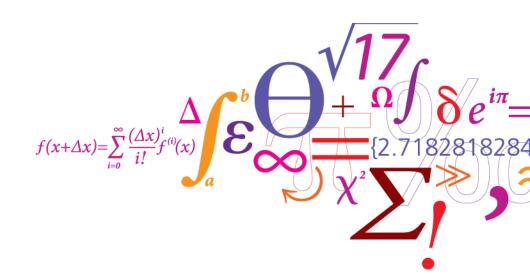


Chemical analysis of fish meal and fish oil

<u>Charlotte Jacobsen</u> and Gonçalo S. Marinho Professor and group leader Research group for Bioactives – Analysis and Application

chja@food.dtu.dk





Agenda

- TVN
- Biogenic amines
- Proteins (Kjeldahl vs Dumas)
- Free fatty acids
- PV (titration vs other spectrophotometric methods)
- AV
- TBARS
- Volatile oxidation products by headspace GC-MS



TVN

 The combined total amount of ammonia, dimethylamine and trimethylamine is called the total volatile base content of the fish (usually expressed as mg-N/I00 g minced fish) and is a commonly used estimate of spoilage

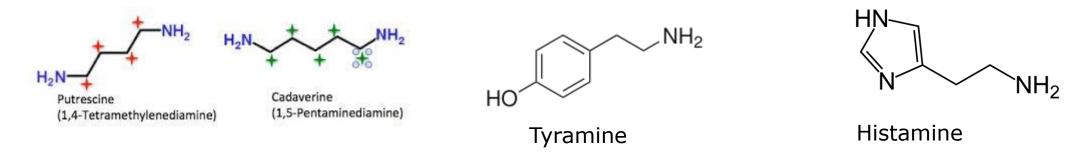
Conway method

- Make an aqueous acidic extract of the material
- Make the extract alcaline to release volatile bases
- Collect the bases in HCl and titrate with NaOH using Andersen indicator
- Can also be determined by steam distillation using Kjeldahl apparatus
- Can alse be determined by capillary electrophoresis



Biogenic amines

• Acidic extraction of biogenic amines (cadaverine, putrescine, tyramine and histamine)



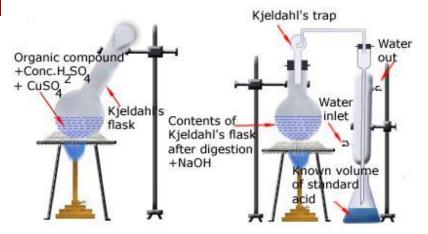
- HPLC analysis
- Analysis by Capillary Electrophoresis (CE) can also be performed



Protein determination in fish meal

Kjeldahl vs Dumas

Kjeldahl



Advantages:



- widely used internationally and is still the standard method for comparison against all other methods
- universality, high precision and good reproducibility have made it the major method for the estimation of protein in foods



Disadvantages:

- ❖ It does not give a measure of the true protein, since all nitrogen in foods is not in the form of protein
- Different proteins need different correction factors because they have different amino acid sequences
- ❖ The use of concentrated sulfuric acid at high temperatures and heavy metal catalysts poses a considerable hazard
- The technique is time consuming to carry-out.

Dumas



Advantages:

❖ It is much faster than the Kjeldahl method (under 4 minutes per measurement, compared to 1-2 hours for Kjeldahl)

Sample in tin capsule

Oxygen -Dosing

CRUCIBLE

Oxidation Catalyst

High Quality Copper

- It doesn't need toxic chemicals or catalysts
- Many samples can be measured automatically
- It is easy to use.

Disadvantages:

Condensation

Drainage Device

H₂0 Trapped

High initial cost

CO, Trapped

GC Column

Trap 1

Trap 2

- ❖ It does not give a measure of the true protein, since all nitrogen in foods is not in the form of protein.
- Different proteins need different correction factors because they have different amino acid sequences

