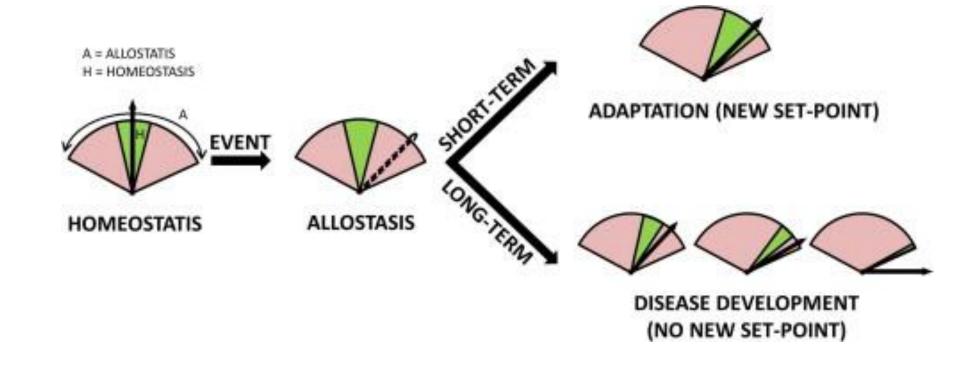
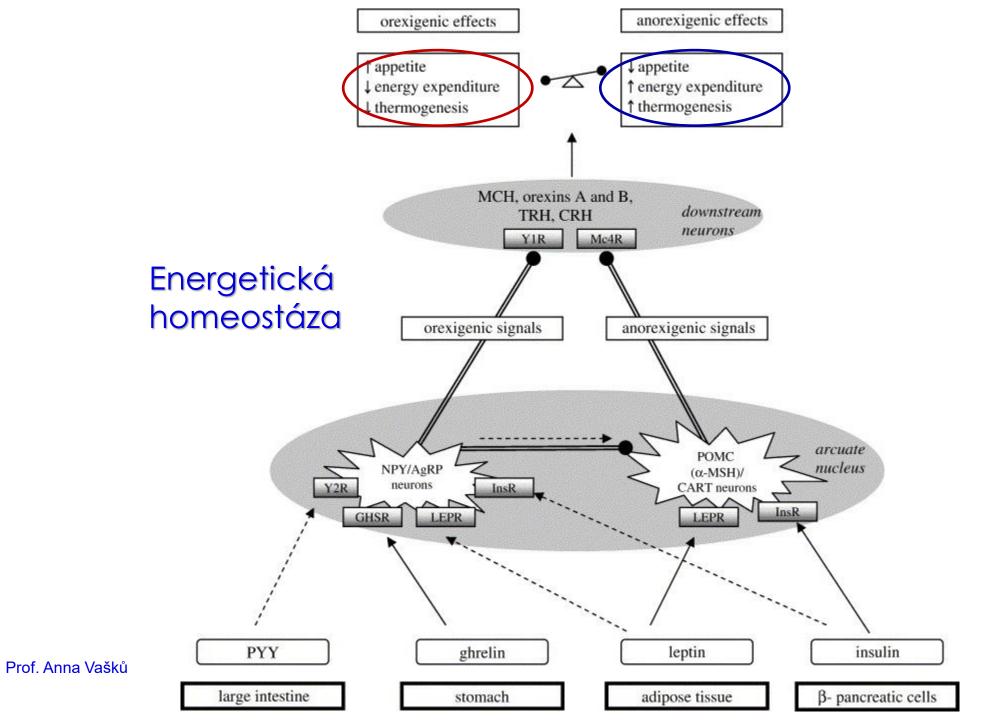


### Energy metabolism

NiP-1 28. 2. 2025



Bienertová-Vašků J, Zlámal F, Nečesánek I, Konečný D, Vasku A. PLoS One. 2016 Jan 15;11(1)

