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Total generalized variation regularization for multi-modal electron tomography†

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In multi-modal electron tomography, tilt series of several signals such as X-ray spectra, electron energy-loss spectra, annular dark-field, or bright-field data are acquired at the same time in a transmission electron microscope and subsequently reconstructed in three dimensions. However, the acquired data are often incomplete and suffer from noise, and generally each signal is reconstructed independently of all other signals, not taking advantage of correlation between different datasets. This severely limits both the resolution and validity of the reconstructed images. In this paper, we show how image quality in multi-modal electron tomography can be greatly improved by employing variational modeling and multi-channel regularization techniques. To achieve this aim, we employ a coupled Total Generalized Variation (TGV) regularization that exploits correlation between different channels. In contrast to other regularization methods, coupled TGV regularization allows to reconstruct both hard transitions and gradual changes inside each sample, and links different channels at the level of first and higher order derivatives. This favors similar interface positions for all reconstructions, thereby improving the image quality for all data, in particular, for 3D elemental maps. We demonstrate the joint multi-channel TGV reconstruction on tomographic energy-dispersive X-ray spectroscopy (EDXS) and high-angle annular dark field (HAADF) data, but the reconstruction method is generally applicable to all types of signals used in electron tomography, as well as all other types of projection-based tomographies.

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Introduction

Electron tomography using a scanning transmission electron microscope (STEM) is a versatile tool for the investigation of nanomaterials in three dimensions down to the single atomic level.^{1,2} Insight into the 3D elemental and chemical make-up of a sample is provided by the spectroscopic signals originating from inelastic scattering in analytical tomography experiments.³ Energy losses of electrons traversing a thin TEM speci-

men can be measured using energy-filtered TEM (EFTEM)^{4–7} or electron energy-loss spectroscopy (EELS)^{8–10} and characteristic X-rays can be recorded using energy-dispersive X-ray spectroscopy (EDXS).^{11–13} Also, EELS and EDXS signals can be acquired simultaneously for combined analytical tomography experiments.¹⁴ Over the last few years, analytical tomography has thereby become an important characterization method in nanoscale materials science.^{13,15–18}

The 3D resolution of a sinogram is generally limited by the number of projections which can be acquired in an experiment and by the achievable tilt range. Reasons for these limitations are the overall dose the sample can sustain, the accuracy of the sample positioning mechanism for very small tilt steps, the geometry of the sample or the sample holder, as well as the limited time for an experiment. This leads to the situation that image reconstruction from an electron tomography experiment usually amounts to solving an ill-posed inverse problem with more unknown parameters than measurements. Especially for analytical tomography data, also the signal-to-noise ratio is very low. Therefore, *a priori* knowledge and regularization are essential to find better solutions for a tomographic reconstruction. In this context sparse recovery has become increasingly important over the last few years.^{19–22} In sparse recovery, the signal of interest is assumed to have sparse representation

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when transformed to a suitable basis. If such a transformation is possible, minimization of the ℓ_1 -norm in this domain enforces sparsity and can lead to an accurate solution.

The most important type of sparsity in electron tomography is sparsity of the image (or volume) gradient.^{19,23,24} Known as Total Variation (TV)²⁵ minimization, ℓ_1 -minimization of the image gradient leads to an image (or volume) with locally constant regions, separated by sharp interfaces. For materials science samples, it can often be assumed that well separated phases exist within a sample, which is why TV minimization has found a range of applications in electron tomography.^{26–29} It has been shown that TV minimization can be especially favorable for analytical electron tomography experiments, where usually only limited and noisy data are available.^{14,30}

Two major limitations can be identified with current TV minimization approaches and their application in analytical tomography. First, the underlying model of TV minimization assumes that the sample only contains sharp interfaces between different phases, severely restricting the range of materials that can be analyzed using TV minimization. All types of gradual changes between different sample regions, such as diffusion at interfaces, are typically not correctly recovered. Second, in analytical tomography and other multi-modal tomography experiments, different signals are acquired at the same time, containing complementary information. But in most approaches this is not exploited but rather each signal is reconstructed independently. Until recently, there was only one attempt to link the information of different signals,³¹ where the intensity of the HAADF signal was expressed as a linear combination of EDXS signals and reconstructed with a conventional reconstruction algorithm.

Here we present an approach for resolving these issues for multi-modal tomography. We apply coupled Total Generalized Variation (TGV)^{32–34} regularization, which allows for both sharp interfaces and gradual intensity changes in the reconstruction and exploits inter-channel correlation. This yields a joint multi-channel reconstruction of HAADF and EDXS data. We demonstrate our approach on phantom data and on an Al-5 wt% Si alloy with 50 ppm Na and 6100 ppm Yb,¹⁴ where the

precipitation of Yb and Si takes place at the nanoscale, but it is generally applicable to analytical electron tomography of all types of nanostructured materials. The basic concept of multi-modal STEM tomography is depicted in Fig. 1. Several tilt series are acquired at the same time in the microscope. In the joint reconstruction all tilt series are used together to obtain a joint multi-modal reconstruction.

We note that in two recent, independent studies,^{35,36} a coupled Total Variation approach was proposed to link data and create smooth reconstructions. While these studies overcome the second limitation of existing approaches as mentioned above, the usage of TV regularization still limits their applicability to piecewise constant densities with sharp interfaces. However, a major benefit of three-dimensional local elemental distributions lies in analyzing local concentration variations. While the compositions of well separated phases are often accessible already by 2D investigations after appropriate sample preparation, local variations are always masked due to projection. A recent example is the unveiling of elemental concentration variations within spherical precipitates in an AlMgScZr-alloy.¹⁷ In such situations it would not be possible to use TV regularization. Beyond overcoming this drawback, a second novelty of our approach is that we developed – and make publicly available³⁷ – a Python³⁸/OpenCL³⁹-based implementation for all components of the reconstruction algorithm, which is crucial in particular for 3D reconstructions as it significantly reduces the computation time. A graphical user interface connected to the code allows for simple and efficient execution. In particular, we developed a custom, GPU-based implementation of the forward operator and its adjoint, ensuring, in contrast to other existing implementations, numerical adjointness of the two, which is in particular crucial for the convergence of iterative minimization algorithms.

We also note that TGV regularization for both single- and multi-channel data has been demonstrated to work well and to be superior to TV regularization in various applications.^{40–45} In particular, in the context of joint magnetic resonance (MR) imaging and positron emission tomography (PET), multi-

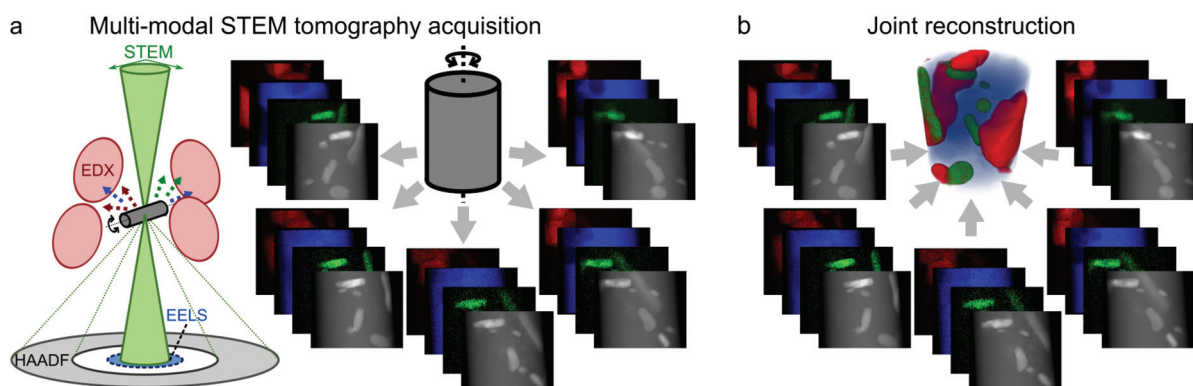


Fig. 1 Basic principles of multi-modal STEM tomography. (a) Simultaneous acquisition of several tilt series (EDXS/EELS+HAADF). (b) Joint reconstruction of all tilt series.



channel TGV regularization was shown to allow for improved recovery of quantitative measurements compared to single-channel approaches.⁴⁶ Also theoretical results on well-posedness and stability for multi-channel regularization are available.⁴⁷ Building on the existing code for MR reconstruction,⁴⁰ a first application of TGV in the context of two-dimensional single-channel electron tomography was also presented in a conference proceedings paper⁴⁸ and shown to be superior to TV regularization.

