

A phase retrieval algorithm for triply periodic minimal surface like structures

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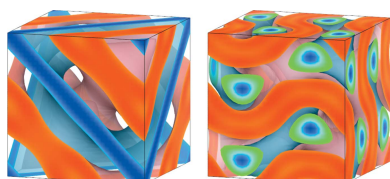
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A method to solve the crystallographic phase problem of materials with triply periodic minimal surface like structures, such as lyotropic liquid crystal bicontinuous cubic phases, is reported. In triply periodic minimal surface like structures, the difference between the maximum and minimum electron densities tends to be the smallest for the true phase combination among the possible combinations [Oka (2022). *Acta Cryst. A* **78**, 430–436]. Using this feature, a new iterative phase retrieval algorithm for structure determination was developed. The algorithm modifies electron densities outside the upper and lower thresholds in the iterative Fourier transformation process with fixed amplitudes for the structure factors, and efficiently searches for the structure with the smallest difference between the maximum and minimum electron densities. The proper structure was determined by this algorithm for all tested data for lyotropic liquid crystal bicontinuous cubic phases and mesoporous silicas. Although some cases required constraints such as the volume fraction for structure determination, more than half could be determined without any constraints, including space groups.

1. Introduction

Crystallography serves as a fundamental method to offer structural information for the understanding of materials. This statement applies to both typical, *i.e.* well ordered, crystals, and highly disordered systems such as liquid crystals. However, the application of crystallographic methods to the latter is very difficult because of the limited number of reflections available. The bicontinuous cubic phase of lyotropic liquid crystals (LLC) has a triply periodic minimal surface (TPMS) like structure and three-dimensional periodicity with natural beauty (Hyde *et al.*, 1996). Thus, these systems are good targets for structural studies. The structure of the LLC bicontinuous cubic phase was first determined by the pioneering work of Luzzati *et al.* using X-ray powder diffraction (Luzzati *et al.*, 1988; Mariani *et al.*, 1988). Recently, the present author established a single-crystallization method for the LLC bicontinuous cubic phase (Oka & Hojo, 2014), and performed structural analyses of the LLC bicontinuous cubic phases while considering model structures (Oka, 2017; Oka *et al.*, 2018, 2020). The essential difficulty is the so-called phase problem, which is still unsolved.

Numerous researchers have tackled the phase problem (Sayre, 2015). Direct methods and variants are routinely used for structure determination: it is impossible to imagine the current practice of crystallography without them (Giacovazzo, 2001). These methods are based on the general properties of crystals. The charge-flipping method, which has a simple iterative algorithm for structural determination, utilizes the



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fact that the positive electron density of atoms is concentrated in a small region, while the remaining regions have zero electron density (Oszlányi & Sütő, 2004, 2008; Palatinus, 2013). Its success implies the possibility of using the structural features of highly disordered systems if we can identify suitable expressions for the features.

In a previous paper (Oka, 2022), the author proposed two indicators reflecting the plausibility of phase combinations of experimental data for the LLC bicontinuous cubic phase. The indicators are based on the structural features of materials: the electron density tends to be constant in the direction in which liquid crystal molecules diffuse. This property suggests that the continuity of the density is a good indicator. The difference density between the maximum and minimum (I_ρ) seems to be a good and simple indicator. Another indicator (I_K), which utilizes the Hessian matrix of the electron density, is also acceptable. In the previous paper, the electron density and indicators were calculated for all possible phase combinations for the test data with centrosymmetric space groups. The result showed that the two indicators work well. Although the potential utility of the method based on the two indicators was confirmed for LLC bicontinuous structures, testing all phase combinations becomes impractical with an increase in the number of independent reflections. In addition, the method is only applicable to centrosymmetric space groups.

In this paper, an iterative phase retrieval algorithm for structure determination to overcome these difficulties is proposed. The algorithm was developed with reference to the charge-flipping method (Oszlányi & Sütő, 2008; Palatinus, 2013). It is emphasized that the iterative algorithm opens the possibility of the application to structures without centrosymmetry, which tremendously widens the search space of phase combinations. The method was tested for LLC bicontinuous cubic phases and mesoporous silicas, and structures were successfully determined for all tested data, although additional constraints were necessary in some cases. Notably, the method converged to the proper structures without information on the space group.

2. Phase retrieval method

The algorithm was designed to find the structure or structure-factor phase with the smallest difference between the maximum and minimum electron densities in a unit cell. The electron density was calculated in a unit cell of $32 \times 32 \times 32$ voxels. All calculations were performed using a home-made script in python3.

A finite number of structure-factor amplitudes $|F_{\text{obs}}(\mathbf{h})|$ are observed in crystal diffraction experiments. Unobserved amplitudes, including $|F(\mathbf{0})|$, were set to zero in the iterative process. $|F(\mathbf{0})|$ cannot be determined in principle because no zero-electron-density region is observed in the target sample. Since $F(\mathbf{0})$ is not included, the electron density, $\rho(\mathbf{r})$, in the unit cell has positive and negative values and its mean is 0.

