



Raw material effects and novel control- and optimisation strategies in agri-food valorization

Nils Kristian Afseth, Sileshi Wubshet, Ingrid Måge, Ulrike Böcker, Jens Petter Wold, Kenneth Kristoffersen and Diana Lindberg

Nofima – Norwegian institute of Food, Fisheries and Aquaculture research

Enzymatic protein hydrolysis in agri-food valorization



Enzymatic protein hydrolysis in agri-food valorization

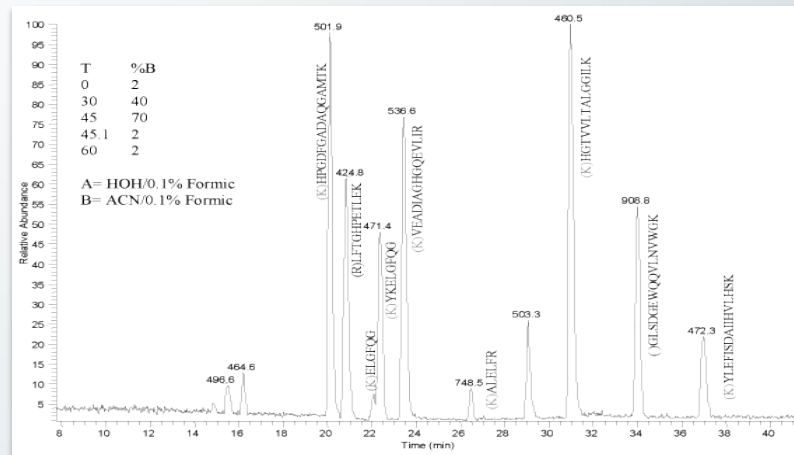


Enzymatic hydrolysis (the simple case)



Single protein digestion:
-Myoglobin (trypsin)

LCMS Chromatogram

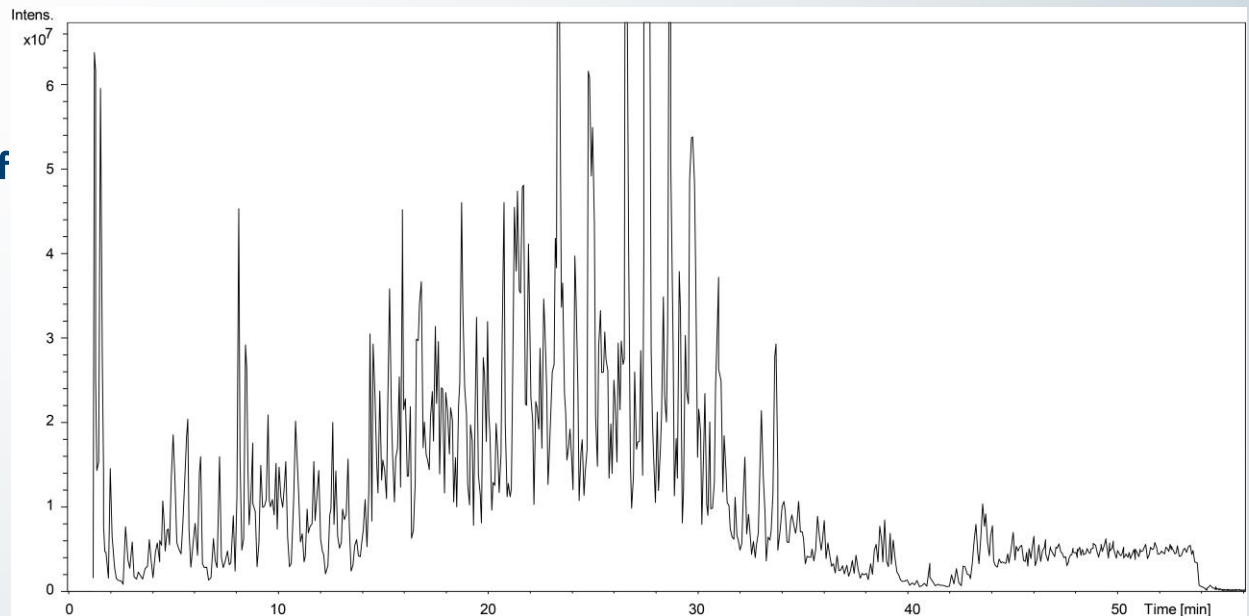


<http://www.ionsource.com>

Enzymatic hydrolysis (the real case)

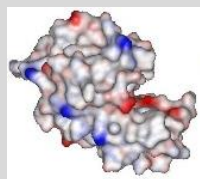
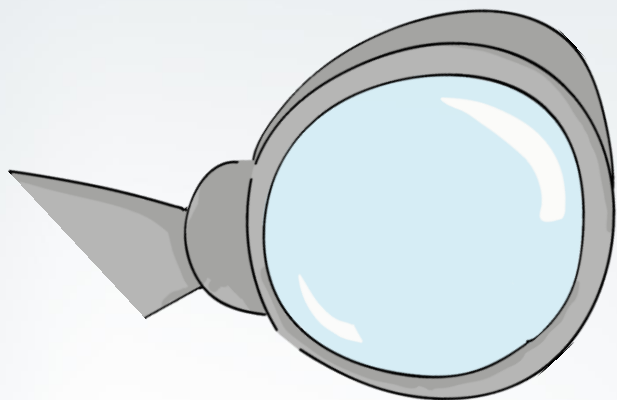


LCMS chromatogram of Hydrolysate from MDC

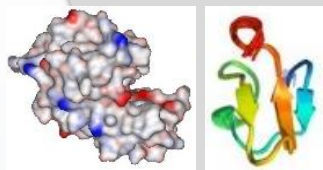


Wubshet, S. G. et al. *Unpublished data, 2016*

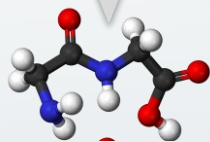
Bioanalytics and protein characterisation



Degree of hydrolysis – OPA / TNBS etc.