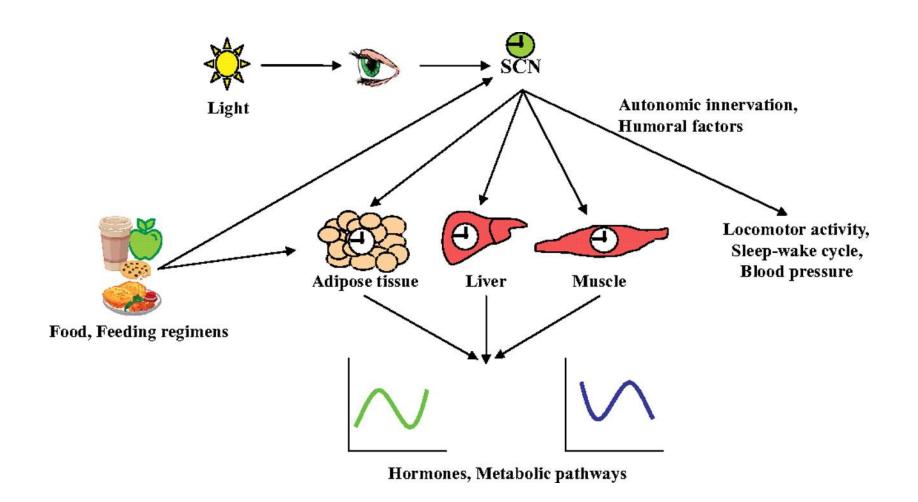


# Energy metabolism

- most food compounds are used as a energy source
- 1g of sugars 17.22kJ
- 1g of lipids 39.06kJ
- 1g of proteins 23.73kJ



### Resetting signals of the central and peripheral clocks.



Froy O Endocrine Reviews 2010;31:1-24



### Regulation of metabolic processes

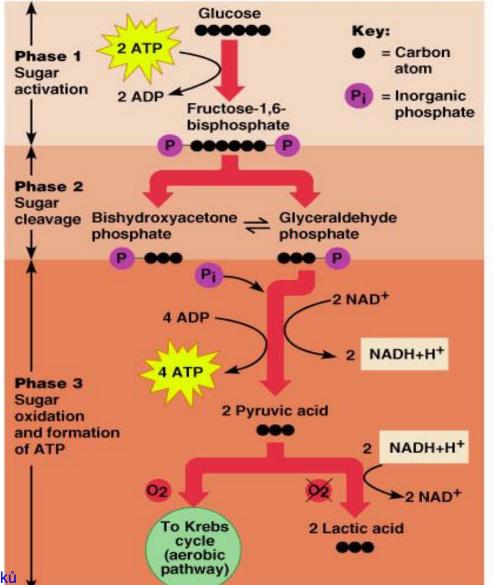
Neuro-immuno-endocrine regulation using hormones: (insulin, glucagon, growth hormone, glucocorticoids, T4 and T3, reproduction hormones:

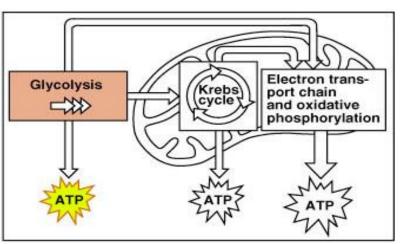
- ✓ Liver
- ✓ Adipose tissue
- ✓ Skin
- √ Kidneys
- ✓ Respiratory and cardiovascular system



Compound	Transcription factor			
Fatty acids Cholesterol	PPARs, SREBPs, LXR, HNF4, ChREBP SREBPs, LXRs, FXR			
Glucose	USFs, SREBPs, ChREBP			
Amino acids	C/EBPs			
Vitamin A Vitamin D Vitamin E	RAR, RXR VDR PXR			
Calcium Iron Zinc	Calcineurin/NF-ATs IRP1, IRP2 MTF1			
Other food components				
Flavonoids Xenobiotics	ER, NFkB, AP1 CAR, PXR	nature REVIEWS GENETICS		
	Fatty acids Cholesterol Glucose Amino acids  Vitamin A Vitamin D Vitamin E Calcium Iron Zinc  nents Flavonoids	Fatty acids PPARs, SREBPs, LXR, HNF4, Che Cholesterol SREBPs, LXRs, FXR  Glucose USFs, SREBPs, ChREBP  Amino acids C/EBPs  Vitamin A RAR, RXR Vitamin D VDR Vitamin E PXR  Calcium Calcineurin/NF-ATs Iron IRP1, IRP2 Zinc MTF1  nents  Flavonoids ER, NFkB, AP1		