In the iterative calculations, the initial phases were given random values in the range of $-\pi$ to π which satisfy the Friedel law. When the space group was used as a constraint on

the initial phases, the phases were assumed to be random to the extent that these satisfy the phase relations in reciprocal space (Shmueli *et al.*, 2010). The initial structure factor is as follows:

$$F(\mathbf{h}) = \begin{cases} |F_{\text{obs}}(\mathbf{h})| \exp[i\phi_{\text{random}}(\mathbf{h})], & \mathbf{h} \in \mathbf{H}_{\text{obs}} \\ 0, & \mathbf{h} \notin \mathbf{H}_{\text{obs}} \end{cases}$$

where \mathbf{H}_{obs} is the set of reciprocal-lattice vectors \mathbf{h} for which the structure factor was observed.

The following calculation steps from (i) to (iv) are repeated:

(i) $\rho(\mathbf{r})$ is calculated by determining the Fourier transform of $F(\mathbf{h})$,

$$\rho(\mathbf{r}) = \frac{1}{V} \sum_{\mathbf{h} \in \mathbf{H}_{\text{obs}}} F(\mathbf{h}) \exp(-2\pi\mathbf{h} \cdot \mathbf{r}).$$

(ii) $\rho(\mathbf{r})$ is modified as follows to obtain $g(\mathbf{r})$:

$$g(\mathbf{r}) = \begin{cases} \rho(\mathbf{r}) - (1 + k_t)[\rho(\mathbf{r}) - (\rho_{\text{shift}} + t_+)], & \rho_{\text{shift}} + t_+ < \rho(\mathbf{r}) \\ \rho(\mathbf{r}), & \rho_{\text{shift}} + t_- \leq \rho(\mathbf{r}) \leq \rho_{\text{shift}} + t_+ \\ \rho(\mathbf{r}) - (1 + k_t)[\rho(\mathbf{r}) - (\rho_{\text{shift}} + t_-)], & \rho(\mathbf{r}) < \rho_{\text{shift}} + t_- \end{cases}$$

where k_t is the flipping parameter and ρ_{shift} is the magnitude of the electron-density shift. When the volume fraction of the positive region v_p is not set, $\rho_{\text{shift}} = 0$. t_+ and t_- are the upper and lower thresholds. $\rho(\mathbf{r}) - (\rho_{\text{shift}} + t_{\text{or-}})$ in the equation is the amount above or below the threshold. The flipping parameter k_t is often set between 0 and 1. When $k_t = 0$, the density outside the upper and lower thresholds is replaced by the threshold value: $g(\mathbf{r}) = \rho_{\text{shift}} + t_{\text{or-}}$. When $k_t = 1$, it is replaced by the threshold minus the amount above or below the threshold: $g(\mathbf{r}) = \rho_{\text{shift}} + t_{\text{or-}} - [\rho(\mathbf{r}) - (\rho_{\text{shift}} + t_{\text{or-}})] = -\rho(\mathbf{r}) + 2(\rho_{\text{shift}} + t_{\text{or-}})$. The upper and lower threshold values are $t_+ = k_t\sigma_+$ and $t_- = -k_t\sigma_-$. When $\rho_{\text{shift}} = 0$, $\sigma_+ = \sigma_- = \sigma$, where σ is the standard deviation of $\rho(\mathbf{r})$. When $\rho_{\text{shift}} \neq 0$, σ_+ is the root-mean-square of $\rho(\mathbf{r}) - \rho_{\text{shift}}$ in the region $\rho(\mathbf{r}) > \rho_{\text{shift}}$, and σ_- is the root-mean-square of $\rho(\mathbf{r}) - \rho_{\text{shift}}$ in the region $\rho(\mathbf{r}) < \rho_{\text{shift}}$. $\rho_{\text{shift}} \neq 0$ only when v_p is set. v_p is the volume fraction in the unit cell where $\rho(\mathbf{r}) > \rho_{\text{shift}}$ and takes values between 0 and 1. The threshold parameter k_t is often set from 0.2 to 1.3.

(iii) $g(\mathbf{r})$ is Fourier transformed to obtain the structure factor, $G(\mathbf{h}) = |G(\mathbf{h})| \exp[i\phi_G(\mathbf{h})]$.

(iv) A new structure factor is obtained as follows:

$$F(\mathbf{h}) = \begin{cases} |F_{\text{obs}}(\mathbf{h})| \exp[i\phi_G(\mathbf{h})], & \mathbf{h} \in \mathbf{H}_{\text{obs}} \\ 0, & \mathbf{h} \notin \mathbf{H}_{\text{obs}} \end{cases}$$

When $k_t = 1$ and $\rho_{\text{shift}} = 0$, the upper and lower threshold values t_+ and t_- are σ , the standard deviation of $\rho(\mathbf{r})$. Thus, most final structures have electron densities outside the upper and lower thresholds. When the parameters k_t and k_p were fixed in the calculation, the rate of obtaining the correct solution was low. Therefore, during the calculation cycle, they were changed. In the j th cycle, the parameters were as follows: $k = \bar{k} + \Delta k \cos(2\pi j/n)$, where \bar{k} , Δk and n are the mean, width and period of a parameter k , respectively. Different

Table 1
Data used in structure determination.

