CORRECTION



Correction to: A comparison of high-throughput plasma NMR protocols for comparative untargeted metabolomics

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Correction to: Metabolomics (2020) 16:64 https://doi.org/10.1007/s11306-020-01686-y

Following publication of the original article, the authors would like to correct a sentence in the paragraph "¹H-NMR spectra were recorded at 298 K…" under the heading "NMR experiments".

The sentence currently reads:

"The LED pulse sequence had the form -RD-901-G1-1801-G1-901-G2-T-901-G1-1801-G1-901-G2-t-901-acquire FID, where RD is a relaxation delay, 901 is a 901 RF pulse, G1 is the pulsed-field gradient that is applied to allow editing, 1801 is a 1801 RF pulse, G2 is a spoil gradient applied to remove unwanted magnetization components. The diffusion delay D is the time during which the molecules are allowed to diffuse—this is the period (901-G1-1801-G1-901-G2-T-); and t is a delay to allow the longitudinal eddy currents caused within the sample to decay (Beckonert et al. 2007)."

The sentence should read:

The original article can be found online at https://doi.org/10.1007/s11306-020-01686-y.

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"The LED pulse sequence had the form -RD-90°-G1-180°-G1-90°-G2-T-90°-G1-180°-G1-90°-G2- τ -90°-acquire FID, where RD is a relaxation delay, 90° is a 90° RF pulse, G1 is the pulsed-field gradient that is applied to allow editing, 180° is a 180° RF pulse, G2 is a spoil gradient applied to remove unwanted magnetization components. The diffusion delay Δ is the time during which the molecules are allowed to diffuse—this is the period (90°-G1-180°-G1-90°-G2 T-); and τ is a delay to allow the longitudinal eddy currents caused within the sample to decay (Beckonert et al. 2007)."

This has been corrected with this erratum.

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