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Estimating Contrast Transfer Function and Associated Parameters by Constrained Nonlinear Optimization

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Abstract

The three-dimensional reconstruction of macromolecules from two-dimensional single-particle electron images requires determination and correction of the contrast transfer function (CTF) and envelope function. A computational algorithm based on constrained nonlinear optimization is developed to estimate the essential parameters in CTF and envelope function model simultaneously and automatically. The application of this estimation method is demonstrated with focal series images of amorphous carbon film as well as images of ice-embedded icosahedral virus particles suspended across holes.

1. Introduction

The 2-D images collected from an electron microscope are not perfect 2-D projections of the 3-D structure. Each experimentally collected image can be treated as a modulated projection with noises. The modulation of the image is determined by a number of factors that are related to the electron microscope settings and imaging conditions. The modulation process has been modeled mathematically as the *contrast transfer function* (Thon, 1971, Erickson & Klug, 1970) and *envelope functions* (Hanszen, 1967). Each of these functions contains a number of parameters affecting the image contrast and quality. In this work, we assume that CTF modulation is considered invariant in the entire micrograph, which is valid in many of the single particle cryo-EM studies.

The CTF parameter estimation problem is essentially a non-linear curve-fitting problem. A number of schemes have been proposed to solve this problem for single particle imaging (Zhu *et al.*, 1997, Conway & Steven, 1999, Ludtke *et al.*, 1999, Huang *et al.*, 2003, Velazquez-Muriel *et al.*, 2003, Sander *et al.*, 2003, Mallick *et al.*, 2005). However, most of these schemes involve some *ad hoc* or manual fitting steps instead of making use of the state of the art numerical optimization algorithms that can be done objectively and accurately. As a result, parameter determination becomes difficult

especially when the experimental images are collected near focus where only one or two CTF rings are apparent.

This paper describes the use of efficient and accurate numerical optimization techniques to estimate these parameters by treating the estimation problem as a constrained nonlinear optimization problem. Such an approach was perceived as infeasible or too computationally demanding in the past. Our experimental results demonstrate that this can be done reliably, efficiently and automatically.

2. Problem Formulation

Thin biological specimen, consisting of mostly low atomic elements (C, N and O) can be approximated as weak phase objects for transmission electron microscopy. For the weak phase objects, the mathematical model that describes the relationship between the object potential function and the observed image has been well established (Thon, 1971, Erickson & Klug, 1970, Hanszen, 1971b). In order to demonstrate our approach, we will describe both the well-known as well as the derived formulations.

2.1 Mathematical Model for Image Contrast

In the image contrast theory, the 2-D Fourier transform of an image, which we denote by $\mathbf{I}(\mathbf{s})$, can be related linearly to the structure factor of the specimen, $\mathbf{F}(\mathbf{s})$, through the expression

$$\mathbf{I}(\mathbf{s}) = \mathbf{F}(\mathbf{s})\mathbf{H}(\mathbf{s}) + \mathbf{N}(\mathbf{s}), \quad (1)$$

where $\mathbf{H}(\mathbf{s})$ is the modulation function characterizing the instrument and experimental conditions, and $\mathbf{N}(\mathbf{s})$ is the noise function originating from various sources including surrounding buffer, electron inelastic scattering and recording media. Here, the bold faced \mathbf{s} denotes a 2-D frequency vector. This is to be distinguished from the non-bold faced s which denotes a 1-D spatial frequency scalar.

Note that $\mathbf{I}(\mathbf{s})$, $\mathbf{F}(\mathbf{s})$ and $\mathbf{N}(\mathbf{s})$ are all complex valued functions. In this paper, we assume that the microscope optics is well aligned during image acquisition so that $\mathbf{H}(\mathbf{s})$ is a real valued function. The computational problem to be solved is to construct $\mathbf{H}(\mathbf{s})$ and $\mathbf{N}(\mathbf{s})$ given $\mathbf{I}(\mathbf{s})$ and $\mathbf{F}(\mathbf{s})$. Analytical expressions for $\mathbf{H}(\mathbf{s})$ exist (Thon, 1971, Erickson & Klug, 1970, Hanszen, 1971b). These expressions contain a number of unknown parameters that can be determined through a numerical fitting procedure (Saad *et al.*, 2001). If one assumes that the projection image is not correlated with the background noise, then it follows from (1) that

$$I_2(\mathbf{s}) = F_2(\mathbf{s})H_2(\mathbf{s}) + N_2(\mathbf{s}), \quad (2)$$

where $I_2(\mathbf{s})$, $F_2(\mathbf{s})$ and $N_2(\mathbf{s})$ are the power spectra of the observed projection image, the structure factor and the background noise, respectively. Here, the power spectrum of an image is defined as the expectation value of the Fourier intensity of the image. The subscript 2 in (2) is used to indicate that functions $I_2(\mathbf{s})$, $F_2(\mathbf{s})$ and $N_2(\mathbf{s})$ describe mappings from the 2-D frequency space to the set of real numbers. These functions are to be distinguished from the functions $I(s)$, $F(s)$ and $N(s)$, which are defined in subsequent sections to describe mappings from 1-D frequency to real numbers.

Equation (2) can be written in polar coordinates as

$$I_2(s, \theta) = F_2(s)H_2(s, \theta) + N_2(s, \theta). \quad (3)$$

Note that $F_2(s)$, which corresponds to the rotationally averaged value of the structure factor associated with the specimen, is a 1-D rotationally invariant function. Such rotationally averaged 1-D structure factor can be measured in an X-ray scattering experiment of a solution suspension of the specimen (Thuman-Commike *et al.*, 1999, Schmid *et al.*, 1999, Saad *et al.*, 2001). Alternatively, structure factor can also be estimated at low resolution directly from particle images and at high resolution from a model (Ludtke *et al.*, 1999). When $F_2(s)$ is known, the parameter estimation problem becomes well defined.

The analytical function used to describe the background noise term $N_2(s, \theta)$ in (3) is somewhat arbitrary and less well defined in the image contrast theory. In this paper, we extend the model defined previously (Saad et al., 2001) by including the azimuthal dependence of the functions i.e., we set

$$N_2(s, \theta) = n_3(\theta) e^{-n_4(\theta)s^2 - n_2(\theta)s - n_1(\theta)\sqrt{s}}, \quad (4)$$

where $n_i(\theta)$ ($i = 1, 2, 3, 4$) are unknown parameters to be determined.

The modulation function $H_2(s, \theta)$ can be defined (Thon, 1971, Erickson & Klug, 1970, Hanszen, 1971b) as

$$H_2(s, \theta) = \alpha^2 CTF^2(s, \theta) \cdot E^2(s, \theta). \quad (5)$$

where

$$CTF(s, \theta) = -\left(\sqrt{1-Q^2} \sin \gamma(s, \theta) + Q \cos \gamma(s, \theta)\right), \quad (6)$$

$$\gamma(s, \theta) = 2\pi \left(-\frac{C_s \lambda^3 s^4}{4} + \frac{\Delta z(\theta) \lambda s^2}{2} \right), \quad (7)$$

The $\sin[\gamma(s, \theta)]$ and $\cos[\gamma(s, \theta)]$ terms in the CTF function (6) are known as the *phase* and *amplitude contrast transfer functions*, respectively (Erickson & Klug, 1970). The wavelength (λ) and the spherical aberration (C_s) are known constants. The unknown parameters to be estimated in (5) are the defocus (Δz), the amplitude contrast ratio (Q), the amplitude coefficient (α), and the envelope function ($E(s, \theta)$). We should point out that Q is, in principle, dependent on the spatial frequency and the atomic composition of the specimen. However, for weak phase objects, the variation of Q with respect to these factors is so small that it can be considered as a constant parameter. The defocus $\Delta z(\theta)$ is anisotropic in general. It can be represented by

$$\Delta z(\theta) = \Delta z_0 + \Delta z_1 \sin(2(\theta - \theta_0)), \quad (8)$$

where Δz_0 is the mean defocus of the sample, Δz_1 is the focal difference due to axial astigmatism and θ_0 represents the reference angle of axial astigmatism (Thon, 1971). When astigmatism is present, the power spectrum often exhibits elliptically shaped CTF rings.

