# Close examination of the structure and dynamics of HC(NH<sub>2</sub>)<sub>2</sub>PbI<sub>3</sub> by MD simulations and group theory

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Supporting Information

Rendering of the supercell corresponding to the simulation of the 444 system at 300 K, for t=50.7 ps. The thin pale lines display the Pb-I bonds to facilitate the understanding of the overall structure.



Time evolution of the cell box lengths (A) and cell angles (B) from the first principles MD simulation of FAPI at **T=300 K**. The curves are actually running averages over 100 points and each point is registered every 1 fs. The cell lengths are divided by 4 in order to represent the lattice constants. The dashed horizontal line in black in panel A is calculated by taking the cubic root of the volume averaged in the lapse that goes from 25 ps to 50 ps, which then results in an average lattice parameter for the cubic cell of <a = 6.39144 Å.

Average supercell parameters calculated for the interval [25,50] ps. All lengths in Å

AX	=	25.583	379601	609736	3		
AY	=	-3.197	77461	129045	20E-002		
AZ	=	6.304	196043	263489	15E-002		
ΒX	=	1.399	933686	753812	15E-002		
BY	=	25.550	94899	340402	.7		
ΒZ	=	-2.938	30995	995350	63E-002		
CX	=	-6.910	)18552	507026	98E-002		
CY	=	-1.019	960780	776709	88E-002		
CZ	=	25.564	26411	562397	8		
VOL			=	16710	.021694	296240	
VOL**	*(1.	/3.)	=	25	.565779	9494771	43



Histograms corresponding to the cell dimensions (A) and cell angles (B) from the first principles MD simulation of FAPI at **T=300 K**. The histograms are calculated taking every frame from 25 ps to 50 ps. The dashed lines are fits with a normal distribution. For panel A,  $\langle a \rangle = 6.39144$  Å.

## Cell Dimensions:

Fitting formula: y = 1/sqrt(2\*3.14159)/A1\*exp(-((x-A0)^2)/(2\*A1^2)) Tolerance = 1e-05 A: Computed values: A0 = 0.00691737 A1 = 0.0259778 Chi-square: 17.4181 Correlation coefficient: 0.998318 Theil U coefficient: 0.0487635

```
B:
Computed values:
     A0 = 0.00691737
     A1 = 0.0259778
Chi-square: 17.4181
Correlation coefficient: 0.998318
Theil U coefficient: 0.0487635
C:
Computed values:
     A0 = 0.000503465
     A1 = 0.0278935
Chi-square: 23.5542
Correlation coefficient: 0.997488
Theil U coefficient: 0.0586736
Angles:
alpha:
Computed values:
     A0 = 90.1038
     A1 = 0.512617
Chi-square: 0.112141
Correlation coefficient: 0.996805
Theil U coefficient: 0.0638638
```

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beta:
```

```
gamma:
Computed values:
      A0 = 90.0504
      A1 = 0.54451
Chi-square: 0.129474
Correlation coefficient: 0.996000
```

```
Theil U coefficient: 0.0703622
```



Time evolution of the cell box lengths (A) and cell angles (B) from the first principles MD simulation of FAPI at **T=450 K**. The curves are actually running averages over 100 points and each point is registered every 1 fs. The cell lengths are divided by 4 in order to represent the lattice constants. The dashed horizontal line in black in panel A is calculated by taking the cubic root of the volume averaged in the lapse that goes from 30 ps to 58.9 ps, which then results in an average lattice parameter for the cubic cell of <a > = 6.49966 Å.



Histograms corresponding to the cell dimensions (A) and cell angles (B) from the first principles MD simulation of FAPI at **T=450 K**. The histograms are calculated taking every frame from 230 ps to 58.9 ps. The dashed lines are fits with a normal distribution. For panel A,  $\langle a \rangle = 6.39144$  Å.

## Cell Dimensions:

Fitting formula: y = 1/sqrt(2\*3.14159)/A1\*exp(-((x-A0)^2)/(2\*A1^2)) Tolerance = 1e-05 A: Computed values: A0 = -0.00379283 A1 = 0.0329111 Chi-square: 33.8019 Correlation coefficient: 0.997231 Theil U coefficient: 0.0626726