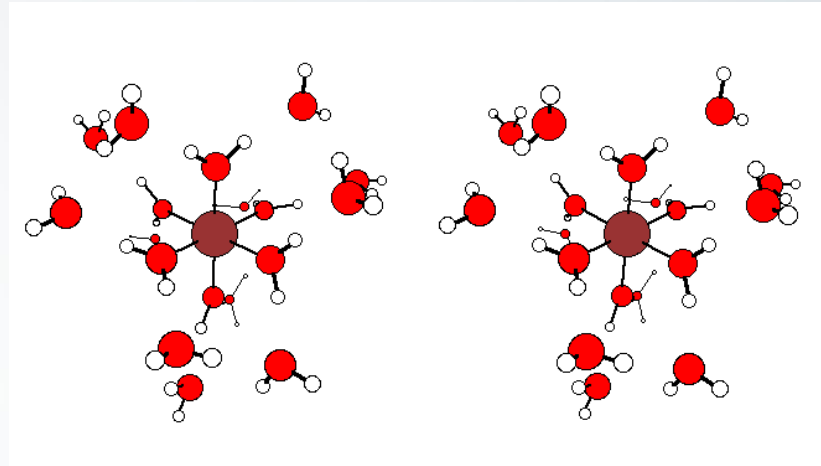


Size distributions – size exclusion chromatography



Detailed characterization – LCMS / NMR

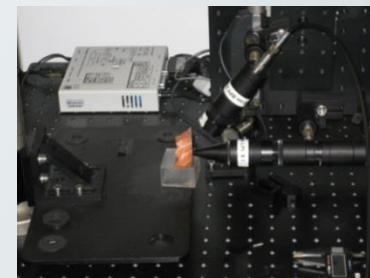
Rapid and non-destructive spectroscopic measurements



**NIR
spectroscopy**

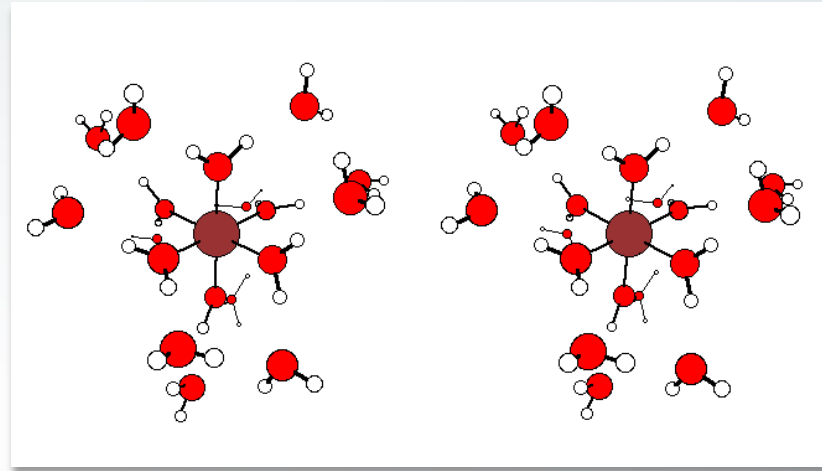


**FTIR
spectroscopy**

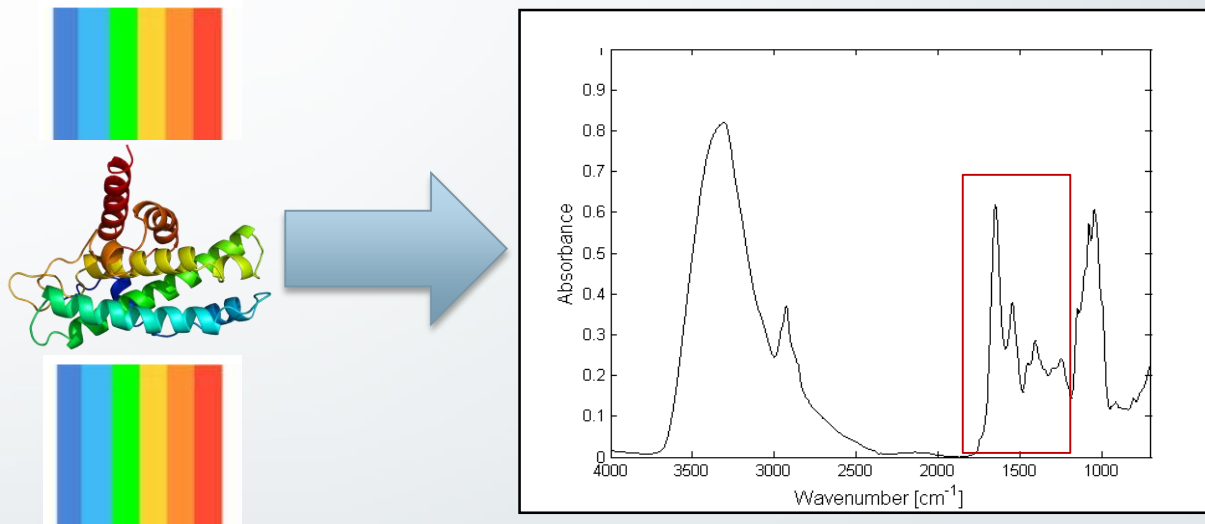


**Raman
spectroscopy**

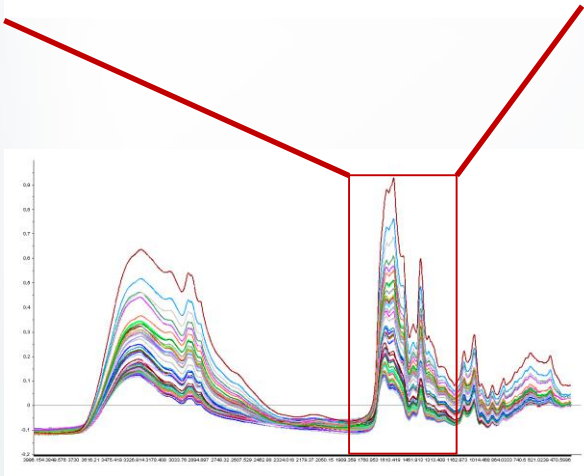
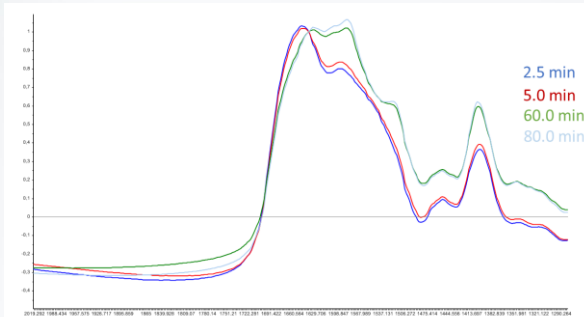
FTIR provides a biochemical fingerprint



