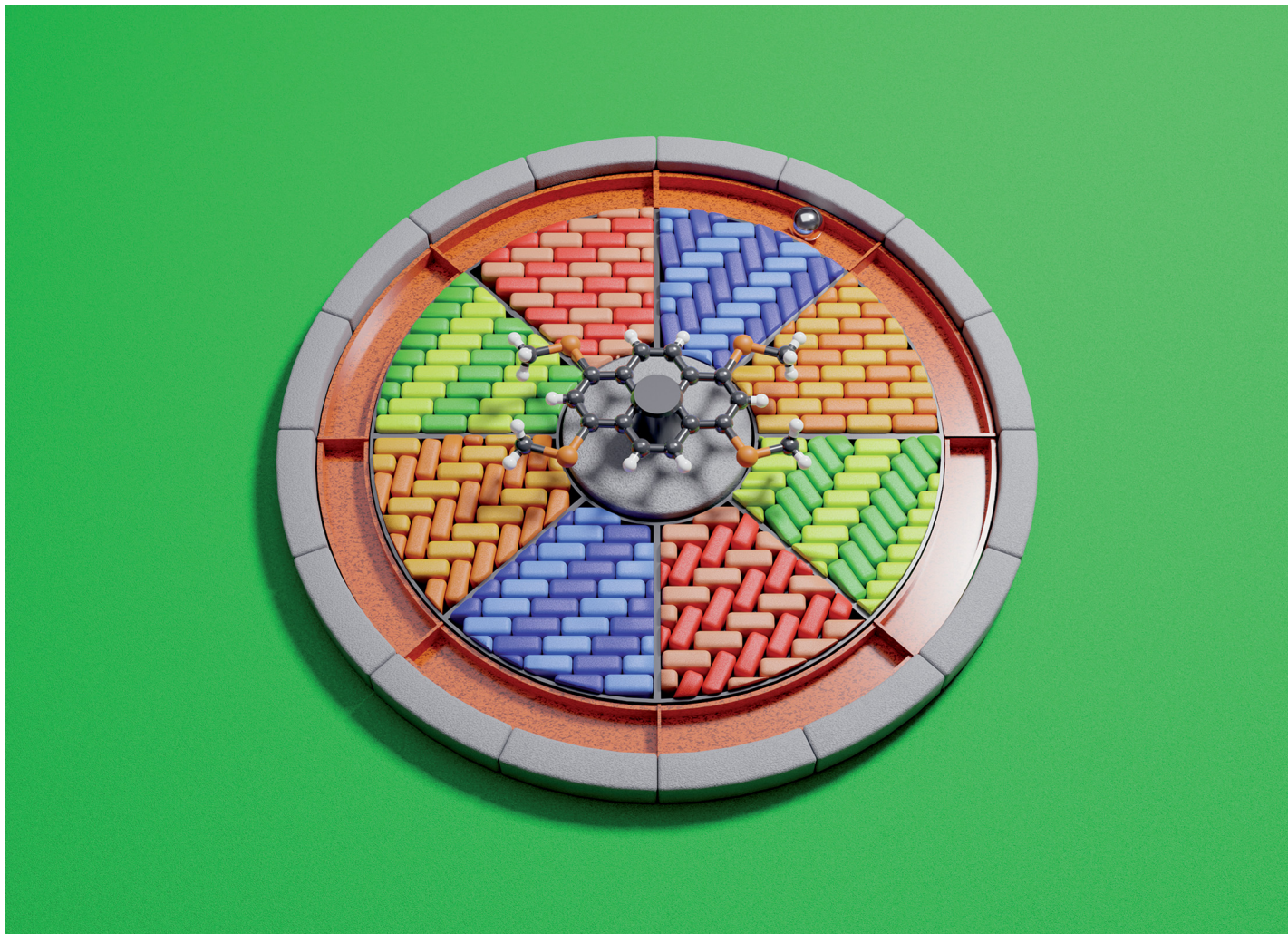


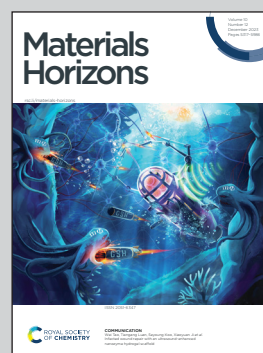


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Crystal-structure simulation of molecular semiconductors:
brickwork-related crystal structures of methylthiolated
peri-condensed polycyclic aromatic hydrocarbons

Better than prediction by chance are novel computational
simulation methods for brickwork-related crystal structures,
which allow us to evaluate the likelihood of the appearance
of different polymorphs and to find new polymorphs
experimentally.