### **Glycolysis**





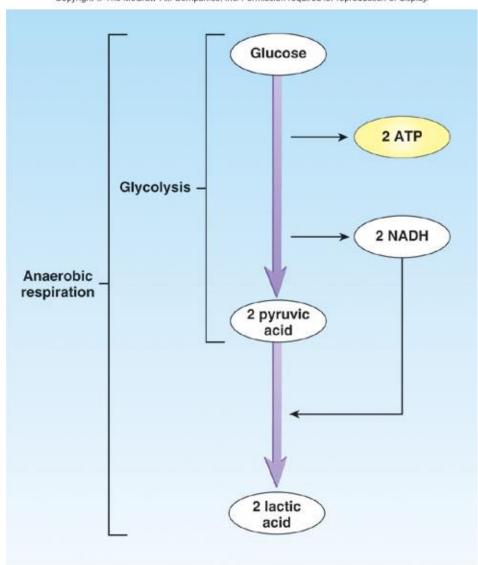


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TABLE 3–8 Characteristics of Glycoly	rsis
Entering substrates	Glucose and other monosaccharides
Enzyme location	Cytosol
Net ATP production	2 ATP formed directly per molecule of glucose entering pathway can be produced in the absence of oxygen (anaerobically)
Coenzyme production	2 NADH + 2 H <sup>+</sup> formed under aerobic conditions
Final products	Pyruvate—under aerobic conditions Lactate—under anaerobic conditions
Net reaction	
Aerobic:	Glucose + 2 ADP + 2 $P_i$ + 2 $NAD^+ \longrightarrow$ 2 pyruvate + 2 ATP + 2 $NADH$ + 2 $H^+$ + 2 $H_2O$
Anaerobic:	Glucose + 2 ADP + 2 $P_1 \longrightarrow 2$ lactate + 2 ATP + 2 $H_2O$

### Anaerobic respiration

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$$CH_3 \longrightarrow C=O+CoA-SH \longrightarrow CH_3$$

