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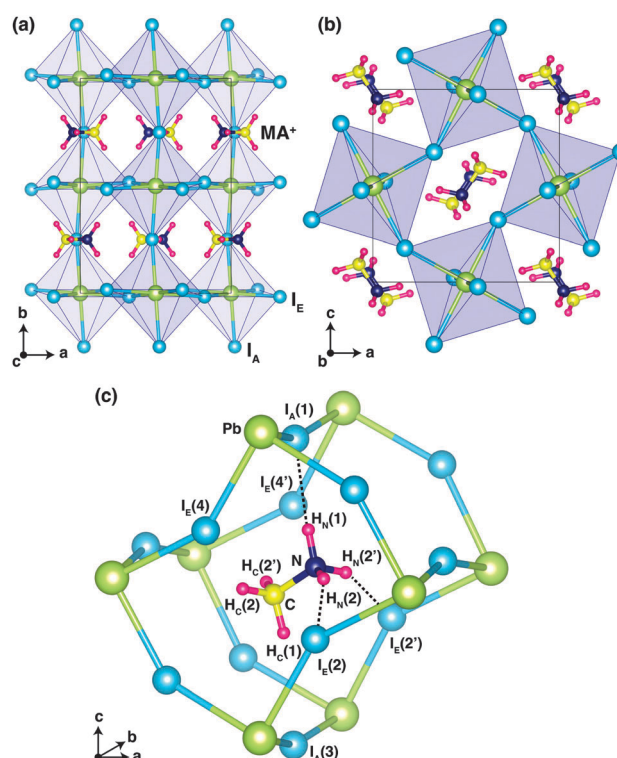
# Role of hydrogen-bonding and its interplay with octahedral tilting in $\text{CH}_3\text{NH}_3\text{PbI}_3$

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**First principles calculations on the hybrid perovskite  $\text{CH}_3\text{NH}_3\text{PbI}_3$  predict strong hydrogen-bonding which influences the structure and dynamics of the methylammonium cation and reveal its interaction with the tilting of the  $\text{PbI}_6$  octahedra. The calculated atomic coordinates are in excellent agreement with neutron diffraction results.**

$\text{CH}_3\text{NH}_3(\text{MA})\text{PbI}_3$  is one of the most extensively studied hybrid halide perovskites due to its impressive power conversion efficiency for solar cell applications.<sup>1</sup> In spite of numerous studies on its structural properties,<sup>2–8</sup> relatively little progress has been made in our understanding on the role of hydrogen-bonding in the orthorhombic phase (*o*-MAPbI<sub>3</sub>). In this study we have used density functional theory (DFT) calculations to examine the thermodynamic consequences of hydrogen-bonding and its interplay with the octahedral rotations that are ubiquitous in perovskites. The computed atomic positions, including those of the hydrogen atoms, are compared to a recent powder neutron diffraction study.<sup>9</sup>

The DFT calculations were performed using the generalized gradient approximation (GGA) and projector augmented wave (PAW)<sup>10</sup> pseudopotentials as implemented in the Vienna ab initio Simulation Package (VASP).<sup>11–13</sup> The effects of spin-orbit coupling and van der Waals (vdW) interactions<sup>14</sup> were included during structural and electronic relaxation. We adopted (i) a  $4 \times 3 \times 4$  Monkhorst-Pack *k*-point mesh centered at  $\Gamma$ ,<sup>15</sup> (ii) a 500 eV plane-wave kinetic energy cutoff, and (iii) the tetrahedron method with the Blöchl corrections for the Brillouin zone integrations.<sup>16</sup> We explicitly treated 14 valence electrons for Pb ( $5d^{10}6s^26p^2$ ), 7 for I ( $5s^25p^5$ ), 4 for C ( $2s^22p^2$ ), 5 for N ( $2s^22p^3$ ), and 1 for H ( $1s^1$ ). The ions were relaxed until the forces on them were less than  $0.01 \text{ eV } \text{\AA}^{-1}$ . The experimentally



**Fig. 1** (a) Optimized crystal structure of *o*-MAPbI<sub>3</sub>, viewed in the *a*-*b* and, (b) *a*-*c* planes.  $\text{PbI}_6$  octahedra are highlighted in purple. (c) Local cage structure of *o*-MAPbI<sub>3</sub> is characterized by hydrogen-bonding between the three H atoms in the  $\text{NH}_3$  group and one axial I [ $I_A(1)$ ] atom and two equatorial I [ $I_E(2)$  and  $I_E(2')$ ] atoms. Hydrogen-bonding interactions are indicated by dotted lines.