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Crystal-structure simulation of molecular semiconductors: brickwork-related crystal structures of methylthiolated *peri*-condensed polycyclic aromatic hydrocarbons†

Kirill Bulgarevich^a and Kazuo Takimiya^{*,abc}

Despite the critical importance to carrier transport properties, studies on the control and prediction of crystal structures of molecular semiconductors have not been well-matured. To tackle this issue, we have developed “*in silico* crystallization” (ISC) protocols for simulating the brickwork (BW) crystal structures of methylchalcogenolated polycyclic aromatic hydrocarbons (PAHs). In this study, by carefully analyzing a BW-related polymorph of experimental crystal structures, an inclined brickwork (iBW) structure, we further extend the ISC protocol to simulate various BW-related crystal structures including iBW structures. Rational conditional branching in the simulation not only makes it possible to simulate

New concepts

Prediction of crystal structures of organic molecules is one of the critical issues in many fields such as the pharmaceutical industry and materials science. Focusing on the relative simplicity of brickwork (BW) crystal structures of molecular semiconductors based on methylchalcogenolated *peri*-condensed polycyclic aromatic hydrocarbons (PAHs), we have developed an “*in silico* crystallization” (ISC) protocol, in which simple calculations of bimolecular intermolecular interaction energies with the Gaussian program package and translation-based grid search to the energy minimum of molecular clusters are the key. This simple yet useful method for simulating the BW structures is further extended, by employing three conditional branching, to simulate different polymorph candidates of BW-related crystal structures. Furthermore, the calculation of the sum of intermolecular interaction energies enables us to evaluate the relative energetic stability, *i.e.*, the likelihood of the appearance, of these polymorph candidates. Although the scope of the ISC protocols is still limited at present, the concepts of the ISC protocols can be further extended to more complicated crystal structures such as herringbone and γ -structures, which would pave the way to simulate crystal structures of various molecular semiconductors prior to their actual synthesis.

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Kazuo Takimiya

It is a great honor to contribute this article to the special issue celebrating ten years of Materials Horizons. Since the launch of the journal in 2013, I have been one of the advisory board members of the journal and enjoyed the cutting-edge sciences in the field of materials science, in particular, new concepts and approaches, on which the journal mainly focuses. Congratulations on the great milestone of ten years and contribution to the community,

and I am looking forward to seeing more breakthroughs.

eight polymorph candidates of methylchalcogenolated PAHs but also helps understand the relationship between the polymorphs. Furthermore, the relative favorability of each polymorphic candidate, *i.e.*, the likelihood of the appearance among the polymorph candidates, can also be evaluated.

1. Introduction

The carrier transport properties, one of the most important properties of molecular semiconductors, are heavily dependent on both the molecular electronic structure and the crystal structure.^{1–3} In contrast to the molecular electronic structure accurately predictable by quantum chemical calculations, the prediction of the crystal structure of molecular semiconductors is still a challenging issue.^{4–7} In this relationship, the recent progress in crystal structure prediction (CPS) would help predict the crystal structure of molecular semiconductors,⁸ though

the computational methods and protocols are not easily accessible to experimental chemists. Intensive research efforts in the last two decades, on the other hand, have accumulated correlations between the molecular structure and crystal structure and thus carrier transport properties;^{9–12} the most representative one for promising transport properties is the herringbone structure, represented by pentacene,¹³ [1]benzothieno[3,2-*b*][1]benzothiophene (BTBT) derivatives,^{14,15} and so on.^{16–18} Other promising crystal structure types are the pitched π -stacking structure realized by rubrene¹⁹ and methylthiolated linear arenes,^{20–25} and the brickwork (BW) structure seen in the crystal structures of (triisopropyl)silylethynyl (TIPS)-pentacene²⁶ and its related compounds.^{27–30}

Recently, we found that the BW crystal structure can be realized by methylchalcogenolation on pyrene, the smallest *peri*-condensed polycyclic aromatic hydrocarbon (PAH).³¹ Furthermore, methylthiolated pyrene (MT-pyrene)³² showed ultra-high mobility of up to $32 \text{ cm}^2 \text{ V}^{-1} \text{ s}^{-1}$ on single-crystal field-effect transistors,³³ which reinforced the potential of the BW crystal structure for the development of further outperforming molecular semiconductors. Inspired by these results, we have developed a simple protocol for the simulation of BW crystal structures of unknown methylchalcogenolated *peri*-condensed PAHs.³⁴ This protocol, which we call “*in silico* crystallization (ISC)” helped us to select a synthetic target, and in fact, thus selected and then synthesized MT-peropyrene (Fig. 1) turned out to be another ultra-high mobility molecular semiconductor ($\mu \sim 30 \text{ cm}^2 \text{ V}^{-1} \text{ s}^{-1}$).³⁴

From these results, it is safe to say that methylchalcogenolated *peri*-condensed PAHs generally afford the BW crystal structure. In fact, recently synthesized methylthiolated perylene (MT-ptylene, Fig. 1) also crystallized into a BW structure,³⁵ and the experimental crystal structure was eventually well-matched with the simulated one by the ISC protocol (*vide infra*). Although the performance of MT-ptylene-based SC-FETs was not promising ($\mu \sim 0.2 \text{ cm}^2 \text{ V}^{-1} \text{ s}^{-1}$), the experimental studies on MT-ptylene gave insightful results on the crystal structures of methylchalcogenolated *peri*-condensed PAHs; MT-ptylene afforded a polymorph classified into an inclined brickwork (iBW) structure,³⁵ similar to that of one of the polymorphs observed for methylselenolated pyrene (MS-pyrene).³¹ Although

the relationship between the BW and iBW structures of MS-pyrene was not clear at all, the two polymorphs, BW and iBW structures, of MT-ptylene seemed to be closely related to each other, as the molecular arrangement, in other words, the mutual position of molecules, in the respective BW layers, seemed to be identical at a glance.

