A common lines approach for ab-initio modeling of molecules with tetrahedral and octahedral symmetry

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Abstract

A main task in cryo-electron microscopy single particle reconstruction is to find a three-dimensional model of a molecule given a set of its randomly oriented and positioned noisy projection-images. In this work, we propose an algorithm for ab-initio reconstruction for molecules with tetrahedral or octahedral symmetry. The algorithm exploits the multiple common lines between each pair of projection-images as well as self common lines within each image. It is robust to noise in the input images as it integrates the information from all images at once. The applicability of the proposed algorithm is demonstrated using experimental cryo-electron microscopy data.

1 Introduction

Cryo-electron microscopy (cryo-EM) is a method for determining the high-resolution three-dimensional structure of biomolecules [5]. The method involves imaging frozen copies of the investigated molecule by an electron-microscope, with each copy assuming some unknown random orientation fixed at the moment of freezing. Due to the low electron dose that can be applied to the imaged molecules, the projection-images produced by cryo-EM are very noisy. Once the orientation of each of the imaged copies of the molecule has been determined, a low-resolution ab-initio model of the molecule may be recovered from the acquired projection-images by tomographic reconstruction algorithms. An accurate ab-initio model is crucial for obtaining a high-resolution model, which is determined by iterative procedures from the set of raw input projection-images. The task of finding the orientation of the molecule giving rise to each projection-image is known as the “orientation assignment problem”, and is the main objective of this work.

Formally, if we denote the electrostatic potential of the molecule by $\psi : \mathbb{R}^3 \to \mathbb{R}$, and consider a set of $N$ rotation matrices

$$R_i = \begin{pmatrix} R_{i1} & R_{i2} & R_{i3} \\ \end{pmatrix} \in SO(3), \quad i \in [N] = \{1, \ldots, N\},$$

(1)

where $SO(3)$ is the group of all rotations in $\mathbb{R}^3$, then the projection-image $P_{R_i}, i \in [N]$, that was generated by imaging $\psi$ rotated by $R_i$, is given by the line integrals of $\psi$ along the lines parallel to $R_{i3}$ (the third column of $R_i$), namely

$$P_{R_i}(x, y) = \int_{-\infty}^{\infty} \psi(R_{i3}r)dz = \int_{-\infty}^{\infty} \psi(xR_{i1} + yR_{i2} + zR_{i3})dz, \quad r = (x, y, z)^T.$$  

(2)

The “orientation assignment problem” is defined as finding a set of $N$ rotation matrices $\{R_i\}_{i=1}^N$ such that (2) holds for all $i \in [N]$, given only the set of projection-images $\{P_{R_i}\}_{i=1}^N$.

An inherent ambiguity in cryo-electron microscopy stems from the fact that the handedness (chirality) of the molecule cannot be resolved from its projection-images. This ambiguity is referred to as the handedness ambiguity. Consequently, any projection-image is compatible with two distinct orientations as follows. We denote by $J = \text{diag}(1, 1, -1)$ the reflection
matrix through the $xy$-plane, and define by $\tilde{\psi}(r) = \psi(Jr)$ the mirror image of the molecule $\psi(r)$, $r = (x, y, z)^T$. Since $J^2 = I$, $\psi(r) = \psi(J^2r) = \tilde{\psi}(Jr)$, and along with (2) we have

$$P_{R_i}(x, y) = \int_{-\infty}^{\infty} \tilde{\psi}(R_ir)dz = \int_{-\infty}^{\infty} \tilde{\psi}(JR_ir)dz = \int_{-\infty}^{\infty} \tilde{\psi}(JR_iJJr)dz.$$  

By noting that $Jr = (x, y, -z)^T$ and using the change of variables $z \to z'= -z$ we have

$$P_{R_i}(x, y) = \int_{-\infty}^{\infty} \tilde{\psi}((JR_iJ)JR)dz = \int_{-\infty}^{\infty} \tilde{\psi}((JR_iJ)(x, y, -z)^T)dz$$

$$= \int_{-\infty}^{\infty} \tilde{\psi}((JR_iJ)(x, y, z')^Td' = \tilde{P}_{JR_iJ}(x, y),$$

where $\tilde{P}$ is a projection-image generated from $\tilde{\psi}$. Equation (3) shows that a projection-image of the molecule $\psi$ at orientation $R_i$ is identical to a projection-image of its mirror image molecule $\tilde{\psi}$ at orientation $JR_iJ$. Thus, both sets of orientations assignments $\{R_i\}_{i=1}^{N}$ and $\{JR_iJ\}_{i=1}^{N}$ are consistent with the same set of projection-images $\{P_{R_i}\}_{i=1}^{N}$. Biologically, only the model reconstructed using the orientations $\{R_i\}_{i=1}^{N}$ is valid, yet distinguishing whether a reconstruction corresponds to $\{R_i\}_{i=1}^{N}$ or $\{JR_iJ\}_{i=1}^{N}$ is impossible without utilizing other structural information.

In this work, we propose an algorithm for solving the “orientation assignment problem” for molecules with tetrahedral or octahedral symmetry [6], denoted by $T \subset SO(3)$ and $O \subset SO(3)$, respectively. To present these symmetries, we denote by $C_n$ the group of all rotations by $2\pi/n$ radians around some fixed axis (rotational symmetry of order $n$). Then, the elements of the tetrahedral symmetry group $T$ are the identity, the elements of 4 $C_3$ rotation groups whose axes pass through each vertex of the regular tetrahedron (see Fig. 1) and the corresponding midpoint of the opposite face, and the elements of 3 $C_2$ rotation groups whose axes pass through the midpoints of two of its opposite edges. In total, the tetrahedral group $T$ has 12 elements. The elements of the octahedral symmetry group $O$ are the identity, the elements of 3 $C_4$ rotation groups whose axes pass through two opposite vertices of the regular octahedron (see Fig. 1), 4 $C_3$ rotation groups whose axes pass through the midpoints of two of its opposite faces, and 6 $C_2$ rotation groups whose axes pass through the midpoints of two of its opposite edges. In total, the octahedral group $O$ has 24 elements.

Since the structure of a molecule is independent of its coordinate system, we choose without loss of generality a coordinate system in which the rotational axes mentioned above coincide with the axes listed in Table 1. In this coordinate system, the symmetry group elements of a molecule with tetrahedral symmetry are given in Appendix A.1 and the symmetry group elements of a molecule with octahedral symmetry are given in Appendix A.2.
| symmetry | axes | angles |
|----------|------|--------|
| $T$      | $[1,1,1], [-1,-1,1], [-1,1,-1], [1,-1,-1]$ | $2\pi/3, 4\pi/3$ |
|          | $[1,0,0], [0,1,0], [0,0,1]$ | $\pi$ |
| $O$      | $[1,1,1], [-1,1,1], [1,-1,1], [1,1,-1]$ | $\pi/2, \pi, 3\pi/2$ |
|          | $[1,1,0], [-1,1,0], [1,0,1], [-1,0,1], [0,1,1], [0,-1,1]$ | $2\pi/3, 4\pi/3$ |

Table 1: The nontrivial elements of the tetrahedral and octahedral symmetries.

To see the effect of symmetry on the orientation assignment problem, we denote by $g^{(k)}$ the $k$-th symmetry group element of the symmetry group $T$ or $O$, $k \in [n]$, where $n$ is the number of elements in the symmetry group. Mathematically, a molecule $\psi$ has symmetry $G$ ($G = T$ or $G = O$) if

$$
\psi(r) = \psi(g^{(k)}r), \quad k \in [n],
$$

for any $r = (x, y, z)^T$. Together with (2), it holds that for any $R_i \in SO(3)$ and any $r = (x, y, z)^T$,

$$
P_{R_i}(x, y) = \int_{-\infty}^{\infty} \psi(R_i r) dz = \int_{-\infty}^{\infty} \psi(g^{(k)} R_i r) dz = P_{g^{(k)} R_i}(x, y),
$$

for all $k \in [n]$, implying that the $n$ projection-images $\{P_{g^{(k)} R_i}\}_{k=1}^{n}$ are identical. Hence, equation (5) reveals another ambiguity of the set of projection-images $\{P_{R_i}\}_{i=1}^{N}$, referred to as the symmetry ambiguity, in which all orientation assignments of the form $\{g_i R_i\}_{i=1}^{N}$, where $g_i \in G$ is an arbitrary symmetry group element, are consistent with the same set of images $\{P_{R_i}\}_{i=1}^{N}$.

Combining the symmetry ambiguity with the handedness ambiguity described in (3), the orientation assignment problem can be stated as finding either one of the sets of orientations $\{R_i\}_{i=1}^{N}$ or $\{J R_i J\}_{i=1}^{N}$, where each $R_i$ may be replaced by $g_i R_i$, with $g_i \in G$ being an arbitrary symmetry group element, independently for each $i$ (that is independently for each rotation).

Solving the orientation assignment problem, in its broadest sense, amounts to relating between images and rotation matrices. In this work, we introduce a common lines based method for solving the orientation assignment problem for molecules having either tetrahedral or octahedral symmetry. The common lines, defined in the following and discussed in detail in subsequent sections, reveal useful relations between images and rotation matrices.

To define common lines, we recall the Fourier projection slice theorem [9], which provides an important relation between the Fourier transform of $\psi$ and the Fourier transform of $P_{R_i}$ (see (2)). We define the two-dimensional Fourier transform of a projection-image (2) by

$$
\hat{P}_{R_i}(\omega_x, \omega_y) = \iiint_{\mathbb{R}^2} P_{R_i}(x, y) e^{-i(x\omega_x + y\omega_y)} dx dy,
$$

and the three-dimensional Fourier transform of the molecule by

$$
\hat{\psi}(\omega_x, \omega_y, \omega_z) = \iiint_{\mathbb{R}^3} \psi(x, y, z) e^{-i(x\omega_x + y\omega_y + z\omega_z)} dx dy dz.
$$

4
Using this notation, the Fourier projection slice theorem states that

$$\hat{P}_{R_i}(\omega_x, \omega_y) = \hat{\psi}(\omega_x R_{i1}^1 + \omega_y R_{i2}^2), \quad (\omega_x, \omega_y) \in \mathbb{R}^2,$$

where $R_{i1}$ and $R_{i2}$ are the first and second columns of $R_i$, respectively. In words, the two-dimensional Fourier transform of any projection-image $P_{R_i}$ is equal to the restriction of the three-dimensional Fourier transform of the molecule $\psi$ to the plane through the origin spanned by $R_{i1}$ and $R_{i2}$, or equivalently, to the central plane whose normal coincides with $R_{i1}^3$.

As any two central planes intersect along a single line through the origin (as long as the central planes do not coincide), the central planes corresponding to any pair of Fourier-transformed projection-images $\hat{P}_{R_i}$ and $\hat{P}_{R_j}$ intersect along such a line, and therefore, both (Fourier transformed) images share a pair of lines on which their Fourier transforms coincide, thus referred to as common lines. Given that $\hat{P}_{R_i}$ and $\hat{P}_{R_j}$ are images of a molecule with tetrahedral or octahedral symmetry, each $\hat{P}_{g^{(k)} R_j}, k \in [n]$, is identical to $\hat{P}_{R_i}$ (see (5)). In addition, each $\hat{P}_{g^{(k)} R_j}, k \in [n]$, also shares a common line with $\hat{P}_{R_i}$. Since the rotations $g^{(k)} R_j, k \in [n]$, are in general different from each other, the planes spanned by their first two columns are also different. Thus, $\hat{P}_{R_i}$ and $\hat{P}_{R_j}$ have $n$ common lines altogether.

## 2 Related work

Common lines methods for ab-initio reconstruction of macromolecules have originated with the angular reconstitution method by Van Heel [23]. It is a sequential method in which given a triplet of projection-images $\{P_{R_i}, P_{R_j}, P_{R_k}\}$, the set of relative rotations $\{R_{i1}\ R_{j1}, R_{i2}\ R_{j2}, R_{i3}\ R_{j3}\}$ is first estimated by detecting common lines between $P_{R_i}$, $P_{R_j}$, and $P_{R_k}$. Then, setting $R_i = I$ without loss of generality, determines $R_j$ and $R_k$ from $R_{i1} R_{j1}$ and $R_{i2} R_{j2}$. By applying this method sequentially to each triplet $\{R_{i1} R_{j1}, R_{i2} R_{j2}, R_{i3} R_{j3}\}$ where $l \neq i, j$, the orientation $R_l$ of the image $P_{R_l}$ is determined from $R_{i1} R_{l1}$ simply by $R_l = R_{i1} R_{l1}$.

Detecting common lines between a pair of images is typically done by finding the pair of central lines in the Fourier transforms of the images that have the highest correlation [19]. In cryo-EM, the images are contaminated with high levels of noise, thus making the detection of common lines error prone. Consequently, the relative rotations in the angular reconstitution method are estimated with errors, which render the method not robust to noise.

A common lines based approach for molecules without symmetry that is robust to noisy input images is the synchronization method [17, 11]. In this approach, all relative rotations $\{R_{i1}^l R_{j1}^l\}_{i < j \in [N]}$ are first estimated using common lines (robust estimation of common lines is described in [19, 7]). Then, the rotations $\{R_{i1}\}_{i \in [N]}$ are estimated simultaneously, by constructing a $3N \times 3N$ matrix whose $(i,j)$ block of size $3 \times 3$ contains the estimate for $R_{i1}^l R_{j1}^l$, and factorizing this matrix using SVD. However, this method is not applicable to symmetric molecules due to the symmetry ambiguity described by (5). Specifically, consider a pair of images $P_{R_i}$ and $P_{R_j}$, $i < j \in [N]$, of a molecule with tetrahedral or octahedral symmetry. By the discussion above, there are $n$ pairs of common lines between the images, corresponding to $n$ pairs of projection planes, but it is unknown which pair of common lines
corresponds to which pair of projection planes. As a result, the best one can estimate from a single pair of common lines between the images is the relative rotation $R^T_i g_{ij} R_j$, where $g_{ij}$ is an unknown arbitrary symmetry group element. In such a case, factorizing the $3N \times 3N$ matrix whose blocks are $R^T_i g_{ij} R_j$ does not give $R_i$ (more precisely $g_i R_i$ for some arbitrary $g_i$), unless we are able to carefully choose $g_{ij}$.

