Effect of electrostatic properties of IRMOFs on VOCs adsorption: a density functional theory study

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Abstract Metal-organic frameworks (MOFs) are of great promises for the adsorption of volatile organic compounds (VOCs). The adsorption of methanol, aldehyde and one on IRMOF-1 and an amino-functionalized framework, IR-MOF-NH₂, were investigated by density functional theory. The adsorption mechanisms and the effects of amino functionalization were clarified with comparison of binding energies, optimized configurations, atomic partial charges and the electrostatic potentials. The calculated results revealed that the electron-donating effect of amino groups had great influence on the electrostatic properties of the framework. The adsorption of methanol was greatly enhanced on organic linkers of IRMOF-NH2 compared to IRMOF-1, but weakened on the metal corners, different from the adsorption behaviors of aldehyde and acetone. A plausible interaction mechanism was inferred as the VOCs adsorption might be governed by the electrostatic interaction between methyl groups of the adsorbates and the binding sites of the framework, while the polar groups of adsorbates only interacted with framework through weak interaction like H-bonding. Besides, the steric hindrance effects of the inserted amino groups should also be taken into consideration, depending on the configuration and the electrostatic properties of the adsorbates. The binding mechanisms of VOCs on IRMOF-1 and IRMOF-NH₂, provide fundamental insights into the oriented design of MOFs for the adsorption of VOCs.

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1 Introduction

Volatile organic compounds (VOCs) are a set of major air pollutants in the upper troposphere, including methanol, aldehyde and ketone (Su et al. 2013) etc., which are not only considered as harmful compounds to health (Wolkoff 2013), but also induces some serious environmental problems, e.g. the photochemical smog, destruction of ozone layer (Laaksonen et al. 1997). Hence the development of novel technologies for VOCs treatment have triggered increasing attention to comply with the environmental regulations. Among various technologies for removing VOCs, adsorption has been generally considered as one of the most effective strategy as its industrial maturity and no chemical degradation (Dou et al. 2011). Hence, experimental and theoretical studies have increasingly focused on the adsorption of VOCs on various porous materials such as activated carbon (AC) (Xuemin et al. 2011), AC fiber (Le Cloirec 2012) and zeolite (Kim and Ahn 2012) etc., however their pore blocking and lack of regeneration ability of these materials prevent them from industrial application of adsorbing VOCs.

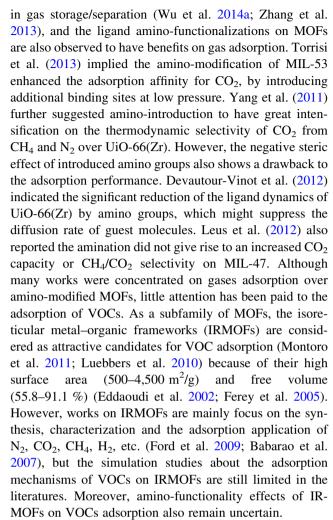
Metal-organic frameworks (MOFs) (Lin et al. 2012), as known as a new family of coordination polymers, are a set of regular crystalline solids with well-organized structures constructed by the coordination of metal corners with organic linkers. MOFs are known for their well-defined porosity, high surface area, adjustable pore geometry and chemical functionality (Furukawa et al. 2013), which are recognized as the potential alternatives in VOCs capture. Montoro et al. (2011) reported a novel MOF-5 type



framework to adsorb VOCs even in ambient moisture, and proposed the VOC-MOF interaction was dominated by pore size and surface hydrophobicity rather than the unsaturated metal sites. Luebbers et al. (2010) experimentally examined the adsorption of over 30 VOCs on IRMOF-1, indicating great intensification of the binding energies (BEs) by the structural degradation of the frameworks, but lack of thermodynamic interaction on organic linkers. Nevertheless, the direct observation of VOCs on the preferential adsorption sites of MOFs remains a challenging and time-consuming task, and the unclarified structure–property relationship of newly-constructed MOFs may impede their developments and applications for VOCs adsorption.

