



# WILEY

*Publishers Since 1807*

111 River Street, Hoboken, NJ 07030

**\*\*\*IMMEDIATE RESPONSE REQUIRED\*\*\***

Your article will be published online via Wiley's EarlyView® service ([www.interscience.wiley.com](http://www.interscience.wiley.com)) shortly after receipt of corrections. EarlyView® is Wiley's online publication of individual articles in full text HTML and/or pdf format before release of the compiled print issue of the journal. Articles posted online in EarlyView® are peer-reviewed, copyedited, author corrected, and fully citable via the article DOI (for further information, visit [www.doi.org](http://www.doi.org)). EarlyView® means you benefit from the best of two worlds—fast online availability as well as traditional, issue based archiving.

Please follow these instructions to avoid delay of publication.

#### **READ PROOFS CAREFULLY**

- This will be your only chance to review these proofs. **Please note that once your corrected article is posted online, it is considered legally published, and cannot be removed from the Web site for further corrections.**
- Please note that the volume and page numbers shown on the proofs are for position only.

#### **ANSWER ALL QUERIES ON PROOFS** (Queries for you to answer are attached as the last page of your proof.)

- Mark all corrections directly on the proofs. Note that excessive alterations may ultimately result in delay of publication and extra costs may be charged to you.

#### **CHECK FIGURES AND TABLES CAREFULLY**

- Check size, numbering, and orientation of figures.
- All images in the PDF are downsampled (reduced to lower resolution and file size) to facilitate Internet delivery. These images will appear at a higher resolution and sharpness in the printed article.
- Review figure legends to ensure that they are complete.
- Check all tables. Review layout, title, and footnotes.

**RETURN PROOFS**  
**CTA (If you have not already signed one)**

**RETURN IMMEDIATELY AS YOUR ARTICLE WILL BE POSTED ONLINE SHORTLY AFTER RECEIPT.**  
**E-MAIL TO: [MRM@wiley.com](mailto:MRM@wiley.com)**

**QUESTIONS?**

**MRM Production Team**  
E-mail: [MRM@wiley.com](mailto:MRM@wiley.com)

---

Refer to journal acronym and article production number  
(i.e., MRM 00-00000).

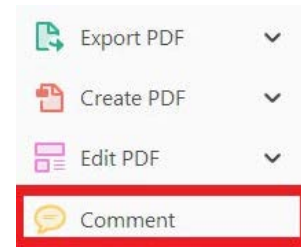
USING e-ANNOTATION TOOLS FOR ELECTRONIC PROOF CORRECTION

Required software to e-annotate PDFs: Adobe Acrobat Professional or Adobe Reader (version 11 or above). (Note that this document uses screenshots from Adobe Reader DC.)


The latest version of Acrobat Reader can be downloaded for free at: <http://get.adobe.com/reader/>

Once you have Acrobat Reader open on your computer, click on the Comment tab (right-hand panel or under the Tools menu).


This will open up a ribbon panel at the top of the document. Using a tool will place a comment in the right-hand panel. The tools you will use for annotating your proof are shown below:

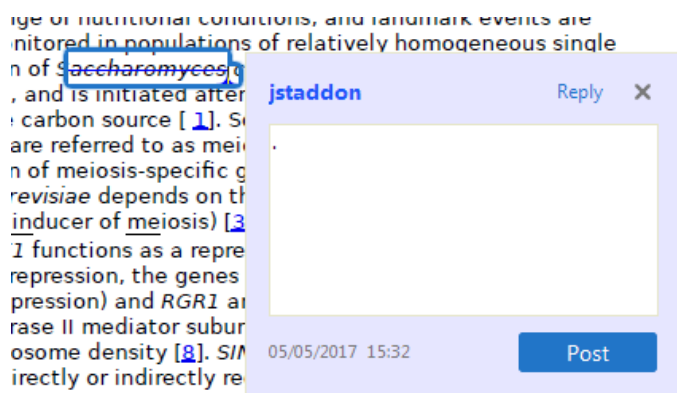


1. **Replace (Ins) Tool** – for replacing text.


 Strikes a line through text and opens up a text box where replacement text can be entered.

**How to use it:**


- Highlight a word or sentence.
- Click on .
- Type the replacement text into the blue box that appears.



2. **Strikethrough (Del) Tool** – for deleting text.

 Strikes a red line through text that is to be deleted.

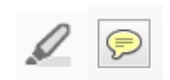
**How to use it:**

- Highlight a word or sentence.
- Click on .
- The text will be struck out in red.



experimental data if available. For ORFs to be had to meet all of the following criteria:

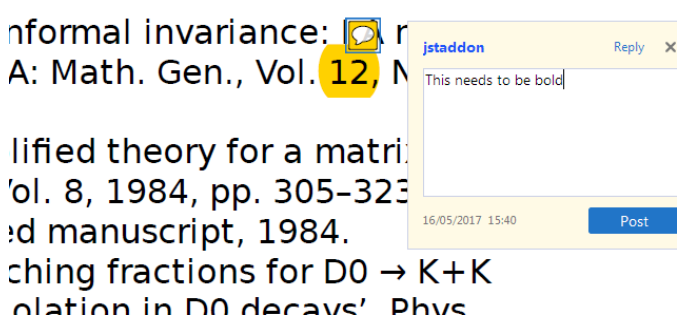
1. Small size (35-250 amino acids).
2. Absence of similarity to known proteins.
3. Absence of functional data which could not be the real overlapping gene.
4. Greater than 25% overlap at the N-terminus terminus with another coding feature; over both ends; or ORF containing a tRNA.

3. **Commenting Tool** – for highlighting a section to be changed to bold or italic or for general comments.


 Use these 2 tools to highlight the text where a comment is then made.

**How to use it:**


- Click on .
- Click and drag over the text you need to highlight for the comment you will add.
- Click on .
- Click close to the text you just highlighted.
- Type any instructions regarding the text to be altered into the box that appears.

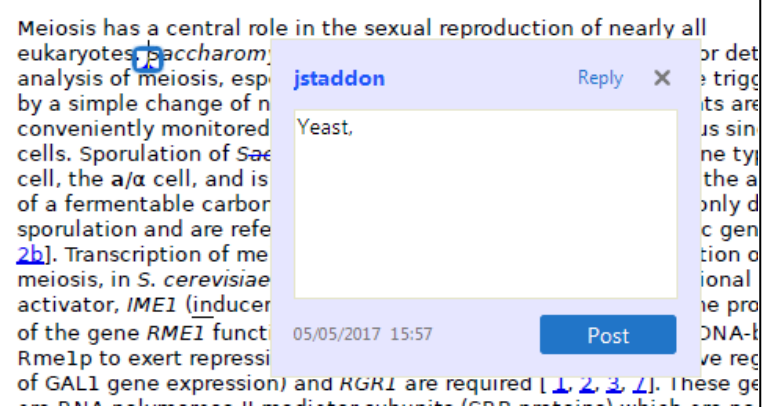


4. **Insert Tool** – for inserting missing text at specific points in the text.


 Marks an insertion point in the text and opens up a text box where comments can be entered.

**How to use it:**


- Click on .
- Click at the point in the proof where the comment should be inserted.
- Type the comment into the box that appears.



**5. Attach File Tool – for inserting large amounts of text or replacement figures.**

 Inserts an icon linking to the attached file in the appropriate place in the text.


**How to use it:**

- Click on .
- Click on the proof to where you'd like the attached file to be linked.
- Select the file to be attached from your computer or network.
- Select the colour and type of icon that will appear in the proof. Click OK.


The attachment appears in the right-hand panel.

chondrial preparator  
ative damage injury  
re extent of membra  
i, malondialdehyde (TBARS) formation.  
used by high perform

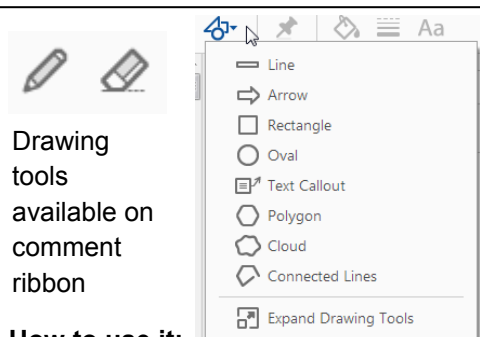
**6. Add stamp Tool – for approving a proof if no corrections are required.**

 Inserts a selected stamp onto an appropriate place in the proof.

**How to use it:**

- Click on .
- Select the stamp you want to use. (The **Approved** stamp is usually available directly in the menu that appears. Others are shown under *Dynamic*, *Sign Here*, *Standard Business*).
- Fill in any details and then click on the proof where you'd like the stamp to appear. (Where a proof is to be approved as it is, this would normally be on the first page).

of the business cycle, starting with the  
on perfect competition, constant ret  
production. In this environment goods  
extra costs should be set to zero for the  
he market. The model is determined by the model. The New-Key  
otaki (1987), has introduced produc  
general equilibrium models with nomin  
and real variables. Most of this literat

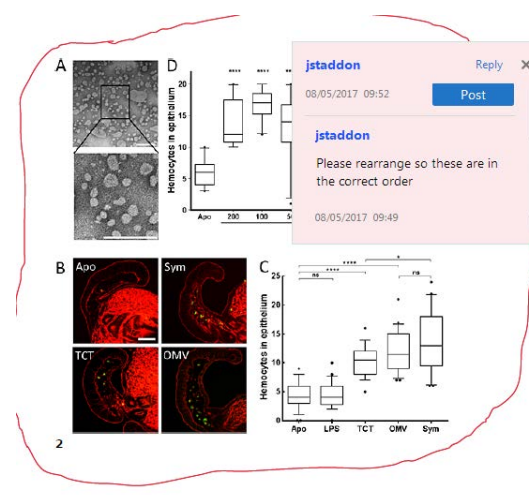


**How to use it:**

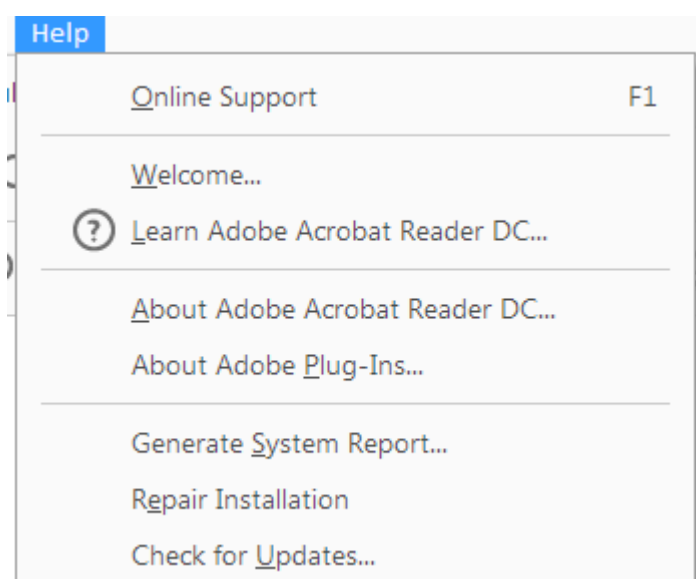
- Click on one of the shapes in the **Drawing Markups** section.
- Click on the proof at the relevant point and draw the selected shape with the cursor.
- To add a comment to the drawn shape, right-click on shape and select *Open Pop-up Note*.
- Type any text in the red box that appears.

**7. Drawing Markups Tools – for drawing shapes, lines, and freeform annotations on proofs and commenting on these marks.**

Allows shapes, lines, and freeform annotations to be drawn on proofs and for comments to be made on these marks.



For further information on how to annotate proofs, click on the **Help** menu to reveal a list of further options:



# WILEY

## **Additional reprint purchases**

Should you wish to purchase additional copies of your article, please click on the link and follow the instructions provided:

<https://caesar.sheridan.com/reprints/redirect.php?pub=10089&acro=MRM>

Corresponding authors are invited to inform their co-authors of the reprint options available.

Please note that regardless of the form in which they are acquired, reprints should not be resold, nor further disseminated in electronic form, nor deployed in part or in whole in any marketing, promotional or educational contexts without authorization from Wiley. Permissions requests should be directed to mail to: [permissionsus@wiley.com](mailto:permissionsus@wiley.com)

For information about 'Pay-Per-View and Article Select' click on the following link: [wileyonlinelibrary.com/aboutus/ppv-articleselect.html](http://wileyonlinelibrary.com/aboutus/ppv-articleselect.html)

## Author Query Form

Journal: MRM

Article: 27143

Dear Author,

During the copyediting of your manuscript the following queries arose.

Please refer to the query reference callout numbers in the page proofs and respond to each by marking the necessary comments using the PDF annotation tools.

Please remember illegible or unclear comments and corrections may delay publication.

