A program for phase identification using diffractograms obtained from TEM structure images

R. Galicia\textsuperscript{a}, R. Herrera\textsuperscript{a,∗}, J. L. Rius\textsuperscript{b}, C. Zorrilla\textsuperscript{a}, and A. Gómez\textsuperscript{a}

\textsuperscript{a}Departamento de Materia Condensada, Instituto de Física, Universidad Nacional Autónoma de México, 20-364, México, 01000, D.F., México.
\textsuperscript{b}Departamento de Sistemas Complejos, Instituto de Física, Universidad Nacional Autónoma de México, 20-364, México, 01000, D.F., México.

\textsuperscript{∗}e-mail: rherrera@fisica.unam.mx

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In this work a computer program for the indexing of diffractograms is presented. The diffractograms are obtained by means of a discrete Fourier transform from high resolution electron microscope images. The program requires the use of x-ray diffraction data files together with a fast Fourier transform program, for this purpose we used the Digital Micrograph software.

Keywords: Electron microscopy; X-ray diffraction; difractograms; indexing.

Se presenta un programa de cómputo que ayuda a identificar los planos cristalográficos (indexar difractogramas) que aparecen al aplicar la Transformada Rápida de Fourier (FFT) a una imagen digital de alta resolución obtenida con un microscopio electrónico de transmisión de alta resolución. El programa requiere el uso de los archivos de Rayos X de uso común y de un programa que obtenga la FFT, para este caso se utiliza el software de Digital Micrograph.

Descriptores: Microscopía electrónica; rayos X; difractogramas; indexación.

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1. Introduction

Diffraction is doubtless one of the most powerful techniques for the elucidation of condensed matter structure. While x-ray, neutron, and electron diffraction all provide basic information on symmetry and atomic arrangements, electron diffraction is unique in its ability to identify nanometric crystal phases within a matrix or on a substrate. The atomic spacing information in a high resolution TEM image of a crystal can be extracted by taking the Fourier transform of the image. This yields a diffractogram of spots (points) in an arrangement similar to an electron diffraction pattern of the same specimen; however, the diffractogram can be obtained from regions of specimen on the order of a nanometer rather than the micrometer typical of conventional selected area electron diffraction.

To identify a small crystalline phase, the specimen is usually tilted so that the electron beam is directed along a major zone axis in the crystal, thus yielding a symmetrical array of diffraction spots. The electron diffraction pattern provides the interplanar spacings of the (hkl) planes belonging to a given [uvw] zone. Indexing consists of assigning the specific (hkl) indices to the various spots. Measured distances from the diffractogram center (that corresponds to the transmitted beam in the diffraction pattern) to the (hkl) diffraction spots yield the $d_{hkl}$-spacings for each spot in the electron diffraction pattern or in the Fourier transform of the lattice image or structure image (the diffractogram). The set of $d$-spacings obtained in this way and the angles between pairs of spots allow identification of the unknown phase by comparison with structures of known substances.

Once the structure of a phase is known, the major task is to index the spots in the diffraction pattern or diffractogram. A survey of related procedures for phase identification and diffraction spot indexing can be found in the Refs. 1 to 6.

This article presents a program for identifying phases using the diffractogram derived from high-resolution TEM structure images that show atomic columns. The program is based on the zone axis method of crystallography [6], and it compares the diffraction pattern data from the unknown sample with that of known compounds and indexes the unknown spots when the best fit is obtained. This article describes the steps involved in the operation of the software and gives a practical example.

It is important to stress that we work with actual electron micrographs and not with simulations obtained, for example, with JEMS or related programs. In the diffractogram the “spots” are identified; the rest of the spatial frequencies are not used.

2. Program Description

A program for the analysis of a diffractogram, obtained from the high-resolution image by means of the Fast Fourier transform (FFT) algorithm, has been developed following a standard architecture in which the Front End (Client), requests services from the Back-end (server).

