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(54) Title: LOW DOSE SINGLE STEP GRATING BASED X-RAY PHASE CONTRAST IMAGING

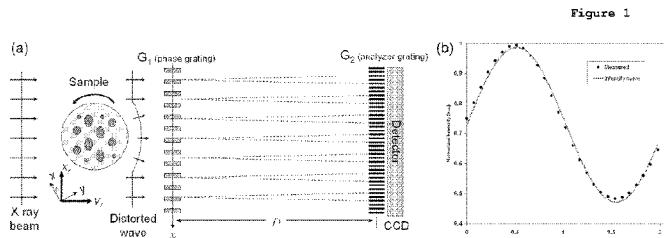


Figure 1

(57) Abstract: Phase sensitive X-ray imaging methods can provide substantially increased contrast over conventional absorption based imaging, and therefore new and otherwise inaccessible information. The use of gratings as optical elements in hard X-ray phase imaging overcomes some of the problems that have impaired the wider use of phase contrast in X-ray radiography and tomography. So far, to separate the phase information from other contributions detected with a grating interferometer, a phase-stepping approach has been considered, which implies the acquisition of multiple radiographic projections. Here, an innovative, highly sensitive X-ray tomographic phase contrast imaging approach is presented based on grating interferometry, which extracts the phase contrast signal without the need of phase stepping. Compared to the existing phase step approach, the main advantage of this new method dubbed "reverse projection" is the significantly reduced delivered dose, without degradation of the image quality. The new technique sets the pre-requisites for future fast and low dose phase contrast imaging methods, fundamental for imaging biological specimens and in-vivo studies.

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Low dose single step grating based X-ray phase contrast imaging

5 The present invention relates of a method and a system for low dose single step grating based X-ray phase contrast imaging.

It is well known that, differently from conventional visible light optics, the refractive index in X-ray optics 10 is very close to and smaller than unity since the X-ray photon energy is often much larger than the atomic resonance energies. In first approximation, for small and negligible anisotropy in the medium, the index of refraction characterizing the optical properties of a 15 tissue can be expressed - including X-ray absorption - with its complex form: $n=1-\delta-i\beta$ where δ is the decrement of the real part of the refractive index, characterizing the phase shifting property, while the imaginary part β describes the absorption property of the sample. In conventional 20 absorption-based radiography, the X-ray phase shift information is usually not directly utilized for image reconstruction. However, at photon energies greater than 10 keV and for light materials (made up of low-Z elements), the phase shift term plays a more prominent role than the 25 attenuation term because δ is typically three orders of magnitude larger than β . As a consequence, phase-contrast modalities can generate significantly greater image contrast compared to conventional, absorption-based imaging. Furthermore, far from absorption edges, δ is 30 inversely proportional to the square of the X-ray energy whilst β decreases as the fourth power of energy. A significant consequence of this mechanism is that phase signals can be obtained with much lower dose deposition than absorption, a very important issue when radiation 35 damage has to be taken into account such as in biological samples or in living systems.

Several approaches have been developed in order to record the phase signal. They can be classified as interferometric methods (with crystals), phase propagation methods, techniques based on an analyzer crystal or on grating
5 interferometry.

In the prior art the feasibility of two-grating interferometry in the hard X-ray region using a pair of transmission gratings made by gold stripes on glass plates
10 has been demonstrated. This work has since been extended to achieve a three-dimensional tomographic phase reconstruction using a hard X-ray two-gratings interferometer. Recently, three-grating interferometry in the hard X-ray region with low-brilliance tube-based X-ray
15 sources has been demonstrated. This laboratory-based instrument is of great interest for applications in biology, medicine and for non-destructive testing.

A grating interferometer setup is mechanically robust, is easy to align, has low sensitivity to mechanical drift and
20 its requirements on temporal coherence ($\Delta E/E \sim 0.1-0.2$) and spatial coherence (few microns) are moderate: as a consequence the instrument can be easily scaled up to large fields of view, an important asset when used in combination with a conventional X-ray tube.

25 These characteristics make grating interferometry superior to other phase contrast approaches and set the prerequisites for a broad use of phase contrast X-ray radiography and tomography.

30 To separate the phase information from other contributions, a phase-stepping approach is normally adopted. One of the gratings is displaced transversely to the incident beam whilst acquiring multiple projections. The intensity signal
35 at each pixel in the detector plane oscillates as a function of the displacement and the phase of this intensity oscillation can be directly linked to the wave-

front phase profile and to the decrement of the real part δ of the object's refractive index.

Obviously, this approach is loaded with the limitation of both (long) data acquisition time and severe dose released to specimen.

SUMMARY

It is therefore the objective of the present invention to provide a method and a system for the extraction of the phase information, which does not require a stepping procedure, thus overcoming limitations of both data acquisition time and dose imparted to the specimen.

According to an aspect of the present disclosure, there is provided an imaging set-up for reverse projection to obtain quantitative (hard) X-ray images from a sample and to quantitatively extract

both absorption and phase information from the sample, comprising:

- a. an X-ray beam generated by an X-ray source;
- b. a beam splitter grating (G1) and an analyzer grating (G2) having their respective lines parallel to each other, wherein the beam splitter grating (G1) is a line grating and the analyzer grating is a line absorption grating with high x-ray absorption; wherein a mechanism is comprised to place the sample to be investigated either between the X-ray source and the beam splitter grating (G1) or between the beam splitter grating (G1) and the analyzer grating (G2);
- c. a position-sensitive detector (PSD) with spatially modulated detection sensitivity having a number of individual pixels;
- d. means for recording the images of the detector, wherein a series of M images is collected by continuously or stepwise rotating from 0 to π or 2π either sample or the gratings (G1, G2) and the X-ray source relative to the sample, wherein each image taken at an angle $0 \leq \Phi \leq \pi$ comprises a corresponding reverse projection image taken at an angle $\pi \leq \Phi + \pi \leq 2\pi$, yielding in total a number of $M/2$ pairs of specular images;
- e. means for calculating pixel-wise an absorption image M and an refraction angle θ_r image out of the pairs of specular images - without the need of phase stepping according to

$$\ln \left(\frac{2S \left(\frac{x_g}{D} \right) I_0}{I(x_r, \phi, z) + I(-x_r, \phi + \pi, z)} \right) = M(x_r, \phi, z) = \int_{-\infty}^{\infty} \mu(x, y, z) dy,$$

$$\frac{1}{C} \frac{I(x_r, \phi, z) - I(-x_r, \phi + \pi, z)}{I(x_r, \phi, z) + I(-x_r, \phi + \pi, z)} = \theta_r(x_r, \phi, z) = - \int_{-\infty}^{\infty} \frac{\partial \delta(x, y, z)}{\partial x_r} dy,$$

wherein a shifting curve is determined by scanning without the sample the phase grating or the analyzer grating along the transverse direction x_g over one period of the analyzer grating and record the normalized intensity I/I_0 on the detector versus the angle x_g/D wherein x_g is the relative displacement between the beam splitter and the analyser grating along the direction perpendicular to both the incoming beam and the line of gratings and D is the distance between the beam splitter and the analyser grating.