As a result, experimental studies are expected to be complemented by computational simulations. In the computational studies of gas adsorption, the positions of adsorbates are usually adjusted to find the most favorable conformation so as to investigate the preferential adsorption sites of MOFs as well as their performance for specific applications (i.e. VOCs adsorption/separations). In these cases, the refinements of adsorbate-adsorbent complexes are commonly obtained by geometric optimization, and the binding energy calculations are further carried out to clarify the interaction strength. Molecular dynamics (MD) simulation is known as a rapid method for geometric optimization. Since the electronic charges used for describing electrostatic interactions are usually kept static, the MD simulations is allowed to refine the configurations on very large MOFs systems [e.g. MIL-101 (Hu et al. 2013)]. However, the final quality of MD optimization significantly depends on the force fields adopted in the calculations. Since developing accurate and transferable forcefields are extremely challenging (Fang et al. 2014), general forcefields [e.g. UFF (Rappe et al. 1992), Dreiding (Mayo et al. 1990)] are still the most widely used in the recent studies, but they are less reliable to describe the more complex adsorption systems. In addition to MD simulation, density functional theory (DFT) method (Cicmanec et al. 2013; Yang et al. 2013) gives another way to theoretically obtain the geometric equilibrium of MOF-based adsorption systems. Contrary to MD simulations, DFT procedure is an electronic ground-state theory that defines the energy of atoms at prescribed positions from corresponding electron density (Young 2004). By using either localized or delocalized basis functions, DFT method allows great efficiency in geometric optimization. Although DFT approach is more time-consuming than MD simulations, it has been considered to be more capable of providing predictions with a satisfactory trade-off between accuracy and speed, and widely employed in studying both homogeneous (Liang et al. 2009) and heterogeneous (Liu et al. 2011) systems.

To date, DFT-based computational studies on various functionalities of MOFs have showed potential applications



In this paper, DFT calculations were carried out for investigating the adsorption mechanisms of methanol, aldehyde and acetone on IRMOF-1 and an amino-functionalized framework, IRMOF-NH₂. The selected VOCs with varied electrostatic properties were expected to give a distinction of adsorption behaviors. Geometric optimization was performed to search the binding sites of the frameworks, following by the BEs calculations of the selected configurations so as to indicate the interaction strength between adsorbate-adsorbent complexes. Finally, the effects of amino-functionalization on VOCs adsorption were clarified by comparing the BEs of different energyminimum configurations, and the electrostatic potentials (ESPs) of the parent and functionalized IRMOFs.

2 Models and computational details

2.1 Structures construction

IRMOF-1, also known as MOF-5 (Yaghi et al. 2003), has a *Fm-3m* crystal space group with a lattice constant of



25.832 Å, a crystal density of 0.593 g/cm³ and a free volume of 79.2 % (Eddaoudi et al. 2002). For simulation, the crystal structure of the framework was taken from experimental X-ray diffraction data (Eddaoudi et al. 2002), as shown in Fig. 1a. IRMOF-1 was a 3-dimensional (3D) porous cubic framework that constructed with eight metaloxide corners of Zn₄O connected by twelve 1,4-benzenedicarboxylate (BDC) organic linkers. Since the varied inclination angle of BDC linkers in the structure, the pores were divided into small and big cells, denoted as S and B in Fig. 1b, respectively. On the other hand, IRMOF-NH₂ was constructed on the basis of its parent IRMOF-1, with the H atoms of BDC organic linker symmetrically replaced by four -NH2 groups, which was expected to prevent the textural non-symmetry effects for adsorption interactions, as denoted in Fig. 2. This version of amino functionality was reported by Mu et al. (2010), suggesting the introduction of -NH2 groups could verify the pore size and electron-combining properties of IRMOFs, but retaining the topological structure of the framework unchanged.

2.2 Geometry optimization

The structures of the frameworks and the adsorption complexes were optimized in a two-step process. In the first step, the unit cells of the IRMOFs were optimized with full periodic boundary conditions in order to retain the structure stability of IRMOFs. All the framework atoms were set as flexible to adjusted the energetically minimization of the structure. Similar as previous works (Liu et al. 2013; Liu et al. 2012; Mu et al. 2010), the gradient corrected (GGA) correlation functional of Perdew and Wang (PW91) (Perdew and Wang 1992) was employed with the double numerical plus (DNP) polarization basis set (Delley 1990, 1991), carried out by Dmol 3 code (Delley 2000). All electron was employed in the nucleus, the orbital global cutoff quality was set as fine, and the convergence threshold parameters for optimization were 1×10^{-5} for energy.