In the last column (volume fraction), the value is the volume fraction of the low-electron-density region in the LLC type II, while it is the volume fraction of the high region in the other samples.

| Classification | Sample | TPMS | Space group | Lattice constant (nm) | Min. interplanar distance (nm) | Independent reflections | Volume fraction (TPMS side) |
|---|---|----------|----------------------|-------------------------------|--------------------------------|-------------------------|-----------------------------|
| LLC type II | Monoolein/water (Oka, 2017) | P | $Im\bar{3}m\uparrow$ | 14.19 | 2.90 | 12 | 0.43 |
| | | D | $Pn\bar{3}m\uparrow$ | 11.26 | 2.40 | 14 | 0.44 |
| | | G | $Ia\bar{3}d\uparrow$ | 14.65 | 2.87 | 8 | 0.54 |
| | Phytantriol/water (Oka <i>et al.</i> , 2018) | D | $Pn\bar{3}m\uparrow$ | 6.474 | 1.202 | 21 | 0.57 |
| | | G | $Ia\bar{3}d\uparrow$ | 8.748 | 1.169 | 21 | 0.66 |
| | | G | $Ia\bar{3}d\uparrow$ | 11.32 | 1.512 | 21 | 0.72 |
| LLC type I Mesoporous silica | $C_{12}EO_6$ /water (Oka <i>et al.</i> , 2020) | G | $Ia\bar{3}d\uparrow$ | 9.6 | 1.35 | 18 | 0.412 |
| | AMS-10 (Gao <i>et al.</i> , 2006) | D | $Pn\bar{3}m\uparrow$ | 23.8 | 2.62 | 25 | 0.283 |
| | $EO_{20}PO_{70}EO_{20}$ templated (Sakamoto <i>et al.</i> , 2004) | G | $Ia\bar{3}d\uparrow$ | 9.661 | 1.030 | 39 | 0.25 |
| | MCM-48 (A-SY) (Solovyov <i>et al.</i> , 2005) | G | $Ia\bar{3}d\uparrow$ | 9.661 | 1.030 | 39 | 0.25 |
| | IBN-9 (Zhang <i>et al.</i> , 2011) | H | $P6_3/mcm\uparrow$ | $a = b = 8.84,$ $c = 8.43$ | 1.91 | 12 | 0.387 |
| PEO_{117} -b- PS_{77} -PtBA $_{179}$ templated (Cao <i>et al.</i> , 2016) | G | $I4_132$ | 70 | 6.2 | 50 | 0.3‡ | |

† Centrosymmetric space groups. ‡ Volume fraction of silica on one gyroid network region. The other gyroid network region (including TPMS) is void.

values were set for n_f and n_p so that the periods of parameter changes of k_f and k_p do not coincide.

If the structure is known to have centrosymmetry, it is possible to add the constraint that the structure factor be real. In this case, the structure factor in step (iv) becomes

$$F(\mathbf{h}) = \begin{cases} |F_{\text{obs}}(\mathbf{h})| \frac{\text{Re}(\exp[i\phi_G(\mathbf{h})])}{|\text{Re}(\exp[i\phi_G(\mathbf{h})])|}, & \mathbf{h} \in \mathbf{H}_{\text{obs}} \\ 0, & \mathbf{h} \notin \mathbf{H}_{\text{obs}} \end{cases}$$

where $\text{Re}()$ is a function that extracts the real part of the value in the parentheses. The progress of the iterative calculation can be monitored by the difference between the maximum and minimum electron densities in the unit cell, $I_\rho = \max\{\rho(\mathbf{r})\} - \min\{\rho(\mathbf{r})\}$ (Oka, 2022). Calculation steps are repeated a set number of iterations, and the final structure is the one with the minimum I_ρ during the calculation. Multiple independent calculations may produce different results. Also, the origin-shifted structures are outputted even if the structures are completely equivalent.

The output results were evaluated using I_ρ and I_K . I_K , an indicator based on the convexity of the electron density, is described in detail in the previous paper (Oka, 2022) and briefly here. In materials with TPMS-like structures, convex regions, *i.e.* regions with closed isoelectron-density surfaces, are considered to be small. The convex regions can be determined by the eigenvalues of the Hessian matrix of the electron density, $\rho(\mathbf{r})$:

$$\mathbf{H}(\mathbf{r}) = \begin{pmatrix} \rho_{xx} & \rho_{xy} & \rho_{xz} \\ \rho_{yx} & \rho_{yy} & \rho_{yz} \\ \rho_{zx} & \rho_{zy} & \rho_{zz} \end{pmatrix}$$

where the subscripts indicate partial derivatives. If the signs of the eigenvalues of the Hessian matrix are all the same, then

the region is strictly convex (Rockafellar & Wets, 2010). Let C be the electron-density regions that are strictly convex. Then, the indicator I_K is defined by $I_K = \int_C |\det[\mathbf{H}(\mathbf{r})]| \, d\mathbf{r}$. Both indicators have been shown to be useful in the structure determination of LLC bicontinuous cubic phases (Oka, 2022). When both of these indicators are small and close values are obtained in several independent calculations, it can be presumed that the proper structure is found.