According to another aspect of the present disclosure, there is provided a method for reverse

0 projection to obtain quantitative (hard) X-ray images from a sample and to quantitatively extract both absorption and phase information from the sample, comprising the steps of:

- 5 a. providing an X-ray source (X-ray);
- b. providing a beam splitter grating (G1) and an analyzer grating (G2) having their respective lines parallel to each other, wherein the beam splitter grating (G1) is a line grating, either an absorption grating with high X-ray absorption or a phase grating with low X-ray absorption, and the analyzer grating (G2) is a line absorption grating with high X-ray absorption;
- c. providing a position-sensitive detector (PSD) with spatially modulated detection sensitivity having a number of individual pixels;
- 15 d. positioning at least one of the gratings, such as G1 and G2 relative to the probe in a direction (x_g) substantially perpendicular to both the incoming beam and the orientation of the lines of grating to make the imaging set-up on the center of the linear region of the shifting curves $S(x_g/D)$;
- e. placing the sample to be investigated either between the X-ray source and the beam splitter grating (G1) or between the beam splitter grating (G1) and the

analyzer grating (G2), applying shots of the X-ray source to the sample and recording the images of the detector (PSD);

f. recording the images of the detector, wherein a series of M images is collected by continuously or stepwise rotating from 0 to pi or 2pi either the sample or the gratings (GO, G1, G2) and the X-ray source relative to the sample, wherein each image taken at an angle $0 \leq \Phi \leq \pi$ comprises a corresponding reverse projection image taken at an angle $\pi \leq \Phi + \pi \leq 2\pi$, yielding in total a number of $M/2$ pairs of specular images;

g. means for calculating pixel-wise an absorption image M and an refraction angle θ_r image out of the pairs of specular images-without the need of phase stepping according to:

$$\ln \left(\frac{2S \left(\frac{x_g}{D} \right) I_0}{I(x_r, \Phi, z) + I(-x_r, \Phi + \pi, z)} \right) = M(x_r, \Phi, z) = \int_{-\infty}^{\infty} \mu(x, y, z) dy_r$$

$$\frac{1}{C} \frac{I(x_r, \Phi, z) - I(-x_r, \Phi + \pi, z)}{I(x_r, \Phi, z) + I(-x_r, \Phi + \pi, z)} = \theta_r(x_r, \Phi, z) = - \int_{-\infty}^{\infty} \frac{\partial \delta(x, y, z)}{\partial x_r} dy_r$$

wherein a shifting curve is determined by scanning without the sample the phase grating or the analyzer grating along the transverse direction x_g over one period of the analyzer grating and record the normalized intensity I/I_0 on the detector versus the angle x_g/D wherein x_g is the relative displacement between the beam splitter and the analyser grating along the direction perpendicular to both the incoming beam and the line of gratings and D is the distance between the beam splitter and the analyser grating.

20 The invented system and method therefore present an innovative, highly sensitive X-ray tomographic phase contrast imaging approach based on grating interferometry, which extracts the phase contrast signal without the need of phase stepping (PS). Compared to the existing phase step approach, the main advantage of this invention dubbed "reverse projection (RP)" is the significantly reduced delivered dose, without degradation of the image quality. The new
25 technique sets the pre-requisites for future fast and low dose phase contrast imaging methods, fundamental for imaging biological specimens and in-vivo studies. Typically, the beam splitter

grating may be a line grating, preferably a phase grating; that is, a grating with low X-ray absorption, but considerable X-ray phase shift(Φ), the latter preferably of either

$$\Phi \in \left((2l-1)\frac{\pi}{2} - \arcsin 0.8, (2l-1)\frac{\pi}{2} + \arcsin 0.8 \right) \text{ or}$$
$$\Phi \in \left((2l-1)\pi - \arcsin 0.8, (2l-1)\pi + \arcsin 0.8 \right), \text{ where } l = 1, 2, 3, \dots.$$

A further preferred embodiment of the present invention may provide the phase grating that acts as the beam splitter is made by deep etching into silicon, a polymer or similar material.

A further preferred embodiment of the present invention may provide the analyzer grating with one-dimensional grating structure being integrated into the detector, the pixel of the detector is in range of 2 to 10 times the size of the period of the grating, half lines with sensor in a pixel are sensitive to X-ray and half lines without sensor let X-ray go through. In this way the analyzer grating with 100% absorption can be achieved without needing to make heavy metal absorption gratings with high aspect ratio, in particular it is possible to avoid gold gratings.

A further preferred embodiment of the present invention may provide an analyzer grating having a one-dimensional grating structure with high X-ray absorption contrast, its period is the same as that of the self image of the phase grating, placed closely in front of the detector, with its lines parallel to those of the phase grating; preferably this analyzer grating serves as an anti-scatter grid, or an anti-scatter grid is used as a modulation mask.

Dimensioning the interferometer is fundamental for the present invention. Advantageously, the distance between the beam splitter grating and the analyzer grating is chosen to be an odd fractional Talbot distance, given by the equation

$$D_{n,sph} = \frac{L \cdot D_n}{L - D_n} = \frac{L \cdot n \cdot p_1^2 / 2\eta^2 \lambda}{L - n \cdot p_1^2 / 2\eta^2 \lambda}, \text{ where } n = 1, 3, 5, \dots, \text{ and}$$

$$\eta = \begin{cases} 1 & \text{if the phase shift of } G_1 \text{ is } (2l-1)\frac{\pi}{2}, \quad p_2 = \frac{L+D_{n,sph}}{L} p_1 \\ 2 & \text{if the phase shift of } G_1 \text{ is } (2l-1)\pi, \quad p_2 = \frac{L+D_{n,sph}}{L} \frac{p_1}{2} \end{cases}, \text{ where}$$

$l=1,2,3,\dots$, D_n is an odd fractional Talbot distance when the parallel X-ray beam is used, while $D_{n,sph}$ is that when the fan or cone X-ray beam is used, L is the distance between

5 the source and the phase grating.

Further, the position of half slope on the shifting curve may be achieved by positioning at least one of the beam

10 splitter grating and the analyzer grating relative to the probe in a direction substantially perpendicular to the orientation of the lines in at least one of the two gratings.

15 In order to establish a rather simple set-up of the present interferometer, a mechanism can be comprised to place the sample to be investigated between the source and the beam splitter grating or between the beam splitter grating and the analyzer grating being rotated from 0 to π or to 2π .

20

A further preferred embodiment of the present invention may provide a collimator placed between the source and the beam splitter grating limiting the spatial extent of the illuminating X-rays to a fan beam; a line-array detector is

25 used, and a mechanism is comprised that allows to rotate (either stepwise or continuously) the sample relative to the rest of the apparatus, the rotational axis being perpendicular to the opening angle of the fan, and preferably at same time allows to translate (either

30 stepwise or continuously) the sample relative to the rest of the apparatus along the direction parallel to the rotational axis.

Alternatively, a collimator placed between the source and the beam splitter grating may limit the spatial extent of the illuminating X-rays to a cone beam, a pixel-array 5 detector is used, and a mechanism is comprised that allows to rotate the sample relative to the rest of the apparatus, perpendicular to the opening angle of the fan.

Excellent results with respect to the quality of the image 10 can be achieved when an analysis procedure is implemented for reverse-projection data that comprises the steps of calculating, for each element of the detector, the absorption signal M and the refraction angle θ_r according to the following equations (8) and (9) resp.: 15

$$\ln \left(\frac{2S \left(\frac{x_g}{D} \right) I_0}{I(x_r, \phi, z) + I(-x_r, \phi + \pi, z)} \right) = M(x_r, \phi, z) = \int_{-\infty}^{\infty} \mu(x, y, z) dy_r \quad (8)$$

$$\frac{1}{C} \frac{I(x_r, \phi, z) - I(-x_r, \phi + \pi, z)}{I(x_r, \phi, z) + I(-x_r, \phi + \pi, z)} = \theta_r(x_r, \phi, z) = - \int_{-\infty}^{\infty} \frac{\partial \delta(x, y, z)}{\partial x_r} dy_r \quad (9)$$

20 Preferred embodiments of the present invention are hereinafter described in more detail thereby referring to the attached drawings.

25 Brief description of the drawings:

Figure 1 (a) shows the working principle of the grating interferometer: through the Talbot effect, a periodic interference pattern (known as self image) is formed behind 30 the phase grating (G1), in the plane of the analyzer grating (G2). Figure 1(b) is a plot of the intensity oscillation (shifting curve) as a function of the grating position x_g for a detector pixel over one period of the

analyzer grating. The dots corresponds to the measured values (normalized to unit) while the gray line shows a sinusoidal fit.