Results

The main principles of multi-modal TGV reconstruction

The proposed tomographic reconstruction framework is based on a convex variational method employing volumetric coupled second order Total Generalized Variation (TGV) regularization, and the Kullback–Leibler (KL) divergence as a discrepancy measure. Given some measured multi-channel† data $f = (f^1, \dots, f^C)$, a reconstruction $u = (u^1, \dots, u^C)$ is obtained as a solution to the convex minimization problem

$$\min_{u \geq 0} \left[\sum_{c=1}^C \mu^c D_{\text{KL}}(T^c u^c, f^c) \right] + \text{TGV}_\alpha^2(u). \quad (1)$$

Here, $D_{\text{KL}}(\tilde{u}, \tilde{f}) := \int \tilde{u} - \tilde{f} \log(\tilde{u})$ corresponds (up to constants and non-negativity constraints) to the Kullback–Leibler divergence, which is an appropriate data fidelity measure in the presence of Poisson noise. The operators T^c map each image channel to the corresponding data space and comprise a slice-wise Radon transform and a possible change of resolution. For the sake of simplicity we assume that for $c = 1, \dots, C$ the forward operators are identical, *i.e.* $T^c = T$, although the theory is not limited to such restrictions. The regularization parameters μ^c ensure the trade-off between data fidelity and regularization for each channel. The functional TGV_α^2 is given as

$$\text{TGV}_\alpha^2(u) = \min_w \alpha_1 \|\nabla u - w\|_{\text{frob}} + \alpha_0 \|\mathcal{E}w\|_{\text{frob}}, \quad (2)$$

where ∇u denotes the Jacobian of the multi-channel image u , w is a matrix field with the same dimensions as ∇u and $\mathcal{E}w$ denotes a symmetrized Jacobian of w . The norms $\|\cdot\|_{\text{frob}}$ are pointwise Frobenius matrix and tensor norms and $\|\cdot\|_1$ denotes the one-norm, *i.e.*, the sum of moduli over all image voxels. For solving the minimization problem (1), we employ a duality-based algorithm for which we developed a Python³⁸/OpenCL³⁹ implementation. Further details of the algorithm and the reconstruction approach are provided in the section Materials and methods.

† It should be noted that as one channel we denote one type of dataset to be reconstructed, such as the HAADF signal or a single EDXS elemental map, meaning if we consider sinograms for HAADF, EDX aluminum, EDX silicon and EDX ytterbium, then $C = 4$. This should not be confused with spectral channels in EDXS or EELS.

We can identify two main features of the proposed coupled TGV regularization functional: first, the functional itself optimally balances between a penalization of the first and second order derivatives. Choosing the auxiliary variable w to be locally zero or equal to ∇u , it is possible to penalize either the one-norm of ∇u or the one-norm of $\mathcal{E}\nabla u$, where $\mathcal{E}\nabla u$ corresponds to the Hessian of u . For this reason the proposed method can be expected to recover both sharp interfaces and gradual intensity changes. Second, the pointwise matrices and tensors resulting from differentiation of u and w are coupled with a Frobenius norm, *i.e.*, the two-norm over all entries. This enforces joint sparsity of the corresponding quantities; in particular, the edge set of each channel can be expected to have small support and to largely coincide with the ones of the other channels.

Reconstruction of phantom data

To illustrate the capability of the joint TGV algorithm, we first test it in an artificial example. For this purpose we created a multi-channel phantom object. The phantom is inspired by the sample investigated in our experiments and is assumed to consist of three elements (Yb, Si and Al), with both abrupt and gradual concentration changes in the sample. For this phantom, we calculate the sinograms of EDX signals for Yb, Si and Al and of the HAADF signal assuming typical experimental parameters for an analytical tomography experiment and add Poisson noise to the calculated sinograms (see the Materials and methods section). To investigate how a full 3D implementation of the reconstruction algorithm compares to a 2D implementation, we replicate the same slice 60 times rotating each slice slightly compared to the previous to obtain a 3D volume with changes along all three dimensions. Fig. 2 shows the phantom and the sinograms with and without noise. For each slice, sinograms are calculated with a size of 305 pixels over a range of 180° with steps of 5° (see the Materials and methods section for more details of the phantom).

For reconstruction, we compare the standard method SIRT⁴⁹ with TV and TGV regularization, where for the latter two we provide the results for 2D slice-wise and volumetric 3D regularization as well as uncoupled and coupled regularization of the different channels. The SIRT reconstructions were obtained using the Astra toolbox^{50,51} and for the TV and TGV results, we used different variants of our reconstruction algorithm presented in the Materials and methods section. Reconstruction parameters for all methods are provided in ESI Table 1.†

In Fig. 3 we provide the reconstruction results, where three orthogonal slices for each method for both the HAADF and the EDXS signals are compared to the original phantom (Fig. 3a). The SIRT reconstruction (Fig. 3b) of the low-noise HAADF data is quite acceptable, showing only some typical artifacts stemming from the limited number of projections. Due to the low signal of the EDXS data however, these reconstructions are dominated by the Poisson noise of the sinograms and hardly anything can be discerned.



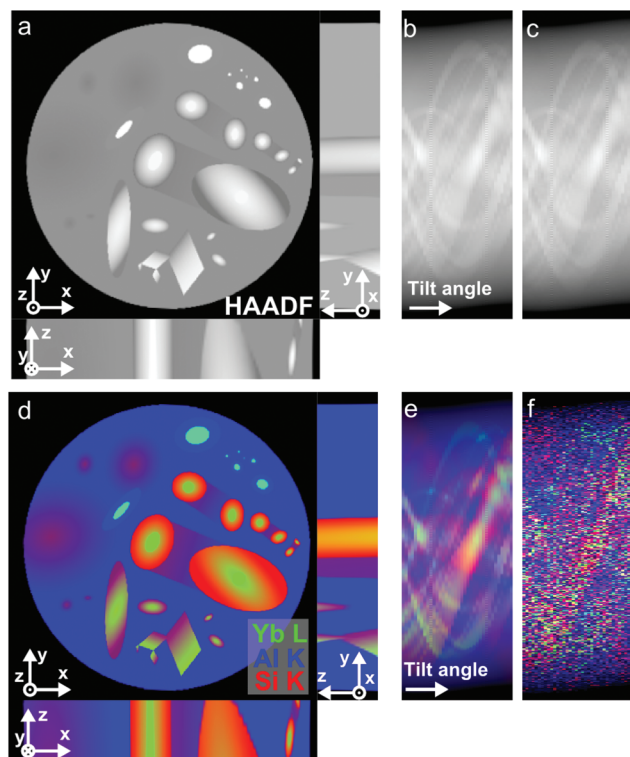


Fig. 2 Multi-channel 3D phantom: (a) HAADF slices along three orthogonal directions, (b) HAADF sinogram without noise and (c) with noise (peak signal-to-noise ratio (PSNR): 55 dB). (d) EDX slices, (e) EDX sinograms without noise and (f) with noise (PSNR for Yb: 7.77 dB, for Si: 9.25 dB, for Al: 18.21 dB). Yb is shown in green, Si in red and Al in blue.

Introducing variational regularization reduces the impact of noise on the reconstruction, but it can be observed that the results strongly depend on the chosen regularizer. In an uncoupled 2D TV reconstruction (Fig. 3c) severe staircasing artifacts (*i.e.*, artificial piecewise constant regions) are present; these artifacts are removed by using the TGV² norm (Fig. 3d), but in both these reconstructions small features in the sample can completely disappear due to the low signal-to-noise ratio of the sinograms.

This situation can be significantly improved by coupling the reconstructions for both TV (Fig. 3e) and TGV² (Fig. 3f). With the transfer of information between the channels the presence and shape of smaller features become clear also in the reconstructions of EDXS data. In the TGV² reconstruction, both sharp and gradual concentration changes can be recovered and also in the TV reconstruction, staircasing artifacts are reduced.

By linking all slices in the third spatial dimension, the 3D implementation of the algorithm is even more robust to noise for both the uncoupled (Fig. 3g & h) and the joint reconstruction (Fig. 3i & j). The 3D joint reconstruction methods make full use of data correlation, along all three spatial dimensions as well as between the different channels and therefore give the best reconstruction for both TV and TGV². With TGV², constant areas and hard transitions are equally well reconstructed as with TV, but, by promoting piecewise linear functions in

addition to sharp interfaces, TGV² is superior to TV when it comes to gradual intensity changes. In the TGV² reconstruction all but some of the very smallest features can be recovered.

These trends in improvement can also be quantified by considering the corresponding peak signal-to-noise ratios between the original phantom images and the reconstructions, as shown in Table 1. We see, for example, that the PSNR of the ytterbium reconstructions strongly benefits from the coupling of channels both in 2D and 3D. Also, all channels greatly benefit from the change from 2D to 3D. Note that, even though visually one can observe qualitative improvements with TGV compared to TV, both methods behave similarly in terms of PSNR. This may be explained using the fact that PSNR is not well suited to quantify such qualitative differences, but also by the fact that the parameters for the reconstruction of Fig. 3 were selected for visually optimal results. Indeed, reducing the amount of regularization slightly can be expected to yield an improvement in terms of PSNR for all methods, but would decrease the visual image quality. Nevertheless, we see that also in terms of PSNR, the improvement obtained with the proposed coupled 3D TGV reconstruction is quite significant compared to 2D uncoupled reconstructions or the SIRT method. In this context, we also refer the reader to the ESI† of this paper, where a mean-to-standard-deviation evaluation of our method on 100 different noise realizations also revealed that a significant improvement was obtained with TGV both for the mean over all reconstructions from different noise realizations and for the standard deviation.

Experimental reconstruction

In an analytical tomography experiment we acquired tilt series from a needle-shaped sample prepared from an Al-5 wt% Si alloy with 50 ppm Na and 6100 ppm Yb. HAADF STEM data and EDXS spectrum images were acquired every 4° with a pixel size of 0.76 nm. HAADF data and elemental maps of Yb, Si and Al at different tilt angles are shown in Fig. 4.

