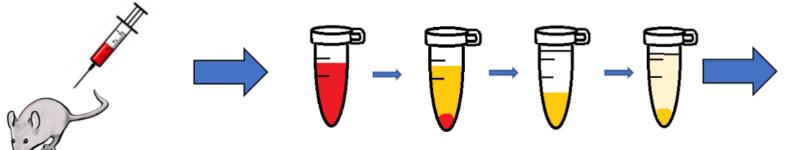


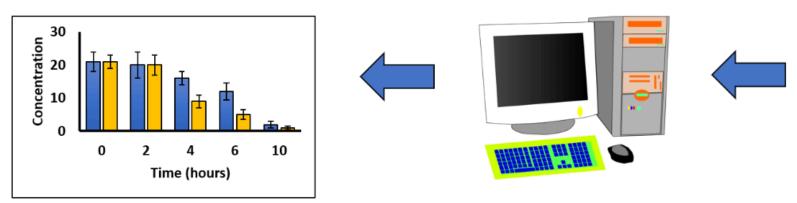
Table 2: Biomolecules purified in ATPS by affinity.

Biomolecule	Ligands attached to PEG	Recovery (%)	Purific. factor	Reference
Lactate dehydrogenase	Tryazine dye- Cibacron Blue F3G-A	81.3	7.4	LIN <i>et al.</i> (1998)
β-galactosidase	p-amino phe nyl-β-D- thiogal actopiran oside — (APGP)	83	6	SILVA <i>et al.</i> (1997)
Protein A	lgG human	87	-	SUZUKI <i>et al.</i> (1995)
Lactate dehydrogenase	Eudrogit-Cibacron Blue	54	11.7	GUOQIANG <i>et al.</i> (1994)
Penicillin acylase	Trimethylamina	97	25.7	GUAN <i>et al.</i> (1992)
Trypsin	Trypsin inhibitor	82	-	LUONG & NGUYEN (1990)

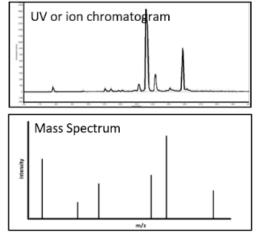


Sample collection





Results Data analysis



Data acquisition



Environmental and Food Analysis

Asst.Prof.Dr.Woravith Chansuvarn

Timeline: Analytical Process

Select method Sample preparation

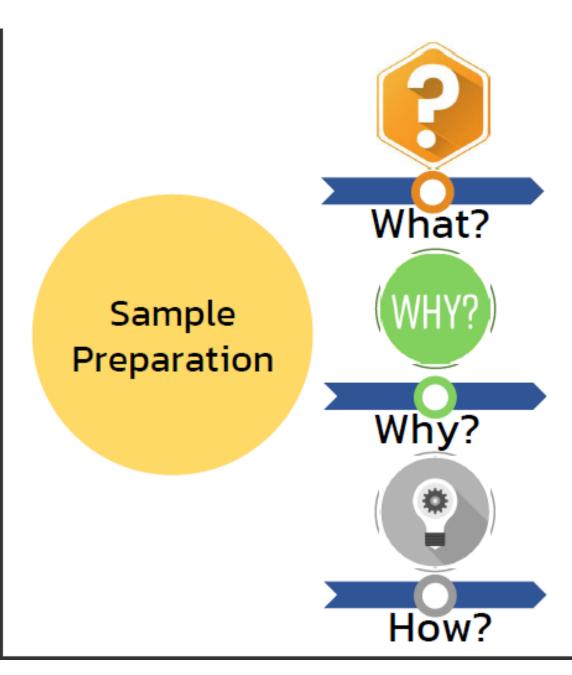
Evaluation



Sampling / Sample preservation

Measurement





- Standard method
- Protocols
- Validation method
- Limit permission and regulations
- CRM/SRM/TRM

Sample Preparation Perspective

Sample preparation is a process required for the transformation of a sample to make it amenable for chemical analysis or to improve the analysis.

The major goal of sample preparation is to prepare the sample for the separation/detection part of the analysis.