5 Figure 2 illustrates phase contrast tomographic reconstructions of a demineralised mouse joint, acquired at a voxel size of $3.5 \times 3.5 \times 3.5 \mu\text{m}^3$. Sub-Figures a1 to a3 show the data obtained with the classical phase stepping (PS) protocol, while Sub-Figures b1 to b3 the
10 reconstruction using the reverse projection (RP) method. a1 and b1 shows an axial slice: b1 is sharper than a1 and there are no ring artifacts (see text below). a2 and b2 depict a coronal slice through the joint, clearly showing that the RP protocol is less sensitive to typical
15 horizontal stripes artifacts observed with the PS method (see enlarged inset). a3 and b3 show a sagittal view through the joint. The dotted lines mark the locations where the axial views (a1 and b1) have been taken. Scale bar is 500 microns.

20 Figure 3 represents a phase contrast reconstructed coronal slice of a rat brain, obtained after tomographic reconstruction using the PS- (a) and the RP-protocol (b). Qualitatively, both reconstructions are very similar. In
25 (b) the effects of the grating imperfection (ring artifacts), as expected, are more evident. Figure (c) shows a quantitative comparison of two line profiles extracted at the position marked by colour bars (hippocampus region). Scale bar is 1 mm.

30 Figure 4(a) shows differential phase contrast radiography of a rat paw (7 stacks, RP-protocol). Figures 4(b1-2) and (c1-2) show axial and coronal slices through the paw acquired with the PS and RP protocol, respectively.
35 Structural details of both soft (muscles, fat) and hard tissue (bone) are well visible. Scale bars are 2 mm in (a) and 1 mm for (b1-2) and (c1-2).

Figure 5(a1-2) is a tomographic reconstruction of a rat brain - (a1-2) obtained with the PS protocol, (b1-2) obtained with the RP-protocol using Eq. 11 to calculate the 5 map of the index of refraction. Scale bar is 1 mm in (a1,b1) and 2 mm in (a2,b2).

Table 1 summarizes the experimental parameters for the 10 tomographic scans of the three investigated samples: a rat brain (4% PFA, paraffin embedded), a (demineralized) mouse joint in PBS and a rat paw (4% PFA). All experiments have been performed at 25 keV and at the 3rd Talbot distance. Visibility of the interferometer was ~ 30%.

15 With reference to the above-mentioned figures, an innovative approach for the extraction of the phase information is presented which does not require a stepping procedure, thus overcoming limitations of both data acquisition time and dose released to specimens.

20 This novel approach relies on the physical similarities between a crystal analyzer based system and a grating interferometer. Both techniques record refraction angle signals and, analogously to the rocking curve of a crystal 25 analyzer, the properties of the shifting curve (see Figure 1) can be exploited to fully describe the performance of a grating interferometer. The refraction angle, i.e., the phase information of the sample, can be extracted by setting the grating interferometer in the central position 30 where the intensity curve follows a linear behavior.

According to the aforementioned analogy, the intensity I recorded by a detector positioned after the grating interferometer can be expressed as:

35

$$I = I_0 \cdot \exp \left[- \int_{-\infty}^{\infty} \mu(x, y, z) dy_r \right] \cdot S \left(\frac{x_g}{D} + \theta_r \right), \quad [1]$$

where μ is the linear absorption coefficient, x_g denotes the relative displacement between the phase grating and the analyzer grating along the direction perpendicular to both the incoming beam and the line of gratings, θ_r is the refraction angle, D is the distance between the phase and the analyzer grating, $S \left(\frac{x_g}{D} \right)$ is the shifting curve. For the sake of simplicity, the scattering contribution - which would induce a weak increment of the background noise - is neglected. (x_r, y_r, z) are the coordinates of the reference frame associated to the X-ray beam and (x, y, z) those associated with the sample. The two frames are linked by the rotation matrix

$$\begin{pmatrix} x \\ y \end{pmatrix} = \begin{pmatrix} \cos\phi & -\sin\phi \\ \sin\phi & \cos\phi \end{pmatrix} \begin{pmatrix} x_r \\ y_r \end{pmatrix} \quad [2]$$

being ϕ the rotation angle between the x_r and the x -axis around the z -axis.

With a good approximation, the behaviour of the shifting curve near its half slope may be considered linear so that, if p_2 is the period of the analyzer grating, $\theta_r \leq \frac{p_2}{4D}$ can be replaced by a first-order Taylor expansion. Further we can write:

$$S \left(\frac{x_g}{D} + \theta_r \right) = S \left(\frac{x_g}{D} \right) + \frac{dS \left(\frac{x_g}{D} \right)}{d\theta} \theta_r = S \left(\frac{x_g}{D} \right) (1 + C \theta_r) \quad [3]$$

where $C = \frac{1}{S \left(\frac{x_g}{D} \right)} \frac{dS \left(\frac{x_g}{D} \right)}{d\theta}$ is a constant.

25

The refraction angle in the X-Y plane (Fig. 1(a)) is determined by the line integral of the first-order

derivative of the refractive index along the X-ray path and it may be written as:

$$\theta_r = - \int_{-\infty}^{\infty} \frac{\partial \delta}{\partial x_r} dy_r , \quad [4]$$

where δ corresponds to the decrement of the real part of the refractive index of the sample as mentioned in the introduction. Substituting Eqs. 3 and 4 into Eq. 1, the projected image for a grating interferometer can be described by:

$$10 \quad I(x_r, z) = I_0 \exp \left\{ - \int_{-\infty}^{\infty} \mu(x, y, z) dy_r \right\} S \left(\frac{x_g}{D} \right) \left[1 - C \int_{-\infty}^{\infty} \frac{\partial \delta(x, y, z)}{\partial x_r} dy_r \right] \quad [5]$$

μ is a scalar and therefore rotational-invariant, while $\frac{\partial \delta}{\partial x_r}$ strongly depends on the direction along which it is measured.

15 The projected image at the rotation angles ϕ and its corresponding reverse image at $\phi + \pi$ can be written as:

$$I(x_r, \phi, z) = I_0 \exp \left\{ - \int_{-\infty}^{\infty} \mu(x, y, z) dy_r \right\} S \left(\frac{x_g}{D} \right) \left[1 - C \int_{-\infty}^{\infty} \frac{\partial \delta(x, y, z)}{\partial x_r} dy_r \right] \quad [6]$$

$$I(-x_r, \phi + \pi, z) = I_0 \exp \left\{ - \int_{-\infty}^{\infty} \mu(x, y, z) dy_r \right\} S \left(\frac{x_g}{D} \right) \left[1 + C \int_{-\infty}^{\infty} \frac{\partial \delta(x, y, z)}{\partial x_r} dy_r \right] \quad [7]$$

The absorption signal can be obtained from the two projected images by the sum of Eq. 6 and 7 and solving the Beer-Lambert relationship, i.e.,

20

$$\ln \left(\frac{2S \left(\frac{x_g}{D} \right) I_0}{I(x_r, \phi, z) + I(-x_r, \phi + \pi, z)} \right) = M(x_r, \phi, z) = \int_{-\infty}^{\infty} \mu(x, y, z) dy_r \quad [8]$$

In the same way, the angle of refraction can be obtained by a proper combination of Eq. 6 and 7, as shown in the following expression:

$$5 \quad \frac{1}{C} \frac{I(x_r, \phi, z) - I(-x_r, \phi + \pi, z)}{I(x_r, \phi, z) + I(-x_r, \phi + \pi, z)} = \theta_r(x_r, \phi, z) = - \int_{-\infty}^{\infty} \frac{\partial \delta(x, y, z)}{\partial x_r} dy_r \quad [9]$$

According to fundamentals of computed tomography reconstruction, i.e., the Fourier Slice Theorem, the absorption coefficient as well as the refractive index can 10 be easily obtained by the inverse Fourier Transform and a Hilbert filter:

$$\mu(x, y, z) = \int_0^{\pi} d\phi \int_{-\infty}^{\infty} [M(x_r, \phi, z) * F^{-1}(|\rho|)] \cdot \delta(x \cos \phi + y \sin \phi - x_r) dx_r \quad [10]$$

$$\delta(x, y, z) = - \int_0^{\pi} d\phi \int_{-\infty}^{\infty} \left[\theta_r(x_r, \phi, z) * F^{-1} \left(\frac{|\rho|}{2\pi j \rho} \right) \right] \cdot \delta(x \cos \phi + y \sin \phi - x_r) dx_r \quad [11]$$

where ρ is the spatial frequency and F^{-1} denotes the inverse Fourier transform.