Reconstructions of the HAADF and EDXS data using the SIRT method as well as uncoupled and coupled TGV regularization are shown in Fig. 5, where for the SIRT method we used 100 and 20 iterations for the HAADF and EDXS data, respectively, and we refer the reader to the section Materials and methods for the parameter choice for TGV. Fig. 5a depicts a conventional SIRT reconstruction and Fig. 5b an uncoupled TGV² reconstruction. For both the HAADF and the EDXS data the quality improvements of TGV reconstruction are evident. In the HAADF reconstruction of the shown slice, two bright features are visible, and in the larger one, dark enclosures are present. These are already visible in the SIRT reconstruction, but are much clearer in the TGV² reconstruction. Furthermore, in the TGV² reconstruction, it becomes evident that at the interfaces of the bright features, both abrupt and gradual transitions are present at different locations. For the reconstructions of the EDXS data, TGV² significantly reduces the impact of noise on the reconstruction, but smaller features, such as the dark enclosures of the HAADF image, are not visible in the



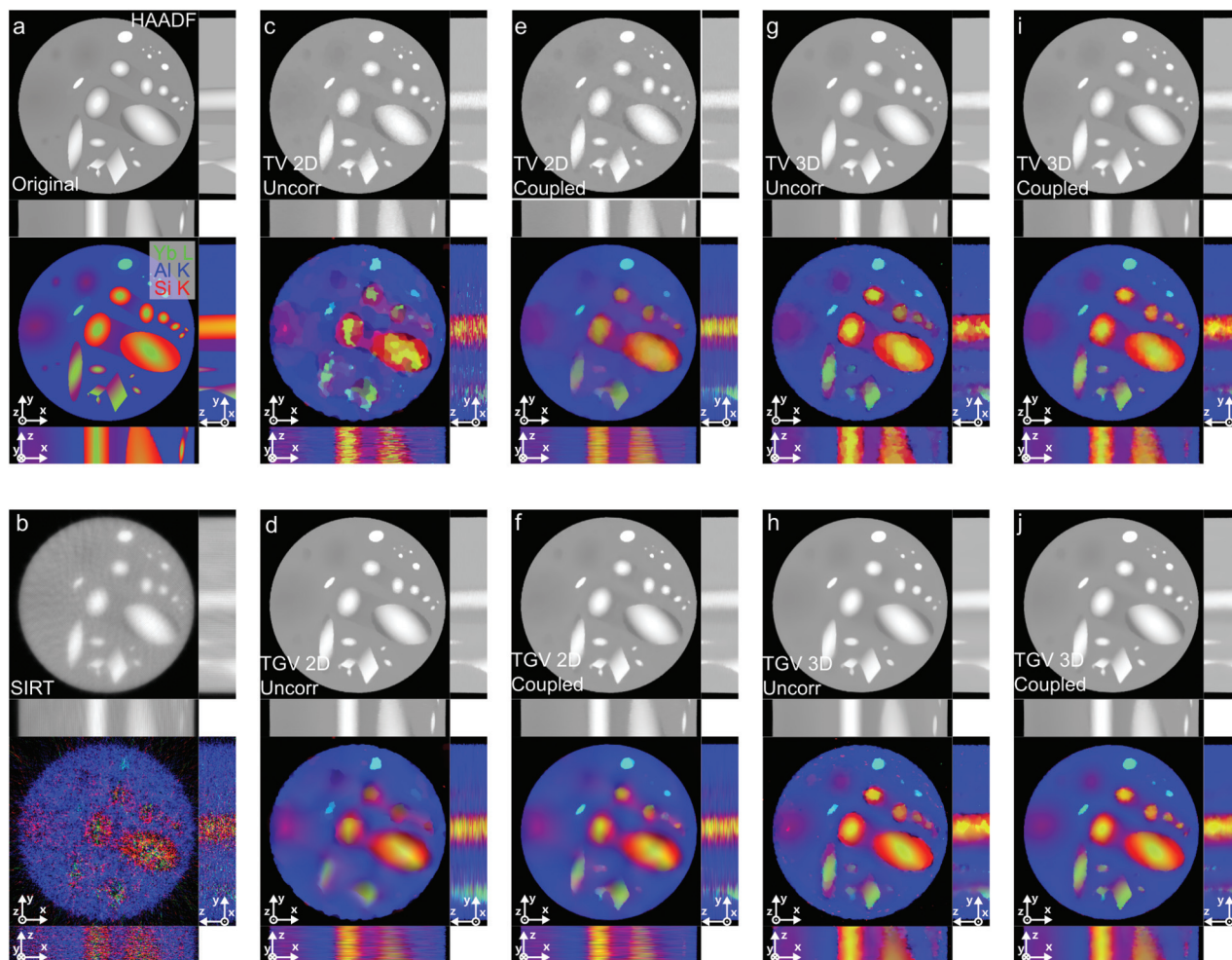


Fig. 3 Reconstructions of a phantom from noisy projection data. In the images, Al is depicted in blue, Si in red and Yb in green. (a) Original phantom, (b) SIRT reconstruction, (c) 2D TV reconstruction without coupling, (d) 2D TGV² reconstruction without coupling, (e) 2D coupled TV reconstruction, (f) 2D coupled TGV² reconstruction, (g) 3D TV reconstruction without coupling, (h) 3D TGV² reconstruction without coupling, (i) 3D coupled TV reconstruction, and (j) 3D coupled TGV² reconstruction.

Table 1 Peak signal-to-noise ratio in dB of the phantom reconstructions depicted in Fig. 3

	SIRT	TV uncorr. 2D	TV coupled 2D	TV uncorr. 3D	TV coupled 3D
HAADF	19.05	34.56	34.59	35.15	35.05
Ytterbium	2.72	6.22	10.14	10.71	13.42
Aluminum	8.26	14.83	15.80	18.42	18.78
Silicon	3.19	9.17	9.94	12.09	13.18
	TGV uncorr. 2D	TGV coupled 2D	TGV uncorr. 3D	TGV coupled 3D	
HAADF	35.10	35.06	35.21	35.23	
Ytterbium	7.92	10.74	11.40	14.09	
Aluminum	14.66	15.25	18.03	18.28	
Silicon	9.13	9.85	10.98	12.44	

EDXS data neither in the SIRT reconstruction nor in the uncoupled TGV² reconstruction. As Yb is by far the heaviest element present in the sample, HAADF contrast is governed by the local Yb concentration, and therefore at the locations of the enclosures, the EDXS signal of Yb should decrease.

This issue can be resolved in the joint TGV² reconstruction depicted in Fig. 5c. Due to the coupling, the reconstructions of the elemental maps show much better defined interfaces and transitions. It is now evident that the dark enclosures visible in the HAADF image correspond to areas with a low concen-



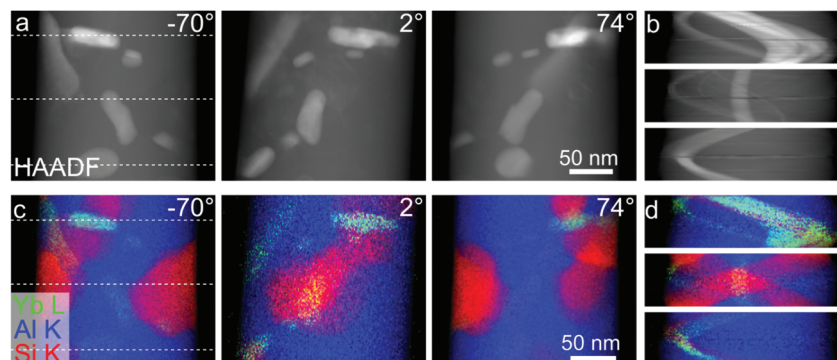


Fig. 4 (a) HAADF STEM projections and (b) sinograms of the HAADF STEM signal at three different heights as indicated by dashed lines in (a). (c) Projections of Yb, Si and Al EDXS signals at different tilt angles and (d) sinograms of the Yb, Si and Al EDXS signals at three different heights as indicated by dashed lines in (c). The individual signals of Yb, Si and Al are shown in the ESI Fig. 2.†

tration of Yb and a high concentration of Si. This is a result of the coupling of the signals, as the transition locations present in the HAADF signals are transferred to the reconstructed elemental maps. In the coupled reconstruction it can also be observed that gradual transitions between different areas are present at some of the interfaces between Yb-rich and Si-rich regions and near the edge of the sample, where beam damage during sample preparation plays a role. In contrast, all interfaces between Yb-rich regions and the Al-matrix are abrupt.

It should be noted that for both the TV and TGV² reconstruction, regularization parameters have been chosen to weight the impact of the norms. A comparison of the effect of different regularization parameters for experimental data is given in Fig. 6. In Fig. 6a the regularization parameter for the HAADF data is varied, while in Fig. 6b the parameter for the HAADF data is kept constant, and the parameters for the EDXS data are changed by the same factor. By visual comparison of different parameters, good parameters can be chosen, which do not lead to a loss of information, while minimizing the impact of noise on the reconstruction. In the section Materials and methods we expand in more detail on how to choose these parameters. Note that also for SIRT and similar methods, a parameter (typically the number of iterations) must be chosen for each channel. The impact of SIRT parameters is depicted in ESI Fig. 3.† In this context, and in contrast to SIRT, one should also note that the number of iterations for our method does not play the role of an additional parameter but rather just needs to be chosen sufficiently high such that convergence is ensured.