The envelope function $E(s, \theta)$ in (5) is used to account for the spatial and temporal coherence effects, specimen drift and other signal decay factors such as the modulation function of the recording medium in $H_2(s, \theta)$. Analytical expressions for some of these factors in the envelope function have been described previously (Frank, 1969, Frank, 1976, Hanszen, 1967, Hanszen, 1971a). In practice, it has been empirically observed that the envelope function for images of ice-embedded particles with subnanometer resolution ($<6 \text{ \AA}$) data from most of the electron microscopes can be approximated by a single Gaussian function of the form

$$E(s, \theta) = e^{-B(s, \theta)s^2}, \quad (9)$$

where the non-negative parameter $B(s, \theta)$ has been called the experimental B -factor (Saad et al., 2001). The techniques we employ to solve the parameter estimation problem allows alternative formulations of the envelope function. In particular, we have experimented with using a more general envelope function of the form

$$E(s, \theta) = e^{-B_1(s, \theta)s - B_2(s, \theta)s^2 - B_3(s, \theta)s^3}$$

to model decay of the power spectrum from low to high frequencies for some data sets.

2.2 Parameter Estimation via Constrained Nonlinear Optimization

We measure the discrepancy between the analytical model (3) and the experimentally measured power spectrum $\hat{I}_2(s, \theta)$ by the residual function

$r_2(s, \theta; x) = I_2(s, \theta; x) - \hat{I}_2(s, \theta; x)$ where $x = (\alpha, \Delta z_0, \Delta z_1, \theta_0, B, Q, n_1, n_2, n_3, n_4)$. To determine the optimal value of x , we propose to minimize the nonlinear objective function

$$\rho_2(x) = \|r_2(s, \theta; x)\|^2, \quad (10)$$

where $\|r_2(s, \theta; x)\|$ is defined as the standard 2-norm of $r_2(s, \theta; x)$, i.e.,

$$\|r_2(s, \theta; x)\| = \sqrt{\int_{s_{\min}}^{s_{\max}} \int_0^{2\pi} r_2^2(s, \theta; x) d\theta ds}, \quad (11)$$

for some low and high cutoff frequencies s_{\min} and s_{\max} .

When $CTF(s, \theta)$, $E(s, \theta)$ and $N_2(s, \theta)$ are independent of θ (i.e. astigmatism and drift are ignored, which is the case for good-quality experimental images), we can simplify the notation to obtain

$$I(s) = \alpha^2 F(s) E^2(s) CTF^2(s) + N(s). \quad (12)$$

When the structure factor $F(s)$ is available, one can determine the parameters $x = (\alpha, \Delta z_0, B, Q, n_1, n_2, n_3, n_4)$ by minimizing the function

$$\rho(x) = \|r(s; x)\|^2, \quad (13)$$

where $r(s; x)$ is the one-dimensional (1-D) residual function that measures the discrepancy between the 1-D analytical model (12) and the rotationally averaged power spectrum $\hat{I}(s)$. The norm of $r(s; x)$ is defined as

$$\|r(s; x)\| = \sqrt{\int_{s_{\min}}^{s_{\max}} r^2(s; x) ds} \approx \sqrt{\sum_{j=j_{\min}}^{j=j_{\max}} r^2(s_j; x)}. \quad (14)$$

Note that the objective function (14) is evaluated on the interval $[s_{\min}, s_{\max}]$. The reason for imposing such a restriction is to eliminate the unreliable and noisy data at both low and high frequencies.

In general, the objective function defined in (14) has many local minima. To narrow the search range and avoid being trapped at an undesirable local minimum, we impose explicit constraints. In most cases, the defocus value of Δz can be estimated from the experimentally intended imaging conditions to be within $[\Delta z_{\min}, \Delta z_{\max}]$. The valid values for Q are between 0 and 1 (Erickson & Klug, 1970). However, in practice, the upper bound for Q is generally believed to be much smaller than one (Toyoshima *et al.*, 1993). Since the experimental B-factor is always positive as defined in (Saad *et al.*, 2001), the inequality of the type $0 \leq B \leq B_{\max}$, for some constant B_{\max} , is a natural constraint. Similarly, to ensure that the intensity of the background noise never falls below zero, we impose $n_3 \geq 0$.

In addition to these bound constraints, we also impose a set of nonlinear inequality constraints in the form of $N(s_j) \leq \hat{I}(s_j)$, for $j = j_{\min}, \dots, j_{\max}$. These constraints are developed to ensure that the noise background term $N(s)$ is always less than $\hat{I}(s)$.

Because the intensity of the background signal typically decreases from low to high spatial frequencies, it is desirable to include constraints of the following type:

$$\frac{\partial N(s_j)}{\partial s} \leq 0 \text{ for } j = j_{\min}, \dots, j_{\max}. \quad (15)$$

In summary, when $CTF(s, \theta)$, $E(s, \theta)$ and $N_2(s, \theta)$ are independent of θ , as found in many experimental cases (e.g., (Jiang *et al.*, 2003, Ludtke *et al.*, 2008, Jiang *et al.*, 2008, Liu *et al.*, 2007, Ludtke *et al.*, 2004)), we can estimate the unknown parameters for the CTF function that characterize the image modulation process by solving the following constrained nonlinear optimization problem.

$$\min_x \rho(x) \quad (16)$$

subject to

$$\Delta z_{\min} \leq \Delta z \leq \Delta z_{\max} \quad (17)$$

$$0 \leq B \leq B_{\max} \quad (18)$$

$$Q_{\min} \leq Q \leq Q_{\max} \quad (19)$$

$$0 \leq n_3 \quad (20)$$

$$N(s_j) \leq \hat{I}(s_j) \text{ for } j = j_{\min}, \dots, j_{\max} \quad (21)$$

$$\frac{\partial N(s_j)}{\partial s} \leq 0 \text{ for } j = j_{\min}, \dots, j_{\max}, \quad (22)$$

where $\rho(x)$ is defined in (13).

In practice, the magnitude of $I(s)$ may vary by several orders of magnitudes between the low and high frequencies as seen in the X-ray solution scattering of the single particle suspension (Thuman-Commike et al., 1999, Ludtke *et al.*, 2001). In this case, applying a non-linear optimization solver to (16)-(22) directly may result in an approximate solution that produces more accurate low frequency fit at the expense of severe misfit at the intermediate and high frequency ranges of the power spectrum. Because the defocus parameter Δz , the most important parameter in the CTF model, is largely determined by the intermediate to high frequency part of the power spectrum, such a misfit is likely to be detrimental in subsequent computations.

To overcome this problem, one may introduce a set of weights ω_j in (13) that vary with respect to the frequency s_j . That is, one can define the objective function in (13) to be

$$\rho(x) = \sum_{s_{\min} \leq s_j \leq s_{\max}} [\hat{I}(s_j) - I(s_j)]^2 \omega_j. \quad (23)$$

However, choosing a set of appropriate weights is not a trivial task.

An alternative strategy for mitigating problems associated with the large magnitude variation in $\hat{I}(s)$ is to estimate the desired parameters by fitting $\log(\hat{I}(s))$ instead. Since the log function is monotonically increasing on $(0, \infty)$, minimizing (13) is equivalent to minimizing

$$\eta(x) = \left\| \log(\hat{I}(s)) - \log(I(s)) \right\|^2. \quad (24)$$

Note that $I(s)$ also depends on the parameter x to be estimated. In this formulation, we may need to impose additional constraints

$$I(s_j) > 0, \text{ for } s_{\min} \leq s_j \leq s_{\max}, \quad (25)$$

to ensure that the second log term in (24) is well defined.