The graphical user interface and the operational algorithm have been programmed under Delphi 6 (in the client), a platform providing a development environment integrated to the Windows environment. It is an object-oriented language that has considerable support for databases.
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The services are requested from an SQL (Structured Query Language) Server 2005 Express database, a product freely distributed, designed as a data base platform that is easy to use where one has stored procedures (SQL statements) that will be called by client procedures (Delphi).

For image analysis, it is necessary to know some of the parameters involved in image acquisition via DigitalMicrograph: the electron wavelength, camera length, accelerating voltage, magnification of the image, etc. All the images were obtained with a JEOL FEG 2010 microscope equipped with image capture software.

The program can be obtained freely from the authors. Although at present the program uses the DigitalMicrograph software, it is possible to use other platforms such as Matlab for the purpose of calculating the Fourier transforms.

3. User Screens

The program contains five screens. In the first one (Fig. 1), the user keys in the data provided by the DigitalMicrograph (DM) program in order to calculate the reference parameters such as the magnitudes of vectors defining the positions (with respect to the coordinate origin) of each of the spots in the diffractogram. The program requires at least two points and the angles between the points relative to the center spot.

The second screen (Fig. 2) introduces the lattice parameter data taken from the X-ray crystallographic data tables relevant to the sample under consideration. These data are stored in a database for future use unless the user deletes them.

In the third screen (Fig. 3), the actual calculations are performed. The user selects the X-ray crystallographic database and gives a tolerance for the maximum error allowed between calculated angles and measured ones (this parameter is called Delta in the program). This part is based upon the so-called zone-axis method; it uses the standard formulas for interplanar angles and spacings as functions of Miller indices \[6\] for cubic, tetragonal, orthorhombic, and hexagonal crystals. Delta is the difference between the measured angle on the diffractogram and the angle as calculated by the program for a pair \((h_1,k_1,l_1)\) and \((h_2,k_2,l_2)\) (taken as a possible solution). While the program requires at least two points, a typical analysis of a diffractogram will use three or more points.

The fourth (Fig. 4) and fifth screens assist the user in selecting the best solution, starting from the zone axis or using

![Figure 1. First screen used for data identification and input.](image1)

![Figure 2. Second screen used for the input of x-ray crystallographic data for various materials.](image2)

![Figure 3. Third screen where the actual calculations are performed.](image3)

![Figure 4. Fourth screen where the possible solutions are displayed.](image4)
the strategy followed to reach a plausible solution can be summarized as follows:

(a) Normally the user has an idea of what sorts of compounds are worth examining. In order to do this the user has to make sure that the relevant X-ray cards have been captured beforehand; otherwise he will have to capture the data using the second screen (Fig. 2). In case the user has no idea at all concerning the nature of his sample he will have to run the program over all the entries in the database, but of course this will not warrant that the “right” compound is there.

(b) From the X-ray crystallographic data table, two sets with seven Miller indices \{h,k,l\} are selected. Each set is selected so as to correspond to the closest \(d_{hkl}\) spacing (in Å) to \(d_1\) and \(d_2\), respectively.

Consider the example given in Fig. 5, the selected spots on the FFT correspond to distances \(d_1 = 2.33\) Å, \(d_2 = 1.97\) Å and \(d_3 = 2.28\) Å. Upon examination of the corresponding x-ray crystallographic data card for Fe\(_{0.902}\)O (cubic), we see (Table I) that the distance closest to \(d_1 = 2.33\) Å is \(d = 2.3552\) Å; for \(d_2 = 1.97\) Å is \(d = 1.9672\) Å and for \(d_3 = 2.28\) Å is \(d = 2.2804\) Å.

Given that the distances measured on the FFT depend on the cursor position (set manually by the user), there is some associated error. In order to take this into account, we take three values above and three values below as “tolerance” and in this way we end up with seven values to consider. The program takes next the Miller indices and proceeds as described below. Should it happen that the chosen distance is the first in the card, the program takes this and the next six values. A similar procedure is followed if the chosen distance is the last one in the card.