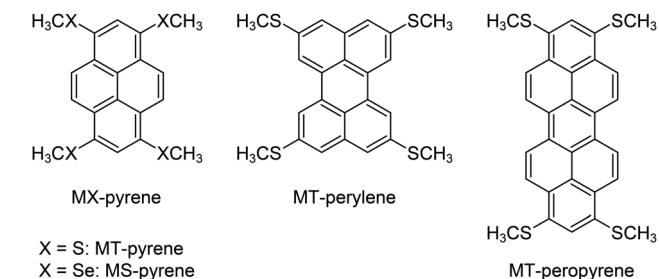
In this work, we first analyze the relationship between the BW and iBW structures of MT-ptylene by using the notation of molecules in the crystal structures developed in the ISC protocol. Then, rigorous analyses of the relationship between a BW structure and an iBW structure are carried out, and then the results are adapted to the extension of the ISC protocol to the iBW structures. Furthermore, these analyses on the iBW structure of MT-ptylene hinted to us to rationally describe the iBW structure of MS-pyrene, which further allows us to extend the ISC protocol enabling us to bring eight BW-related polymorph candidates by three conditional branches.

2. Results and discussion

2.1 BW and iBW structures of MT-ptylene

The BW and iBW crystal structures of MT-ptylene are shown in Fig. 2a and b, respectively. As the BW structure is classified into the triclinic $P\bar{1}$ space group with Z (the number of molecules in the unit cell) = 1, all the molecules in the BW crystal structure are related by the translation operation, which means that the relationship between the neighboring BW layers is also the translation (Fig. 2a bottom).³⁴ On the other hand, in the iBW structure with the monoclinic $P2_1/n$ space group, the relationship between the neighboring BW layers is the glide operation. Note that all the molecules in each BW layer in the iBW structure are related by the translation operation similar to that of the BW structure.

In the protocol of ISC simulation for the BW structure, the molecules in the Cartesian system are designated, as shown in Fig. 3; three directions of the molecule at the origin (designated as molecule O), “face”, “side”, and “end” directions, are first



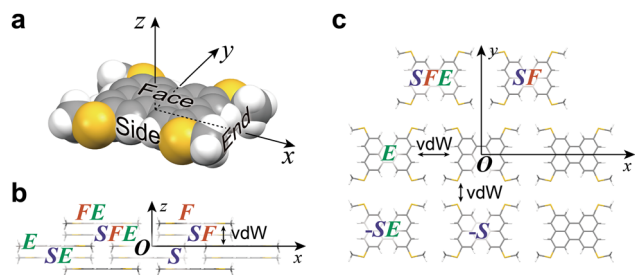


Fig. 3 (a) Definition of “face”, “side”, and “end” directions. (b) Definition of molecules in the BW layer. (c) Definition of molecules in the neighboring BW layer.

defined (Fig. 3a), and then the neighboring molecules in these directions are defined as F, S, and E, respectively. With these definitions and a combination of the definitions, all the molecules surrounding O are distinguished. For example, in each BW layer, molecule O is surrounded by molecules F, E, and FE (a molecule in the end-to-end direction from F), and their equivalents in the negative directions ($-F$, $-F-E$, and $-E$) (Fig. 3b). Similarly, in the neighboring BW layer, four molecules contacting O, namely, molecule S, (the nearest neighbor to O), SF (a molecule in the face-to-face direction from S), SFE (a molecule in the end-to-end direction from SF), and SE (a molecule in the end-to-end direction from S) are defined (Fig. 3c). Table 1 summarizes the coordinates of three molecules (F, E, and FE) in the BW layer of both the BW and iBW structures experimentally determined. Intriguingly, the coordinates of these molecules are quite similar indicating that the mutual position of molecules in the BW layers in different polymorphs are very similar to each other. It should be noted that by using the Cartesian system, the molecular arrangements in the different polymorphs can be quantitatively compared.

2.2 Relationship between the BW and iBW structures of MT-perylene

By using the above definition of molecules in the BW layer, the BW layer can be further simply expressed by two vectors, *i.e.*, $\mathbf{p} = \text{O}-\text{F}$ and $\mathbf{q} = \text{O}-\text{E}$ (Fig. 2). For constructing the three-dimensional crystal structures, these two vectors and the corresponding vectors in the neighboring layers, *i.e.*, $\mathbf{p}' = \text{S}-\text{SF}$ and $\mathbf{q}' = \text{S}-\text{SE}$, shall be identical. Obviously, the translation operation, namely, the BW structure, satisfies this condition. In addition to the translation operation, the glide operation also satisfied this condition as experimentally shown in the iBW structure of MT-perylene. In the Cartesian system, this can be expressed by a

