

Bayes–Turchin approach to XAS analysis

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Modern analysis of X-ray absorption fine structure (XAFS) is usually based on a traditional least-squares fitting procedure. Here an alternative Bayes–Turchin method is discussed which has a number of advantages. In particular the method takes advantage of *a priori* estimates of the model parameters and their uncertainties and avoids the restriction on the size of the model parameter space or the necessity for Fourier filtering. Thus the method permits the analysis of the full X-ray absorption spectra (XAS), including both XAFS and X-ray absorption near-edge spectra (XANES). The approach leads to a set of linear equations for the model parameters, which are regularized using the ‘Turchin condition’. Also, the method naturally partitions parameter space into relevant and irrelevant subspaces which are spanned by the experimental data or the *a priori* information, respectively. Finally we discuss how the method can be applied to the analysis of XANES spectra based on fits of experimental data to full multiple-scattering calculations. An illustrative application yields reasonable results even for very short data ranges.

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1. Introduction

The goal of X-ray absorption spectra (XAS) analysis is to extract various experimental parameters, *e.g.* interatomic distances R , coordination numbers N , vibrational amplitudes σ^2 *etc.* This requires the solution to an inverse problem, *i.e.* the determination of the physical parameters, which we denote by a vector \vec{x} , that best fit a model XAS spectrum $\mu^0(\vec{x}, \omega)$ to a set of K measurements μ_k , $k = 1 \dots K$, *i.e.* by inverting the relation $\mu_k = \mu^0(\vec{x}, \omega_k)$. Most modern approaches for the analysis of X-ray absorption fine structure (XAFS) are based on standard least-squares fitting algorithms (Lytle *et al.*, 1988; Newville *et al.*, 1995), either in k - or R -space, where $k = (E - E_0)^{1/2}$ is the photoelectron wavenumber measured from threshold and R is its Fourier complement. In this approach one minimizes the normalized mean square error,

$$\chi^2 \equiv \sum_{k=1}^K \left[\frac{\mu_k - \mu^0(\vec{x}, \omega_k)}{\Delta\mu_k} \right]^2, \quad (1)$$

with respect to each model parameter x_i , where $\Delta\mu_k$ denotes the experimental error at each data point k . A number of software packages for this purpose are available, *e.g.* *FEFFIT* (Newville, 2001), *EXAFSPAK* (George & Pickering, 2000), *EXCURVE* (Binsted, Campbell *et al.*, 1991; Binsted, Strange *et al.*, 1991) and *GNXAS* (Filipponi & Di Cicco, 1995). The least-squares approach has also been applied recently to the analysis of X-ray absorption near-edge structure (XANES) using the *MSXAN* package (Benfatto *et al.*, 2001).

There are a number of problems with the least-squares approach, however. First it is rarely obvious *a priori* which of the model parameters x_i or how many of them can be included in the fits. Indeed, there are usually many more model parameters than can be represented by the data, which leads to an ill-conditioned inverse problem, as discussed below. The problem is typically avoided in practice by Fourier filtering and a judicious selection of model parameters. For example, the Nyquist criterion $N_p = (2/\pi)\Delta k\Delta R$ gives an estimate of the number of independent variables that span a given range in k - and R -space. This procedure, however, is subject to user experience and subjectivity and hence difficult to automate. Also the effects of systematic errors both in theory and experiment are not easily taken into account. Moreover, many parameters are strongly correlated, leading to large error bars in the results without the imposition of appropriate constraints.

To address these problems we discuss an alternative method which permits the analysis of the full XAS spectrum, including the XANES and/or the XAFS. The method is based on the Bayes–Turchin approach introduced into XAFS analysis by Krappe & Rossner (2000, 2002). The approach has several advantages over traditional data-analysis procedures. In particular, the method avoids the restriction on the size of the model parameter space by making use of Bayes’ theorem. Given suitable restraints based on *a priori* parameters and their errors, the Bayes–Turchin method leads to a stable set of linear equations for the model parameters. Moreover, the

approach naturally partitions parameter space into relevant and irrelevant subspaces, which are determined primarily by the experimental data or the *a priori* information, respectively. Thus only relevant parameters in a given model parameter space are affected by a fit, while irrelevant parameters are essentially pinned at their *a priori* values.