(c) For each pair \((h_1,k_1,l_1)\) and \((h_2,k_2,l_2)\) (and all equivalent permutations and sign changes), \(\cos(\theta_{12})\) calculated and compared to the value measured from the diffractogram (taken from the first screen). If \(\Delta \theta_{12}\) is larger than the given tolerance, the program rejects the solution.

(d) If \(\Delta \theta_{12}\) is less than or equal to the tolerance, the solution is stored in a file and the program calculates \(d_{12}\), the distance between consecutive planes in the same family \{hkl\}, and the corresponding zone axis [uvw].

(e) The procedure is repeated with another pair of indices until all possible permutations and/or sign changes are included. The program will repeat steps (c) and (d) more than 100 thousand times, but the number of possible solutions will depend on the given \(\Delta \theta_{12}\).

(f) The program repeats (b) but only for the third spot and takes as possible solutions the \((h_3,k_3,l_3)\) stored in the file (step e) for the second spot.

(g) With these new sets, steps (c) and (d) take into account not just the value of \(\Delta \theta_{12}\) but also the zone axis, which should be the same as the one found for the first two spots.

(h) The solutions are stored in a new file. It may happen that there is no solution, in which case the program will indicate that the search is over, but the solution screen will remain blank.

5. An application of the program

Figure 5 shows an example of this procedure applied to a 3 nm nanoparticle. The particle was synthesized by means of bioreduction using alfalfa as a reducing agent in distilled water with FeNH\(_4\)(SO\(_4\))\(_2\)12H\(_2\)O and a buffer solution at pH=10. The high resolution image was obtained with a JEOL2010-FEG microscope. Given that in this case we have followed all the steps necessary to obtain iron oxide nanoparticles we will search for our solution among the X-ray cards [7] for Fe\(_2\)O\(_4\) (Magnetite, cubic, PDF#890688), Fe\(_{0.902}\)O (Wuestite, cubic, PDF#862316), FeO (Wuestite, cubic, PDF#890687), FeOOH (Lepidocrocite, orthorhombic, PDF#741877) and Fe\(_2\)O\(_3\) (Maghemite, cubic, PDF#391346).

The FFT was obtained from the image with the Digital Micrograph (DM) software, (Fig. 5), three spots were selected for analysis. The positions of these points with respect to the 000 spot were \(P_1 = (1.64, -3.96)\), \(P_2 = (4.92, -1.23)\), and \(P_3 = (3.41, 2.73)\), and from them the program calculated the corresponding interplanar spacings, \(d_1 = 2.33 \) Å, \(d_2 = 1.97 \) Å and \(d_3 = 2.28 \) Å. The angles between vectors from the center spot to the other spots were \(\theta_{1,2} = 53.47^\circ\) and \(\theta_{2,3} = 52.72^\circ\).
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FIGURE 5. Example of a Fe₂O₃ nanoparticle: (left) structure image of 3 nm particle and (right) indexed diffractogram showing the points (spots) used for analysis.

After running the program using the x-ray crystallographic data from the table for the five chosen candidate compounds, it was found that the best fit was with \( \Delta 1_2 = 1.07^\circ \), \( \angle 1_2 = 52.4^\circ \), \( \Delta 2_3 = 0.32^\circ \), \( \angle 2_3 = 52.4^\circ \). For a Zone Axis of \([1,0,1]\), the indexing of the three planes was:

- Plane 1 = \((-1,1,1)\)
- Plane 2 = \((0,1,0)\)
- Plane 3 = \((1,1,-1)\)

Which yielded interplanar spacings of \( d_1 = 2.36\ \text{Å} \), \( d_2 = 1.93\ \text{Å} \), and \( d_3 = 2.36\ \text{Å} \). These values suggest that the unknown nanoparticle is FeOOH with an orthorhombic structure as shown in Table II.