Comparison of Dumas and Kjeldahl on different samples at DTU Food

1: Microalgae

2: Fish

3: Oat beer

4: Barley beer

5: Malt beer



Protein content (g/100 g; N x conversion factor)

| Samples | DTU (Kjeldahl) | Elementar (Dumas) | LECO 628 (Dumas) |
|------------------|----------------|----------------------|---------------------|
| Microalgae | 44,1±1,1 | 51,92±0,08 | 48,21±0,09 |
| Fish | 21,43±0,07 | 22,48±0,53 | 22,09±0,03 |
| Malt beer (B1) | 0,25 – 0,28 | | 0,31±0,003 |
| Barley beer (B2) | 0,15 - 0,31 | 0,24±0,008 * | |
| Oat beer (B3) | 0,51 – 0,60 | | 0,18±0,003 |





Free fatty acids

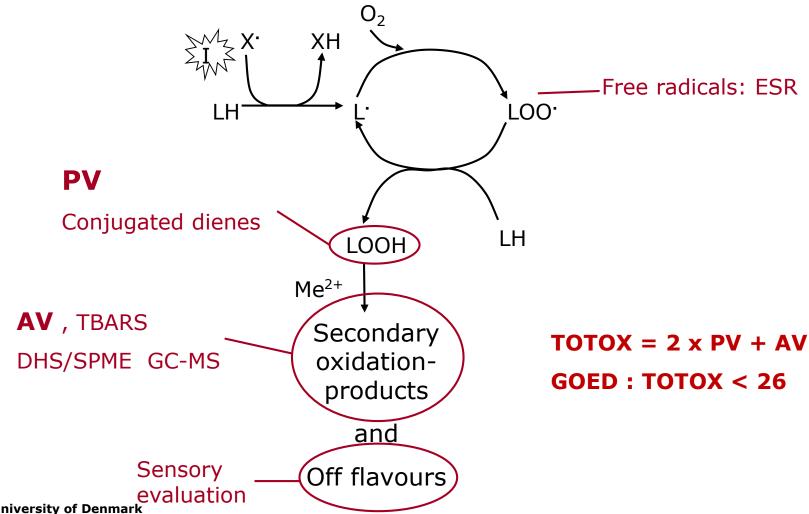
• Free fatty acids are titrated with NaOH with phenolphthalein as indicator

$$R-COOH + OH- \longrightarrow R-COO- + H2O$$

• pKa for fatty acids: 4-5



Measurement of lipid oxidation





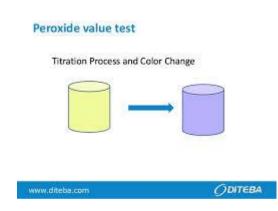
Peroxide value

For fish meal: Lipid extraction by chloroform and methanol to obtain lipid extract before PV analysis

PV (titration - colour change) (Standard method)

$$ROOH + 2I^{-} \rightarrow ROH + I_{2}$$

$$I_2 + 2S_2O_3^{2-} \rightarrow 2I^- + S_4O_6^{2-}$$





Different PV methods

| | Modif IDF/ferro | Titration | Micro | FOX2 |
|---|---|--|--|---|
| 1 | Ox of ferro-salts to ferri ions | Ox of iodide to free iodine | Ox of iodide to free iodine | Ox of ferro-salts to ferri ions |
| 2 | Production of red colour after addition of SNC- | Titration with thiosulfate | Production of blue colour Iodine-starch complex | Production of blue colour complex (Ferri-Xyl-orange) |
| 3 | ROOH + Fe ²⁺ →Fe ³⁺ Fe ³⁺ + 3SNC ⁻ →complex | ROOH + $2I^{-} \rightarrow ROH$ + I_{2} $I_{2} + 2S_{2}O_{3}^{2-} \rightarrow 2I^{-}$ + $S_{4}O_{6}^{2-}$ | ROOH+ $2I^{-} \rightarrow ROH + I2$ $I_2 + \text{ starch } \rightarrow$ Incl.complex | ROOH+ Fe ²⁺ → Fe ³⁺ Fe ³⁺ + Xyl-or → complex |
| 4 | A 500nm | | A 565nm | A 560nm |
| 5 | 0.01-0.3 g / 0.1g | 1 g | 0.02-0.08 g | 0.01-0.3 g |

4: Absorption maximum coloured product; 5: Sample amount (oil); 6:

DTU Food, T Solvent volume, incl.complex: inclusion complex; Xyl-or: xylenol orange



AV - standard method oil industry

AV (spectrophotometric):

$$H_3C\text{-}O\text{-}\bigg[\hspace{-1mm}\big] \text{-}NH_2 + OHC\text{-}CH = \hspace{-1mm}CH\text{-}CH_3H_7 \hspace{1mm} \longrightarrow \hspace{1mm} H_2O + H_3C\text{-}O\text{-}\bigg[\hspace{-1mm}\big] \hspace{-1mm}\big[\hspace{-1mm}\big] \text{-}N = \hspace{-1mm}CH\text{-}CH = \hspace{-1mm}CH\text{-}CH_3H_7$$

p-anisidine + aldehyde (fx: 2-hexenal) \rightarrow coloured product

Colour intensity depends on the structure of the aldehydes!!