The use of the objective function (13) is not appropriate when astigmatism and drift are present in the micrograph. When $CTF(s, \theta)$, the envelope function $E(s, \theta)$ and the background noise $N_2(s, \theta)$ all vary with respect to θ , one must resort to the most general form of the objective function defined in (10). Because parameters α , B , Q , and n_i ($i=1,2,3,4$) are all assumed to have angular dependency in this case and the defocus is now parameterized by three parameters Δz_0 , Δz_1 and θ_0 that appear in (8), the number of unknown parameters to be estimated becomes $7m_\theta + 3$, where m_θ is the number of angular samples used in the evaluation of (10). The angular dependency of the parameters to be estimated also introduces angular dependency in the constraints defined by (18) – (22). As a result, the total number of nonlinear constraints (eqns. 21-22) in the constrained nonlinear optimization model will increase by a factor of m_θ . The increased number of unknowns and constraints makes the optimization problem much more

difficult to solve. Hence, we need to seek other alternatives that are computationally more efficient.

When the angular dependency of (3) is caused solely by astigmatism, i.e., the defocus $\Delta z(\theta)$ is the only parameter that varies with respect to θ , integrating the right hand side of (5) with respect to θ yields a closed form expression which we will not show here. Such an expression allows us to again reduce the 2-D fitting problem to a 1-D fitting problem. Unfortunately, the residual norm (13) associated with this 1-D fitting problem has far too many local minima within the domain defined by the constraints (18)-(22). Hence, it is difficult to compute the optimal estimation of the desired parameters in practice.

When $CTF(s, \theta)$, $E(s, \theta)$ and $N_2(s, \theta)$ vary slowly with respect to θ , which is the case for good quality images, a simple and practical strategy that one can use to reduce the complexity of the computation is to divide the 2-D power spectrum evenly into k angular sectors for some k that is relatively small (e.g., between 8 and 10). This strategy is discussed in (Frank, 1996) and implemented in (Huang et al., 2003). All parameters are assumed to be rotationally invariant within each sector. Rotational averaging of the power spectrum is performed within each sector to produce k averaged 1-D profiles. The unknown parameters associated with (3) are estimated separately within each sector by solving (16)-(22) within that sector. This procedure returns k defocus values $\Delta z^{(j)}$, $j = 1, 2, \dots, k$. These defocus values can be used to estimate the parameters Δz_0 , Δz_1 and θ_0 by solving a constrained nonlinear least squares (NLS) problem

$$\min_{\delta_0, \delta_1, \theta_0} \frac{1}{2} \sum_{j=1}^k \left[\Delta z_0 + \Delta z_1 \sin\left(2j \frac{2\pi}{k} - 2\theta_0\right) - \Delta z^{(j)} \right]^2 \quad (26)$$

subject to

$$\Delta z_0^{\min} \leq \Delta z_0 \leq \Delta z_0^{\max} \quad (27)$$

$$\Delta z_1^{\min} \leq \Delta z_1 \leq \Delta z_1^{\max} \quad (28)$$

$$\theta_0^{\min} \leq \theta_0 \leq \theta_0^{\max}. \quad (29)$$

We will demonstrate that this strategy works very well for images that contain a modest level of astigmatism.

When the experimental images contain significant amount astigmatism, one may need to divide the power spectrum of each image into a larger number of angular sectors in order to accurately determine the astigmatism parameters. The potential pitfall of this approach is that the signal to noise (SNR) ratio associated with the 1-D rotationally averaged power spectrum within each sector is likely to be very low, hence the defocus, the experimental B-factor and other parameters associated with each sector may not be reliably estimated. We argue that in this case, the collected images should be discarded anyway. However, such a decision calls for an image analysis tool that can automatically make a distinction between images that contain mild astigmatism and images that are too distorted to be useful. We have developed such a tool based on the active contour model (ACM) algorithm (Blake, 1998). The main idea behind the ACM algorithm is to use a special contour tracing technique to identify concentric Thon rings in the power spectra of each image. The ratio between the radii associated with the major and minor axes of these elliptically shaped rings are estimated through a least squares procedure. When the estimated ratio is much larger than one, the image would be excluded for subsequent image processing and reconstruction.

3. Numerical Methods

In this section, we describe numerical algorithms and software we use to tackle the constrained nonlinear minimization problem formulated above. We focus on the 1-D curve fitting formulation shown in (16)-(22), which can be used directly to estimate the unknown defocus, and the parameters for the envelope and noise functions when astigmatism and drift are negligible. When both astigmatism and anisotropic

experimental B-factor are present in the data, we divide the power spectrum into several angular sectors and perform separate 1-D curve fittings within each sector.

The constrained minimization problem described by (16)-(22) can be solved in a number of ways. Algorithms for solving general constrained nonlinear optimization problem include the quadratic penalty method, the log barrier method, the augmented Lagrangian method, and the sequential quadratic programming (SQP) method (Nocedal, 1999). We have chosen the SQP method because recent studies (Gould, 2004) indicate that the SQP method is the most effective one for small to medium sized problems, i.e., problems with less than a thousand variables and constraints.

To simplify the notation in the discussion that follows, we denote the set of nonlinear constraint functions in (21) and (22) by $q(x)$, where x is a column vector representation of the unknown parameters to be estimated, i.e.,

$$x = (\alpha, B, \Delta z, Q, n_1, n_2, n_3, n_4). \quad (31)$$

In this notation, all nonlinear constraints in (21) and (22) can be conveniently represented by a single vector inequality $q(x) \geq 0$.

The SQP algorithm searches for an optimal solution to (16)-(22) iteratively. In SQP, the approximate solution x_k is updated, at each step, by

$$x_{k+1} \leftarrow x_k + \tau_k p_k, \quad (32)$$

where the search direction p_k is obtained by solving a quadratic minimization problem of the form

$$\min_{p_k} \frac{1}{2} p_k^T H_k p_k + \nabla \rho(x_k)^T p_k \quad (33)$$

subject to the same bound constraints as defined in (17)-(20) and also the linearized constraint

$$\nabla q_j(x_k)^T p_k + q_j(x_k) \geq 0. \quad (34)$$

The matrix H_k in (33) is an approximate Hessian of the Lagrangian function

$$L(x, \mu) = \rho(x) - \sum_{j=1}^n \mu_j q_j(x), \quad (35)$$

evaluated at the k -th iterate x_k , and $\mu = (\mu_1, \mu_2, \dots, \mu_n)$ denotes a set of Lagrangian multipliers associated with the nonlinear constraints $q_j(x)$. The step length τ_k in (32) is chosen to minimize some merit function while keeping the approximate solution x_{k+1} within the bound constraints (Nocedal, 1999).

To use a constrained nonlinear optimization solver, one must provide procedures for calculating the objective function (13) or (24) and the constraint function $q(x)$. One may also provide a procedure for computing the gradient of the objective function and the constraint with respect to the unknown parameters in x . Because the derivatives of the objective and constraint functions with respect to the unknown parameters are easy to compute for the problem defined in (16)-(22), we carry out these operations explicitly. If the procedures for the gradient calculation are not supplied by the user, most software packages have the capability to compute approximate gradient through the technique of finite difference.

Several software packages have been developed to solve nonlinear constrained optimization problems using SQP. Among the most well known are NPSOL (Gill, 1986, Schittkowski, 1986, Gill, 2002) and the fmincon function in MATLAB (MathWorks, 2004). We use the MATLAB fmincon function for our implementation.

4. Results and Discussion

4.1. Image Data

Two types of datasets were generated to demonstrate the applicability of the proposed method for the microscope parameter determination. One data set was the focal series images of an amorphous carbon film, which was evaporated on freshly-cleaved mica surface and then transferred onto a holey grid. The images were recorded at 200 kV in a JEM2010F electron microscope onto the Gatan 4k×4k CCD camera (US4000) at an effective magnification of 110,400 ×. To assess the reliability and accuracy of our computational estimation scheme, we collected images of carbon film in a broad range of defocus settings from 0.2 to 5.0 μm underfocus.

Carbon film was chosen as the test specimen for our algorithm because of the ease of detecting the CTF rings in the power spectra of the images. The images were taken with pre-determined defocus so that we can assess the accuracy of the proposed computational procedure. Figure 1 shows a focal series of carbon-film images and their corresponding power spectra. In this case, the astigmatism was well adjusted to negligible level prior to the data collection. These represent the best type of data that could be recorded. To test the capability of astigmatism estimation, we purposely introduced a mild level of stigmatism in the image which power spectrum is shown in Figure 2. The structure factor for the carbon film data set was estimated from electron diffraction pattern of carbon film (courtesy of Dr. Jaap Brink).