Two robust common lines based methods which are applicable to symmetric molecules are described in [12] for molecules with $C_n$ symmetry and in [16] for molecules with $D_2$ symmetry. In both methods, all common lines between each pair of images are utilized to estimate a set $\{R^T_i g_{ij} R_j\}_{i<j} \in [N]$, with $g_{ij}$ being an unknown symmetry group element of the $C_n$ or $D_2$ symmetry groups. Once the set $\{R^T_i g_{ij} R_j\}_{i<j} \in [N]$ has been estimated, the methods exploit the symmetry group properties to obtain the set of rotation matrices $\{R_i\}_{i \in [N]}$, with each rotation matrix satisfying $R_i \in \{g_k R_i\}_{k=1}^n$. Unfortunately, the methods in [12, 16] are not applicable to molecules with $T$ or $O$ symmetries. Specifically, the method in [12] for molecules with $C_n$ symmetry uses the property that the average of the group elements of $C_n$ is the matrix diag(0, 0, 1). This property doesn’t hold for molecules with $T$ or $O$ symmetry, as the average over all group elements of the groups $T$ and $O$ is the zero matrix. As for the method in [16] for molecules with $D_2$ symmetry, this method assumes that the rotational symmetry axes of the molecule coincide with the $x$, $y$ and $z$ axes, which does not hold for $T$ and $O$ symmetries.

While we propose a method that is based on common lines, there exist algorithms for finding ab-initio models that are based on casting the reconstruction problem as an optimization problem [13, 26, 22, 3]. Such algorithms use some general-purpose optimization algorithm (such as stochastic gradient descent, stochastic hill climbing, projection-matching, and simulated annealing, to name a few), on a subset of the data or its class averages. All these methods boil down to a non-convex optimization, which is susceptible to getting stuck in a local minimum corresponding to a structure that is inconsistent with the investigated molecule. Yet, these methods are widely used in practice and many times produce satisfactory initial models. In Section 5 we show an example of both a success as well as a failure of such a method.

The paper is organized as follows. In Section 3, we formally define the common lines, introduce the notion of self common lines and derive basic properties of common lines and self common lines. In Section 4, we describe our algorithm for estimating the orientations of a given set of projection-images. Then, in Section 5, we report some numerical experiments we conducted using simulated and experimental data sets, demonstrating the robustness and effectiveness of our proposed method. Finally, in Section 6, we discuss possible future work.

\section{Common lines and self common lines}

Formally, for each $k \in [n]$, the unit vector

$$q_{R_i, R_j}^k = \frac{R^3_i \times g^{(k)} R^3_j}{\|R^3_i \times g^{(k)} R^3_j\|}$$

(7)
gives the direction of the common line between the central planes of \( \hat{P}_{R_i} \) and \( \hat{P}_{g^{(k)}R_j} \), since it is perpendicular to the normal vectors of both of them. We can express \( q^{k}_{R_i,R_j} \) using its local coordinates on both central planes by

\[
q^{k}_{R_i,R_j} = \cos(\alpha^{k,1}_{R_i,R_j}) R^1_i + \sin(\alpha^{k,1}_{R_i,R_j}) R^2_i = \cos(\alpha^{k,2}_{R_i,R_j}) g^{(k)} R^1_j + \sin(\alpha^{k,2}_{R_i,R_j}) g^{(k)} R^2_j, \tag{8}
\]

where \( \alpha^{k,1}_{R_i,R_j} \) and \( \alpha^{k,2}_{R_i,R_j} \) are the angles between \( q^{k}_{R_i} \) and the local \( x \)-axes of the planes. Using this notation along with (6), we have that for any \( \xi \in \mathbb{R} \) and \( k \in [n] \),

\[
\hat{P}_{R_i}(\xi \cos(\alpha^{k,1}_{R_i,R_j}), \xi \sin(\alpha^{k,1}_{R_i,R_j})) = \hat{\psi}(\xi \cos(\alpha^{k,1}_{R_i,R_j}) R^1_i + \xi \sin(\alpha^{k,1}_{R_i,R_j}) R^2_i) = \hat{\psi}(\xi q^k_{R_i,R_j}) = \hat{\psi}(\cos(\alpha^{k,2}_{R_i,R_j}) g^{(k)} R^1_j + \sin(\alpha^{k,2}_{R_i,R_j}) g^{(k)} R^2_j) = \hat{P}_{g^{(k)}R_j}(\xi \cos(\alpha^{k,2}_{R_i,R_j}), \xi \sin(\alpha^{k,2}_{R_i,R_j})) = \hat{P}_{R_j}(\xi \cos(\alpha^{k,2}_{R_i,R_j}), \xi \sin(\alpha^{k,2}_{R_i,R_j})). \tag{9}
\]

Following (9), we express the set of common lines between the pair of images \( \hat{P}_{R_i} \) and \( \hat{P}_{R_j} \) by the set of local coordinates \( \{(\alpha^{k,1}_{R_i,R_j}, \alpha^{k,2}_{R_i,R_j})\}_{k \in [n]} \). In particular, \( \alpha^{k,1}_{R_i,R_j} \) and \( \alpha^{k,2}_{R_i,R_j} \) may be recovered from the entries of \( R^T_i \) \( g^{(k)} R_j \) using

\[
\alpha^{k,1}_{R_i,R_j} = \text{atan2} \left( (R^T_i \ g^{(k)} R_j)_{1,3}, -(R^T_i \ g^{(k)} R_j)_{2,3} \right), \\
\alpha^{k,2}_{R_i,R_j} = \text{atan2} \left( -(R^T_i \ g^{(k)} R_j)_{3,1}, (R^T_i \ g^{(k)} R_j)_{3,2} \right). \tag{10}
\]

Note that in (7), the vector \( R^3_i \) is multiplied by the symmetry group element \( g^{(k)} \), \( k \in [n] \), while \( R^3_i \) is not. Since this choice is arbitrary, we show in the following that we get the same common lines if we multiply \( R^3_i \) by \( g^{(k)} \) instead. In other words, we show in the following that the set of local coordinates \( \{(\alpha^{k,1}_{R_i,R_j}, \alpha^{k,2}_{R_i,R_j})\}_{k \in [n]} \) for the common lines between the pair of images \( \hat{P}_{R_i} \) and \( \hat{P}_{R_j} \) is well defined.

Similarly to (7), for each \( l \in [n] \), the unit vector

\[
\tilde{q}^{k}_{R_i,R_j} = \frac{g^{(l)} R^3_i \times R^3_j}{\|g^{(l)} R^3_i \times R^3_j\|} \tag{11}
\]
gives the direction of the common line between the central planes of the Fourier transformed images \( \hat{P}_{g^{(l)}R_i} \) and \( \hat{P}_{R_j} \). As \( G \) is a group, for each \( l \in [n] \) there is \( k \in [n] \) such that \( (g^{(k)})^T = g^{(l)} \). Then, using (7), it holds that

\[
(g^{(k)})^T q^{k}_{R_i,R_j} = (g^{(k)})^T \frac{R^3_i \times g^{(k)} R^3_j}{\|R^3_i \times g^{(k)} R^3_j\|} = \frac{g^{(l)} R^3_i \times R^3_j}{\|R^3_i \times g^{(k)} R^3_j\|} = \frac{g^{(l)} R^3_i \times R^3_j}{\|g^{(l)} R^3_i \times R^3_j\|} = \tilde{q}^{k}_{R_i,R_j}, \tag{12}
\]

where the second equality follows since for any rotation \( R \) it holds that \( R(a \times b) = (Ra) \times (Rb) \), and the third equality follows from the latter property along with the anti-commutative
property of the cross product, i.e., \( a \times b = -(b \times a) \), and the invariance of the 2-norm to orthogonal transformations. By multiplying (8) by \((g^{(k)})^T\) from the left we get using (12)
\[
q^T_{R_i,R_i} = \cos(\alpha_{R_i,R_i}^{k,1}) g^{(i)} R_i^1 + \sin(\alpha_{R_i,R_i}^{k,1}) g^{(i)} R_i^2 = \cos(\alpha_{R_i,R_i}^{k,2}) R_i^1 + \sin(\alpha_{R_i,R_i}^{k,2}) R_i^2.
\]
Equation (13) implies that \(\alpha_{R_i,R_i}^{k,1}\) and \(\alpha_{R_i,R_i}^{k,2}\) are also the angles between \(q^T_{R_i,R_i}\) and the local \(x\)-axes of the planes of the Fourier transformed images \(\hat{P}_{g^{(i)} R_i}\) and \(\hat{P}_{R_i}\). Then, similarly to (9),
\[
\hat{P}_{R_i}(\xi \cos(\alpha_{R_i,R_i}^{k,1}), \xi \sin(\alpha_{R_i,R_i}^{k,1})) = \hat{P}_{g^{(i)} R_i}(\xi \cos(\alpha_{R_i,R_i}^{k,1}), \xi \sin(\alpha_{R_i,R_i}^{k,1}))
\]
\[
= \hat{P}_{R_i}(\xi \cos(\alpha_{R_i,R_i}^{k,2}), \xi \sin(\alpha_{R_i,R_i}^{k,2})).
\]
Thus, the set of local coordinates for the common lines is well defined, as the same set is obtained from the two equivalent definitions (7) and (11).

Another important property of projection-images of symmetric molecules, and in particular of molecules with tetrahedral or octahedral symmetry, is the existence of self common lines, which are common lines between any two (identical) images \(\hat{P}_{R_i}\) and \(\hat{P}_{g^{(k)} R_i}\), \(k \in \{2, \ldots, n\}\). The direction vector of the self common line between \(\hat{P}_{R_i}\) and \(\hat{P}_{g^{(k)} R_i}\) is
\[
q_{R_i,R_i}^k = \frac{R_i^3 \times g^{(k)} R_i^3}{||R_i^3 \times g^{(k)} R_i^3||}.
\]
When expressing \(q_{R_i,R_i}^k\) by the local coordinates \((\alpha_{R_i,R_i}^{k,1}, \alpha_{R_i,R_i}^{k,2})\), we get similarly to (9) that
\[
\hat{P}_{R_i}(\xi \cos(\alpha_{R_i,R_i}^{k,1}), \xi \sin(\alpha_{R_i,R_i}^{k,1})) = \hat{P}_{R_i}(\xi \cos(\alpha_{R_i,R_i}^{k,2}), \xi \sin(\alpha_{R_i,R_i}^{k,2})),
\]
and \(\alpha_{R_i,R_i}^{k,1}\) and \(\alpha_{R_i,R_i}^{k,2}\) may be recovered from the entries of \(R_i^T g^{(k)} R_i\) using
\[
\alpha_{R_i,R_i}^{k,1} = \text{atan2} \left( (R_i^T g^{(k)} R_i)_{1,3}, -(R_i^T g^{(k)} R_i)_{2,3} \right),
\]
\[
\alpha_{R_i,R_i}^{k,2} = \text{atan2} \left( -(R_i^T g^{(k)} R_i)_{3,1}, (R_i^T g^{(k)} R_i)_{3,2} \right).
\]
Thus, the set of self common lines of the image \(\hat{P}_{R_i}\) is expressed by the set of local coordinates \(\{ (\alpha_{R_i,R_i}^{k,1}, \alpha_{R_i,R_i}^{k,2}) \}_{k \in \{2, \ldots, n\}}\).

4 Algorithm

In this section, we derive our method for solving the orientation assignment problem for molecules with tetrahedral or octahedral symmetry. Throughout this section, we denote by \(G\) either the group \(\mathbb{T}\) or the group \(\mathbb{O}\). Our method consists of two steps; first, we assign to each pair of projection-images \(P_{R_i}\) and \(P_{R_j}\) (see (2)), \(i < j \in [N]\), of a molecule whose
symmetry group is \( G \), a pair of rotation matrices \((\tilde{R}_{ij}, \tilde{R}_{ji})\) which is an estimate to a pair of rotation matrices \((R_{ij}, R_{ji})\) which satisfies

\[
\{R^T_{ij} g^{(k)} R_{ji}\}_{k=1}^n = \{R^T_i g^{(k)} R_j\}_{k=1}^n, \quad g^{(k)} \in G.
\] (18)

Then, we estimate the orientations of all projection-images \(\{P_{R_i}\}_{i \in [N]}\) from the set of rotation matrices \(\{(\tilde{R}_{ij}, \tilde{R}_{ji})\}_{i \neq j \in [N]}\).

To find the rotations \(\tilde{R}_{ij}\) and \(\tilde{R}_{ji}\) which estimate \(R_{ij}\) and \(R_{ji}\) of (18), we follow the maximum likelihood approach described in [12, 16] as follows. First, we construct a function \(\pi_{ij}(Q_r, Q_s)\), which for any two rotations \(Q_r, Q_s \in SO(3)\) computes a score that indicates how well \(\{Q_r T g^{(k)} Q_s\}_{k=1}^n\) approximates \(\{R^T_i g^{(k)} R_j\}_{k=1}^n\). Since it is impossible to find efficiently the optimum of \(\pi_{ij}\) over \(SO(3) \times SO(3)\), we show in Appendix C how to construct a finite subset \(SO_G(3) \subset SO(3)\) on which we search for the optimum of \(\pi_{ij}\). In particular, the subset \(SO_G(3)\) takes advantage of the fact that the underlying molecule is symmetric, to reduce the number of rotations in \(SO_G(3)\), while maintaining high accuracy of our algorithm. We use the pair \((Q_r, Q_s) \in SO_G(3) \times SO_G(3)\) that attains the highest score \(\pi_{ij}\) as our estimate \((\tilde{R}_{ij}, \tilde{R}_{ji})\) for \((R_{ij}, R_{ji})\) of (18).

