



PRESENT AND FUTUR OF METABOLOMICS BY NMR

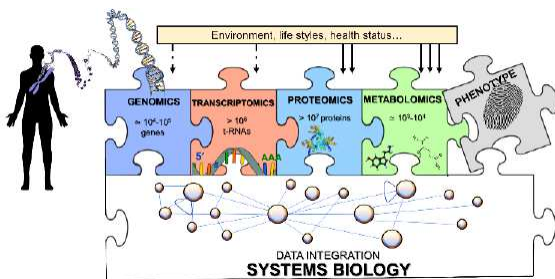
GUEST LECTURE by



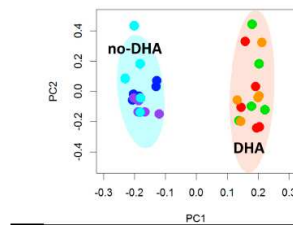
Prof. Dr. Claudio Luchinat
Center of Magnetic Resonance (CERM)
and Department of Chemistry,
University of Florence, Italy

Friday, 05.10.2018
10:00

Lecture Hall MC.05, MED Campus
(Neue Stiftingtalstrasse 6, ground floor), MUG

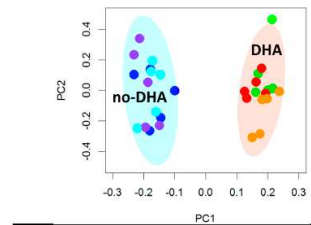


The flow of information in systems biology from „High-throughput metabolomics by 1D NMR.“ Vignoli et al. (2018) Angew Chem Int Ed Engl. in press



Confusion Matrix		
P \ A	DHA	No-DHA
DHA	93.3	6.7
No-DHA	13.3	86.7

Discrimination accuracy = 90%



Confusion Matrix		
P \ A	DHA	No-DHA
DHA	100	0
No-DHA	0	100

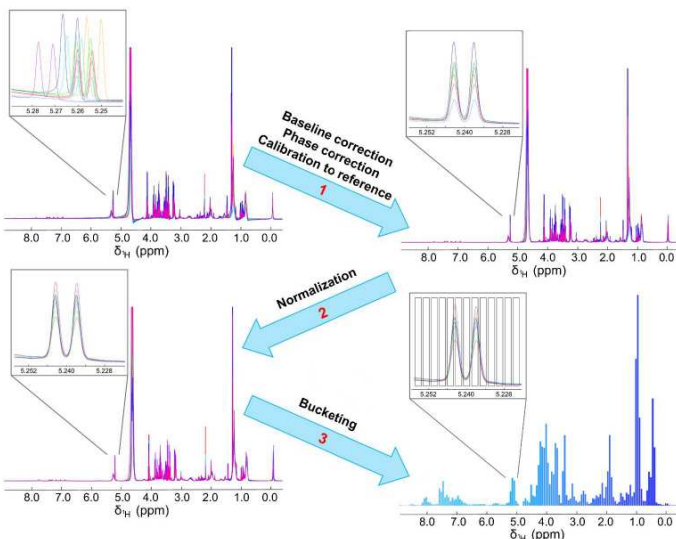
Discrimination accuracy = 100%

(a)

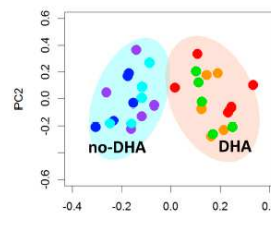
(b)

Lipidomic phenotyping by NMR analysis.

from „Evidence of a DHA Signature in the Lipidome and Metabolome of Human Hepatocytes.“ Ghini et al. (2017) Int J Mol Sci. 18(2). pii: E359



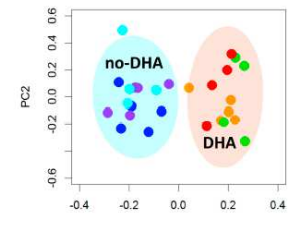
Key stages of NMR spectral processing. from „High-throughput metabolomics by 1D NMR.“ Vignoli et al. (2018) Angew Chem Int Ed Engl. in press



Confusion Matrix		
A \ P	DHA	no-DHA
DHA	86.7	13.3
no-DHA	6.7	93.3

Discrimination accuracy = 90%

(a)



Confusion Matrix		
A \ P	DHA	no-DHA
DHA	92.9	7.1
no-DHA	7.7	92.3

Discrimination accuracy = 92.6%

(b)

Cytoplasmic metabolomic phenotyping by NMR analysis.

from „Evidence of a DHA Signature in the Lipidome and Metabolome of Human Hepatocytes.“ Ghini et al. (2017) Int J Mol Sci. 18(2). pii: E359