Many thanks for your assistance.

| Query References | Query  | Remarks |
|------------------|--|---------|
| AQ1              | AUTHOR: AUTHOR: Please confirm that given names (red) and surnames/family names (green) have been identified correctly.  |         |
| AQ2              | AUTHOR: AUTHOR: Please review all affiliations and confirm or amend. Note that journal style asks that you provide site/location name, city, state/province, and country where needed for each affiliation per journal style.  |         |
| AQ3              | AUTHOR: AUTHOR: Please confirm that (1) the "Funding information" footnote is correct as set; (2) that the correspondence footnote is correct as set; and (3) if desired, please provide the corresponding author's Twitter handle (@handle).  |         |
| AQ4              | AUTHOR: AUTHOR: Would you please rework "...as by contrast to the latter it benefits from the SNR advantages arising from Fourier integration." for clarity  |         |
| AQ5              | AUTHOR: AUTHOR: Please clarify if you are referring to Figure 2 or Figure 3.   |         |
| AQ6              | AUTHOR: AUTHOR: Please rework the sentence, "While in their original single-shot implementation, both SPEN and xSPEN were affected by a progressive T <sub>2</sub> spatial weighting, phase-encoded xSPEN is devoid from such spatially-dependent contrast effects by virtue of its FT-based nature." for clarity. |         |
| AQ7              | AUTHOR: AUTHOR: Please provide manufacturer's location (i.e., city and state/country) at first mention for all manufacturer's per journal style.   |         |
| AQ8              | AUTHOR: AUTHOR: Please confirm that (1) this article contains Supporting Information and (2) the item(s) is/are cited in the correct locations.  |         |
| AQ9              | AUTHOR: AUTHOR: Please rework the sentence, "Note that because of the full sampling, the multi-scan sequence performs of the k-domain, these data corresponds to a lower resolution image, rotated in space" for clarity.  |         |
| AQ10             | AUTHOR: AUTHOR: Please define "FTd" at first mention.  |         |
| AQ11             | AUTHOR: AUTHOR: Would you please rework "...so would cropping a small length of the y' data allows one to reinstate the original one-to-one y/y' correspondence, and hence the correctness of the procedure introduced in Figures 2B and 3." for clarity   |         |
| AQ12             | AUTHOR: Author: If desired, please provide the corresponding author's Twitter handle (@handle).  |         |
| AQ13             | AUTHOR: Please confirm that given names (red) and surnames/family names (green) have been identified correctly.  |         |

## Funding Info Query Form

Please confirm that the funding sponsor list below was correctly extracted from your article: that it includes all funders and that the text has been matched to the correct FundRef Registry organization names. If a name was not found in the FundRef registry, it may be not the canonical name form or it may be a program name rather than an organization name or it may be an organization not yet included in FundRef Registry. If you know of another name form or a parent organization name for a not found item on this list below, please share that information.

| <b>FundRef name</b>  | <b>FundRef Organization Name</b> |
|--|----------------------------------|
| Israel Science Foundation  | Israel Science Foundation        |
| EU through ERC-2016-PoC  | [NOT FOUND IN FUNDREF REGISTRY]  |
| Federal German Ministry for Education and Research                 | [NOT FOUND IN FUNDREF REGISTRY]  |
| Kimmel Institute for Magnetic Resonance; Perlman Family Foundation | [NOT FOUND IN FUNDREF REGISTRY]  |
| Minerva Foundation   | Minerva Foundation               |
| Weizmann Institute of Science                                      | Weizmann Institute of Science    |
| European Commission  | European Commission              |





# Phase-encoded xSPEN: A novel high-resolution volumetric alternative to RARE MRI

Zhiyong Zhang<sup>1</sup> | Michael Lustig<sup>2</sup> | Lucio Frydman<sup>1</sup>

<sup>1</sup>Department of Chemical and Biological Physics, Weizmann Institute of Science, Rehovot, Israel

<sup>2</sup>Department of Electrical Engineering and Computer Sciences, University of California, Berkeley, California

## Correspondence

Lucio Frydman, Department of Electrical Engineering and Computer Sciences, University of California, Berkeley, CA 94720.

Email: lucio.frydman@weizmann.ac.il.

## Funding information

Israel Science Foundation, Grant/Award Number: 795/13; EU through ERC-2016-PoC, Grant/Award Number: 751106; Federal German Ministry for Education and Research, Grant/Award Number: 712277; Kimmel Institute for Magnetic Resonance; Perlman Family Foundation

**Purpose:** To develop a rapid, non-CPMG high-resolution volumetric imaging approach, exhibiting a speed and in-plane resilience to field inhomogeneities comparable to RARE/turbo-spin-echo (TSE) while endowed with unique downsampling characteristics.

**Methods:** A multi-scan extension of cross-term spatiotemporal encoding (xSPEN) is introduced and analyzed. The method simultaneously yields  $k_y/k_z$  data containing low and high frequency components, as well as transposed, low-resolution  $z/y$  images. This dual  $k$ -/spatial-domain information is captured by a multi-scan procedure that phase-encodes  $k_y$  while simultaneously slice-selecting  $z$ . A reconstruction scheme converting this information into high resolution 3D images with fully multiplexed volumetric coverage is introduced and exemplified.

**Results:** Phase-encoded xSPEN was tested by human brain imaging at sub-mm resolutions. The method exceeded 2D TSE's sensitivity by factors of  $\approx 3$ – $4$ , while providing similar resolution and SNR as 3D TSE in  $\approx 50\%$  acquisition times. The method's contrast is dominated by  $T_2$  and is free from "bright-fat" effects associated to spin-echo trains. Further acceleration is enabled by the method's downsampling abilities. Tradeoffs between encoding time, number of measurements, spatial resolution, SNR, and artifact levels are also laid out.

**Conclusion:** A new MRI strategy is introduced delivering high in- and through-plane resolutions while enjoying full Fourier multiplexing, leading to fast acquisitions with high SNR.

## KEYWORDS

cross-term spatiotemporal encoding, downsampling, high definition resolution enhancement, 3D MRI, turbo spin-echo

## 1 | INTRODUCTION

Numerous MRI approaches aim at achieving the highest possible anatomical resolution in minimal scanning times, including fast spin- and gradient-echo techniques, steady-state low flip angle acquisitions, and multi-echo/multi-gradient approaches.<sup>1</sup> The latter in particular form the basis of multi-slice 2D turbo spin-echo (TSE), routinely used in MRI exams of animals and humans as optimal compromise

between image quality and scanning time. These methods usually afford higher in- than through-plane resolutions because of their reliance on slice-selective pulses. Extensions to 3D TSE and fast spin-echo experiments can make up for this deficiency,<sup>2–4</sup> yet they are less widespread than their slice-selective counterparts because of their demand for long-echo-trains, associated with image blurring or mixed contrast. 3D TSE's high SAR also results in lengthier acquisition times. Imaging methods can be further accelerated by echo

## 2 | Magnetic Resonance in Medicine

48 planar strategies,<sup>5,6</sup> yet the relatively low bandwidth(s) of the  
49 phase-encoded dimension(s) that these experiments involve  
50 makes them susceptible to field distortions. Recent years  
51 have seen the introduction of methods that, based on a spa-  
52 tiotemporal encoding (SPEN), can deliver single-scan single-  
53 slice images with higher robustness to field inhomogeneities  
54 than their EPI counterparts.<sup>7–12</sup> Unlike what happens in con-  
55 ventional  $k$ -based scans, SPEN experiments read out their  
56 images in direct, physical space. To do so, swept pulses and  
57 gradients are used to imprint a non-linear phase during an  
58 initial encoding process, which then provides a focal point  
59 for a subsequent, gradient-driven image probing. In their  
60 original SPEN implementation this phase involved a quad-  
61 ratic,  $\phi = Cy^2$  modulation,<sup>10,13</sup> for which the signal from  
62 anywhere, but the center of the parabola is dephased; more  
63 recently, an alternative cross-term (xSPEN) procedure was  
64 introduced whereby the phase modulation takes a hyperbolic,  
65  $\phi = Cy \times z$  profile.<sup>12,14</sup> As a result of such pre-acquisition  
66 phase the signal received will be dominated by the saddle  
67 point at the center of an hyperbola. An unusual situation then  
68 arises whereby applying either  $G_a^y$  or  $G_a^z$  acquisition gradients  
69 will spatially translate this saddle point and unravel, respec-  
70 tively, the  $\rho(z)$  or the  $\rho(y)$  spatial profiles through the excited  
71 slab/slice. The xSPEN sequence on which we hereby focus  
72 relates to the second of these options; i.e., on the application  
73 of a  $z$ -axis gradient to unravel a  $\rho(y)$  profile. The fact that the  
74 exact geometry of the  $z$ -gradient plays a secondary role in  
75 unraveling  $\rho(y)$  serves as basis of xSPEN as a single-shot 2D  
76 imaging approach that avoids noticeable in-plane chemical  
77 shift or other field inhomogeneity distortions. This capability  
78 is valuable when considering single-shot 2D acquisitions  
79 in the presence of  $B_0$  heterogeneities,<sup>15</sup> known to severely  
80 distort basic functional and diffusivity studies.<sup>16,17</sup>

81 The physical basis of how the application of an acquisi-  
82 tion gradient allows one to read, in direct space and without  
83 the need for a Fourier transform (FT), the spins' profile along  
84 an orthogonal axis, was introduced in Zhang et al.<sup>12</sup> The  
85 image domain sensitive-point analysis in that study is intu-  
86 itive, but does not portray the whole story. For example, in  
87 the case of the original SPEN quadratic encoding, the result-  
88 ing spatially encoded image was blurred by a quadratic phase  
89 kernel that extended beyond the usual *sinc*-kernel of  $k$ -based  
90 encoding, requiring super-resolution/deconvolution proce-  
91 dures to achieve an optimal image resolution.<sup>11,18–23</sup> Simi-  
92 larly, xSPEN will have an associated point-spread function  
93 mixing  $y$  and  $z$  features around its hyperbolic saddle point,  
94 leading to the fact that data in the  $k_y/k_z$  plane will appear as a  
95 transposed, low-resolution  $z/y$  image. This dual spatial-/ $k$ -  
96 domain information content of xSPEN is important because  
97 it highlights certain “complementarity” in the method,  
98 whereby the application of a gradient will probe the spin  
99 density  $\rho$  along its own direction ( $k$ -domain sampling) but  
100 also reveal a profile along an orthogonal direction (spatial-

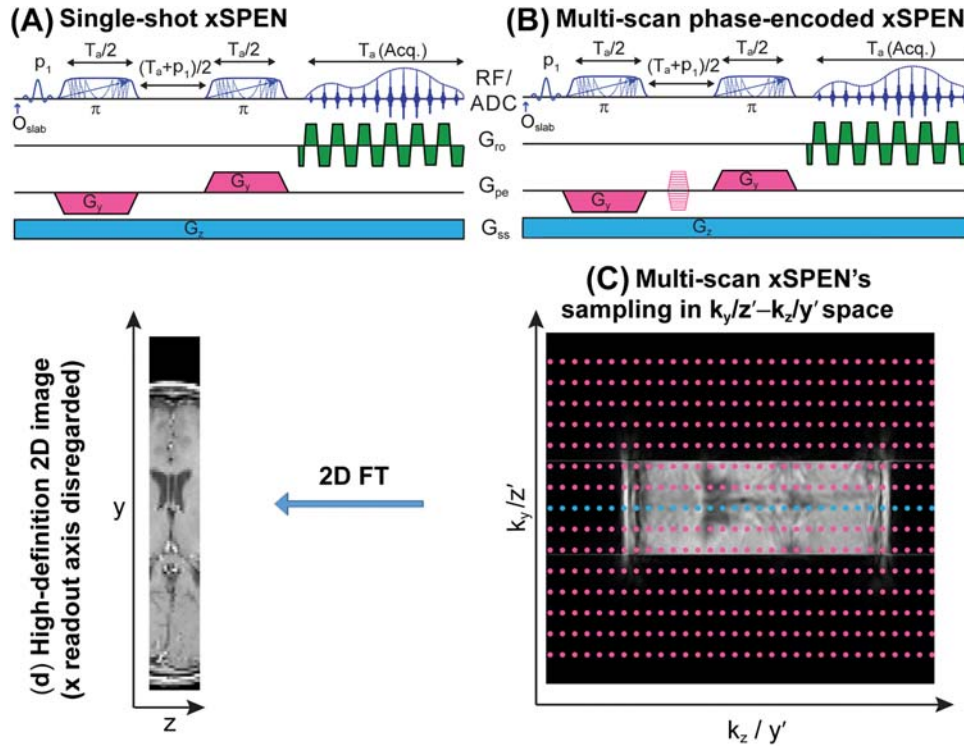
domain representation). As a result of this complementarity,  
it follows that (1) to capture image information in  $y$  or  $z$  it is  
necessary to sample a suitably dense and wide grid in  $k_z$  and  
 $k_y$ , as these variables will also act as spatially-sensitive 3D  
imaging points, (2)  $k$ -space sampling requirements can still  
be used to derive the final resolution and image sensitivity  
based on traditional Nyquist criteria, and (3) unusual rela-  
tions will emerge in terms of scanning time, resolution,  
FOV, and SNR among the  $y$ - and  $z$ -domains, particularly on  
downsampling the  $k$ -values. In this work, we further develop  
these topics. The following section introduces a multi-scan,  
phase-encoded 3D version of xSPEN and a novel reconstruc-  
tion scheme that exploits this experiment's hyperbolic phase  
encoding. We then discuss the sensitivity, FOV, resolution,  
contrast, and downsampling characteristics of the new  
method. The Results section provides a series of high-  
resolution multi-scan xSPEN imaging examples and analyzes  
their sensitivity vis-à-vis commonly used multi-slice 2D fast  
spin-echo (FSE/TSE/RARE) and multi-slab 3D fast spin-  
echo MRI techniques. We conclude with a discussion of the  
limitations and further potential of this new imaging  
technique.