The phases of the outputted structure can be compared with that of the true structure by the following R_p value (Oka, 2022):

$$R_p = \frac{\sum_{\mathbf{h} \in \mathbf{H}_{\text{obs}}} |F_{\text{obs}}(\mathbf{h})| |\Delta\phi(\mathbf{h})|}{\sum_{\mathbf{h} \in \mathbf{H}_{\text{obs}}} |F_{\text{obs}}(\mathbf{h})| \left(\frac{\pi}{2}\right)}$$

where $\Delta\phi(\mathbf{h}) = \phi_{\text{true}}(\mathbf{h}) - [\phi_{\text{output}}(\mathbf{h}) + 2\pi\mathbf{r}_{\text{shift}} \cdot \mathbf{h}] [-\pi < \Delta\phi(\mathbf{h}) \leq \pi]$. The origin shift $\mathbf{r}_{\text{shift}}$ of the outputted structure was obtained by minimization of $\sum_{\mathbf{h} \in \mathbf{H}_{\text{obs}}} |F_{\text{obs}}(\mathbf{h})|^2 \sin^2[\Delta\phi(\mathbf{h})/2]$. R_p approaches 0 as the phase agreement becomes higher. According to Babinet's principle, when the electron density of a sample is inverted, the diffraction intensity is equivalent to that before the inversion. For this reason, a structure and a density-inversed structure are equivalent, and the smaller R_p of the two structures is adopted as the R_p of the structure. Thus, R_p is a value between 0 and 1.

3. Structure determination examples

3.1. Phase retrieval with a centrosymmetric space-group constraint

Table 1 lists the 11 data sets used in the structure determination. The six LLC bicontinuous cubic phases correspond to X-ray diffraction data from single crystals we have measured previously (Oka, 2017; Oka *et al.*, 2018, 2020). These

Table 2

Structure determination results using centrosymmetric space groups as initial random phase constraints.

Structure factors are restricted to real numbers during the calculation in all structure determinations. The penultimate column gives the number of structures out of the 100 obtained that satisfy $R_p < 0.1$.

| Sample | Space group | $\bar{k}_i \pm \Delta k_i (n_i)$ | $\bar{k}_f \pm \Delta k_f (n_f)$ | Iterations | Additional constraint | $R_p < 0.1$ | Min. R_p |
|-----------------------------------|--------------|----------------------------------|----------------------------------|------------|-----------------------|-------------|------------|
| Monoolein/water | $Im\bar{3}m$ | 0.25 ± 0.25 (17) | 0.6 ± 0.4 (13) | 200 | – | 100† | 0 |
| | $Pn\bar{3}m$ | 0.75 ± 0.25 (17) | 0.6 ± 0.4 (13) | 200 | – | 100† | 0.080 |
| | $Ia\bar{3}d$ | 0.75 ± 0.25 (17) | 0.75 ± 0.25 (13) | 200 | – | 100† | 0 |
| Phytantriol/water | $Pn\bar{3}m$ | 0.75 ± 0.25 (17) | 0.75 ± 0.25 (13) | 200 | – | 100† | 0.015 |
| | $Ia\bar{3}d$ | 0.75 ± 0.25 (17) | 0.75 ± 0.25 (13) | 200 | – | 100 | 0 |
| $C_{12}EO_6$ /water | $Ia\bar{3}d$ | 0.25 ± 0.25 (17) | 0.75 ± 0.25 (13) | 200 | $v_p = 0.75$ | 19 | 0.005 |
| AMS-10 | $Pn\bar{3}m$ | 0.75 ± 0.25 (17) | 0.75 ± 0.25 (13) | 200 | – | 100† | 0.003 |
| $EO_{20}PO_{70}EO_{20}$ templated | $Ia\bar{3}d$ | 0.75 ± 0.25 (17) | 0.75 ± 0.25 (13) | 200 | – | 100 | 0.018 |
| MCM-48 | $Ia\bar{3}d$ | 0.75 ± 0.25 (29) | 0.6 ± 0.4 (19) | 700 | $v_p = 0.25$ | 100 | 0 |
| IBN-9 | $P6_3/mcm$ | 0.75 ± 0.25 (17) | 0.75 ± 0.25 (13) | 200 | – | 100† | 0.056 |

† Only one solution was obtained.

are considered to be accurate with regard to the phase of the structure factors, which is obtained by optimizing the model to the X-ray diffraction data. As examples other than the LLC bicontinuous cubic phases, the structures of mesoporous silicas were determined. Four of the mesoporous silica data sets were obtained using high-resolution transmission electron microscopy (Sakamoto *et al.*, 2004; Gao *et al.*, 2006; Zhang *et al.*, 2011; Cao *et al.*, 2016), and the structure factors, including phase, are considered reliable. The data for MCM-48 were obtained by powder X-ray diffraction at a synchrotron radiation facility, and the phase of the structure factor was obtained by model optimization (Solovyov *et al.*, 2005). Therefore, the phases are reliable.

Materials with TPMS-like structures generally have large structural disorder, and the spatial resolution of the data obtained is lower than for solid crystals. Also, due to their high symmetry, the number of independent reflections is not large (Table 1). In the type-I LLC bicontinuous cubic phase, polar regions with high electron density gather on the TPMS and non-polar regions with low electron density gather on the network side; in type II, the positions of polar and non-polar regions are opposite to those in type I (Hyde *et al.*, 1996). In all mesoporous silicas in Table 1, with one exception, the silica walls are located on the TPMS and the network sides are vacant. Only PEO₁₁₇-b-PS₇₇-PtBA₁₇₉ templated mesoporous silica has a single-gyroid structure with silica located on only one of the two srs nets, the other being vacant (Cao *et al.*, 2016). Therefore, this structure is chiral and not centrosymmetric, whereas all the other structures are centrosymmetric. The volume fraction of the TPMS side of each sample is shown in Table 1.