$$C=O+CoA-SH \longrightarrow C=O+CO_2$$

$$COOH$$

$$S-CoA$$

### Pyruvic acid

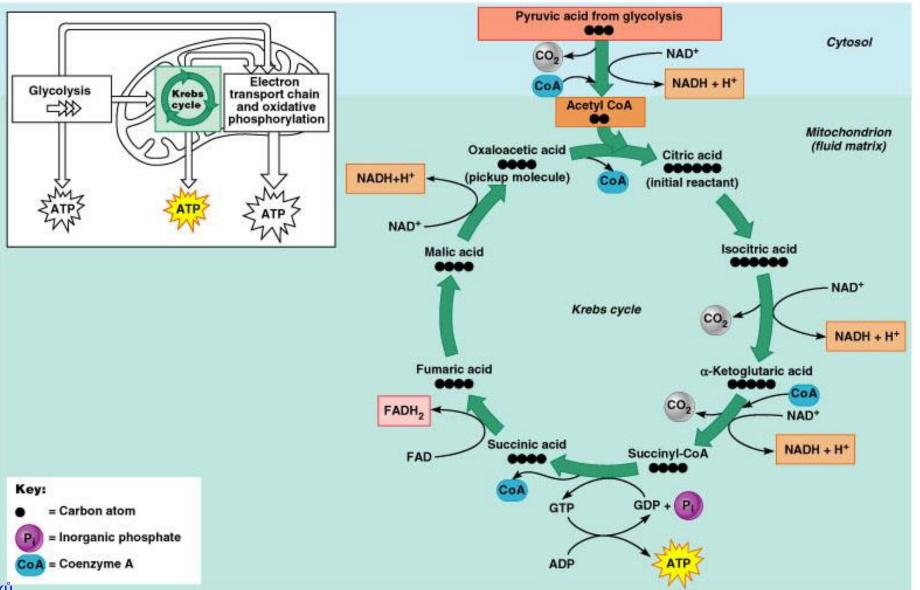
vate

Each transition of pyruvate to acetyl coenzyme A yields one NADH and one CO<sub>2</sub>.

The acetyl coenzyme A then enters the Krebs cycle.