## 2 | METHODS

### 2.1 | Principles of phase-encoded multi-scan xSPEN MRI.

Figure 1A illustrates the xSPEN strategy that we introduced  
for single-shot 2D acquisitions free from noticeable in-plane  
offset distortions, over  $x$  and  $y$  fields-of-view  $FOV_x$ ,  $FOV_y$ .  
This sequence applies a  $G_z$  gradient for the initial excitation  
of a slab-width  $L_z$ . Two linearly-swept adiabatic inversion  
pulses<sup>13,24</sup> of bandwidth  $BW$  and duration  $T_p = T_a/2$ —with  
 $BW = |\gamma G_z L_z| + |\gamma G_y FOV_y|$  and  $T_a$  denoting the eventual  
acquisition time—are then applied in conjunction with a  
bipolar  $\pm G_y$  gradient. This leads to the  $\Phi_e = -Cy \times z$  hyper-  
bolic phase modulation characterizing xSPEN, with  $C =$   
 $\frac{T_a \times \gamma G_y \times \gamma G_z}{\pi \times BW}$  a spatiotemporal encoding constant under the  
experimentalist's control. The saddle-shaped phase profile  
associated to this encoding leaves solely a sensitive region  
fulfilling  $\frac{\partial \phi}{\partial y} = \frac{\partial \phi}{\partial z} = 0$  where substantial cancelation among  
neighboring spins does not occur; over the course of the  
signal acquisition the continued action of  $G_z$  displaces this  
saddle-shaped profile, progressively probing  $\rho(y)$ . The  
mechanism by which the constant application of a  $G_z$  gradi-  
ent delivers, in direct physical space, an image free from  
inhomogeneities along an orthogonal axis, has been dis-  
cussed in detail elsewhere.<sup>12</sup> In brief, this behavior follows  
from calculating the time-domain signal that will be origi-  
nated in such experiment, which for a thin enough slice can  
be written as





**FIGURE 1** (A) Original single-shot xSPEN sequence mapping  $\rho(x,y')$  for a single  $z$ -slab. (B) Phase-encoded multi-scan xSPEN sequence introduced in this work. (C) Simulated  $k$ -domain data arising from the sequence in (B) along the  $(k_y, k_z)$ -domain; notice the low-resolution rendition of the brain being imaged. Blue dots indicate what would be collected in a single-shot xSPEN acquisition, corresponding to the  $y'$  profile for a particular  $z$  slice position; magenta dots indicate the fully phase-encoded xSPEN space sampled by the multi-scan sequence (B). (D) Image arising on applying a 2D FT to the signal  $S(k_y, k_z)$ , followed by a magnitude calculation. Note that because of the full sampling, the multi-scan sequence performs of the  $k$ -domain, these data corresponds to a lower resolution image, rotated in space

AQ9

$$S(k_z) \propto L_z \int_r \rho(x, y) \cdot \text{sinc} \left[ (-Cy + k_z) \frac{L_z}{2} \right] dx dy. \quad (1)$$

150 The *sinc* modulation in Eq. (1) means that the  $S$ -signal  
 151 will probe, linearly in the wavenumber  $k_z = \gamma G_z t$ , the spin  
 152 density along  $y$ . In the full single-shot 2D experiment in  
 153 Figure 1A, this 1D probing is executed in conjunction with  
 154 an oscillating  $\pm G_x$  gradient applied along an orthogonal  
 155 readout dimension, that samples the  $k_x$ -axis in an echo planar  
 156 manner. Given the conventional way in which this  $x$ -axis is  
 157 involved, all normal manipulations (partial sampling, weight-  
 158 ing, etc.) can be applied. Because of its standard nature we  
 159 ignore this dimension for most of the remaining treatment.

160 When considering the extension of these principles to  
 161 multi-scan volumetric scanning, a number of options arise.  
 162 One is multi-slicing the original single-shot, slice-selective  
 163 2D technique. A problem associated with doing so relates to  
 164 the limited resolution that one would obtain: along the  $z$  axis  
 165 this would equal the pulse-selected slice width  $L_z$ , whereas  
 166 along the  $y$  direction resolution would be given by the width  
 167 of the *sinc* in Eq. (1). This *sinc* essentially acts as the point-  
 168 spread-function of the  $y$ -axis sampling, for which it can be  
 169 shown that resolution will be given by  $FOV_y$  divided by the  
 170 time-bandwidth product  $Q = T_p \cdot BW$  of the frequency-swept

171 encoding pulses.<sup>12</sup> Therefore, unless ready to afford long  
 172 acquisition times (with concomitant diffusion- and  
 173 relaxation-driven sensitivity losses) or large excitation band-  
 174 widths (with concurrent SAR penalties) there is a limit to  
 175 how large  $Q$  can be made. One can thereby visualize  $k_z$  as  
 176 sampling a “low-resolution” rendering of  $\rho(y)$ , a  $\rho(y' = k_z/C)$   
 177 profile whose definition we would like to increase. One  
 178 possibility to do so could arise from the addition of a  
 179 conventionally looped, phase-encoding  $G_y$  gradient, into the  
 180 sequence (Figure 1B), a multi-scan phase-encoded (PE)  
 181 xSPEN variant that will be the focus of this study. FT along  
 182 the  $k_y$  dimension resulting from incrementing this phase-  
 183 encoding then provides  $FOV_y$ s and resolutions  $\Delta y$  along the  
 184 usual guidelines of Nyquist criteria. However, the bilinear  
 185 kernel  $e^{iCyz}$  that is also encoding all spins in this variant,  
 186 leads to another peculiar situation. In the same manner as it  
 187 was shown that a  $k_z$  wavenumber could read out the  $y'$  pro-  
 188 file, this 2D xSPEN scanning of a  $k_y$  axis leads to an analog  
 189 of Eq. (1)

$$S(k_z) \propto FOV_y \int_r \rho(x, y) \cdot \text{sinc} \left[ (-Cz + k_y) \frac{FOV_y}{2} \right] dx dz, \quad (2)$$

and thereby to the sampling, in direct physical space, of a  
 190 low resolution  $\rho(z' = k_y/C)$  profile. In fact the resolution of  
 191

COLOR IN ONLINE AND PRINT

192 this indirectly-sampled  $z'$  image will be given, like its direct-  
 193 domain sampling  $y'$  analog, by the time-bandwidth character-  
 194 istics of the same encoding pulses. This ability to sample  
 195 low-resolution  $y'/z'$  spatial dimensions using  $G_z/G_y$  gradients  
 196 is a consequence of the acquisition kernel imposed by the  
 197  $e^{iCyz}$  hyperbolic phase modulation: unlike what happens in  
 198 conventional MRI, where such kernel is given by a well-  
 199 localized 2D *sinc*  $k$ -domain function where orthogonal spa-  
 200 tial dimensions are decoupled, xSPEN entangles the  $y$ - and  
 201  $z$ -domain information and spreads the  $k$ -space sampling from  
 202 traditional MRI's well-localized situation, into a 2D box-like  
 203 shape. Figure 1C illustrates an unusual consequence of this  
 204 feature on a 2D human brain scan: note the “swapping” of  
 205 the sampling  $k$ -axes and of the post-Fourier spatial data, with  
 206 the former strongly resembling a rotated version of the latter.

## 207 2.2 | Processing the phase-encoded 208 xSPEN data

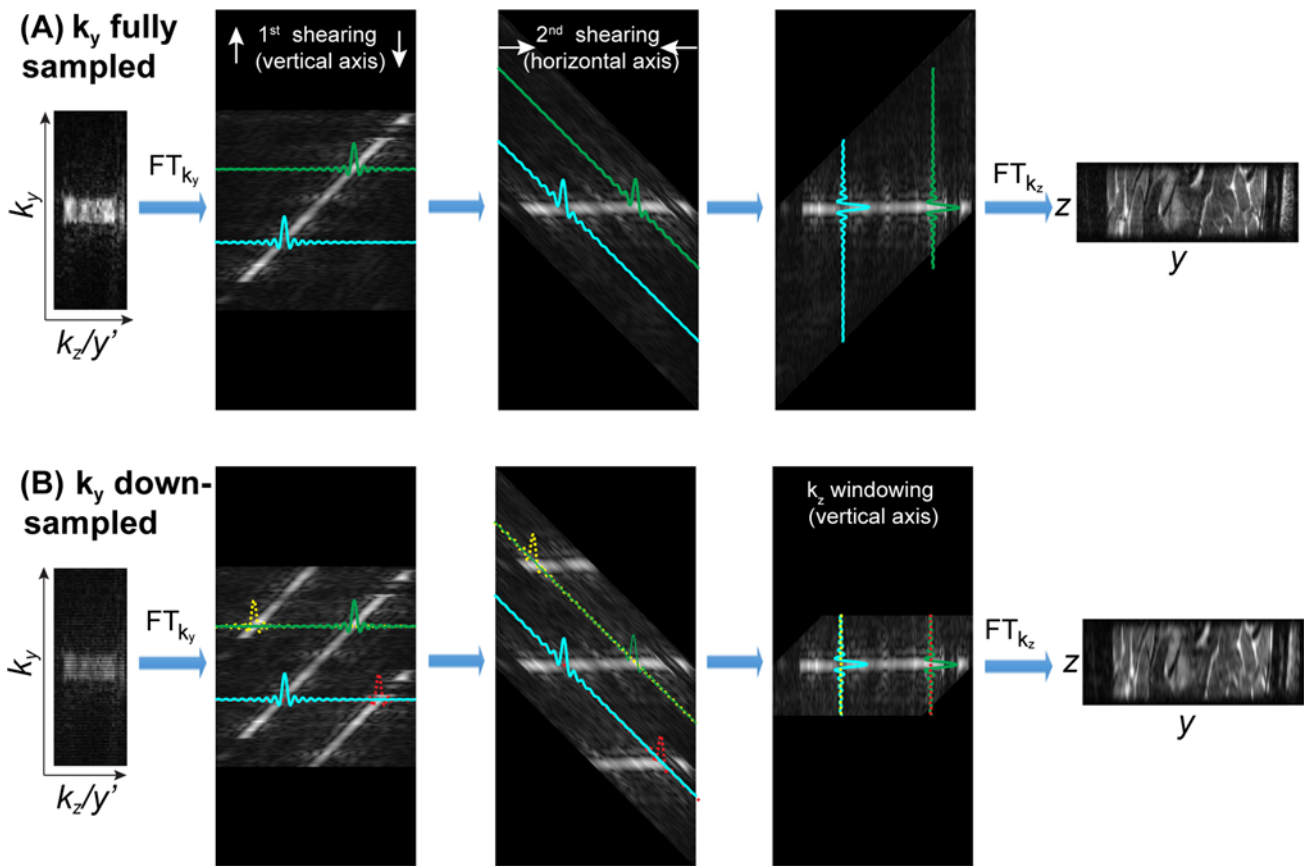
209 The fact that in this experiment the  $k$ -domain data is similar  
 210 to the spatial 2D image is reminiscent to what happened with  
 211 the original SPEN quadratic kernel.<sup>13</sup> This notwithstanding,  
 212 the fact remains that the encoding carried out by  $G_y$  and  $G_z$   
 213 in the sequence shown in Figure 1B, will also actively  
 214 sample the 2D  $k_y/k_z$ -space. Hence, aside from an overall  $e^{iCyz}$   
 215 phase factor deriving from xSPEN's hyperbolic phase modu-  
 216 lation, 2D FT of data that have been suitably sampled in the  
 217  $k_y/k_z$ -plane, will provide a conventional 2D  $\rho(y,z)$  MR image.  
 218 Figure 1D evidences this with the data arising by 2D FT  
 219 (+magnitude calculation to remove the residual hyperbolic  
 220 phase) of the low-resolution  $\rho(y' = k_z/C, z' = k_y/C)$  “image”  
 221 introduced in Figure 1C. Notice the higher resolution of the  
 222 FT-derived image both along the PE as well as along the  
 223 slice/slab-selective (SS) dimension, which is dissected into  
 224  $Q$  points by the  $G_z$  gradient. Besides a higher resolution,  
 225 this FT-originated image is also characterized by a better  
 226 sensitivity than the xSPEN-rasterized image, as by contrast  
 227 to the latter it benefits from the SNR advantages arising from  
 228 Fourier integration.

