

NMR-Based Metabolomics of the Lipid Fraction of Organic and Conventional Bovine Milk

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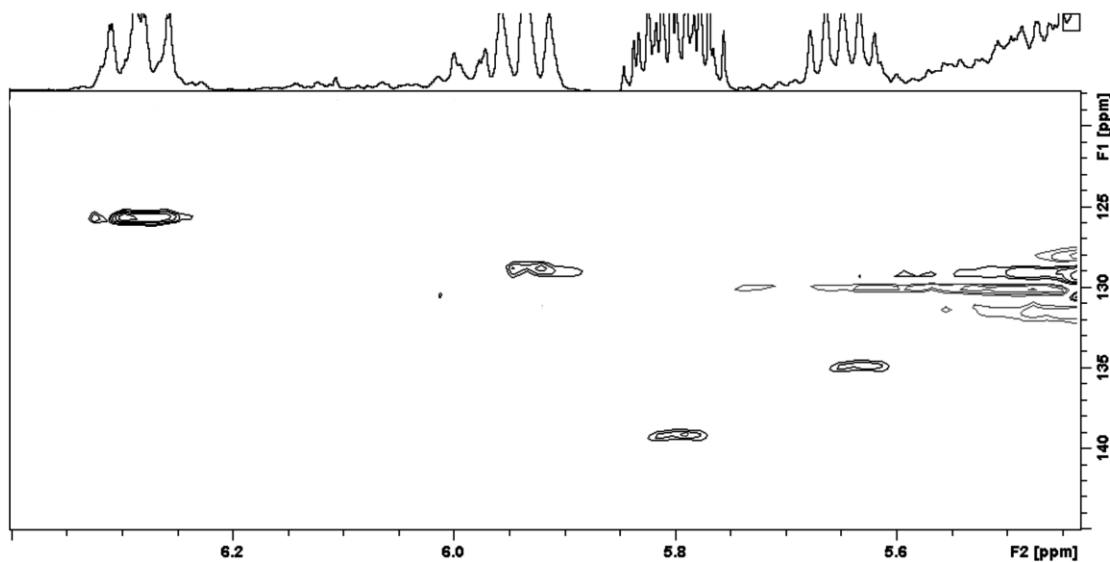


Figure S1. Selected region of 500 MHz ^1H - ^{13}C HSQC spectrum of the lipid fraction of a lyophilized bovine milk sample in CDCl_3 illustrating ^1H - ^{13}C connectivities of conjugated (9-*cis*,11-*trans*)18:2 linoleic acid (CLA) and caproic acid; T, 298 K; 40 repetitions of 256 increments, total experimental time 4 h 35 min.