First, the structural determination was tried using the constraint of centrosymmetry and using as few other constraints as possible. The initial phases were set to random values within the range satisfying the phase relationship in each centrosymmetric space group (Shmueli *et al.*, 2010). Since the space groups are centrosymmetric, the constraint that the structure factor be real in the iterative processes was used. A structure lacking centrosymmetry was not tried here because of the complexity of the initial constraints derived from the

space group. The results of the structure determinations are summarized in Table 2. For eight data sets, out of 100 independent calculations without additional constraints, structures with sufficiently small R_p were obtained for all of them. The R_p were calculated between the obtained structures and the previously determined true structures.

Fig. 1(a) shows the changes in the indicator I_ρ and the parameters k_t and k_f during the iterative calculation process of

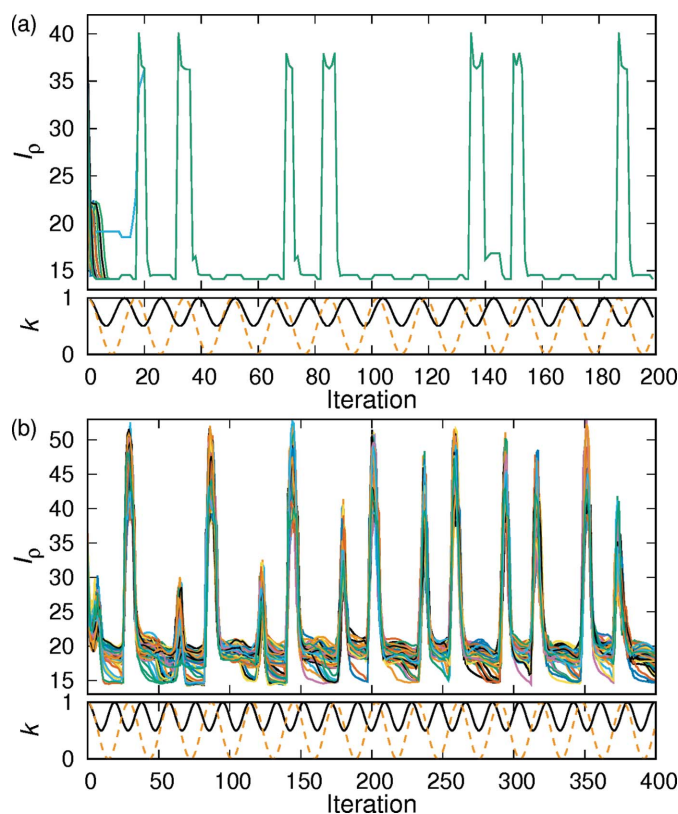


Figure 1 Changes of the indicator I_ρ and the parameters k_t and k_f during the calculation of structure determination in phytantriol $Pn\bar{3}m$. Parameter settings are listed in Table 2 (a) and Table 3 (b). The tops of the figures show the changes in I_ρ for 100 independent calculations in various colors. At the bottom of the figures, the changes in k_t are shown as a solid black line and k_f as a dashed orange line.

structure determination in the data of phytantriol $Pn\bar{3}m$. I_ρ was useful as an indicator to show the progress of structure determination, since I_ρ appeared to be minimized in the iterative process. The top part of the figure shows the change in I_ρ over 100 independent calculations. The I_ρ values for those 100 independent calculations converge to two value traces when the number of iterations is less than 10, and to a single trace at about 20 iterations. Thereafter, the value of I_ρ may increase temporarily in response to changes in the parameters k_f and k_t , but it remains close to the lowest value in many regions. Although the value of I_ρ fluctuated during the iterative calculations, the structure with minimum I_ρ was adopted as the final solution. The lower part of the figure shows changes in the parameters k_f and k_t , which were changed periodically in the iterative process. The reason for the periodic changes of both parameters was to perform

calculations with different combinations of parameters and to avoid staying in local minima. Fixing k_f and k_t in the iterative process often resulted in convergence to several different local minima, which frequently did not result in the proper structure.

The I_ρ and I_K values for the structures obtained from the phytantriol $Pn\bar{3}m$ and $Ia\bar{3}d$ data are shown in Figs. 2(a) and 2(b). The true and obtained structures have very close or coincident I_ρ and I_K values. The previous study showed that the I_ρ and I_K values of the true structure have values close to the minimum (Oka, 2022). The structure obtained in this study [Fig. 2(a)] is consistent with the structure with the minimum values of both I_ρ and I_K previously obtained for phytantriol $Pn\bar{3}m$. The two obtained structures in phytantriol $Ia\bar{3}d$ [Fig. 2(b)] also agree with the first and second minimum structures previously obtained.

Each data set for monoolein in three space groups showed a single final structure (Table 2). For $Im\bar{3}m$ and $Ia\bar{3}d$, the R_p value was 0, which is in perfect agreement with the true structure. On the other hand, $Pn\bar{3}m$ had $R_p = 0.080$, the highest minimum R_p among the data used in this study. This is probably due to the fact that the true structure in $Pn\bar{3}m$ is I_K -minimal but not I_ρ -minimal, as shown in a previous paper (Oka, 2022). The mesoporous silica, except for MCM-48, yielded a final solution close to the true structure (Table 2).