Qualitative and quantitative information on protein structure



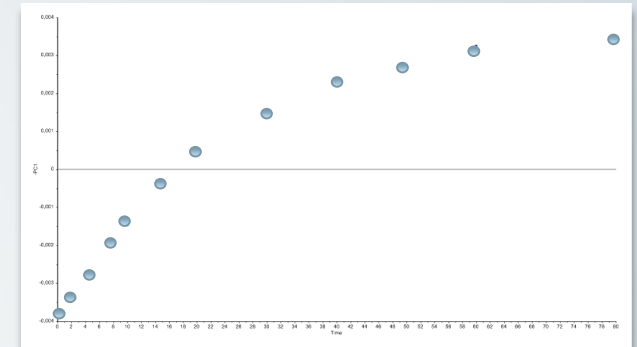
FTIR for monitoring enzymatic protein hydrolysis



FTIR spectra



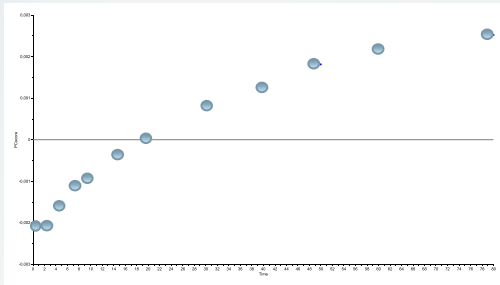
Degree of hydrolysis (DH%)



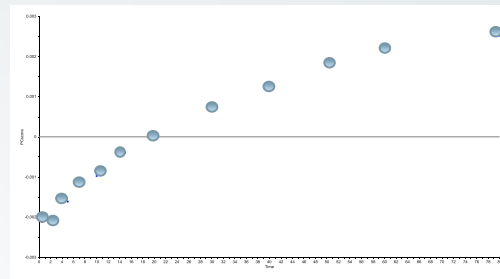
Time

Average protein size
estimation /
Degree of Hydrolysis (DH%)

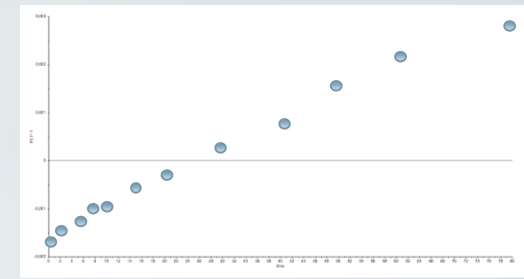
FTIR for monitoring enzymatic protein hydrolysis



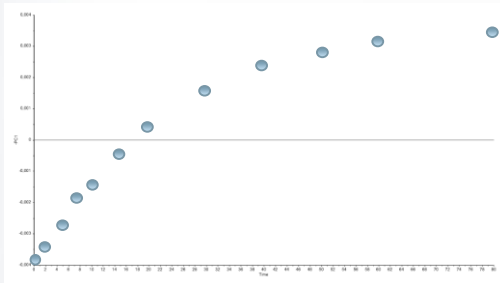
Chicken muscle



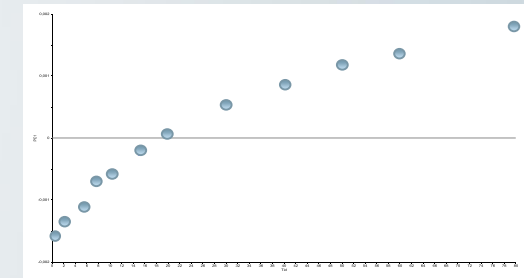
Salmon muscle



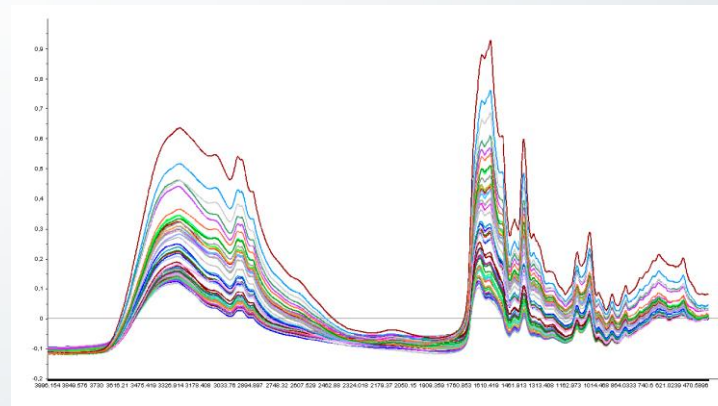
Salmon heads



Chicken by-products



Salmon backs

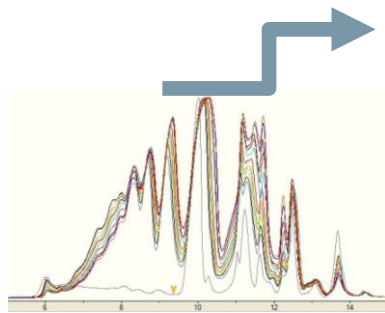


FTIR spectra - Chicken by-products

Ref:

Böcker, U., Wubshet, S., Måge, I., Lindberg, D., and Afseth, N.K. Fourier-transform Infrared spectroscopy for qualitative characterization of protein chain reductions in enzymatic hydrolysis, submitted manuscript.