15

Based on Eq. 10 and 11, we introduce here a novel acquisition protocol, dubbed "reverse projection" (RP) method.

20 It can be described in five steps:

- (i) without sample, scan the phase grating or the analyzer grating along the transverse direction x_g over one period of the analyzer grating and record the normalized intensity $\frac{I}{I_0}$ on the detector versus the angle $\frac{x_g}{D}$, i.e., get the shifting curve $S\left(\frac{x_g}{D}\right)$,

- (ii) set the grating interferometer at the center of the linear region of the shifting curve by positioning the phase grating or the analyzer grating at $x_g = p_2/4$ or $x_g = -p_2/4$,
- 5 (iii) put the sample in front of or behind the phase grating, collect m angular projections of the sample over a rotation of 360° ,
- (iv) extract M and θ , according to Eq. 8 and Eq. 9 and finally
- 10 (v) reconstruct either the absorption coefficient or the refractive index using the filtered back-projection. Therefore, the total number of acquired projection images is m .

15 On the contrary, the Phase Stepping (PS) acquisition protocol can be described in four steps:

- (i) put the sample in front of or behind the phase grating, scan one of the two gratings along the transverse direction x_g (k points over one period of the analyzer grating) and record one projection image at each point,
- 20 (ii) repeat step (i) for a total of $m/2$ times over a sample rotation of 180° ,
- (iii) extract the gradient signal via the Fourier analysis of the intensity signal and
- 25 (iv) reconstruct the phase via a filtered back-projection.

For this second method, the total number of acquired projections images is $k*m/2$.

30 As a consequence, the total number of projections required by the RP protocol is reduced by a factor of $k/2$ compared to the PS.

The method was validated by performing both phase stepping (PS) and reverse projection (RP) experiments using the grating interferometer installed at the TOMCAT beamline of the Swiss Light Source at the Paul Scherrer Institute, 5 Villigen, Switzerland. The energy was tuned at 25 keV and the interferometer was operated in the 3rd Talbot distance. In this configuration, the visibility has been measured to be 30%. Additional details on the grating interferometer installed at TOMCAT can be found in public documentation 10 related to this installation with the Paul Scherrer Institute.

In a first case study, we investigated two different samples: a rat brain first fixed in 4% paraformaldehyde 15 (PFA) and then embedded in paraffin and a demineralized mouse joint, fixed only in a phosphate buffer solution (PBS) (no embedding). We used the mouse joint and the rat brain to test the reconstruction method both on small (< 4 mm) and large (> 10 mm) samples. Both specimens are weakly 20 absorbing objects and therefore ideal candidates for phase contrast imaging. Reconstructions based on the two methods (PS and RP) are shown in Fig. 2 and 3 while experimental parameters are summarized in Table 1.

25 Fig. 2 shows axial, sagittal and coronal views of a mouse joint obtained with both PS and RP protocols (Table 1). The joint was immersed and fixed in an Eppendorf vial containing PBS to avoid any movements during the acquisition. A qualitative comparison of the images clearly 30 shows that RP-reconstructions are comparable to those obtained with the PS approach. Moreover, looking at the inset shown in Figure (2,a2) and (2,b2) the RP-slice appears to be sharper than the PS-reconstructions. This can be explained by the fact that the shifting curve is 35 directly proportional to the refraction angle and that this - in the RP protocol - is obtained by simple subtraction of a reference image (with no sample) from the paired images

described in Eq. 9. In addition, since with the RP method no phase-stepping is required, the system is less sensitive to mechanical instabilities.

5 The largest investigated sample, a rat brain, was mounted vertically on the flat surface of the sample support to match the horizontal field of view of the detector system. The vertical sample arrangement also enabled a direct reconstruction of coronal slices through the sample, an
10 approach very useful when trying to identify anatomical brain regions (Fig. 3). The height of the sample was larger than the vertical height of the beam and therefore four scans have been collected along the vertical direction to image the whole brain. To achieve phase matching between
15 sample and surroundings, we used an aquarium bath filled with room temperature liquid paraffin (chemical formula C_nH_{2n-2} where $n = 5-17$, density $\approx 0.7 \text{ g cm}^{-3}$). For large samples too, a qualitative comparison of the images clearly shows that the RP-reconstruction is as good as the one
20 obtained with the PS-approach. In addition, a line profile taken at the level of the hippocampus, see Fig. 3c, shows a quantitative good agreement between RP and PS approaches.

In the second case, the novel method has been validated
25 using a more realistic sample, namely a specimen containing both soft and hard tissue. For this purpose, we investigated a rat paw (containing both bone and muscles) which was only fixed in 4% PFA. This fixation procedure is frequently used to maintain biological samples in a status
30 as close as possible to their natural, original conditions. The rat paw was also mounted vertically in order to best match the horizontal field of view of the detector. Seven stacked scans were necessary to image the full sample volume.
35 The measurement of the rat paw was the most challenging experiment since the sample has been measured in air. This usually causes large phase jumps at the air-specimen

interface and explains the "star" artifacts visible in Fig. 4b-1 and, less serious, in 4c-1. This is because the shifting curve is saturated when $\theta_r \geq \frac{p_2}{4D}$ and, as a consequence, the RP method is not very sensitive to large 5 refraction angles. This is not the case for the PS method, which has to cope with angles as large as $\theta_r \leq \frac{p_2}{2D}$.

Our invention introduces a novel approach for fast and low dose extraction of both the absorption coefficient and the 10 refractive index of a sample using a grating interferometer is introduced. It is demonstrated that this new approach yields comparable information to the established phase stepping technique but with a reduction factor of $k/2$ in the total dose delivered to the sample. Moreover, the 15 reverse projection approach makes high-sensitivity phase contrast computed tomography (CT) as straightforward as conventional, absorption based CT. It is first shown that this new method works well with parallel beam geometries but it is not difficult to generalize it to either cone or 20 fan beam setups, making it accessible also to X-ray tube-based applications.

In particular, the significant decrease of the dose and the straight forward acquisition protocol does no affect image 25 quality, while representing a major advancement in hard X-ray phase contrast tomography for synchrotron radiation and laboratory X-ray sources, enabling experiments impossible so far.

30 The next and probably most challenging application of the RP-protocol will be *in-vivo phase contrast* imaging. With the advent of new, high efficient and high speed detectors it will be possible to acquire the same amount of data within a fraction of a second. We estimate that it will be 35 realistic to obtain a full tomographic data set with the RP

protocol with a total exposure time of 2-3 seconds. This, together with the ongoing efforts regarding robust and reliable iterative reconstruction algorithms, requiring a significant smaller amount of projections, can push the

5 total acquisition time below 1s and hence opening up the possibility of phase contrast tomographic microscopy of small living animals.

Another very challenging application of the RP-protocol
10 will be the quantitative 3D description of the scattering signal. This image contrast is generated by small-angle scattering within the sample and it provides complementary and otherwise inaccessible structural information at micrometer and sub-micrometer length scale. However, the
15 signal is not rotational-invariant and therefore it will be very challenging to quantitatively obtain such information in 3D.

Further developments will concern the manufacturing of
20 optimized gratings for high X-ray energies leading to the implementation of the RP-protocol in new medical X-ray CT scanners that would offer a significant increase in soft tissue sensitivity, a characteristic now provided (at much lower resolutions however) only by much more expensive
25 techniques such as magnetic resonance imaging.

Finally, we would like to point out that this approach is not limited to X-ray imaging and may be easily generalized to other methods such as grating based neutron phase
30 imaging and visible light differential interference contrast (DIC) microscopy where a similar shifting curve is considered and a quantitative phase description appears possible.