The second set of data was the P22 mature phage particles recorded onto photographic films (Kodak SO-163) in the JEM3000SFF electron microscope operated at 300 kV and specimen temperature of 4.2K. Images between 0.5 to 3 μm underfocus were used in this test (e.g., Figure 3). In this dataset, no carbon film was used to determine the CTF and associated parameters because the ice-embedded virus particles are suspended across holes with no support film. The data were digitized with a Nikon scanner at a scanning interval of 1.06 Å/pixel. The structure factor associated with P22 mature phage was obtained from the modified X-ray solution scattering curve (Thuman-Commike et al., 1999) to yield the best fit for a broad spectra of spatial frequencies to the cryo-EM data.

4.2. CTF and associated parameter estimation of carbon film images with negligible astigmatism

Using the estimated structure factor, we applied the constrained nonlinear minimization algorithm (the MATLAB `fmincon` function) discussed in Section 3 to each individual power spectrum image shown in Figure 1. The bound constraints for each of the parameters are listed in Table 1. The cutoff frequencies defined in (eqn. 11) were chosen to be $s_{\min} = 0.02 \text{ \AA}^{-1}$ and $s_{\max} = 0.2 \text{ \AA}^{-1}$. Our initial guesses for the B-factors, amplitude contrast ratio, and noise parameters were set to: $B=100$, $Q=0.1$, $n_1 = n_2 = n_3 = n_4 = 1.0$, respectively. Because our dataset contains images taken under a wide range of defocus settings, we tried five different starting guesses for the defocus value ($dz = 1.0, 3.0, 5.0, 7.0, 9.0 \text{ \mu m}$) for each of the runs. The CTF, envelope and noise parameters associated with minimum final objective function value (13) among the five runs were chosen to be our optimal estimation of the parameters. Note that all these procedures are implemented as part of the fitting processes, there is no need for user to provide initial guess for different micrographs and to repeat the runs.

Table 2 shows typical convergence history associated with each `fmincon` run. The first column of the table lists the iteration number. Column 2 gives the total number of function evaluations performed up to the k -th iteration. The progress of the convergence is measured by the value of the objective function (column 3), the magnitude of the directional derivative along the search direction (column 4), and the norm of the Lagrangian gradient (column 5) which provides the necessary first order optimal condition for the constrained nonlinear optimization problem defined in (16). The minimization procedure was terminated when the norm of the Lagrangian gradient is less than 0.05. The final objective function attains the value of 0.08 indicating a good match between the computational model defined by the estimated parameters and the power spectrum data. In Figure 4, we plot both the 1-D rotationally averaged power spectrum (the red curve) and the intensity curve defined by the function (12) using the optimal parameters returned from the constrained minimization procedure (blue curve). It is apparent that the difference between the experimental data (red curve) and the fitted data

(blue curve) is negligible in the frequency domain of interest. This suggests that our constrained optimization procedure successfully identified the global minimum of the objective function defined in (eqn. 16).

In Table 3, we list the optimal parameters associated with these carbon film images. The first row of the table gives the intended defocus values during the data collection where the defocus of the first image was determined using the DigitalMicrograph software (Gatan, Inc.) and the rest of the image defocuses were digitally set using the JAMES software (Booth *et al.*, 2004). Clearly, our estimations of the defocus values (the second row) match very well with the intended defocuses. This suggests that our fitting procedure can be used reliably to estimate the CTF parameters associated with images taken under a wide range of defocus settings.

4.3. The Importance of constraints

We shall emphasize the importance of constraints in the formulation of the minimization problem (16)-(22). Removing bound and/or nonlinear constraints from the problem formulation turns the CTF parameter estimation problem into a standard nonlinear least squares (NLSQ) problem which can be solved efficiently using a Gauss-Newton type of method (Dennis, 1981, More, 1984, Nocedal, 1999). However, unless the starting guess used by an NLSQ solver is sufficiently close to the optimal solution, one may obtain a solution that is physically wrong.

To illustrate this point, we use the power spectrum associated with the carbon film micrograph as an example. The image is taken under roughly $1.0 \mu\text{m}$ defocus (Figure 1(b)). We plot the contour of the function

$$\zeta(\Delta z, B) = \rho(\bar{\alpha}, B, \Delta z, \bar{Q}, \bar{n}_1, \bar{n}_2, \bar{n}_3, \bar{n}_4) \quad (36)$$

where ρ is the objective function defined in (13), and $\bar{\alpha}$, \bar{Q} and \bar{n}_i are fixed at the optimal values obtained from a manual fit. This function is the restriction of ρ to a 2-D subspace (spanned by B and Δz) with $\alpha = \bar{\alpha}$, $Q = \bar{Q}$, and $n_i = \bar{n}_i$, for $i = 1, 2, 3, 4$.

The contour plot shown in Figure 5 indicates that $\zeta(\Delta z, B)$ has two local minima within $[-2, 2] \times [0, 300]$. The desired local minimum is marked by a plus sign on the left half of the figure. If the update of the approximate minimizer in an NLSQ solver is not restricted to ensure that the underdefocus is used, the optimization procedure may converge to a local minimum that is entirely infeasible. Figure 5 also shows that the convergence of the optimization algorithm is less sensitive to the starting guess for B because $\zeta(\Delta z, B)$ appears to be convex in the direction of B within the neighborhood of interest.

To demonstrate the importance of the nonlinear constraints, we applied a NLSQ solver to (16) alone without additional constraints using a starting guess close to the optimal solution. Figure 6 shows that without the nonlinear constraints, the NLSQ solver converged to an infeasible solution in which the background term $N_2(s)$ in (12) becomes larger than the measured power spectrum at the second and the third CTF zeros. The quality of the fitting curve is considerably worse than that obtained from constrained nonlinear optimization shown in Figure 9(b).

4.4. Multiple starting guesses for the defocus parameter

The bound and nonlinear constraints established in (17)-(22) do not completely remove all undesirable local minima of (16). When particle images are collected under a broad range of defocus settings (i.e., the difference between Δz_{\min} and Δz_{\max} is large), the objective function in (16) may still have multiple local minima within the range of defocus of interest. Figure 7 shows the change of the objective function (16) with respect to different defocus values along the line segment defined by $x = (\bar{\alpha}, \bar{B}, \Delta z, \bar{Q}, \bar{n}_1, \bar{n}_2, \bar{n}_3, \bar{n}_4)$, where $\bar{\alpha}$, \bar{B} , \bar{Q} , and \bar{n}_i ($i = 1, 2, 3, 4$) are optimal parameters determined in advance and $\Delta z \in [0, 10]$ (μm). Clearly, the objective function $\rho(x)$ contains two local minima within this interval. The global minimum (marked by the circle) is located at $\Delta z = 0.53 \mu\text{m}$, which is the desired defocus value associated with this particular data set. However, if one chooses the initial guess of the defocus to be

around $8.0 \mu\text{m}$, for example, SQP may converge to an incorrect defocus value which corresponds to the undesirable local minimum located near $7.8 \mu\text{m}$.

To prevent SQP from converging to the wrong local minima within the defocus range of interest, we solve (16)-(22) with multiple starting guesses evenly distributed between Δz_{\min} and Δz_{\max} . Because the number of local minima within the defocus range of interest is typically small, we normally need to try only 3-5 different starting guesses.

4.5. Astigmatism estimation

When images contain a mild level astigmatism such as the one shown in Figure 2, we apply the practical procedure discussed in Section 2.5 to estimate all angular dependent parameters. To test this procedure on the power spectrum shown in Figure 2, we divided power spectrum evenly into eight angular sectors. Each sector was rotationally averaged to produce a 1-D curve to be fitted with the constrained nonlinear model described in (16)-(22). Figure 8(a) shows the 1-D curves generated from different angular sectors of the power spectrum shown in Figure 2. These curves differ slightly in the positions of their peaks and valleys, implying the variation of the defocus along different radial directions.