Fig. 7 shows several slices through the linked reconstruction of all signals and a volume rendering of the different signals. As the HAADF signal is governed by the local Yb-concentration, a close correlation between these signals can be observed. By linking the reconstructions, the Yb-elemental map becomes well defined and shows almost the same features and transitions as the HAADF reconstruction, where Yb-rich precipitates are visible. These precipitates are also visible in the Si-elemental map, showing a higher Si-concentration

than the matrix and in the Al-elemental maps, where they appear dark, indicating a low Al concentration. In the Si- and Al-elemental maps a second type of precipitate with a high concentration of Si and a moderate concentration of Al is visible.

Discussion

In both simulations and experiments, we could observe that TGV² reconstruction is well suited for reconstructing gradual transitions inside a sample, in addition to sharp interfaces. Thereby it extends TV reconstruction, making sparse reconstruction applicable to a wider range of samples. In the investigated sample such gradual transitions are present between Yb-rich and Si-rich precipitates.

Linking different signals in the reconstruction significantly improves the quality of analytical STEM reconstructions by transferring morphological information present in the HAADF STEM data to the analytical data and in between the analytical datasets.

Since this transfer is a key feature of our approach, we want to quickly illustrate its inner workings. For simplicity, assume that we have only two channels and consequently the reconstruction approach reduces to

$$\min_{(u^1, u^2) \geq 0} \mu^1 D_{\text{KL}}(Tu^1, f^1) + \mu^2 D_{\text{KL}}(Tu^2, f^2) + \text{TGV}_\alpha^2(u),$$

i.e. we look for u with a low corresponding value, improperly referred to as “cheap”. Note that due to the nature of TGV_α^2 it is cheaper for TGV_α^2 to align edges or slopes in the two different channels at common positions. However, this might come at the expense of the reconstruction fitting the data less, making the data terms more expensive. Thus a compromise between fitting and regularity might be ideal, and the concrete behavior therefore depends on various factors.

Let us assume that f^1 contains far less noise than f^2 . Then one would naturally choose μ^1 greater than μ^2 , as then f^1 is more reliable and thus we make deviation from f^1 more expen-



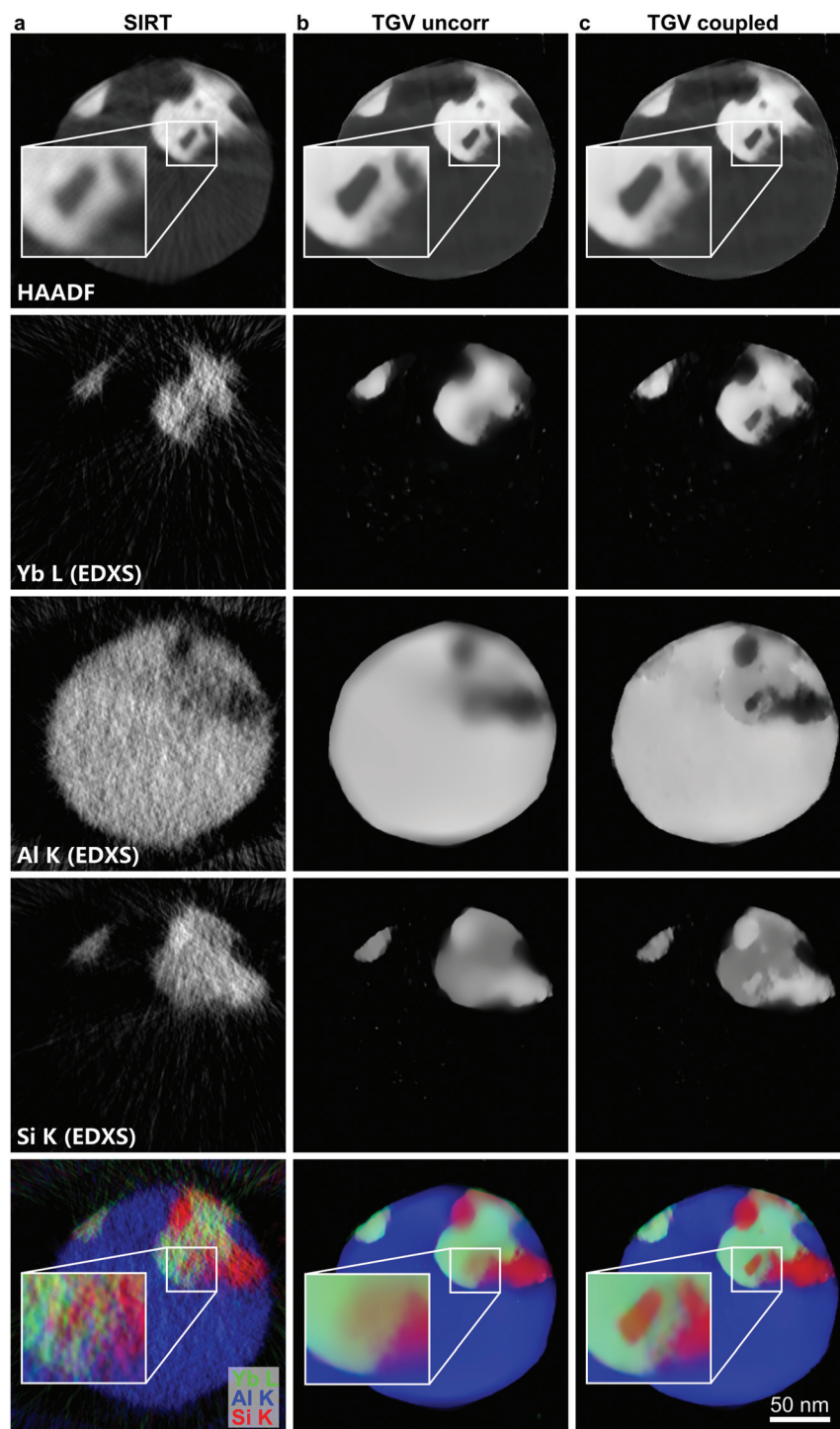


Fig. 5 Reconstruction of HAADF data and Yb, Si and Al elemental maps. (a) Uncorrelated SIRT reconstruction, (b) uncorrelated TGV reconstruction, and (c) joint TGV reconstruction.

sive. Therefore, the approach finds u^1 fitting f^1 well, and rather adjusts u^2 to obtain common features. This is exactly the effect we observed between the HAADF and Yb reconstructions in the experimental data, where Yb was strongly adjusted to HAADF, but HAADF remained unchanged and in particular was not negatively impacted.

On the other hand, if both channels have the same noise level, and consequently the weights are chosen to be of similar magnitude, deviation from data is equally expensive for both channels, leading to a more balanced compromise in adjusting the reconstructions to obtain common features. This, for example, happens in the reconstruction of the experimental



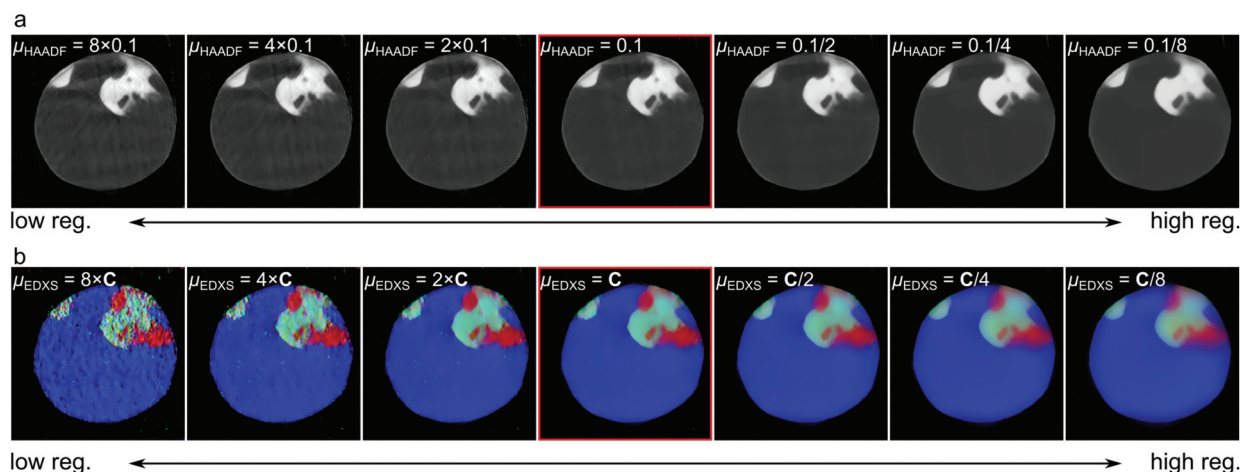


Fig. 6 Comparison of different regularization parameters for TGV reconstructions of the experimental data: (a) for the HAADF data and (b) for the EDXS data. In (b), $\mu_{\text{HAADF}} = 0.1$ for all reconstructions and for the central image $C = (\mu_{\text{Al}}, \mu_{\text{Si}}, \mu_{\text{Yb}}) = (0.0024, 0.0014, 0.001)$.