Table 1

| | Mouse joint | | Rat brain | | Rat paw | |
|----------------------------|-------------------|-----------------------|-----------------------------|-----------------------|-------------------|-----------------------|
| | Phase Stepping | Reverse Projection | Phase Stepping | Reverse Projection | Phase Stepping | Reverse Projection |
| Rotation | 0-180° | 0-360° | 0-180° | 0-360° | 0-180° | 0-360° |
| Pixel size [μm] | 3.5 × 3.5 | 3.5 × 3.5 | 11.2 × 11.2 | 11.2 × 11.2 | 7.4 × 7.4 | 7.4 × 7.4 |
| Field of view [mm] | 3.58 × 3.58 | 3.58 × 3.58 | 11.45 × 3.6 ¹ | 11.45 × 3.6 | 15.5 × 3.6 | 15.5 × 3.6 |
| Angl. proj. | 181 | 361 | 361 | 721 | 501 | 1001 |
| Phase steps | 9 | 1 | 9 | 1 | 9 | 1 |
| Single exposure [ms] | 200 | 200 | 200 | 200 | 60 | 60 |
| Total exposure (s) | 325 | 72 | 650 | 144 | 270 | 60 |

¹ The optical system always produces field of views with a square shape. However, along the vertical direction the size of the field of view is reduced and limited by the vertical size of the beam (approximately 3.6 mm at 25 keV and at 25 m from the source for the TOMCAT beamline of the Swiss Light Source).

SUPPLEMENTARY MATERIAL I

Ring-like artefacts due to grating imperfections are discussed hereinafter.

5

A careful study of the reconstructed images reveals that the reverse projections (RP) protocol imposes more stringent requirements on gratings, i.e., it needs better full-field uniformity and lower local grating imperfections
10 than gratings suitable for the phase stepping (PS) method. In particular, imperfections may induce small ring-like artefacts in the RP-reconstructed slices, which are less evident or missing in PS-reconstructions.

15 Ring artifacts are clearly visible in Figure 5, for both coronal (b1) and sagittal (b2) cuts. Due to the averaging effect associated to the phase stepping extraction, the PS-protocol is significantly less sensitive to grating defects and therefore the rings artifact are less pronounced, see
20 Figure 5 a1-2.

25

CLAIMS

1. An imaging set-up for reverse projection to obtain quantitative (hard) X-ray images from a sample and to quantitatively extract both absorption and phase information from the sample, comprising:

- a. an X-ray beam generated by an X-ray source;
- b. a beam splitter grating (G1) and an analyzer grating (G2) having their respective lines parallel to each other, wherein the beam splitter grating (G1) is a line grating and the analyzer grating is a line absorption grating with high x-ray absorption; wherein a mechanism is comprised to place the sample to be investigated either between the X-ray source and the beam splitter grating (G1) or between the beam splitter grating (G1) and the analyzer grating (G2);
- c. a position-sensitive detector (PSD) with spatially modulated detection sensitivity having a number of individual pixels;
- d. means for recording the images of the detector, wherein a series of M images is collected by continuously or stepwise rotating from 0 to pi or 2pi either sample or the gratings (G1, G2) and the X-ray source relative to the sample, wherein each image taken at an angle $0 \leq \Phi \leq \pi$ comprises a corresponding reverse projection image taken at an angle $\pi \leq \Phi + \pi \leq 2\pi$, yielding in total a number of $M/2$ pairs of specular images;
- e. means for calculating pixel-wise an absorption image M and an refraction angle θ_r image out of the pairs of specular images - without the need of phase stepping according to:

$$\ln \left(\frac{2S \left(\frac{x_g}{D} \right) I_\phi}{I(x_r, \Phi, z) + I(-x_r, \Phi + \pi, z)} \right) = M(x_r, \Phi, z) = \int_{-\infty}^{\infty} \mu(x, y, z) dy_r$$

$$\frac{1}{C} \frac{I(x_r, \Phi, z) - I(-x_r, \Phi + \pi, z)}{I(x_r, \Phi, z) + I(-x_r, \Phi + \pi, z)} = \theta_r(x_r, \Phi, z) = - \int_{-\infty}^{\infty} \frac{\partial \delta(x, y, z)}{\partial x_r} dy_r$$

wherein a shifting curve is determined by scanning without the sample the phase grating or the analyzer grating along the transverse direction x_g over one period of the analyzer grating and record the normalized intensity I/I_0 on the detector versus

the angle x_g/D wherein x_g is the relative displacement between the beam splitter and the analyser grating along the direction perpendicular to both the incoming beam and the line of gratings and D is the distance between the beam splitter and the analyser grating.

2. The imaging setup according to claim 1 wherein the intensity I recorded by the PSD is expressed as:

$$I = I_0 \cdot \exp \left[- \int_{-\infty}^{\infty} \mu(x, y, z) dy_r \right] \cdot S \left(\frac{x_g}{D} + \theta_r \right),$$

where μ is the linear absorption coefficient, x_g denotes the relative displacement between the phase grating and the analyzer grating along the direction perpendicular to both the incoming beam and the line of gratings, θ_r is the refraction angle, D is the distance between the phase and the analyzer grating, $S(x_g/D)$ is the shifting curve (see page 10, lines 1 to 7).

3. The imaging set-up according to claim 1 or 2, wherein an analyzer grating having a one-dimensional grating structure (G2) with high X-ray absorption contrast is placed closely in front of the position sensitive (PSD) detector with its lines parallel to those beam splitter grating (G1).

4. The imaging set-up according to any of the preceding claims, wherein an analyzer grating having a one-dimensional grating structure with high X-ray absorption contrast, its period is the same as that of the self-image of the beam splitter grating (G1), is placed closely in front of the detector (PSD) with its lines parallel to those of the beam splitter grating (G1).

5. The imaging set-up according to any of the preceding claims, wherein the distance (D) between the beam splitter grating (G1) and the analyzer grating (G2) is chosen to be an odd fractional Talbot distance, given by the equation

$$D_{n,spk} = \frac{L \cdot D_n}{L - D_n} = \frac{L \cdot n \cdot p_1^2 / 2\eta^2 \lambda}{L - n \cdot p_1^2 / 2\eta^2 \lambda}, \text{ where } n = 1, 3, 5, \dots, \text{ and}$$

$$\eta = \begin{cases} 1 & \text{if the phase shift of } G_1 \text{ is } (2l-1)\frac{\pi}{2}, \quad p_2 = \frac{L + D_{n,spk}}{L} p_1 \\ 2 & \text{if the phase shift of } G_1 \text{ is } (2l-1)\pi, \quad p_2 = \frac{L + D_{n,spk}}{L} \frac{p_1}{2} \end{cases}, \text{ where}$$

$l=1,2,3,\dots$, D_n is an odd fractional Talbot distance when the parallel X-ray beam is used, while D_{ngsph} is that when the fan or cone X-ray beam is used, L is the distance between the source and the phase grating.

6. The imaging set-up according to any of the preceding claims, wherein the beam splitter grating (G1) is a line phase grating with low x-ray absorption, but with considerable X-ray phase shift (Φ), the latter preferably of either

$$\Phi \in \left((2l-1) \frac{\pi}{2} - \arcsin 0.8, (2l-1) \frac{\pi}{2} + \arcsin 0.8 \right) \text{ or}$$

$$\Phi \in ((2l-1)\pi - \arcsin 0.8, (2l-1)\pi + \arcsin 0.8), \text{ where } l = 1, 2, 3, \dots.$$

7. The imaging set-up according to any of the preceding claims wherein if the beam splitter grating (G1) is a line phase grating with low X-ray absorption, it will be made into silicon, polymer or similar materials.

8. The imaging set-up according to any of the preceding claims, wherein the analyzer grating (G2) is either placed closely in front of the detector (PSD) or with its one-dimensional grating structure integrated into the detector, the pixel of the detector is from 2 to 10 times the size of the period of the grating, half lines with sensor in a pixel are sensitive to X-ray and half lines without sensor let X-ray go through.

9. The imaging set-up according to any of the preceding claims, wherein a collimator placed between the X-ray source and the beam splitter grating (G1) limits the spatial extent of the illuminating X-rays to a fan beam, a line array detector is used, and a mechanism is included that allows to rotate (either stepwise or continuously) the sample relative to the rest of the apparatus, the rotational axis being perpendicular to the opening angle of the fan, and preferably at the same time allows to translate (either stepwise or continuously) the sample relative to the rest of the apparatus along the direction parallel to the rotational axis.