Acetyl coenzyme A

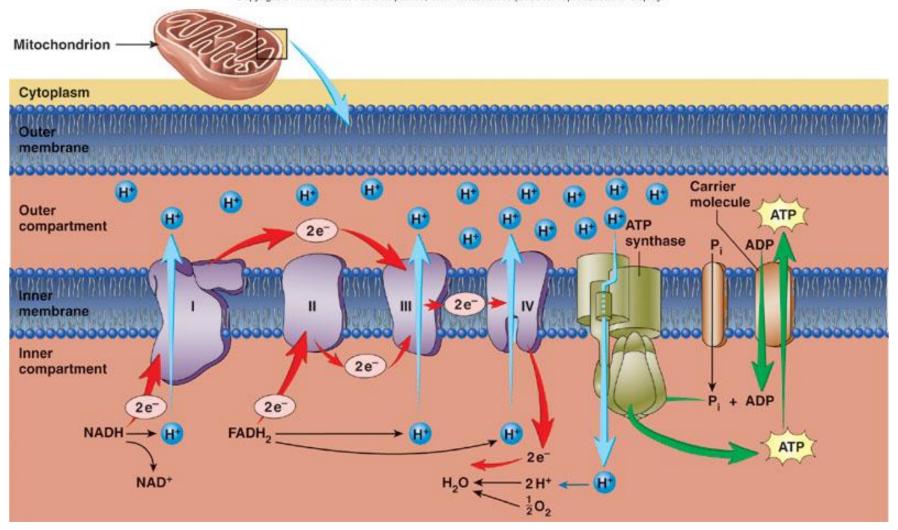
### Krebs cycle





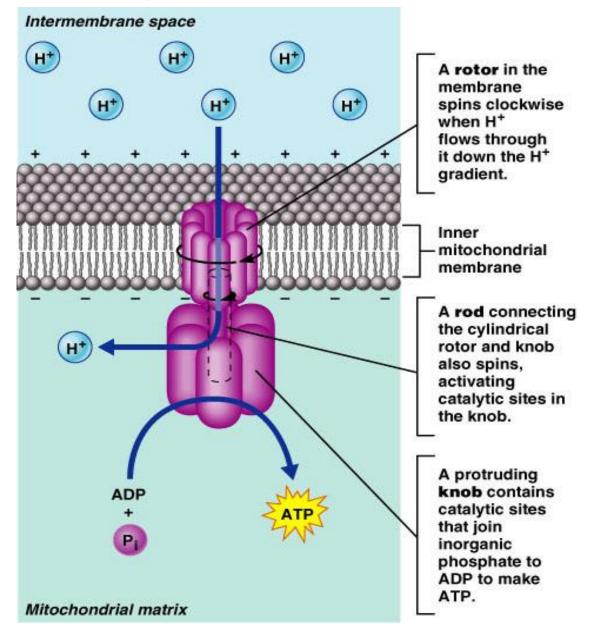
### Oxidative phosphorylation

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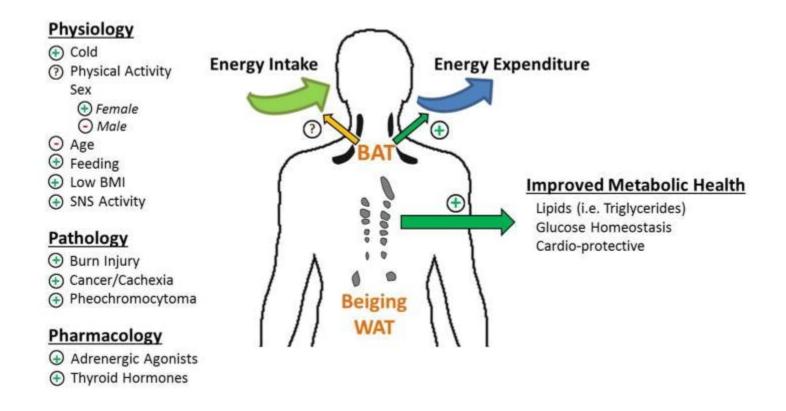




### ATP synthase



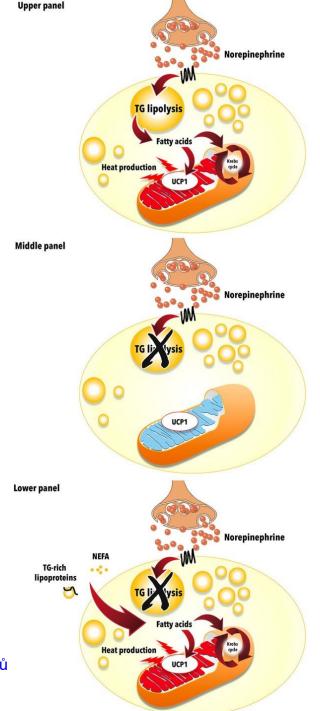




#### Mediators of BAT (or Beige) Amount/Activity and the Impact on Metabolism

There are several physiological, pathological, and pharmacological conditions that are known to mediate the amount of BAT and its activity (on the left). Specifically, "+" and "-" indicates that BAT is "stimulated" or "depressed" by the following condition or stimuli, respectively. On the right are the potential effects of BAT on energy intake, energy expenditure, and cardiometabolic health.





Uncoupling protein 1 (UCP1)-mediated brown adipose tissue thermogenesis.

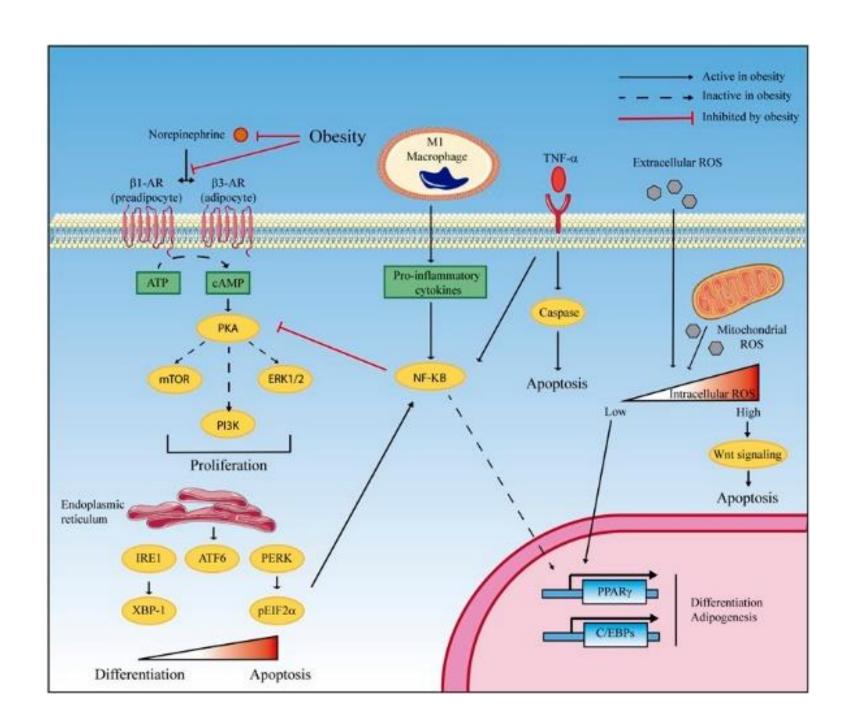
**Upper panel:** Brown adipose tissue UCP1-mediated thermogenesis is activated by fatty acids produced via norepinephrine-induced intracellular triglyceride (TG) lipolysis during cold exposure.