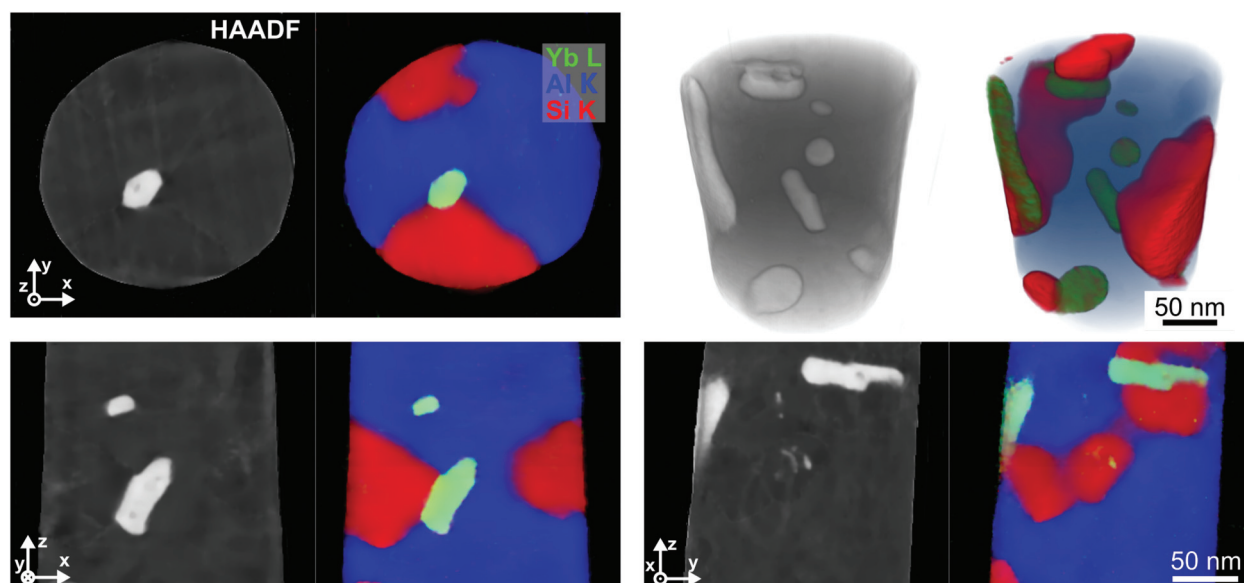


Fig. 7 Different slices through correlated reconstructions of the HAADF signal, Yb-, Si- and Al-elemental maps and 3D volume rendering.

data at the edges between Al and Si, leading to better defined edges for both channels.

Of course, if attaining common features comes at too high a cost, in particular if there actually are no common features in reality, the algorithm will not force common features. So no adjustment appears where it makes no sense to adjust.

In its current implementation, the algorithm will work well for all types of samples which fully remain in the field of view at all tilt angles and is robust to a limited tilt range, *cf.* ESI Fig. 4.† As such, the algorithm is also readily applicable to samples on a thin-film support, for example, nanoparticles. Only if the sample extends beyond the field of view, problems

may arise, similar to other reconstruction algorithms such as SIRT.

The algorithm can be readily applied to any type of signal fulfilling the projection requirement; in particular, its application to EELS data or to combined EDXS/EELS datasets is straightforward.

For the material, the observed gradual transition at the interface between Yb-rich and Si-rich precipitates indicates that a side by side nucleation of the Yb and eutectic Si particles might happen. The diffusion interface between Yb and Si leads to stoichiometry changes (*i.e.* gradual transitions) of both phases depending on the solute and/or solid diffusion of Yb atoms.



Conclusion

Without the TV-induced bias towards piecewise constant solutions, TGV² reconstruction achieves high-quality results – and is therefore applicable – for a wide range of materials and will help in understanding their morphology, composition and functionality. The presented linkage of different signals is an ideal approach for obtaining most information out of the available data and can be used for any type of signal used for tomographic reconstruction in a TEM, but the approach is not limited to electron tomography, but can be applied to every type of projection-based tomography. In analytical electron tomography, optimized usage of the available data is mandatory, as the quality and quantity of data which can be acquired in an analytical tomography experiment are severely limited by potential beam damage and acquisition time.

Materials and methods

Numerical simulations

In this chapter we give a more detailed description of the phantom data shown in the section Reconstruction of phantom data. To demonstrate the functionality of the joint TGV reconstruction algorithm, we created a multi-channel phantom object. This phantom consists of three channels, which we attribute to the atomic fraction c_A of Yb, Al and Si, and has an original size of 2048×2048 pixels, with the pixel size being set to 0.1 nm. To obtain a realistic estimation of the number of detector counts for both the EDXS signal and the HAADF signal, we calculated local signal creation probabilities.

For the EDXS signal, we used

$$I_A = \frac{\rho \cdot t \cdot m_A \cdot D_e}{\zeta_A} \quad (3)$$

with the local density ρ , the sample thickness t , the local concentration of the element A m_A (mass fraction), the electron dose D_e and the zeta-factor for the element A ζ_A .⁵² To calculate the local density, we weighted the elemental densities of Yb ($\rho_{Yb} = 6.9 \text{ g cm}^{-3}$), Al ($\rho_{Al} = 2.7 \text{ g cm}^{-3}$) and Si ($\rho_{Si} = 2.3 \text{ g cm}^{-3}$) by their local atomic fraction. The thickness t equals the phantom pixel size of 0.1 nm.

The dose D_e was calculated for an electron current of $I_e = 2 \text{ nA}$, a pixel dwell time of $t_{px} = 2 \text{ ms}$ and the elemental charge of $q = 1.602 \times 10^{-19} \text{ C}$ using

$$D_e = \frac{I_e}{t_{px} \cdot q} \quad (4)$$

For the zeta-factors, we used $\zeta_{Si} = \frac{500}{4} \text{ kg m}^{-2}$ for Si, $\zeta_{Al} = \frac{500}{4} \text{ kg m}^{-2}$ for Al and $\zeta_{Yb} = \frac{240}{4} \text{ kg m}^{-2}$ for Yb. Zeta-factors for Si and Al were measured for our microscope; the zeta-factor for Yb is an estimation based on theoretical k -factors for the investigated elements.

To obtain an estimate of the HAADF signal, we used tabulated values of the differential cross section $\frac{\partial \sigma}{\partial \Omega}$ for elastic scattering from the NIST database,⁵³ integrating them over the angular detector range $[\theta_{min}, \theta_{max}]$ with^{54,55}

$$\sigma_{[\theta_{min}, \theta_{max}]} = 2\pi \int_{\theta_{min}}^{\theta_{max}} \frac{\partial \sigma}{\partial \Omega} \sin \theta d\theta. \quad (5)$$

For the inner and outer detector angles, we used $\theta_{min} = 94.9 \text{ mrad}$ and $\theta_{max} = 188 \text{ mrad}$, which was measured in our system for the Fischione HAADF detector at a camera length of 58 mm.

For each element A, the scattering factor f_A was calculated as

$$f_A = e^{-\frac{N_A}{m_A} c_A \cdot \sigma_A \cdot \rho t} \quad (6)$$

with the Avogadro constant $N_A = 6.022 \times 10^{23} \text{ mol}^{-1}$, the atomic mass m_A , the local atomic fraction c_A and the scattering cross section σ_A for the element A, as well as the local density ρ and the pixel size t .

The total HAADF signal was calculated using the scattering factors for all elements and the electron dose D_e according to

$$I_{HAADF} = D_e (1 - f_{Al} \cdot f_{Si} \cdot f_{Yb}). \quad (7)$$

The electron dose is calculated from eqn (4) with $I_e = 2 \text{ nA}$ and a dwell time of $t_{px} = 2 \text{ ms}$. To account for detector efficiency, we reduce the HAADF signal by a factor of 0.7.

Finally, the phantom data are downsampled from 2048×2048 pixels to 305×305 pixels using bilinear interpolation, corresponding to a pixel size of 0.67 nm, close to the experimental pixel size. After downscaling, the pixel values are multiplied by a factor of 0.67 to account for the larger thickness of each pixel.

Electron tomography experiments

In this section, we give a more detailed description of the experimental data used in the Experimental reconstruction section. The investigated sample stems from an Al-5 wt% Si alloy with 50 ppm Na and 6100 ppm Yb; a similar sample has been previously analyzed by analytical electron tomography.¹⁴ From this material a needle-shaped sample containing Yb- and Si-rich precipitates is prepared using a dual-beam focused ion beam/scanning electron microscope (FIB/SEM) instrument. STEM images are acquired on a probe-corrected FEI Titan³ G2 60-300 microscope operated at 300 kV and equipped with a FEI Super-X four quadrant silicon drift detector (SDD) EDXS system. EDXS spectrum images were acquired every 4° between -74° and $+78^\circ$ at a pixel size 0.76 nm with 276×296 pixels. The pixel time was 2 ms, leading to a frame time of 4 min 36 s. The tilt range was limited by the sample holder available at the time of the experiment (a Fischione 2020 Advanced Tomography Holder). Elemental maps of the Al and Si K-lines and of the Yb L-lines are extracted from the EDXS spectrum images. For this purpose, the background is fitted with a Kramers model⁵⁶ and Gaussian peaks are fitted at all peak positions.