10. A method for reverse projection to obtain quantitative (hard) X-ray images from a sample and to quantitatively extract both absorption and phase information from the sample, comprising the steps of:

- a. providing an X-ray source (X-ray);

- b. providing a beam splitter grating (G1) and an analyzer grating (G2) having their respective lines parallel to each other, wherein the beam splitter grating (G1) is a line grating, either an absorption grating with high X-ray absorption or a phase grating with low X-ray absorption, and the analyzer grating (G2) is a line absorption grating with high X-ray absorption;
- c. providing a position-sensitive detector (PSD) with spatially modulated detection sensitivity having a number of individual pixels;
- d. positioning at least one of the gratings, such as G1 and G2 relative to the probe in a direction (x_g) substantially perpendicular to both the incoming beam and the orientation of the lines of grating to make the imaging set-up on the center of the linear region of the shifting curves $S(x_g/D)$;
- e. placing the sample to be investigated either between the X-ray source and the beam splitter grating (G1) or between the beam splitter grating (G1) and the analyzer grating (G2), applying shots of the X-ray source to the sample and recording the images of the detector (PSD);
- f. recording the images of the detector, wherein a series of M images is collected by continuously or stepwise rotating from 0 to π or 2π either the sample or the gratings (G0, G1, G2) and the X-ray source relative to the sample, wherein each image taken at an angle $0 \leq \Phi \leq \pi$ comprises a corresponding reverse projection image taken at an angle $\pi \leq \Phi + \pi \leq 2\pi$, yielding in total a number of $M/2$ pairs of specular images;
- g. means for calculating pixel-wise an absorption image M and an refraction angle θ_r image out of the pairs of specular images-without the need of phase stepping - according to:

$$\ln \left\{ \frac{2S \left(\frac{x_s}{D} \right) I_0}{I(x_r, \Phi, z) + I(-x_r, \Phi + \pi, z)} \right\} = M(x_r, \Phi, z) = \int_{-\infty}^{\infty} \mu(x, y, z) dy,$$

$$\frac{1}{C} \frac{I(x_r, \Phi, z) - I(-x_r, \Phi + \pi, z)}{I(x_r, \Phi, z) + I(-x_r, \Phi + \pi, z)} = \theta_r(x_r, \Phi, z) = - \int_{-\infty}^{\infty} \frac{\partial \delta(x, y, z)}{\partial x_r} dy,$$

wherein a shifting curve is determined by scanning without the sample the phase grating or the analyzer grating along the transverse direction x_g over one period of the analyzer grating and record the normalized intensity I/I_0 on the detector versus the angle x_g/D wherein x_g is the relative displacement between the beam splitter and the analyser grating along the direction perpendicular to both the incoming beam and the line of gratings and D is the distance between the beam splitter and the analyser grating.

11. The method according to claim 10, wherein if the beam splitter grating (G1) is a line phase grating with low X-ray absorption, the thickness of the grating line will be with considerable X-ray phase shift (Φ), the latter preferably of either

$$\Phi \in \left((2l-1)\frac{\pi}{2} - \arcsin 0.8, (2l-1)\frac{\pi}{2} + \arcsin 0.8 \right) \text{ or}$$
$$\Phi \in ((2l-1)\pi - \arcsin 0.8, (2l-1)\pi + \arcsin 0.8), \text{ where } l = 1, 2, 3, \dots$$

12. The method according to claim 10 or 11 wherein if the beam splitter grating (G1) is a line phase grating with low X-ray absorption, it will be made into silicon, polymer or similar material.

13. The method according to any of the claims 10 to 12, wherein an analyzer grating (G2) having a one-dimensional grating structure with high X-ray absorption contrast, its period is the same as that of the image of the beam splitter grating, is placed closely in front of the detector (PSD) with its lines parallel to those of the phase grating; preferably this grating structure serves as an anti-scatter grid, or an anti-scatter grid is used as the modulation mask.

14. The method according to any of the preceding claims 10 to 13, wherein the distance between the beam splitter and the analyzer is chosen to be an odd fractional Talbot distance, given by the equation

$$D_{n,sph} = \frac{L \cdot D_n}{L - D_n} = \frac{L \cdot n \cdot p_1^2 / 2\eta^2 \lambda}{L - n \cdot p_1^2 / 2\eta^2 \lambda}, \text{ where } n = 1, 3, 5, \dots, \text{ and}$$

$$\eta = \begin{cases} 1 & \text{if the phase shift of } G_1 \text{ is } (2l-1)\frac{\pi}{2}, \quad p_2 = \frac{L + D_{n,sph}}{L} p_1 \\ 2 & \text{if the phase shift of } G_1 \text{ is } (2l-1)\pi, \quad p_2 = \frac{L + D_{n,sph}}{L} \frac{p_1}{2} \end{cases}, \text{ where}$$

$l = 1, 2, 3$, D_n is an odd fractional Talbot distance when the parallel X-ray beam is used, while $D_{n,sph}$ that when the fan or cone X-ray beam is used, L is the distance between the source and the phase grating.

15. The method according to any of the preceding claims 10 to 14, wherein a collimator placed between the source and the beam splitter grating (G1) limits the spatial extent of the illuminating X-rays to a fan beam, a line-array detector is used, and a mechanism is comprised that allows to rotate (either stepwise or continuously) the sample relative to the rest of the apparatus, the rotational axis being perpendicular to the opening angle of the fan, and preferably at the same time allows to translate (either stepwise or continuously) the sample relative to the rest of the apparatus along the direction parallel to the rotational axis.

16. The method according to any of the preceding claims 10 to 15, wherein a collimator placed between the source and the beam splitter grating (G1) limits the spatial extent of the illuminating X-rays to a cone beam, a pixel-array detector is used, and a mechanism is comprised that allows to rotate the sample relative to the rest of the apparatus, perpendicular to the opening angle of the fan.

17. The method according to any of the preceding claims 10 to 16, wherein the analyzer grating (G2) is either placed closely in front of the detector (PSD) or with its one-dimensional grating structure integrated into the detector, the pixel of the detector is from 2 to 10 times the size of the period of the grating, half lines with sensor in a pixel are sensitive to X-ray and half lines without sensor let X-ray go through.

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Patent Attorneys for the Applicant/Nominated Person

