Refinement of anomalous dispersion correction parameters in single-crystal structure determinations

Florian Meurer,a Oleg V. Dolomanov,b Christoph Hennig,c,d Norbert Peyerimhoff,∗ Florian Kleemiss,a Horst Puschmannb,e and Michael Bodensteinera∗

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Correcting for anomalous dispersion is part of any refinement of an X-ray diffraction crystal structure determination. The procedure takes the inelastic scattering in the diffraction experiment into account. This X-ray absorption effect is specific to each chemical compound and is particularly sensitive to radiation energies in the region of the absorption edges of the elements in the compound. Therefore, the widely used tabulated values for these corrections can only be approximations as they are based on calculations for isolated atoms. Features of the unique spatial and electronic environment that are directly related to the anomalous dispersion are ignored, although these can be observed spectroscopically. This significantly affects the fit between the crystallographic model and the measured intensities when the excitation wavelength in an X-ray diffraction experiment is close to an element’s absorption edge. Herein, we report on synchrotron multi-wavelength single-crystal X-ray diffraction, as well as X-ray absorption spectroscopy experiments which we performed on the molecular compound Mo(CO)6 at energies around the molybdenum K edge. The dispersive ($f_0$) and absorptive ($f_00$) terms of the anomalous dispersion can be refined as independent parameters in the full-matrix least-squares refinement. This procedure has been implemented as a new feature in the well-established OLEX2 software suite. These refined parameters are in good agreement with the independently recorded X-ray absorption spectrum. The resulting crystallographic models show significant improvement compared to those employing tabulated values.

1. Introduction

With the increasing capabilities of X-ray diffraction equipment, the deficiencies of the conventional crystallographic model become more and more apparent. This model is based on the measured intensities of each Bragg reflection according to Equations (1) and (2), where the structure factor $F$ is calculated by a sum over $N$ atoms in the unit cell with the respective atomic form factor $f_n$ and the anisotropic displacement parameters $T_{ij}$.

$$F(hkl) = \sum_{n=1}^{N} f_n e^{2\pi i (hX_n + kY_n + lZ_n)} T_{ij}$$

Resonant scattering, i.e. discrete electron transitions in the form of photoabsorption of incident radiation, is treated with the parameters $f'$ and $f''$ – the real (dispersive) and imaginary (absorptive) part of the dispersion correction. Those are added to the atomic form factor $f_{n,0}$ according to Equation (3).
\[ f_n = f_{n,0} + f'_n + if''_n \]  

(3)

The impact of these parameters depends on the excitation energy, the type of atoms and their specific chemical bonding situation in the respective crystal structure. The original values for dispersion corrections commonly used so far are taken from tabulated libraries and based on calculations by Cromer (1965), Cromer & Liberman (1970) and Hönl (1933). They have been improved over several decades and are in good agreement with the experimental data of isolated atoms (Cromer & Mann, 1968). However, the tabulated literature values only consider the excitation energy and the atom type, completely ignoring the uniqueness of the atomic environment.

Fig. 1 shows an example of the effects of these correction terms for the atomic form factor of molybdenum at its K absorption edge at 20 000 eV. The real part of the correction \( f' \) can be understood as a direct and the imaginary part \( f'' \) as a phase-shifted contribution to the scattering power, which is constant across the entire range of resolution [see Fig. 1(c)]. As shown in Fig. 1(a), the amplitude of the effective atomic form factor \( f \) is significantly diminished by these dispersion corrections. At the K absorption edge of molybdenum \((Z = 42)\), the resulting scattering power is reduced by as much as 9.6 electrons relative to a pure Thomson scattering model. At this energy, the corrected atomic form factor closely resembles the form factor of germanium \([Z = 32; \text{Fig. 1(b)}]\). In contrast to molybdenum, the germanium atomic form factor is only slightly affected by a dispersion correction at this energy. This effect even allows a wrongly assigned atom type to yield a similar or even better crystallographic model (Guss et al., 1989).

The frequency of an absorption edge is mostly determined by the given element. It has been shown that this frequency is

\[ f''(\omega) = \frac{m_e c \omega^4}{4 \pi N e^2} \]  

(4)

\[ f'(\omega) = \frac{2}{\pi} \int_{\omega_K}^{\infty} \frac{\omega f''(\omega')}{\omega^2 - \omega'^2} d\omega' \]  

(5)

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![Figure 1](image_url)

Dispersions corrections for the phase-dependent \( f(000) \) for (a) molybdenum and (b) comparison to germanium at 20 000 eV; (c) effect of \( f' \) and \( f'' \) on the atomic form factor \( f_{\text{Ge}} \); (d) resolution-dependent difference between \( f_{\text{Ge}} \) and \( f \).
further affected by the charge of the respective atom (Ankudinov, 1998). Spatzal et al. (2016) applied their spatially resolved anomalous dispersion (SpReAD) refinement to determine the individual oxidation states of the Fe atoms in nitrogenase from diffraction data (Einsle et al., 2007). To the best of our knowledge, such an experimental determination of anomalous dispersion parameters was practically never carried out in small molecule or solid-state crystallography.