229 The redundancy that in terms of information, content  
 230 arises between the phase-encoded signal  $S(k_y)$  and the  $k_z$ -  
 231 based decoding of the  $\rho(y')$  image, opens an option to speed  
 232 up these multi-scan acquisitions. To appreciate this, consider  
 233 the lengthier sampling occurring along the PE  $k_y$  axis, whose  
 234 resolution will be given by the largest  $k_y^{max}$  value sampled,  
 235 and whose span is given by  $\Delta k_y = 1/FOV_y$ . Even disregard-  
 236 ing the possibility of accelerating experiments by parallel  
 237 MRI,<sup>25–28</sup> the fact that the full  $FOV_y$  is evaluated in a  
 238 folding-free, lower resolution fashion throughout the direct-  
 239 domain  $k_z/C = y'$  acquisition opens unusual downsampling  
 240 possibilities. Indeed, skipping  $k_y$  points will lead to folding  
 241 over among different regions throughout the  $FOV_y$  range.  
 242 The fact that this information is available along the

orthogonal  $k_z/C = y'$  axis makes it possible to unfold these 243  
 artifacts and recover the full  $y$  range—while still enjoying 244  
 the full resolution arising along this axis from the  $k_y^{max}$  sam- 245  
 pling. Although there is more than 1 avenue to perform this 246  
 kind of unfolding, Figure 2 shows an alternative that we 247  
 chose for highlighting the potential as well as the compro- 248  
 mises of this approach. Consider first a scenario involving a 249  
 full dense sampling of the relevant 2D  $k$ -space. As discussed 250  
 in Figure 1, the simplest avenue to obtain from these data a 251  
 high definition image is by performing a 2D FT. Figure 2A 252  
 shows an alternative way of arriving at the same image in a 253  
 series of steps that explicitly remove the hyperbolic phase by 254  
 a number of shearing transformations. Involved in the first 255  
 step, is a 1D FT along  $k_y$ , leading to a correlation between a 256  
 high-definition  $y$ -axis and a low-resolution image sampled 257  
 by the  $k_z/C = y'$  variable. This is represented in the second 258  
 left-most image of Figure 2A, possessing a clear diagonal 259  
 arising from the ensuing  $y/y'$  image correlation (for ease of 260  
 representation, we assume a zero-padding of the  $k_z$  axis that 261  
 places this diagonal along a unity-value slope). Colored 262  
 traces showing different  $k_z$ -domain echoes, which in this 263  
 mixed-phase interferogram will encode the spatial informa- 264  
 tion along the slice-selected domain, are also illustrated in 265  
 this experimental panel. The high-resolution  $y$ -image arising 266  
 along the  $y/y'$  diagonal can be placed along a horizontal 267  
 dimension via a shearing transformation of the 2D data set 268  
 (Figure 2A, center). Although this will result in losing the 269  
 “pure”  $k_z$ -encoding along the orthogonal axis, this can be 270  
 remedied by a second, orthogonal shearing transformation 271  
 leading to the echo disposition illustrated in the second right- 272  
 most interferogram of Figure 2A. From this data set, an addi- 273  
 tional 1D FT along  $k_z$  (now placed along the vertical axis 274  
 because of the shearing processes) will afford the high- 275  
 definition 2D image being sought. 276

277 Consider now the extension of the same procedure to a 278  
 downsampled case, where phase-encoded lines were periodi- 279  
 cally skipped for the sake of speeding up the multi-scan 280  
 acquisition (Figure 2B). Following the first FT versus  $k_y$ , a 281  
 PE that has been downsampled by a factor  $R_y$  will lead to 282  
 foldovers along the resulting  $y$  axis (Figure 2B, left, red and 283  
 yellow echoes). Notice, however, that it will incur in no reso- 284  
 lution sacrifices along this axis because  $k_y^{max}$  has not 285  
 changed. Normally such  $R_y$ -folded images cannot be disen- 286  
 tangled without additional help (i.e., from multiple receiving 287  
 coils). Additional information, however, is present in the 288  
 xSPEN acquisition, because when correlated against the  $k_z/ 289  
 C = y'$  information only the folding-free data will lie along a  
 unity diagonal. The first of the shearing transformations 290  
 introduced in Figure 2A can clearly disentangle this informa- 291  
 tion, providing the folding-free profile in the middle horizon- 292  
 tal line in the center panel of Figure 2B. A second shearing 293  
 transformation followed by a cropping out of the spurious 294

COLOR IN ONLINE AND PRINT



**FIGURE 2** Reconstruction diagram proposed for processing phase-encoded xSPEN with or without downsampling, leading to the same kind of Fourier magnitude image. (A) Reconstruction for a fully sampled data set. (B) Reconstruction of a data set that has been downsampled in  $k_y$  (notice that every other  $k_y$  line is missing in the left-most starting set). The foldovers that this originates are represented by dashed yellow and red echoes, which are removed by the indicated procedures. See the main text for further details

295 folded information and by a second FT therefore leads to a  
296 full-definition  $y$ -axis information.

297 While exploiting its additional  $k_z/y'$  information for  
298 obtaining a folding-free image despite the downsampling,  
299 Figure 2 shows that skipping PE lines will bring a penalty—  
300 not along the  $y$  but along the SS axis. At first sight, this may  
301 seem counterintuitive because this was not the downsampled  
302 axis. Notice, however, that as a result of the  $k_y$  downsampling,  
303 echoes that were overlapping along their  $k_z$ -domain  
304 will be cropped out by the processing in Figure 2B; this will  
305 reduce their support by a factor  $R_y$ , and hence lower their  $z$ -  
306 axis resolution by this same factor. The “xSPEN picture”  
307 that assigns the sampling of the  $k_y/k_z$  axes to a rasterization  
308 of the  $z/y$  domains makes this prediction, as missing elements  
309 in the former equates to skipping positions in the latter. An  
310 experimental verification of this can be appreciated by comparing  
311 the leftmost panels in Figure 2, where the loss in  $z$   
312 resolution on downsampling  $k_y$  is evident. In fact, there is  
313 another  $z$ -imaging penalty associated to the downsampling  
314 that may also lead to a special kind of  $k_z$ -domain foldover  
315 involving high frequency components aliased from non-  
316 diagonal  $y$  positions that are not eliminated by the data cropping  
317 (see color-coding of the  $k_z$  echoes in Figure 2B for a

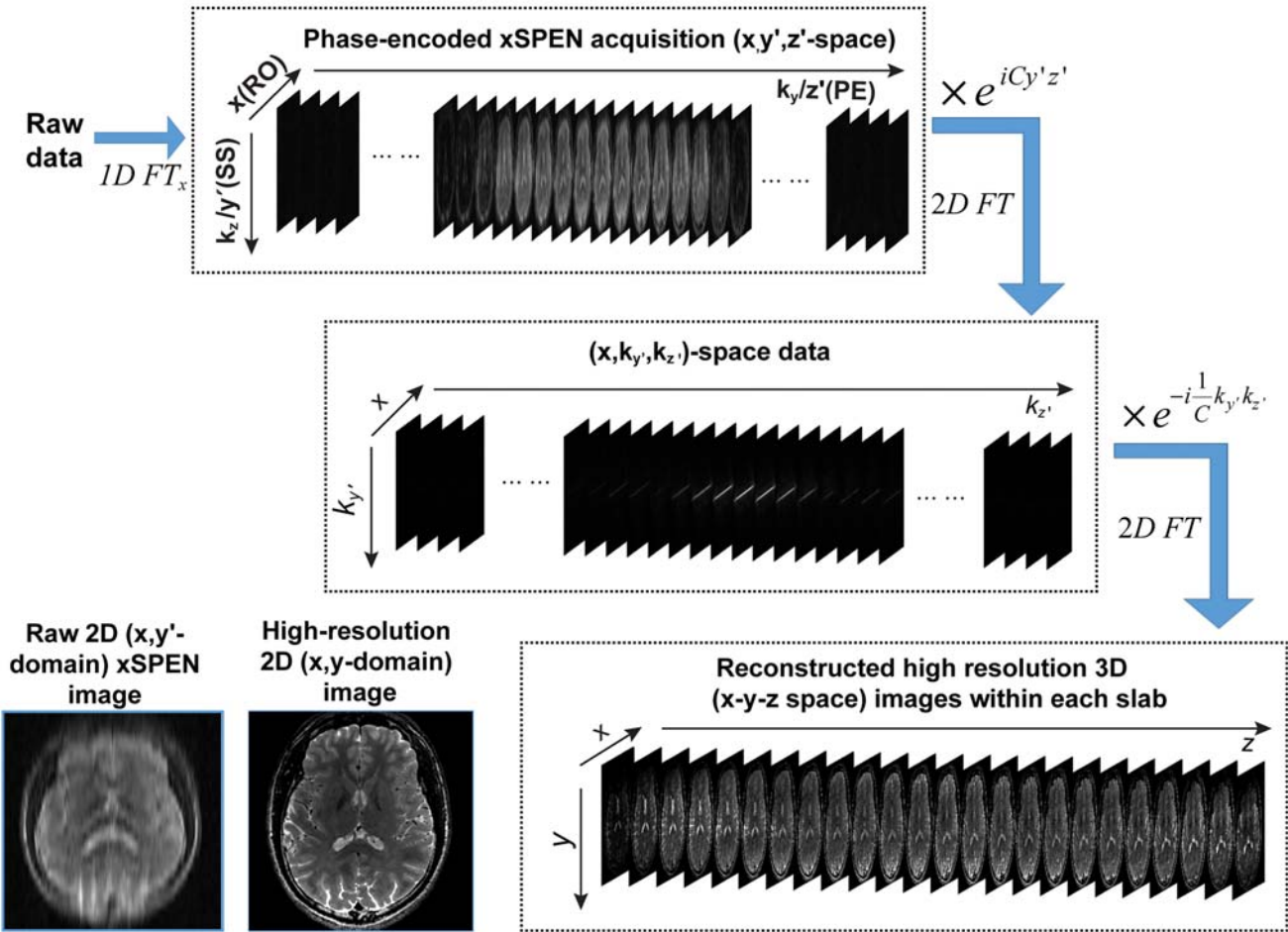
318 graphic explanation). The precise artifacts arising from this  
319  $k_z$  overlap will depend on the features within the  $z$ -domain  
320 slab and on the “tightness” of the hyperbolic encoding kernel.  
321 Specifically, if a targeted  $z$  slab is to be defined by a limited  
322 number of Fourier coefficients  $N_{slab}$ , then it can be  
323 proven that no compromises will be made as long as  
324  $N_{slab} \leq Q/R_y$ , where  $Q$  is both the aforementioned time-  
325 bandwidth product defining the hyperbolic encoding as well  
326 as the number of  $\pm$ readout echoes collected (i.e., the number  
327 of points ideally sampled along  $k_z$ ). As further illustrated  
328 below, this will be the lower limit over which downsampling  
329 can be executed while leaving a nearly lossless  
330 reconstruction.

331 Figure 3 presents additional details regarding the procedure  
332 introduced in Figure 2, this time taking into consideration  
333 the presence of the third, readout domain. As mentioned,  
334 this involves an EPI-like  $\pm k_x$  train of gradient oscillations  
335 on which a regular FT is applied to map the  $x$  profile  
336 information. The  $y'$  and  $z'$  axes associated to the  $k_z$   
337 and  $k_y$  domains are also indicated in the figure. Notice that  
338 the 2 shearing transformations introduced in Figure 2 are  
339 carried out by 2 separate phase corrections imparted in their  
340 conjugate Fourier domains—hence the mentioning of 2 2D

AQ5



Single-slab data reconstruction in multiscan PE 3D xSPEN MRI



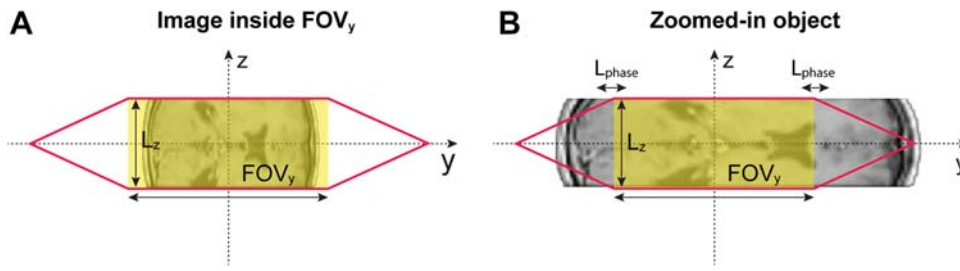
**FIGURE 3** Details of the reconstruction procedure introduced in Figure 2, illustrated for a given xSPEN phase-encoded slab acquired with a spatiotemporal encoding factor  $C$ . To obtain the ensuing high resolution images, the readout dimension (encoded by  $G_x$ ) is first FTd, and the illustrated steps are used to arrive to the final high resolution 3D image within each slab. These steps reflect the shearing procedures illustrated in Figure 2, realized here via  $C$ -dependent phase multiplications in the relevant mixed  $k/r$ -spaces. The bottom left images compare the resulting high resolution images with the low resolution counterpart arising in the ( $y' = k_z/C, x$ )-domain

341 FT steps, instead of the 1D FTs as in Figure 2. The zero padding mentioned as the initial step in Figure 2 and the windowing/cropping mentioned in the last step of that figure are omitted in Figure 3 for simplification, but they are actually required as part of the processing.

346 **2.3 | FOV and resolution in multi-scan**  
347 **xSPEN MRI.**

348 As was the case with its single-shot counterpart,<sup>12</sup> the FOVs  
349 that will be observed in multi-scan PE xSPEN will be  
350 defined by the characteristics of the encoding swept pulses.  
351 At the same time, additional FOV and resolution considerations  
352 will be dictated by the sampling and the FTs implemented on the various  $k$ -variables. Further effects will arise  
353 from potential  $k_y$  downsampling procedures. To consider all  
354 this, we neglect, as before, the conventional RO domain and  
355 focus on events defining images in the  $y/z$  plane. As

discussed in Figure 4 of Zhang et al.,<sup>12</sup> the initial SS plus the  
2 ensuing swept pulses will define, in unison with the various  
 $G_y/G_z$  gradients, a diamond-like targeted shape in this  
plane. Red polygons are shown in Figure 4 under idealized  
conditions that disregard field inhomogeneity or chemical/  
susceptibility shifts, as well as non-idealities of the adiabatic  
sweeps that are considered in more detail in the Supporting  
Information S4. In the original single-shot implementation,  
the  $y$ -axis information emerged on sampling  $k_z$ , and therefore  
“zooming in” could be carried out trivially. In the present  
case, however, different scenarios emerge depending on  
whether the object to be imaged is targeted by the  $FOV_y$ , arising  
from the  $k_y$  encoding, or it is not (Figure 4). If the imaged  
object is entirely located within the predefined  $FOV_y \times L_z$   
region (Figure 4A), all the previously derived multi-scan  
arguments will hold. By contrast, if zooming on an  
object that extends beyond the predefined  $FOV_y$  (Figure 4B),  
foldover artifacts may arise along the  $y$ -axis. This reflects the



**FIGURE 4** Two scenarios arising in a multi-scan xSPEN MRI experiment targeting a  $FOV_y \times L_z$  region (yellow box). The red polygon shows the slab defined by the RF pulses themselves,<sup>12</sup> whereas the grey images illustrate a putative object. Notice that in common imaging cases,  $FOV_y \geq 10 \times L_z$ . (A) The whole object is located within the  $FOV_y \times L_z$  region defined by the  $k_y/k_z$  sampling parameters. (B) “Zoomed-in” scan where the imaged object extends beyond the predefined  $FOV_y \times L_z$  region. In this case, a slightly larger  $FOV = L_{phase} + FOV_y$  is required for imaging the desired  $FOV_y$  without folding artifacts, where  $L_{phase} = FOV_y/Q$  is generally much smaller than  $FOV_y$  for common ( $Q \approx 20-40$ ) settings

fact that the xSPEN  $k_z/y'$  dimension will sample  $y$  information arising from object components outside the  $FOV_y$ , without folding; hence the occurrence of a  $k_y$ -driven folding would make the procedure introduced in Figure 2 inconsistent.