SPRUSON & FERGUSON

Figure 1

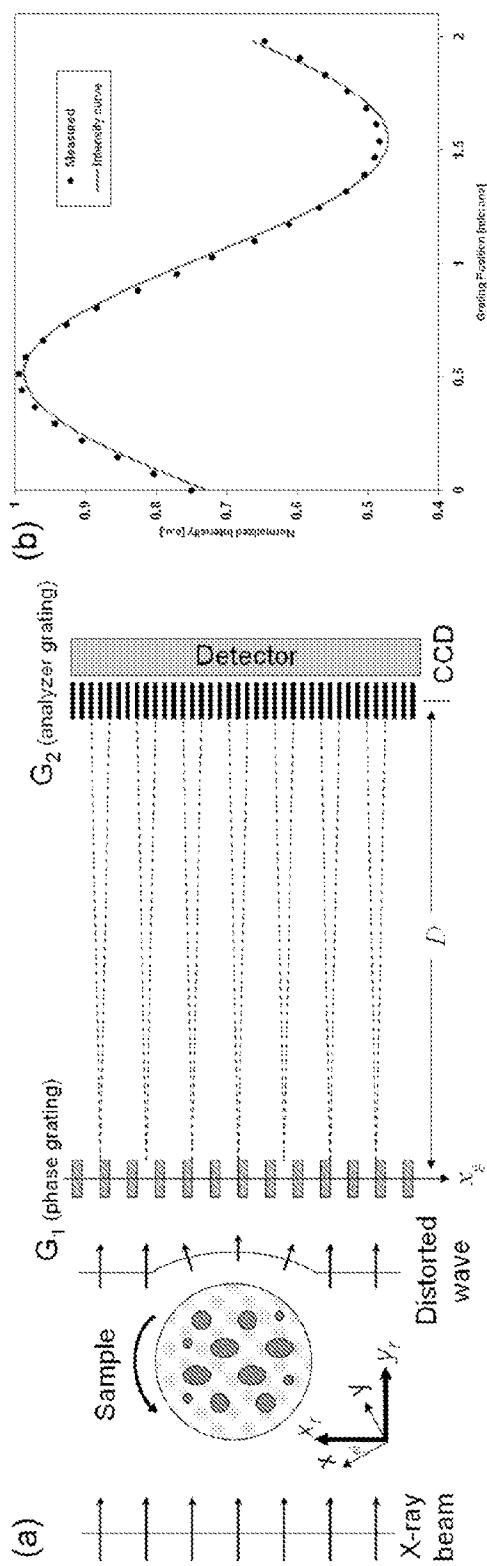


Figure 2

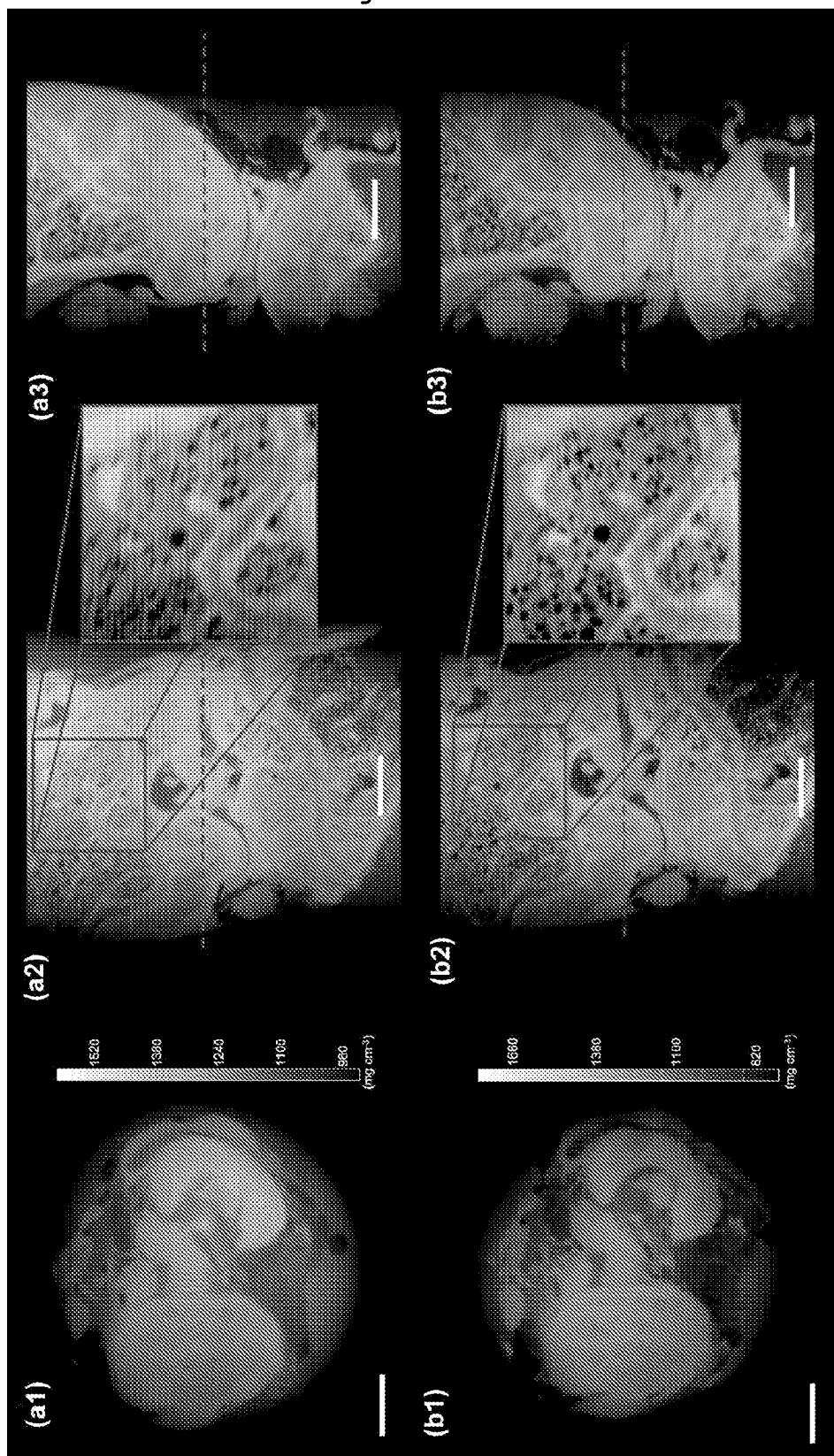


Figure 3