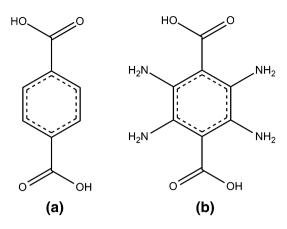


Fig. 2 Organic linkers of IRMOFs: a BDC of IRMOF-1 and b $\rm N_4BDC$ of IRMOF-NH $_2$

In the second step, the supercell models of IRMOFs were generated from the optimized unit cell by $1 \times 2 \times 1$ primitive cells to proceed the optimization of the adsorbate-adsorbent complexes, as shown in Fig. 3. This calculation, also using DFT-GGA/PW91 method, was conducted to determine the energy minimum orientations and positions of the adsorbate molecules on the frameworks. Because of large amount of atoms in a unit cell. periodic supercell model was performed, which not only maintains the local environment of the binding sites but also reduces computational demanding. In this step, all the framework atoms were set as fixed to prevent deformations of the structure, while the atoms of adsorbates were allowed to relax in order to satisfy the certain minimumenergy criteria. Previous work (Ford et al. 2009) had indicated the negligible atomic flexibility of IRMOF structures when the kinetic diameters of the adsorbed molecules were greatly smaller than pore size; and/or the frameworks did not sustain any strong guest-host interactions in the adsorption system. The purpose of this step is to relax the energy of the adsorption system and prepare for the following calculations of BEs.

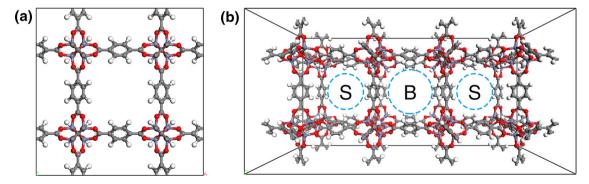
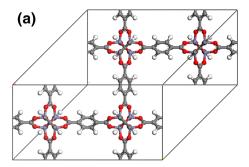


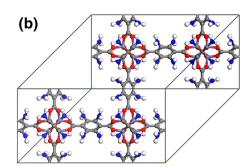
Fig. 1 The a crystal structure and b the pores division of IRMOF-1 (S represents small cells, while B represents big cells)



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Fig. 3 The supercell models of IRMOFs: **a** IRMOF-1 **b** IRMOF-NH₂





After geometry optimization, a variety of possible configurations were found, but only those in the energy minimum locations (binding sites) were discussed in detail. Four major binding sites, denoted as site IS (small cell of the metal corner), site IB (big cell of the metal corner), site IIS (small cell of the linker), respectively, were discussed. In addition, two types of orientations of VOC molecules on each binding site, *cis*-form orientation (the oxygen atoms of VOCs turn to the binding sites, noted as 1) and anti-form orientation ($-CH_3$ groups turn to the binding sites, noted as 2), were also taken into account, as shown in Fig. 4. Thus, eight (4×2) adsorption configurations were considered in this work.

2.3 Binding energies calculations

BE is regarded as a significant parameter in indicating the interaction strength of adsorbate adsorption on MOFs, hence the BE calculations of the selected adsorption configurations were processed after geometry optimization, also using GGA-level of PW91/DNP procedure. Only one VOC molecule per supercell was considered and the BE was evaluated as Eq. 1:

$$BE = E_{MOF/VOC} - (E_{MOF} + E_{VOC}) \tag{1}$$

where $E_{MOF,VOC}$ is the total energy of the IRMOFs/VOCs adsorption complexes in equilibrium state, and E_{MOF} and E_{VOC} are the total energy of the guest-free IRMOFs structures and the chosen VOC adsorbate, respectively. The negative value of binding energy shows the exothermic binding of the VOC adsorbates on IRMOFs, a higher absolute value of BE indicates a stronger adsorption strength.