A principle aim of the paper is to show that this approach is also advantageous for the analysis of XANES spectra. This requires a treatment of the smooth atomic background absorption as well as the fine structure and fast calculations of full-multiple-scattering theory as in the FEFF8 code (Ankudinov *et al.*, 1998) which are now possible using parallel algorithms (Ankudinov *et al.*, 2002). Such an approach is important, *e.g.* in the analysis of biostructures such as metalloproteins. For such systems the XANES often contains the best signal-to-noise ratio and is easiest to collect experimentally. However, the region is typically eliminated by the Fourier filtering in conventional analyzes.

The remainder of this paper is as follows. In §2 we discuss the problems with the least-squares method, and in §3 we outline the Bayes–Turchin approach. The method is illustrated with an application to the XANES of GeCl₄ in §4. Finally, §5 contains some conclusions.

2. Least-squares analysis

We begin by showing formally that the usual least-squares approach leads to an ill-conditioned inverse problem (Krappe & Rossner, 2002). Assuming that the theoretical model for the XAS $\mu^0(\vec{x}, \omega)$ is a smooth function of the parameters $\vec{x} = (x_1, x_2, \dots, x_N)$ near their true values \vec{x}_0 , one can approximate the model near any data point ω_k as

$$\mu_k^0(\vec{x}) = \mu_k^0(\vec{x}_0) + \sum_{i=1}^N \frac{\partial \mu_k^0}{\partial x_i} x_i, \quad (2)$$

where x_i denotes a normalized residual, *e.g.* $x_R \equiv (R - R_0)/\Delta R$ and ΔR is a nominal scaling factor chosen so that x_R is of order unity in fits. Inserting this approximation into the definition of χ^2 , we obtain

$$\chi^2 = \left| \vec{\Delta} - \sum_i^N \vec{G}_i x_i \right|^2, \quad (3)$$

where the vector $\vec{\Delta}$ denotes the signal components,

$$\Delta_k = \frac{\mu_k - \mu_k^0}{\Delta \mu_k} \equiv \bar{\mu}_k - \bar{\mu}_k^0. \quad (4)$$

Here $\bar{\mu} = \mu/\Delta\mu$ represents the dimensionless XAS, and $G_{ki} \equiv \vec{G}_i$ are the components of the normalized model gradients \vec{G}_i , *i.e.*

$$G_{ki} = \frac{1}{\Delta \mu_k} \frac{\partial \mu_k^0}{\partial x_i}. \quad (5)$$

With these definitions, minimizing χ^2 with respect to the parameters x_i leads to a set of N linear equations,

$$\sum_{j=1}^N Q_{ij} x_j = b_i, \quad i = 1 \dots N, \quad (6)$$

where Q_{ij} are components of the $N \times N$ information matrix \mathbf{Q} ,

$$Q_{ij} = \vec{G}_i \cdot \vec{G}_j = G_{ik}^T G_{kj}, \quad (7)$$

and b_i are the normalized signal components,

$$b_i = \vec{\Delta} \cdot \vec{G}_i. \quad (8)$$

Since the dot product is invariant under coordinate rotations, the information matrix does not depend on the space (*e.g.* k or R) in which it is evaluated.