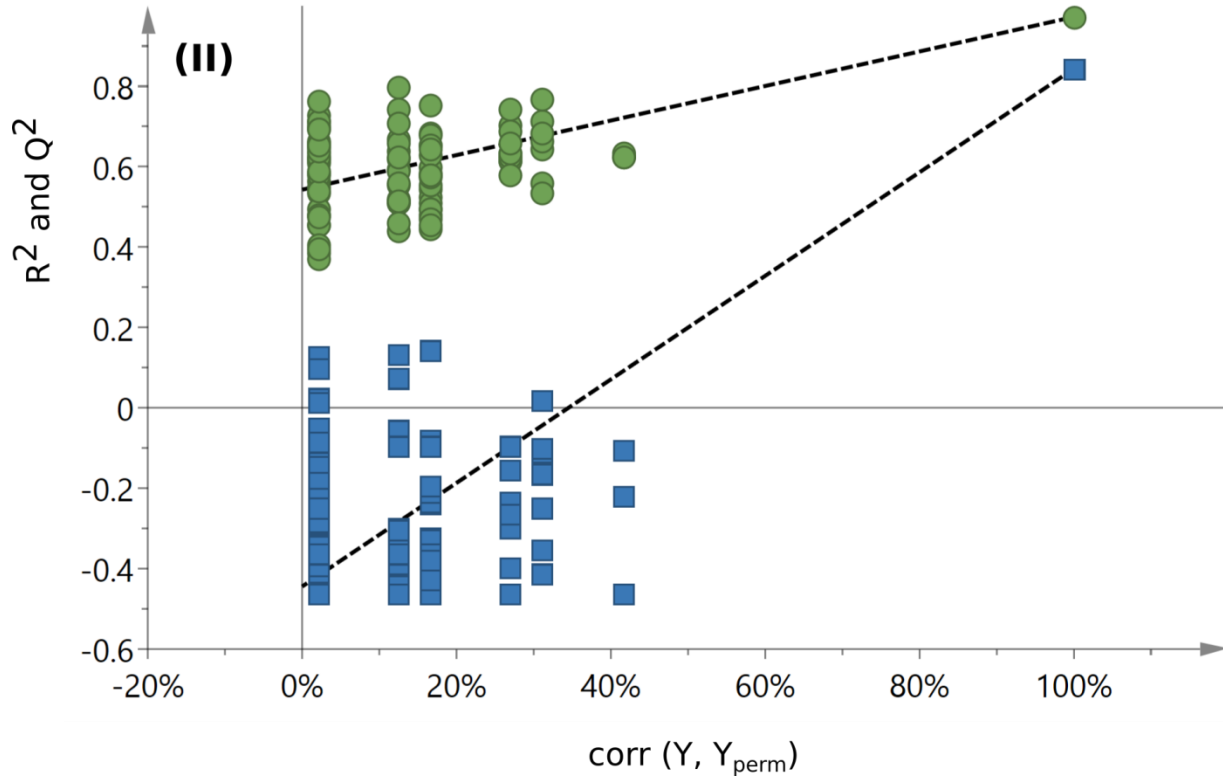
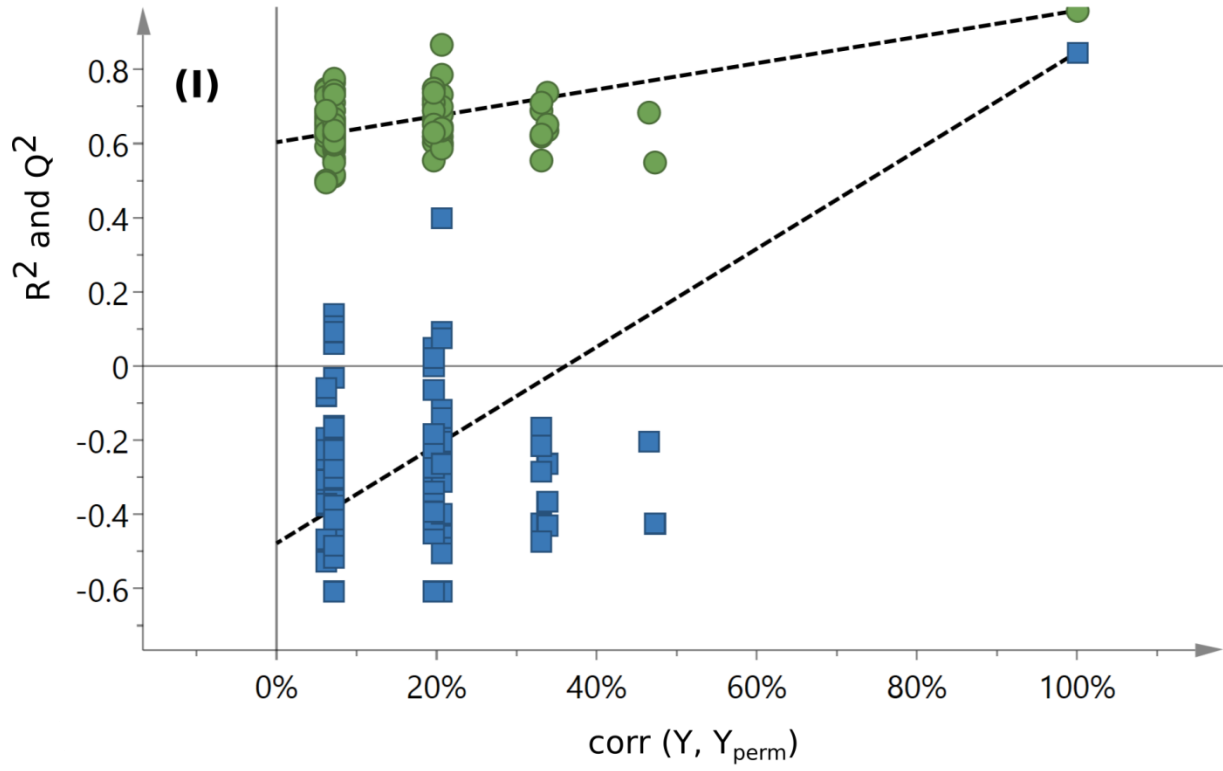


Figure S2. Permutation test of the P samples with PLS-DA: (I) UNor method, (II) NorCont method.