Thus, we do not really know what we measure

More sensitive and specific methods are therefore required, particularly for measurements of secondary oxidation products



TBARS

• Thiobarbituric acid reactive substances "TBA(RS)":

• TBA reacts with malondialdehyde, but pigment (535nm) is also formed with many other compounds (non-specific and interferences!)

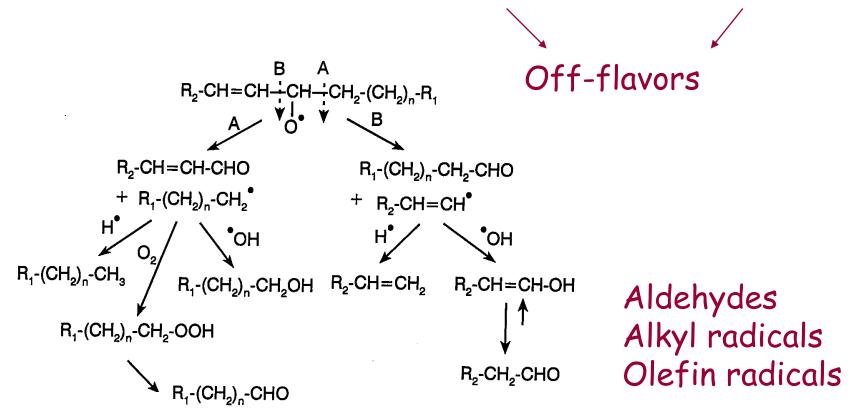
Lipid hydroperoxide decomposition



Metal ions catalyzes this reaction

+ R₁-(CH₂)₂₋₁-CH₂ + HCHO

 $R_2 - \downarrow - CH(O \cdot) - \downarrow - R_1 \rightarrow Aldehydes + other volatiles$