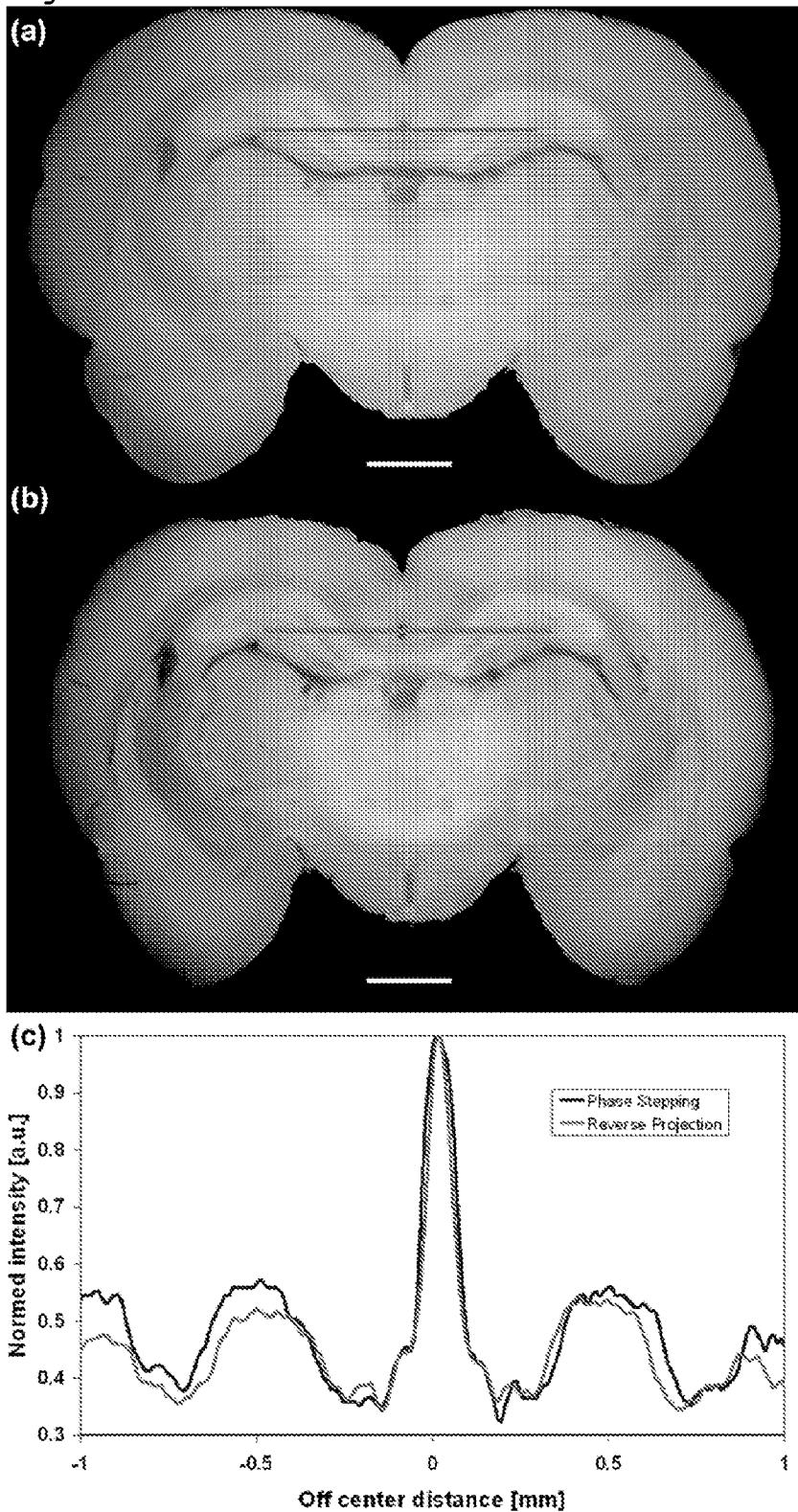
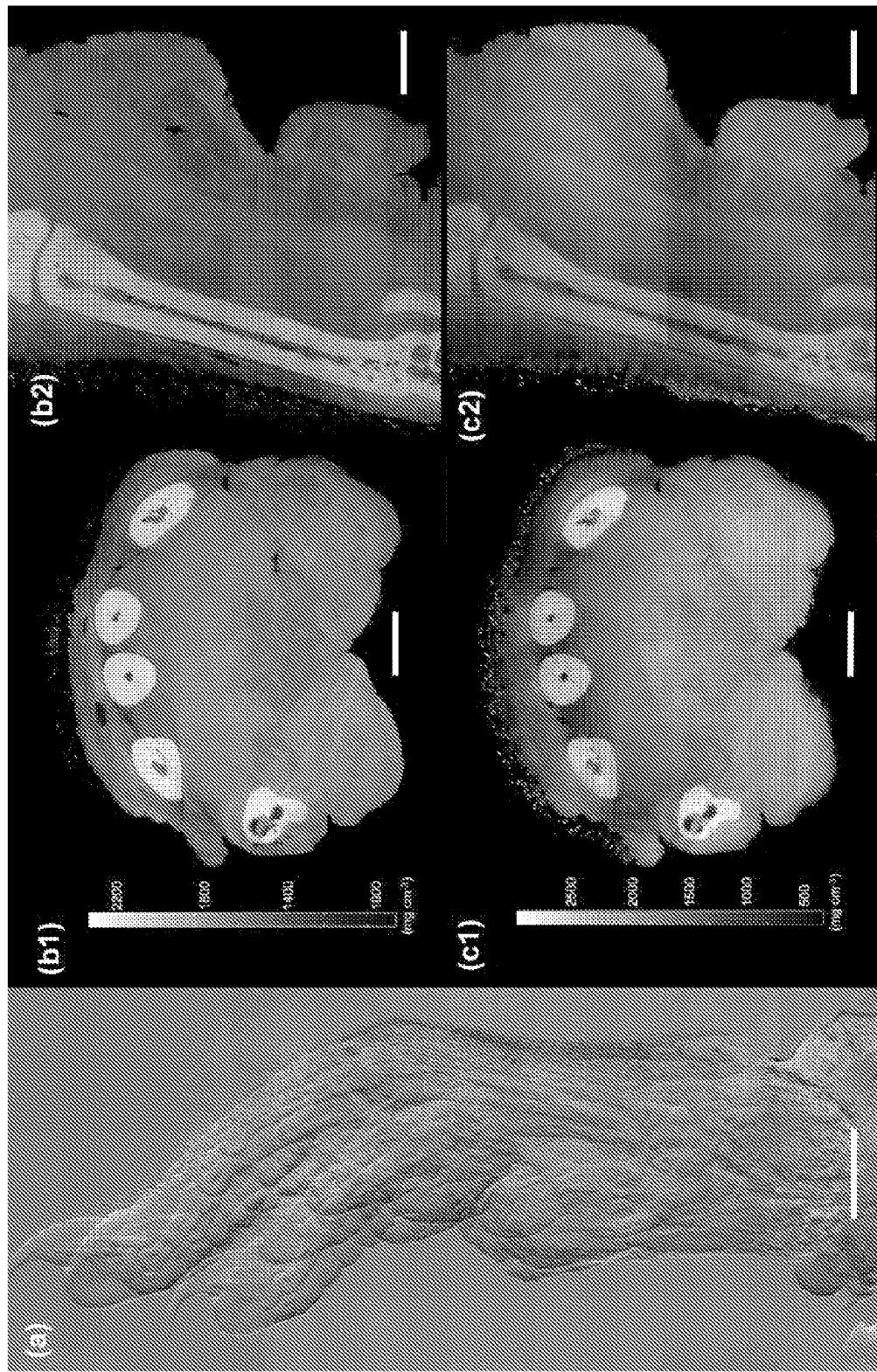


Figure 4

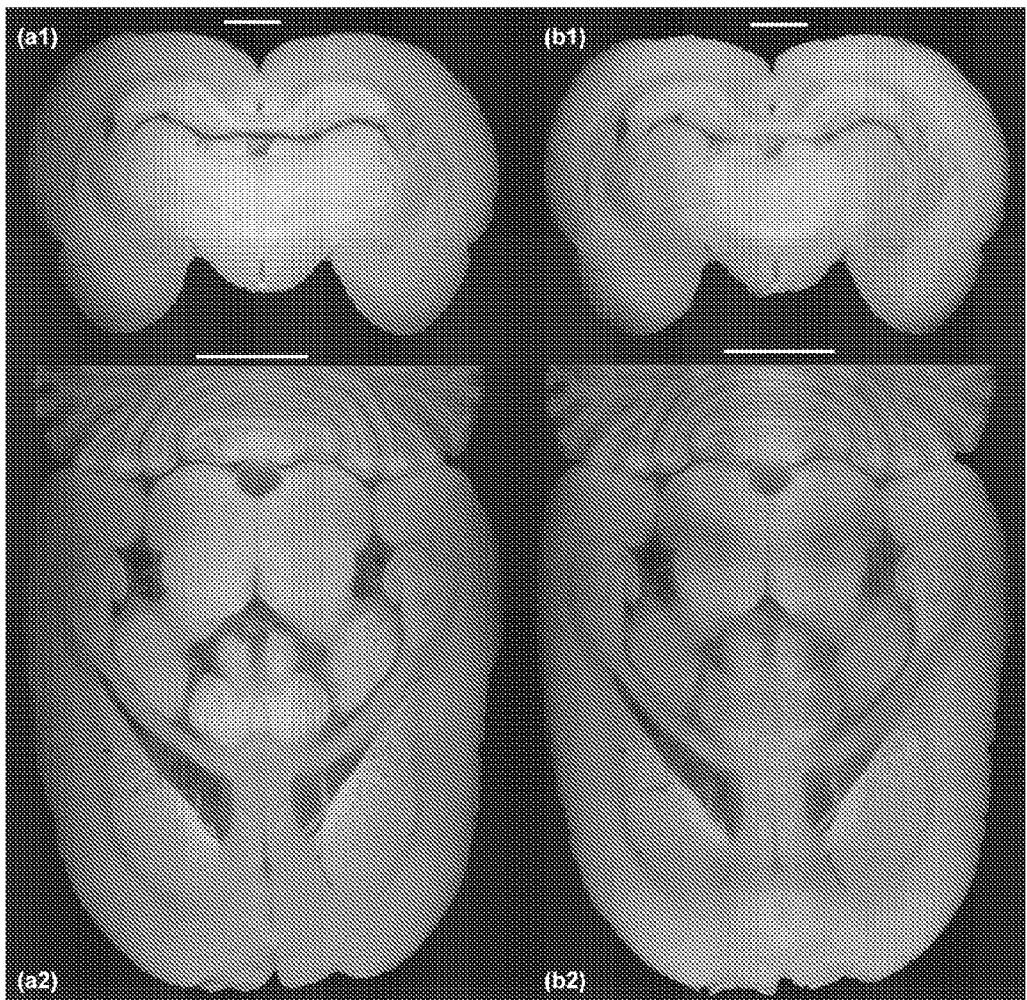


Figure 5