The  $y$ -axis folding effect will be limited by the selectivity of the spatiotemporal encoding to a spatial extent  $L_{phase} = FOV_y/Q$ , which is generally much smaller than  $FOV_y$  for commonly used ( $Q \approx 20-40$ ) settings. Expanding by an additional factor  $1/Q$ , the  $FOV_y$  extent targeted by the  $k_y$  phase encoding will solve this folding problem, alternatively, so would cropping a small length of the  $y'$  data allows one to reinstate the original one-to-one  $y/y'$  correspondence, and hence the correctness of the procedure introduced in Figures 2B and 3. Figure 4B shows that even with  $k_y$  oversampling, the profile along the  $y$  axis may be slightly affected by the non-uniform thickness of the  $z$  profile; this, however, is a negligible effect for the relatively thin slabs targeted in most imaging experiments. It is worth noting that from a practical perspective, a small oversampling along the  $k_z$  acquisition is also usually useful in dealing with non-idealities in the slab selection process.

The resolution and line shape characteristics of multi-scan xSPEN for a fully-sampled  $k_y$  acquisition can be appreciated from Figure 2A. This shows that the  $k_z$  echoes will appear shifted for different  $y$  positions, with a minimal extent of  $|k_z|$  sampling occurring for  $y=0$  and a prediction of dispersive-free line shapes and a  $\Delta z = L_z/Q$  resolution for this symmetric-echo case. For other  $y$ -values, the echoes will be asymmetrically placed thereby leading to potential dispersive components; however, given that an object's image should be real, these components can be eliminated by a complex conjugation that restores the full symmetry to the  $k_z$  echoes.<sup>29,30</sup> Further, by doing this asymmetric-to-symmetric echo conversion, the  $z$  resolution will be improved in a  $y$ -dependent fashion, ranging from the aforementioned  $\Delta z$  at the center of  $FOV_y$ , to  $\Delta z = L_z/2Q$  at the edges. Conversely, should the data be processed so that the  $k_y$  echoes appear shifted for different  $z$  positions, the  $y$  resolution would span from a minimum  $\Delta y = 1/k_y^{max}$  value at the center of the

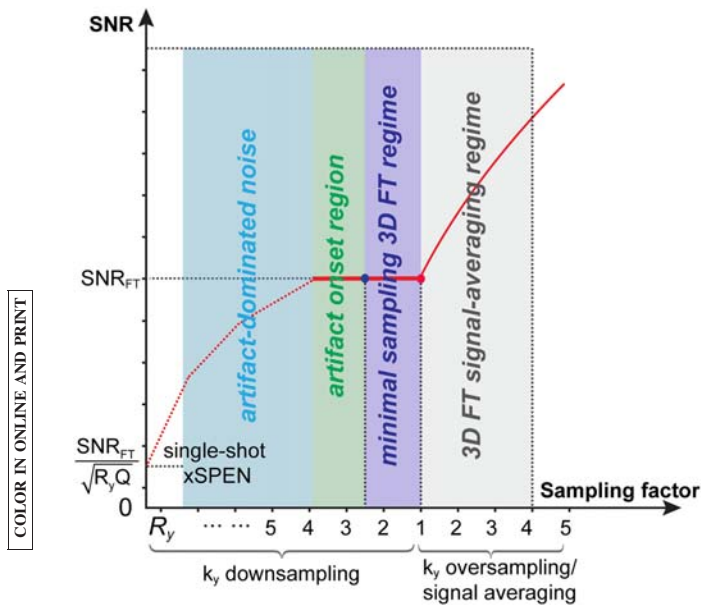
$z$ -slab, to an improved  $\Delta y = 1/(k_y^{max} + Q/FOV_y)$  value (after a complex-conjugate constrained reconstruction) at the edges of the  $z$  axis. In the case of a downsampled  $k_y$  acquisition, the unfolded  $y$  information will still be accessible from the reconstruction shown in Figure 2B. The  $y$ -axis resolution will then stay identical as with the full  $k_y$  sampling counterpart, whereas the  $z$  resolution will be decreased and potential artifacts may arise because of the aforementioned  $k_z$ -folding effects. This worsening in the  $z$  resolution will be  $y$ -independent, and artifacts will depend on the object's characteristics at other  $y$  position(s). In general, resolution losses and  $z$ -axis artifacts will be negligible if the  $z$  features change smoothly, as will usually be the case when dealing with natural objects; alternatively, spurious artifacts will arise if dealing with strong, sharp features. In most objects that we have scanned, the  $k_z$ -folding effects brought about by  $k_y$  downsampling yields images whose appearance are akin to their fully  $k_y$ -sampled counterpart, apart from a “skipping” of intra-slab  $z$ -slices. This is again in accordance with the picture of the  $k_y$  variable as sampling physical positions along the  $z$  direction: downsampling in the former axis will then lead to missing slices in the latter.

## 2.4 | SNR and contrast considerations

As shown in Figure 1, multi-scan xSPEN can, for a full  $k_y$  sampling, reconstruct a conventional  $y$ - $z$  image (aside from a phase  $e^{iCyz}$ ) by 2D FT. This magnitude image will have the same SNR as a fully 3D  $k$ -encoded FT-based counterpart ( $SNR_{FT}$  in Figure 5), because the FT implemented along xSPEN's  $k_z$  axis reinstates a multiplexing advantage that was absent on 2D multi-slice experiments collected under identical resolution and FOV conditions. Increasing the number of scans beyond this value by either repeating scans or by oversampling the phase-encoded axis, will bring about a conventional  $\sqrt{N_{scans}}$  (or  $\sqrt{N_{oversampling}}$ ) dependence on the number of averaged scans (or oversampled points). On downsampling the  $k_y$  axis, a number of scenarios will arise. As long as artifacts from the aforementioned  $k_z$  foldover effects do not

COLOR IN ONLINE AND PRINT





**FIGURE 5** Sensitivity (SNR) considerations affecting multi-scan xSPEN MRI as a function of its over/downsampling.  $SNR_{FT}$  represents the sensitivity that—if ignoring for simplicity field inhomogeneity,  $T_2$ , and diffusion-decay effects—will characterize a fully-sampled 3D FT MRI image under identical resolution and FOV conditions. See text for the meaning of the various colors and regimes

## 2.5 | Experimental

479

Human volunteers were scanned on Weizmann's 3T Siemens TrioTIM platform using a 32-channel head coil. All experiments were approved by the Internal Review Boards of the Wolfson Medical Center (Holon, Israel) and the Weizmann Institute and were collected after obtaining informed, suitably written consents. The purposes of the experiments were 2-fold: to test the various features derived above for multi-scan 3D xSPEN and to compare the performance of this sequence against alternatives that can provide comparable 3D images in terms of resolution, contrast, and sensitivity. Three sequences were chosen for this comparison, all taken from the scanner's library: 1 based on 2D slab-selective EPI, and the other 2 on turbo spin-echo (TSE) MRI. Given the mm and sub-mm resolutions being sought, the EPI sequence had to be run using multi-scan phase-encoding along the SS axis, as well as multi-shot interleaving along the low-bandwidth (PE) dimension; although these procedures worked on phantoms, they gave consistent artifacts under in vivo conditions (despite the collection and reliance on navigator scans) and hence its resulting images are not presented here. Both multi-slice 2D and phase-encoded 3D versions of TSE were successfully run at the desired resolutions and compared against home-written xSPEN acquisition sequences. In none of the experiments were parallel acquisition, fat suppression, partial FT, or simultaneous multi-slice capabilities implemented: such optimizations have not yet been fully developed for xSPEN, and hence they were left out of the comparisons. Two sets of scan parameters are presented in this study; details on one are shown in Table 1, whereas the multi-scan parameters for the second set included  $TR = 2s$ ,  $TE = 81$  ms,  $FOV_x \times FOV_y = 192 \times 192$  mm<sup>2</sup>, in-plane matrix size =  $192 \times 192$ ,  $z$ -slab thickness = 12 mm, number of slabs = 16 (i.e.,  $FOV_z = 192$  mm), xSPEN time-bandwidth product  $Q = 24$ ,  $G_z = 1.57$  mT/m,  $G_y = 0.10$  mT/m, echo train length = 29 ( $\approx 1.2 \times Q$ ; i.e., a 20% oversampling to better resolve the slices within each slab at a  $12\text{ mm}/24 = 0.5$  mm resolution), echo spacing = 1.03 ms, and an initial 15 ms long SLR  $90^\circ$  excitation pulse. The overall scan times of these experiments without downsampling was 6 min 40 s. To explore zooming-in capabilities, small FOV imaging xSPEN experiments were also recorded using the same parameters as above except that  $G_y = 0.20$  mT/m,  $FOV_y = 96$  mm, number of phase encodings = 96, and overall scan time = 3 min 20 s.

## 3 | RESULTS

523

Figure 6 illustrates high-resolution features of multi-slab, phase-encoded xSPEN MRI. These images arise from a multi-shot 3D xSPEN acquisition that covered the whole brain using an in-plane resolution of  $1 \times 1$  mm<sup>2</sup> and a

451 arise, the SNR will remain constant—even if the  $z$ -axis resolu-  
 452 tion will start to decrease owing to the  $z$ -skipping effects  
 453 mentioned above (purple region in Figure 5). At some point,  
 454 however, the  $k_y$  downsampling will start to cause an overlap  
 455 between the tails of the various  $k_z$  acquisitions, leading to  
 456 artifacts that may eventually overcome noise as the main fac-  
 457 tor controlling the image's sensitivity (Figure 5, in green).  
 458 Eventually, in the  $k_y = 0$  limit, the experiment reverts to  
 459 single-shot xSPEN; this acquisition does not attempt to  
 460 resolve anymore the intra-slab  $z$  structure, and because all  
 461 multiplexing advantages are lost the sensitivity decreases  
 462 down to a factor  $\sqrt{R_y Q}$ .

AQ6 463 While in their original single-shot implementation, both  
 464 SPEN and xSPEN were affected by a progressive  $T_2$  spatial  
 465 weighting, phase-encoded xSPEN is devoid from such  
 466 spatially-dependent contrast effects by virtue of its FT-  
 467 based nature. There will still be an overall  $T_2$ -weighting  
 468 arising from the pre-acquisition delays—but they will  
 469 affect the in-plane images uniformly. A diffusion contrast  
 470 will also affect the images arising from the  $b \approx 100$  s/mm<sup>2</sup>  
 471 weighting associated with xSPEN's encoding gradients—  
 472 yet this will also be a constant. The sole progressive effect  
 473 will be as a function of the readout echo number (i.e.,  
 474 along  $k_z$ ). This should result in an intra-slab blurring along  
 475  $z$  although this is relatively minor and was not noticed in  
 476 the experiments—probably as a result of the relatively  
 477 short duration and weak  $G_z$  gradients associated to the data  
 478 acquisition process.

**TABLE 1** Sequence parameters for 3D xSPEN, 3D TSE, and 2D TSE experiments

| Parameters                   | PE xSPEN scan 1             | PE xSPEN scan 2             | 3D TSE                      | 2D TSE                      |
|------------------------------|-----------------------------|-----------------------------|-----------------------------|-----------------------------|
| Echo spacing (ms)            | 1.03                        | 1.03                        | 9.88                        | 13.2                        |
| Echo train length            | 29                          | 29                          | 29                          | 29                          |
| TR (ms)                      | 2000                        | 4000                        | 4000                        | 8200                        |
| TE (ms)                      | 99                          | 99                          | 99                          | 92                          |
| Slice/slab thickness (mm)    | 12                          | 12                          | 12                          | 0.5                         |
| Slices/slabs ( $N$ )         | 13                          | 13                          | 13                          | 312                         |
| Volume ( $\text{mm}^3$ )     | $192 \times 192 \times 156$ | $192 \times 192 \times 156$ | $192 \times 192 \times 156$ | $192 \times 192 \times 156$ |
| Resolution ( $\text{mm}^3$ ) | $1 \times 1 \times 0.5$     | $1 \times 1 \times 0.5$     | $1 \times 1 \times 0.5$     | $1 \times 1 \times 0.5$     |
| SAR (W/kg)                   | 0.699                       | 0.349                       | 0.719                       | 0.428                       |
| Total acquisition time       | 6 min 24 s                  | 12 min 48 s                 | 16 min 54 s <sup>a</sup>    | 16 min 30 s <sup>b</sup>    |

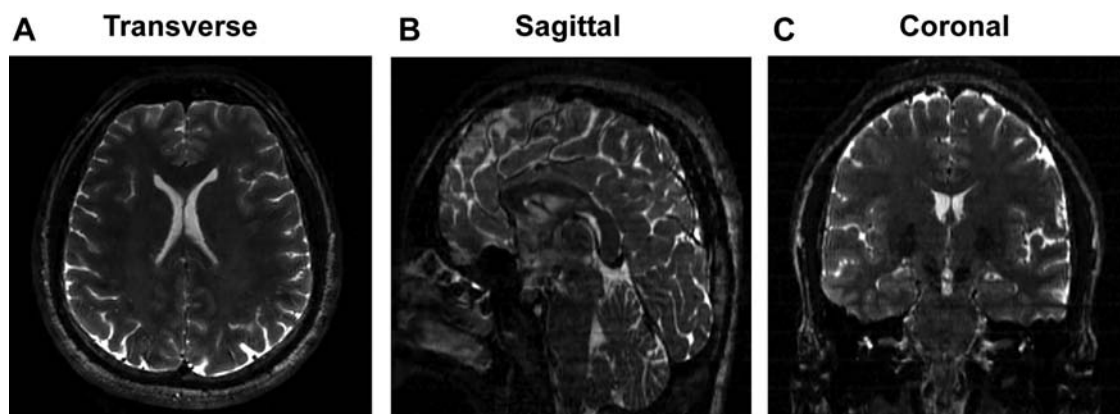
<sup>a</sup>An oversampling factor of 1.5 was applied along the slab dimension of this acquisition to avoid slice folding effects.