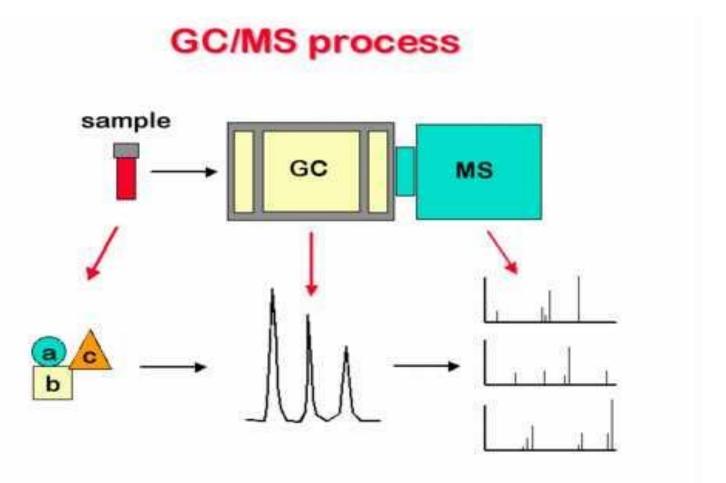


Frankel 1998



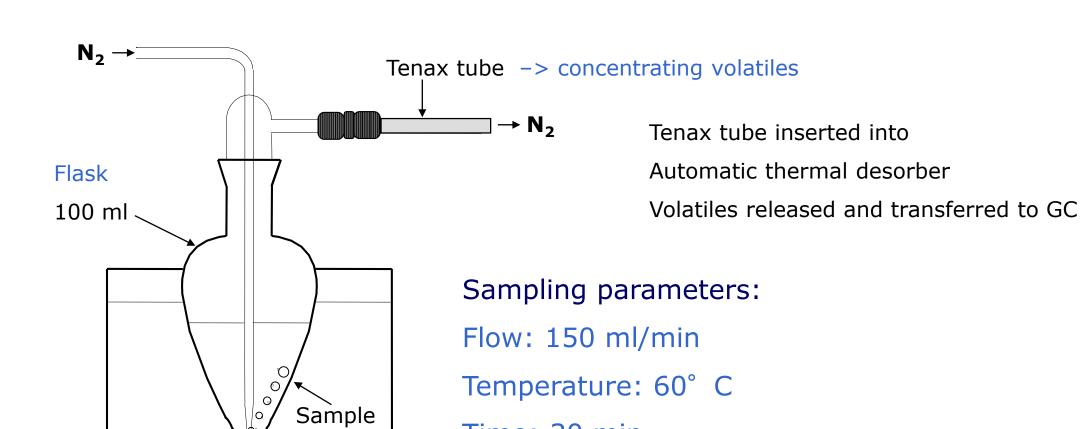
Gas chromatography - Mass spectrometry

- GC: separation of the different compounds -> chromatogram
- MS: analysis of the different compounds -> spectrum







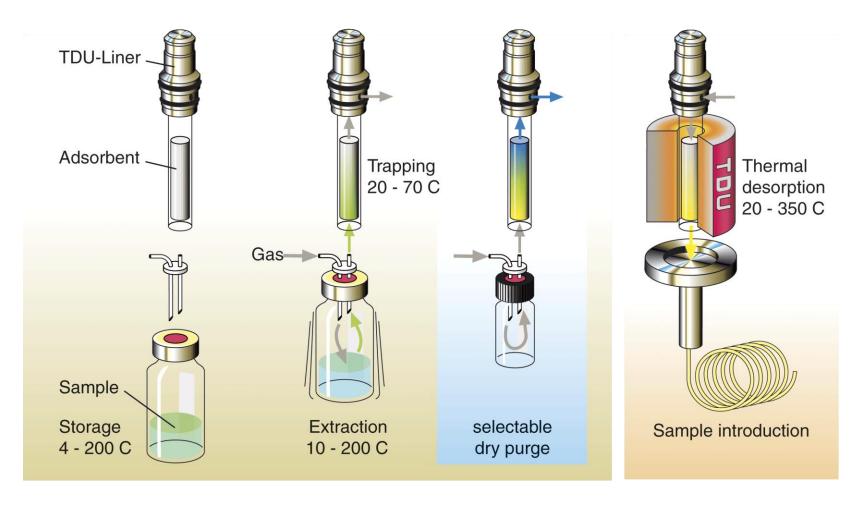


Time: 30 min

Waterbath T=45° C



New automated TDU/DHS method



Courtesy: Gerstel GmbH & Co. KG

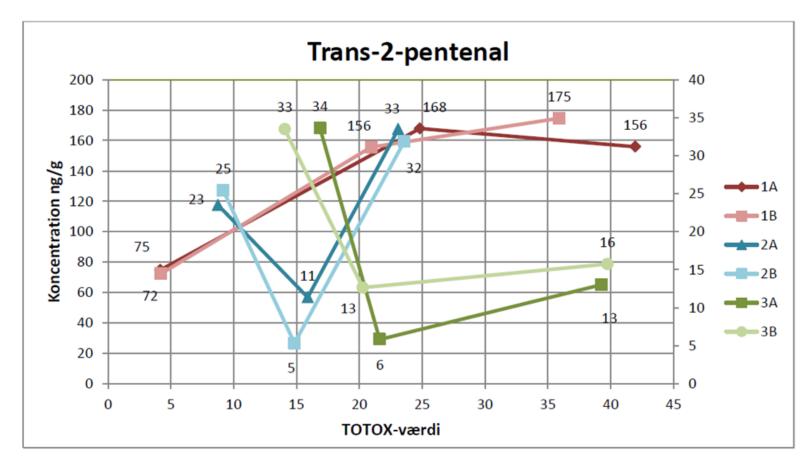


SPME and TDU sampling robot











Challenges and research needs

- For fish meal
 - Standard method for protein determination using Dumas principle?

- For fish oil (for human consumption):
 - An alternative to the AV method is needed
 - For headspace GC-MS there is no standard method and labs are doing the analysis in many different ways



Thank you for your attention!

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Research Group for Bioactives – Analysis and Application Charlotte Jacobsen (Professor & Group Leader) chja@food.dtu.dk