Sample preparation is the way in which a sample is treated to prepare for analysis. Sample preparation is carefully critical in analytical chemistry to accurately generate either a standard or unknown sample for a chemical measurement.

Sample preparation is common in many analyses and is developed to allow or to improve a specific analysis. This step may be the most time-consuming in an analysis and affects significantly the analytical information.

— The Ultimate Goals

- Transform analyte from a non-compatible environment into the instrumental compatible with analytical techniques
- Remove unwanted matrix components that may interfere with the analysis of the desired compound
- Improve limits of detection and/or quantitation (Enrichment analyte concentration)
- Separate/isolate of individual components from complex mixtures or matrix

Standard method



International standard



Regional standard



National standard



Association standard



Society standard



Publication

Developing method

วิธีมาตรฐานระดับนานาชาติ (International standard)

เป็นมาตรฐานที่ได้จากข้อตกลงร่วมกัน ของประเทศสมาชิกต่างๆ ที่มีความ สนใจร่วมกัน

วิธีมาตรฐาน ISO
 (International Standards
 Organization)



ISO/TS 15495 | IDF/RM 230:2010, Milk, milk products and infant formulae-Guidelines for the quantitative determination of melamine and cyanuric acid by LC-MS/MS

วิธีมาตรฐานระดับภูมิภาค (Regional standard)

เป็นมาตรฐานที่เกิดขึ้นจากการ ประชุม ปรึกษาหารือกันระหว่าง ประเทศในภูมิภาคเดียวกัน

• วิธีมาตรฐานของสหภาพยุโรป (European standard)



EN 1233:1996 Water quality-Determination of chromium-Atomic absorption spectrometric methods

EN 14084:2003 Foodstuffs-Determination of trace elements-Determination of lead, cadmium, zinc, copper and iron by AAS after microwave digestion

วิธีมาตรฐานระดับประเทศ (National standard)

เป็นมาตรฐานที่ได้จากการประชุมหารือ เพื่อหาข้อตกลงร่วมกันของ ผู้เกี่ยวข้องหลายภาคส่วนในประเทศ

- Japanese Industrial Standard (JIS)
- British Standard (BS)
- USEPA
- APHA
- AWWA
- NIOSH
- OSHA
- มอก.











วิธีมาตรฐานระดับสมาคม (Association standard)

เป็นมาตรฐานที่กำหนดขึ้นจากกลุ่ม บริษัทที่อยู่ในวงการค้าเดียวกัน หรือ เกิดจากข้อตกลงของกลุ่มบริษัท หรือโรงงานที่มีกิจกรรมของ อุตสาหกรรมเป็นอย่างเดียวกัน หรือ มีการผลิตของชนิดเดียวกัน

 AATCC (American Association of Textile Chemists and Colorists)



AATCC 112:2014

Formaldehyde release from fabric, Determination of: sealed Jar method

วิธีมาตรฐานรับรองโดยองค์กรทางวิชาการ

ปัจจุบันมีหลายองค์กรที่เป็นที่ยอมรับใน ระดับนานาชาติ

- ASTM
- IAOAC exércitable entratura écuations
- AOCS
- APPA
- AWWA
- EPA







ASTM D5630-13 Standard test method for ash content in plastics

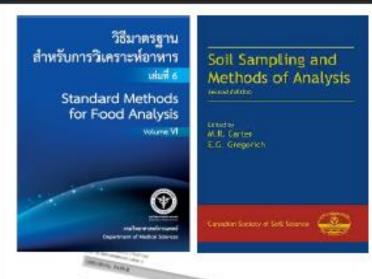
AOAC Official Method 999.10 lead, copper, zinc and iron in foods: Atomic absorption spectrometry after microwave digestion

AOCS official method Ca 12-55 Phosphorus

วิธีเผยแพร่ในตำรา/วารสารวิจัย (Publication method)

ส่วนใหญ่เป็นวิธีที่พัฒนาขึ้นใหม่

- วิธีมาตรฐานสำหรับการวิเคราะห์อาหาร โดยกรมวิทยาศาสตร์การแพทย์
- Analytical Chemistry
- Journal of Chromatography A
- Analytica Chimica Acta
- Food Chemistry
- Analyst
- Microchimica Acta
- Analytical Letters







AOAC Official Method 931.08 Formaldehyde in Food

First Action 1931

(See also 964.21 [see 44.5.14].)