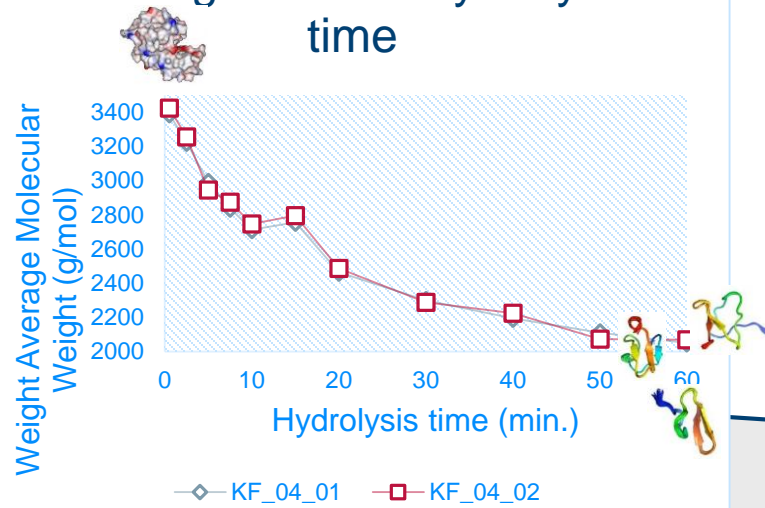
Quantitative monitoring of enzymatic protein hydrolysis