<sup>b</sup>Although a single slice image took  $\approx 400$  ms to collect, achieving a 156 mm coverage along the SS dimension demanded a large number of slices (312) to be acquired (in 15 concatenations) leading to a minimum TR of 8200 ms.

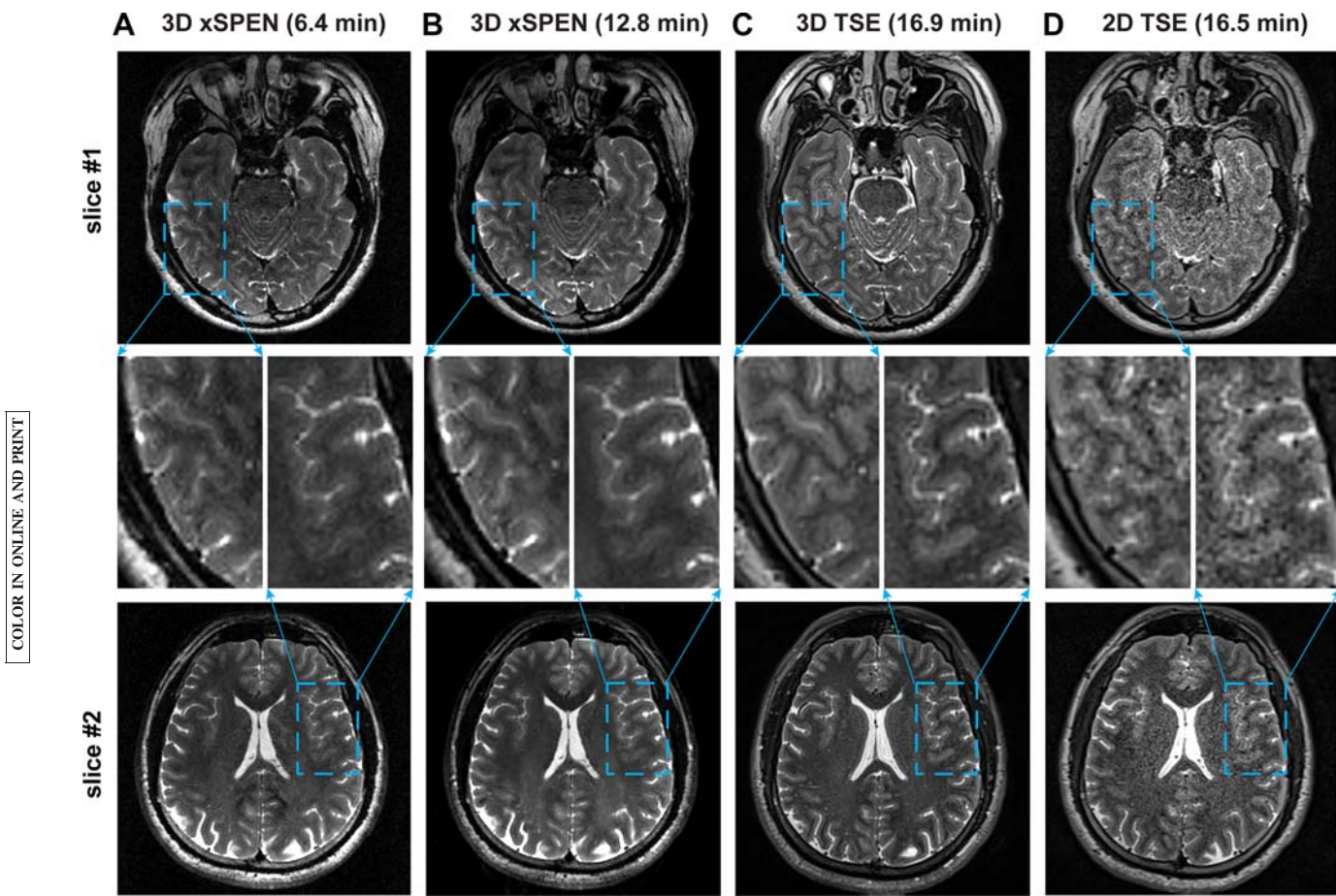
0.5 mm resolution across the slab dimension. The data were collected without downsampling, using a full  $k_x/k_y/k_z$  data set and processed according to the recipe in Figure 3. These images are dominated by a  $T_2$  contrast and lack any noticeable susceptibility or fat–water shift in-plane distortions despite the absence of multiple spin echoes owing to xSPEN’s built-in compensation characteristics.<sup>12</sup> Nearly absent as well, are “stitching effects” along the SS axis, owing to the cleanness with which the SLR slab-selective excitation and the subsequent inversion pulses addressed each of the “diamonds” introduced in Figure 4. To ensure this cleanness throughout the full  $FOV_y$ ,  $FOV_z$  extents, the Q-values of the WURST-shaped<sup>24</sup> encoding pulses were increased by  $\sim 35\%$  (see the Supporting Information S4 for further details on the slice selection). An interleaved slab acquisition procedure also ensured long TRs the absence of

noticeable  $T_1$ -weighting effects (that could be included by shortening the TRs).

Figure 7 and Table 1 compare representative performances of 3D multi-slab PE xSPEN acquisitions against 3D multi-slab PE TSE and 2D multi-slice TSE results collected under as-close-as-possible acquisition conditions. Important criteria maintained throughout these comparisons were equal FOV and resolution along all dimensions; similar  $T_2$ -based contrasts among the images were also sought. At first glance, the various whole-brain images shown in Figure 7 look similar, even if slight differences—particularly concerning the fat-derived signals—arise because of the different slab/slice excitation bandwidths of the different experiments. The most noticeable differences among both sets of images arise from “bright fat” artifacts<sup>31</sup> of the kind that usually accompany TSE experiments acquired with long echo trains, effects that



**FIGURE 6** Three-plane high-resolution images acquired using multi-slab phase-encoded xSPEN imaging. Multi-slabs were combined based on their weighted slab profiles<sup>35,36</sup> to get sagittal and coronal planes from multiple transverse acquisitions



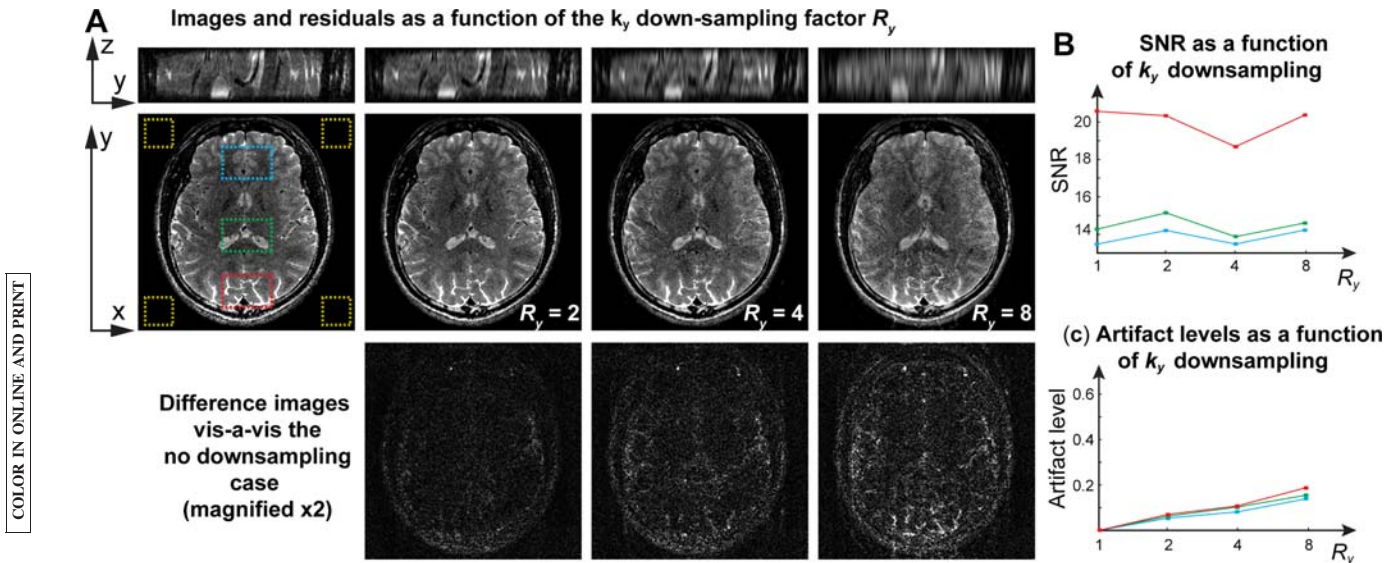
**FIGURE 7** Comparison between xSPEN and TSE acquisitions covering a volunteer’s whole brain, shown for 2 representative slices and indicating their total scan times. (A and B) Multi-slab 3D PE xSPEN images collected using TRs of 2 and 4 s, respectively. (C) Multi-slab 3D PE TSE images (TR = 4 s). (D) Multi-slice 2D TSE images (TR = 8.2 s). Zoomed images shown in the middle row arise from the light blue marked regions. No downsampling was done in any of the experiments. Further acquisition parameters are detailed in Table 1

560 are absent in the case of xSPEN. As can be appreciated from  
 561 the zoomed images in the middle panels of Figure 7, arising  
 562 from the regions marked by light-blue dashes in the whole-  
 563 brain slices, a remarkably superior sensitivity characterizes  
 564 all 3D methods over 2D TSE, thanks to their use of a FT  
 565 along the SS dimension. The longer acquisition times that  
 566 2D TSE needs to achieve the 0.5-mm slice thickness deliv-  
 567 ered by the 3D methods are also remarkable, reflecting the  
 568 large number of slices leads needed for covering the whole  
 569 brain. Moreover, although the sensitivities of 3D TSE and  
 570 3D xSPEN are comparable, the minimum TR required by  
 571 3D TSE (~3800 ms) is substantially longer than its xSPEN  
 572 counterpart (~1600 ms) for identical slab settings covering  
 573 the whole brain. This reflects the longer per-slice acquisition  
 574 times and the heavier SAR deposition required by 3D TSE  
 575 per unit time, to cover these large FOVs at these high resolu-  
 576 tions (Table 1). By contrast, xSPEN’s EPI-based acquisition  
 577 and its full  $T_2^*$ -refocused nature can achieve the same effects,  
 578 without requiring TSE’s trains of rewinding gradients and  
 579 refocusing pulses.

580 A quantitative in vivo comparison of SNR among the  
 581 various PE methods is challenged by the influence of

582 multichannel coil operation,<sup>32,33</sup> as well as because of poten-  
 583 tially different motion and diffusion effects. To account for  
 584 this, the protocol’s SNRs were measured on a NiSO<sub>4</sub>-doped  
 585 water phantom by repeating them several times identical  
 586 conditions, without relying on parallel receiving. SNRs  
 587 were then evaluated as  $mean_{r \in ROI}(S(r, \kappa_i) + S(r, \kappa_{i+1})) /$   
 588  $stddev_{r \in ROI}(S(r, \kappa_i) - S(r, \kappa_{i+1}))$ ,<sup>32</sup> where the functions  
 589  $mean()$  and  $stddev()$  return the mean value and the SD over  
 590 selected regions of interests, and the  $\kappa_i$  represent repeated  
 591 measurements. Results of these analyses are presented as  
 592 Supporting Information S1. These results validate Figure 7  
 593 and show that: (1) although an SNR advantage of  $\approx 1.28 \times$  is  
 594 expected for 3D TSE versus 3D xSPEN because of their dif-  
 595 ferent oversampling factors along the slab dimensions (1.5  
 596 vs. 1.2) and RO bandwidths (789 vs. 1042 Hz/pixel), a  
 597 slightly higher gain (1.36 $\times$ ) is experimentally observed, and  
 598 (2) although an SNR advantage of 4.7 $\times$  is expected for  
 599 multi-scan xSPEN over 2D TSE, reflecting the former’s PE  
 600 steps along slab/slice dimension (29 vs. 1) and different RO  
 601 bandwidths (1042 vs. 789 Hz/pixel), a slightly lower experi-  
 602 mental gain is observed (3.4 $\times$ ). Both of these effects can be  
 603 traced to xSPEN’s  $\exp(-b \times D)$  diffusivity losses, which for





**FIGURE 8** Results arising on downsampling the phase-encoded axis of an xSPEN acquisition. (A) Images in the y-z and x-y planes, including residuals arising on implementing xSPEN MRI with different  $k_y$  downsampling factors. The latter are shown as differences (magnified  $\times 2$ ) arising on subtracting each of the latter from the fully sampled reference. (B) SNRs arising from the blue/green/red squares indicated on the leftmost image, calculated by dividing these signals by the noise's SD arising from the yellow squares for in-plane images arising with different downsampling factors. (C) Artifact levels evaluated by taking the ratios between the signals in the various difference images, and the signals arising in the fully sampled image, focusing again on the blue/green/red squares

604 the imaging gradients used involved  $b_{xSPEN} \approx 120$  s/mm<sup>2</sup>  
 605 values and thereby lead to  $\approx 25\%$  signal attenuations beyond  
 606 the T<sub>2</sub> losses.

