

Monitoring cellular responses upon fatty acid exposure by Fourier Transform infrared spectroscopy and Raman spectroscopy

Heidi Najbjerg^a, Nils Kristian Afseth^b, Jette F. Young^{*a}, Hanne C. Bertram^c, Mona E. Pedersen^b,
Stine Grimmer^b, Gjermund Vogt^b, Achim Kohler^{b,d}

^a Department of Food Science, Aarhus University, Blichers Allé 20, 8830 Tjele, Denmark

^b Nofima Mat, Osloveien 1, 1430 Ås, Norway

^c Department of Food Science, Aarhus University, Kirstinebjergvej 10, 5792 Aarslev, Denmark

^d Centre for Integrative Genetics (CIGENE), Dept. of Mathematical Sciences and Technology, Norwegian University of Life Sciences, 1432 Ås, Norway

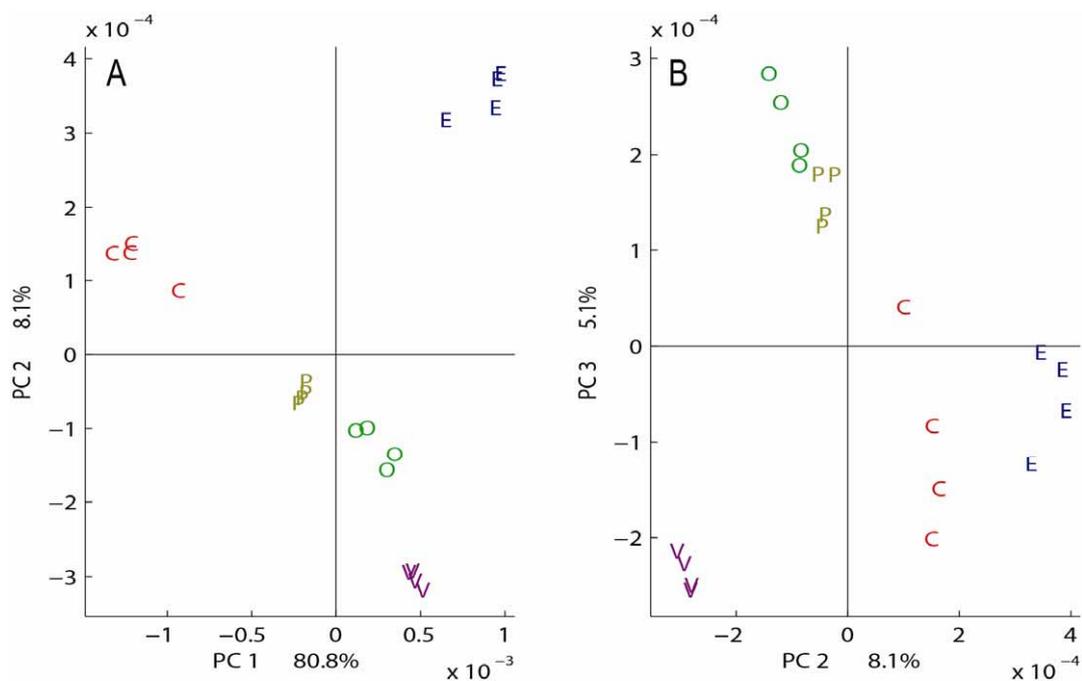


Fig. S1 Principal component analysis (PCA) of FTIR spectral data of HepG2-SF cells exposed to four different fatty acids: palmitic acid (16:0), oleic acid (18:1cis-9), elaidic acid (18:1trans-9) and vaccenic acid (18:1trans-11) and control medium. A) PCA score plot showing principal component (PC) 1 and 2, explaining 80.8% and 8.1 % of the sample variance, respectively. B) PCA score plot PC 2 and 3, with PC 3 explaining 5.1 % of the sample variance. The raw spectra are pre-processed using second derivative and Extended Multiplicative Signal Correction (EMSC). Second derivative minima are multiplied by -1. C: control, P: palmitic, O: oleic, E: elaidic, V: vaccenic.