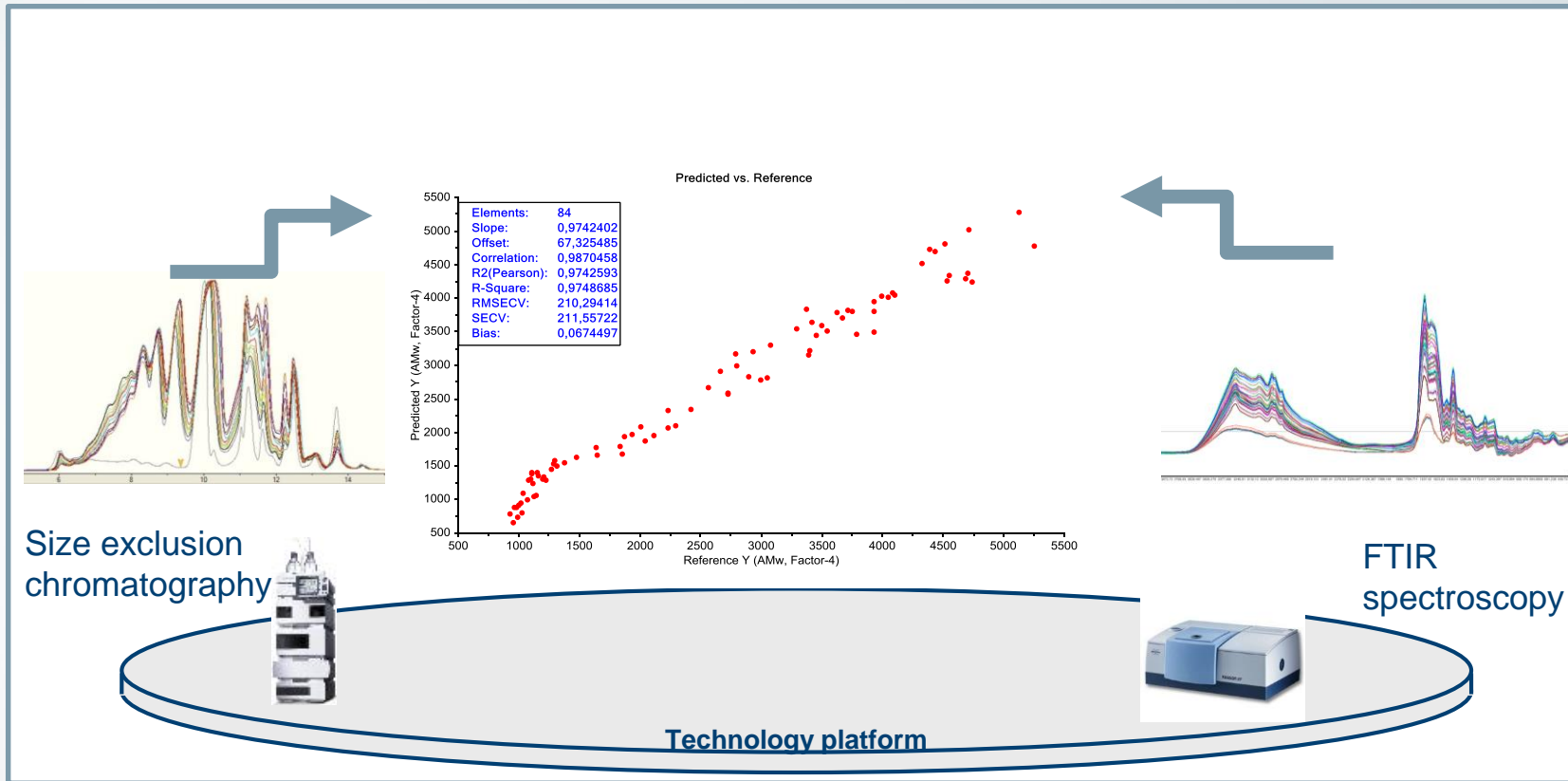
Size exclusion chromatography



Average MW vs Hydrolysis time



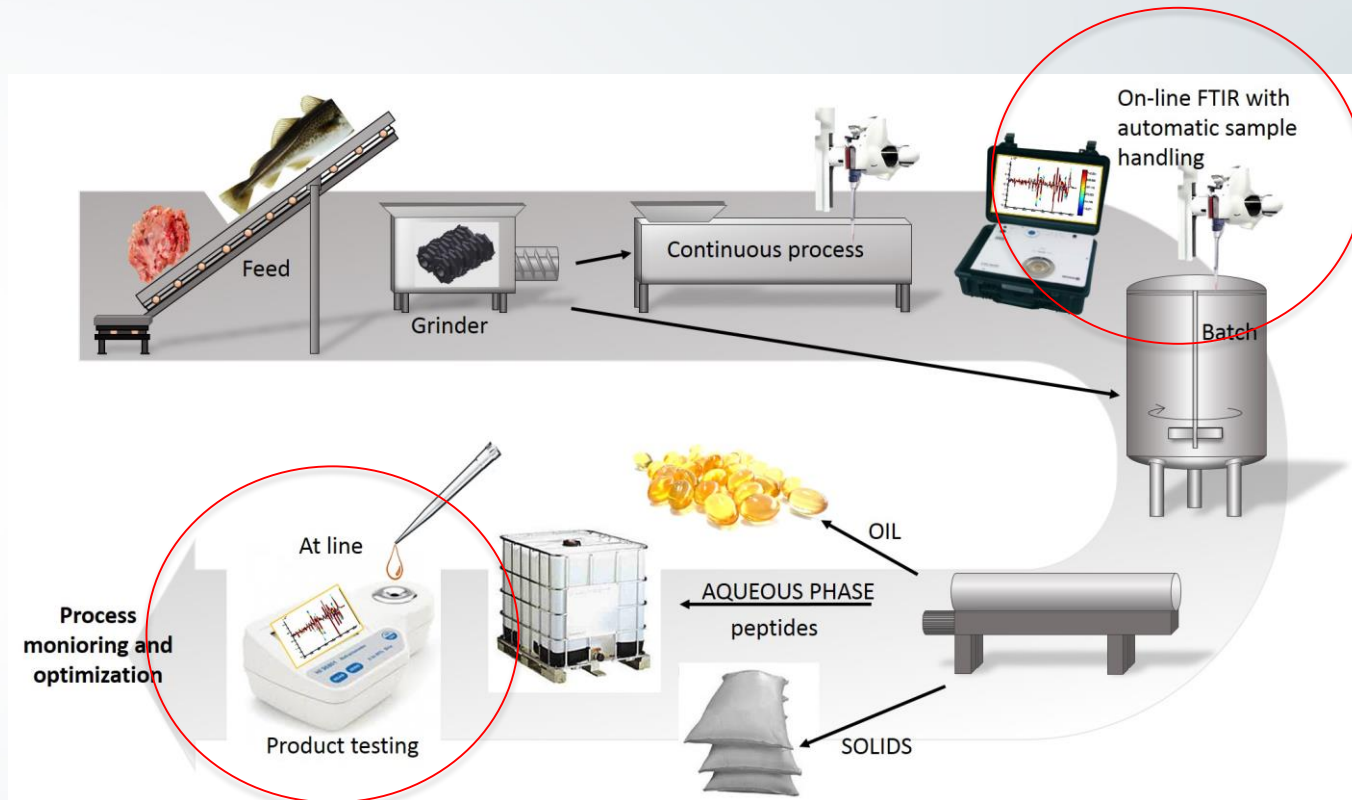
Quantitative monitoring of enzymatic protein hydrolysis



Refs:

Wubshet, S., Måge, I., Bocker, U., Lindberg, D., Knutsen, S.H., Rieder, A., Airado-Rodriguez, D. & Afseth, N.K.. FTIR as a rapid tool for monitoring enzymatic protein hydrolysis of food processing by-products, submitted manuscript.

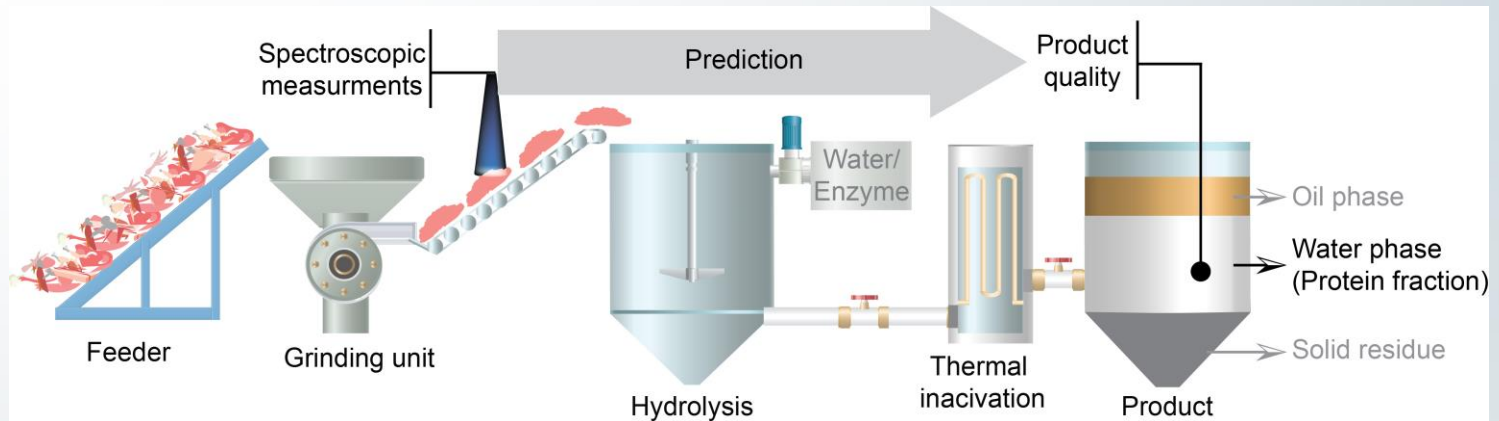
A spectroscopic platform for control and optimisation of biotech processes



