

A Simulation Study of the Flexible TWIST View Sharing Impact on the Breast DCE MRI

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Target Audience Radiologists, MRI physicists and scientists.

Introduction TWIST (Time-resolved angiography With Stochastic Trajectories) view-sharing has received increasing attention in studies of breast DCE-MRI¹⁻⁵. Simulation studies is very helpful in finding out how accurate the enhancement curve/pattern is represented in images acquired with under-sampling techniques such as TWIST, and the most appropriate under-sampling strategy for certain types of application⁶⁻⁸. This study aims to provide such information for the optimization and error estimations in breast DCE-MRI with TWIST.

Methods A digital 'phantom' of $36 \times 36 \times 13 \text{ cm}^3$ ($448 \times 448 \times 162$) was generated with three spherical uniform 'lesions' of 5 mm diameter with typical 'persistent', 'plateau' and 'wash-out' type of enhancement respectively, and one composite lesion of 10mm diameter with a mixture of three types of enhancement (Fig. 1), all embedded in non-enhancing 'breast tissue'⁹. A modified TWIST technique with more flexible view sharing (Fig.2) was simulated. The simulation method is similar to a previously published kidney DCE-MRI study⁸. TR = 5.6 ms, GRAPPA factor = 2, phase/slice resolution= 80%/ 70%; partial Fourier in frequency/phase/slice direction = 80%. K-space views were shared from the nearest available time point while backward sharing was preferred. The percentage of the central k-space region size was A = 10%, 20%, 30%, 40%, 50% and 100%; the peripheral region k-space update rate was B=10%, 12.5%, 20%, 25%, 33% and 50%. Each set of images was reconstructed to $448 \times 448 \times 162$. To calculate the error, all measured enhancement curves were interpolated onto the same grid of time points. Average RMS error of the enhancement curve: $RMS_{all} = \frac{1}{N} \sum_{n=1}^N (1/T \sqrt{\sum_{t=1}^T (enhancement_{measured}(t) - enhancement_{true}(t))^2})$ (N being the number of voxel inside the lesion), and the RMS error at the peak of the wash-out curve:

$RMS_{peak} = (1/N \sqrt{\sum_{n=1}^N (enhancement_{measured}(t) - enhancement_{true}(t))^2})$ were measured for each lesion.



Figure 1. The middle section of the composite lesion with three partially overlapping regions with (a) persistent; (b) plateau; (c) wash-out enhancement. The whole lesion (d), simulated acquisition with A =100% (e) and simulated acquisition with A = 10% and B = 10% (f) at the peak of wash-out curve.

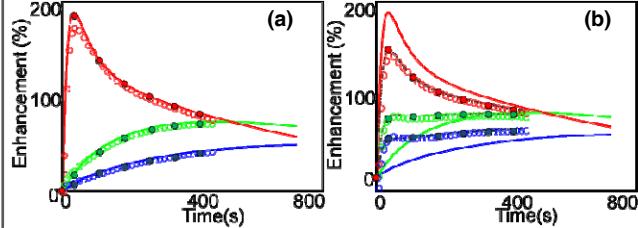


Figure 3. Measured enhancement in (a) three 5 mm spherical lesions with persistent, plateau and wash-out type of enhancement, and (b) the persistent, plateau and wash-out part of the 10 mm composite lesion. Circles are acquired with A = 10% and B = 10%, solid points are with A = 100%. Solid lines show the 'true' enhancement. Dash lines in (b) show the theoretical weighted average enhancement, which almost overlapped with the measured data.

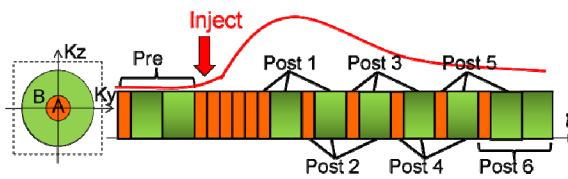


Figure 2. A full k-space ($AB_1B_2..B_n$) pre-contrast acquisition , a series of (A) only before the peak of enhancement, several (AB_i) after the peak of enhancement and another full k-space acquisition at the end (In this Figure we have 6 post-contrast as an example).

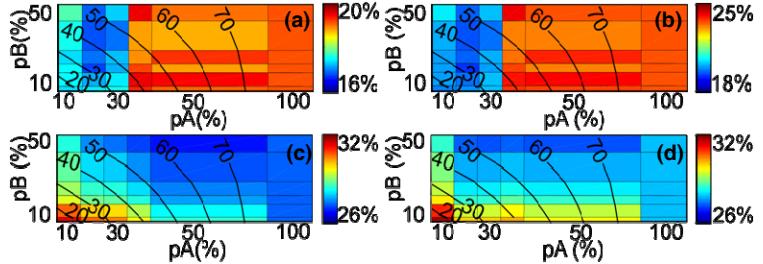


Figure 4. RMS_{all} of (a) composite and (b) 5 mm wash-out lesion both reached a minimum at A = 20% and B = 20%; while the RMS_{peak} in (c) composite and (d) 5 mm wash-out lesion have a lower error region with high A and B (A = 50%, B=50%). Contours show the time needed for each partial acquisition (unit: s).

Results Fig.1e and 1f show a decrease of spatial resolution with A=10% compared with A=100%. A decrease of the peak enhancement value on the wash-out curve with A=10% and B=10% is shown in Fig. 3, while the measured curves for 'plateau' and 'persistent' types of enhancement were almost identical between different As and Bs. The RMS error in the 5 mm lesion is slightly higher than that of the composite lesion. RMS_{all} was the lowest with A =20% and B =20% (Fig. 4 a-b), while the RMS_{peak} was the lowest with A=50% and B = 50% (Fig. 4 c-d).

Discussions In this study we assumed that the first TWIST partial acquisition (AB_i) is timed accurately at the peak of the washout curve. The error may increase if the peak cannot be accurately determined. The representation of the wash-out type of enhancement and the enhancement in the composite tumor were impacted more by the selection of A and B than that for the persistent and plateau type of enhancement. Multiple local minima and maxima of the error exist in the tested range for A and B; which may be caused by the difference in the temporal foot print for the acquisition of each A or B region along the curve. RMS_{all} and RMS_{peak} allow assessment of the accuracy of the enhancement kinetics and morphological patterns, therefore can be used to guide the selection of optimal under-sampling parameters such as A and B.

Reference

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