In Figure 8(b), we plot the defocus values derived from the constrained nonlinear minimization procedure (applied to each angular sector) against $\hat{\theta}_k$, where $\hat{\theta}_k$ is the angle formed by the bisector of the k -th angular sector and the horizontal axis. The estimated defocus values are marked by circles. An NLS algorithm was used to fit these defocus values to the analytical expression $\Delta z(\theta) = \Delta z_0 + \Delta z_1 \sin(2(\theta - \theta_0))$. The fitting procedure produces $\Delta z_0 = 0.546 \mu\text{m}$, $\Delta z_1 = 0.017 \mu\text{m}$ and $\theta_0 = 0.059$.

We should point out that multiple starting guesses are typically required to solve the constrained NLS problem (16)-(22) in at least one of the angular sectors. Once the parameter estimation problem has been solved for that angular sector, the estimated parameters associated with that particular angular sector can be used as the starting guesses for the minimization procedure applied to other angular sectors. Since the

variation of the CTF, envelope and noise background parameters are typically small for images that contain mild astigmatism and drift, the use of this starting guess often enables the optimization routine to converge in a few iterations.

It is worth pointing out that typical cryo-EM images to be used for image reconstruction have a negligible level of astigmatism. A level that is much smaller than that shown in this test case. In fact, single particle cryo-EM studies have been able to obtain near atomic resolution ($\sim 4\text{\AA}$) 3-D reconstructions without the need of considering astigmatism in the images (Jiang et al., 2008, Ludtke et al., 2008). While we have shown here that this method can successfully handle the images with a significant level of astigmatism, the use of this functionality is rarely necessary in practice for single particle cryo-EM study.

4.6. Parameters Estimation with Images of Ice-Embedded Particles

Often the ice-embedded particles are suspended across holes without any carbon substrate as shown in Figures 3(a) and 3(b). The incoherent average of the power spectrum of the boxed-out particles has been used to determine the CTF and associated parameters by manual fitting procedure (Saad et al., 2001). Figure 9 shows that the fitting curves produced by the optimization procedure match extremely well with the 1-D rotationally averaged power spectra of the micrographs. Furthermore, the determined parameters compare well with those determined manually. These data show that even without the carbon support film, our fitting method works equally well with authentic ice-embedded particle images.

4.7. CPU Requirements

The average amount of CPU time required to fit a micrograph of 240MB size on a 1.8 Ghz Pentium 4 laptop is under a minute. With a meticulous choice of the convergence tolerance and maximum iteration number, the computational time can be further reduced. In practice, the range of defocus values associated with the experimental data is often much less than $10\ \mu\text{m}$. Thus, we may either tighten the bound constraints associated with the defocus or reduce the number of initial guesses to further speed up the computation.

4.8. General Accessibility of the software

The algorithms and techniques described here have been implemented as a standard-alone Python script, which is available on the NCMI Web site (<http://ncmi.bcm.edu/software/fitctf>). It runs on all the major computer platforms (Linux, Windows, MacOS X). It allows the user to perform CTF estimation in a fully automated fashion once the 2-D single particles have been boxed out from the micrograph. The resulting parameters are formatted to be compatible to single particle image processing software package EMAN.

5. Conclusions

An accurate determination of the CTF and associated parameters are essential in 3-D structural determination of biological samples. Though manual fitting methods for these determinations have been used successfully, it is time consuming and subject to human errors. The proposed constrained nonlinear minimization algorithm has provided not only objective and accurate but also automated protocol. The examples shown here demonstrate its utility not only on images of carbon film but also on ice-embedded biological particles. A unique feature of this algorithm is the ability of determining images taken at smaller defocus (i.e. 0.5 μm), which is desirable for high resolution structure determination (Jiang et al., 2008, Liu et al., 2007). For such a small defocus image, it is generally difficult to estimate its CTF with confidence by a manual fitting method. This algorithm has been successfully applied to determining the CTF and associated parameter in images used for 3D reconstruction in a broad range of resolutions (e.g. Chang et al, 2006, Jiang et al., 2006, 2008).

Though the extension (Figures 8 and 9) of our nonlinear optimization based fitting procedure can handle images with astigmatism and anisotropic B-factor, the accuracy of their determination may not be very high because of the need of dividing the power spectrum into multiple sectors resulting in poor signal-to-noise ratio. On the other hand, many structures of single particles have been determined to 4-9 \AA resolution without considering astigmatism and anisotropic B-factor by excluding those images with

apparent astigmatism and/or drift (Zhou *et al.*, 2001, Jiang *et al.*, 2003, Ludtke *et al.*, 2004, Ludtke *et al.*, 2005). More recently, structure of epsilon15 phage has been solved to 4.5 Å (Jiang *et al.*, 2008) with images of which the CTF and associated parameters were determined with this procedure. Therefore, our proposed algorithm will be of immediate usage for data up to this resolution range at which the resulting structure is interpretable in terms of protein backbone trace.

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Table 1. The bound constraints for the CTF, envelope and noise parameters to be estimated.

Parameters	Lower bound (carbon film)	Upper bound (carbon film)	Lower bound (P22 phage)	Upper bound (P22 phage)
Δz	0	9.0	0	4.0
B	0	∞	0	∞
α	0	∞	0	∞
Q	0	0.2	0	0.1
n_1	∞	∞	∞	∞
n_2	∞	∞	∞	∞
n_3	0.0	∞	0.0	∞
n_4	∞	∞	∞	∞

Table 2. A typical convergence history of `fmincon` when it is applied to CTF parameter estimation of a carbon film. The first column gives the iteration number. The second column gives the total number of function evaluations at the end of the k -th iteration. The third column lists the relative norm of the residual. The fourth column gives the directional derivative at the k -th iteration. The last column gives the first-order optimality of the constrained optimization problem.

k	f-count	$\eta(\alpha, \beta, \Delta z, Q, \{n_i\})$	$\nabla \eta^T s_k$	$\ L\ $
1	9	453.9	-1440	3160
11	126	32.82	-0.33	14.4
21	232	28.71	0.002	6.02
31	341	28.34	-0.06	4.77
41	453	24.00	-0.2	18.4
51	559	21.12	-0.39	25.8
61	662	3.538	1.07	85.4
71	762	0.09	-0.02	3.63
76	812	0.08	$-7e-7$	0.04

Table 3. The CTF, envelope function, and noise background parameters returned from the constrained nonlinear minimization procedure (the MATLAB *fmincon* function) for carbon film images taken at different defocus settings. The first row (bold faced numbers) of the table shows the intended defocus under which each image is taken.

Intended Δz (μm)	0.2	0.5	1.0	1.5	2.0	3.0	4.0	5.0	9.0
Determined Δz (μm)	0.23	0.54	1.04	1.55	2.05	3.06	4.10	5.13	9.36
B (\AA^2)	119	104	104	104	106	110	116	126	186
α	12.0	8.14	6.66	5.91	5.43	4.99	4.72	4.63	4.39
Q	0.03	0.0	0.01	0.02	0.03	0.03	0.03	0.03	0.02
n_1	-2.26	26.0	16.4	13.5	23.2	31.6	33.0	27.9	33.5
n_2	17.0	-33.6	-10.1	-2.8	-22.5	-39.8	-4.2	-27.9	-34.7
n_3	5.8	343	86.3	56.9	197	547	719	435	1402
n_4	-0.7	4.6	-2.5	-1.3	3.4	4.5	4.6	-3.4	3.6

Figure Captions

Figure 1. 200-kV CCD frames of carbon-film (a-d) and their corresponding power spectra (e-h) taken under different defocus settings: (a) $0.5 \mu\text{m}$ (b) $1 \mu\text{m}$ (c) $3 \mu\text{m}$ (d) $4 \mu\text{m}$.

Figure 2. Power spectrum of a carbon film image with a mild astigmatism.

Figure 3. 300-kV CCD frames of P22 mature phage (a) and (b) taken under different defocus settings and the corresponding power spectra (c) and (d). The estimated defocus is $0.41 \mu\text{m}$ for (a) and $1.14 \mu\text{m}$ for (b).

Figure 4. Comparing the 1-D rotationally averaged power spectrum data (the red dots) with the CTF fitting curves (the solid blue curves) generated by the constrained nonlinear minimization on a series of carbon-film images shown in Figures 1(a)-(d). The dash-dotted curves in (a)-(d) show the noise background estimated from solving equations (16)-(22).