observed ratio of the orthorhombic unit cell vectors<sup>2</sup> was imposed when we optimized the lattice parameters by calculating the Kohn-Sham (K-S) energy as a function of volume.<sup>17</sup>

Fig. 1 shows the optimized crystal structure of *o*-MAPbI<sub>3</sub> with space group *Pnma*. The computed lattice parameters are  $a = 8.844 \text{ \AA}$ ,  $b = 12.592 \text{ \AA}$ , and  $c = 8.563 \text{ \AA}$ , in good agreement with the experimental values at 100 K ( $a = 8.8362 \text{ \AA}$ ,  $b = 12.5804 \text{ \AA}$ ,

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**Table 1** Calculated fractional coordinates for *o*-MAPbI<sub>3</sub> compared with experimental values. The H<sub>C</sub>(1) and H<sub>C</sub>(2) atoms are connected to C while the H<sub>N</sub>(1) and H<sub>N</sub>(2) atoms are connected to N in the MA cation. ND and XD denote neutron powder diffraction data<sup>9</sup> and X-ray diffraction data,<sup>2</sup> respectively

	Wyckoff site	This work				Experiment		
		<i>x</i>	<i>y</i>	<i>z</i>		<i>x</i>	<i>y</i>	<i>z</i>
Pb	4b	0.500	0.000	0.000	ND	0.500	0.000	0.000
					XD	0.500	0.000	0.000
I <sub>A</sub> (1)	4c	0.476	0.250	0.938	ND	0.484	0.250	0.944
					XD	0.486	0.250	0.947
I <sub>E</sub> (2)	8d	0.176	0.017	0.172	ND	0.189	0.015	0.184
					XD	0.190	0.017	0.186
N	4c	0.953	0.750	0.023	ND	0.942	0.750	0.030
					XD	0.932	0.750	0.029
C	4c	0.908	0.250	0.075	ND	0.937	0.250	0.058
					XD	0.913	0.250	0.061
H <sub>C</sub> (1)	4c	0.941	0.250	0.198	ND	0.937	0.250	0.187
H <sub>C</sub> (2)	8d	0.842	0.178	0.048	ND	0.866	0.170	0.029
H <sub>N</sub> (1)	8d	0.111	0.183	0.003	ND	0.128	0.189	−0.009
H <sub>N</sub> (2)	4c	0.978	0.750	0.142	ND	0.954	0.750	0.146

and  $c = 8.5551$  Å for the X-ray study<sup>2</sup> and  $a = 8.8657$  Å,  $b = 12.6293$  Å, and  $c = 8.5769$  Å for the neutron study<sup>9</sup>). Note that the calculated values lie between the experimental ones. The perovskite structure exhibits corner-linked PbI<sub>6</sub> octahedra forming a three-dimensional network. The *Pnma* phase displays a particular pattern of rotations of the octahedra involving either in-phase or out-of-phase rotations around the three I–Pb–I octahedral axes, denoted by  $a^-b^+a^-$  in Glazer's notation.<sup>18</sup> Regarding the organic MA ion, the C–N bonds are aligned approximately along the [101] and [10 $\bar{1}$ ] directions with a head-to-tail configuration as shown in Fig. 1. Our optimized atomic positions for this structure are listed in Table 1, where H atoms that are connected to N atoms are denoted by H<sub>N</sub>, and H atoms that are connected to C atoms are denoted by H<sub>C</sub>. The subscripts A and E on the iodine atoms refer to axial and equatorial configurations, respectively, as shown in Fig. 1(c). Our optimized atomic positions agree very well with the previous experimental values. In particular, the coordinates of the H atoms are in very good agreement with the recent powder neutron diffraction study.<sup>9</sup>

Fig. 1(c) shows the ground-state configuration of the H atoms in the surrounding inorganic Pb–I cage. The positions of the H atoms conform with the *Pnma* space group. If we assume that hydrogen-bonding is important when  $H \cdots I < 3$  Å, there are three hydrogen-bonds in the Pb–I cage between H<sub>N</sub>(1), H<sub>N</sub>(2), and H<sub>N</sub>(2') and I<sub>A</sub>(1), I<sub>E</sub>(2), and I<sub>E</sub>(2'), respectively.