UV-Vis method

A. Preparation of Test Portion

If food is solid or semisolid, macerate 100 g with 100 mL H2O in mortar. Transfer to 800 mL Kjeldahl flask, acidify with H3PO4, add 1 mL excess, connect with condenser through trap, and slowly distil 50 mL. For milk, dilute 100 mL with 100 mL H2O, and acidify and distil as for solids. With other liquid foods, acidify 200 mL and distil as for solids.

- B. Chromotropic Acid Test
- (a) Reagent.—Prepare saturated solution of 1,8-dihydroxynaphthalene-3,6-disulfonic acid (ca 500 mg/100 mL) in ca 72% H2SO4 (pour 150 mL H2SO4 into 100 mL H2O and cool). Solution is light straw-colored.
- (b) Test.—Place 5 mL reagent in test tube and add, with mixing, 1 mL distillate, A. Place in boiling H2O bath 15 min, and observe during heating period. Presence of HCHO is indicated by appearance of light to deep purple (depth of color depending on amount of HCHO present).

Reference: Z. Anal. Chem. 110, 22(1937).

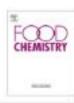
Source: http://files.foodmate.com/2013/files_2990.html



Contents lists available at ScienceDirect

Food Chemistry

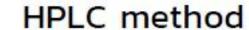




Analytical Methods

Determination of formaldehyde in food and feed by an in-house validated HPLC method

P. Wahed a, Md.A. Razzaq a, S. Dharmapuri b, M. Corrales b, a



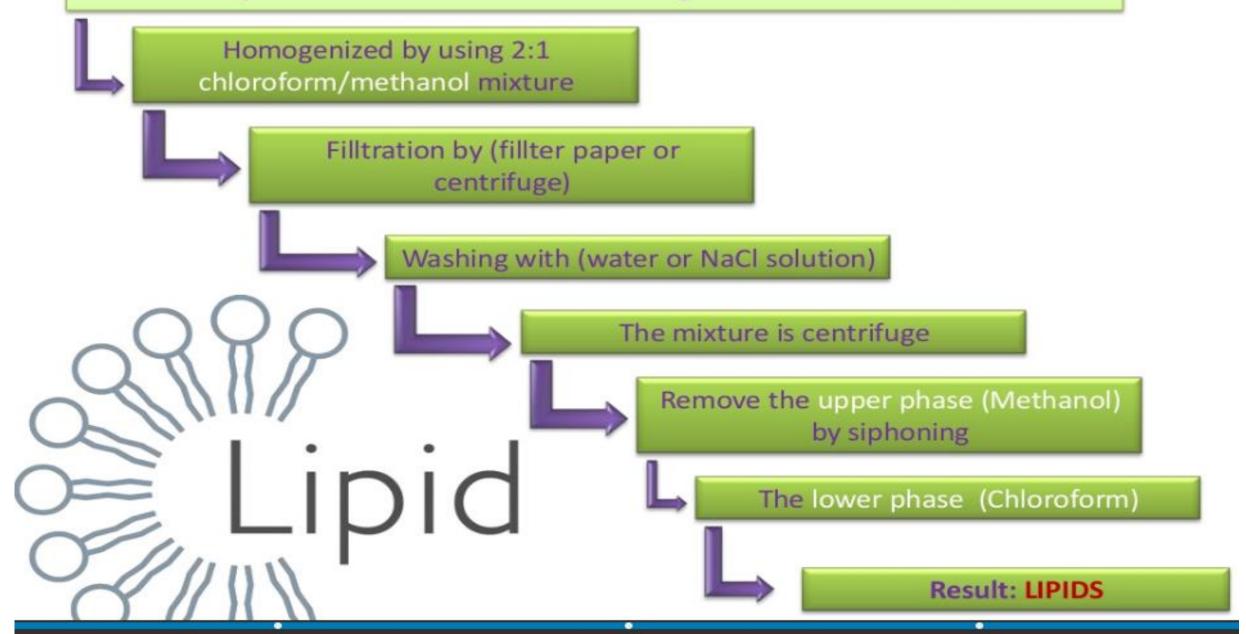
Derivatization kinetics followed the procedure described by Claeys et al. (2009) but was slightly modified. Edible parts of the food; fruit flesh and fish fillets were used for the analysis. For derivatization kinetics, mango samples were ground, homogenized and spiked with 10 mg/L of formaldehyde standard. To sample aliquots of 5 g, 5 mL of acetonitrile were added, and the sample vortexed and then sonicated for 30 min. The samples were centrifuged at 5000 rpm for 5 min and the supernatant was passed through a 90 mm diameter Whatman® 541 (Hardened Ashless) filter paper (SIGMA-Aldrich, Buchs SG, Switzerland). Two and half milliliter of 2,4 DNPH was added to the extract and mixed well. Samples were incubated at 40 °C for 30, 60, 90 and 120 min in a shaking water bath (model BS-11, Oxon, UK). Formaldehyde was quantitatively converted to its Schiff base in 60 min. In all experiments, derivatization time was set to 60 min. After incubation, the acetonitrile layer was collected, membrane filtered (0.45 µm) and injected into the HPLC.