Figure 5. The contour of $\zeta(\Delta z, B)$ associated with the image of carbon film taken under roughly $1.0 \mu\text{m}$ defocus. This function has two local minima. The desired local minimum is marked by a plus sign on the left half of the contour plot.

Figure 6. Without imposing the nonlinear constraints (21) and (22), applying a NLSQ fitting procedure to the P22 mature phage particle image shown in Figure 3(b) returns a solution in which the background term $N(s)$ in (12) (the black dash-dotted curve) is larger than the power spectrum (red dots) near 0.06 and 0.14 \AA^{-1} .

Figure 7. Variation of the objective function (16) with respect to the defocus values along the line segment defined by $x = (\bar{\alpha}, \bar{B}, \Delta z, \bar{Q}, \bar{n}_1, \bar{n}_2, \bar{n}_3, \bar{n}_4)$, where $\bar{\alpha}$, \bar{B} , $Q = \bar{Q}$, and

\bar{n}_i ($i = 1,2,3,4$) are optimal parameters determined in advance. Clearly, (16) has two local minima in $[0,10]\mu\text{m}$. The desired global minimum is marked by a circle near $0.5 \mu\text{m}$.

Figure 8. (a) Variation of the 1-D power spectra obtained from rotationally averaging the 2-D power spectrum shown in Figure 2 among 8 even divided angular sectors. (b) Variation of the defocus along different radial directions. The circles represent the defocus value estimated from each angular sector of the power spectrum. The curve corresponds to the function $\Delta z(\theta) = \Delta z_0 + \Delta z_1 \sin(2(\theta - \theta_0))$, where $\Delta z_0, \Delta z_1, \theta_0$ are estimated by a nonlinear least squares fitting procedure.

Figure 9. Comparing the 1-D rotationally averaged power spectrum data (the red dots) with the CTF fitting curves (the solid blue curves) generated by the constrained nonlinear minimization on the P22 mature phage images shown in Figure 3(a) and (b). The dash-dotted curves in (a) and (b) show the noise background estimated from solving equations (16)-(22).

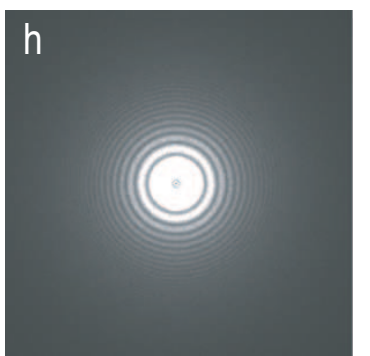
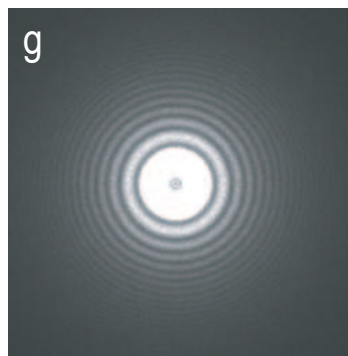
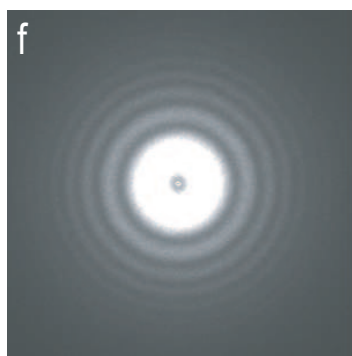
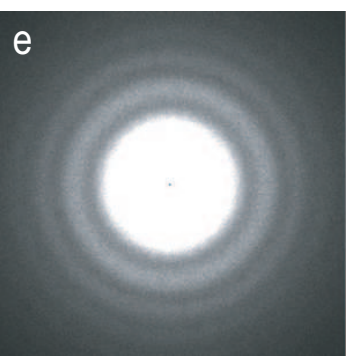
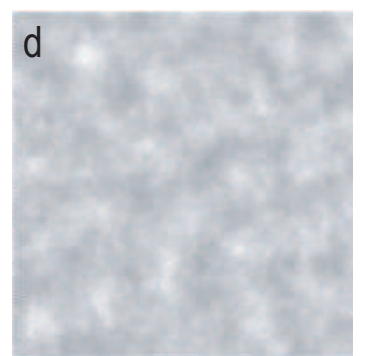
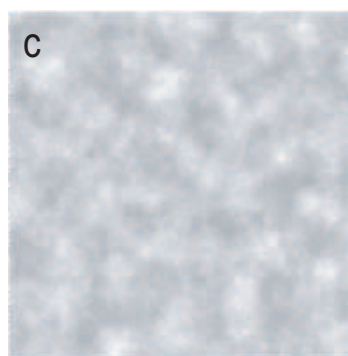
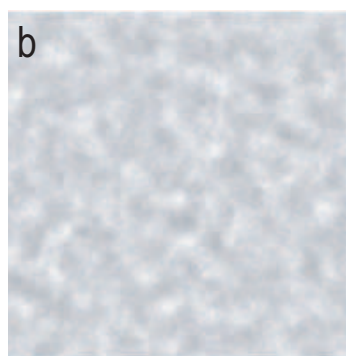
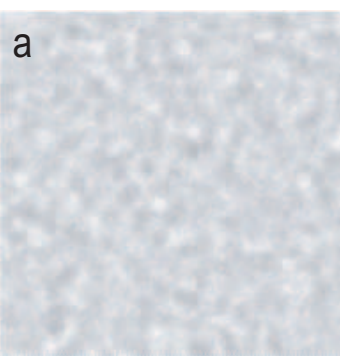
References

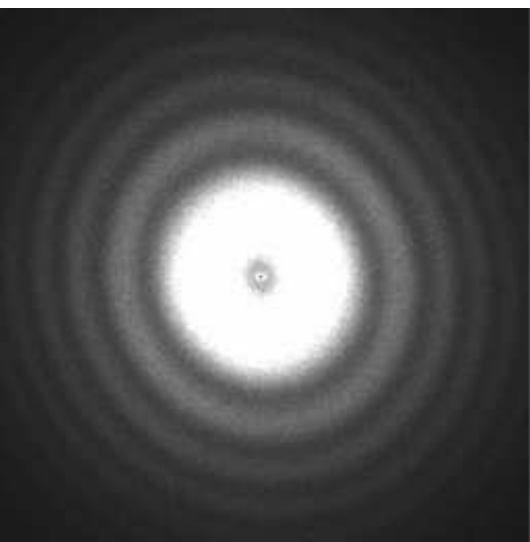
- Blake, A. a. I., M. (1998) *Active Contours*, Springer Verlag, New York.
- Booth, C. R., Jiang, W., Baker, M. L., Zhou, Z. H., Ludtke, S. J. & Chiu, W. (2004) A 9 angstroms single particle reconstruction from CCD captured images on a 200 kV electron cryomicroscope. *J Struct Biol*, **147**, 116-127.
- Chang, J., Weigele, P., King, J., Chiu, W. and Jiang, W. (2006). Cryo-EM asymmetric reconstruction of bacteriophage P22 reveals organization of its DNA packaging and infecting machinery. *Structure* **14**, 1073-82.
- Conway, J. F. & Steven, A. C. (1999) Methods for reconstructing density maps of "single" particles from cryoelectron micrographs to subnanometer resolution. *J Struct Biol*, **128**, 106-118.
- Dennis, J. E., Gay, D. M., and Welsch, R. E. (1981) NL2SOL - an adaptive nonlinear least squares algorithm. *ACM Trans. Math. Software*, **7**, 369-383.
- Erickson, H. P. & Klug, A. (1970) The Fourier transform of an electron micrograph: effects of defocussing and aberrations, and implications for the use of underfocus contrast enhancement. *Phil. Trans. Roy. Soc. Lond. B*, **261**, 105-118.
- Frank, J. (1969) Nachweis von objektbewegungen im lichtoptischen diffraktogramm von elektronmikroskopischen aufnahmen. *Optik*, **30**, 171-180.
- Frank, J. (1976) Determination of source size and energy spread from electron micrographs using the method of Young's fringes. *Optik*, **44**, 379-391.
- Frank, J. (1996) *Three-dimensional electron microscopy of macromolecular assemblies*, Academic Press, San Diego.
- Gill, P. E., Murray, W. and Saunders, M. A. (2002) SNOPT: An SQP algorithm for large-scale constrained optimization. *SIAM J. Optim*, **12**, 979-1006.
- Gill, P. E., Murray, W., Saunders, M. A. and Wright, M. H. (1986) User's guide for NPSOL version 4.0: A FORTRAN package for nonlinear programming. Stanford University.
- Gould, N. I. M., Orban, D. and Toint, P. L. (2004) GALAHAD, a library of thread-safe FORTRAN 90 packages for large-scale nonlinear optimization. *ACM Trans. Math. Software*, **29**, 353-372.