Data preprocessing

For the variational reconstruction to work efficiently, the raw sinogram data f need to be consistent with the model. Therefore, the data require suitable preprocessing, and failing to do so could result in artifacts in the reconstructed images. In the ESI† we also discuss the effect of model inconsistencies, which may arise for HAADF STEM data.

To correct for sample drift during spectrum image acquisition, the spectrum image is corrected using a linear drift model (constant speed and direction). Drift parameters are calculated by minimizing the difference between the HAADF data acquired during SI (Spectrum Image) acquisition and the fast scan image acquired before. The calculated parameters are used to correct all SI data.⁵⁷ Alignment of the tilt series is done using the HAADF STEM data. For alignment in the direction of the tilt axis, common-line alignment is applied,⁵⁸ in the direction perpendicular to the tilt axis the center of mass of each projection is used for alignment and for tilt axis alignment, rotational centers are calculated based on the center of mass.⁵⁹ The alignment parameters calculated for the HAADF STEM data are applied to all datasets. Drift correction and alignment are done using MATLAB.

Further pre-processing steps can be done using the Python software presented here. Intensity fluctuations can appear between different tilt angles. There are two main sources for these effects: partial shadowing of the detector and residual diffraction contrast. To reduce intensity variations between different projections, we rescale all projections to be non-negative and to have a common mean. This also compensates for partial shadowing of the EDXS detectors. Additionally, projections with very strong diffraction contrast, containing unsatisfactory information, are removed completely.

To correct for non-zero signal values in background regions with no mass (e.g. due to Gaussian noise stemming from the electronics), the corresponding pixels are masked *via* thresholding and their values are set to zero. A detailed investigation of the impact of data inconsistencies is provided in the ESI.†

Variational modeling

In this section we provide details of the proposed variational model for electron tomography reconstruction, which comprises the solution of (1). For an easier understanding, we describe the reconstruction of a single signal first.

Forward model. In our electron tomography measurement, a focused electron beam is directed at the sample orthogonal to a fixed imaging plane, and raster scanned over the region of interest in the sample. For the HAADF signal, the number of electrons scattered to a specific annular range is measured, while for EDXS maps, characteristic X-rays for specific elements are measured for each position of the electron beam. For each beam position, this corresponds to measuring (at least approximately) the density of either a single element (EDXS) or a weighted sum of all elements (HAADF) integrated along the line of the electron beam that intersects the sample orthogonally to the imaging plane. Then, the imaging

plane is rotated and the experiment is repeated for several tilt angles.

Mathematically, this setup can be described as follows: given an angle $\theta \in S^1$ (the unit sphere in \mathbb{R}^2), the imaging plane is defined as $P_\theta = \{(x, y, z) \in \mathbb{R}^3 | (x, y) \cdot \theta = 0\}$ and, for given parameters s and z describing the horizontal and vertical offset, respectively, the measurement line $L_{s,\theta,z}$ orthogonal to this imaging plane is given as $L_{s,\theta,z} = \{(x', y', z') \in \mathbb{R}^3 | (x', y') \cdot \theta^\perp = s, z' = z\}$, where $\theta^\perp = (\theta_1, \theta_2)^\perp = (\theta_2, -\theta_1)$ is a clockwise rotation of θ by 90° . The measurement $Tu(s, \theta, z)$ for the parameters (s, θ, z) of a density described by $u: \mathbb{R}^3 \rightarrow \mathbb{R}$ is then given as the integration of u over the measurement line $L_{s,\theta,z}$, i.e.,

$$Tu(s, \theta, z) := \int_{L_{s,\theta,z}} u.$$

For any fixed height z , this corresponds to the 2D Radon transform⁶⁰ of the density u in the plane $P_z = \{(x, y, z) \in \mathbb{R}^3 | x, y \in \mathbb{R}\}$. Thus, the measurement operator T corresponds to a slice-wise Radon transform where the image space is parameterized by the height z , the normal of the imaging plane θ and the position in the plane s . The tomographic reconstruction problem then corresponds to solving the equation

$$Tu = f$$

for given sinogram data $f = f(s, \theta, z)$. In a practical realization, both the measured data and the unknown density are defined on a discrete grid and we employ a discrete Radon transform that is based on a weighted sum of points close to the line of integration as follows: for discrete position parameters (s, θ, z) , we define the Radon transform of a discrete density u as

$$Tu(s, \theta, z) = \sum_{(x,y): |(x,y) \cdot \theta^\perp - s| < \Delta s} \left(1 - \frac{|(x,y) \cdot \theta^\perp - s|}{\Delta s} \right) u(x, y, z), \quad (8)$$

where the sum is taken over all positions (x, y) in the discrete pixel grid. We note that, naturally, this is not the only possible discretization,⁶¹ but we chose this one since it comprises a suitable balance between approximation accuracy and computational effort. In particular, it allows a very efficient computation of the numerically exact adjoint operation (backward projection) which will be required in the optimization algorithm. This is of particular importance, as a non-exact adjoint can significantly hinder optimization both in convergence speed and in accuracy; see the ESI.†

Noise model. For both the HAADF and the EDXS signal, the main contribution to noise, especially at low count rates, comes from the counting of electrons or X-rays at the detector, which follows Poisson statistics.^{62,63} Additionally, the signal can also be affected by Gaussian noise stemming from the electronics. Omitting Gaussian noise (see also the section Data



preprocessing for related preprocessing steps), the noise on the data is typically assumed to be Poisson distributed.

Hence, from a Bayesian perspective, the appropriate data discrepancy term in a variational model is the Kullback-Leibler divergence,⁶⁴ which is for non-negative densities \tilde{u}, f (up to constants) given as

$$D_{\text{KL}}^f(\tilde{u}) = D_{\text{KL}}(\tilde{u}, f) = \int \tilde{u} - f \log(\tilde{u}), \quad (9)$$

where we use the convention $0 \log(0) = 0$ and $-a \log(0) = \infty$ for $a > 0$. For discrete measurements, the integral is replaced by a sum.

Regularization. In order to achieve a regularized inversion of the Radon transform corresponding to the forward operator of an electron tomography experiment, we incorporate the Total Generalized Variation (TGV)^{32,33} as a regularization term. The TGV functional realizes an ℓ^1 -type penalization of higher order derivatives *via* a cascadic decomposition of the unknown. Such a procedure preserves the advantage of the popular Total Variation (TV) functional²⁵ of allowing reconstruction of sharp interfaces while overcoming its defects, particularly its tendency to foster piecewise constant density distributions (the *staircasing effect*). TGV can be defined for an arbitrary order of differentiation $k \in \mathbb{N}$ and for $k = 1$ coincides with the TV functional. In this work, we employ the second order TGV functional for regularization ($k = 2$), which, for a single-channel image (or volume) u and parameters $\alpha_0, \alpha_1 \in (0, \infty)$, is given as

$$\text{TGV}_\alpha^2(u) = \min_w \alpha_1 \|\nabla u - w\|_1 + \alpha_0 \|\mathcal{E}w\|_{\text{frob}}, \quad (10)$$

where w is a vector field and $\mathcal{E}w = \frac{1}{2}(\nabla w + (\nabla w)^T)$ denotes the symmetrized Jacobian matrix. The pointwise norms $|\cdot|$ and $|\cdot|_{\text{frob}}$ are the standard Euclidean vector norm and the Frobenius matrix norm, *i.e.*, the root of the sum of squares of all entries of the vector or matrix, respectively.

Thus, TGV_α^2 minimizes the norm of the gradient while allowing subtraction of a function w , but penalizing this *via* the norm of the symmetrized Jacobian of w . In particular, in areas with constant u , w can be chosen equal to 0, and in areas with constant ∇u , meaning u is linear, w can be chosen equal to ∇u , such that those areas do not contribute to the TGV_α^2 functional. Therefore, in reconstructions, TGV_α^2 promotes piecewise linear solutions, which allow sharp edges to remain intact while allowing linear density gradients, hence maintaining the advantages of TV while overcoming its defects.