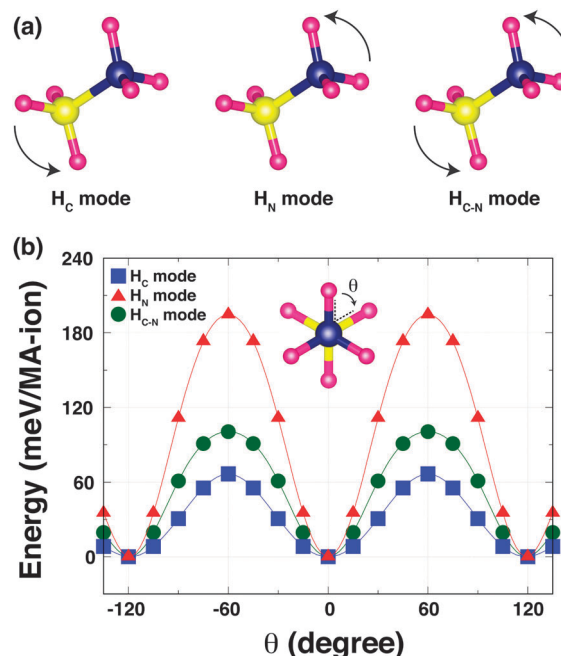
These three  $H \cdots I$  bonds per MA ion are controlled by both the particular inorganic  $a^-b^+a^-$  tilt pattern and the organic MA conformation. Among these, the H<sub>N</sub>(1)  $\cdots$  I<sub>A</sub>(1) hydrogen bond is the shortest (2.565 Å) and presumably the strongest. Accordingly, the N–H<sub>N</sub>(1)  $\cdots$  I<sub>A</sub>(1) angle is almost 180° (Table 2). These results are in good agreement with the recent powder neutron diffraction study, as well as the structure of ammonium iodide (NH<sub>4</sub>I)<sup>19</sup> where hydrogen-bonding is known to play a key role in structural stabilization.<sup>20</sup> Details of the computed bond lengths and bond angles for the principal interactions are given in Table 2 and compared with the experimental neutron diffraction values. The N  $\cdots$  I distances are much shorter than those of C  $\cdots$  I and also the angle of C–H<sub>C</sub>(1)  $\cdots$  I<sub>A</sub>(3) is smaller than that

**Table 2** Calculated H  $\cdots$  I, N  $\cdots$  I, and C  $\cdots$  I bond lengths and N–H  $\cdots$  I, C–H  $\cdots$  I and Pb–I–Pb bond angles compared with NH<sub>4</sub>I<sup>19</sup> and ND data for MAPbI<sub>3</sub>

	This work	NH <sub>4</sub> I <sup>19</sup>	Experiment <sup>9</sup>
Bond type (Å)			
H <sub>N</sub> (1) $\cdots$ I <sub>A</sub> (1)	2.565	2.454	2.613
H <sub>N</sub> (2) $\cdots$ I <sub>E</sub> (2)	2.611		2.808
H <sub>C</sub> (1) $\cdots$ I <sub>A</sub> (3)	3.137		3.190
H <sub>C</sub> (2) $\cdots$ I <sub>E</sub> (4)	3.094		3.006
N $\cdots$ I <sub>A</sub> (1)	3.606	3.473	3.611
N $\cdots$ I <sub>E</sub> (2)	3.566		3.681
C $\cdots$ I <sub>A</sub> (3)	4.221		4.298
C $\cdots$ I <sub>E</sub> (4)	4.032		4.090
Angle type (°)			
N–H <sub>N</sub> (1) $\cdots$ I <sub>A</sub> (1)	177	178	174
N–H <sub>N</sub> (2) $\cdots$ I <sub>E</sub> (2)	152		146
C–H <sub>C</sub> (1) $\cdots$ I <sub>A</sub> (3)	170		172
C–H <sub>C</sub> (2) $\cdots$ I <sub>E</sub> (4)	144		148
Pb–I <sub>A</sub> (1)–Pb	159		162
Pb–I <sub>E</sub> (2)–Pb	145		151

of N–H<sub>N</sub>(1)  $\cdots$  I<sub>A</sub>(1). Accordingly, the hydrogen-bond interactions mainly originate from H atoms on nitrogen.