We next describe the construction of the function \(\pi_{ij} : SO_G(3) \times SO_G(3) \rightarrow [0, 1]\) for each pair of images \(P_{R_i}\) and \(P_{R_j}\). We denote by

\[
\nu_{i, \theta}(\xi) = \hat{P}_{R_i}(\xi \cos \theta, \xi \sin \theta), \quad \xi \in (0, \infty)
\] (19)

the half line (known as a Fourier ray) in the direction which forms an angle \(\theta\) with the \(x\)-axis of the Fourier transformed image \(\hat{P}_{R_i}, i \in [N]\), and by

\[
\rho_{ij}(\theta, \phi) = \Re \int_0^\infty \frac{(\nu_{i, \theta}(\xi))^* \nu_{j, \phi}(\xi) d\xi}{||\nu_{i, \theta}(\xi)||_2 ||\nu_{j, \phi}(\xi)||_2}
\] (20)

the real part of the normalized cross correlations between \(\nu_{i, \theta}(\xi)\) and \(\nu_{j, \phi}(\xi)\). Note that due to (9), it holds that \(\rho_{ij}(\alpha_{R_i, R_j}^{k,1}, \alpha_{R_i, R_j}^{k,2}) = 1\) for all \(k \in [n]\), where \(\{(\alpha_{R_i, R_j}^{k,1}, \alpha_{R_i, R_j}^{k,2})\}_{k \in [n]}\) is the set of common lines between \(\hat{P}_{R_i}\) and \(\hat{P}_{R_j}\).

Now, consider a pair of rotations \(Q_r, Q_s \in SO_G(3)\). Analogously to (10), we compute the set of local coordinates \(\{(\alpha_{Q_r, Q_s}^{k,1}, \alpha_{Q_r, Q_s}^{k,2})\}_{k \in [n]}\) from the set \(\{Q_r^T g^{(k)} Q_s\}_{k=1}^n\) using

\[
\alpha_{Q_r, Q_s}^{k,1} = \arctan2 \left((Q_r^T g^{(k)} Q_s)_{1,3}, -(Q_r^T g^{(k)} Q_s)_{2,3}\right),
\]
\[
\alpha_{Q_r, Q_s}^{k,2} = \arctan2 \left(-((Q_r^T g^{(k)} Q_s)_{3,1}, (Q_r^T g^{(k)} Q_s)_{3,2}\right).
\] (21)

If \(\{Q_r^T g^{(k)} Q_s\}_{k=1}^n\) is an estimate to \(\{R^T_i g^{(k)} R_j\}_{k=1}^n\), then (21) along with (10) imply that the set of local coordinates \(\{(\alpha_{Q_r, Q_s}^{k,1}, \alpha_{Q_r, Q_s}^{k,2})\}_{k \in [n]}\) is the set of common lines of the pair of images \(P_{R_i}\) and \(P_{R_j}\), i.e., it is equal to \(\{(\alpha_{R_i, R_j}^{k,1}, \alpha_{R_i, R_j}^{k,2})\}_{k \in [n]}\). The score function \(\pi_{ij}\) is thus defined as

\[
\pi_{ij}(Q_r, Q_s) = \prod_{k \in [n]} \rho_{ij}(\alpha_{Q_r, Q_s}^{k,1}, \alpha_{Q_r, Q_s}^{k,2}),
\] (22)
satisfying $\pi_{ij}(Q_r, Q_s) = 1$ if $\{Q_r^T g^{(k)} Q_s\}_{k=1}^n = \{R_i^T g^{(k)} R_j\}_{k=1}^n$. We note that we define $\pi_{ij}$ in (22) as a product to enforce that all $n$ correlations are large simultaneously. Each $\rho_{ij}(\alpha_{Q_r, Q_s}^{k,1}, \alpha_{Q_r, Q_s}^{k,2})$ is a proxy to the probability that $(\alpha_{Q_r, Q_s}^{k,1}, \alpha_{Q_r, Q_s}^{k,2})$ is a common line between $P_{R_i}$ and $P_{R_j}$, and we want all these probabilities to be large simultaneously.

In practice, since $\pi_{ij}$ is computed using noisy images, and since $SO_G(3)$ is only a finite subset of $SO(3)$, $\pi_{ij}(Q_r, Q_s)$ is unlikely to be exactly 1. Thus, the pair of candidates $Q_r, Q_s \in SO_G(3)$ with the highest score $\pi_{ij}(Q_r, Q_s)$ is used to construct $\{Q_r^T g^{(k)} Q_s\}_{k=1}^n$, which serves as an estimate for $\{R_i^T g^{(k)} R_j\}_{k=1}^n$.

In order to achieve a more robust estimate of $\{R_i^T g^{(k)} R_j\}_{k=1}^n$, we also combine the set of self common lines into the score function $\pi_{ij}$ of (22). Specifically, as $\{(\alpha_{R_i,R_i}^{k,1}, \alpha_{R_i,R_i}^{k,2})\}_{k \in \{2, \ldots, n\}}$ and $\{(\alpha_{R_j,R_j}^{k,1}, \alpha_{R_j,R_j}^{k,2})\}_{k \in \{2, \ldots, n\}}$ are the sets of self common lines of $P_{R_i}$ and $P_{R_j}$ respectively (see (16) and its following paragraph), we define

$$\pi_{ij}(Q_r, Q_s) = \prod_{k \in [n]} \rho_{ij}(\alpha_{Q_r, Q_s}^{k,1}, \alpha_{Q_r, Q_s}^{k,2}) \prod_{k \in \{2, \ldots, n\}} \rho_{ii}(\alpha_{Q_r, Q_s}^{k,1}, \alpha_{Q_r, Q_s}^{k,2}) \rho_{jj}(\alpha_{Q_r, Q_s}^{k,1}, \alpha_{Q_r, Q_s}^{k,2}).$$

Using the score function $\pi_{ij}$ of (23), we choose for each $i < j \in [N]$ the pair $(\tilde{R}_{ij}, \tilde{R}_{ji})$ that satisfies

$$\langle \tilde{R}_{ij}, \tilde{R}_{ji} \rangle = \arg \max_{(Q_r, Q_s) \in SO_G(3) \times SO_G(3)} \pi_{ij}(Q_r, Q_s),$$

and use it as an estimate for $(R_{ij}, R_{ji})$. The procedure for computing the set $\{(\tilde{R}_{ij}, \tilde{R}_{ji})\}_{i < j \in [N]}$ is summarized in Algorithm 1.

**Algorithm 1** Computing $\{(\tilde{R}_{ij}, \tilde{R}_{ji})\}_{i < j \in [N]}$ for molecules with $T$ or $O$ symmetry

1: **Input:** (i) $SO_G(3)$ (ii) Images $\hat{P}_{R_i}$, $i \in [N]$
2: **for** $i < j = 1, \ldots, N$ **do**
3: \hspace{1cm} $(\tilde{R}_{ij}, \tilde{R}_{ji}) \leftarrow \arg \max_{(Q_r, Q_s) \in SO_G(3) \times SO_G(3)} \pi_{ij}(Q_r, Q_s)$ \hspace{1cm} $\triangleright$ See (24)
4: **end for**
5: **Output:** $\{(\tilde{R}_{ij}, \tilde{R}_{ji})\}_{i < j \in [N]}$

The Fourier projection-slice theorem (6) relies on the stipulation that the centers of all projection-images coincide with the center of the three-dimensional molecule. In practice, it is unlikely that all projection-images are simultaneously aligned with respect to a common three-dimensional origin, making the Fourier projection-slice theorem, as stated in (6), not applicable to pairs of experimental projection-images. Thus, the procedure for detecting common lines between projection-images needs to be modified in order to handle the presence of unknown shifts, as we now describe. Since each input image is centered differently, we do not observe $P_{R_i}(x, y)$ that follows (2), but rather $P_{R_i}(x - \Delta x_i, y - \Delta y_i)$ for some unknown $(\Delta x_i, \Delta y_i)$. This results in the Fourier ray (19) in the direction of a common line being multiplied by some phases that correspond to a one-dimensional shift along the common
line, though the directions of the common lines (10) do not change (see [20] for a detailed derivation). Thus, we replace in (23) each $\rho_{ij}$ (see (20)) by

$$\rho_{ij}(\theta, \phi) = \max_s \Re \int_0^\infty \frac{(\nu_{i,\theta}^s(\xi))^* \nu_{j,\phi}(\xi) d\xi}{||\nu_{i,\theta}^s(\xi)||_{L_2} ||\nu_{j,\phi}(\xi)||_{L_2}},$$

where $\nu_{i,\theta}^s(\xi)$ is the Fourier ray $\nu_{i,\theta}(\xi)$ (see (19)) multiplied by the phases that correspond to a shift of the common line by $s$. In practice, we maximize over a finite set of values of $s$ (for example, up to 10% of the size of the image in steps of 1 pixel). We make a similar change in $\rho_{ii}$ and $\rho_{ji}$ in (23).

Due to the inherent handedness ambiguity of (3), the images $\hat{P}_{R_i}$ and $\hat{P}_{JR_iJ}$ are identical. Thus, the common line between each pair of projection-images $\hat{P}_{R_i}$ and $\hat{P}_{g^{(k)}R_j}$, $i < j \in [N]$, $k \in [n]$, is identical to the common line between the pair of projection-images $\hat{P}_{JR_iJ}$ and $\hat{P}_{g^{(k)}R_j}$. Since by a direct calculation it can be shown for the symmetry group $G$ that $\{Jg^{(k)}J\}_{k=1}^n = \{g^{(k)}\}_{k=1}^n$, it holds that

$$\{(Jr_iJ)^T g^{(k)} JR_jJ\}_{k=1}^n = \{JR_iT g^{(k)} R_j J\}_{k=1}^n.$$

According to (10) and (17), we note that the set $\{JR_iT g^{(k)} R_j J\}_{k=1}^n$ produces the same set of local coordinates as the set $\{R_i^T g^{(k)} R_j\}_{k=1}^n$, and thus also maximizes $\pi_{ij}$ of (23). As a result, the pair $(\hat{R}_{ij}, \hat{R}_{ji})$ estimates either the pair $(R_{ij}, R_{ji})$ or the pair $(JR_{ij}J, JR_{ji}J)$, yet it is impossible to distinguish between the two. Moreover, the estimate for each pair of indices $(i, j)$ is independent from other pairs of indices. Therefore, we apply the handedness synchronization procedure [11] in order to partition the set $\{(\hat{R}_{ij}, \hat{R}_{ji}) \mid i < j \in [N]\}$ into two subsets, given by

$$A = \{(\hat{R}_{ij}, \hat{R}_{ji}) \mid (\hat{R}_{ij}, \hat{R}_{ji}) \text{ estimates } (R_{ij}, R_{ji})\},$$

$$B = \{(\hat{R}_{ij}, \hat{R}_{ji}) \mid (\hat{R}_{ij}, \hat{R}_{ji}) \text{ estimates } (JR_{ij}J, JR_{ji}J)\}. \quad (25)$$

Then we choose either one of the subsets, and replace each estimate $(\hat{R}_{ij}, \hat{R}_{ji})$ in it with $(JR_{ij}J, JR_{ji}J)$. Since $J^2 = I$, all estimates are now consistent with the same hand. From now on, we assume without loss of generality that each pair $(\hat{R}_{ij}, \hat{R}_{ji})$ computed by Algorithm 1, $i < j \in [N]$, estimates $(R_{ij}, R_{ji})$.

Once we have computed $(\hat{R}_{ij}, \hat{R}_{ji})$ for all $i < j \in [N]$ (using Algorithm 1), which estimate $(\hat{R}_{ij}, \hat{R}_{ji})$ satisfying (18), in the second step of the proposed method, we estimate the orientations of all projection-images $\{P_{R_i}\}_{i \in [N]}$. This step relies on the following two propositions and corollary, showing that $R_{ij}$ and $R_{ji}$ which satisfy (18) are not unique, and may be expressed by $R_i$ and $R_j$ of (18) up to a symmetry group element in $O$. As a result, the pair $(\hat{R}_{ij}, \hat{R}_{ji})$ of (24) is also not unique, in the sense that it may estimate any pair $(R_{ij}, R_{ji})$ which satisfies (18).
We start by recalling that the normalizer of a subgroup $\tilde{G}$ in a group $\tilde{H}$ ($\tilde{G} \subseteq \tilde{H}$) is given by

$$N_{\tilde{H}}(\tilde{G}) = \{ \tilde{h} \in \tilde{H} : \tilde{h}^T \tilde{G} \tilde{h} = \tilde{G} \}.$$ 

**Proposition 1.** Let $R_{ij}, R_{ji}$ and $R_i, R_j$ be two pairs of rotations satisfying (18), $i < j \in [N]$. Then $R_{ij}, R_{ji}$ and $R_i, R_j$ satisfy

$$R_{ij} = h_{ij}R_i \text{ and } R_{ji} = h_{ji}R_j, \quad h_{ij}, h_{ji} \in N_{SO(3)}(G).$$

**Proof.** Since $R_{ij}, R_{ji}, R_i, R_j \in SO(3)$, there exist $h_{ij}, h_{ji} \in SO(3)$ such that

$$R_{ij} = h_{ij}R_i \text{ and } R_{ji} = h_{ji}R_j. \quad (27)$$

Substituting (27) into (18) and multiplying both sides of the resulting equation by $R_i$ from the left and by $R_j^T$ from the right results in

$$\{ h_{ij}^T g^{(k)} h_{ji} \}_{k=1}^n = \{ g^{(k)} \}_{k=1}^n. \quad (28)$$

Since $I \in G$ (the identity element of $G$), we deduce from (28) that there exists $g_{ji} \in G$ such that

$$h_{ij}^T Ih_{ji} = h_{ij}^T h_{ji} = g_{ji}$$

and thus

$$h_{ji} = h_{ij}g_{ji}, \quad (29a)$$

$$h_{ij}^T = g_{ji} h_{ji}^T. \quad (29b)$$

Plugging (29a) and (29b) into (28) results in

$$\{ h_{ij}^T g^{(k)} h_{ji} \}_{k=1}^n = \{ g^{(k)} \}_{k=1}^n. \quad (30a)$$

$$\{ g_{ji} h_{ji}^T g^{(k)} h_{ji} \}_{k=1}^n = \{ g^{(k)} \}_{k=1}^n. \quad (30b)$$

Since $G$ is a finite group, it holds that $\{ g_{ji} h_{ji}^T g^{(k)} \}_{k=1}^n = \{ g_{ji} h_{ji}^T g^{(k)} \}_{k=1}^n = \{ g^{(k)} \}_{k=1}^n$. Thus, by multiplying (30a) by $g_{ji}^T$ from the right and by multiplying (30b) by $g_{ji}^T$ from the left, we get

$$\{ h_{ij}^T g^{(k)} h_{ji} \}_{k=1}^n = \{ g^{(k)} \}_{k=1}^n,$$

$$\{ h_{ji}^T g^{(k)} h_{ji} \}_{k=1}^n = \{ g^{(k)} \}_{k=1}^n. \quad (31)$$

Equation (31) implies that $h_{ij}$ and $h_{ji}$ belong to the normalizer of the group $G$ in $SO(3)$, i.e, $h_{ij}, h_{ji} \in N_{SO(3)}(G)$. \hfill \blacksquare

**Proposition 2.** $N_{SO(3)}(T) = \emptyset$ and $N_{SO(3)}(O) = \emptyset$.

The proof of Proposition 2 is given in Appendix D.
Corollary 3. Let $R_{ij}, R_{ji}$ and $R_i, R_j$ be two pairs of rotations satisfying (18), $i < j \in [N]$. For the symmetry group $T$ it holds that

$$R_{ij} = h_{ij}g_{ij}R_i \text{ and } R_{ji} = h_{ji}g_{ji}R_j, \quad g_{ij}, g_{ji} \in T, \quad h_{ij} \in \Omega.$$  \hfill (32)

For the symmetry group $\Omega$ it holds that

$$R_{ij} = g_{ij}R_i \text{ and } R_{ji} = g_{ji}R_j, \quad g_{ij}, g_{ji} \in \Omega.$$  \hfill (33)

Proof. For the symmetry group $T$, by (29a) and Proposition 2, there exist $g_{ji} \in T$ and $h_{ji}, h_{ij} \in \Omega$ such that $h_{ji} = h_{ij}g_{ji}$. Thus by (26), $R_{ji} = h_{ji}R_j = h_{ij}g_{ji}R_j$ and $R_{ij} = h_{ij}R_i = h_{ij}g_{ij}R_i$ with $g_{ij} = I \in T$, implying (32). For the symmetry group $\Omega$, (33) follows directly from (26) and Proposition 2.