Table 1 Cartesian coordinates of molecules (centers of mass) [Å] between the origin (O) and various neighboring molecules in the experimentally determined BW and iBW crystal structures of MT-perylene at 100 K

MT-perylene, BW structure				MT-perylene, iBW structure			
Mol.	x	y	z	Mol.	x	y	z
F	3.351	1.051	3.363	F	3.382	0.834	3.351
E	-12.787	-3.724	0.182	E	-12.775	-3.574	0.213
FE	-9.436	-2.673	3.545	FE	-9.393	-2.740	3.564

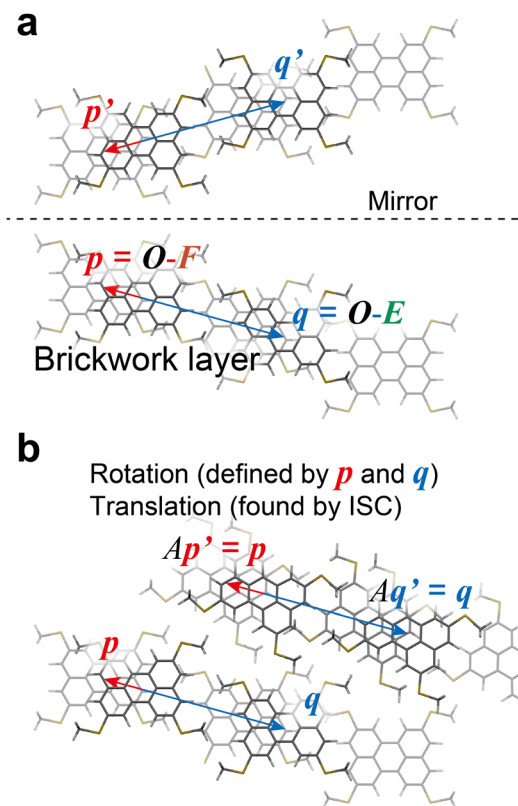


Fig. 4 Construction of the neighboring layer in the iBW structure of MT-perylene from the BW structure. The mirror plane can be chosen arbitrarily and is $y = 8.5$ Å in the present case (a). Rotation matrix A in (b) is defined in eqn (1) to ensure $A\mathbf{q}' = \mathbf{q}$ (and $A\mathbf{p}' = \mathbf{p}$).

mirror operation of the initial BW layer against any plane followed by a specific rotation that aligns the mirrored vectors back to being parallel to $\mathbf{p}$ and $\mathbf{q}$, and also translation (Fig. 4). The rotation that aligns the mirrored vectors is dependent on the mirror plane; in the case of the mirror plane parallel to the xz plane in the Cartesian system, the mirrored vectors of $\mathbf{p} = (p_x, p_y, p_z)$ and $\mathbf{q} = (q_x, q_y, q_z)$ are expressed as $\mathbf{p}' = (p_x, -p_y, p_z)$ and $\mathbf{q}' = (q_x, -q_y, q_z)$. The rotation matrix A thus shall satisfy $A\mathbf{p}' = \mathbf{p}$ and $A\mathbf{q}' = \mathbf{q}$. Without loss of generality, equations $A\mathbf{p}' = \mathbf{p}$ and $A\mathbf{q}' = \mathbf{q}$ can be solved to obtain the rotation matrix uniquely based on $\mathbf{p}$ and $\mathbf{q}$:

$$A = \frac{1}{D} \begin{pmatrix} 2r_x^2 - D & -2r_x r_y & 2r_x r_z \\ 2r_x r_y & -2r_y^2 + D & 2r_y r_z \\ 2r_x r_z & -2r_y r_z & 2r_z^2 - D \end{pmatrix} \quad (1)$$

where $(r_x, r_y, r_z) = \mathbf{p} \times \mathbf{q}$ and $D = \mathbf{p}^2 \mathbf{q}^2 - (\mathbf{p} \cdot \mathbf{q})^2$.

2.3 Simulation of the iBW structure by the ISC protocol

The original ISC protocol for the BW structures is as follows (Fig. 5): the molecules F, E, and FE are placed temporarily at van der Waals contacts with O. In step 1, the relative position of two stacking molecules to O, F and FE, is searched by minimizing the sum of intermolecular interaction energies (E_{int})³⁶ between O-F and O-FE while moving F and FE as a cluster with respect to O and by using the Gaussian16 program.³⁷ This

two-molecular (2M) stacking optimization defines a BW layer consisting of six molecules surrounding O: F, FE, and their symmetrically translated molecules, E, -F, -F-E, and -E. Next, in step 2, the BW layers are piled up in the positive side-to-side direction by optimizing the positions of S, SF, and SFE and in the negative side-to-side direction by -S, -S-F, and -S-F-E. Note that there are two ways to pile the neighboring layers up in step 2, which corresponds to polymorph candidates of the BW structures (*vide infra*). Then, molecule E and molecules related to E (FE, SE, SFE, *etc.*), which were temporarily defined in step 1, are finely optimized in step 3. These steps are repeated typically three times to determine the final BW structure with high quality (Fig. 5, ISC_{BW}). In the case of MT-erylene, the simulated BW structure nicely reproduced the experimental BW structure as in the case of MT-pyrene and MT-peropyrene (Table 2).