For $C_{12}EO_6$ and MCM-48, the proper structure could not be obtained without additional constraints. The proper structure was obtained when the parameter v_p was set. When $v_p = 0.75$ was used in $C_{12}EO_6$, structures close to the true structure were obtained 19 times out of 100 independent calculations. Fig. 2(c) shows the distribution of I_ρ and I_K for the 100 structures obtained from the $C_{12}EO_6$ calculation; the 19 indicator points [overlapped by a single point in Fig. 2(c)] almost overlap with the point of the true structure. However, the I_ρ and I_K of the other points are widely distributed, but many of them have smaller I_ρ than the I_ρ of the true structure. Many points are also distributed around the indicator values of the structure determined without using the v_p constraint. This result is consistent with a previous study showing that the I_ρ of the true structure is not minimal in $C_{12}EO_6$. Therefore, I_ρ cannot be used as an indicator of the true structure in $C_{12}EO_6$. On the other hand, as shown in previous studies, I_K is valid as an indicator, and the structure with the minimum I_K among the obtained structures is close to the true structure. The true structure was always obtained when $v_p = 0.25$ in MCM-48. The I_ρ of the obtained structure was larger than that when v_p was not set, but I_K was smaller (data not shown). It has been previously shown that when the volume fraction of the TPMS side is above 0.7, I_ρ is less valid as an indicator of structure and I_K is superior to I_ρ (Oka, 2022). These examples suggest that if the volume fraction of the high (or low) electron-density region is far from 0.5, it is better to set an additional constraint v_p . It is also better to focus on I_K rather than I_ρ in determining the final structure when v_p needs to be set. A reminder about v_p here: v_p determines the zero position of the electron density when calculating the standard deviation, and is one of the parameters similar to k_f and k_t . It is recommended to set a

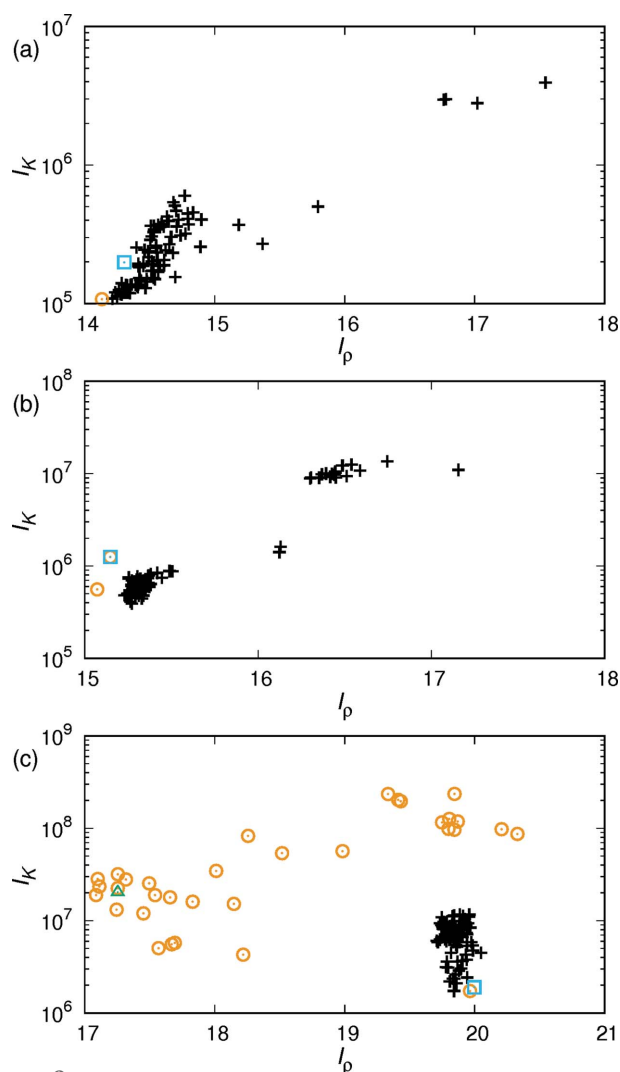


Figure 2

The scatter plot of I_ρ and I_K for the structures obtained from 100 independent calculations in phytantriol $Pn\bar{3}m$ (a) and $Ia\bar{3}d$ (b) and $C_{12}EO_6$ (c). The orange circles were obtained with the parameters in Table 2, and the black crosses were obtained with those in Table 3. The sky-blue squares are the values of the true structures obtained previously. The green triangle in (c) was obtained with the parameters in Table 2 other than v_p .

Table 3

Structure determination results without using the space group as a constraint on the initial phase.