We now describe the second step of our proposed method, i.e. how to estimate the orientations of the projection-images $\{P_{R_i}\}_{i \in [N]}$ from the set of rotations $\{\tilde{R}_{ij}, \tilde{R}_{ji}\}_{i < j \in [N]}$ computed by Algorithm 1. This step fundamentally relies on the choice of axes in Table 1, which implies that the matrices corresponding to the group elements of $T$ (Appendix A.1) and $\Omega$ (Appendix A.2) all have exactly one nonzero entry in each row and each column which is equal to either 1 or $-1$. A key property of these symmetry group elements is that they may be represented uniquely using addition and subtraction of single entry matrices, defined as follows.

Definition 4. A single-entry matrix, denoted by $e_{ij} \in \mathbb{R}^{3\times 3}$, is a matrix whose $(i, j)$ element is one and the rest of its elements are zero. Moreover, we define

$$e_{(-i)(-j)} = e_{ij}, \quad e_{(-i)j} = e_{i(-j)} = -e_{ij}.$$  

Definition 5. Given a symmetry group element $g$ from Appendix A.1 for $T$ or Appendix A.2 for $\Omega$, we define the one-line notation of $g$ by the vector $\sigma = (\sigma(1) \sigma(2) \sigma(3))$ given by

$$\sigma^T = g \begin{pmatrix} 1 \\ 2 \\ 3 \end{pmatrix}.$$  

In words, we multiply the matrix that corresponds to a group element by the vector $(1, 2, 3)^T$.

Lemma 6. Each symmetry group element $g$ from Appendix A.1 for $T$ or Appendix A.2 for $\Omega$ may be represented uniquely by the sum

$$g = e_{1\sigma(1)} + e_{2\sigma(2)} + e_{3\sigma(3)},$$

where $\sigma$ is the one-line notation of $g$, and $e_{m\sigma(m)}$, $m = 1, 2, 3$, are given in Definition 4.

Proof. By a direct calculation using the representation given in Appendix A.1 for $T$ and Appendix A.2 for $\Omega$.  

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Recall that due to the symmetry ambiguity (discussed in Section 1), all orientations assignments of the form \( \{g_i R_i\}_{i=1}^N \), where \( g_i \in G \) is an arbitrary symmetry group element, are consistent with the same set of images \( \{P_{R_i}\}_{i \in [N]} \). Hence, there are \( n^N \) valid assignments while only one of them is required. Therefore, our method is designed to obtain one arbitrary valid assignment \( \{g_i R_i\}_{i=1}^N \) from this set of \( n^N \) valid assignments.

The key idea of our method for obtaining a valid assignment \( \{g_i R_i\}_{i=1}^N \) is to estimate one of the rows of all matrices \( g_i R_i \) simultaneously, then another row, and finally the last row, and then assemble the matrices \( g_i R_i \) from these estimations. Note that we do not know which row of the matrices (first, second, or third) we estimate at each step, as explained below.

Let \( \{g_i R_i\}_{i=1}^N \) be any valid assignment to the set of projection-images \( \{P_{R_i}\}_{i \in [N]} \), and let \( m = 1, 2, 3 \). We denote the one-line notation of each \( g_i \in G \), \( i \in [N] \), from \( \{g_i R_i\}_{i=1}^N \) by \( \sigma_i \) (see Definition 5), and define \( \sigma = (\sigma_1, \sigma_2, \ldots, \sigma_N) \). We also denote the \( m \)th row of \( R_i \) by \( v_i^{(m)} \), and for simplicity, we denote by \( v_i^{(-m)} \) the vector \(-v_i^{(m)}\), i.e., the minus of the \( m \)th row of \( R_i \). In particular, \( v_i^{(\sigma_i(m))} \) is the \( m \)th row of \( g_i R_i \) since

\[
g_i R_i = (e_{1\sigma_i(1)} + e_{2\sigma_i(2)} + e_{3\sigma_i(3)}) \begin{pmatrix} -v_i^{(1)} & - & - \\ -v_i^{(2)} & - & - \\ -v_i^{(3)} & - & - \end{pmatrix} = \begin{pmatrix} -v_i^{(\sigma_i(1))} & - & - \\ -v_i^{(\sigma_i(2))} & - & - \\ -v_i^{(\sigma_i(3))} & - & - \end{pmatrix},
\]

where the first equality follows by expressing \( g_i \) using Lemma 6 and the second equality follows by a direct calculation. Moreover, let \( \tau \) denote a one-line notation corresponding to an arbitrary symmetry group element \( g_\tau \in \mathcal{G} \). Then \( v_i^{(\sigma_i(\tau(m)))} \) is the \( \tau(m) \)th row of \( g_i R_i \).

Finally, denote by \( v_{\sigma,\tau(m)} \) the concatenation (as a row vector of length \( 3N \)) of the \( \tau(m) \)th row of each \( g_i R_i \), \( i \in [N] \), i.e.

\[
v_{\sigma,\tau(m)} = (v_i^{(\sigma_i(\tau(m)))}, \ldots, v_N^{(\sigma_N(\tau(m)))}),
\]

and define

\[
H_{\sigma,m} = v_{\sigma,\tau(m)}^T v_{\sigma,\tau(m)}.
\]

Then \( H_{\sigma,m} \) is a rank-1 \( 3N \times 3N \) block matrix whose \((i, j)\) \( 3 \times 3 \) block is given by the rank-1 matrix \( v_i^{(\sigma_i(\tau(m)))} v_j^{(\sigma_j(\tau(m)))} \).

Once we construct the three matrices \( H_{\sigma,m}, m = 1, 2, 3 \), then factorizing each matrix \( H_{\sigma,m} \) using SVD results in either the vector \( v_{\sigma,\tau(m)} \) or the vector \(-v_{\sigma,\tau(m)}\), which we denote by \( s_m \cdot v_{\sigma,\tau(m)} \), \( s_m \in \{-1, 1\} \). Note that

\[
\begin{pmatrix} -s_1 \cdot v_{\sigma,\tau(1)} & - \\ -s_2 \cdot v_{\sigma,\tau(2)} & - \\ -s_3 \cdot v_{\sigma,\tau(3)} & - \end{pmatrix} = S \cdot \begin{pmatrix} -v_{\sigma,\tau(1)} & - \\ -v_{\sigma,\tau(2)} & - \\ -v_{\sigma,\tau(3)} & - \end{pmatrix} = S \cdot g_\tau \cdot \begin{pmatrix} -v_{\sigma,1} & - \\ -v_{\sigma,2} & - \\ -v_{\sigma,3} & - \end{pmatrix} = S \cdot g_\tau \cdot \begin{pmatrix} v_1^{(\sigma_1(1))}, \ldots, v_N^{(\sigma_N(1))} \\ v_1^{(\sigma_1(2))}, \ldots, v_N^{(\sigma_N(2))} \\ v_1^{(\sigma_1(3))}, \ldots, v_N^{(\sigma_N(3))} \end{pmatrix},
\]

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where $S = \text{diag}(s_1, s_2, s_3)$. The latter equation together with (34) and (35) means that if we are able to construct the matrices $H_{\sigma,m}$, then factorizing each of these matrices gives us simultaneously the rows of all matrices $g_i R_i$, up to multiplication by the matrix $O = S \cdot g_r$, $O \in O(3)$. If $\det(O g_i R_i) = -1$, we simply multiply $O g_i R_i$ by $-1$, and thus, we may assume without loss of generality that $O$ is a rotation. The matrix $O$ is an inherent degree of freedom of the orientation assignment problem, with $\{O g_i R_i\}_{i=1}^N$, being a valid solution.

Next, we describe how to construct three rank-1 matrices $H_{\sigma,m}$, $m = 1, 2, 3$, given a set of matrices $\{(R_{ij}, R_{ji})\}_{i<j\in[N]}$ where each $(R_{ij}, R_{ji})$ satisfies (18). Since by Proposition 1 there are many sets of matrices $\{(R_{ij}, R_{ji})\}_{i<j\in[N]}$ satisfying (18), there are also many possible triplets of matrices $H_{\sigma,m}$, $m = 1, 2, 3$, and our algorithm will return one of these triplets. Overall, we will obtain a valid assignment to the set of projection-images $\{P_{R_i}\}_{i\in[N]}$.

We start by showing how to recover from the pair $(R_{ij}, R_{ji})$ of (18) the rank-1 $3 \times 3$ matrices $(v_i^{(\sigma_{ij}(\tau_{ij}(m)))})^T v_j^{(\sigma_{ji}(\tau_{ij}(m)))}$, $m = 1, 2, 3$, $i < j \in [N]$, where $v_i^{(\sigma_{ij}(\tau_{ij}(m)))}$ and $v_j^{(\sigma_{ji}(\tau_{ij}(m)))}$ are the $\tau_{ij}(m)$th rows of $g_{ij} R_i$ and $g_{ji} R_j$, with $g_{ij}, g_{ji} \in G$, whose one-line notations are $\sigma_{ij}$, $\sigma_{ji}$ respectively, and $\tau_{ij}$ is a one-line notation corresponding to a symmetry group element in $\mathcal{O}$. These matrices are the building blocks from which the three rank-1 $3N \times 3N$ block matrices $H_{\sigma,m}$, $m = 1, 2, 3$, will be constructed. We will use the following lemma, whose proof is given in Appendix B.

**Lemma 7.** Let $g_1, g_2$ be any two symmetry group elements from Appendix A.1 for $\mathbb{T}$ or Appendix A.2 for $\mathbb{O}$, with $\sigma_1, \sigma_2$ being their one-line notations, respectively. Then,

\[ g_i^T e_{mm} g_2 = e_{\sigma_i(m)\sigma_2(m)}, \quad m = 1, 2, 3, \quad (36) \]

\[ g_i^T e_{mm} g_1 = e_{\sigma_i(m)\sigma_1(m)}, \quad m = 1, 2, 3, \quad (37) \]

\[ \{-g_i^T e_{mr} g_2\}^3_{m,r=1} = \{-e_{mr}\}^3_{m,r=1}. \quad (38) \]

Using Lemma 7, we prove the following proposition, which relates $(R_{ij}, R_{ji})$ of (18) with the rank-1 $3 \times 3$ matrices $(v_i^{(\sigma_{ij}(\tau_{ij}(m)))})^T v_j^{(\sigma_{ji}(\tau_{ij}(m)))}$, $m = 1, 2, 3$.

**Proposition 8.** Let $R_{ij}, R_{ji}$ and $R_i, R_j$ be two pairs of rotations satisfying (18), $i < j \in [N]$. Then, for $m = 1, 2, 3$,

\[ R_{ij}^T e_{mm} R_{ji} = (v_i^{(\sigma_{ij}(\tau_{ij}(m)))})^T v_j^{(\sigma_{ji}(\tau_{ij}(m)))} \quad (39) \]

where the matrices $e_{mm}$, $m = 1, 2, 3$, are single entry matrices defined in Definition 4, $v_i^{(\sigma_{ij}(\tau_{ij}(m)))}$ and $v_j^{(\sigma_{ji}(\tau_{ij}(m)))}$ are the $\tau_{ij}(m)$th rows of $g_{ij} R_i$ and $g_{ji} R_j$, with $g_{ij}, g_{ji} \in G$, whose one-line notations are $\sigma_{ij}$ and $\sigma_{ji}$ respectively, and $\tau_{ij}$ is a one-line notation corresponding to a symmetry group element in $\mathcal{O}$ (see Definition 5 for the definition of a one-line notation).

**Proof.** For any two symmetry group elements $g_{ij}, g_{ji} \in G$, it holds by (36) that for $m = 1, 2, 3$

\[ g_i^T e_{mm} g_{ji} = e_{\sigma_{ij}(m)\sigma_{ji}(m)}, \quad (40) \]

where $\sigma_{ij}, \sigma_{ji}$ are the one-line notations of $g_{ij}, g_{ji} \in G$, respectively. For any $h_{ij} \in \mathcal{O}$, it holds by (37) that for $m = 1, 2, 3$

\[ h_i^T e_{mm} h_{ij} = e_{\tau_{ij}(m)\tau_{ij}(m)}, \quad (41) \]
where \( \tau_{ij} \) is the one-line notation of \( h_{ij} \).