^{*}National Food Safety Laboratory, Institute of Public Health, Mohakhali, Dhaku, Songladesh

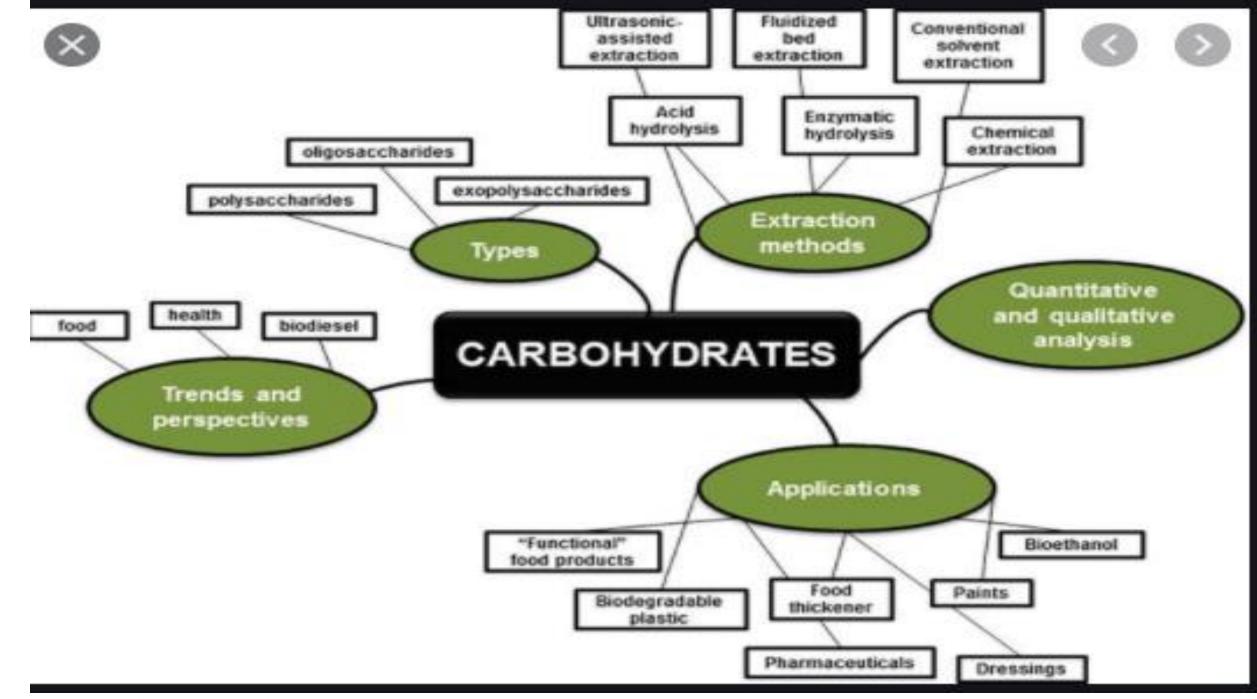
⁸ Food and Agriculture Organization of the United Nations, Dhanmondi, Dhaka, Bangialesk