The individual absorption spectrum exhibits additional features around the absorption edge which originate from electronic transitions into half-occupied and unoccupied orbitals, as well as into the continuum above the Fermi level (Hennig, 2007). Measuring these spectral features is the subject of the X-ray absorption near-edge structure (XANES) technique, and also of extended X-ray absorption fine structure (EXAFS) spectroscopy. The spectral fine structure at excitation energies above the absorption edge depends on the distances to neighbouring atoms and is therefore unique to each structure. None of these spectral features are considered in the tabulated dispersion values. Therefore, the application of these literature parameters is insufficient for X-ray crystallography, especially near absorption edges. Structures obtained with an incident photon energy near the absorption edge of a given atom show artefacts and result in poor crystallographic models (Dittrich et al., 2015). This gave rise to a common practice of avoiding single-crystal diffraction measurements near absorption edges.

2. Results and discussion

Since the exact energy of the absorption edge and the spectral features in the near vicinity to the edge are unique to the atom in its specific chemical environment, a simple approximation using calculations based on independent neutral atoms is always incorrect and leads to problems in the crystallographic model. Artefacts occur in the Fourier map and affect the

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**Figure 2**

(a) X-ray absorption spectrum of Mo(CO)₆ (top black line) and $f'$ calculated from it (bottom black line), Sasaki’s tabulated (x symbol blue) and refined values (+ symbol red) for $f''$ and $f'$, and the resulting quality parameters (b) $R_1$ and (c) $wR_2$ of refined structure models.
Single-crystal X-ray diffraction experiments were performed at these energies, as well as at 19 900 and 20 100 eV as reference data well below and above the absorption edge. The diffraction data were corrected for absorption by employing the semi-empirical multi-scan routine, which is commonly applied for redundant synchrotron data (Blessing, 1995). Since \( \mu \) depends on \( f'' \) according to Equation (4), no reasonable additional face-indexed absorption correction can be applied. The crystallographic software \texttt{OLEX2} was employed using \texttt{SHELX} for the initial structure solution and \texttt{olex2.refine} as the refinement engine (Dolomanov et al., 2009; Sheldrick, 2015; Bourhis et al., 2015). Nonspherical atomic form factors were calculated with \texttt{NoSpherA2} (Kleemiss et al., 2021), performing Hirshfeld-Atom-Refinement (HAR) employing a level of theory of DKH2-PBE0/x2c-TZVP within \texttt{ORCA} (Version 5.0; Capelli et al., 2014; Jayatilaka & Dittrich, 2008; Neese et al., 2020). HAR uses tailor-made form factors \( f_0 \) computed from electron densities after a single-point wavefunction calculation for the current model, partitioned by Hirshfeld stockholder partitioning (Hirshfeld, 1976, 1977). The obtained atomic form factors are subsequently used in the refinement of the crystallographic model parameters, repeating form factor calculation and refinement until all parameters reach a convergence threshold.

To compare the effects of different dispersion parameters on the structure refinement, models were created using the values for molybdenum from the tables of Henke et al. (1993) and Sasaki (1989) for the respective energies. The latter closely resemble the values computed according to Brennan & Cowan (1992) (see Fig. S13 in the supporting information).

The most commonly used source for dispersion correction parameters from the \textit{International Tables for Crystallography} (Vol. C) are given for only a few energies used on standard laboratory diffractometers (Creagh & McAuley, 1992). The refinement of the dispersion correction for molybdenum was performed by the \texttt{olex2.refine} engine, which introduces \( f' \) and \( f'' \) as independent scalar parameters in the matrix least-squares refinement procedure (Bourhis et al., 2015). The results of these calculations around the K edge of molybdenum and the obtained quality indicators are compared to those using the tabulated values in Fig. 2.

At 19 900 eV, Sasaki’s tabulated \( f' \) and \( f'' \) values agree well with both those obtained from the XAS data and the refined ones. The strongest deviation between the literature values and the refined dispersion parameters is observed at 20 001 eV. At this energy, the K edge is already exceeded in the tabulated values, while this is not yet the case for Mo(CO)_6 according to the X-ray absorption spectrum. Above the absorption edge, the refined values follow the observed spectral fine structure. These features are specific to the individual crystal structure and cannot be captured by precalculated dispersion corrections. However, the values for \( f' \) and \( f'' \) can reliably be determined from the diffraction data. It is remarkable that here the smallest refined value of \( f' \) is 3.75 electrons (e) above that of the tabulated values. The standard uncertainties of the dispersion values obtained by refinement is in the range 0.03–0.06 e (see Tables S1–S7 in the supporting information).
correlation between \( f' \) and \( f'' \) is low although they are related via Equation (5). This is due to the fact that the mutual dependence of \( f' \) and \( f'' \) includes the integral of a wide energy range, whereas the refinement is performed at a specific energy.