607 A remarkable aspect stressed in Fig. 2b for multi-scan  
 608 xSPEN MRI, was its ability to deal with  $k_y$ - downsamplings.  
 609 The folding that this will incur along the y axis can be  
 610 undone by the available low-resolution  $y' = k_z/C$  profile—at  
 611 the expense of resolution along the z axis and of eventually  
 F8 612 increasing artifacts. Figure 8 demonstrates this feature by  
 613 comparing fully  $k_y$ -sampled results with images arising from  
 614 downsampled experiments. As can be appreciated from the  
 615 y-z plane images, only a substantial downsampling of  $k_y$   
 616 brings about strong effects in the resolution along z—that ini-  
 617 tially is mostly affected by a reduction in the number of  
 618 image points. To better gauge these downsampling effects,  
 619 Figure 8 also focuses on how SNR and artifacts affect  
 620 images in the x-y plane for the same z-resolved center slice.  
 621 At first glance, all these in-plane images look similar, and  
 622 artificial structures are only visible when the downsampling  
 623 reaches  $R_y = 8$ . SNR estimations arising from different  
 624 regions in the image also lack a clear trend as a function of  
 625  $R_y$  (Figure 8B), even if artifact levels evaluated by taking dif-  
 626 ferences against the images arising from a fully sampled  
 627 experiment do evidence a monotonic increase with down-  
 628 sampling (Figure 8C). The artifacts arising in these x-y  
 629 images are dependent on the z-axis features for the various y-  
 630 positions and, in this instance, show no particularly evident  
 631 structure.

F9 632 Figure 9 demonstrates another feature of multi-scan  
 633 xSPEN MRI: its built-in ability to zoom into the phase-

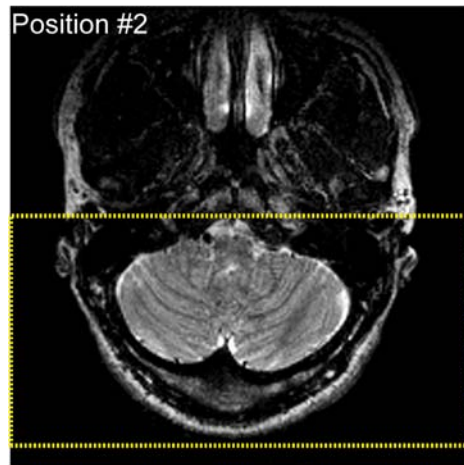
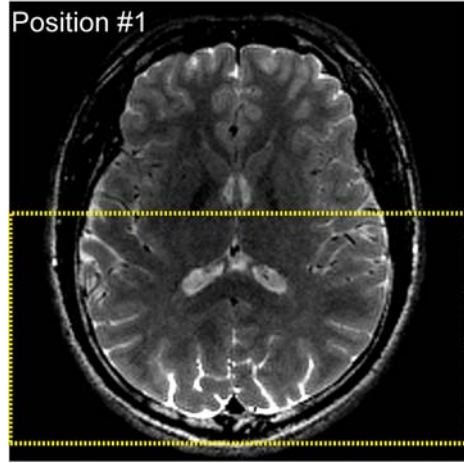
634 encoded  $FOV_y$ , without suffering from major folding. Indeed, 634  
 635 as was the case with its single-shot counterpart,<sup>12</sup> the range of 635  
 636 the latter  $FOV$  will also be dominated by the encoding chirped 636  
 637 pulses. As discussed in connection to Figure 4, however, edge 637  
 638 effects that were absent in the single-shot counterparts will 638  
 639 arise in the multi-scan phase encoding case. These effects are 639  
 640 examined in Figure 9 on slices centered on the back of the 640  
 641 brain (position 1) and on the cerebellum (position 2). Whereas 641  
 642 no artifacts arise on targeting a full  $FOV_y = 192$  mm (Figure 642  
 643 9), zooming to a half  $FOV_y = 96$  mm (yellow-dotted regions 643  
 644 in Figure 9) shows the onset of aliasing effects. These are not 644  
 645 as severe as those often seen in conventional phase encoding 645  
 646 methods and involve 2 small edge regions extending over 646  
 647  $L_{phase}/2 \approx 2$ mm.

## 4 | DISCUSSION

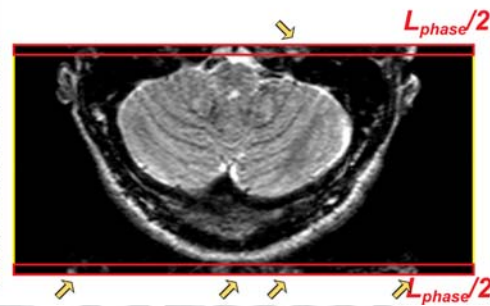
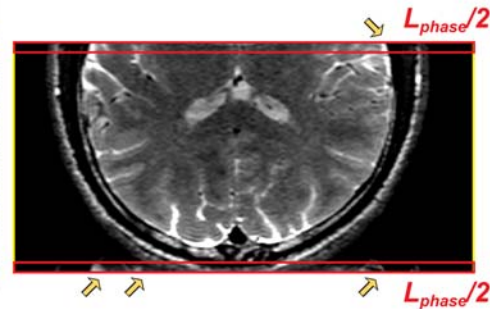
648

649 The present study explored basic features that arise when 649  
 650 extending single-shot xSPEN to multi-scan acquisitions. 650  
 651 Main rationales for implementing this extension included 651  
 652 enhancing the resolution limitations exhibited in all axes by 652  
 653 the single-shot technique, making up for the sensitivity losses 653  
 654 of the original non-Fourier method, and at the same time 654  
 655 seeking to exploit the resilience that single-shot xSPEN 655  
 656 showed to field and shift heterogeneities. A solution capable 656  
 657 of achieving these aims was devised by adding a phase- 657  
 658 encoding loop to the original implementation. Owing to the 658  
 659 pre-acquisition hyperbolic phase encoding of the spins, this 659  
 660 multi-scan xSPEN variant ended up exhibiting a dual-axis 660

**A Fully FOV<sub>y</sub> xSPEN acquisitions**



**B “Zoomed-in” (FOV<sub>y</sub> /2) xSPEN acquisitions**



COLOR IN ONLINE AND PRINT

**FIGURE 9** Restricted FOV imaging in multi-scan xSPEN MRI. (A) Reference images arising from a full FOV<sub>y</sub> (192 mm) xSPEN acquisition. (B) “Zoomed-in” images arising on halving FOV<sub>y</sub> to 96 mm (selected yellow dot colored regions in A) exemplifying the folding phenomena introduced in Figure 4. Red rectangles mark the  $L_{phase}/2$  regions, exhibiting minor foldovers marked by the yellow arrows

661 redundancy whereby  $k_z/k_y$ , not only encoded the imaging  
 662 information in reciprocal space but also provided low-  
 663 resolution  $y/z$  images in direct spatial space. The first of these  
 664 features endowed the resulting experiment with resolution  
 665 and sensitivity without relinquishing on xSPEN’s fully refo-  
 666 cused acquisition nature, while fully sampling a 3D  $k$ -space  
 667 using a single PE loop. At the same time, the redundant  
 668 information makes the experiment tolerant to downsampling  
 669 along the PE axis and hence to further acquisition speed ups.  
 670 In conventional MRI, such downsampling leads to image ali-  
 671 asing, requiring multiple receivers for its unfolding. Folding  
 672 will also occur in xSPEN, yet this can be undone thanks to  
 673 redundant information along the  $k_z/y'$  axis. PE downsampling  
 674 may still incur losses of resolution along the SS axis; as  
 675 explained above, these losses will depend on the degree of  
 676 features present in the selected slab. Interestingly, it appears  
 677 that parallel imaging procedures can also be introduced inde-  
 678 pendently from the unfolding considerations above to further  
 679 accelerate phase-encoded xSPEN MRI scans. A full charac-  
 680 terization of this behavior is under investigation.

In terms of SNR/unit\_time, multi-scan xSPEN performs 681  
 most competitively against TSE counterparts when seeking 682  
 sub-mm 3D spatial resolutions. In the specific cases pre- 683  
 sented here, we scanned the  $z$  axis by interleaving the acqui- 684  
 sition of multiple slabs, each with a 12-mm thickness. This 685  
 choice of slab thickness  $L_z$  was somewhat arbitrary and arose 686  
 as a trade-off between the resolution and SNR being sought, 687  
 the number of readout echo trains, and the tolerable 688  
 immunity to field inhomogeneity along the  $z$  axis. Indeed, it 689  
 follows from our theoretical derivations that xSPEN’s resolu- 690  
 tion along the  $z$  axis will be given by  $\Delta z = L_z/Q$ , where  $Q$  is 691  
 both the time-bandwidth product of the encoding pulses as 692  
 well as the number of elements defining resolution along  $z$ . 693  
 $Q$  generally needs to be  $\geq 15$  to keep the adiabaticity condi- 694  
 tions underlying the frequency-swept-pulse assumption; on 695  
 the other hand,  $Q$  cannot be too large because this is also the 696  
 number of RO echoes and increasing it excessively will end 697  
 up in long echo trains and diminished sensitivities.  $\Delta z$  698  
 resolution can also be manipulated by changing the  $L_z$  slab 699  
 thickness, yet this is also constrained: very thin slabs will 700



require increased  $G_z$  gradients associated to signal losses because of diffusion,<sup>12</sup> whereas thicker slabs will achieve a more efficient coverage of  $FOV_z$  yet be subject to distortions caused by field inhomogeneity acting over the course of the slice-selective excitation (that is not subject to xSPEN's full refocusing). In summary, there is a range of  $L_z$ , TE, and  $Q$  combinations that can lead to similar coverage and resolution characteristics but may have its optimal setting depending also on sensitivity and on the artifact levels that can be tolerated. Supporting Information 2 presents a range of choices that, by themselves or in combination with denoising algorithms, worked well in a variety of brain scans with voxel sizes in the 0.6–1 mm<sup>3</sup> range.

In terms of their data acquisition modules, the main difference between xSPEN and TSE acquisitions is the fact that the former uses a gradient-echo train whereas the latter uses spin-echoes. Normally the  $T_2^*$  and field heterogeneity effects in these 2 kinds of echo trains would be dramatically different; xSPEN's in-plane refocusing, however, manages to keep its acquisitions distortion-free without using RF. This, in turn, enables the use of much shorter echo spacings: in the above-mentioned examples these times were 1.03 for xSPEN versus 9.88/13.2 ms for 2D/3D TSE, with the latter 2 delays defined by the selective RF pulses, PE gradient rewinders, and crusher gradients. These shorter acquisition times free xSPEN of blurring phenomena like those arising in TSE because of  $T_2$ -related effects (see Supporting Figure S1 for an examples of this).<sup>34</sup> Interestingly,  $T_2$ -related effects affecting the signal intensity along the  $k_z$  axis may also influence the xSPEN images in the form of a spatially progressive weighting along the orthogonal  $y$ -axis. Not relying on RF pulse trains also endowed multi-scan xSPEN with ~50% lower SARs than 3D PE TSE counterparts; this enabled PE xSPEN acquisitions to use ~50% the TRs needed by 3D TSE experiments targeting similar numbers of slabs and of phase-encoding steps, leading to shorter overall volumetric acquisitions. (As for the 2D TSE scans: these should in principle have taken a shorter duration than their 3D TSE counterparts; yet to achieve the high SS definition sought while fulfilling whole brain coverage, a large number of slices had to be collected, leading again to long volumetric scan times). A final consequence arising from xSPEN's RF-free acquisitions is their lack of "bright fat" artifacts,<sup>31</sup> associated in spin echo trains to RF-driven spin-locking effects. Further comparisons on the contrast-to-noise ratios characterizing TSE and xSPEN scans in brain are discussed and illustrated in Supporting Information 3.

Another important acquisition difference arises from xSPEN's demand for a  $G_z$  acquisition gradient acting throughout its course. This results in a relatively high diffusion-related  $b$ -weighting, which makes this 3D imaging technique ~25% less sensitive than comparable 3D TSE counterparts on a per-scan basis. The  $z$ -axis imaging that this

constant gradient makes over the course of the echo planar train also demands that the slab-selection bandwidth be made equal to the inverse of the echo train spacing (i.e., to  $\Delta k_z$ ). This requirement is absent in TSE, and for long RO trains it could introduce water-fat displacement problems along the SS dimension; although not used here, fat suppression and/or partial FT<sub>RO</sub> procedures can alleviate this limitation.