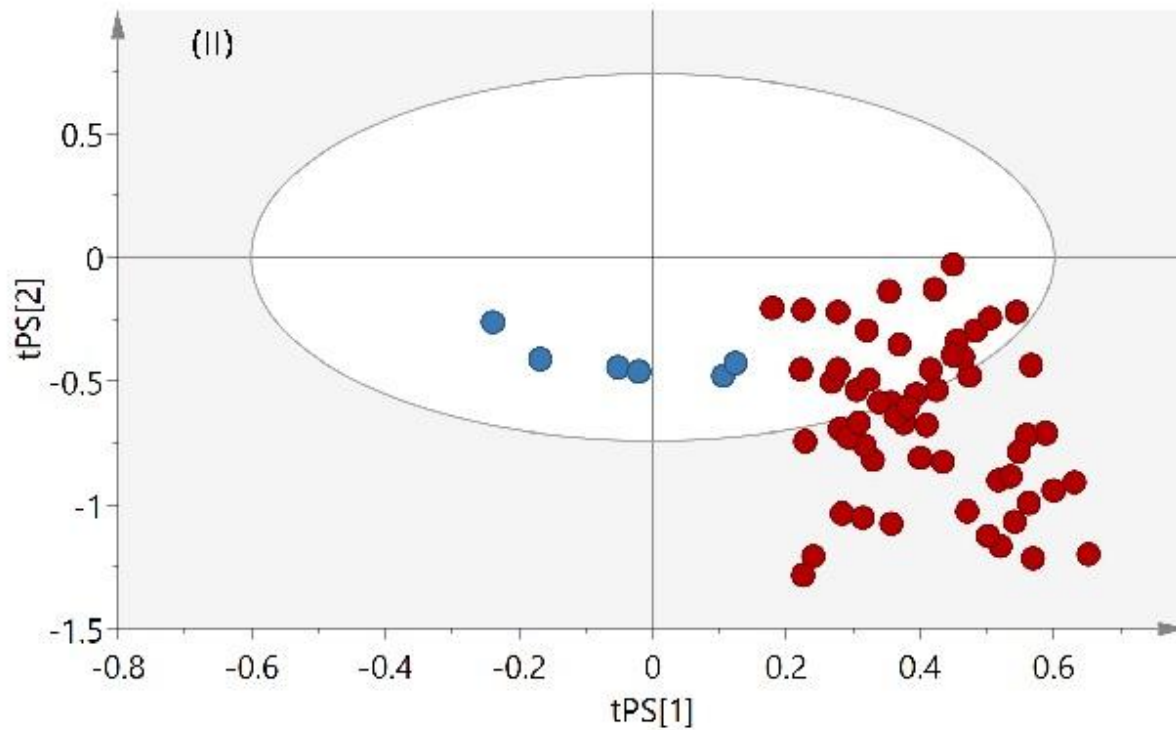
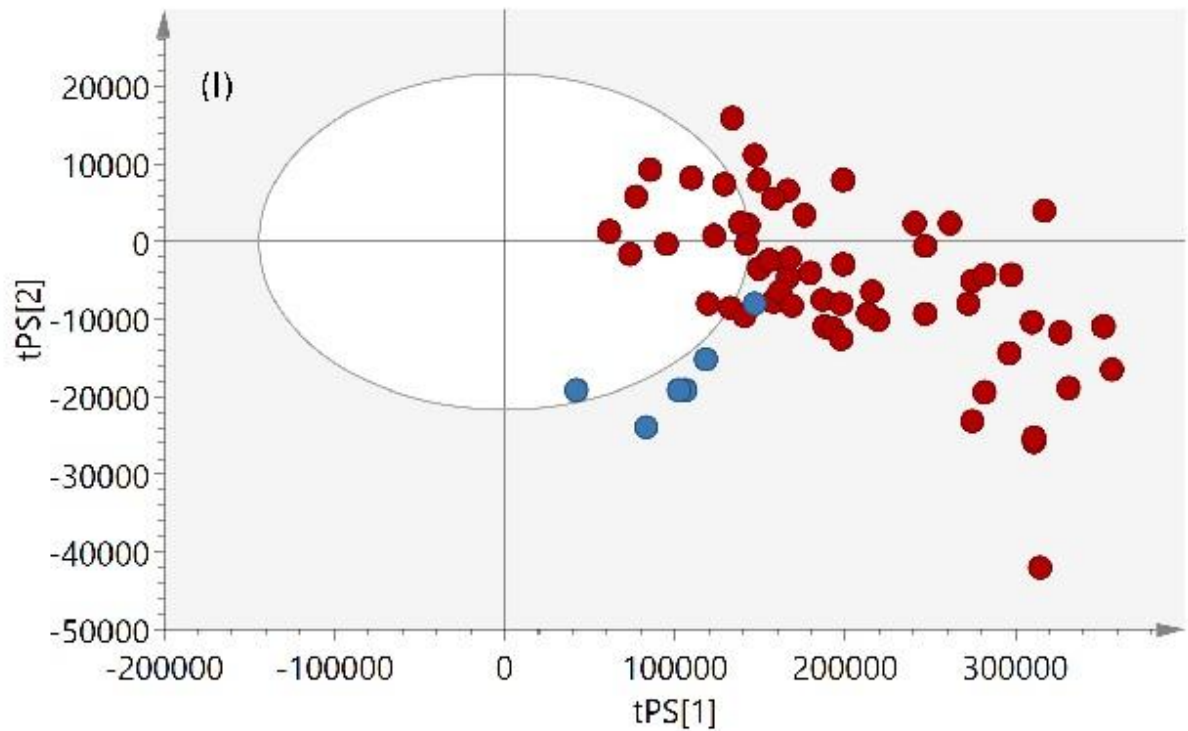