Middle panel: Acute pharmacological inhibition of intracellular TG lipolysis blunts brown adipose tissue thermogenesis via reduction of intracellular fatty acids availability.

**Lower panel:** Genetic deletion-mediated issues leads to increased reliance on circulating nonepinephrine inhibition of intracellular TG lipolysis in brown adipose non esterified fatty acids (NEFA) and triglyceride (TG)-rich lipoproteins to sustain UCP1mediated thermogenesis.



# Depletion of BAT induced by obesity

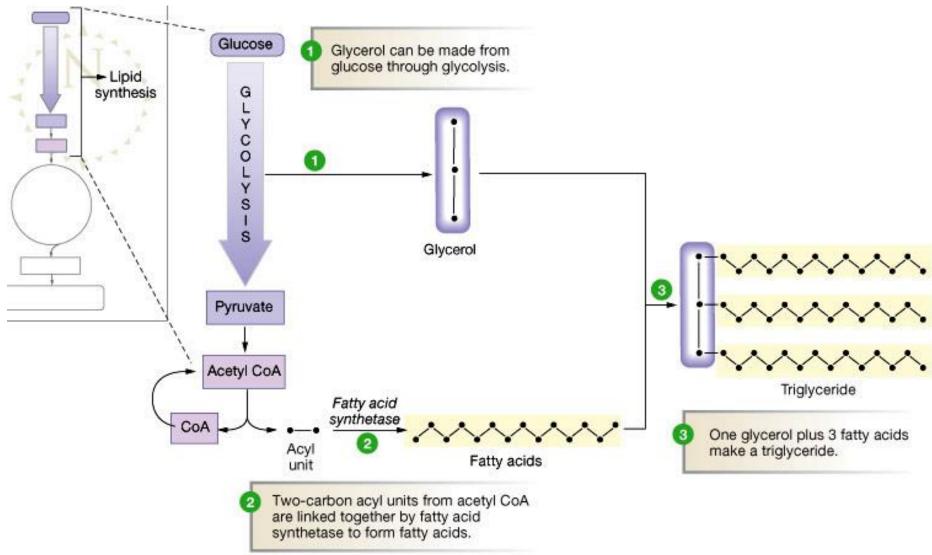


### Triglyceride Glucose Triglycerides break down into glycerol and 3 fatty acids. → Glycerol-Glycerol becomes a glycolysis substrate. COLYSIS Fatty acid Pyruvate Cytosol β-oxidation chops 2-carbon acyl units off the fatty acids. β-oxidation \*CO2 Acetyl CoA -Acyl units become acetyl CoA and can be used in the citric acid cycle. CoA -CITRIC ACID CYCLE Mitochondrial matrix

## Lipolysis



### Synthesis of lipids





### **Proteins**

- Main building material of cells and tissues, enzymes, some hormones and compounds of pigments
- Food sources: animal(contain all AAs) and plant (not all contain essential AAs)proteins ⇒ digestion in stomach pepsin (from pepsinogen, activated by HCL) and small intestine trypsin, aminopeptidases (cleave N-terminal AAs), carboxypeptidases (cleave C-terminal AAs), dipeptidases (cleave dipeptides) and AAs ⇒ absorption to v. portae in presence of vit. B6
   Condition for adequate protein digestion- they must denaturized (cooking)



# Protein metabolism

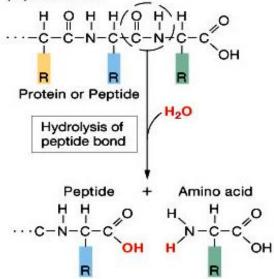
- Non essential amino acids can be formed by transamination (= transfer of amino acid group to keto acid).
- When using for energy formation, they undergo oxidative deamination. NH3 and keto acids are side products of the reaction. NH3 is changed to urea and excreted by urine.
- Amino acids are not stored in the body.