For the simulation of the iBW structure of MT-erylene, the final BW structure by the ISC_{BW} protocol was used; two vectors, **p** and **q**, extracted from the simulated BW structure, are used to define the rotation matrix **A** according to eqn (1). Then, three molecules in the neighboring layers, -S, SF, and SFE, are mirrored by the *xz* plane and rotated by matrix **A**, and then, as described above for step 2, the positions of these molecules optimized by searching the minimum of the sum of E_{int} s ($\sum E_{\text{int}}$) between O-S, O-SF, and O-SFE are calculated, and molecule -S is placed at the position of energy minimum

Table 2 Comparison of cell parameters between the predicted and experimental crystal structures of MT-erylene

MT-erylene	Brickwork (BW)		Inclined brickwork (iBW)	
	Simulated	Experimental ^a	Simulated	Experimental ^a
Crystal system	(Triclinic) ^a	Triclinic	(Triclinic) ^a	Monoclinic
Space group	(<i>P1</i>) ^b	<i>P1</i>	(<i>P1</i>) ^b	<i>P2₁/n</i>
<i>a</i> /Å	4.98	4.8621(3)	4.98	4.8336(2)
<i>b</i> /Å	9.17	9.3471(5)	21.93	21.9293(8)
<i>c</i> /Å	11.17	11.0114(8)	9.17	9.3596(3)
α /°	89.6	90.500(5)	89.1	90
β /°	101.8	100.650(5)	92.4	91.596(3)
γ /°	92.4	91.424(4)	91.2	90
<i>V</i> /Å ³	499.4	491.61(5)	1001.2	991.71(6)
$E_{\text{int}}(15)$ /kcal mol ⁻¹ ^c	-100.7	-107.9	-101.8	-110.5
Ref.	34	35	This work	35

^a Data obtained at *T* = 100 K. ^b Space group was not explicitly determined. ^c $E_{\text{int}}(15)$: sum of intermolecular interaction energies (E_{int}) between O and its 14 neighboring molecules.

(Fig. 5, step 2 (inclined) in ISC_{iBW}). Note that the translation operation after the mirror and rotation operations define the initial position of the molecules for searching for the energy minimum, and the glide symmetry is not assumed in the simulation.

Table S3 (ESI[†]) summarizes the centers of mass of molecules of the simulated iBW structure of MT-erylene in comparison to the experimental iBW structure. It should be emphasized that the molecular positions are virtually identical, and the root-mean-square deviation of the 15 molecules (evaluated atom-wise, RMSD₁₅) is as small as 0.27 Å, which is comparable to the search step size in the simulation.³⁸ The triclinic cell parameters extracted from the optimized molecular clusters are summarized in Table 2 (see also the crystallographic information files (CIFs); ESI[†]). Note that although the glide symmetry of the *P2₁/n* space group was not assumed, the crystallographic α and γ angles of the simulated iBW structure of MT-erylene are almost 90°, which is a characteristic of a monoclinic *P2₁/n* space group. The other crystallographic parameters including the cell axes (*a*, *b*, and *c*) and the angle (β) were also successfully reproduced. The relative stability of the polymorphs can be evaluated by lattice energy, and in the present case, we calculated the sum of intermolecular interaction energies ($E_{\text{int}}(15)$) between O and its 14 neighboring molecules, which is a similar parameter to the lattice energy.³⁹ The experimental iBW structure of MT-erylene is energetically slightly more stable than the BW structure, which was also reproduced in the present ISC simulations (Table 2).

2.4 iBW structure of MS-pyrene

As shown in Fig. 6, MS-pyrene also has an iBW structure polymorph like MT-erylene. However, the BW layers in the BW and iBW polymorphs are noticeably different. The **p** and **q** vectors in the experimental BW structure, **p**_{BW} = (5.510, 0.254, 3.581) and **q**_{BW} = (-14.316, 1.748, -0.045), do not match the **p** and **q** vectors in the iBW structure, **p**_{iBW} = (3.835, -1.653, 3.534) and **q**_{iBW} = (-14.400, 1.241, 0.183). The **p**_{BW} and **q**_{BW} in the