The penultimate column gives the number of structures out of the 100 obtained that satisfy $R_p < 0.1$.

| Sample | Space group | $\bar{k}_f \pm \Delta k_f (n_f)$ | $\bar{k}_i \pm \Delta k_i (n_i)$ | Iterations | Additional constraint | $R_p < 0.1$ | Min. R_p |
|---|--------------|----------------------------------|----------------------------------|------------|---------------------------------|-------------|------------|
| Monoolein/water | $Im\bar{3}m$ | $0.5 \pm 0.5 (29)$ | $0.75 \pm 0.25 (19)$ | 700 | Real F^\dagger | 96 | 0 |
| | $Pn\bar{3}m$ | $0.5 \pm 0.5 (29)$ | $0.6 \pm 0.4 (19)$ | 700 | Real F^\dagger | 60 | 0.065 |
| | $Ia\bar{3}d$ | $0.5 \pm 0.5 (29)$ | $0.75 \pm 0.25 (19)$ | 700 | – | 76 | 0.001 |
| Phytantriol/water | $Pn\bar{3}m$ | $0.5 \pm 0.5 (29)$ | $0.75 \pm 0.25 (19)$ | 700 | – | 56 | 0.042 |
| | $Ia\bar{3}d$ | $0.5 \pm 0.5 (29)$ | $0.75 \pm 0.25 (19)$ | 700 | – | 80 | 0.021 |
| $C_{12}EO_6$ /water | $Ia\bar{3}d$ | $0.5 \pm 0.5 (17)$ | $0.75 \pm 0.25 (13)$ | 400 | $v_p = 0.75$ | 100 | 0.041 |
| AMS-10 | $Pn\bar{3}m$ | $0.5 \pm 0.5 (29)$ | $0.6 \pm 0.4 (19)$ | 700 | – | 92 | 0.002 |
| $EO_{20}PO_{70}EO_{20}$ templated | $Ia\bar{3}d$ | $0.5 \pm 0.5 (29)$ | $0.75 \pm 0.25 (19)$ | 700 | – | 92 | 0.020 |
| MCM-48 | $Ia\bar{3}d$ | $0.5 \pm 0.5 (29)$ | $0.65 \pm 0.35 (19)$ | 7000 | Real F^\dagger , $v_p = 0.25$ | 72 | 0 |
| IBN-9 | $P6_3/mcm$ | $0.5 \pm 0.5 (29)$ | $0.75 \pm 0.25 (19)$ | 700 | – | 56 | 0.060 |
| PEO ₁₁₇ -b-PS ₇₇ -PtBA ₁₇₉ templated | $I4_132$ | $0.5 \pm 0.5 (29)$ | $0.65 \pm 0.35 (19)$ | 7000 | $v_p = 0.40$ | 96‡ | 0.050 |

† Structure factors are restricted to real numbers during the calculation. ‡ Sum of two chiral (mirror image) structures.

value close to the real volume fraction, but it is not necessary to set it to the same value. It would be better to adjust v_p based on the results of iterative calculations.

3.2. Phase retrieval without a space-group constraint

Next, the structure was determined without using the space-group constraint and with as few other constraints as possible. The results are summarized in Table 3. Under the condition of no constraints, proper structures with $R_p < 0.1$ were obtained with more than half probability for six data sets. To obtain the proper structures in the other data, constraints were necessary.

An example of data for which a proper structure was obtained under completely unconstrained conditions is shown for phytantriol $Pn\bar{3}m$. The change in I_ρ during the iterative process is shown in Fig. 1(b). The value traces of I_ρ repeatedly increase and decrease, but some of them reach a minimum value around $I_\rho = 14$ after a large increase in I_ρ . Calculations with fixed parameters k_f and k_p were also tried: I_ρ was often trapped at a certain value and I_ρ did not decrease further even if the number of calculations was increased. On the other hand, when k_f and k_p were changed periodically, the search for a solution was efficient. In addition, the calculation time was acceptable for the current condition. Therefore, the parameters k_f and k_p were changed periodically. The data of phytantriol $Pn\bar{3}m$ yielded 56 structures with $R_p < 0.1$ in 100 independent calculations over 700 iterations (Table 3). Scatter plots of I_ρ and I_K are shown in Fig. 2(a). The values of the obtained structures are distributed around that of the true structure. The distribution of the obtained values appears to be in the process of converging to the value of the structure using the space group as a constraint. When the number of iterations was set to 2800, the number of structures with $R_p < 0.1$ increased to 92 out of 100 calculations. If the number of iterations is set to a larger value, the distribution will probably converge to one or a few points. For phytantriol $Ia\bar{3}d$, 80 of 100 independent calculations resulted in $R_p < 0.1$, with a minimum R_p of 0.021 (Table 3). The scatter plot, Fig. 2(b), shows that the I_ρ and I_K distributions for the final structure are concentrated near the true structure.

For the monoolein $Im\bar{3}m$ and $Pn\bar{3}m$ data, the structure factor had to be constrained to a real number because without the constraint the proper structure could not be obtained. Compared with the other data, the result of monoolein may have been affected by the small number of independent reflections. On the other hand, monoolein $Ia\bar{3}d$, with the smallest number of independent reflections, converged to a nearly true structure without constraints. The ease of convergence may differ for each space group.

For the volume fraction of the LLC bicontinuous cubic phase, $C_{12}EO_6$ has the largest value of 0.72 (Table 1). A volume fraction constraint was necessary for $C_{12}EO_6$. When calculated with $v_p = 0.75$, the distribution of I_ρ and I_K for the obtained structure is shown in Fig. 2(c). The values of the obtained structures are distributed around that of the true structure. Also, under this condition, all calculations yielded structures with $R_p < 0.1$. In the calculation with $v_p = 0.72$, the structure with $R_p < 0.1$ was obtained in 60 out of 100 independent calculations (Table 3). Better results were obtained for $C_{12}EO_6$ by changing v_p slightly from the actual volume fraction.