The chosen functional PW91, combining with DNP basis set, has been successfully used by researchers for geometry optimization and BE calculations of H-bonding and electrostatic-dominated interactions on MOFs (Wu et al. 2010; Venkataramanan et al. 2009). Nevertheless, GGA/PW91 simulations have somewhat limitation in describing dispersion energies effects (Pakarinen et al. 2009; Wu et al. 2001), and may slightly underestimates the absolute values of BEs when treating long-range

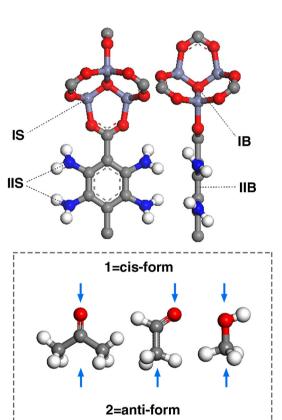


Fig. 4 The binding sites in IRMOFs and the orientation of VOCs to the binding sites. *IS* the small cell at the corner, *IB* the big cell at the corner, *IIS* the small cell at the linker, *IIB* the big cell at the linker. *I* represents the *cis*-form orientation of adsorbates, while 2 represents the anti-form orientation

interactions. However, the purpose of this work is to study the role of amino-functionalization on VOC adsorptions by comparing the calculated BEs among the different configurations, and between different adsorption sites. It has been reported that the adsorbate-framework dispersive interactions on IRMOF-1 and IRMOF-NH₂ were comparative, and the amino introduction did not have significant effects on it (Mu et al. 2010). Moreover, the previous work (Torrisi et al. 2009) also indicated a good agreement between PW91 and MP2 method in calculating the BE and configuration of benzene/CO₂ molecular complex, which



was governed by weak van der Waals interactions. It is, therefore, the deviation caused by dispersion effects could be offset when making a relative comparison. Actually, the GGA/PW91 simulations have been successfully employed by many researchers to study weak van der Waals (vdW) interactions on MOFs (Yan et al. 2014; Wu et al. 2014b).

2.4 Atomic partial charge calculations

The atomic partial charges of IRMOFs were computed to quantitatively clarify the amino-functionality effects on charge transfer and ESP of the frameworks. Similar as the previous works (Xu and Zhong 2010; Wu et al. 2012; Priya et al. 2014), the nonperiodic cluster models were used to calculate the ESP derived charges of the atoms in the frameworks, which were treated as the atomic partial charges in this paper. The cluster models were illustrated in Fig. 5. To maintain the correct hybridization, the dangling bonds on the fragmented clusters were terminated by methyl (-CH₃) groups. On the basis of the ChelpG method (Breneman and Wiberg 1990), DFT calculations were performed by using unrestricted B3LYP functional. The basis set LANL2DZ was used for metal atom Zn, while 6-31+G* (Hariharan and Pople 1972) was used for the rest of the atoms. For complexes involving heavy metal atoms, the approach of effective core potential basis set (e.g. LANL2DZ) for heavy atoms was commonly adopted to mixed with all-electron basis sets for rest atoms. LANL2DZ is a collection of double-ξ basis sets (Foguet Albiol et al. 2005) that contains effective pseudopotentials to represent the core potential for valence electrons, which can significantly reduce the computational cost. These calculations were performed using the GAUSSIAN 09 suite of programs (Frisch et al. 2009), and the results were listed in Table 1.

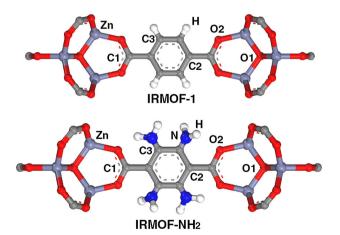


Fig. 5 Cluster models used in the atomic partial charges calculations for IRMOF-1 and IRMOF-NH $_2$

Table 1 The atomic partial charges (eV) of IRMOFs

Atom types	IRMOF-1	IRMOF-(NH ₂) ₄
C1	+0.73	+1.03
C2	+0.05	-0.52
C3	-0.14	+0.27
Н	+0.15	+0.35
N	_	-0.79
O1	-1.53	-2.34
O2	-0.66	-0.86
Zn	+1.28	+1.75

3 Results and discussion

3.1 Electrostatic properties of IRMOFs

Table 1 lists the atomic partial charges of IRMOF-1 and IRMOF-NH₂ (with the atom types denoted in Fig. 5). As can be seen in Table 1, some atomic charges greatly increase after amino-functionalization, e.g. the charges of C3 atoms increase to be +0.27 eV from -0.14 eV, the coordinatively unsaturated site (CUS) of Zn atoms also have a more positive charge of +1.75 eV on IRMOF-NH₂. This indicates an electrons deficit of these atoms that results from the high electronegativity of amino groups. Conversely, the charges on the C2 atoms of IRMOF-NH₂ decrease to a negative one of -0.52 eV from IRMOF-1, and the charges of O1 and O2 atoms also slightly decrease to a more negative value. These results reveal the great influence of the electron-donating groups (-NH2) on the charge distributions of IRMOFs, inducing the electron transfer and electrostatic density redistribution on the whole framework, which may consequently change the VOCs adsorption performance. The detail of amino-functionalization effects on IRMOFs are studied by comparing the BEs in the following sections.

