POSTER PRESENTATION

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Detection of myocardial oedema with the use of diffusion-weighted imaging in acute myocardial infarction

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Introduction

MR diffusion-weighted imaging is an important application for oedema detection in various tissues. Evaluation of the area at risk in reperfused acute myocardial infarction (AMI) is currently performed with STIR T2-weighted and LGE sequences.

Purpose

The aim of the study was to find practical application for a new diffusion-weighted sequences in evaluation myocardial oedema and compare it with routinely used STIR-T2 techniques.

Methods

In preliminary study myocardial oedema in 15 patients (13 male) with STEMI within 2-4 days were evaluated . The CMR examination was performed on a 1,5 T scanner (Magnetom Avanto; Siemens; Germany; Erlangen;) using a 8-channel phased-array coil. The parameters of the diffusion-weighted EPI sequence (DWI) were as follows: slice thickness 10mm, repetition time (depending on patient breath cycle) 3-4s, echo time 78ms, bandwidth 1,736 Hz/Px. The DW sequence was ECG-gated and synchronized to the respiratory cycle using PACE technique. Each slice was acquired with b = 50 s/mm2, 400s/mm2 and 800 s/mm2 with three perpendicular directions of the diffusion gradient. DW, STIR T2-weighted and LGE images were obtained in 2-chamber, 4-chamber or short-axis planes. Images were analysed quantitatively, contrast to noise ratio (CNR) of high signal (oedema) to healthy myocardium (CNR1)

Figure 1 Myocardial oedema in 2 chamber view evaluated by A) STIR T2-weighted image; B) DW EPI sequence.

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and high signal to blood (CNR2) were calculated. For statistical analysis a non parametric Wilcoxon test with significance level of p<0,05 was used.

Results

The CNR were significantly higher on DWI than on STIR T2-weighted images: CNR1 (22 ± 7 vs 12 ± 8 p= 0,004, respectively)and CNR2 (28 ± 10 vs 21 ± 9 , p=0,02, respectively).

Conclusions

Our study confirms DW EPI is feasible sequence for myocardial oedema detection with even better contrast to noise ratio than standard STIR T2 sequences.

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