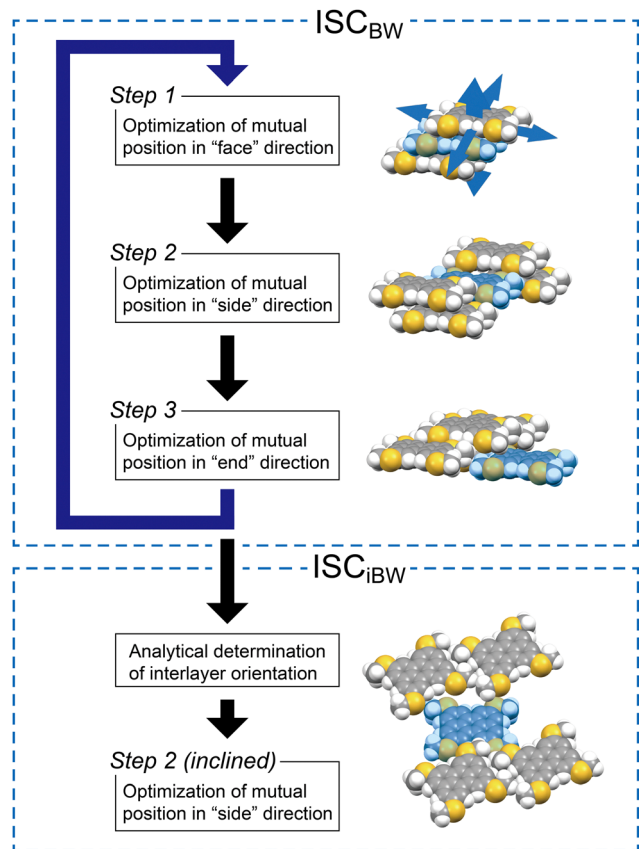


Fig. 5 Schematic representation of the ISC protocol for BW structures (ISC_{BW}) and renewed ISC covering iBW structures (ISC_{iBW}).

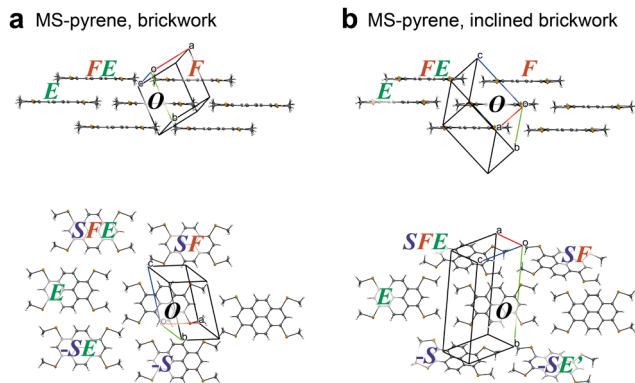


Fig. 6 Crystal structures of MS-pyrene featuring BW (a) and iBW (b) structures. The structures are shown from the side (top) and top (bottom) of the molecular plane.

simulated structure of MS-pyrene by the ISC_{BW} protocol (Fig. 5) match the $\mathbf{p}_{\text{BW}}$ and $\mathbf{q}_{\text{BW}}$, not the $\mathbf{p}_{\text{iBW}}$ and $\mathbf{q}_{\text{iBW}}$ of the experimental structures. This strongly implies that there are other ways to construct the BW layers. Close inspection of two BW layers of MS-pyrene revealed that the intermolecular interaction energy between FE and O is much smaller than that between F and O in the iBW structure. This means that the iBW structure of MS-pyrene is more one-dimensional (1D) than the BW structure. To simulate such a 1D-like stacking structure, we tested the positional search in step 1 by just using one stacking molecular pair of O and F. This resulted in an optimal position of $\mathbf{F} = (3.50, -1.50, 3.50)$, which afforded a $\mathbf{p}$ vector roughly matching $\mathbf{p}_{\text{iBW}}$. Although such a BW structure with a 1D-like stacking structure was so far not found in the actual crystals, performing the ISC with such a one-molecular (1M) stacking in step 1 results in the determination of the finalized $\mathbf{p} = (3.70, -1.60, 3.49)$ and $\mathbf{q} = (-14.59, 1.20, 0.22)$, which matches the $\mathbf{p}_{\text{iBW}}$ and $\mathbf{q}_{\text{iBW}}$ of the experimental structure with good accuracy. We then proceeded to step 2 for iBW structures. The RMSD₁₅ of the simulated structure with respect to the experimental one was as low as 0.27 Å, indicating that the iBW structure of MS-pyrene could also be simulated. Note that for MS-pyrene the simulated crystallographic α and γ angles (see Table 3) were very close to 90°, reproducing the monoclinic cell, similar to the case of MT-perylene. As already mentioned, the ISC_{iBW} protocol (as well as the ISC_{BW} protocol) does not assume the monoclinic cell, yet the resulting simulated structures are close to the monoclinic cells, indicating the utility of the protocols.

2.5 Conditional branching in the simulation of BW-related crystal structures

In step 2 of the ISC_{BW} protocol (Fig. 5), there generally exist two ways to pile the BW layers up, as already reported in the previous paper.³⁴ Fig. 7 shows the case of MT-pyrene as an example. These two structures can be distinguished by the molecular arrangement in the crystallographic bc plane: a polymorph with a molecular array that is “cleavable” along the crystallographic ac -plane, and another polymorph with a not cleavable molecular array, as the molecules are “zipped” by the methylthio groups

Table 3 Comparison of cell parameters between the predicted and experimental inclined brickwork structures of MS-pyrene (1M-ZL-iBW) and MT-peropyrene (1M-ZL-iBW)