For the symmetry group \( \mathbb{T} \), we get by (32) that for \( m = 1, 2, 3 \)

\[
R^{T}_{ij}e_{mm}R_{ji} = R^{T}_{i}g^{T}_{ij}h^{T}_{j}e_{mm}h_{ij}g_{j}R_{j} = R^{T}_{i}g^{T}_{i}\epsilon_{\tau_{ij}(m)\tau_{ij}(m)}g_{j}R_{j}
\]

\[
= R^{T}_{i}\epsilon_{\sigma_{ij}(\tau_{ij}(m))\sigma_{ij}(\tau_{ij}(m))}R_{j} = (v^{(\sigma_{ij}(\tau_{ij}(m)))}_{i})^{T}v^{(\sigma_{ji}(\tau_{ij}(m)))}_{j},
\]

where the second equality follows from (41), the third equality follows from (40), and the last equality follows by a direct calculation. For the symmetry group \( \mathbb{O} \), we get by (33) that for \( m = 1, 2, 3 \)

\[
R^{T}_{ij}e_{mm}R_{ji} = R^{T}_{i}g^{T}_{ij}e_{mm}g_{ji}R_{j} = R^{T}_{i}\epsilon_{\sigma_{ij}(\sigma_{ij}(m))\sigma_{ji}(\tau_{ij}(m))}R_{j} = (v^{(\sigma_{ij}(\tau_{ij}(m)))}_{i})^{T}v^{(\sigma_{ji}(\tau_{ij}(m)))}_{j},
\]

where the second equality follows from (40), and the last equality follows by a direct calculation. For convenience only, we write for the symmetry group \( \mathbb{O} \)

\[
R^{T}_{ij}e_{mm}R_{ji} = (v^{(\sigma_{ij}(\tau_{ij}(m)))}_{i})^{T}v^{(\sigma_{ji}(\tau_{ij}(m)))}_{j} = (v^{(\sigma_{ij}(\tau_{ij}(m)))}_{i})^{T}v^{(\sigma_{ji}(\tau_{ij}(m)))}_{j},
\]

where \( \tau_{ij}(m) = m \), so we use consistent notation for both symmetry groups \( \mathbb{T} \) and \( \mathbb{O} \).

We next construct the \( 3N \times 3N \) matrices \( H_{\sigma,m} \), \( m = 1, 2, 3 \), by setting their \( 3 \times 3 \) blocks one by one using Proposition 8, making sure at each step that each \( H_{\sigma,m} \) is a rank-1 matrix satisfying \( H_{\sigma,m} = v^{T}_{\sigma,\tau(m)}v_{\sigma,\tau(m)} \), where \( v_{\sigma,\tau(m)} \) is the concatenation of the \( \tau(m) \)th row of \( g_{i}R_{i} \), \( i \in [N] \), each \( g_{i} \in G \) is an arbitrary symmetry group element, and \( \tau \) is a one-line notation corresponding to an arbitrary symmetry group element in \( \mathbb{O} \).

We start by setting the \((1,2)\) blocks of the matrices \( H_{\sigma,m} \), \( m = 1, 2, 3 \), though the method may be adjusted to start with any other block \((i,j)\), \( i < j \in [N] \). We set the \((1,2)\) block of the matrix \( H_{\sigma,m} \) to be

\[
H^{(1,2)}_{\sigma,m} := R^{T}_{12}e_{mm}R_{21}.
\]

By Proposition 8, it holds that \( H^{(1,2)}_{\sigma,m} = (v^{(\sigma_{12}(\tau_{12}(m)))}_{1})^{T}v^{(\sigma_{21}(\tau_{12}(m)))}_{2} \) where \( v^{(\sigma_{12}(\tau_{12}(m)))}_{1} \) and \( v^{(\sigma_{21}(\tau_{12}(m)))}_{2} \) are the \( \tau_{12}(m) \)th rows of \( g_{12}R_{1} \) and \( g_{21}R_{2} \), respectively, with \( \sigma_{12}, \sigma_{21} \) being the one-line notations corresponding to \( g_{12}, g_{21} \in G \), and \( \tau_{12} \) is a one-line notation corresponding to a symmetry group element in \( \mathbb{O} \). Thus, the \((1,2)\) block of \( H_{\sigma,m} \) is a rank-1 \( 3 \times 3 \) matrix which encodes the \( \tau_{12}(m) \)th row of \( g_{12}R_{1} \) and \( g_{21}R_{2} \).

Once the \((1,2)\) blocks of the matrices \( H_{\sigma,m} \), \( m = 1, 2, 3 \) have been set, to ensure that the matrices \( H_{\sigma,m} \) are of rank-1, the \((1,i)\) and \((2,i)\) blocks of each matrix \( H_{\sigma,m} \), \( i = 3, \ldots, N \), must be of the form \( (v^{(\sigma_{12}(\tau_{12}(m)))}_{1})^{T}v^{(\sigma_{21}(\tau_{12}(m)))}_{2} \) and \( (v^{(\sigma_{21}(\tau_{12}(m)))}_{2})^{T}v^{(\sigma_{12}(\tau_{12}(m)))}_{1} \), respectively, where \( \sigma_{i} \) is a one-line notation corresponding to a symmetry group element \( g_{i} \in G \). Without loss of generality, we continue by setting the \((1,i)\) blocks of the matrices \( H_{\sigma,m} \), \( i = 3, \ldots, N \) (i.e., we could also continue by setting the \((2,i)\) blocks instead).

By Proposition 8, we can compute from \( R_{1i}, R_{1} \) which satisfy (18), \( i = 3, \ldots, N \), the three rank-1 \( 3 \times 3 \) matrices \( (v^{(\sigma_{12}(\tau_{12}(r)))}_{1})^{T}v^{(\sigma_{21}(\tau_{12}(r)))}_{2} \), \( r = 1, 2, 3 \), where \( v^{(\sigma_{12}(\tau_{12}(r)))}_{1} \) and \( v^{(\sigma_{21}(\tau_{12}(r)))}_{2} \) are the \( \tau_{12}(r) \)th rows of \( g_{12}R_{1} \) and \( g_{21}R_{2} \), respectively, with \( \sigma_{12}, \sigma_{21} \) being the one-line notations corresponding to \( g_{12}, g_{21} \in G \), and \( \tau_{12} \) is a one-line notation corresponding to a symmetry
group element in \( G \). Since each product \((v_i^{(\sigma_1(\tau_i(r)))})^T v_i^{(\sigma_1(\tau_i(r)))}\) is some row of \( R_1 \) (or minus some row of \( R_1 \)) times some row of \( R_i \) (or minus some row of \( R_i \)), it must belong to one of the \( H_{\sigma,m} \). We therefore show how to find which matrix \((v_i^{(\sigma_1(\tau_i(r)))})^T v_i^{(\sigma_1(\tau_i(r)))}\), \( r = 1, 2, 3 \), belongs to which \( H_{\sigma,m} \), \( m = 1, 2, 3 \).

By noting that \( v_i^{(s)} \), \( s = 1, 2, 3 \), are the rows of the orthogonal matrix \( R_1 \), we have that for \( m, r = 1, 2, 3 \)

\[
H_{\sigma,m}^{(1,2)} r_{1i}^T e_{rr} R_{i1} = (v_1^{(\sigma_1(\tau_1(m)))})^T v_1^{(\sigma_1(\tau_1(r)))} = v_2^{(\sigma_1(\tau_1(m)))} v_2^{(\sigma_1(\tau_1(r)))},
\]

where \( H_{\sigma,m} \) has already been set by (42), and thus

\[
\|H_{\sigma,m}^{(1,2)} r_{1i}^T e_{rr} R_{i1}\|_F \neq 0 \iff v_1^{(\sigma_1(\tau_1(m)))} v_1^{(\sigma_1(\tau_1(r)))} = \pm v_1^{(\sigma_1(\tau_1(m)))} v_1^{(\sigma_1(\tau_1(r)))},
\]

(44)

Take \( r \in \{1, 2, 3\} \) for which \( \|H_{\sigma,m}^{(1,2)} r_{1i}^T e_{rr} R_{i1}\|_F \neq 0 \). Then

\[
R_{1i}^T e_{rr} R_{i1} = v_1^{(\sigma_1(\tau_1(r)))} v_1^{(\sigma_1(\tau_1(r)))} = v_1^{(\sigma_1(\tau_1(m)))} v_1^{(\sigma_1(\tau_1(r)))},
\]

where the first equality follows by Proposition 8 and the second equality follows by (44).

Next, by (44), \( \tau_1(r) = \pm \sigma_{1i}^{-1}(\sigma_{12}(\tau_1(m))) \), where \( \sigma_{1i}^{-1} \) denotes the one-line notation corresponding to \( g_{1i}^{-1} \in G \), and thus, \( \sigma_{1i}^{-1}(\sigma_{12}) \) is the one-line notation corresponding to \( g_{1i}^{-1} g_{12} \). Then

\[
R_{1i}^T e_{rr} R_{i1} = v_1^{(\sigma_{12}(\tau_1(m)))} v_1^{(\pm \sigma_{1i}(\tau_1(r)))} = v_1^{(\sigma_{12}(\tau_1(m)))} v_1^{(\pm \sigma_{1i}(\sigma_{1i}^{-1}(\sigma_{12}(\tau_1(m)))})},
\]

where \( \sigma_{1i}(\sigma_{1i}^{-1}(\sigma_{12})) \) is the one-line notation corresponding to \( g_{1i}^{-1} g_{12} \in G \). In order to simplify the notation, we define for \( i = 3, \ldots, N \), \( g_i^* = g_{1i}^{-1} g_{12} \in G \), and denote by \( \sigma_i^* \) its one-line notation. Thus

\[
R_{1i}^T e_{rr} R_{i1} = v_1^{(\sigma_{12}(\tau_1(m)))} v_1^{(\sigma_i^*(\sigma_{12}(\tau_1(m)))}) = v_1^{(\sigma_{12}(\tau_1(m)))} v_1^{(\sigma_i^*(\tau_1(m)))},
\]

implying that for \( r \in \{1, 2, 3\} \) which satisfies \( \|H_{\sigma,m}^{(1,2)} r_{1i}^T e_{rr} R_{i1}\|_F \neq 0 \), the matrix \( R_{1i}^T e_{rr} R_{i1} \) is the \( 3 \times 3 \) rank-1 matrix which is the \( (1, \bar{i}) \) block of the matrix \( H_{\sigma,m} \).
Hence, we set the \((1, i)\) blocks of the matrices \(H_{\sigma,m}, m = 1, 2, 3, i = 3, \ldots, N\), as
\[
H^{(1,i)}_{\sigma,m} := \arg \max_{M \in \{R^T_{i} e_r R_{1}\}_{r=1,2,3}} \|H^{(1,2)}_{\sigma,m} M\|_F. \tag{45}
\]
Thus, for \(i = 2, \ldots, N\), each \((1, i)\) block of \(H_{\sigma,m}\) is given by the rank-1 \(3 \times 3\) matrix
\[
v_1^{(\sigma_1^*(\tau_{12}(m)))} v_i^{(\sigma_i^*(\tau_{12}(m)))} (\text{for notation consistency, we also denote by } \sigma_1^* \text{ and } \sigma_2^* \text{ the one-line notations } \sigma_{12} \text{ and } \sigma_{21}, \text{ respectively}).
\]
At this point, we note that as each \(H_{\sigma,m}, m = 1, 2, 3\), must be of rank-1, the \((i, j)\) block of \(H_{\sigma,m}, i, j \in [N]\), must be equal to the \(3 \times 3\) rank-1 matrix
\[
v_i^{(\sigma_i^*(\tau_{12}(m)))} v_j^{(\sigma_j^*(\tau_{12}(m)))}.
\]
This implies that the \((i, 1)\) block of \(H_{\sigma,m}, i = 2, \ldots, N\), is defined as
\[
H^{(i,1)}_{\sigma,m} := H^{(i,1)}_{\sigma,m}, \tag{46}
\]
the \((1, 1)\) block of \(H_{\sigma,m}\) is defined as
\[
H^{(1,1)}_{\sigma,m} := H^{(1,1)}_{\sigma,m}, \tag{47}
\]
and the \((i, j)\) block of \(H_{\sigma,m}, i, j = 2, \ldots, N\), is defined as
\[
H^{(i,j)}_{\sigma,m} := H^{(i,j)}_{\sigma,m}. \tag{48}
\]
Overall, we showed that given the set \(\{(R_{ij}, R_{ij})\}_{i<j\in[N]}\), we can construct three rank-1 \(3N \times 3N\) block matrices \(H_{\sigma,m}, m = 1, 2, 3\), with each \(H_{\sigma,m}\) satisfying \(H_{\sigma,m} = v_{\sigma,\tau_{12}(m)}^T v_{\sigma,\tau_{12}(m)}\)
where
\[
v_{\sigma,\tau_{12}(m)} = (v_1^{(\sigma_1^*(\tau_{12}(m)))}, \ldots, v_i^{(\sigma_i^*(\tau_{12}(m)))}, \ldots, v_N^{(\sigma_N^*(\tau_{12}(m)))}).
\]
Note that the construction of \(H_{\sigma,m}, m = 1, 2, 3\), described above uses only part of the data \(\{(R_{ij}, R_{ij})\}_{i<j\in[N]}\), namely, it uses only the pairs \(\{(R_{1i}, R_{1i})\}_{i=2}^{N}\) to construct the matrices \(H_{\sigma,m}, m = 1, 2, 3\). Since in practice the set \(\{(\tilde{R}_{ij}, R_{ji})\}_{i<j\in[N]}\) computed by Algorithm 1 is only an estimate of the set \(\{(R_{ij}, R_{ji})\}_{i<j\in[N]}\), we only obtain an estimate to each matrix \(H_{\sigma,m}, m = 1, 2, 3\), which we denote by \(\tilde{H}_{\sigma,m}\). Thus, we next show how to modify our construction such that all pairs \(\{(R_{ij}, R_{ji})\}_{i<j\in[N]}\) are considered. This way, constructing \(\tilde{H}_{\sigma,m}, m = 1, 2, 3\), using \(\{(\tilde{R}_{ij}, R_{ji})\}_{i<j\in[N]}\) would result in more robust estimates to the orientations of the projection-images \(\{R_{k1}\}_{i=1}^{N}\). The main idea of the modification is to compute the \((i, j)\) blocks of \(H_{\sigma,m}, m = 1, 2, 3\) from \((R_{ij}, R_{ji}), i, j = 2, \ldots, N, i \neq j\), as follows.