To understand the strength of the hydrogen-bonding in *o*-MAPbI<sub>3</sub>, we have examined the energetics as a function of the H positions. The most stable structure shows that the MA cation adopts a staggered conformation [Fig. 2(b) inset], as expected and in agreement with the neutron results; this is denoted by  $\theta = 0^\circ$  in the following calculations in which we rotate the NH<sub>3</sub> and CH<sub>3</sub> groups around the C–N axis for all four MA-ions in the unit cell. We have calculated the Kohn–Sham (K–S) energies as a function of the torsion angle  $\theta$ , fixing all



**Fig. 2** (a) Schematic view of three MA rotational modes (see text). (b) The computed Kohn–Sham (K–S) energies as a function of the torsion angle  $\theta$  for the three rotational modes and a staggered conformation (inset) with the torsion angle  $\theta = 0^\circ$ .



other atoms, for three rotational modes as shown in Fig. 2(a): (i) in the  $H_C$  mode, the  $NH_3$  group is held rigid while the  $CH_3$  group is rotated around the C–N bond axis. This mode reveals the stability of the staggered conformation at  $\theta = 0^\circ$  compared with the eclipsed conformation at  $\theta = 60^\circ$  [Fig. 2(b)]. The staggered conformation is  $\sim 66$  meV per MA-ion more stable than the eclipsed structure. This suggests that the methyl groups may be rotating at high temperatures, but becomes ordered when the material is cooled to lower temperatures, as is seen in  $[(CH_3)_2NH_2]Mn(HCOO)_3$ .<sup>21</sup> (ii) In the  $H_N$  mode, the  $NH_3$  group rotates around the C–N bond axis. This mode reflects the stability of both the staggered conformation and the hydrogen-bonding [Fig. 2(b)], and is the stiffest mode. (iii) In the third mode,  $H_{C-N}$ , the MA cation is held in the staggered conformation and rotated around the C–N axis. This mode is dominated by the strength of the hydrogen-bonding, which represents  $\sim 102$  meV per MA-ion. This value is quite close to that obtained by NMR for the corresponding rotation in the dimethylammonium perovskite formates (88 meV per DMA-ion).<sup>21</sup> Note that the difference in the energy barriers between the  $H_N$  and  $H_C$  modes is similar to the energy barrier for the  $H_{C-N}$  mode. The three K–S energy curves plotted as a function of the torsion angle  $\theta$  for the three rotational modes have  $120^\circ$  periodicity in common. This is because the most stable structure represents not only the staggered conformation but also the optimised hydrogen-bonding in the Pb–I cage.

We have also examined the relationship between hydrogen-bonding and the octahedral tilting in *o*-MAPbI<sub>3</sub>. We began by fixing the  $H_{C-N}$  modes at  $\theta = 0^\circ$  (hydrogen bonding maximised) and  $60^\circ$  (hydrogen bonding minimised) but allowing the inorganic Pb–I cage to relax. We found that the resulting structure and octahedral rotations were almost unaltered. This suggests, unsurprisingly in perovskites with tolerance factors less than 1,<sup>22,23</sup> that the octahedral rotations are robust.

To understand how the robust octahedral rotations (which change amplitude and pattern with temperature), can affect the hydrogen bonding, we performed a second computational experiment. Here, we computed the K–S energies as a function of the torsion angle  $\theta$  for the same three rotational modes but in a hypothetical high-symmetry structure. This high-symmetry structure is characterized by no octahedral tilting, but with the MA ions in the same configuration as the orthorhombic phase [Fig. 3(a)]. When we calculated the energy of the high-symmetry structure, we relaxed only the MA units within this configuration to optimise any hydrogen-bonding in the Pb–I cage.

In the high-symmetry structure, interestingly, the strength of hydrogen-bonding is remarkably reduced. This is seen for the  $H_N$  and  $H_{C-N}$  modes where the energy barriers are significantly shallower in the high-symmetry structure [Fig. 3(b)]. The energy barrier of the  $H_N$  mode is  $\sim 90$  meV per MA-ion, while that of  $H_{C-N}$  mode is  $\sim 12$  meV per MA-ion. The latter is an order of magnitude smaller than the corresponding barrier in the orthorhombic (ground state) phase with octahedral tilting [Fig. 3(b) inset]. On the other hand, the energy barrier of the  $H_C$  mode ( $\sim 96$  meV per MA-ion) of the high-symmetry structure slightly increases. This suggests that the MA cation is held in the staggered conformation and easily rotated around the C–N