The large differences between \( f' \) and \( f'' \) are directly reflected in the structural models. The overall agreement factors vary between \( 1.30 < R_1 < 3.62\% \) and \( 3.62 < wR_2 < 11.86\% \) for the models with tabulated dispersion values. In contrast, the models obtained with refined dispersion corrections resulted in consistent agreement factors between \( 1.29 < R_1 < 1.51\% \) and \( 3.59 < wR_2 < 4.03\% \) (see Tables S1–S7 in the supporting information). Therefore, only the refinement of dispersion values leads to a robust model within the energy range investigated.

Furthermore, the deviations of the agreement factors are strongly correlated with the disagreement of the applied dispersion values to the X-ray absorption spectrum. This large deviation shows how drastically an incorrect dispersion correction can affect the crystallographic model if the incident energy is in the range of the absorption edge of an element involved.

The effect on the crystallographic models can also be observed in the atomic displacement parameters (ADPs). Fig. 3 shows the models and difference Fourier maps at 20 001 eV using Sasaki tabulated dispersion values for molybdenum and refined ones, respectively. The unreasonably small displacement parameters of the metal atom in Fig. 3(a) are caused by the inadequate dispersion treatment according to the Sasaki tables, since the effective scattering power is overestimated. As a result, the residual electron-density map shows excessive electron density around the molybdenum position and electron depletion around the carbonyl ligands. This showcases the impact of one incorrect atomic form factor on the entire structure model. The biggest effect, however, is observed in the vicinity of the metal centre.

A similar observation regarding the ADP size is known when an atom type is assigned to a position that contains more electrons than the modelled atom type provides. In the case of insufficient dispersion treatment, such a wrong assignment of an atom type leads to more reasonable anisotropic displacement parameters and refinement indicators. In contrast, the model with refined dispersion values in Fig. 3(b) shows none of these artefacts. The fractal dimension plots of the models (see Figs. S2 and S3 in the supporting information) according to Meindl & Henn (2008) also show a much better agreement. In addition, the precision of the C—O bond is best in all cases with the refined parameters (see Figs. S2 and S3). A more detailed discussion of the misassignment of atomic types in the Mo(CO)\(_6\) structure of this study and its implications for the atomic form factors is given in the supporting information.

3. Conclusion

We have shown that better treatment of the anomalous dispersion correction is an important improvement to the conventional crystal structure determination. The inclusion of dispersion parameters \( f' \) and \( f'' \) in the least-squares refinement is simple and reliable, with low correlations and errors. The refined absorptive term \( f'' \) follows the fine structure of the independently measured X-ray absorption of Mo(CO)\(_6\). The resulting crystallographic models are characterized by low \( R_1 \) and \( wR_2 \) values for the excitation energies chosen around the Mo K absorption edge. Conversely, the models using tabulated dispersion values are substantially worse. The atomic displacement parameters, as well as the residual electron densities, are strongly affected by the incorrect dispersion treatment. The resulting effective form factor is incorrect by up to 3.7 e at 20 001 eV for \( f' \) and 1.3 e for \( f'' \) relative to the literature values.

We assume that the described method of dispersion refinement also influences crystal structures measured with laboratory diffractometers, especially when the available radiation falls near an absorption edge of a heavy element present in the compound. For instance, X-ray diffraction experiments with Cu \( K\alpha \) radiation containing late first-row \( d \)-block metals or lanthanides will be affected. The latter elements even have three L absorption edges that occur in a broad energy range of about 2000 eV. In addition, laboratory diffractometers are operated with mixed \( K\alpha_1,2 \) radiation that differs by 20 eV for Cu, 105 eV for Mo and even 172 eV for Ag \( K\alpha \) radiation, respectively. These energy differences would require two sets of dispersion parameters, which are individual for a crystal structure, to perform a proper correction for this effect. Since the \( f'' \) parameter of the anomalous dispersion correction is directly related to the absorption coefficient \( \mu \), a resulting insufficient absorption correction can further worsen the results.

Therefore, a reconsideration of anomalous dispersion treatment will lead to a significant improvement of routine home source crystal structure determinations containing heavy elements.

4. Related literature

The following references are cited in the supporting information for this article: Dyadkin et al. (2016); Parsons et al. (2012); Ramseshan & Abrahams (1975).

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