First, for \(i, j = 2, \ldots, N, i \neq j\), we have that \(H^{(i,j)}_{\sigma,m}\) as defined in (48) is equal to the \(3 \times 3\) rank-1 matrix
\[
v_i^{(\sigma_i^*(\tau_{12}(m)))} v_j^{(\sigma_j^*(\tau_{12}(m)))} (\text{as the set of } 3 \times 3 \text{ rank-1 matrices } \{\pm(v_i^{(k)})^T v_j^{(l)}\}_{k,l=1}^3 \text{ consists of all possible products between the rows of } R_i \text{ and the rows of } R_j).\]
In the following proposition, we show how to obtain the set \(\{\pm(v_i^{(k)})^T v_j^{(l)}\}_{k,l=1}^3\) from \((R_{ij}, R_{ji})\).
Proposition 9. Let \( R_{ij}, R_{ji} \) and \( R_i, R_j \) be two pairs of rotations satisfying (18), \( i < j \in [N] \). Then
\[
\{ \pm R_{ij}^{T} e_{kl} R_{ji} \}_{k,l=1}^3 = \{ \pm (v_i^{(k)})^{T} v_j^{(l)} \}_{k,l=1}^3,
\]
where the matrices \( e_{kl} \), \( k, l = 1, 2, 3 \), are single entry matrices defined in Definition 4, \( v_i^{(k)} \) is the \( k \)th row of \( R_i \) and \( v_j^{(l)} \) is the \( l \)th row of \( R_j \).

Proof.
\[
\{ \pm R_{ij}^{T} e_{kl} R_{ji} \}_{k,l=1}^3 = \{ \pm R_i^{T} e_{kl} R_j \}_{k,l=1}^3 = \{ \pm R_i^{T} e_{kl} R_j \}_{k,l=1}^3 = \{ \pm (v_i^{(k)})^{T} v_j^{(l)} \}_{k,l=1}^3,
\]
where the first equality is due to (26), the second equality is due to (38), and the last equality follows by a direct calculation.

Thus there exist \( k, l \in \{1, 2, 3\} \) such that either \( R_{ij}^{T} e_{kl} R_{ji} \) or \( -R_{ij}^{T} e_{kl} R_{ji} \) is equal to \( v_i^{(\sigma_i^{(\tau_1(m))})^{T}} v_j^{(\sigma_j^{(\tau_2(m))})} \). As the current \((i, j)\) blocks of \( H_{\sigma,m} \), \( m = 1, 2, 3 \), are given by \( H_{\sigma,m}^{(1,i)} H_{\sigma,m}^{(1,j)} \) (equation (48)), we replace them with
\[
H_{\sigma,m}^{(i,j)} := \arg \min_{M \in \{ \pm R_{ij}^{T} e_{kl} R_{ji} \}_{k,l=1}^3} \| M - H_{\sigma,m}^{(1,i)} H_{\sigma,m}^{(1,j)} \|_F,
\]
where \( H_{\sigma,m}^{(1,i)}, H_{\sigma,m}^{(1,j)} \) are defined in (45).

Finally, we describe how to construct the three matrices \( \tilde{H}_{\sigma,m} \), \( m = 1, 2, 3 \), from all estimates \( \{ (\tilde{R}_{ij}, \tilde{R}_{ji}) \}_{i<j \in [N]} \) computed by Algorithm 1 (replacing \( \{ (R_{ij}, R_{ji}) \}_{i<j \in [N]} \) used above). Denoting by \( \tilde{H}_{\sigma,m}^{(i,j)} \) the estimate of the \( 3 \times 3 \) block \( H_{\sigma,m}^{(i,j)} \), \( m = 1, 2, 3, i,j \in [N] \), we have by (42)
\[
\tilde{H}_{\sigma,m}^{(1,2)} := \tilde{R}_{12}^{T} e_{m,m} \tilde{R}_{21}.
\]
by (45)
\[
\tilde{H}_{\sigma,m}^{(1,i)} := \arg \max_{M \in \{ R_i^{T} e_{rr} R_i \}_{r=1,2,3}} \| \tilde{H}_{\sigma,m}^{(1,2)} M \|_F, \quad i = 3, \ldots, N,
\]
by (46)
\[
\tilde{H}_{\sigma,m}^{(i,1)} := \tilde{H}_{\sigma,m}^{(1,i)^T}, \quad i = 2, \ldots, N,
\]
and by (49)
\[
\tilde{H}_{\sigma,m}^{(i,j)} := \arg \min_{M \in \{ \pm R_{ij}^{T} e_{kl} R_{ji} \}_{k,l=1}^3} \| M - \tilde{H}_{\sigma,m}^{(1,i)^T} \tilde{H}_{\sigma,m}^{(1,j)} \|_F, \quad i, j = 2, \ldots, N, \quad i \neq j.
\]
Lastly, we note that for \( i \in [N] \), each \( \tilde{H}_{\sigma,m}^{(i,j)} \) \( \tilde{H}_{\sigma,m}^{(j,i)} \), where \( \tilde{H}_{\sigma,m}^{(j,i)} \) was computed by (53), is an estimate to \( H_{\sigma,m}^{(i,j)} \), \( j \in [N] \setminus \{i\} \). Thus, we get a more robust estimate for the \((i, i)\) block of \( H_{\sigma,m} \) by computing the average
\[
\tilde{H}_{\sigma,m}^{(i,i)} := \frac{1}{N-1} \sum_{j=1, j \neq i}^{N} \tilde{H}_{\sigma,m}^{(j,i)^T} \tilde{H}_{\sigma,m}^{(j,i)}
\]
followed by computing the best rank-1 approximation of each \((i, i)\) block of \(\tilde{H}_{\sigma, m}\) using SVD.

To conclude, we estimated three \(3N \times 3N\) block matrices \(\tilde{H}_{\sigma, m}, m = 1, 2, 3\), whose \((i, j)\) \(3 \times 3\) block is an estimate to the rank-1 matrix \(v_i^{(\sigma^* (\tau_{12}(m)))^T} v_j^{(\sigma^* (\tau_{12}(m)))}\) which is estimated from \((\tilde{R}_{ij}, \tilde{R}_{ji})\). We then factorize each matrix using SVD and obtain the estimates \(\{Og^*_i R_i\}_{i=1}^N\) for the orientations of the projection-images \(\{P_{R_i}\}_{i=1}^N\), where \(O\) is a rotation and \(g^*_i \in G\).

The construction of \(\tilde{H}_{\sigma, m}, m = 1, 2, 3\), and the estimation of the orientations of all projection-images \(\{P_{R_i}\}_{i=1}^N\) from the set \(\{(\tilde{R}_{ij}, \tilde{R}_{ji})\}_{i<j \in [N]}\) is summarized in Algorithm 2.

The computational complexity of Algorithm 1 is quadratic in both the number of images as well as in the size of \(SO_G(3)\) (constructed in Appendix C). The computational complexity of Algorithm 2 is quadratic in the number of images.

## 5 Experimental results

We implemented the proposed algorithm in MATLAB, and tested it on both simulated and experimental data. We start with testing the algorithm on simulated data in Section 5.1, to assess its robustness to noise. Then, in Section 5.2, we test the algorithm on experimental cryo-electron microscopy data. All tests were executed on a dual Intel Xeon E5-2683 CPU (32 cores in total), with 768GB of RAM running Linux, and one nVidia GTX TITAN XP GPU (used for Algorithm 1). The implementation of the algorithms is available as part of the ASPIRE software package [1]. To assess the actual memory consumption of the algorithm, we monitored it through the operating system during its execution. The maximal amount of memory used by the algorithm is on the order of storing the reconstructed volume.

### 5.1 Simulated data

To test the performance of our algorithm in the presence of noise, we applied it to noisy simulated projection-images as follows. For \(O\) symmetry, we downloaded from EMDB the map EMD-4905 [10], generated from it clean projection-images of size \(129 \times 129\) pixels, and added to the clean images Gaussian noise with zero mean and variance that results in signal to noise ratio (SNR) of the images the is equal to 1000 (considered as clean images for reference), 1, 1/2, and 1/4 (the signal to noise ratio is defined as the ratio between the energy of the signal in the image and the energy of the noise). Figure 2 shows several examples of projection-images of EMD-4905 at these noise levels.

We then applied our algorithm on sets of \(N = 25, 50, 100, 200\) images at these noise levels. For each group of \(N\) images, we plot in Fig. 3 the Fourier shell correlation curve (FSC) [24] of the volume reconstructed by our algorithm relative to the ground truth volume. In a nutshell, the FSC measures the size of the smallest feature in the reconstructed volume that can be resolved. Figure 3 shows that increasing the number of input images improves the performance of the algorithm in terms of the achieved resolution. Moreover, we see that the algorithm fails for some of the noise levels (achieved resolution worse than 30 Å), but as we increase the number of images, the algorithm successfully reconstructs a three-dimensional model of the molecule, with resolution of about 20 Å. To further demonstrate this point, we
Algorithm 2 Estimating $R_i$, $i = 1, \ldots, N$, for molecules with $T$ or $O$ symmetry.

1: **Input:** $\{(\tilde{R}_{ij}, \tilde{R}_{ji})\}_{i<j \in [N]}$ estimated by Algorithm 1.
2: **Initialize:** Matrices $\tilde{H}_{\sigma,m}$, $m = 1, 2, 3$, of size $3N \times 3N$, with all entries set to zero.
3: $\{(\tilde{R}_{ij}, \tilde{R}_{ji})\}_{i<j \in [N]} \leftarrow$ handedness synchronization($\{(\tilde{R}_{ij}, \tilde{R}_{ji})\}_{i<j \in [N]}$)
4: for $m = 1, 2, 3$ do
5: $\tilde{H}_{\sigma,m}^{(1,2)} = \tilde{R}_{12}^T e_m \tilde{R}_{21}$. \(\triangleright\) $\tilde{H}_{\sigma,m}^{(i,j)}$ denotes the $(i, j)$ $3 \times 3$ block of $\tilde{H}_{\sigma,m}$. See (50).
6: for $i = 3, \ldots, N$ do
7: $\tilde{H}_{\sigma,m}^{(i,i)} = \arg\max_{M \in \{\tilde{R}_{i1}^T e_r \tilde{R}_{i1}\}_{r=1,2,3}} \|\tilde{H}_{\sigma,m}^{(1,2)^T} M\|_F$. \(\triangleright\) See (51).
8: end for
9: for $i < j = 2, \ldots, N$ do
10: $\tilde{H}_{\sigma,m}^{(i,j)} = \arg\min_{M \in \{\pm \tilde{R}_{ij}^T e_k \tilde{R}_{ji}\}_{k,l=1,2,3}} \|M - \tilde{H}_{\sigma,m}^{(1,2)^T} \tilde{H}_{\sigma,m}^{(1,j)}\|_F$. \(\triangleright\) See (53).
11: end for
12: $\tilde{H}_{\sigma,m} \leftarrow \tilde{H}_{\sigma,m} + \tilde{H}_{\sigma,m}^T$
13: for $i = 1, \ldots, N$ do
14: $\tilde{H}_{\sigma,m}^{(i,i)} = \frac{1}{N-1} \sum_{j=1, j \neq i}^{N} \tilde{H}_{\sigma,m}^{(j,i)^T} \tilde{H}_{\sigma,m}^{(i,j)}$ \(\triangleright\) See (54).
15: end for
16: $V_m = \arg\max_{\|v\|=1} v^T \tilde{H}_{\sigma,m} v$ \(\triangleright\) $V_m$ is the eigenvector of the leading eigenvalue of $\tilde{H}_{\sigma,m}$.
17: end for
18: for $i = 1, \ldots, N$ do
19: for $m = 1, 2, 3$ do
20: $v_i^{(m)} = \frac{V_m(3i-2:3i)}{||V_m(3i-2:3i)||}$ \(\triangleright\) $v_i^{(m)}$ is the $m$th row of the orthogonal matrix $\tilde{R}_i$.
21: end for
22: $\tilde{R}_i \leftarrow \begin{pmatrix}
- v_i^{(1)} & - \\
- v_i^{(2)} & - \\
- v_i^{(3)} & - 
\end{pmatrix}$
23: if $\det \tilde{R}_i < 0$ then
24: $\tilde{R}_i \leftarrow -\tilde{R}_i$
25: end if
26: end for
27: **Output:** $\tilde{R}_i$, $i = 1, \ldots, N$. 
Figure 2: Examples of simulated projection-images of EMD-4905 [10] (© symmetry) with signal to noise ratio of (from left to right) 1000, 1, 1/2, 1/4.
| N  | 25  | 50  | 100 | 200 |
|----|-----|-----|-----|-----|
| Time (sec) | 282 | 852 | 2,842 | 11,177 |

Table 2: Timing (in seconds) for $\mathbb{O}$ symmetry.

| N  | 25  | 50  | 100 | 200 |
|----|-----|-----|-----|-----|
| Time (sec) | 397 | 1,352 | 5,178 | 20,228 |

Table 3: Timing (in seconds) for $\mathbb{T}$ symmetry.

show in Figure 4 the Fourier shell correlation curves for a fixed SNR and a variable number of images. As before, increasing the number of images improves the achieved resolution. The timing (in seconds) of the algorithm is summarized in Table 2. These timings were computed by averaging for each $N$ the timing results for all SNRs (as the running time is independent of the noise level).

The results of the same experiment for $\mathbb{T}$ symmetry with projections generated from EMD-10835 [14] are shown in Figs. 5, 6 and 7. The timing for $\mathbb{T}$ symmetry is summarized in Table 3.