To show that these equations are ill-conditioned, consider the eigenvalues q_α of the symmetric matrix \mathbf{Q} , listed in order of decreasing magnitude. Since \mathbf{Q} is directly related to the model gradients, it is clear that parameters on which the model depends strongly correspond to large eigenvalues and, conversely, those which have little effect on the model correspond to small eigenvalues. The conditioning number Z_Q is defined as the ratio q_1/q_N , which can grow large as the number of parameters is increased. For the applications to XAFS discussed by Krappe & Rossner (2002), Z_Q can be up to 10^{16} . Thus in the eigen space spanned by the eigenvectors of the matrix \mathbf{Q} , the solutions to the linear equations are formally given by

$$x_\alpha = b_\alpha / q_\alpha. \quad (9)$$

However, the signal components b_α are generally limited by the experimental noise and only include one factor of the gradients, while the eigenvalues q_α involve a product of gradients and can be much smaller. Thus, as a result of noise, the weak eigen-components are intrinsically unstable. Thus the standard least-squares approach becomes unstable as the number of parameters is increased, necessitating some form of regularization. Various methods of regularization exist, *e.g.* Wiener filtering (Bijaoui, 2002); however, many of these are more or less *ad hoc*. In the next section we show how the Bayesian approach naturally regularizes the inversion.

3. Bayes–Turchin approach

The Bayes–Turchin approach of Krappe & Rossner (2002) addresses the ill-posed nature of the inverse problem by making use of known or *a priori* information on the model parameters \vec{x} . Here we will only summarize the key results, as the original papers provide much more detail. For simplicity we will assume that the parameters x_i have Gaussian distributions, which may be correlated, *i.e.* they are characterized by the *a priori* probability distribution,

$$P_{\text{prior}}(\vec{x}) = N \exp[-(1/2)\chi_{\text{prior}}^2], \quad (10)$$

where N is a normalization factor and χ_{prior}^2 is a quadratic form,

$$\chi_{\text{prior}}^2 = \sum_{ij} A_{ij} x_i x_j, \quad (11)$$

with a kernel \mathbf{A} determined by the inverse of the cross-correlation matrix

$$A_{ij}^{-1} = \langle x_i x_j \rangle. \quad (12)$$

Thus, for example, the *a priori* variance $\sigma_{\text{prior}}^2(R)$ of the coordinate R yields $A_{R,R}^{-1} = \sigma_{\text{prior}}^2(R)/\Delta_R^2$.

Secondly, as discussed by Krappe & Rossner (2000), the method explicitly takes into account theoretical errors by considering the dependence of the model $\mu^0[\vec{x}, \vec{y}(\vec{x}), \omega]$ on both the model parameters \vec{x} and on the theoretical parameters, *e.g.* the phase shifts, scattering amplitudes, mean free paths *etc.*; these are denoted by a dimensionless vector $\vec{y} = y_1, y_2, \dots$, the components of which may also depend on the model parameters \vec{x} .

The incorporation of the *a priori* information is carried out using Bayes theorem for the conditional probability distribution, given the *a priori* information. This yields the *a posteriori* probability distribution for the model parameters x given $\vec{\mu}$,

$$P_{\text{post}}(x|\vec{\mu}) = \frac{P_{\text{prior}}(x) P_{\text{cond}}(\vec{\mu}|x)}{\int P_{\text{prior}}(x) P_{\text{cond}}(\vec{\mu}|x) d^N x}. \quad (13)$$

Here the conditional probability is

$$P_{\text{cond}} = \exp[-(1/2)\chi_{\text{cond}}^2], \quad (14)$$

where in tensor notation (*i.e.* writing $G_{ij} \equiv \mathbf{G}$),

$$\chi_{\text{cond}}^2 = \mathbf{x}^T \mathbf{Q} \mathbf{x} - 2\mathbf{b}^T \mathbf{x} + (\vec{\mu} - \vec{\mu}^0)^T \mathbf{C} (\vec{\mu} - \vec{\mu}^0). \quad (15)$$

Here the $K \times K$ matrix \mathbf{C} appears,

$$\mathbf{C} \simeq [1 + \mathbf{B}^{-1} + \mathbf{T} \mathbf{D}^{-1} \mathbf{T}^T]^{-1}, \quad (16)$$

which characterizes the errors in the theory. In particular, \mathbf{B} characterizes the truncation error arising, for example, from a finite cluster size of the model, while \mathbf{D} involves errors in the theoretical parameters (*e.g.* the phase shifts) and \mathbf{T} their gradients. If the off-diagonal terms are neglected, the effect of \mathbf{C} is to renormalize the mean-square error Δ_k at each point k by the sum of the squares of the experimental, model and truncation errors.