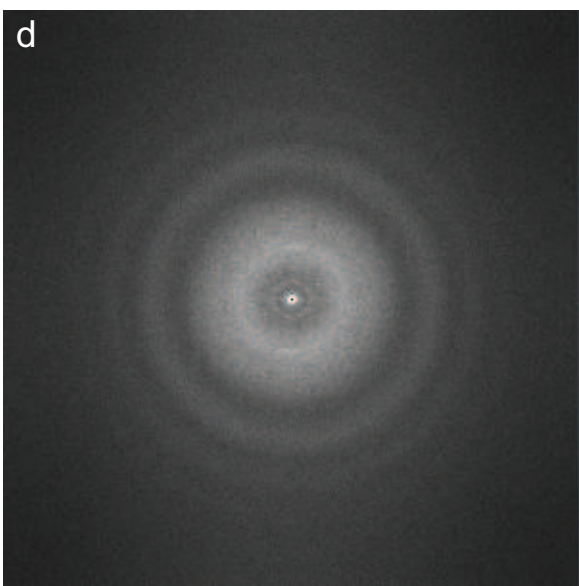
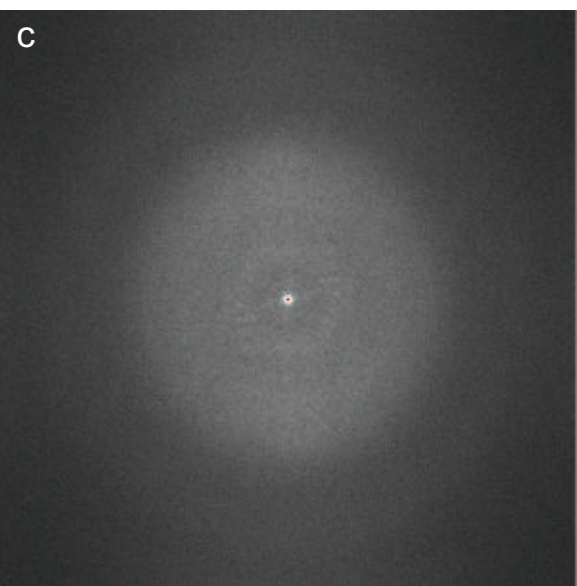
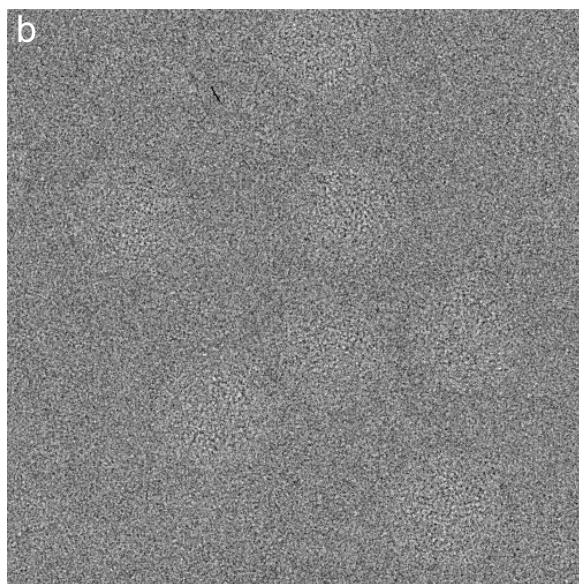
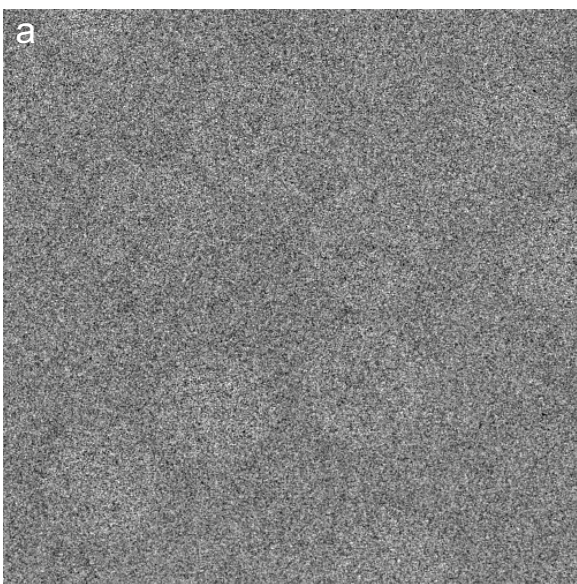
- Hanszen, K. J. (1967) New knowledge on resolution and contrast in the electron microscope image. *Naturwissenschaften*, **54**, 125-133.
- Hanszen, K. J. (1971a) The optical transfer theory of the electron microscope: Fundamental principles and applications. In: *Advances in optical and electron microscopy* (ed. R. a. C. V. E. Barer). Academic Press, New York.
- Hanszen, K. J. a. T., L. (1971b) Die kontrastübertragung im Elecktonmikroskop bei partiell koharenter beleuchtung. *Optik*, **33**, 166-182.
- Huang, Z., Baldwin, P. R., Mullapudi, S. & Penczek, P. A. (2003) Automated determination of parameters describing power spectra of micrograph images in electron microscopy. *J Struct Biol*, **144**, 79-94.
- Jiang, W., Baker, M. L., Jakana, J., Weigele, P. R., King, J. & Chiu, W. (2008) Backbone structure of the infectious epsilon15 virus capsid revealed by electron cryomicroscopy. *Nature*, **451**, 1130-1134.
- Jiang, W., Li, Z., Zhang, Z., Baker, M. L., Prevelige, P. E. & Chiu, W. (2003) Coat protein fold and maturation transition of bacteriophage P22 seen at sub-nanometer resolutions. *Nat Struct Biol*, **10**, 131-135.
- Liu, X., Jiang, W., Jakana, J. & Chiu, W. (2007) Averaging tens to hundreds of icosahedral particle images to resolve protein secondary structure elements using a Multi-path Simulated Annealing optimization algorithm. *J Struct Biol*, **160**, 11-27.
- Ludtke, S. J., Baker, M. L., Chen, D. H., Song, J. L., Chuang, D. T. & Chiu, W. (2008) De Novo Backbone Trace of GroEL from Single Particle Electron Cryomicroscopy. *Structure*, **16**, 441-448.
- Ludtke, S. J., Baldwin, P. R. & Chiu, W. (1999) EMAN: semiautomated software for high-resolution single-particle reconstructions. *J Struct Biol*, **128**, 82-97.
- Ludtke, S. J., Chen, D. H., Song, J. L., Chuang, D. T. & Chiu, W. (2004) Seeing GroEL at 6 Å resolution by single particle electron cryomicroscopy. *Structure (Camb)*, **12**, 1129-1136.
- Ludtke, S. J., Jakana, J., Song, J. L., Chuang, D. T. & Chiu, W. (2001) A 11.5 Å single particle reconstruction of GroEL using EMAN. *J Mol Biol*, **314**, 253-262.