3.2 Adsorption thermodynamics of methanol on IRMOFs

3.2.1 Methanol adsorbed on IRMOF-1

Figure 6 shows the geometric equilibrium configurations and the corresponding BEs on each binding sites of IR-MOF-1. It is noted that the BEs of the configurations on site I are significantly higher than those on site II, indicating more adsorption affinity of methanol on CUS sites, which resembles the adsorption of other adsorbates like benzene (Zeng et al. 2012) and small gases (Dubbeldam et al. 2007) on IRMOF-1 verified both experimentally and theoretically. Due to the electronic-accepting effects of CUS sites (Mu et al. 2010), the adsorption on site I is



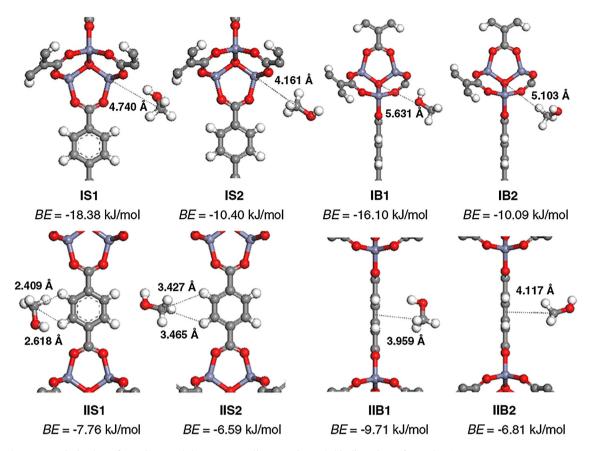


Fig. 6 The DFT optimized configurations and the corresponding BEs in each binding sites of IRMOF-1

governed by electrostatic interactions, which is greatly stronger than the vdW interactions on site II. Besides, the calculated BEs of IRMOF-1 show different trends on site I and site II. The BE on site I follows the order of site IS > site IB, but it turns to be site IIB > site IIS on site II reversely, suggesting varied interaction mechanism on site I and site II.

On site I, larger amount of Zn atoms on the big cell (site IB) give rise to the stronger positive electrostatic field, which will repulsively interacted with adsorbates possessed similar positive field. Contrarily, these adsorbates can attractively interact with site IS, which exhibits relatively weaker positive ESP. It is noted that the BEs of methanol on site IB are weaker than site IS (e.g. -16.10 and 18.38 kJ/mol for IB1 and IS1, respectively), suggesting site IB may repulsively interacts with -CH₃ group of methanol, since the -CH₃ group possesses positive electrostatic field because of the electron-donating effects of the –OH group. Correspondingly, -CH₃ group of methanol is able to interact with site IS preferably. On site II, however, the adsorption behavior of the framework differs from that on site I. The BEs on site IIB are even slightly higher than site IIS (-9.71 and -7.76 kJ/mol for IIB1 and IIS1, respectively), suggesting a π system-dominated interaction on the organic linker. The conjugated π -system with stronger negative electrostatic field presents above both sides of the linker (site IIB), which interacts with the –CH₃ group of methanol preferably, while methyl-linker interaction is weaker at the side of aromatic ring (site IIS) due to the lack of π system delocalization.