As a result, one obtains an *a posteriori* distribution given by

$$P_{\text{post}}(\vec{x}) = \exp[-(1/2)\chi_{\text{post}}^2], \quad (17)$$

where $\chi_{\text{post}}^2 = \chi_{\text{prior}}^2 + \chi_{\text{cond}}^2$ is given by the quadratic form,

$$\chi_{\text{post}}^2 = \mathbf{x} \mathbf{A} \mathbf{x} + (\Delta - \mathbf{G} \mathbf{x})^T \mathbf{C} (\Delta - \mathbf{G} \mathbf{x}). \quad (18)$$

The *a posteriori* expectation values of the model parameters,

$$\langle \vec{x} \rangle = \int \vec{x} P_{\text{post}}(\vec{x}) d^N x, \quad (19)$$

are obtained by minimizing χ_{post}^2 with respect to x_i . This yields the regularized linear equations

$$\sum_{j=1}^N [\bar{Q}_{ij} + A_{ij}] x_j = \bar{b}_i, \quad i = 1 \dots N, \quad (20)$$

where the renormalized information matrix is

$$\mathbf{Q} = \mathbf{G}^T \mathbf{C} \mathbf{G}, \quad (21)$$

and the renormalized signal coefficients are given by

$$\bar{b}_i = \Delta \mathbf{C} \vec{G}_i. \quad (22)$$

Clearly the *a priori* information in the matrix \mathbf{A} regularizes the inversion, since in the eigen-space of $\mathbf{Q} + \mathbf{A}$ the solutions are then always stable. That is, the formal solutions for the model parameters are

$$x_\alpha = \frac{b_\alpha}{q_\alpha + a_\alpha}, \quad (23)$$

which is well behaved even when $q_\alpha \rightarrow 0$.

Moreover, it is seen that the *a priori* information naturally partitions the data into relevant and irrelevant subspaces, \mathbf{R} and \mathbf{P} , respectively, where

$$x_\alpha = b_\alpha / q_\alpha, \quad \alpha \in \mathbf{R}, \quad (24)$$

$$x_\alpha = b_\alpha / a_\alpha, \quad \alpha \in \mathbf{P}. \quad (25)$$

Thus only the parameters in the relevant subspace are significantly fit by the data, while the parameters in the irrelevant subspace do not deviate significantly from their *a priori* values. An important finding is that the dimension N_R of the relevant subspace \mathbf{R} is significantly smaller than that given by the Nyquist criterion, *i.e.* $N_R < N_p = (2/\pi)\Delta k \Delta R$, since N_R takes into account the effects of experimental noise and systematic error.

These conditions can be satisfied by setting $A_{nn'} = \alpha \delta_{nn'}$ for an appropriate cut-off α (Krappe & Rossner, 2000). In particular, two methods for determining α were introduced by Turchin *et al.* (1971). The first fixes α^* such that the effective number of degrees of freedom is

$$K_{\text{eff}} = K - \text{Tr} \mathbf{Q}(\mathbf{Q} + \mathbf{A})^{-1}. \quad (26)$$

This criterion ensures that the information in the data is not distorted by *a priori* information. A second criterion, for situations when one would like to decrease the effect of the *a priori* information, gives a value $\alpha'^* < \alpha^*$.