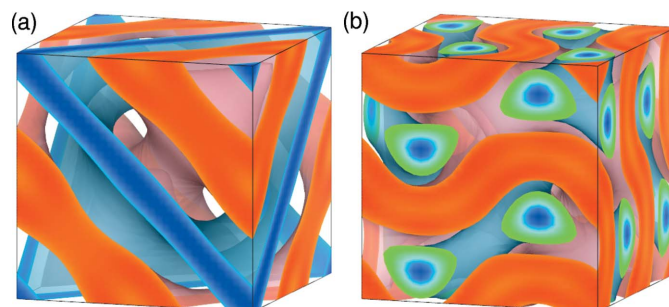


Figure 3 Electron densities of (a) phytantriol/water ($Pn\bar{3}m$) and (b) $C_{12}EO_6$ with the minimum I_K among those calculated under the conditions in Table 3. They are translationally shifted so that the origins of the unit cells are the same as those of the expected space groups. Isodensity surfaces are drawn with a volume fraction of (a) 0.25 and (b) 0.55 on high-density sides (pale red) and 0.25 on low-density sides (pale blue). In the cross section, the highest electron-density regions are depicted in red and the lowest in blue. Electron densities were drawn using VESTA (Momma & Izumi, 2011).

The three mesoporous silicas, AMS-10, EO₂₀PO₇₀EO₂₀ templated and IBN-9, yielded final structures close to the true structure without constraints. The volume fractions of silica in these structures ranged from 0.283 to 0.412, relatively close to 0.5 compared with the value of 0.25 for MCM-48 (Table 1). MCM-48 required two constraints: a real number for the structure factor and a set volume fraction. When the number of iterations was 700, only five out of 100 independent calculations yielded a structure with $R_p < 0.1$. Then, when the number of iterations was set to 7000, the number increased to 72 (Table 3). It seems that as iterations are increased, the number of final structures that are close to the true structure increases. For PEO₁₁₇-b-PS₇₇-PtBA₁₇₉ templated, 100 independent calculations yielded 96 structures with $R_p < 0.1$. Since the structure is chiral, this value includes the mirror image structure; excluding the mirror image, the value is 45. This calculation required the constraint of v_p ; the best I_K values for the final structures were obtained when $v_p = 0.40$. The reason for the necessity of the v_p constraint is probably due to the large number of independent reflections.

The obtained electron densities of phytantriol/water ($Pn\bar{3}m$) and C₁₂EO₆ are shown in Fig. 3. The electron densities are the translationally shifted ones obtained by phase retrieval. These are in good agreement with the previously determined electron densities (Figs. S1 and S2 in the supporting information), although the space group was not used.

Without the use of space groups, nearly true structures were obtained without any constraints in more than half of the cases. Even when iteration did not work without constraints, the constraints of a real structure factor or set v_p yielded a nearly true structure. Judging from this result, this structure determination method can be used for a TPMS-like structure even when the space group is not known.

In this paper, the electron density was calculated in a unit cell of $32 \times 32 \times 32$ voxels. The minimum grid spacing required for the charge-flipping method is $d_{\min}/2$ (Oszlányi & Sütő, 2004), where d_{\min} is the minimum interplanar distance. Thus, the minimum required number of divisions per one edge of the unit cell is $d_{\min}/2a$, where a is a lattice constant. In the data used in this paper, the minimum required number is 10–22. Since a small number of divisions results in a poor representation of the electron density and affects the values of the indicators I_ρ and I_K , 32 divisions were used.

4. Conclusion

An algorithm for determination of TPMS-like structures from diffraction data was developed and shown to be effective. Although additional constraints were necessary in some cases, structure determination was achieved for all tested data sets. The obtained structures were close to the true structure, and R_p , which indicates phase difference, was close to zero. The algorithm appears to efficiently find the structure with the smallest I_ρ by cutting or inverting the electron density outside the upper and lower thresholds. Constraints on v_p were necessary when the volume fraction was far from 0.5 and when

the number of independent reflections was large. For some parameter settings, several different structures were obtained, but by using I_K in conjunction with I_ρ , the proper structure could be determined. Previous studies have shown that I_K is superior to I_ρ in determining TPMS-like structures. Therefore, when multiple structure candidates are obtained, I_K should also be used as a reference.

The determination was more efficient when information on centrosymmetric space groups was used. Therefore, when using this algorithm to determine unresolved structures, it appears better to use known information on space groups. On the other hand, this is not essential in structure determination. Thus, it is possible to determine a structure for which the space group is unknown. The number of independent reflections in the data sets for the space group $Ia\bar{3}d$ was 8 for monoolein and 39 for MCM-48, so the relative spatial resolution was low for monoolein and high for MCM-48. Thus, the method appears to be applicable independently of spatial resolution and the number of independent reflections. The previously reported method requires a long computation time when the number of independent reflections is large, so the iterative method reported here should be used in such cases.

Although the LLC bicontinuous cubic phase and mesoporous silica were used here, this method could be applied to TPMS-like structures such as thermotropic liquid crystals and polymers. If there exists a three-dimensional periodic structure with bicontinuous or polycontinuous regions different from the TPMS-like structure, the method may be applicable to such a system as well.

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