On the other hand, the orientation and distance of -CH₃ group of methanol to the specific binding site also shows varied trends for site I and site II on IRMOF-1. On both site IS and site IB, methanol with cis-form orientation (1) has a greater adsorption affinity (BE) and a farther distance from binding sites than anti-form (2). In this case, the final interaction strength of methanol on site I can be controlled by either the electrostatic-dominated interaction between -CH₃ group and binding sites or the steric hindrance of methanol molecule, or both of them, depending on the properties of the adsorbate molecule, such as ESP, textural properties, etc. The behaviors of methanol on site I suggests that the negative steric hindrance effects of -CH₃ group is relatively stronger than the electrostatic interaction with binding sites of the framework. In addition, the H-bonding interaction between -OH group of methanol and the neighboring O2 atoms also contributes to methanol adsorption with cis-form orientations (H-O2 distance: 2.33 and 2.50 Å for site IS1 and IB1, and O-H-O₂ angles are less than 20° for both sites). H-bonding interaction should



be considered when two conditions are satisfied: the H-bond distance less than 3.5 Å, and the hydrogen-donor–acceptor angle less than 30° (Luzar and Chandler 1996). On site II, however, the BEs become more exothermic in a closer binding distance with *cis*-form configuration, e.g. methanol has a greater adsorption affinity on site IIB1 in a closer binding distance (3.959 Å) than site IIB2 (4.117 Å). It suggests the negative steric hindrance effects becomes less important on site II and the adsorption turns to be controlled by methyl- π system interactions, resulting from the larger binding environment around the organic linker than the metal corner.

Combining the adsorption behaviors and the electrostatic properties of methanol and IRMOF-1, a plausible adsorption mechanism is deduced that the adsorption of methanol on IRMOF-1 is likely dominated by the electrostatic interactions between the binding sites and the –CH₃ group of methanol, while the polar group (–OH) of methanol only interacts with the framework through H-bonding interactions. This will be further verified with the comparative study of aldehyde and acetone in the following section.

3.3 Methanol adsorbed on IRMOF-NH₂

In order to understand the interaction mechanism of methanol over IRMOFs, electron-donating –NH₃ groups are introduced to modify the electrostatic properties of the framework for comparison. The adsorption configurations

and the corresponding BEs on each binding sites of IRMOF-NH₂ are calculated in Fig. 7. As have discussed earlier, amino-functionalization has a great influence on the charge distribution of the framework, which is expected to improve methanol adsorption performance, similar as the adsorption of CO₂ (Mu et al. 2010) on amino-modified IRMOFs. The BEs on site II are indeed greatly intensified by -NH₂ groups, especially on site IIS1 and IIB1 (13.12) and 7.57 kJ/mol higher than IRMOF-1, respectively). Besides, methanol tends to stabilize in a closer binding distance with cis-form orientation on site II, similar as IRMOF-1, indicating amino-functionality does not significantly change the steric environment around the linker, since the site II interaction on IRMOF-NH2 is still dominated by π system. However, the BEs on site I (site IS1/2) and IB1/2) of IRMOF-NH₂ are less exothermic compared to IRMOF-1. It suggests varied effects of amino-functionalization that both enhance the methyl- π system interactions on site II and weaken the interaction strength on site I.

To investigate the varied effects of amino-functionalization on site I and site II, the contour maps of the ESPs of IRMOFs are further calculated in Fig. 8. Similar as the previous work (Mu et al. 2010), the ESPs are obtained using BLYP exchange–correlation functional of GGA level, and DNP basis set is also adopted. As shown in Fig. 8, IRMOF-NH₂ has an ESP with larger gradient and higher absolute values above the two sides of the aromatic ring, suggesting a stronger electrostatic field on site II of

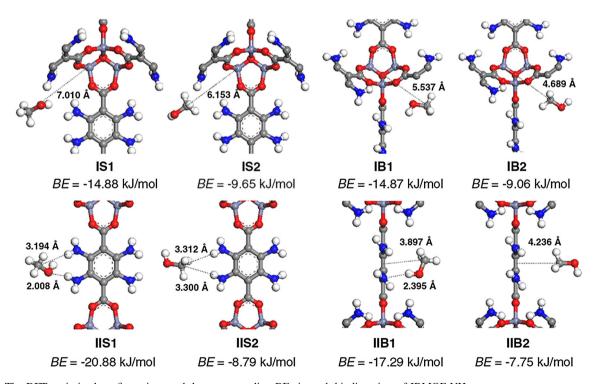


Fig. 7 The DFT optimized configurations and the corresponding BEs in each binding sites of IRMOF-NH₂



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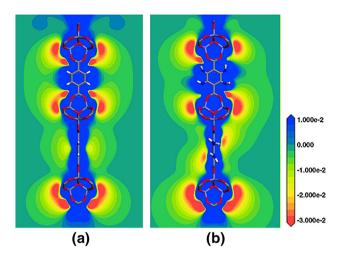