	MS-pyrene (1M-ZL-iBW)		MT-peropyrene (1M-CL-iBW)	
	Simulated	Experimental ^a	Simulated	Experimental ^a
Crystal system	(Triclinic) ^b	Monoclinic	(Triclinic) ^b	Triclinic
Space group	($P1$) ^b	$P2_1/n$	($P1$) ^b	$P\bar{1}$
$a/\text{\AA}$	5.33	5.4709(2)	5.15	5.3250(2)
$b/\text{\AA}$	16.52	16.2641(8)	13.99	13.4178(6)
$c/\text{\AA}$	10.37	10.1056(5)	16.78	16.3151(4)
$\alpha/^\circ$	90.3	90	109.4	108.191(3)
$\beta/^\circ$	91.8	93.349(4)	90.1	91.312(3)
$\gamma/^\circ$	90.1	90	99.9	99.886(4)
$V/\text{\AA}^3$	913.2	897.65(7)	1120.9	1087.36(8)
$E_{\text{int}}(15)/\text{kcal mol}^{-1c}$	−103.6	−112.8	−123.7	−132.4
Ref.	This work	31	This work	This work

^a Data obtained at $T = 100$ K. ^b Space group was not explicitly determined. ^c $E_{\text{int}}(15)$: sum of intermolecular interaction energies (E_{int}) between O and its 14 neighboring molecules.

groups.³¹ Thus, we call the first one “cleavable layer (CL)” and the other “zipped layer (ZL)” in the interlayer structures.

In most cases, the CL polymorph is energetically more favorable than the ZL polymorph in the ISC simulations. However, the experimentally elucidated structures of MT-pyrene and 1,3,6,8-tetramethoxypyrene (MO-pyrene) are classified into the ZL polymorph. Therefore, it is important to simulate both polymorphs by the ISC protocols. Out of MO-, MT-, MS-pyrene, MT-perylene, and MT-peropyrene, the CL (ZL) polymorph was noticeably more stable for MT-perylene and MT-peropyrene (MO-pyrene). For MT- and MS-pyrene, the $E_{\text{int}}(15)$ of the CL and ZL polymorphs were competing. With these results in mind, we tried to find the CL polymorph of MT-pyrene and the ZL polymorph of MS-pyrene (*vide infra*).

To summarize the updated ISC protocols, it is now possible to simulate BW structures with one-/two-molecular (1M/2M) stacking configurations in step 1, two different interlayer configurations of the BW layers (CL/ZL) in step 2, and two different interlayer orientations of the BW layers (BW/iBW) in the final step. These three branching points can potentially result in eight BW-related crystal structures (polymorph candidates):

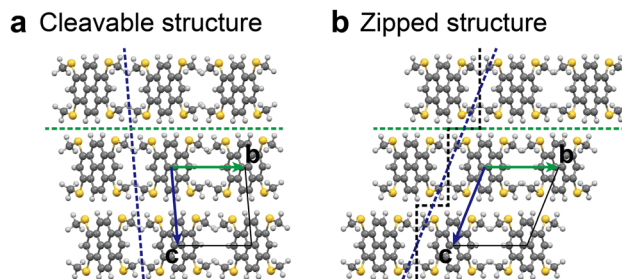


Fig. 7 Two polymorph candidates of MT-pyrene that arise in step 2 in the original ISC protocol: (a) “cleavable” structure along the crystallographic ac -plane; (b) another structure, where the molecular array is not cleavable as the molecules are “zipped” by the methylthio groups.