Lipid preparation and extraction

The lipid extraction by Folch method



Carbohydrate preparation and extraction









REACTIONS OF SUCROSE

TEST	OBSERVATION	INFERENCE	
Molisch Test Purple ring at the junction of liquids		f two Sucrose is a carbohydrate	
Benedict's Test	No color change	It is a non reducing carbohydrate	
Barfoed's test No change in color		It is not a mono saccharide	
Seliwanoff test Cherry red color		Keto hexose containing disaccharide	
Hydrolysis The hydrolytic products give (Inversion) test positive reaction with Benedict's and Barfoed's reagents.		Confirmatory test for Sucrose	
Osazone test No reaction		Sucrose does not form osazone crystals	
	Molisch Test Benedict's Test Barfoed's test Seliwanoff test Hydrolysis (Inversion) test	Molisch Test Purple ring at the junction of two liquids Benedict's Test No color change Barfoed's test No change in color Seliwanoff test Cherry red color Hydrolysis (Inversion) test positive reaction with Benedict's and Barfoed's reagents.	

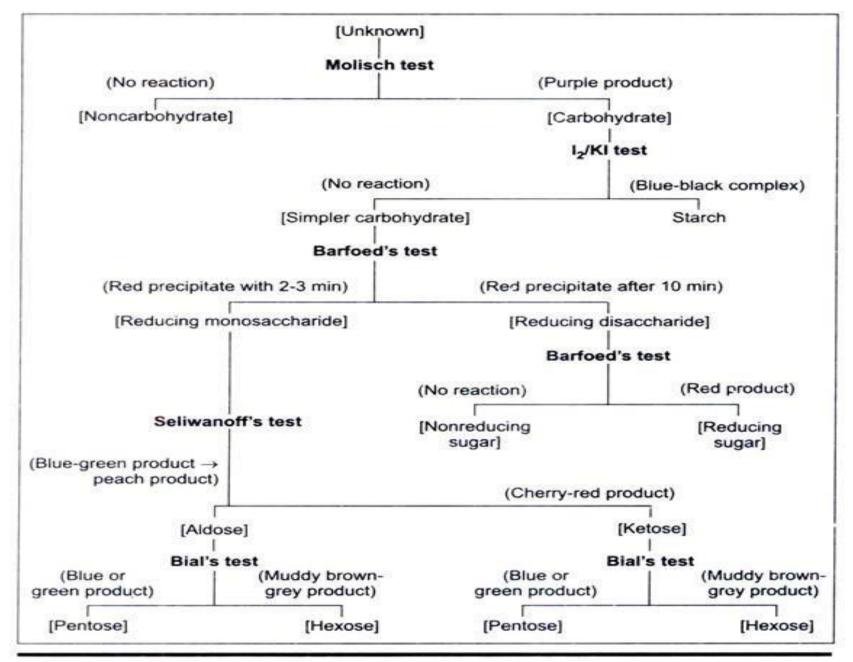
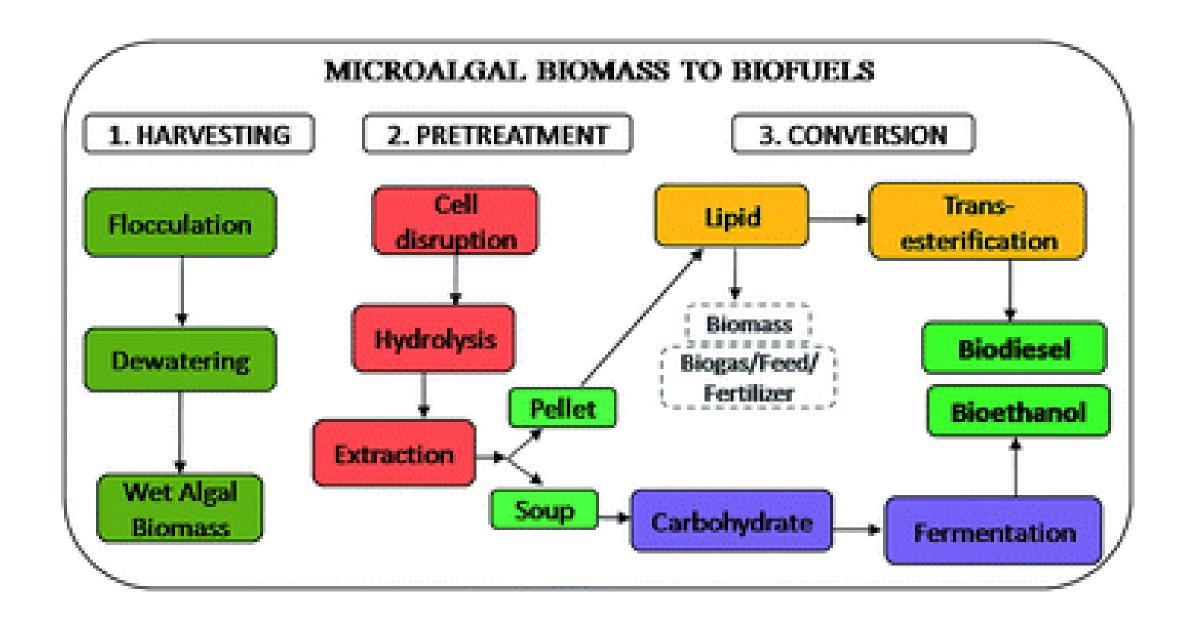
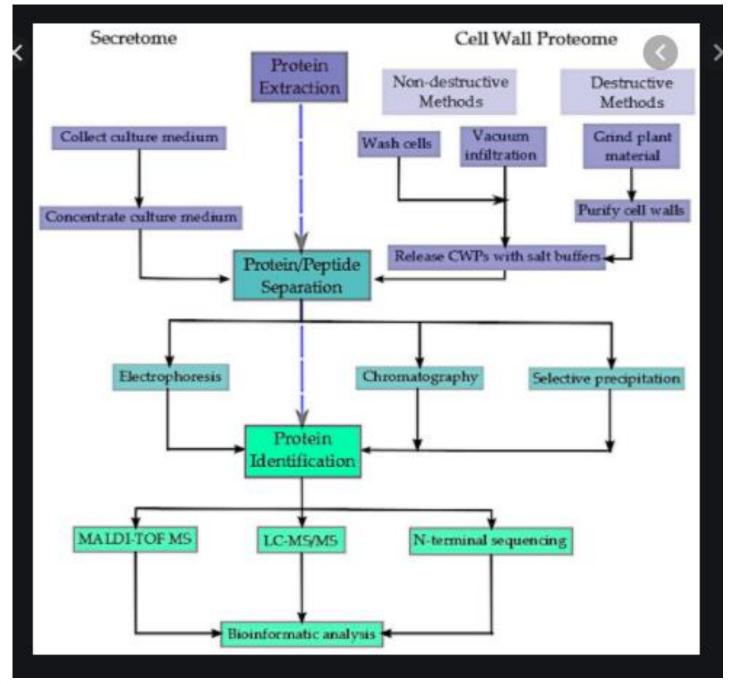
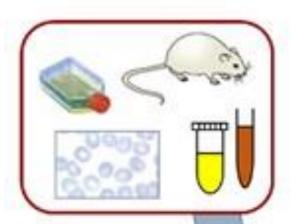


Fig. 18.1: Flow chart for classifying unknown carbohydrate



Protein preparation and extraction





Cell lysis/subcellular fractionation



Protein extraction

Protein enrichment/ fractionation



Enzymatic digestion



Animals

Identification of modulated proteins /PTMS
Identification of new interacting partners
....