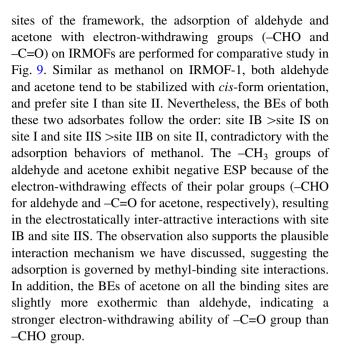
Fig. 8 The contour maps of electrostatic potentials (ESP) for: a IRMOF-1 and b IRMOF-NH₂

IRMOF-NH₂ than that on IRMOF-1. It can be attributed to the conjugation effect of the lone pair electrons in -NH₂ groups, which greatly enhances the π system delocalization and polarized electron density above the aromatic ring. This observation agrees well with the results for the stronger BEs on site II of IRMOF-NH2 than IRMOF-1 (shown in Figs. 6, 7). Meanwhile, weak H-bonding interaction between -OH group of methanol and -NH2 groups forms when these two groups approach each other, which also contributes to adsorbate-adsorbent interactions in addition to the electrostatic interactions. Contrary to site II, the BEs of site I in IRMOF-NH2 are even slightly weaker compared to IRMOF-1 (shown in Figs. 6, 7). It can be readily evidenced by the increased positive electrostatic fields around the CUS of Zn atoms (as shown in Fig. 8), also as noted from the more positive atomic partial charges of Zn atoms [+0.47 eV higher than IRMOF-1 (from +1.28 eV to +1.75 eV), indicating adsorbate-adsorbent interactions are suppressed by the repulsive methyl-CUS interactions. Moreover, the negative steric effect of inserted –NH₂ groups also more hinders the adsorption on site I than site II, since the CUS binding sites represent only a small region of the framework and can be easily blocked by the functional groups. Consequently, the adsorption of methanol on site I of IRMOF-NH₂ is strongly suppressed by both the electrostatically repulsive interaction and the steric inhibition of -NH₂ groups.

3.4 Adsorption thermodynamics of aldehyde and acetone on IRMOFs

3.4.1 Aldehyde and acetone adsorbed on IRMOF-1

To verify the electrostatic-dominated interaction mechanism between -CH₃ group of the adsorbate and binding



3.5 Aldehyde and acetone adsorbed on IRMOF-NH₂

The adsorption of aldehyde and acetone on IRMOF-NH₂ is also performed for a further investigation, as the calculated BEs shown in Fig. 10. As is expected, aldehyde and acetone have stronger BEs on site I of IRNMOF-NH₂ compared to IRMOF-1, resulting from the enhanced positive ESP around the metal corners by amino-functionalization. Meanwhile, the increased BEs are slightly higher on site IB than site IS for IRMOF-NH₂ (e.g. for aldehyde, 2.48 and 0.91 kJ/mol higher than IRMOF-1 on site IB1 and IS1, respectively), implying the enhanced effect of aminofunctionalization on site IB is greater than site IS.

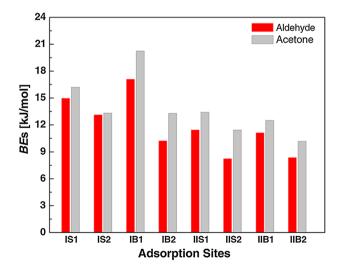


Fig. 9 The BEs of aldehyde, acetone and methanol adsorbed in IRMOF-1



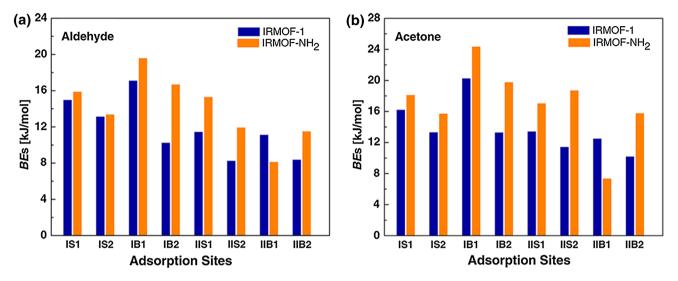


Fig. 10 The binding energies of a aldehyde and b acetone adsorbed in IRMOF-1 and IRMOF-NH₂

Additionally, the BEs of aldehyde and acetone on site IIS1/2 are also stronger than IRMOF-1, suggesting the relatively stronger positive potential field of site IIS to attractively interact with the negative potential $-CH_3$ group, which results from greatly enhanced delocalization of π system above the aromatic ring (shown in Fig. 8b).

