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Correction: Myocardial native T₁ mapping and extracellular volume quantification in asymptomatic female carriers of Duchenne muscular dystrophy gene mutations



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Following publication of the original article [1], we have been notified of several mistakes.

The authors' names were initially published as per below:

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They should be as follows:

Lucia Masárová1,2, Roman Panovský1,2*,Martin Pešl1,2,3, Mary Luz Mojica-Pisciotti1,2, Tomáš Holeček1,5,7, Vladimír Kincl1,2, Lenka Juříková6, Jan Máchal1,4, Lukáš Opatřil1,2 and Věra Feitová1,5

Results section text was as follows (bold and incorrect spacing):

The mean global native T1 relaxation time was similar for DMD-FC and CG (1005.1 ± 26.3 ms vs. 1003.5 ± 25 ms; p-value=0.81) (Fig. 1), as well as the mean global ECV value ($27.92\pm2.02\%$ vs. $27.09\pm2.89\%$; p-value=0.20) (Fig. 2). The representative native and post-gadolinium T1 maps are presented in Fig. 3f or **DMD-FC** and Fig. 4f or **healthy volunteer.**

The text should be corrected as follows:

The mean global native T1 relaxation time was similar for DMD-FC and CG (1005.1 ± 26.3 ms vs. 1003.5 ± 25 ms; p-value=0.81) (Fig. 1), as well as the mean global ECV value ($27.92\pm2.02\%$ vs. $27.09\pm2.89\%$; p-value=0.20) (Fig. 2). The representative native and post-gadolinium T1 maps are presented in Fig. 3 for DMD-FC and Fig. 4 for healthy volunteer.



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Figure 3 was incorrect and with incorrect caption as per below:

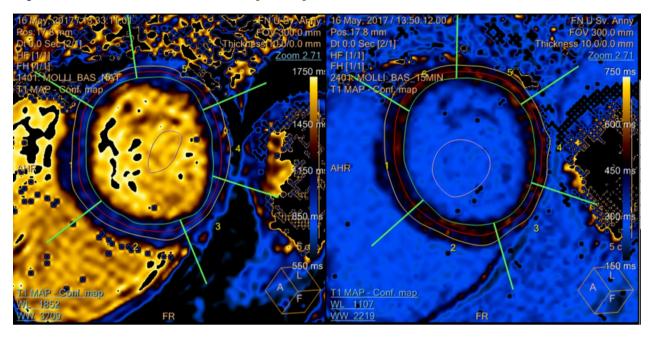


Fig. 3 A representative picture of the native and post-gadolinium T_1 map of DMD-FC. DMD-FC- Female carriers of Duchenne muscular dystrophy gene mutations

Figure 3 and its caption should be as follows:

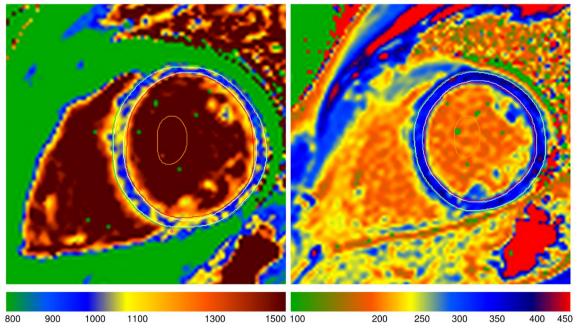


Fig. 3 A representative picture of the native and post-gadolinium T_1 map of DMD-FC. DMD-FC—Female carriers of Duchenne muscular dystrophy gene mutations

Figure 4 was incorrect, and it is shown below:

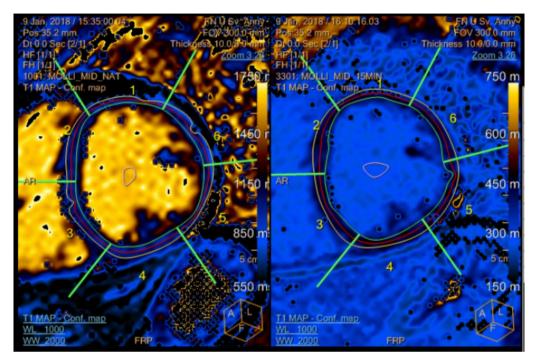


Fig. 4 A representative picture of the native and post-gadolinium T_1 map of a healthy volunteer

Figure 4 should be as follows:

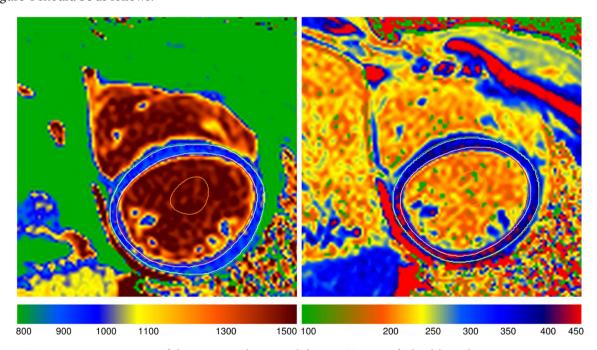


Fig. 4 A representative picture of the native and post-gadolinium T_1 map of a healthy volunteer CMR acquisition section last paragraph had some incorrect spacing, it was as follows:

 T_1 mapping was performed as described previously [17] using a Modified Look-Locker Inversion recovery sequence (MOLLI) with a 5(3) 3 scheme to measure native T_1 (pre-contrast) and a 4(1) 32 scheme for T_1 postgadolinium (15 min after contrast agent administration).

CMR acquisition section last paragraph text should be as follows:

 T_1 mapping was performed as described previously [17] using a Modified Look-Locker Inversion recovery sequence (MOLLI) with a 5(3)3 scheme to measure native T_1 (pre-contrast) and a 4(1)3(1)2 scheme for T_1 post-gadolinium (15 min after contrast agent administration).

The original article was updated.

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Reference

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