<table>
<thead>
<tr>
<th>Plane ( a \rightarrow b )</th>
<th>Compound</th>
<th>( \Delta a \rightarrow b )</th>
<th>( \text{Spacing}_a )</th>
<th>( \text{Spacing}_b )</th>
<th>Zone Axis</th>
</tr>
</thead>
<tbody>
<tr>
<td>1-2 Fe₂O₃</td>
<td>1.266</td>
<td>2.42</td>
<td>2.10</td>
<td>([1,1,0])</td>
<td></td>
</tr>
<tr>
<td>2-3 FeO</td>
<td>2.016</td>
<td>2.10</td>
<td>2.42</td>
<td>([1,1,0])</td>
<td></td>
</tr>
<tr>
<td>1-2 Fe₀.₉₀₂O</td>
<td>1.266</td>
<td>2.24</td>
<td>1.94</td>
<td>([1,1,0])</td>
<td></td>
</tr>
<tr>
<td>2-3 FeO</td>
<td>2.016</td>
<td>1.94</td>
<td>2.24</td>
<td>([1,1,0])</td>
<td></td>
</tr>
<tr>
<td>1-2 FeOOH</td>
<td>1.07</td>
<td>2.36</td>
<td>1.93</td>
<td>([1,0,1])</td>
<td></td>
</tr>
<tr>
<td>2-3 Fe₂O₃</td>
<td>0.32</td>
<td>1.93</td>
<td>2.36</td>
<td>([1,0,1])</td>
<td></td>
</tr>
<tr>
<td>1-2 Fe₂O₃</td>
<td>1.266</td>
<td>2.41</td>
<td>2.08</td>
<td>([0,1,1])</td>
<td></td>
</tr>
<tr>
<td>2-3 Fe₂O₃</td>
<td>2.016</td>
<td>2.08</td>
<td>2.41</td>
<td>([0,1,1])</td>
<td></td>
</tr>
</tbody>
</table>

Thus, in Table II we have presented the best solutions for the tested structures, and they are very similar. However, in this case for FeOOH we get the lowest Delta values and the best fitting with the measured distances.

6. Discussion

The program requires the use of the DigitalMicrograph (DM) program and X-ray crystallographic data for various known substances. The DM package was used to analyze the images, to Fourier-transform them, and to obtain the coordinates of the various spots. Under DM it is possible to work at \( 64 \times 64 \), \( 128 \times 128 \) and \( 256 \times 256 \) resolutions. This implies varying errors in the spot positions as a function of pixel size, so the program works within a given tolerance. The solutions are understood as best fits. Given the spot positions, the program searches for a fit to interplanar distances and angles between planes. Currently, it works for cubic, tetragonal, orthorhombic, and hexagonal structures.

By considering all the possibilities, for every pair of spots, the program can produce up to 100,000 solutions in general. This number is quickly reduced to a few dozen possibilities by ruling out those where the angles produce matches outside the user-given tolerance. The program produces a list of possible solutions, and the final choice is left to the user.

Some of the virtues of the program are that it is friendly and easy to use, no special expertise is required, only modest memory resources are needed, the search takes only a few seconds, and the lattice parameter data captured can be stored in a data base for future use. Since it is based on a micrograph, smaller regions of the sample (a few nanometers) can be studied than with selected area electron diffraction. Of course our approach is not meant to be a replacement for convergent beam electron diffraction (CBED), but it can be useful not only when CBED is unavailable but also when a set of high resolution micrographs is available and crystallographic information of small areas is desired.

The main limitation of the program is that the user has to “guess” (from the chemical and physical conditions of sample preparation) what are the likely structures and to include the corresponding X-ray data; it is impossible to include all the available cards.

The accuracy of the method is limited by the precision with which the magnification in the micrograph is known. In principle it would be desirable to have a “built in” standard together with the actual sample.

7. Conclusions

Diffractograms obtained as Fourier Transforms from a digital TEM image can be used to identify an unknown crystal in the image. This article describes a program devised to achieve this. The program has a graphical user interface that allows the user to control various windows, input boxes, and processes. Measured data can be compared with data from candidate compounds to arrive at the likely identity of the unknown crystal. Crystals of only a few nanometers in size can be analyzed by this method.

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