On the basis of the interaction mechanism deduced, the adsorption of aldehyde and acetone on both site IIB1 and site IIB2 of IRMOF-NH₂ are expected and should be hindered by amino-insertion, resulting from the repulsive electrostatic interactions between –CH₃ group and the enhanced π system on site IIB, similar to the repulsive adsorption of methanol on site I of IRMOF-NH₂. However, it is interesting to find the exception on site IIB2, where the BEs of both aldehyde and acetone are unexpectedly more exothermic compared to IRMOF-1. As illustrated in Fig. 11, –CH₃ groups of the adsorbates on site IIB have greater adsorption affinity with anti-form orientation (site IIB2), which also has a closer

distance from the binding site than *cis*-form (site IIB1) (1.57 and 2.05 Å closer for aldehyde and acetone, respectively). Figure 8b shows positive potential field exists between the aromatic ring and the π delocalization region, as confirmed from the positive partial charges of aromatic C3 atoms (increased to +0.27 eV from -0.14 eV of IRMOF-1), which is likely to interact with -CH₃ groups of aldehyde and acetone preferably when approaching close enough. Compared to methanol, -CH₃ groups of aldehyde and acetone possess stronger ESP and therefore, is capable of accessing the opposite electrostatic binding site via the negative π system delocalization. These adsorption behaviors of aldehyde and acetone on site IIB2 show the interaction of VOCs with MOFs is complex and that more work is needed to probe the mechanisms of these processes. In addition, acetone is observed to have a slightly weaker adsorption affinity (-7.35 kJ/mol) than aldehyde (-8.12 kJ/mol) on site IIB1 of IRMOF-NH₂. It can attribute to -CH₃ groups of acetone

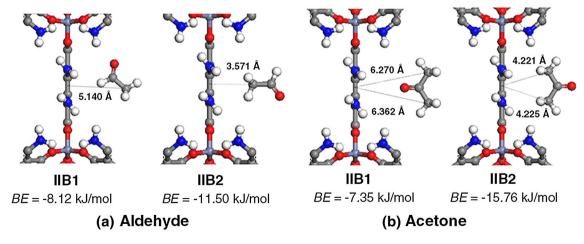


Fig. 11 The DFT optimized structures of a aldehyde and b acetone in site IIB of IRMOF-NH₂



that possesses both stronger negative electrostatic field and a larger steric occupation than aldehyde, inducing the repulsive electrostatic interactions and negative steric hindrance effects on site IIB1, and consequently weaken the adsorbate-adsorbent interactions.

4 Conclusions

DFT calculation has been carried out to study the adsorption behaviors of methanol, aldehyde and acetone on IRMOFs. The BEs and the energy-minimum configurations of these VOCs on IRMOF-1 and a newly-constructed IRMOF-NH₂, are compared to reveal the functionality effects of amino groups on VOCs adsorption. Moreover, the atomic partial charges and ESPs of IRMOFs are also calculated to elucidate the electron transference and distribution on the frameworks caused by the inserted amino groups. The calculated results indicate that the electrondonating effect of amino groups greatly intensifies both the electrostatic delocalization above two sides of the organic linkers (site II), and the positive potential field around the metal corners (site I). Consequently, the adsorption of methanol is greatly enhanced by attractive interactions on site II of IRMOF-NH₂, but weakened by electrostatically repulsive interactions on site I. However, reverse behaviors are observed on the adsorption of aldehyde and acetone, resulting from the varied ESP field of their methyl groups. A plausible adsorption mechanism is proposed that the adsorption of these VOCs on IRMOFs is likely controlled by electrostatic interaction between methyl groups of the adsorbates and the binding sites of the framework, while the polar groups of these adsorbates only interact with framework through weak interaction strength H-bonding). Besides, the inserted amino groups may also sterically hinder the electrostatic interactions, depending on the configurations and the electronic properties of the adsorbates. The underlying binding mechanism elucidated by a detailed electrostatic and structural analysis, provides guidelines to the synthesis of new functional MOFs for the adsorption of VOCs.

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