```
B:
Computed values:
     A0 = 0.00363547
     A1 = 0.0337745
Chi-square: 41.2574
Correlation coefficient: 0.996508
Theil U coefficient: 0.0700441
C:
Computed values:
     A0 = 0.00139695
     A1 = 0.0336066
Chi-square: 51.4399
Correlation coefficient: 0.995783
Theil U coefficient: 0.077564
Angles:
alpha:
Computed values:
     A0 = 90.0101
     A1 = 0.70345
Chi-square: 0.34543
Correlation coefficient: 0.991168
Theil U coefficient: 0.101673
beta:
Computed values:
     A0 = 90.0527
     A1 = 0.647799
Chi-square: 0.256629
Correlation coefficient: 0.994359
Theil U coefficient: 0.0836695
gamma:
Computed values:
     A0 = 89.9542
     A1 = 0.655574
Chi-square: 0.302291
Correlation coefficient: 0.993144
Theil U coefficient: 0.0917068
```

#### SYSTEM 222, 300 K

In order to check the stability of our model system with respect to system size we performed two simulations based on the 96 atoms, 222 system.

In the first case, the setting were exactly the same as in the NPT\_F simulations of the system 444. The magnitude of the fluctuations makes are excessively large to perform any meaningful analysis.

In the second case we restricted the supercell fluctuations to isotropic expansions and/ or contractions by using the NPT\_I keyword in the CP2K input. In this case the model behaves in a more stable way, although showing very large fluctuation resulting directly from the small number of atoms in the system. The summation was extended for a total of 24 ps.

The average lattice parameter resulting from averaging last 10 ps of simulation is <a>=6.6145 Å, which is 3.5 % larger than the value obtained with the 444 system under NPT\_F conditions. The inspection of the radial distribution functions is interesting in order to further asses the importance of the system size and the resulting fluctuation affecting the local organization.



## Figure S5

Radial distribution function for the atoms pairs forming the inorganic framework of the FAPI obtained from NPT\_I simulations on a 222 system at 300 K. <u>Average relaxation of r+ and r-. SYSTEM 444, 300 K and 450 K. Fits.</u>



Time dependent self correlation function for r+ (black lines) and r- (red lines) at 300 K (solid lines) and 450 K (dashed lines). The green lines are fits as described below.

#### <u>r+, 450 K</u>

```
Fitting with formula: y = A0 \exp(-x/A1)
Tolerance = 1e-05
Computed values:
     a0 = 1.0048
     a1 = 0.521643
Chi-square: 1.92515
Correlation coefficient: 0.996082
RMS relative error: 7.42554
Theil U coefficient: 0.0851113
<u>r-, 450 K</u>
Fitting with formula: y = A0 \exp(-x/A1)
Tolerance = 1e-05
Computed values:
     a0 = 1.02171
     a1 = 1.1698
Chi-square: 4.08357
Correlation coefficient: 0.996016
RMS relative error: 1.40925
Theil U coefficient: 0.0814744
```

- RMS relative error: 0.181207
- Theil U coefficient: 0.0347234

## Relaxation of individual molecules, r+, 300 K, System 444.



#### Figure S7

The relaxation of the individual FA molecules, as displayed by the r+ vector, shows heterogeneity in the same way that the MA does, for 300 K.

<u>Coefficients of fitting r+ colormap at 300 and 450 K using cubic harmonics.</u> <u>System 444.</u>

(i)	300 K C_i	400 K C_i
4	0.898123	0.157817
6	0.307109	0.015817
8	0.399998	0.041476
10	0.130961	0.005785
12	0.045936	0.008149

## **COUPLING BETWEEN MOLECULAR ORIENTATIONS AND LATTICE DISTORTIONS**



## Figure S8

Qualitative assessment of the interplay between the deformations of the inorganic cage and the orientation of the molecules. This figure was done by taking the average position, over the whole production run, of the N and C atoms of the molecules. For 300 K, since the rotational dynamics is slow, the average results in configurations that roughly retain the molecular architecture. For the case corresponding to 450 K, since the rotational dynamic is fast, the average position of the atoms are all located approximately in the center of the inorganic cage. The figure displays a section of the system 444 at both temperatures. The molecules are in average oriented parallel to the long axis of the cavities at 300 K, but no particular correlation is observed at 450 K.