Fig. 3 (a) A high-symmetry structure characterised by the removal of octahedral tilting in *o*-MAPbI<sub>3</sub>. (b) The K–S energies plotted as a function of the torsion angle  $\theta$  for three rotational modes in the high-symmetry structure. Inset compares the energy barriers for the  $H_{C-N}$  mode in the high-symmetry structure and the ground-state structure with octahedral tilting.

axis in the high-symmetry structure without invoking the octahedral tilting, in agreement with NMR results.<sup>24</sup> Thus, one can conclude that the octahedron tilting is strongly correlated with the hydrogen-bonding interaction. Our findings are consistent with a recent report of a stiffening of the lattice of a hybrid metal–organic formate  $[NH_4][Zn(HCOO)_3]$  when cooling through a hydrogen bond ordering transition, as revealed by Resonant Ultrasound Spectroscopy.<sup>25</sup> Our results could also explain why the MA-ion ordering does not appear above the orthorhombic transition, but is coincident with the phase transition driven by the robust octahedral rotations.

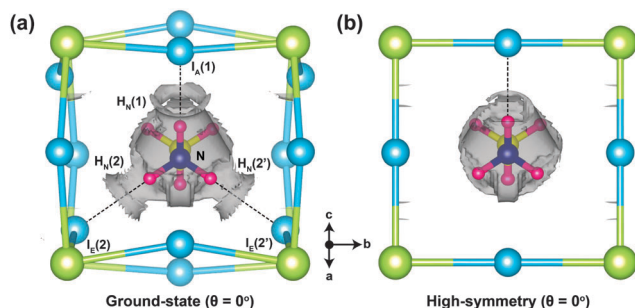


Fig. 4 Non-covalent interaction density isosurfaces for (a) the ground-state ( $\theta = 0^\circ$ ) and (b) high-symmetry structures ( $\theta = 0^\circ$ ). The reduced gradient density cut-off is 1.0 au.





As a way of visualising the hydrogen-bonding in *o*-MAPbI<sub>3</sub>, we present in Fig. 4 the non-covalent interaction isosurfaces calculated using the Critic2 code<sup>26,27</sup> for the ground-state and high-symmetry structures at  $\theta = 0^\circ$ . The ground-state case in Fig. 4(a) shows non-covalent interaction density located between three H atoms [H<sub>N</sub>(1), H<sub>N</sub>(2), and H<sub>N</sub>(2')] and three I atoms [I<sub>A</sub>(1), I<sub>E</sub>(2), and I<sub>E</sub>(2')], which is consistent with hydrogen bonding in the energy minimised structure. On the other hand, there is only one non-covalent interaction density contour between H<sub>N</sub>(1) and I<sub>A</sub>(1) in the high-symmetry structure. The octahedron tilting is clearly coupled to the strength of the hydrogen-bonding. To further quantify this strength we have calculated the hydrogen bond strength index<sup>28</sup> from the N–H stretching frequencies of the MA molecule in isolation and in the perovskite. For the H<sub>N</sub>(1)··I<sub>A</sub>(1) bond, for example, this index is 0.11 in the ground-state structure but only 0.03 in the high-symmetry structure. For the H<sub>N</sub>(2)··I<sub>E</sub>(2) bond the values are 0.05 and 0.02 respectively. The 11% softening of the N–H stretching frequency in *o*-MAPbI<sub>3</sub> is comparable to the softening of the O–H stretching frequency in ice,<sup>29</sup> suggesting similar strengths of hydrogen bonding in both cases.

We have calculated precise positions for the H atoms in *o*-MAPbI<sub>3</sub> and found excellent agreement with a recent powder neutron diffraction study.<sup>9</sup> We have also explored the role of hydrogen bonding in relation to the behaviour of the MA cations and the tilting of the PbI<sub>6</sub> octahedra. The MA cation adopts a staggered conformation in the optimised structure. It is shown that rotation of the staggered MA cation around its C–N bond is quite hindered, with an activation energy of ~102 meV per MA-ion, due to the strong hydrogen-bonding. The strength of the hydrogen-bonding is found to be highly correlated with the octahedral rotations in MAPbI<sub>3</sub>, providing an explanation for why the MA-ordering transition is coincident with the tilting phase transition temperature.

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