4. XAS analysis

4.1. XAFS analysis

In the original papers of Krappe & Rossner (2000, 2002), the Bayes method was applied to the analysis of XAFS data of Cu metal based on a theoretical model using the multiple-scattering path expansion and the FEFF8 code. In an initial test, a large model parameter space of dimension $N = 158$ was used, including the variables, $S_0^2, E_0, R_1 \dots R_{78}, \sigma_1^2 \dots \sigma_{78}^2$. Of these, only a small number turn out to be important. The dimension of the relevant subspace for this model was found to be $N_R = 21$ from the Turchin cut-off α^* and $N_R = 40$ from the cut-off α'^* , both much smaller than the Nyquist estimate, $N_p = 55$. Subsequent investigations for a variety of systems showed that the approach is very robust. In these investigations, the vibrational information was fit to a small set of spring constants κ_s using an efficient theoretical model (Poiarkova & Rehr, 1999), which is much more efficient than fitting pairwise Debye–Waller factors. The third cumulants were also included as model parameters.

4.2. XANES analysis

Here we have attempted to extend the Bayes–Turchin method for XANES analysis. The near edge requires a different fitting procedure for several reasons. First, the measured absorption contains a background contribution μ_{back} (from the other edges) as well as the atomic background μ_0 from a given edge, in addition to the fine structure χ , all of which must be taken into account,

$$\mu = \mu_{\text{back}} + \mu_0(1 + \chi). \quad (27)$$

Second, the path expansion for the fine structure does not always converge well in the XANES region, thus often necessitating full multiple-scattering calculations, which do not have a simple analytical form. Thus numerical gradients of the model parameters are needed,

$$G_{ki} = \frac{\mu(\vec{x} + \delta x_i \hat{x}_i) - \mu(\vec{x} - \delta x_i \hat{x}_i)}{2\delta x_i}, \quad (28)$$

where \hat{x}_i is the i th unit vector. To fit the atomic background, we have adopted a new Bayesian background-subtraction procedure introduced by Krappé & Rossner (2004). In this method, a correction $\delta\mu_0(\omega)$ is added to an *a priori* background absorption $\mu_0(\omega)$ from FEFF, *i.e.* with a small set of spline parameters $\delta\mu_1, \delta\mu_2, \dots, \delta\mu_T$, where in our work T is typically 3–5. Related Bayesian background-subtraction algorithms have also been developed by Klementev (2001).

4.3. Results

The above formalism was tested using high-quality gas-phase GeCl_4 data (Bouldin, 1990) using the FEFF8 code. The model parameters included E_0 , R and between three and five spline parameters $\delta\mu_i$. Thus this example with its rather restricted parameter set should be viewed as a preliminary step towards more extensive future analysis. The pre-edge background μ_{back} was isolated using the *ATHENA* package (Ravel & Newville, 2005) and subtracted from the data prior to fitting. The *a priori* error values of these few parameters were found to have little effect on the fit, that is, all of these parameters were found to be relevant. However, such *a priori* estimates are generally expected to become important when additional parameters, such as distortions, spring constants and the mean free path, are added.

Our results are illustrated in Fig. 1, which shows the result of a fit in k -space of the XANES of GeCl_4 from threshold to $k = 3.89 \text{ \AA}^{-1}$, *i.e.* 11013–11170 eV. Clearly the fit is qualitatively satisfactory, and leads to a reduced χ^2 of $\chi_{\text{red}}^2 \simeq 19$, where $\chi_{\text{red}}^2 = \chi^2/(K - 7)$, where $K - 7$ is the number of data points minus the number of model parameters, or the number of degrees of freedom. The experimental error $\Delta\mu_k$ at all the data points was taken to be 0.01, based on the observed fluctuations in the tail of the experimental spectrum. Theoretical errors were not included, but are typically larger than $\Delta\mu_k$, as discussed by Krappé & Rossner (2002). As seen in Fig. 1, the theoretical error is quite large near 11120 eV, probably due to the neglect of non-spherical corrections and many-electron excitations in the theory. This systematic error

Table 1

Key parameters and their *a priori* and final fitted values and uncertainties.