A discretization of TGV_α^2 is defined according to (10) with the discrete ℓ^1 norm, the pointwise Frobenius norms of a vector and a matrix given in the standard way, and discretized differential operators. Denoting by $U = \mathbb{R}^{N_1 \times N_2 \times N_3}$ the space discrete volumetric densities, the discretized gradient $\nabla: U \rightarrow U^3$ and symmetrized Jacobian $\mathcal{E}: U^3 \rightarrow U^6$ are defined for $u \in U$ and $w \in U^3$, respectively, as $\nabla u = ((\nabla u)^1, (\nabla u)^2, (\nabla u)^3)$, where

$$(\nabla u)^1 = \delta_{1+}u, \quad (\nabla u)^2 = \delta_{2+}u, \quad (\nabla u)^3 = \delta_{3+}u$$

and as $\mathcal{E}w = ((\mathcal{E}w)^1, (\mathcal{E}w)^2, (\mathcal{E}w)^3, (\mathcal{E}w)^4, (\mathcal{E}w)^5, (\mathcal{E}w)^6)$ where

$$\begin{aligned} (\mathcal{E}w)^1 &= \delta_{1-}w^1, & (\mathcal{E}w)^2 &= \delta_{2-}w^2, & (\mathcal{E}w)^3 &= \delta_{3-}w^3, \\ (\mathcal{E}w)^4 &= \frac{\delta_{1-}w^2 + \delta_{2-}w^1}{2}, \\ (\mathcal{E}w)^5 &= \frac{\delta_{1-}w^3 + \delta_{3-}w^1}{2}, \\ (\mathcal{E}w)^6 &= \frac{\delta_{2-}w^3 + \delta_{3-}w^2}{2}, \end{aligned}$$

and

$$\delta_{1+}, \delta_{2+}, \delta_{3+}: U \rightarrow U \text{ and } \delta_{1-}, \delta_{2-}, \delta_{3-}: U \rightarrow U$$

are the standard forward and backward finite difference operators with respect to the 3 space dimensions, using pixel replication at the boundary. Note that for $\mathcal{E}w$ we store each off-diagonal entry only once and, to account for that, count off-diagonal entries twice in the evaluation of the Frobenius norm of $\mathcal{E}w$.

Discrete single-channel model. With the TGV functional in (10), the forward operator in (8) and the discrepancy term in (9), we formulate the reconstruction of a single channel volumetric image as the following convex minimization problem:

$$\min_{u \geq 0} \mu D_{\text{KL}}(Tu, f) + \text{TGV}_\alpha^2(u). \quad (11)$$

Problem (11) balances the discrepancy between the Radon transform Tu and the data f with the $\text{TGV}_\alpha^2(u)$ regularization, weighted by the regularization parameter $\mu > 0$.

Discrete multi-channel model. In order to jointly reconstruct density information and elemental maps and to take advantage of complementary information in these datasets, we propose a joint TGV regularization method for multi-channel data. To achieve this aim, we define the extension of TGV for multi-channel data, as introduced in previous studies,^{34,65} by regarding multiple images with a common resolution as one multi-channel image $u = (u^1, u^2, \dots, u^C)$ and setting

$$\text{TGV}_\alpha^2(u) = \min_w \alpha_1 \|\nabla u - w\|_{\text{frob}} + \alpha_0 \|\mathcal{E}w\|_{\text{frob}}. \quad (12)$$

Note that here, the operators ∇, \mathcal{E} are applied channel-wise, hence $\nabla u - w$ is a matrix field, $\mathcal{E}w$ is a tensor field and we use pointwise Frobenius norms for both of them. This corresponds to an ℓ^1/ℓ^2 -type penalization and promotes joint sparsity of both $\nabla u - w$ and $\mathcal{E}w$, and in particular promotes joint sparsity of the edge sets of all channels. We note that in other studies,⁴⁶ a similar regularization has been proposed for the joint reconstruction of MR and PET data, where a nuclear-norm based coupling of the first order derivatives was carried out. While the authors reported a slight improvement by using the nuclear norm, the performance was comparable to a Frobenius-norm based penalization and for the sake of simplicity, we employ the latter.



With this multi-channel extension of TGV, we propose the following joint reconstruction model for multi-channel data as an extension of (11):

$$\min_{\substack{u = (u^1, \dots, u^C) \\ u^c \geq 0 \ \forall c = 1, \dots, C}} \left[\sum_{c=1}^C \mu^c D_{\text{KL}}(T^c u^c, f^c) \right] + \text{TGV}_\alpha^2(u). \quad (13)$$

Here, different data discrepancies are used for the different measured data sets denoted by f^c and each μ^c weights data fidelity and regularization for the c -th channel. Solving (13) balances the joint regularity of all channels imposed by TGV_α^2 with the individual channel discrepancies, thus promoting joint edge sets while jointly reconstructing the individual channels. The operators T^c model the measurement operation for each channel and comprise the Radon transform as described above. In this context, we also note that in our experiments all data sets were acquired and reconstructed with the same resolution. Since acquisition of HAADF data is typically much faster than EDXS, an alternative would be to acquire the HAADF signal with a higher resolution and reconstruct all channels at this high resolution. This can be realized in our model by including an additional subsampling operator in the EDXS channels after the Radon transform and would allow a joint reconstruction and zooming of EDXS data.

It is worth mentioning that our method can, of course, also be used to include more information such as EELS data, from which one can expect a further improvement due to additional information present in the data. This will be highly beneficial in cases where elemental maps from EELS data have a better signal-to-noise ratio, or if, for example, valence states are mapped by EELS. We note, however, that assuming the forward operators for EDXS and EELS signals to coincide, such a coupled reconstruction is equivalent to first taking the (weighted) average of the EDXS and EELS signal for each material and then carrying out a reconstruction using the averaged data for the different materials together with the HAADF data.

Numerical solution

In this section we present the derivation and implementation of a solution algorithm for (13), of which the single-channel setting (11) is a special case. For further reference, we note that Graptor – our Python/OpenCL^{38,39} implementation of the algorithm – as well as a script reproducing all experiments of the paper is available online.^{37,68} Due to the non-differentiability of the objective functional, we employ a first order primal-dual algorithm⁶⁶ to solve (13). In order to do so, the problem needs to be reformulated as a saddle-point problem to which the primal-dual algorithm is applicable. To achieve this aim, we first note that TGV_α^2 can be re-written as

$$\text{TGV}_\alpha^2(u) = \min_w R(\nabla u - w, \mathcal{E}w),$$

with $R(\varphi, \psi) = \alpha_1 \|\varphi\|_{\text{frob}} + \alpha_0 \|\psi\|_{\text{frob}}$. Also, the constraint $u^c \geq 0$ for all c can be replaced by adding the functional $\mathcal{I}_D(u)$ to the energy in (13), where $\mathcal{D} = \{u = (u^1, \dots, u^C) \mid u^c \geq$

0 for all $c\}$ and for any set M we denote $\mathcal{I}_M(u) = 0$ if $u \in M$ and $\mathcal{I}_M(u) = \infty$ else. Denoting by g^* the convex conjugate of a function g , defined as

$$g^*(v) = \sup_{\xi} (v, \xi) - g(\xi),$$

with (v, ξ) being the scalar product, *i.e.*, the sum of the point-wise products of all entries of v and ξ , we can then rewrite (13) as

$$\begin{aligned} & \min_{u=(u^1, \dots, u^C)} \sum_{c=1}^C \mu^c D_{\text{KL}}(Tu^c, f^c) + \text{TGV}_\alpha^2(u) + \mathcal{I}_D(u) \\ & \Leftrightarrow \min_{u=(u^1, \dots, u^C)} \min_w \sum_{c=1}^C \mu^c D_{\text{KL}}(Tu^c, f^c) \\ & \quad + R(\nabla u - w, \mathcal{E}w) + \mathcal{I}_D(u) \\ & \Leftrightarrow \min_{x=(u, w)} \max_{z=(p, q, r)} (Kx, z) - R^*(p, q) \\ & \quad + \mathcal{I}_D(u) - \sum_{c=1}^C (\mu^c D_{\text{KL}}^c)^*(r^c). \end{aligned} \quad (14)$$

Here, the variables $p \in U^{C \times 3}$, $q \in U^{C \times 6}$ are the dual variables corresponding to TGV_α^2 , $r = (r^1, \dots, r^C) \in V^C$ is a dual variable corresponding to the data terms, and

$$Kx = K(u, w) = (\nabla u - w, \mathcal{E}w, Tu^1, \dots, Tu^C)^T. \quad (15)$$

Hence, problem (13) is equivalent to

$$\min_{x=(u, w)} \max_{z=(p, q, r)} (Kx, z) - F^*(z) + H(x), \quad (16)$$

where F^* summarizes the conjugate functionals in (14) and $H(x) = H(u, w) = \mathcal{I}_D(u)$. With the notion of proximal mappings, *i.e.*,

$$\text{prox}_{g, \gamma}(z) = \arg \min_{\xi} g(\xi) + \frac{\|\xi - z\|_2^2}{2\gamma},$$

the iteration steps of the primal-dual algorithm⁶⁶ employed for the saddle-point problem (16) for (x, \tilde{x}, z) are

$$\begin{cases} z_+ & \leftarrow \text{prox}_{F^*, \sigma}(z + \sigma K \tilde{x}) \\ x_+ & \leftarrow \text{prox}_{H, \tau}(x - \tau K^* z_+) \\ \tilde{x} & \leftarrow 2x_+ - x \\ (x, z) & \leftarrow (x_+, z_+). \end{cases} \quad (17)$$

Here, K^* denotes the adjoint operator of K and, as a direct computation shows, the proximal mapping of F^* decouples component-wise and can be given as

$$\text{prox}_{F^*, \sigma}(p, q, r) = \left(\text{proj}_{\alpha_1}(p), \text{proj}_{\alpha_0}(q), \left(\text{prox}_{(\mu^c D_{\text{KL}}^c)^*, \sigma}(r^c) \right)_{c=1}^C \right),$$

where for any $\nu > 0$ and any discrete spatial coordinate \mathbf{x} ,

$$(\text{proj}_\nu(r))(\mathbf{x}) = r(\mathbf{x}) / \max(1, |r(\mathbf{x})|_{\text{frob}}/\nu),$$



which corresponds to a projection to the set $\{r \mid \|r(x)_{\text{frob}}\| \leq \nu \text{ for all } x\}$. The proximal mapping of the data terms is given as

$$\text{prox}_{(\mu^c D_{\text{KL}}^{f^c})^*, \sigma}(r^c) = r^c - \frac{r^c - \mu^c + \sqrt{(r^c - \mu^c)^2 + 4\sigma\mu^c f^c}}{2},$$

while the proximal mapping of H corresponds again to a projection and is given as

$$\begin{aligned} \text{prox}_{H, \tau}(x) &= \text{prox}_{H, \tau}(u, w) = (\text{proj}_{\mathcal{D}}(u), w) \\ &\text{with } \text{proj}_{\mathcal{D}}(u) = \max(u, 0), \end{aligned}$$

where the maximum is taken point- and component-wise. We also refer the reader to a related study⁴⁶ for further details of these derivations. For the sake of simplicity, we use the notation

$$T^*r = (T^*r^c)_{c=1}^C$$

where T^*r^c denotes the backprojection of r^c .