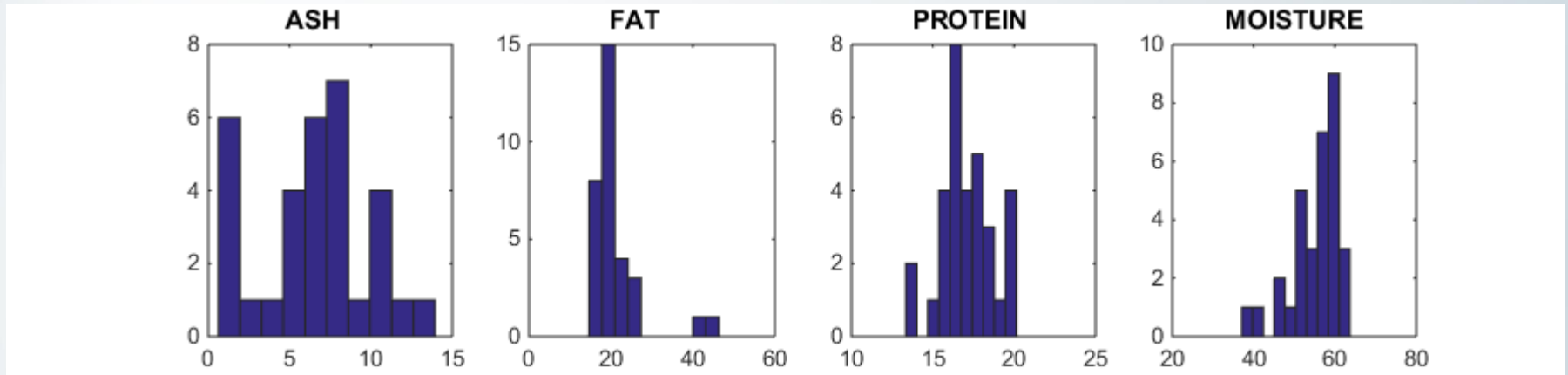
Projects:

- LiqIR – Project leader: Prediktor (2015 – 2017) - Funded by the Regional research fund “Oslofjordfondet”, with Biomega, Nutrinar, VEAS, and SINTEF ICT

Prediction of hydrolysis outcome based on raw material analysis



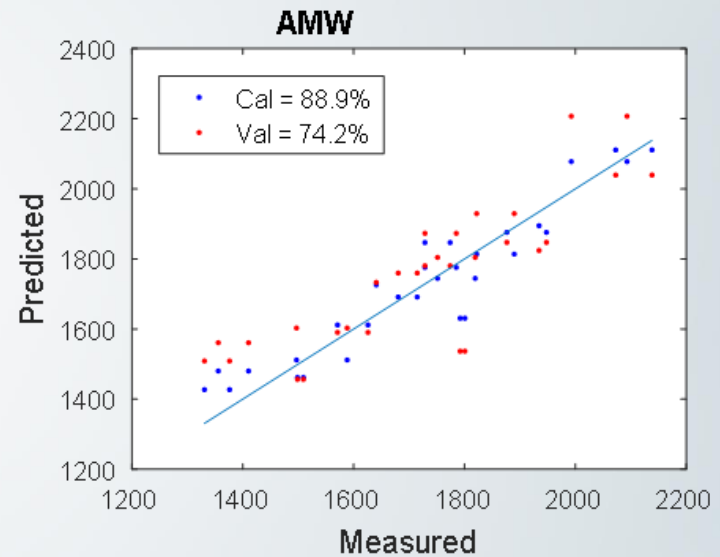
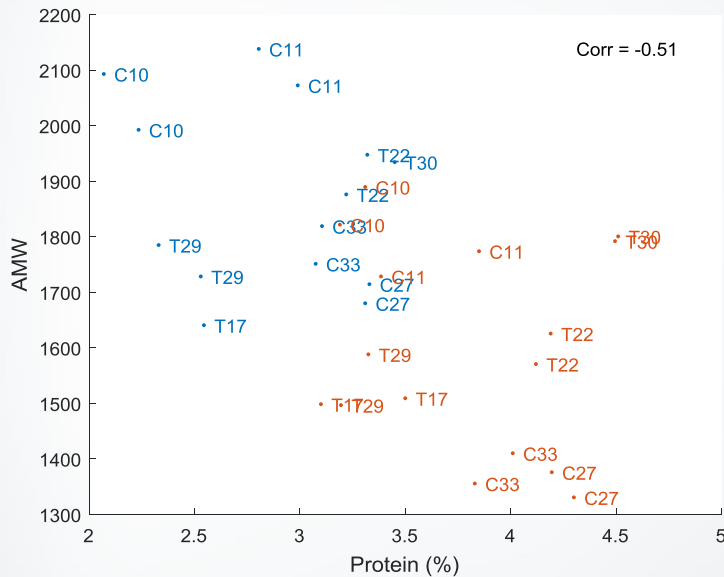
Prediction of hydrolysis outcome – the chicken and turkey case



32 samples of chicken and turkey rest raw materials were collected at Nortura Hærland, spanning variation in chemical composition