Statistical analysis

Bioinformatic processing m/z

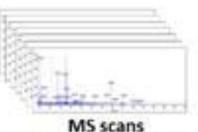


TopN

Data-dependent

analysis





(~ 50 000 scans/ analysis)

MS/M5 scans (~30 000 spectra / analysis)

nanoLC-column

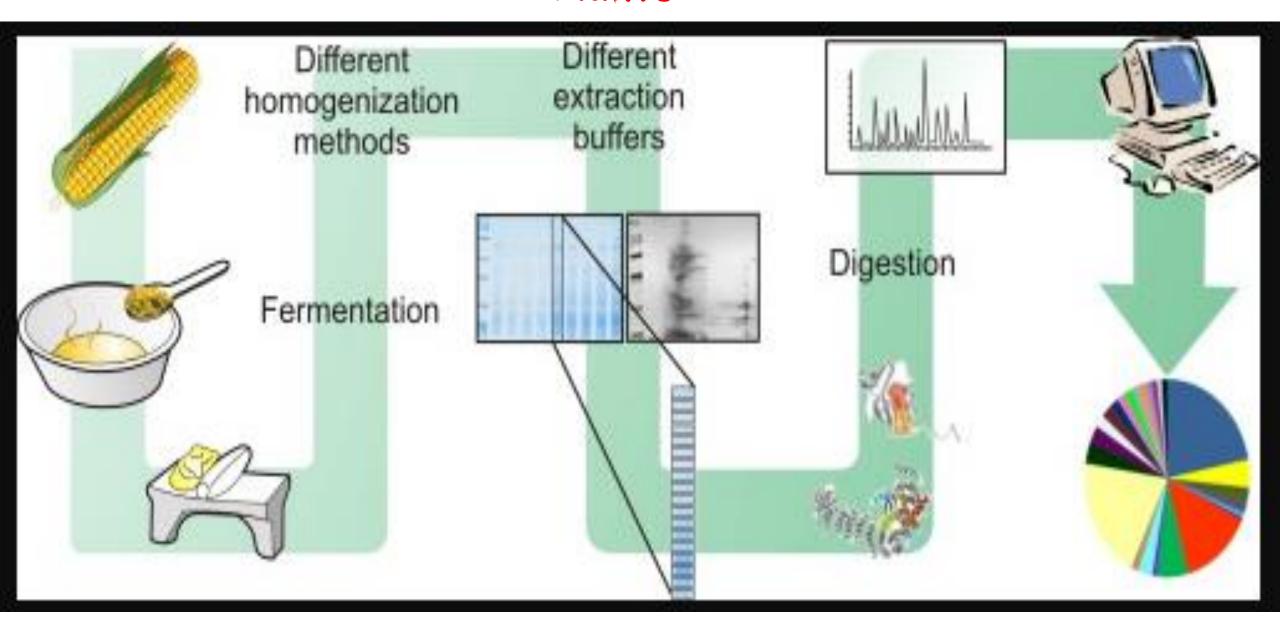
Peptide

enrichment,

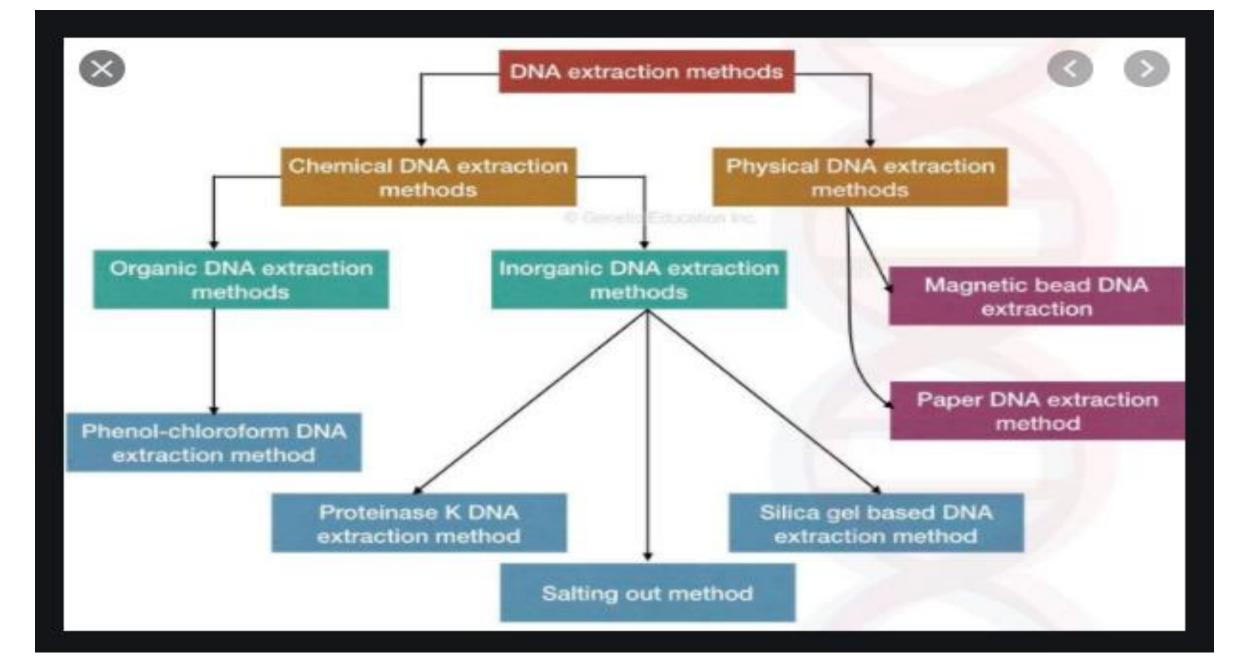
fractionation

Copyright from http://www.profiproteomics.fr/rd-services/

Plants



Nucleic acid preparation and extraction





Two-Step Lysis Method







RBCs lysed with an anionic detergent in the presence of a DNA stabilizer