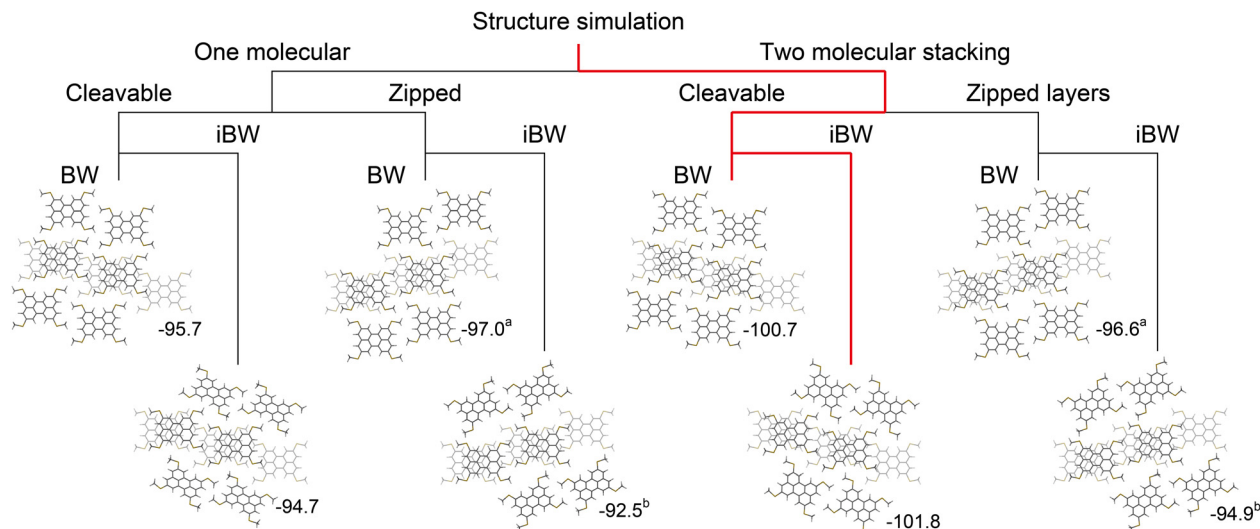


Fig. 8 Eight polymorph candidates of MT-perylene crystal structures simulated through three conditional branching. The energetically favourable two polymorphs, 2M-CL-BW and 2M-CL-iBW, were experimentally confirmed. The values are $E_{\text{int}}(15)$ (kcal mol⁻¹), and the alphabets in the superscript form denote almost identical structures, e.g., the 1M-ZL-BW and 2D-ZL-BW polymorphs are virtually identical.

e.g., 1M-CL-iBW structure, 2M-ZL-BW structure, and so on. As examples, the eight simulated structures of MT-perylene are summarized in Fig. 8 (see also ESI† for MT- and MS-pyrene, and MT-peropyrene). For MT-perylene, the two polymorph candidates, which were simulated to be the most energetically favorable, 2M-CL-BW and 2M-CL-iBW structures, were the experimentally confirmed ones.³⁵

2.6 Search for new polymorphs that were predicted to be energetically favorable

With the above-simulated crystal structures in hand, we have tried to find other polymorphs of these materials that correspond to the simulated polymorph candidates. For all materials, the crystals were grown using the physical vapor transport (PVT) method under a flow of nitrogen or argon gas.⁴⁰ For MT-peropyrene with quite high vaporizing pressure, the pressure was reduced to <20 Pa, while for other materials, PVT was performed under atmospheric pressure. By extensive search, we managed to find the 2M-CL-BW polymorph for MT-pyrene and 1M-CL-iBW polymorph for MT-peropyrene (RMSD₁₅ of corresponding simulated structures were 0.26 and 0.48 Å, respectively). Interestingly, the MT-peropyrene iBW structure, unlike those of MS-pyrene and MT-perylene, was of triclinic $P\bar{1}$ space group with two half-molecules in the asymmetric unit. The 1M-CL-iBW polymorph of MT-peropyrene experimentally found is the first example of an iBW structure without glide symmetry in the present methylchalcogenolated PAH series.

The polymorphs experimentally confirmed so far, the 2M-ZL-BW and 2M-CL-BW polymorphs of MT-pyrene, the 2M-CL-BW and 2M-CL-iBW polymorphs of MT-perylene, and the 2M-CL-BW and 1M-CL-iBW polymorphs of MT-peropyrene, are energetically favorable polymorphs among the eight polymorph candidates simulated by the ISC protocols. On the contrary, the 2M-ZL-BW polymorph of MS-pyrene was not found, although it is estimated to be the most energetically stable out of the

simulated structures. If it exists, the 2M-CL-BW polymorph of MS-pyrene is expected to show better two-dimensionality than the known 2M-CL-BW structure and thus is a more promising structure compared to the existing two polymorphs (2M-CL-BW and 1M-ZL-iBW polymorphs). Nevertheless, it can be said that the renewed ISC protocols can successfully simulate various BW-related structures and even predict the energetical stability (i.e., the likelihood of the appearance) of each polymorph to some extent.

3. Conclusions

Close inspection and analyses of the BW and iBW structures of MT-perylene experimentally found encouraged us to expand our recently developed ISC protocol for the simulation of methylchalcogenolated PAH-based molecular semiconductors. Thus, renewed ISC protocols enable us to simulate a variety of BW-related polymorph candidates through three conditional branching based on the stacking mode in the BW layer, and configuration and orientation of the interlayer structure.

It was found that MT-, MS-pyrene, MT-perylene, and MT-peropyrene all afforded two polymorphs out of (and no polymorphs outside of) the eight simulated ones. These structures include the new polymorphs of MT-pyrene and MT-peropyrene found in this work. The monoclinic nature of the MS-pyrene and MT-perylene iBW structures and the triclinic nature of the MT-peropyrene iBW structures were successfully simulated without assuming a particular symmetry in the simulations. Therefore, the method, “*in silico* crystallization”, is now a powerful tool to simulate the possible crystal structures and understand their relationships in the methylchalcogenolated PAH-based molecular semiconductors. It is also expected that this approach can be expanded to other crystal structure types such as pitched π -stack and herringbone crystal structures of molecules with planar molecular structures.

Author contributions

Kirill Bulgarevich: experiment and computational design, methodology, experiment and calculation, validation, investigation, data curation, writing – original draft. Kazuo Takimiya: validation, investigation, writing – review & editing, funding acquisition, supervision.

Conflicts of interest

There are no conflicts to declare.

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Notes and references

‡ Simulated crystal structures are available as crystallographic information format (see ESI†)

§ Experimental crystallographic data for new polymorphs of MT-pyrene (2M-CL-BW, CCDC 2256594) and MT-peropyrene (1M-CL-iBW, CCDC 2280111).†

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