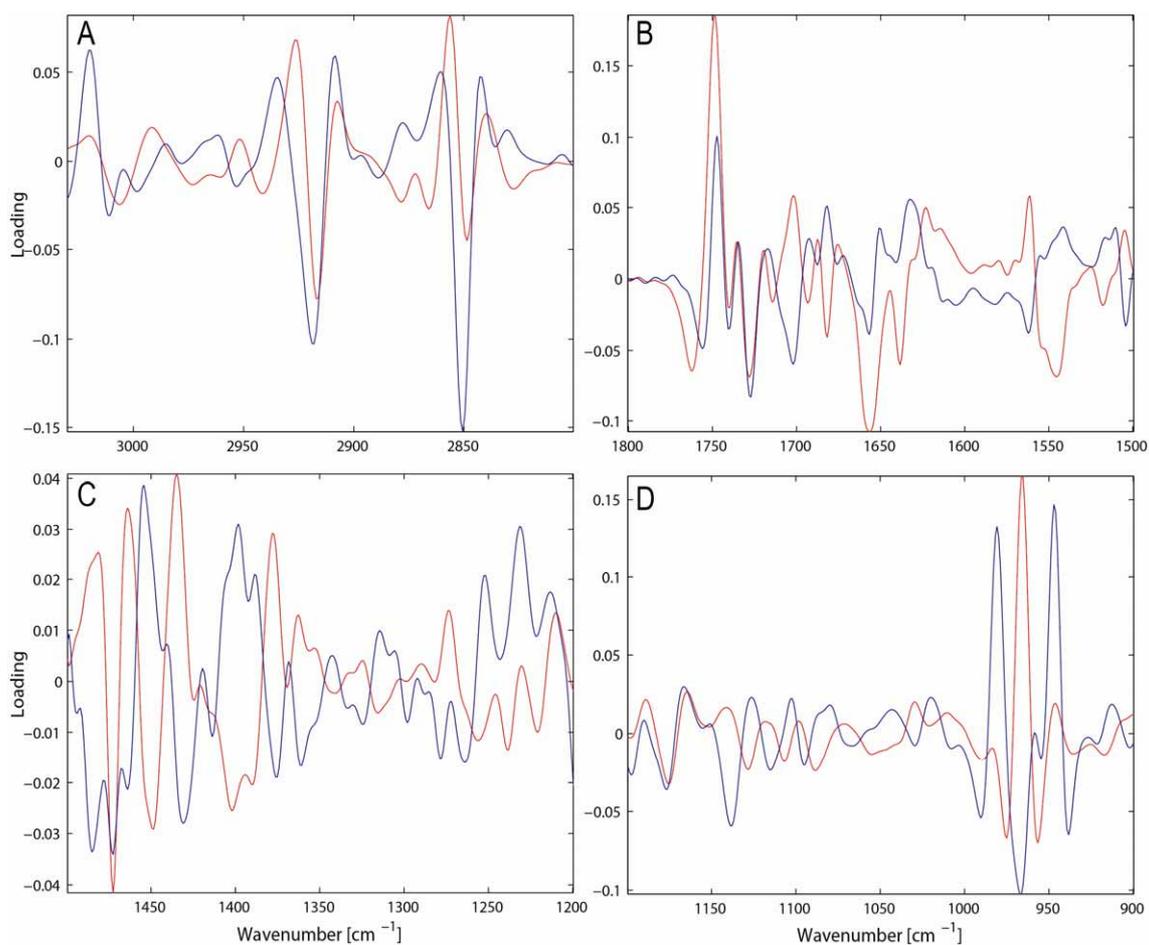


Fig. S2 Line plot of loadings from principal component analysis (PCA) of FTIR spectral data of HepG2-SF cells exposed in three days to four different fatty acids: palmitic acid (16:0), oleic acid (18:1, cis-9), elaidic acid (18:1, trans-9) and vaccenic acid (18:1 trans-11) showing PC1 (Red) and PC2 (Blue). The line plot is divided into four regions: 3050-2800 cm^{-1} (A), 1800-1500 cm^{-1} (B), 1500-1200 cm^{-1} (C), 1200-900 cm^{-1} (D).