Enhanced diffractograms for image sorting and improvement of robustness and accuracy of CTF estimation: Application on high-resolution reconstruction of glutamate synthase and phosphorylase kinase

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PSD:

CTF:

Abstract: ADSTRUCT In a recent work, we developed a method for improving the visibility of diffraction rings computed from cryo-electron micrographs of vitreous ice (without carbon film or high concentration of diffracting material) [1]. We used these enhanced power spectrum densities (enhanced PSDs) to semi-automatically detect and remove micrographs with a strong envelope function or with a low signal-to-noise ratio (e.g., non-diffracting micrographs or their areas) or those that introduce errors in the global 3D map (e.g., drifted micrographs or charged (e.g., non-diffracting micrographs or their areas) or those that infroduce errors in the global 3D may (e.g., drifted micrographs or charged areas). This sorting is based on the normalized cross-correlation between enhanced PSDs and their copies rotated by 90 degrees. In the present work, we show single-particle reconstruction of two macromolecular complexes, using this image sorting method. The two complexes are: 1) bacterial glutamate synthase at resolution of 9.8 Å according to the FSC_{0.5} criterion or 8.3 Å according to the 1/2-bit criterion, and 2) mammalian phosphorylase kinase at resolution of 9.8 Å according to the FSC_{0.5} criterion or 8.4 Å according to the 1/2-bit criterion, we also show the use of enhanced PSDs in improving robustness and accuracy of contrast transfer function (CTF) estimation on images that are kept show the use of enhanced PSDs in improving robustness and accuracy of contrast transfer function (C1F) estimation on images that are kept after sorting. We show that the use of enhanced PSDs can improve the estimation of a full two-dimensional PSD model that takes into account the envelope, such as the one proposed in [2]. Determination of defocus parameters (maximum defocus, minimum defocus, angle of astigmatism) is the most difficult part of the PSD model estimation. This is why the algorithm first estimates roughly as many parameters of the PSD model as possible (i.e., background and envelope parameters), and then, estimates very accurately defocus parameters are refined. The source code of the algorithms for PSD enhancement and CTF estimation is available for implementing in any image processing package, upon request from the authors or as a part of the onen-source package Xminn (http://xminn.en/ bsic.es/) equest from the authors or as a part of the open-source package Xmipp (http://xmipp.cnb.csic.es/).





Sorting based on normalized cross-correlation (NCC) between the enhanced PSDs and their copies rotated by 90°





Sixteen lowest NCC values: Drifted (B-D,G-M), nondiffracting (A,E,F), and strongly astigmatic (N-P) images

> 3. Examples of 3D cryo-EM maps computed after image sorting based on enhanced-PSD sorting

Resolution: 9.8 Å (FSC0 5), 8.3 Å (1/2-bit), 7.7 Å (30)



Bacterial Glutamate synthase





nmalian Phosphorylase kinase Resolution: 9.8 Å (FSC_{0.5}), 8.4 Å (1/2-bit), 7.2 Å (3σ)









4. Power spectrum density model

 $PSD_{theoretical}(\boldsymbol{\varpi}) = \boldsymbol{K}^2 \left| \boldsymbol{H}(\boldsymbol{\varpi}) \right|^2 + \boldsymbol{S}_{N}(\boldsymbol{\varpi})$

 $H(\boldsymbol{\varpi}) = -E_{\Theta_{1,2}}(\boldsymbol{\varpi})(\sin(\chi_{\Theta_{1,2}}(\boldsymbol{\varpi})) + Q_0\cos(\chi_{\Theta_{1,2}}(\boldsymbol{\varpi})))$

 $\rho(x,y)$: Correlation coefficient between x and y

w : Weight determining the influence of the PSD enhancement term on the optimization algorithm (w>0) Ω : Annular mask in the frequency domain

6. Results of PSD estimation on a single micrograph







Literature

- Jonic S, Sorzano CO, Cottevieille M, Larquet E, Boisset N (2007). A novel method for improvement of visualization of power spectra for sorting cryo-electron micrographs and their local areas. J Struct Biol 157(1): 156-67.
- [2] Velazquez-Muriel JA, Sorzano CO, Fernandez JJ, Carazo JM (2003). A method for estimating the CTF in electron microscopy based on ARMA models and parameter adjustment. Ultramicroscopy 96(1): 17-35. [3] Sorzano CO, Jonic S, Nuñez R, Boisset N, Carazo JM. Fast, robust and accurate determination of transmission electron microscopy contrast transfer
- function. (submitted to J Struct Biol, Nº JSB-07-64).