WBCs lysed, gDNA and cellular RNA released into solution, RNA digesting treatment

Proteins removed by salt precipitation

DNA precipitated, washed & resuspended in hydration buffer

Cell lysis 1

Cell lysis 2

Contaminant Removal

DNA recovery

b. One-Step Lysis Method

RBCs & WBCs lysed, nuclei & mitochondria isolated from WBC by centrifugation

Pellet resuspended in denaturation buffer with a chaotropic salt & Protease

DNA precipitated, washed & resuspended in hydration buffer

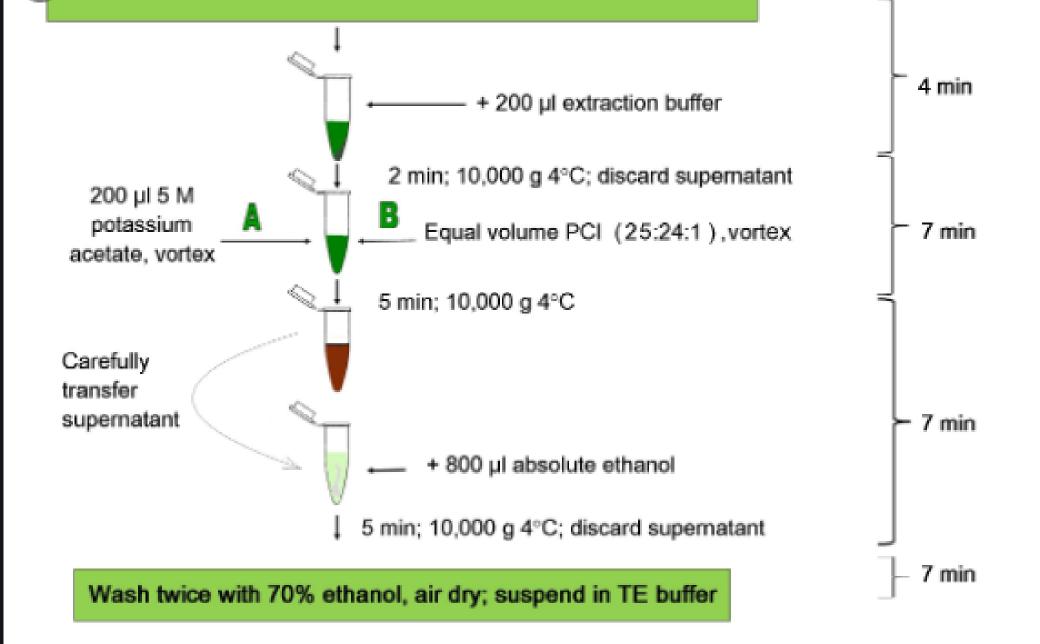


Cell lysis

Contaminant Removal

DNA recovery





The Ends