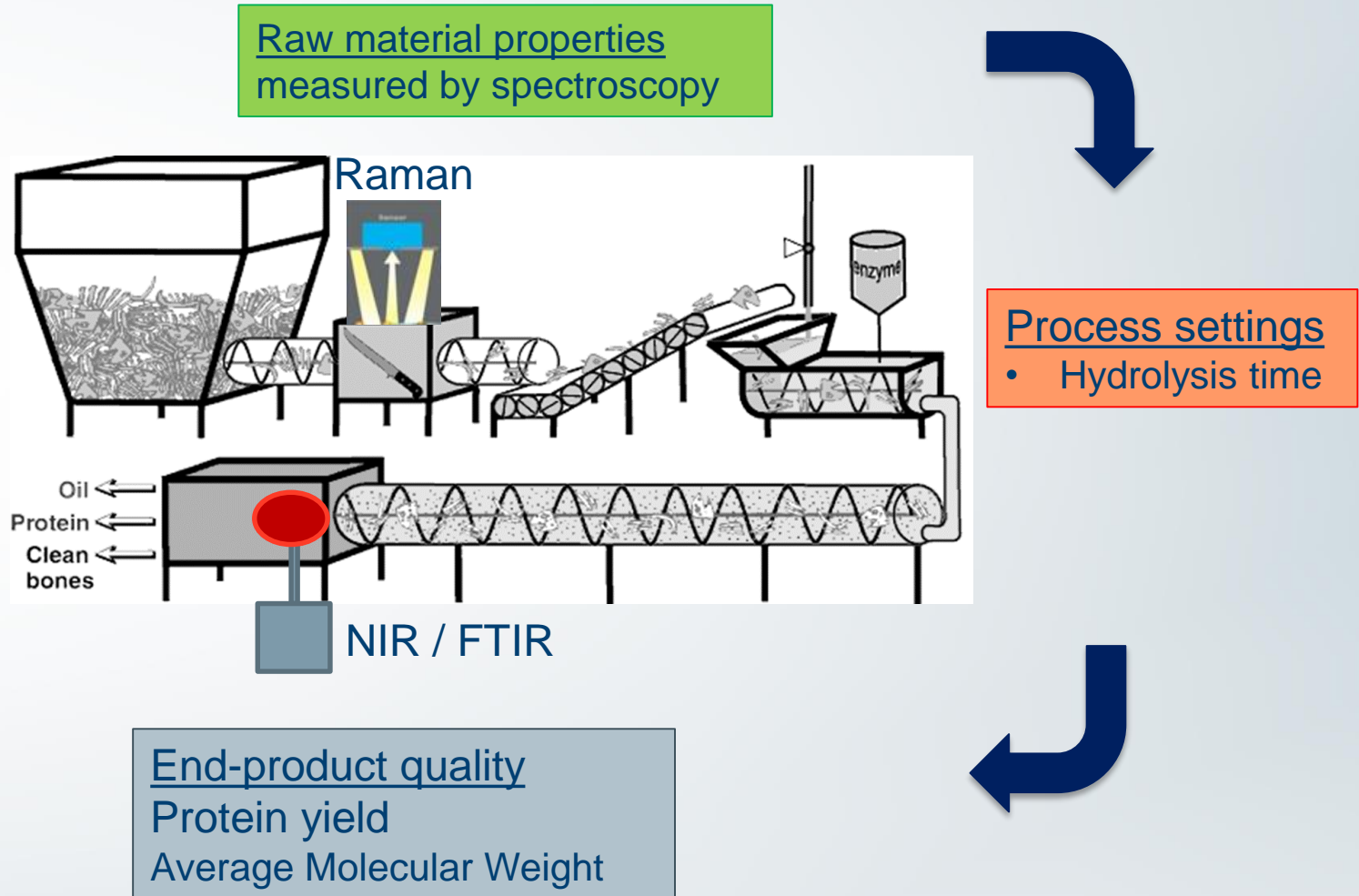
End-product quality prediction

By-product-characterisation (spectroscopy)
+ processing conditions:



(AMV = Average molecular weight measured by SEC)

Future protein hydrolysis – raw material variation can be handled using spectroscopy



Refs:

Wubshet, S. et al., Spectroscopic process monitoring in enzymatic protein hydrolysis of byproducts: A strategy for stable product quality, manuscript.

Future agri-food valorisation



Sensory, consumer
and market



Paula Tomasco



Themis Altintzoglou



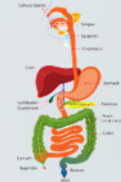
Industry-scale
processing



Ragnhild Whitaker



Åge Oterhals



In vitro digestion
and bioactivity



Bente Kirkhus



Sissel Rønning



Grain utilisation



Stefan Sahlstrøm



Fruits, berries and
vegetables



Grethe Iren Borge



Raw materials quality



Eva Veiseth



Rune Rødbotten



Fermentation



Lars Axelsson



Dimi Tsimorotas



Lipid utilisation



John-Erik Haugen



Sediment utilisation

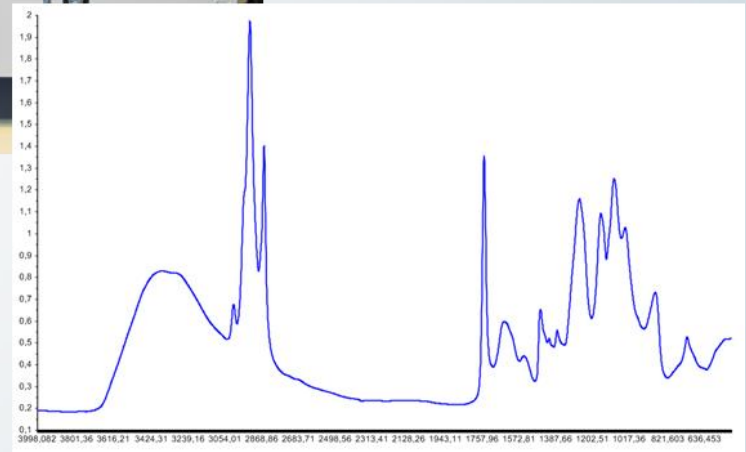
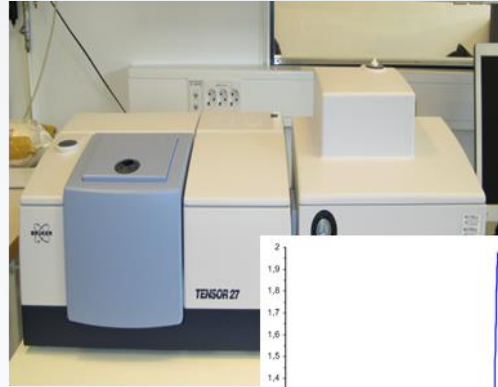
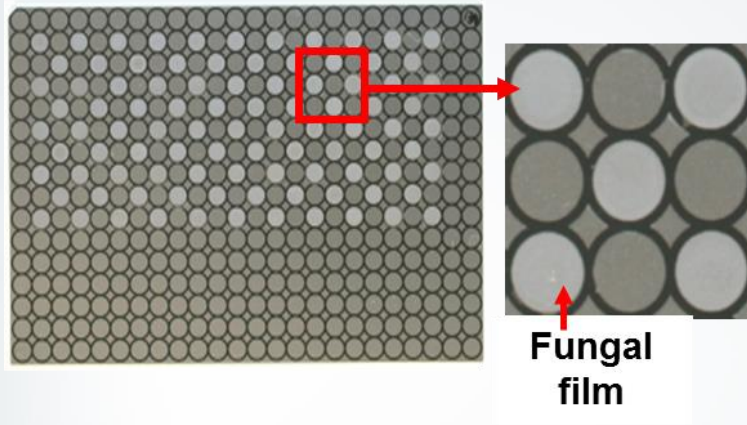


