn2 -08	171.1(2)	03 -Zn2 -08	87.85(12)
3	03- Zn2 -08	87.85(12)	07- Zn2- 06	91.29(15)
4	03 -Zn2- 06	173.17(14)	03- Zn2- 06	89.08(15)

5