Figure S3. Predicted Score Plot of the R samples (as test set) using P samples as a prediction set: (I) UNor method, (II) NorCont method (blue circles: organic milk samples, red circles: conventional milk samples).

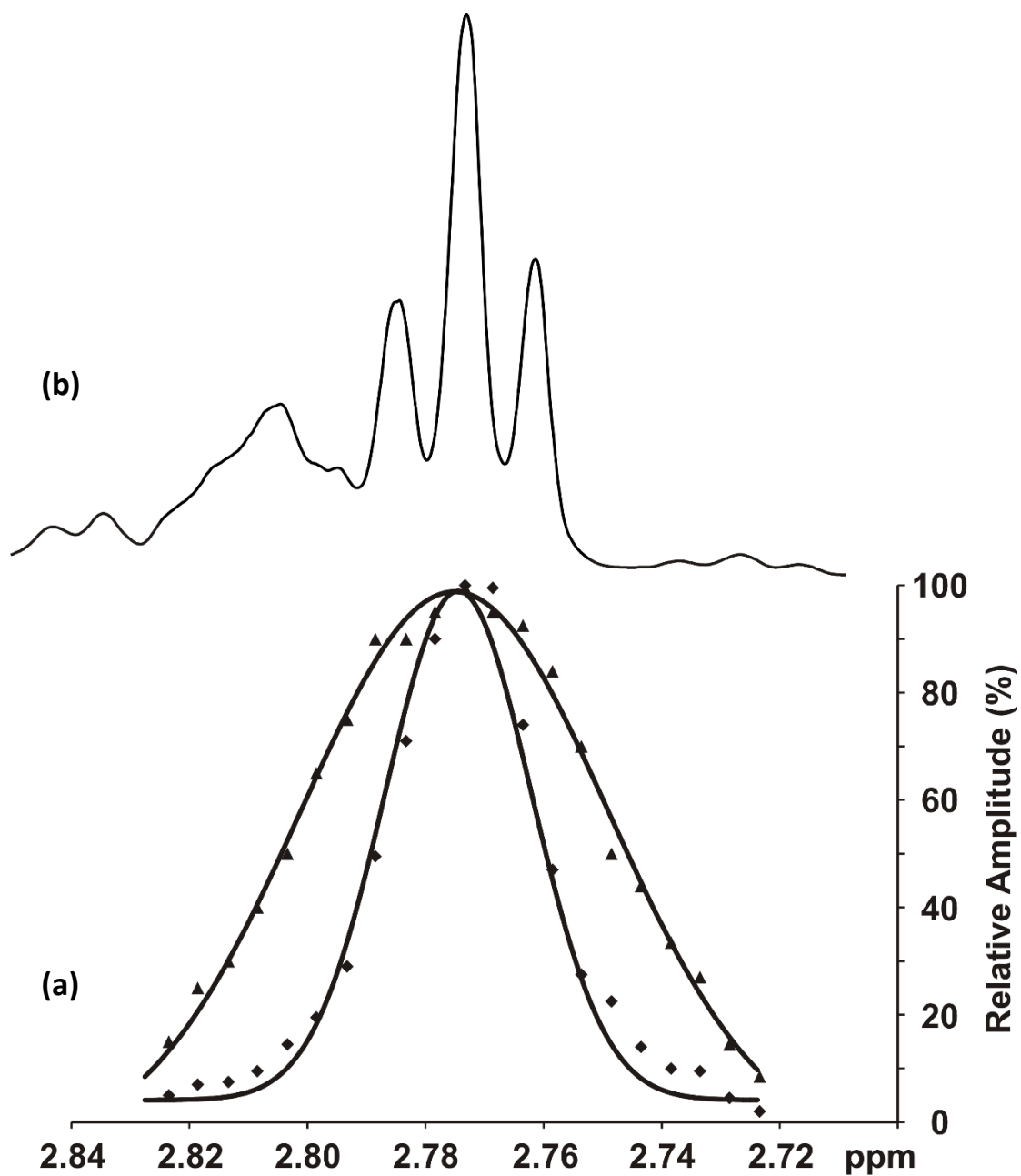


Figure S4. (a) Selectivity of the excitation bandwidth of the shaped pulse of 20 ms (▲) and 80 ms (◆) of the 1D TOCSY experiment. The carrier frequency of the shaped pulse was set in both cases at $\delta = 2.77$ ppm. (b) For