5.2 Experimental data

Next, we applied our algorithm to two experimental data sets – EMPIAR-10272 and EMPIAR-10389 from the EMPIAR repository [8]. The EMPIAR-10272 data set corresponds to EMD-4905 [10] that has $\mathbb{O}$ symmetry, and the EMPIAR-10389 data set corresponds to EMD-10835 [14] that has $\mathbb{T}$ symmetry. For comparison, we also generated an ab-initio models from these data sets using Relion [26].

The EMPIAR-10272 data set consists of 480 micrographs, each comprised of 38 raw unaligned movie frames, with pixel size of 0.65 Å/pixel. We first applied motion correction to the movie frames using MotionCor2 [25], resulting in aligned micrographs, to which we applied CTF estimation [5] using CTFFind4 [15]. All subsequent processing steps were executed in Relion [26]. We used Laplacian auto-picking followed by one round of 2D classification to generate templates for template-based picking. Auto-picking resulted in 80,806 particles, which were subjected to 15 rounds of 2D classification, until 24,540 particles in 13 classes were retained. These 13 classes (Fig. 8) were the input to our algorithm, and resulted in an ab-initio model whose resolution is 6.45 Å (compared to the ground-truth density map EMD-4905 [10]). For comparison, we generated a three-dimensional ab-initio model using Relion, using as an input the same particles that were used to generate the class averages for our algorithm. The resolution of the model estimated by Relion 22.84 Å (also compared to the ground-truth density map EMD-4905 [10]). The Fourier shell correlation curves [24] for the initial models generated by our algorithm and by Relion are shown in Fig. 9. To assess visually the two models, we show in Fig. 10 a two-dimensional view of the ground-truth.
Figure 3: Fourier shell correlation curves for volumes reconstructed from simulated projection-images of EMD-4905 (O symmetry). Each volume is reconstructed from a set of images whose size is specified in the caption of the panels.
Figure 4: Fourier shell correlation curves for volumes reconstructed from simulated projection-images of EMD-4905 (O symmetry). Each panel corresponds to a given SNR, and a number of projections as appears in the legend.
Figure 5: Examples of simulated projection-images of EMD-10835 [14] (T symmetry) with signal to noise ratio of (from left to right) 1000, 1, 1/2, 1/4.
Figure 6: Fourier shell correlation curves for volumes reconstructed from simulated projection-images of EMD-10835 (T symmetry). See Fig. 3 for details.
Figure 7: Fourier shell correlation curves for volumes reconstructed from simulated projection-images of EMD-10835 (T symmetry). See Fig.4 for details.
Figure 8: Class averages used to generate an ab-initio model for EMPIAR-10272.

volume, the volume reconstructed by our algorithm (denoted ASPIRE in the figure), and the volume reconstructed by Relion. It can be observed that for this data set, the initial model generated by Relion is clearly inferior to the one generated by our algorithm.

Next, we tested our algorithm on the EMPIAR-10389 data set, which has $T$ symmetry. The EMPIAR-10389 data set consists of 4,313 dose-weighted micrographs with pixel size of 0.639 Å/pixel. Automatic particle picking was done using the KLT picker [4], resulting in 164,183 particles of size 512 × 512 pixels. (We note that for the EMPIAR-10272 data set discussed above, we used Relion’s particle picker, as it gave superior results.) The particles were then imported into Relion [26], and were subjected to 5 rounds of 2D classification, until 63,057 particles remained in 30 classes. These classes were used as the input to our algorithm, and are shown in Fig. 11. The resolution of the resulting ab-initio model compared to the ground-truth density map EMD-10835 [14] is 7.31 Å. Using Relion’s initial model algorithm on the particles of these 30 classes resulted in a resolution of 10 Å (also compared to the ground-truth density map EMD-10835). The Fourier shell correlation curves for the initial models generated by our algorithm and by Relion are shown in Fig. 12. As before, we show in Fig. 13 two-dimensional views of the ground-truth volume EMD-10835, the volume reconstructed by our algorithm, and the volume reconstructed using Relion.
Figure 9: Fourier shell correlation curves for ab-initio models for EMPIAR-10272.

Figure 10: Ab-initio models for EMPIAR-10272.
Figure 11: 16 of the class averages used to generate an ab-initio model for EMPIAR-10389.

Figure 12: Fourier shell correlation curves for ab-initio models for EMPIAR-10389.
6 Future work

In this work, we proposed a method for estimating the orientations corresponding to a given set of projection-images of a molecule with tetrahedral or octahedral symmetry. The method relies on the observation that the elements of the tetrahedral and octahedral symmetry groups may be represented as rotation matrices with exactly one nonzero entry in each row and each column which is equal to either 1 or -1.

A future extension of this work would be to generalize it to molecules with icosahedral symmetry denoted by $I$. Since the elements of the icosahedral symmetry group cannot be represented as rotation matrices with exactly one nonzero entry in each row and each column which is equal to either 1 or -1, the method suggested in this work is not applicable to this symmetry.

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Appendices

Appendix A Symmetry group elements

A.1 Tetrahedral group $T$

| element | matrix | axis | angle | single-entry sum | one-line notation |
|---------|--------|------|-------|------------------|------------------|

Figure 13: Ab-initio models for EMPIAR-10389.
\begin{align*}
g^{(1)} &= \begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix} \quad \text{any} \quad 0 \quad e_{11} + e_{22} + e_{33} \quad \sigma_1 = (1 \ 2 \ 3) \\
g^{(2)} &= \begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix} \quad [1,1,1] \quad 2\pi/3 \quad e_{13} + e_{21} + e_{32} \quad \sigma_2 = (3 \ 1 \ 2) \\
g^{(3)} &= \begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix} \quad [1,1,1] \quad 4\pi/3 \quad e_{12} + e_{23} + e_{31} \quad \sigma_3 = (2 \ 3 \ 1) \\
g^{(4)} &= \begin{pmatrix} 0 & 0 & -1 \\ 1 & 0 & 0 \\ 0 & -1 & 0 \end{pmatrix} \quad [-1,-1,1] \quad 2\pi/3 \quad -e_{13} + e_{21} - e_{32} \quad \sigma_4 = (-3 \ 1 \ -2) \\
g^{(5)} &= \begin{pmatrix} 0 & 0 & -1 \\ 0 & 1 & 0 \\ -1 & 0 & 0 \end{pmatrix} \quad [-1,-1,1] \quad 4\pi/3 \quad e_{12} - e_{23} - e_{31} \quad \sigma_5 = (2 \ -3 \ -1) \\
g^{(6)} &= \begin{pmatrix} 0 & 1 & 0 \\ -1 & 0 & 0 \\ 0 & 0 & 1 \end{pmatrix} \quad [1,-1,-1] \quad 2\pi/3 \quad -e_{13} - e_{21} + e_{32} \quad \sigma_6 = (-3 \ -1 \ 2) \\
g^{(7)} &= \begin{pmatrix} 0 & 0 & 1 \\ 0 & 1 & 0 \\ -1 & 0 & 0 \end{pmatrix} \quad [1,-1,-1] \quad 4\pi/3 \quad -e_{12} + e_{23} - e_{31} \quad \sigma_7 = (-2 \ 3 \ -1) \\
g^{(8)} &= \begin{pmatrix} 0 & 0 & 1 \\ 0 & -1 & 0 \\ -1 & 0 & 0 \end{pmatrix} \quad [-1,-1,-1] \quad 2\pi/3 \quad e_{13} - e_{21} - e_{32} \quad \sigma_8 = (3 \ -1 \ -2) \\
g^{(9)} &= \begin{pmatrix} 0 & 0 & -1 \\ 0 & 1 & 0 \\ 1 & 0 & 0 \end{pmatrix} \quad [-1,-1,-1] \quad 4\pi/3 \quad -e_{12} - e_{23} + e_{31} \quad \sigma_9 = (-2 \ -3 \ 1) \\
g^{(10)} &= \begin{pmatrix} 0 & -1 & 0 \\ 0 & 0 & -1 \\ 1 & 0 & 0 \end{pmatrix} \quad [1,0,0] \quad \pi \quad e_{11} - e_{22} - e_{33} \quad \sigma_{10} = (1 \ -2 \ -3) \\
g^{(11)} &= \begin{pmatrix} 0 & 1 & 0 \\ 0 & 0 & -1 \\ -1 & 0 & 0 \end{pmatrix} \quad [0,1,0] \quad \pi \quad -e_{11} + e_{22} - e_{33} \quad \sigma_{11} = (-1 \ 2 \ -3) \\
g^{(12)} &= \begin{pmatrix} 0 & -1 & 0 \\ 0 & 0 & 1 \\ -1 & 0 & 0 \end{pmatrix} \quad [0,0,1] \quad \pi \quad -e_{11} - e_{22} + e_{33} \quad \sigma_{12} = (-1 \ -2 \ 3) \\
\end{align*}

### A.2 Octahedral group $\mathcal{O}$

| element | matrix | axis | angle | single-entry sum | one-line notation |
|---------|--------|------|-------|------------------|------------------|
| $g^{(1)}$ | $\begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix}$ | $[1,1,1]$ | $2\pi/3$ | $e_{11} + e_{22} + e_{33}$ | $(1 \ 2 \ 3)$ |
| $g^{(2)}$ | $\begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix}$ | $[1,1,1]$ | $4\pi/3$ | $e_{12} + e_{23} + e_{31}$ | $(2 \ 3 \ 1)$ |
| $g^{(3)}$ | $\begin{pmatrix} 0 & 0 & -1 \\ 1 & 0 & 0 \\ 0 & -1 & 0 \end{pmatrix}$ | $[-1,-1,1]$ | $2\pi/3$ | $-e_{13} + e_{21} - e_{32}$ | $(-3 \ 1 \ -2)$ |
| $g^{(4)}$ | $\begin{pmatrix} 0 & 0 & -1 \\ 0 & 1 & 0 \\ -1 & 0 & 0 \end{pmatrix}$ | $[-1,-1,1]$ | $4\pi/3$ | $e_{12} - e_{23} - e_{31}$ | $(2 \ -3 \ -1)$ |
| $g^{(5)}$ | $\begin{pmatrix} 0 & 1 & 0 \\ -1 & 0 & 0 \\ 0 & 0 & 1 \end{pmatrix}$ | $[1,-1,-1]$ | $2\pi/3$ | $-e_{13} - e_{21} + e_{32}$ | $(-3 \ -1 \ 2)$ |
| $g^{(6)}$ | $\begin{pmatrix} 0 & 0 & 1 \\ 0 & 1 & 0 \\ -1 & 0 & 0 \end{pmatrix}$ | $[1,-1,-1]$ | $4\pi/3$ | $-e_{12} + e_{23} - e_{31}$ | $(-2 \ 3 \ -1)$ |
| $g^{(7)}$ | $\begin{pmatrix} 0 & 0 & 1 \\ 0 & -1 & 0 \\ -1 & 0 & 0 \end{pmatrix}$ | $[-1,-1,-1]$ | $2\pi/3$ | $e_{13} - e_{21} - e_{32}$ | $(3 \ -1 \ -2)$ |
| $g^{(8)}$ | $\begin{pmatrix} 0 & 0 & -1 \\ 0 & 1 & 0 \\ 1 & 0 & 0 \end{pmatrix}$ | $[-1,-1,-1]$ | $4\pi/3$ | $-e_{12} - e_{23} + e_{31}$ | $(-2 \ -3 \ 1)$ |
| $g^{(9)}$ | $\begin{pmatrix} 0 & -1 & 0 \\ 0 & 0 & -1 \\ 1 & 0 & 0 \end{pmatrix}$ | $[1,0,0]$ | $\pi$ | $e_{11} - e_{22} - e_{33}$ | $(1 \ -2 \ -3)$ |
| $g^{(10)}$ | $\begin{pmatrix} 0 & 1 & 0 \\ 0 & 0 & -1 \\ -1 & 0 & 0 \end{pmatrix}$ | $[0,1,0]$ | $\pi$ | $-e_{11} + e_{22} - e_{33}$ | $(-1 \ 2 \ -3)$ |
| $g^{(11)}$ | $\begin{pmatrix} 0 & -1 & 0 \\ 0 & 0 & 1 \\ -1 & 0 & 0 \end{pmatrix}$ | $[0,0,1]$ | $\pi$ | $-e_{11} - e_{22} + e_{33}$ | $(-1 \ -2 \ 3)$ |
\[
\begin{align*}
g^{(1)} &= \begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix}, & \text{any} & 0 & e_{11} + e_{22} + e_{33} & \sigma_{15} = (1 \ 2 \ 3) \\
g^{(2)} &= \begin{pmatrix} 0 & -1 & 0 \\ 1 & 0 & 0 \\ 0 & 0 & 1 \end{pmatrix}, & [0,0,1] & \pi/2 & -e_{12} + e_{21} + e_{33} & \sigma_1 = (-2 \ 1 \ 3) \\
g^{(3)} &= \begin{pmatrix} -1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix}, & [0,0,1] & 3\pi/2 & e_{12} - e_{21} + e_{33} & \sigma_2 = (2 \ -1 \ 3) \\
g^{(4)} &= \begin{pmatrix} 1 & 0 & 0 \\ 0 & 0 & -1 \\ 0 & 1 & 0 \end{pmatrix}, & [1,0,0] & \pi/2 & e_{11} - e_{23} + e_{32} & \sigma_3 = (1 \ -3 \ 2) \\
g^{(5)} &= \begin{pmatrix} 0 & 1 & 0 \\ 0 & 0 & 1 \\ 0 & -1 & 0 \end{pmatrix}, & [1,0,0] & 3\pi/2 & e_{11} + e_{23} - e_{32} & \sigma_4 = (1 \ 3 \ -2) \\
g^{(6)} &= \begin{pmatrix} 0 & 0 & -1 \\ 1 & 0 & 0 \\ 0 & 0 & 1 \end{pmatrix}, & [1,-1,1] & 2\pi/3 & -e_{12} - e_{23} + e_{31} & \sigma_5 = (-2 \ -3 \ 1) \\
g^{(7)} &= \begin{pmatrix} -1 & 0 & 0 \\ 0 & -1 & 0 \\ 0 & 0 & 1 \end{pmatrix}, & [1,-1,1] & 4\pi/3 & e_{13} - e_{21} - e_{32} & \sigma_6 = (3 \ -1 \ -2) \\
g^{(8)} &= \begin{pmatrix} 0 & 0 & 1 \\ -1 & 0 & 0 \\ 0 & 0 & -1 \end{pmatrix}, & [-1,1,1] & 2\pi/3 & -e_{12} + e_{23} - e_{31} & \sigma_7 = (-2 \ 3 \ -1) \\
g^{(9)} &= \begin{pmatrix} -1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & -1 \end{pmatrix}, & [-1,1,1] & 4\pi/3 & -e_{13} - e_{21} + e_{32} & \sigma_8 = (-3 \ -1 \ 2) \\
g^{(10)} &= \begin{pmatrix} 0 & 1 & 0 \\ 0 & 0 & 1 \\ 1 & 0 & 0 \end{pmatrix}, & [-1,-1,-1] & 2\pi/3 & e_{12} + e_{23} + e_{31} & \sigma_9 = (2 \ 3 \ 1) \\
g^{(11)} &= \begin{pmatrix} 1 & 0 & 0 \\ 0 & 0 & 1 \\ 0 & 1 & 0 \end{pmatrix}, & [-1,-1,-1] & 4\pi/3 & e_{13} + e_{21} + e_{32} & \sigma_{10} = (3 \ 1 \ 2) \\
g^{(12)} &= \begin{pmatrix} 0 & 0 & 1 \\ 0 & 1 & 0 \\ -1 & 0 & 0 \end{pmatrix}, & [0,1,0] & \pi/2 & e_{13} + e_{22} - e_{31} & \sigma_{11} = (3 \ 2 \ -1) \\
g^{(13)} &= \begin{pmatrix} 0 & 1 & 0 \\ 0 & 0 & -1 \\ 1 & 0 & 0 \end{pmatrix}, & [0,1,0] & 3\pi/2 & -e_{13} + e_{22} + e_{31} & \sigma_{12} = (-3 \ 2 \ 1) \\
g^{(14)} &= \begin{pmatrix} 0 & 0 & -1 \\ 0 & 1 & 0 \\ -1 & 0 & 0 \end{pmatrix}, & [1,1,-1] & 2\pi/3 & e_{12} - e_{23} - e_{31} & \sigma_{13} = (2 \ -3 \ -1) \\
\end{align*}
\]
Appendix B  Proof of Lemma 7