In summary, we can see that, due to our particular reformulation, all iteration steps of Algorithm (17) can be computed explicitly and fast. Furthermore, convergence of the iterate x to a global optimum of (13) can be guaranteed.⁶⁶ A more explicit formulation of the algorithm can be found in Algorithm 1 below.

Algorithm 1 TGV regularized ET reconstruction

```

1: function TGV-ET RECONSTRUCTION ( $f, \alpha, \mu$ )
2:    $(T, f) \leftarrow \text{normalize}(T, f)$ 
3:    $u \leftarrow T^*f, \tilde{u} \leftarrow u, w \leftarrow 0, \tilde{w} \leftarrow 0, p \leftarrow 0, q \leftarrow 0, r \leftarrow 0$ 
4:   Choose  $\sigma > 0$  and  $\tau > 0$  such that  $\sigma\tau\|K\|^2 < 1$ 
5:   repeat
6:      $p \leftarrow \text{proj}_{\alpha_0}(p + \sigma(\nabla\tilde{u} - \tilde{w}))$ 
7:      $q \leftarrow \text{proj}_{\alpha_0}(q + \sigma\mathcal{E}\tilde{w})$ 
8:      $r^c \leftarrow \text{prox}_{(\mu^c D_{\text{KL}}^{f^c})^*, \sigma}(r^c + \sigma(T\tilde{u})^c)$ , for  $c = 1, \dots, C$ 
9:      $u_+ \leftarrow \text{proj}_{\mathcal{D}}(\tilde{u} - \frac{\sigma}{\tau}(\nabla^*p + T^*r))$ 
10:     $w_+ \leftarrow w - \tau(-p + \mathcal{E}^*q)$ 
11:     $(\tilde{u}, \tilde{w}) \leftarrow 2(u_+, w_+) - (u, w)$ 
12:     $(u, w) \leftarrow (u_+, w_+)$ 
13:  until Stopping criterion fulfilled
14:  return renormalize( $u$ )

```

There, the normalization of the operator T and the corresponding data f in line 2 aims at obtaining an operator with a norm of approximately 1 and data on similar scales, as we found this to improve convergence speed. In order to do so, we estimate the operator norm of T by power iteration,⁶⁷ divide T by this estimate and rescale the data f^c of all channels to the range $[0, 1]$. Note that, by the nature of the Kullback–Leibler discrepancy, these rescaling operations only affect the choice of the regularization parameter but not the solution set and are reverted with the renormalization step at the end of the algorithm. With rescaled T , the norm $\|K\|$ can be estimated from above *via* straightforward computations by $\sqrt{17}$, allowing σ and τ to be chosen such that $\sigma\tau < 1/17$. Although within this constraint, the choice of $\sigma\tau$ is arbitrary in principle and, in fact, can influence the convergence speed of the algorithm significantly,⁴⁴ we experienced good convergence behavior with

the straightforward choice $\sigma = \tau = \frac{1}{\sqrt{17}}$, which was used for all experiments.

As the stopping criterion for Algorithm 1, we used a rather high, predetermined number of 5000 iterations for the experimental and 2000 iterations for the phantom reconstructions such that no relevant change in the iterates was observed beyond this point. A more elaborate stopping criterion that, in some cases, allows to compute the difference of the value of the objective functional to the optimal value could be based on the primal-dual gap of the problem.⁴⁴ However, this would require some additional computations during the algorithm and for the sake of simplicity and speed, we hence used a fixed number of iterations.

In order to further accelerate the computations, we developed an implementation in Python³⁸ *via* OpenCL³⁹ for multi-core CPUs and Graphical Processing Units (GPUs), which allows for highly parallel execution of all major computations. This is of particular relevance for the forward and backward projections, which are major contributors to the computational effort needed for Algorithm 1 and can be executed completely in parallel, leading to significant speedups.

Tables 2 and 3 illustrate the time required for reconstruction on a computer with an Nvidia(R) GeForce(R) GTX 980 graphic card and an Intel(R) Xeon(R) CPU E5-2690 processor in order to show the computational effort.

Regarding the memory requirement, we note that our method in 3D essentially requires to hold, for each channel, 13 variables of image size and two variables of the data size in memory (ignoring temporary storage for computations). In comparison, TV regularization using the same algorithm essentially requires a memory that is 4 times the image size and two times the data size.

Table 2 Reconstruction time in seconds with Algorithm 1 as a function of the number of slices, channels and iterations. Reconstruction of slices with $N \times N$ pixels from $N \times N_\phi$ dimensional sinogram data, where $N_\phi = 39$ denotes the number of projections and the number of detectors is $N = 296$

Number of slices	10			30		
Iterations	500	2000	5000	500	2000	5000
1 Channel	3.03	14.79	86.93	8.13	31.84	109.19
4 Channels	10.98	43.27	126.87	46.51	145.08	346.17

Table 3 Reconstruction time in seconds with Algorithm 1 as a function of the size of data with a single channel and 10 slices. Reconstruction of slices with $N \times N$ pixels from the $N \times N_\phi$ dimensional sinogram

Iterations	500	2000	5000
$N_\phi = 39, N = 296$	3.03	14.79	86.93
$N_\phi = 39, N = 872$	22.92	93.51	235.90
$N_\phi = 155, N = 872$	59.80	261.65	660.78



Parameter choice. Here we comment on our parameter choices for the presented experiments and provide a guideline on how to choose parameters in practice. For the TGV parameters $\alpha = (\alpha_0, \alpha_1)$, we note that only the ratio α_0/α_1 is relevant and balances between the first and second order smoothness. Here, the choice $\alpha = (4, 1)$ is known to be rather robust and to yield reasonable results; hence it can be left unchanged for different applications of our method (as was done for all experiments of the paper).

The choice of the regularization weights $\mu = (\mu^1, \dots, \mu^C)$ is more crucial and depends for instance on the noise level of the different measurements, the image resolution and the number of measured angles. Generally, decreasing those parameters leads to increasingly regular and smoothed reconstructions but reduces details, while increasing them provides more details but also increases noise. Nevertheless, as can be observed in Fig. 6, our method is rather robust against suboptimal choices of μ and hence a rough tuning of those parameters should suffice to achieve good results in practice.

A general guideline for choosing those parameters in multispectral electron tomography applications is given as follows, where the measurement setup of the paper is used as example. Since the HAADF density data usually contain far less noise than EDXS data, one can first choose the parameter μ^c corresponding to the HAADF signal, either using our method to first reconstruct only the HAADF signal separately or using some rough, fixed guess for the other parameters.

For the choice of μ^c for the EDXS channels, remember that these values should roughly correspond to the (unknown) noise levels present in each channel and that channels with similar amounts of noise, e.g., with similar electron counts, should be handled with similar parameters. Motivated by this observation, a heuristic that one can use is to reduce the choice of different μ^c to the choice of a single parameter ν by setting $\mu^c = \nu \xi^c$, where $\xi^c = \max(f^c)$. This second, single parameter ν can then again be chosen by visual inspection.

Using this strategy for our experimental data, we first fixed $\mu_{\text{HAADF}} = 0.1$. Computing the $\max(f^c)$ then resulted in a ratio of $\mu_{\text{Al}} : \mu_{\text{Si}} : \mu_{\text{Yb}} = 2.4 : 1.4 : 1$ and choosing the parameter ν again by visual inspection finally led to the parameters for the EDXS maps given as $(\mu_{\text{Al}}, \mu_{\text{Si}}, \mu_{\text{Yb}}) = (0.0024, 0.0014, 0.001)$. A comparison of different settings for the EDXS parameters using this ratio is given in Fig. 6b.

For the parameters used in the reconstruction of the phantom data shown in Fig. 3, we refer the reader to ESI Table 1,† where we used visual comparison of results to determine the parameters.

Finally, we also refer the reader to the ESI† for more information on the parameter choice and the performance of our method in the case of limited angle tomography.

Conflicts of interest

There are no conflicts of interest to declare.

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