comparison the apparent triplets of the allylic protons of linoleic acid ($\delta = 2.77$ ppm) and α -linolenic acid ($\delta = 2.81$ ppm) of the 1D ¹H NMR spectrum of the lipid fraction of lyophilized bovine milk sample in CDCl₃ are presented.

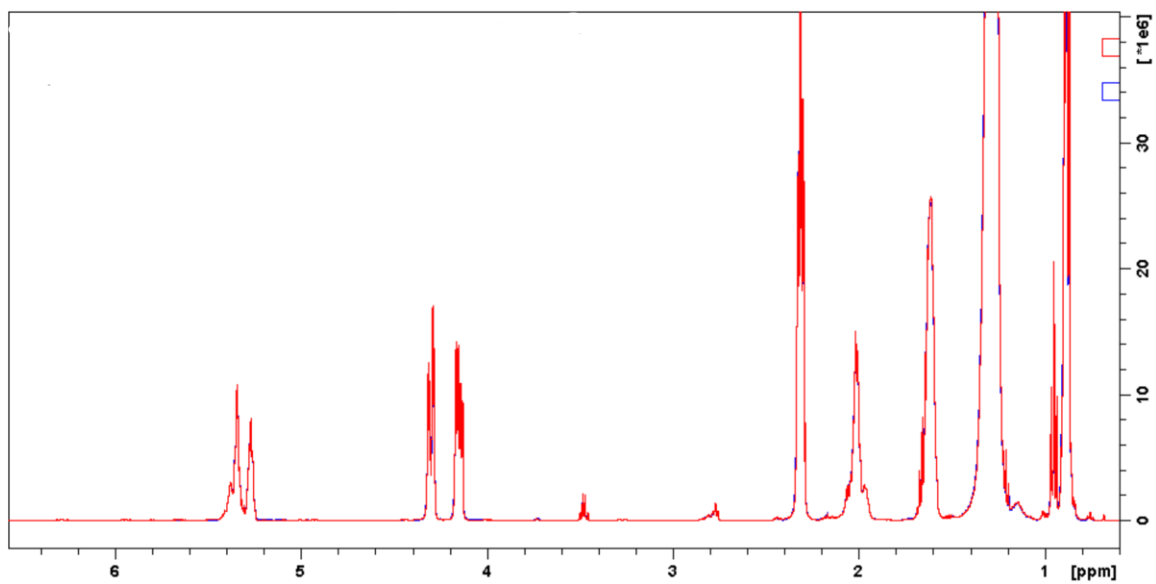


Figure S5. 500 MHz ^1H NMR spectra of the lipid fraction of a lyophilized bovine milk sample in CDCl_3 ; T, 298 K; number of scans, 256 ; recycle time, 9.3 s (blue color) and 14.5 s (red color).