Sissel Albrigtsen



Tone Aspevik

Identification, optimization and screening of microbial oils grown on agrifood rest raw materials



FTIR spectrum of *M. circinelloides* producing microbial oils

Interrest and SingleCellOil projects



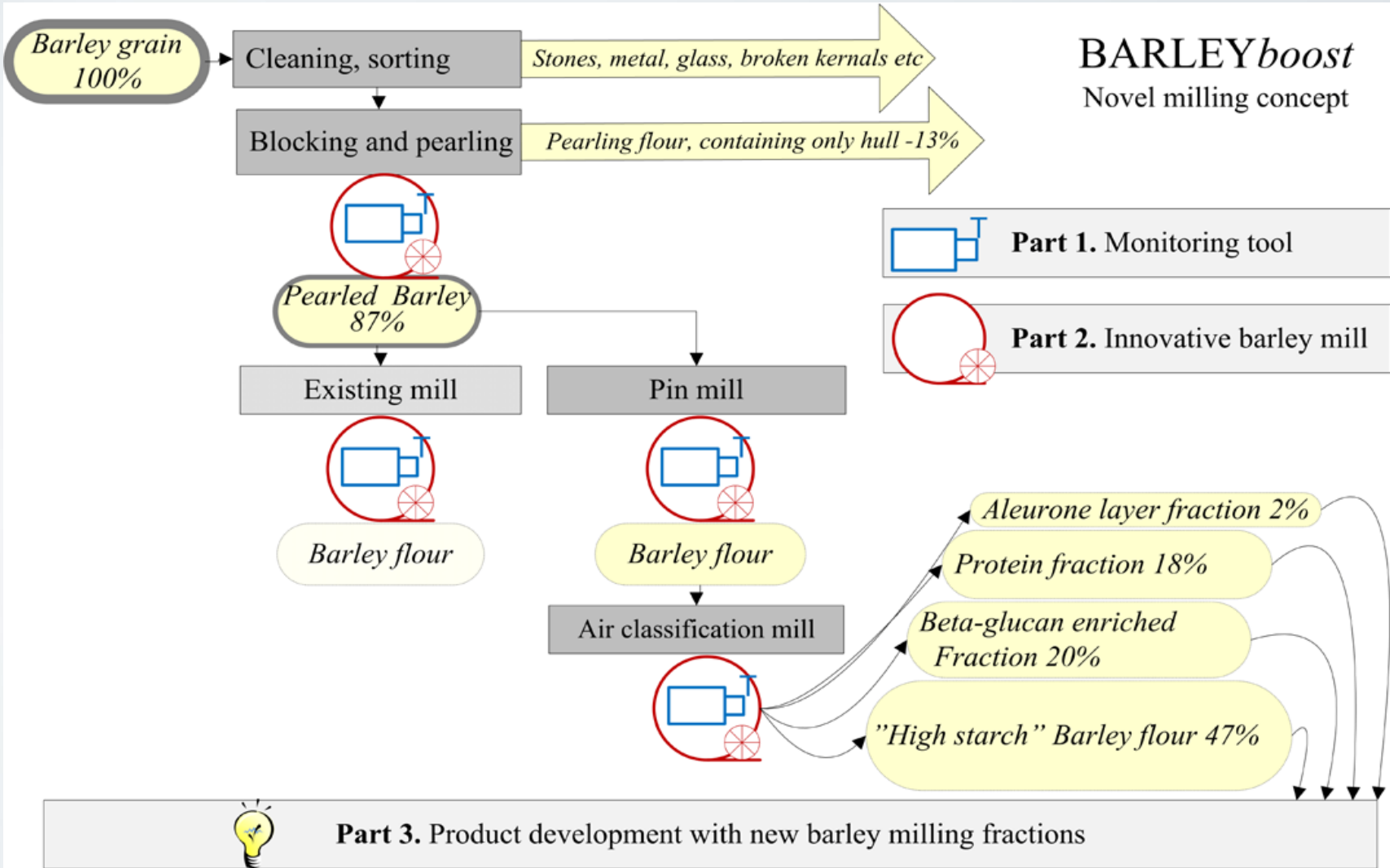
Lars Axelsson



Dimi Tsimorotas

Ref:

Shapaval, Volha; Afseth, Nils Kristian; Vogt, Kjell Gjermund; Kohler, Achim. Fourier transform infrared spectroscopy for the prediction of fatty acid profiles in *Mucor* fungi grown in media with different carbon sources. *Microbial Cell Factories* 2014 ;Volume 13. p. 86



Stefan Sahlstrøm

Jens Petter Wold



Concluding remarks

- Controlling and handling rest raw material complexity and variation is essential for future valorisation
- Industrial analytical tools are key factors in producing protein products of defend qualities (e.g. sensorial, functional or bioactive properties) for higher paying markets:
 - Flexible processing for dealing with rest raw material heterogeneity and variations (e.g. depending on season or pre-production steps)
 - Possibilities for real-time process adjustments



Sustainable use of rest raw materials through development of profitable industrial processes and products

Acknowledgements

- Industry collaboration:
 - Norilia, Nortura, Biomega and Nutrimar
- Research collaboration:
 - SINTEF FA, SINTEF ICT and NMBU
- Funding from The Norwegian Research Council, Oslofjordfondet, the foundation for Research levy on Agricultural Products, and EU:
 - «HydrolyseMonitor», led by Nofima
 - «Chickenlysis», led by Nortura
 - «LiqIR», led by Prediktor
 - «Cycle», led by Sintef FA
 - «Interrest», led by Nofima
 - «SingleCellOil», led by NMBU
 - «BarleyBoost», led by Nofima
 - «Multiblock», led by Nofima
 - «Food quality imaging», led by Nofima

Nofima centre of excellence

SPECTEC

Rapid and non-destructive measurements for process optimisation

Leader: Jens Petter Wold

PEPTEK

A characterisation and processing platform for future protein production

Leader: Ragnhild Whitaker