Proof. First, note that for the matrices $e_{ij}$ and $e_{kl}$ defined in Definition 4, it holds that

$$e_{ij}e_{kl} = \begin{cases} e_{il} & \text{if } k = j, \\ -e_{il} & \text{if } k = -j, \\ 0_{3 \times 3} & \text{else.} \end{cases}$$

(55)
In addition, for any single entry matrix defined in Definition 4, it follows by a direct calculation that $e_{ij}^T = e_{ji}$. By expressing $g_1$ and $g_2$ using Lemma 6, we have

$$g_1 = e_1\sigma_1(1) + e_2\sigma_1(2) + e_3\sigma_1(3), \quad g_2 = e_1\sigma_2(1) + e_2\sigma_2(2) + e_3\sigma_2(3).$$

For (36), we have that for $m = 1, 2, 3$

$$g_1^T e_{mm} g_2 = (e_1\sigma_1(1) + e_2\sigma_1(2) + e_3\sigma_1(3))^T e_{mm} (e_1\sigma_2(1) + e_2\sigma_2(2) + e_3\sigma_2(3))$$

$$= (e_1\sigma_1(1)1 + e_1\sigma_2(2) + e_1\sigma_3(3)) e_{mm} (e_1\sigma_2(1) + e_2\sigma_2(2) + e_3\sigma_2(3))$$

$$= e_1\sigma_1(m)\sigma_2(m),$$

where the last equality follows by (55). For (37), we use (36) and obtain

$$g_1^T e_{mm} g_1 = e_1\sigma_1(m)\sigma_1(m).$$

For (38),

$$\{g_1^T e_{mr} g_2\}_{m, r = 1}^3 = \{(e_1\sigma_1(1) + e_2\sigma_1(2) + e_3\sigma_1(3))^T e_{mr} (e_1\sigma_2(1) + e_2\sigma_2(2) + e_3\sigma_2(3))\}_{m, r = 1}^3$$

$$= \{(e_1\sigma_1(1)1 + e_1\sigma_2(2) + e_3\sigma_3(3)) e_{mr} (e_1\sigma_2(1) + e_2\sigma_2(2) + e_3\sigma_2(3))\}_{m, r = 1}^3$$

$$= \{e_1\sigma_1(m)\sigma_2(r)\}_{m, r = 1}^3,$$

where the last equality follows by (55), and thus

$$\{\pm g_1^T e_{mr} g_2\}_{m, r = 1}^3 = \{\pm e_1\sigma_1(m)\sigma_2(r)\}_{m, r = 1}^3 = \{\pm e_{mr}\}_{m, r = 1}^3.$$

\[\square\]

**Appendix C  Constructing $SO_G(3)$**

We denote by $SO_G(3) \subset SO(3)$ the finite subset of rotations for the symmetry group $G$ on which we search for the optimum of the score function $\pi_{ij}$ of (23). A naive choice for $SO_G(3)$ would be an almost equally spaced grid of rotations from $SO(3)$, denoted as $\tilde{SO}(3)$ and defined below. However, the symmetry of $G$ allows us to significantly reduce the number of rotations in this naive set while maintaining the same accuracy of our algorithm. Note that for any $Q_r, Q_s \in SO(3)$ and $g \in G$, it holds that $\{Q_1^T g^{(k)} Q_s\}_{k=1}^n = \{Q_1^T g^{(k)} g Q_s\}_{k=1}^n$, and so the set of local coordinates $\{(\alpha_{Q_r, Q_s}^{k,1}, \alpha_{Q_r, Q_s}^{k,2})\}_{k=1}^n$ is equal to the set of local coordinates $\{(\alpha_{Q_r, gQ_s}^{k,1}, \alpha_{Q_r, gQ_s}^{k,2})\}_{k=1}^n$. Thus, keeping both $Q_s$ and $g Q_s$ in $SO_G(3)$ is redundant. Consequently, our objective is to find all pairs of rotations $Q_s, Q_l \in \tilde{SO}(3)$ for which there exists $g^{(k)} \in G \setminus I$ such that $Q_s = g^{(k)} Q_l$, and filter either $Q_s$ or $Q_l$ from $\tilde{SO}(3)$. The resulting set would be $SO_G(3)$.

Since $\tilde{SO}(3)$ is finite, an exact equality between $Q_s$ and $g^{(k)} Q_l$ is unlikely. Therefore, the proximity between $Q_s$ and $g^{(k)} Q_l$ is determined up to pre-defined thresholds, based on their representation using viewing direction and in-plane rotation (see [18]) as follows. The viewing
directions of $Q_s$ and $g^{(k)}Q_l$ are given by their third columns $Q_s^3$ and $(g^{(k)}Q_l)^3$, respectively. If $Q_s$ and $g^{(k)}Q_l$ are two rotations with the same viewing direction, i.e., $\langle Q_s^3, (g^{(k)}Q_l)^3 \rangle = 1$, then the rotation matrix $Q_s^T g^{(k)}Q_l$ is an in-plane rotation matrix which has the form
\[
\begin{pmatrix}
\cos(\theta_{sl}^{(k)}) & -\sin(\theta_{sl}^{(k)}) & 0 \\
\sin(\theta_{sl}^{(k)}) & \cos(\theta_{sl}^{(k)}) & 0 \\
0 & 0 & 1
\end{pmatrix},
\] (56)
where $\theta_{sl}^{(k)} \in [0, 360^\circ]$ is the in-plane rotation angle (see [18]). If $\theta_{sl}^{(k)} = 0$, then $Q_s^T g^{(k)}Q_l = I$, and so $Q_s = g^{(k)}Q_l$. Hence, we define two thresholds; the viewing direction threshold $\epsilon_1$, and the in-plane rotation angle threshold $\epsilon_2$. For the viewing direction threshold, we define $\epsilon_1 = 5^\circ$ along with the condition
\[
\langle Q_s^3, (g^{(k)}Q_l)^3 \rangle > \cos(\epsilon_1).
\] (57)
Satisfying condition (57) implies that the rotations $Q_s$ and $g^{(k)}Q_l$ have nearby viewing directions, and so it is reasonable to assume that the angle
\[
\tilde{\theta}_{sl}^{(k)} = \arctan \left( \frac{(Q_s^T g^{(k)}Q_l)_{2,1}}{(Q_s^T g^{(k)}Q_l)_{1,1}} \right)
\] (58)
approximates the in-plane rotation angle $\theta_{sl}^{(k)}$ of (56). We therefore define $\epsilon_2 = 5^\circ$ along with the condition
\[
\tilde{\theta}_{sl}^{(k)} < \epsilon_2.
\] (59)
Once both conditions (57) and (59) hold, the proximity between $Q_s$ and $g^{(k)}Q_l$ is sufficient to remove either $Q_s$ or $Q_l$ from $\tilde{SO}(3)$.

Of course, it is possible to replace the proximity measure we have used above with any other proximity measure. The advantage of the measure we use is its simple geometric interpretation, which allows to easily set and interpret its thresholds.

It remain to show how to construct the set $\tilde{SO}(3)$, which is the input of the above pruning procedure. To that end, we let $L$ be a positive integer, and let $\tau, \theta, \varphi$ denote Euler angles. We construct $\tilde{SO}(3)$ by sampling the Euler angles in equally spaced increments as follows. First, we sample $\tau \in \{0, \ldots, \frac{\pi}{L} \}$ at $\lfloor \frac{L}{4} \rfloor$ points. Then, for each $\tau$, we sample $\theta \in \{0, \ldots, \pi \}$ at $\lfloor \frac{L}{2} \sin(\tau) \rfloor$ points. Finally, for each pair $(\tau, \theta)$, we sample $\varphi \in \{0, \ldots, 2\pi \}$ at $\lfloor \frac{L}{2} \sin(\tau) \sin(\theta) \rfloor$ points. For each $(\tau, \theta, \varphi)$ on this grid, we compute a corresponding rotation matrix $R$ by
\[
R = R_z(\tau)R_y(\theta)R_x(\varphi),
\]
Proof. A classification of the closed subgroups of $SO(3)$ is given in [6], stating that every closed subgroup of $SO(3)$ is conjugate to one of $SO(3)$, $O(2)$, $SO(2)$, $D_n(n \geq 2)$, $C_n(n \geq 2)$, $\mathbb{T}$, $\mathbb{O}$, $\mathbb{I}$ (the icosahedral symmetry), $\mathbb{1}$ (the trivial group). Moreover, $\mathbb{T}$ and $\mathbb{O}$ are closed subgroups of $SO(3)$. Since for topological groups the normalizer of a closed subgroup is closed (Lemma 10 below) and since $SO(3)$ is indeed a topological group, the normalizers of the closed subgroups $\mathbb{T}$ and $\mathbb{O}$ in $SO(3)$, i.e. $N_{SO(3)}(\mathbb{T})$ and $N_{SO(3)}(\mathbb{O})$, are also closed subgroups, thus conjugate to one of the closed subgroups of $SO(3)$.

By definition of the normalizer, $G \subseteq N_{SO(3)}(G)$, which precludes $O(2)$, $SO(2)$, $D_n(n \geq 2)$, $C_n(n \geq 2)$ and $\mathbb{1}$ from being the normalizers of $\mathbb{T}$ or $\mathbb{O}$, since each has at most one symmetry axis of order larger than 2, while both $\mathbb{T}$ and $\mathbb{O}$ have more than one such axis. In addition, $SO(3)$ and $\mathbb{I}$ are simple groups [21, 2], and so have no non-trivial normal subgroups. By definition of the normalizer, $G$ is a normal subgroup of $N_{SO(3)}(G)$. Thus, since $SO(3)$ and $\mathbb{I}$ have no non-trivial normal subgroups, neither $\mathbb{T}$ nor $\mathbb{O}$ are normal subgroups of $\mathbb{I}$ or $SO(3)$, which precludes $SO(3)$ and $\mathbb{I}$ from being the normalizers of $\mathbb{T}$ or $\mathbb{O}$. Since $\mathbb{T}$ is normal in $\mathbb{O}$ [2], we have that $\mathbb{O} \subseteq N_{SO(3)}(\mathbb{T})$ and thus it must hold that $N_{SO(3)}(\mathbb{T}) = \mathbb{O}$ and $N_{SO(3)}(\mathbb{O}) = \mathbb{O}$. ■

Lemma 10. Suppose $\tilde{H}$ is a topological group. Then, the normalizer of a closed subgroup $\tilde{G}$ of $\tilde{H}$

$$N_{\tilde{H}}(\tilde{G}) = \{\tilde{h} \in \tilde{H} : \tilde{h}^{-1}\tilde{G}\tilde{h} = \tilde{G}\}$$

is a closed subgroup.

Proof. Fix $\tilde{g} \in \tilde{G}$ and define the map $f_{\tilde{g}} : \tilde{H} \to \tilde{H}$ by $f_{\tilde{g}}(\tilde{h}) = \tilde{h}^{-1}\tilde{g}\tilde{h}$. Since $\tilde{H}$ is a topological group, $f_{\tilde{g}}$ is continuous as the composition of multiplication and inversion maps. Thus, the preimage of the closed subgroup $\tilde{G}$ under $f_{\tilde{g}}$, defined by $f_{\tilde{g}}^{-1}(\tilde{G}) = \{\tilde{h} \in \tilde{H} : f_{\tilde{g}}(\tilde{h}) \in \tilde{G}\} = \{\tilde{h} \in \tilde{H} : \tilde{h}^{-1}\tilde{g}\tilde{h} \in \tilde{G}\}$, is closed. As any intersection of closed sets is closed, the intersection

$$\bigcap_{\tilde{g} \in \tilde{G}} f_{\tilde{g}}^{-1}(\tilde{G}) = \{\tilde{h} \in \tilde{H} : \tilde{h}^{-1}\tilde{g}\tilde{h} \in \tilde{G} \forall \tilde{g} \in \tilde{G}\} = N_{\tilde{H}}(\tilde{G})$$

is a closed subgroup.
is closed.

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