A final property worth noting is xSPEN's zooming capabilities, because restricting FOVs is often a simple means to increase resolution without acquisition time overheads. In conventional phase-encoded MRI, this generally requires saturation filtering bands that increase both the minimum TR as well as the SAR. Multi-scan xSPEN MRI, by contrast, has a built-in restricted FOV capability thanks to the action of frequency-swept encoding pulses along the PE domain (Figure 9). Because of the differences arising between the imaging FOVs targeted by the  $k_y$  PE variable and the  $k_z/y'$  reading out the SS axis, edge effects arise on zooming that were absent in single-shot counterparts (Figure 4). These spatial compromises, however, are quite small, and could open interesting high-definition applications for imaging small, soft tissues organs.

## 5 | CONCLUSIONS

This study summarized the main features arising when the robustness of single-shot 2D xSPEN are coupled with the additional sensitivity and resolution arising from phase-encoded multi-scanning. From a fundamental standpoint, it is interesting to notice the similarities and differences that, owing to the hyperbolic phase encoding and xSPEN's special spatiotemporal refocusing conditions, arise between the ensuing experiment and 3D  $k$ -based MRI alternatives such as TSE. Whereas the latter were implemented without certain improvements that still need to be worked out for xSPEN counterparts—including partial FT, parallel imaging, and multi-band excitation—it was encouraging to see that in crucial parameters including sensitivity per unit time for a given FOV and resolution, the new method compared well with existing alternatives. Furthermore, the xSPEN sequence exhibits a number of features that could open yet additional opportunities. Each step in the phase-encoded procedure, for instance, entails a single-shot xSPEN acquisition providing a low-resolution 2D image; this could enable the scan-by-scan identification and elimination of motional artifacts, leading to very high definition 3D MRI capabilities. Another promise arises from the relatively short acquisition times involved in the xSPEN gradient echo train, which make it compatible with additional spin-echo combinations. These and other potential improvements of the experiments introduced here are currently under investigation.

803 **ACKNOWLEDGMENTS**

804 We are grateful to Dr. Sagit Shushan (Wolfson Medical Cen-  
805 ter) and the Weizmann MRI team (Edna Furman-Haran,  
806 Fanny Attar, and Nachum Stern) for assistance in the human  
807 scans. Z.Z. thanks Israel's Council of Higher Education and  
808 to the Koshland Foundation for partial postdoctoral fellow-  
809 ships. M.L. acknowledges the Weizmann Institute for a Visit-  
810 ing Faculty Program Fellowship. Financial support from the  
811 Israel Science Foundation (795/13), the EU through ERC-  
812 2016-PoC (751106), Minerva funding (712277) from the  
813 Federal German Ministry for Education and Research, the  
814 Kimmel Institute for Magnetic Resonance, and the generosity  
815 of the Perlman Family Foundation are also acknowledged.

816 **REFERENCES**

- 817 [1] Chavhan GB, Babyn PS, Jankharia BG, Cheng HLM, Shroff  
818 MM. Steady-state MR imaging sequences: physics, classification,  
819 and clinical applications. *Radiographics*. 2008;28:1147-1160.
- 820 [2] Oshio K, Jolesz FA, Melki PS, Mulkern RV. T2-weighted thin-  
821 section imaging with the multislab three-dimensional RARE  
822 technique. *J Magn Reson Imaging*. 1991;1:695-700.
- 823 [3] Yuan C, Schmiedl UP, Weinberger E, Krueck WR, Rand SD.  
824 Three-dimensional fast spin-echo imaging: pulse sequence and in  
825 vivo image evaluation. *J Magn Reson Imaging*. 1993;3:894-899.
- 826 [4] Mugler JP. Optimized three-dimensional fast-spin-echo MRI.  
827 *J Magn Reson Imaging*. 2014;39:745-767.
- 828 [5] Mansfield P. Multi-planar image-formation using NMR spin ech-  
829 oes. *J Phys C*. 1977;10:L55-L58.
- 830 [6] Lutti A, Thomas DL, Hutton C, Weiskopf N. High-resolution func-  
831 tional MRI at 3 T: 3D/2D echo-planar imaging with optimized physi-  
832 ological noise correction. *Magn Reson Med*. 2013;69:1657-1664.
- 833 [7] Shrot Y, Frydman L. Spatially encoded NMR and the acquisi-  
834 tion of 2D magnetic resonance images within a single scan.  
835 *J Magn Reson*. 2005;172:179-190.
- 836 [8] Tal A, Frydman L. Spatial encoding and the single-scan acquisi-  
837 tion of high definition MR images in inhomogeneous fields.  
838 *J Magn Reson*. 2006;182:179-194.
- 839 [9] Schmidt R, Frydman L. New spatiotemporal approaches for fully  
840 refocused, multislice ultrafast 2D MRI. *Magn Reson Med*. 2014;  
841 71:711-722.
- 842 [10] Chamberlain R, Park JY, Corum C, et al. RASER: a new ultra-  
843 fast magnetic resonance imaging method. *Magn Reson Med*.  
844 2007;58:794-799.
- 845 [11] Cai C, Dong J, Cai S, et al. An efficient de-convolution recon-  
846 struction method for spatiotemporal-encoding single-scan 2D  
847 MRI. *J Magn Reson*. 2013;228:136-147.
- 848 [12] Zhang Z, Seginer A, Frydman L. Single-scan MRI with excep-  
849 tional resilience to field heterogeneities. *Magn Reson Med*. 2017;  
850 77:623-634.
- 851 [13] Tal A, Frydman L. Single-scan multidimensional magnetic reso-  
852 nance. *Prog Nucl Magn Reson Spectrosc*. 2010;57:241-292.
- 853 [14] Paquin R, Pelupessy P, Bodenhausen G. Cross-encoded mag-  
854 netic resonance imaging in inhomogeneous fields. *J Magn  
855 Reson*. 2009;201:199-204.
- [15] Solomon E, Liberman G, Zhang Z, Frydman L. Diffusion 856  
MRI measurements in challenging head and brain regions 857  
via cross-term spatiotemporally encoding. *Sci Rep*. 2017;7: 858  
18010. 859
- [16] Bernstein MA, King KF, Zhou XJ. *Handbook of MRI pulse* 860  
*sequences*. Amsterdam: Academic Press; 2004. 1040 p. 861
- [17] Narsude M, Gallichan D, van der Zwaag W, Gruetter R, Mar- 862  
ques JP. Three-dimensional echo planar imaging with controlled 863  
aliasing: a sequence for high temporal resolution functional 864  
MRI. *Magn Reson Med*. 2016;75:2350-2361. 865
- [18] Ben-Eliezer N, Shrot Y, Frydman L. High-definition, single-scan 866  
2D MRI in inhomogeneous fields using spatial encoding meth- 867  
ods. *Magn Reson Imaging*. 2010;28:77-86. 868
- [19] Chen Y, Li J, Qu X, Chen L, et al. Partial Fourier transform 869  
reconstruction for single-shot MRI with linear frequency-swept 870  
excitation. *Magn Reson Med*. 2013;69:1326-1336. 871
- [20] Chen L, Bao L, Li J, Cai S, Cai C, Chen Z. An aliasing 872  
artifacts reducing approach with random undersampling for 873  
spatiotemporally encoded single-shot MRI. *J Magn Reson*. 874  
2013;237:115-124. 875
- [21] Seginer A, Schmidt R, Leftin A, Solomon E, Frydman L. 876  
Referenceless reconstruction of spatiotemporally encoded 877  
imaging data: principles and applications to real-time MRI. 878  
*Magn Reson Med*. 2014;72:1687-1695. 879
- [22] Chen L, Li J, Zhang M, et al. Super-resolved enhancing and 880  
edge deghosting (SEED) for spatiotemporally encoded single- 881  
shot MRI. *Med Image Anal*. 2015;23:1-14. 882
- [23] Ben-Eliezer N, Irani M, Frydman L. Super-resolved spatially encoded 883  
single-scan 2D MRI. *Magn Reson Med*. 2010;63:1594-1600. 884
- [24] Garwood M, DelaBarre L. The return of the frequency sweep: 885  
designing adiabatic pulses for contemporary NMR. *J Magn* 886  
*Reson*. 2001;153:155-177. 887
- [25] Deshmane A, Gulani V, Griswold MA, Seiberlich N. Parallel 888  
MR imaging. *J Magn Reson Imaging*. 2012;36:55-72. 889
- [26] Griswold MA, Jakob PM, Heidemann RM, et al. Generalized 890  
autocalibrating partially parallel acquisitions (GRAPPA). *Magn* 891  
*Reson Med*. 2002;47:1202-1210. 892
- [27] Uecker M, Lai P, Murphy MJ, et al. ESPIRiT—an eigenvalue 893  
approach to autocalibrating parallel MRI: where SENSE meets 894  
GRAPPA. *Magn Reson Med*. 2014;71:990-1001. 895
- [28] Pruessmann KP, Weiger M, Scheidegger MB, Boesiger P. 896  
SENSE: sensitivity encoding for fast MRI. *Magn Reson Med*. 897  
1999;42:952-962. 898
- [29] Haacke EM, Lidskogj ED, Lin W. A fast, iterative, partial- 899  
fourier technique capable of local phase recovery. *J Magn* 900  
*Reson*. 1991;92:126-145. 901
- [30] MacFall JR, Pelc NJ, Vavrek RM. Correction of spatially 902  
dependent phase shifts for partial Fourier imaging. *Magn Reson* 903  
*Imaging*. 1988;6:143-155. 904
- [31] Hardy PA, Henkelman RM, Bishop JE, Poon ECS, Plewes DB. 905  
Why fat is bright in rare and fast spin-echo imaging. *J Magn* 906  
*Reson Imaging*. 1992;2:533-540. 907
- [32] Dietrich O, Raya JG, Reeder SB, Reiser MF, Schoenberg SO. 908  
Measurement of signal-to-noise ratios in MR images: influence 909  
of multichannel coils, parallel imaging, and reconstruction filters. 910  
*J Magn Reson Imaging*. 2007;26:375-385. 911

- 912 [33] Kellman P. Image reconstruction in SNR units: a general method  
913 for SNR measurement. *Magn Reson Med.* 2007;58:211-212.
- 914 [34] Tamir JI, Uecker M, Chen W, Lai P, Alley MT, Vasanawala SS,  
915 Lustig M. T2 shuffling: Sharp, multicontrast, volumetric fast  
916 spin-echo imaging. *Magn Reson Med.* 2017;77:180-195.
- 917 [35] Wu W, Koopmans PJ, Frost R, Miller KL. Reducing slab  
918 boundary artifacts in three-dimensional multislab diffusion MRI  
919 using nonlinear inversion for slab profile encoding (NPEN).  
920 *Magn Reson Med.* 2016;76:1183-1195.
- 921 [36] Parker DL, Yuan C, Blatter DD. MR angiography by multiple  
922 thin slab 3D acquisition. *Magn Reson Med.* 1991;17:434-451.

## 923 SUPPORTING INFORMATION

AQ8 924 Additional Supporting Information may be found in the  
925 online version of this article.

926  
927 **FIGURE S1** SNR performance of the phase-encoded  
928 xSPEN and TSE acquisitions. (A and B) PE xSPEN  
929 images collected using TR = 2 and TR = 4 s, respectively.  
930 (C) Multi-slab 3D TSE images. (D) Multi-slice 2D TSE  
931 images. Yellow dot marks the ROIs on which SNR was  
932 computed, whereas red arrows show the T<sub>2</sub> blurring effects  
933 in TSE experiments. Average SNRs are listing on the bot-  
934 tom of each column. Experimental parameters are detailed  
935 in Table 1 of the main text.

936 **FIGURE S2** (A) Raw image with a resolution of  
937  $1 \times 1 \times 0.5$  (thickness) mm<sup>3</sup> reconstructed from the phase-  
938 encoded xSPEN. (B) Denoised counterpart image arising

939 on using the Block-matching and 4D filtering (BM4D)  
940 algorithm. Zoomed images from the red dotted regions of  
941 (A) and (B) are shown in the middle column.

942 **FIGURE S3** Contrast in PE xSPEN and in 3D TSE  
943 acquisitions (A) PE xSPEN result using TR = 2 s and cov-  
944 ering 11 slabs. (B) Multi-slab 3D TSE result using TR = 2  
945 s and covering 3 slabs. (C) Multi-slab 3D TSE using  
946 TR = 4 s and covering 11 slabs. All slab thicknesses were  
947 12 mm.

948 **FIGURE S4** CNR values calculated between the 2 indi-  
949 cated voxels in each image. CNR was defined as  $|mean(tis-$   
950  $sue1) - mean(tissue2)|/std(noise)$ , where for each image  
951 the 2 red boxes represent the tissue1, tissue2 signals  
952 whereas the yellow boxes were used for calculating the SD  
953 of the noise.

954 **FIGURE S5** Boundary artifacts arising in multiscan  
955 xSPEN if not pushing transition areas outside the targeted  
956 region of interests (right) and explanation of why using  
957 larger  $G_y$  gradients and concurrently large RF sweeping  
958 bandwidths will alleviate these problematic edge effects  
959 without affecting the slab chosen along the  $z$  axis

961 **How to cite this article:** Zhang Z, Lustig M, Frydman  
962 L. Phase-encoded xSPEN: A novel high-resolution vol-  
963 umetric alternative to RARE MRI. *Magn. Reson. Med.*  
964 2018;00:1–15. <https://doi.org/10.1002/mrm.27143>  
965

Author Proof