	<i>A priori</i> value	Fit value
R	2.00 ± 0.02	2.10 ± 0.01
E_0	-4 ± 1	-4.7 ± 0.2

Table 2

Variation of the fitted near-neighbor distance R and error estimate δR versus data range k_{max} .

The initial guess in all cases was set to $R = 2.0 \pm 0.2 \text{ \AA}$.

E_{max} (eV)	k_{max} (\AA^{-1})	R (\AA)	δR
11120	1.14	2.11	± 0.07
11130	1.98	1.92	± 0.03
11140	2.57	2.22	± 0.06
11150	3.06	2.12	± 0.04
11160	3.46	2.16	± 0.02
11170	3.81	2.10	± 0.01
11180	4.14	2.14	± 0.02
11190	4.46	2.13	± 0.01
11200	4.70	2.103	± 0.008

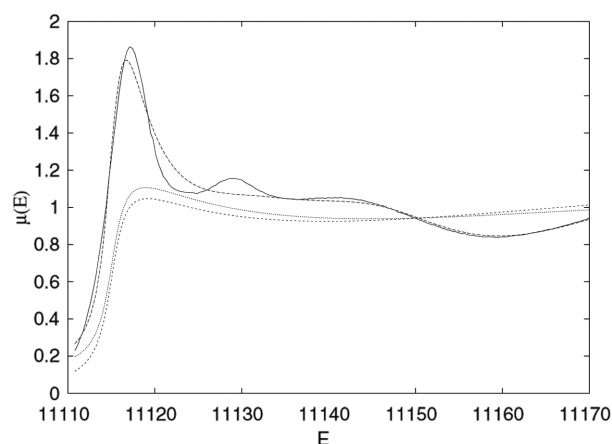


Figure 1

Experimental XAS (solid) and fit (dashes) for GeCl_4 for $k_{\text{max}} = 3.89 \text{ \AA}^{-1}$. The experimental near-neighbor distance is 2.113 \AA . Also shown is the *a priori* atomic background μ_0 from FEFF8 (short dashes) and the corrected μ_0 from a three-point spline fit (dots). The model parameters included in the fit were E_0 , R , an absolute energy shift, an experimental broadening term and three spline parameters.

is likely to be the primary source of error in the fit. The *a priori* and fitted values of the parameters R and the threshold level E_0 are shown in Table 1.

An important question to be considered by these preliminary results is how the fitted parameters vary as the data range is reduced. Remarkably this is found to be reasonably stable, even for very small data ranges. Results for the near-neighbor distance R with respect to the data range $[0, k_{\text{max}}]$ for various values of the maximum wavevector k_{max} are given in Table 2. Note that the results for the fitted distances R tend toward the experimental value 2.113 \AA as the k -range is increased while the errors tend to decrease. The errors in the fit parameters are still not well determined, owing to the lack of a treatment of the theoretical errors, which appear to dominate the fit. The r.m.s. errors in R , obtained from

$(Q_{R,R}^{-1})^{1/2}$, are typically less than about 0.03 Å. However, since our fits did not include the theoretical error, and $\chi_{\text{red}}^2 \simeq 20$, these errors are underestimates.

5. Conclusions

We have found that the Bayes–Turchin approach can be applied to the analysis of XANES data, even over short experimental data ranges. The spline correction to the atomic background μ_0 is found to be stable and thus provides a useful procedure to improve background subtraction in XAS.

Such developments are important, *e.g.* in the analysis of biostructures such as metalloproteins. For such systems the Bayes–Turchin approach has many advantages compared with conventional least-squares fitting methods. In particular, the method avoids the restriction on the size of the model parameter space. Moreover, the method can take advantage of *a priori* estimates of model parameters, as well as their uncertainties and correlations, thus improving the significance of fits. Thus the method can perhaps be automated and has the potential to provide a smart black-box XAS analysis tool. Finally, the method is quite general and can be applied as an add-on to existing XAS analysis techniques by modifying the χ^2 function which is minimized, thus providing a procedure for adding fuzzy constraints on model parameters.

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