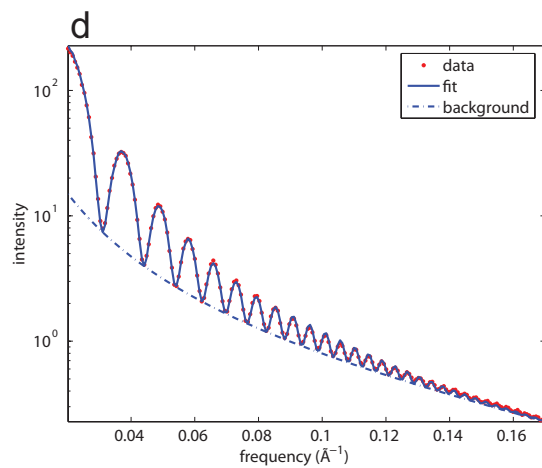
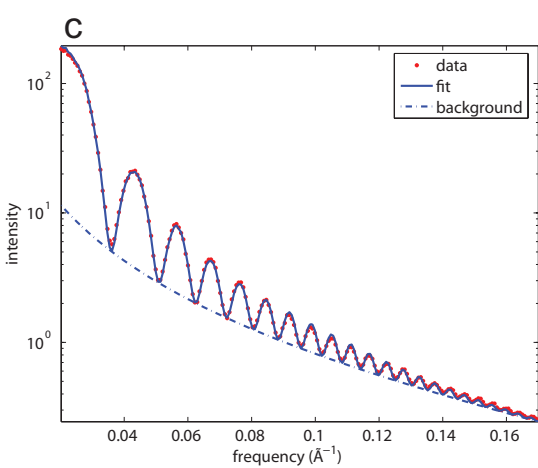
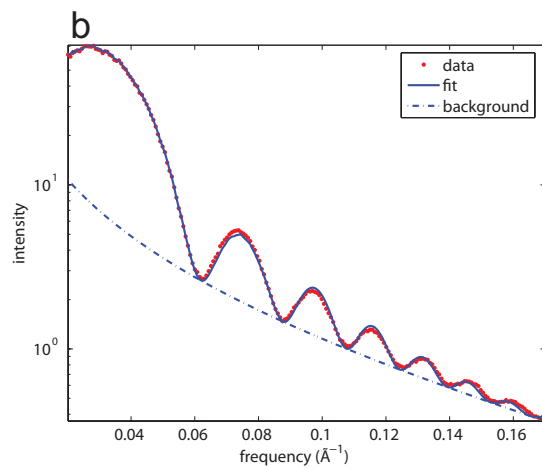
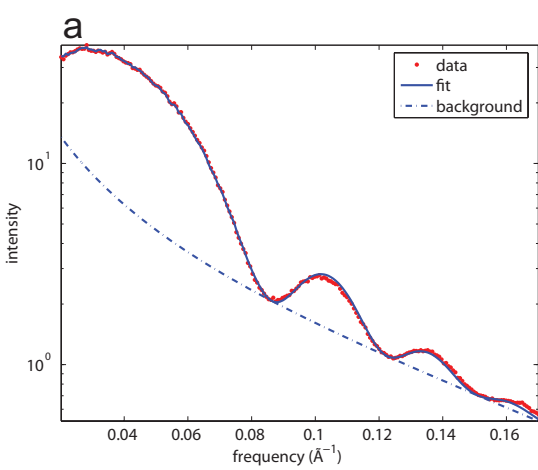
- Ludtke, S. J., Serysheva, II, Hamilton, S. L. & Chiu, W. (2005) The pore structure of the closed RyR1 channel. *Structure (Camb)*, **13**, 1203-1211.
- Mallick, S. P., Carragher, B., Potter, C. S. & Kriegman, D. J. (2005) ACE: automated CTF estimation. *Ultramicroscopy*, **104**, 8-29.
- MathWorks (2004) *Optimization Toolbox User's Guide*, The MathWorks, Inc., Natick, MA.
- More, J. J., Sorensen, D. C., Hillstrom, K. E. and Garbow, B. S. (1984) The MINPACK project. In: *Sources and Development of Mathematical Software* (ed. W. J. Cowell). Prentice Hall, NJ.
- Nocedal, J. a. W., S. J. (1999) *Numerical Optimization*, Springer, New York.
- Saad, A., Ludtke, S. J., Jakana, J., Rixon, F. J., Tsuruta, H. & Chiu, W. (2001) Fourier amplitude decay of electron cryomicroscopic images of single particles and effects on structure determination. *J Struct Biol*, **133**, 32-42.
- Sander, B., Golas, M. M. & Stark, H. (2003) Automatic CTF correction for single particles based upon multivariate statistical analysis of individual power spectra. *J Struct Biol*, **142**, 392-401.
- Schittkowski, K. (1986) NLPQL: A FORTRAN subroutine for solving constrained nonlinear programming problems. *Annals of Operations Research*, **5**, 485-500.
- Schmid, M. F., Sherman, M. B., Matsudaira, P., Tsuruta, H. & Chiu, W. (1999) Scaling structure factor amplitudes in electron cryomicroscopy using X-Ray solution scattering. *J Struct Biol*, **128**, 51-57.
- Thon, F. (1971) *Phase contrast electron microscopy*, Academic Press, New York.
- Thuman-Commike, P. A., Tsuruta, H., Greene, B., Prevelige, P. E., Jr., King, J. & Chiu, W. (1999) Solution x-ray scattering-based estimation of electron cryomicroscopy imaging parameters for reconstruction of virus particles. *Biophys J*, **76**, 2249-2261.
- Toyoshima, C., Yonekura, K. & Sasabe, H. (1993) Contrast transfer for frozen-hydrated specimens II. Amplitude contrast at very low frequencies. *Ultramicroscopy*, **48**, 165-176.

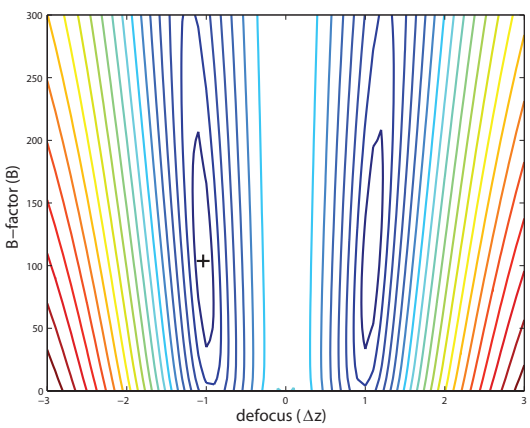
- Velazquez-Muriel, J. A., Sorzano, C. O., Fernandez, J. J. & Carazo, J. M. (2003) A method for estimating the CTF in electron microscopy based on ARMA models and parameter adjustment. *Ultramicroscopy*, **96**, 17-35.
- Zhou, Z. H., Baker, M. L., Jiang, W., Dougherty, M., Jakana, J., Dong, G., Lu, G. & Chiu, W. (2001) Electron cryomicroscopy and bioinformatics suggest protein fold models for rice dwarf virus. *Nat Struct Biol*, **8**, 868-873.
- Zhu, J., Penczek, P. A., Schroder, R. & Frank, J. (1997) Three-dimensional reconstruction with contrast transfer function correction from energy-filtered cryoelectron micrographs: procedure and application to the 70S Escherichia coli ribosome. *J Struct Biol*, **118**, 197-219.



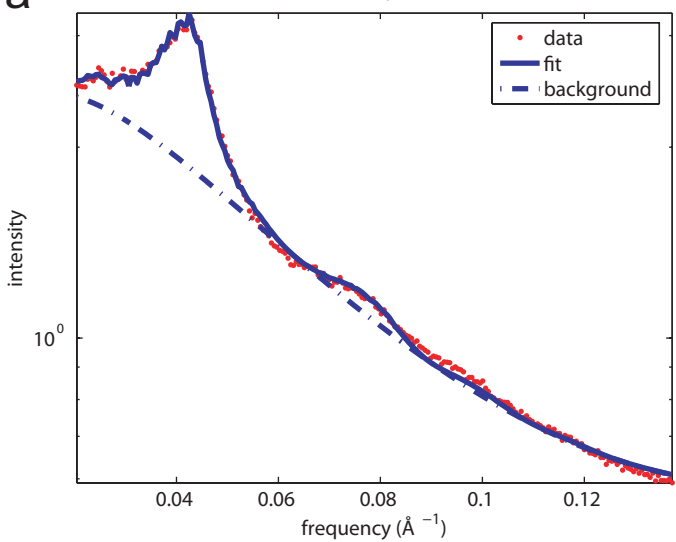








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