### Catabolism of proteins

#### (a) Protein catabolism

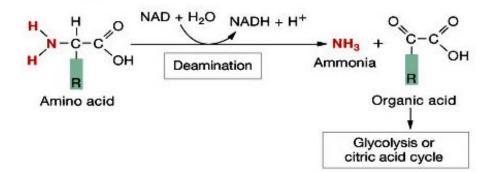
Proteins are broken into amino acids by hydrolysis of their peptide bonds.



(c) Ammonia is toxic and must be converted to urea.

#### (b) Deamination

Removal of the amino group from an amino acid creates ammonia and an organic acid.





### Summary: metabolic reactions of sugars

#### Thumbnail Summary of Metabolic Reactions

#### Carbohydrates

Cellular respiration Reactions that together complete the oxidation of glucose, yielding CO2, H2O, and ATP

Glycolysis Conversion of glucose to pyruvic acid

Glycogenesis Polymerization of glucose to form glycogen

Glycogenolysis Hydrolysis of alycogen to alucose monomers

Formation of glucose from noncarbohydrate precursors Gluconeogenesis

Krebs cycle Complete breakdown of pyruvic acid to CO2, yielding small amounts of ATP and reduced

coenzymes

Electron transport

chain

Energy-yielding reactions that split H removed during oxidations to H+ and e- and create a

proton gradient used to bond ADP to Pi, forming ATP



# Summary: metabolic reactions of lipids and proteins

)	5258 00 WMANN WATERS VIEW DOOR IN 1658
TABLE 24.4 /	Thumbnail Summary of Metabolic Reactions

Lipids

Beta oxidation Conversion of fatty acids to acetyl CoA

Lipolysis Breakdown of lipids to fatty acids and glycerol

Lipogenesis Formation of lipids from acetyl CoA and glyceraldehyde phosphate

**Proteins** 

Transfer of an amine group from an amino acid to α-ketoglutaric acid, thereby transforming Transamination

α-ketoglutaric acid to glutamic acid

Oxidative deamination Removal of an amine group from glutamic acid as ammonia and regenerating α-ketoglutaric

acid (NH3 is converted to urea by the liver)

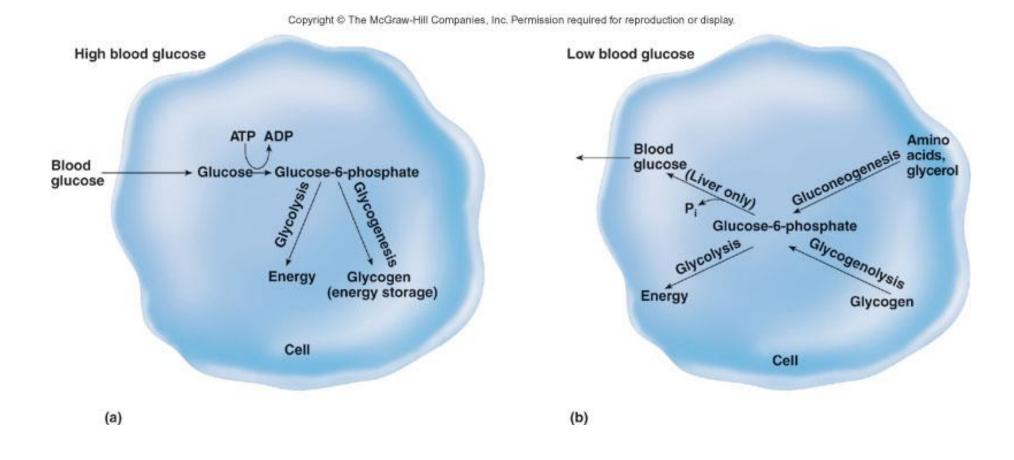


### Metabolism

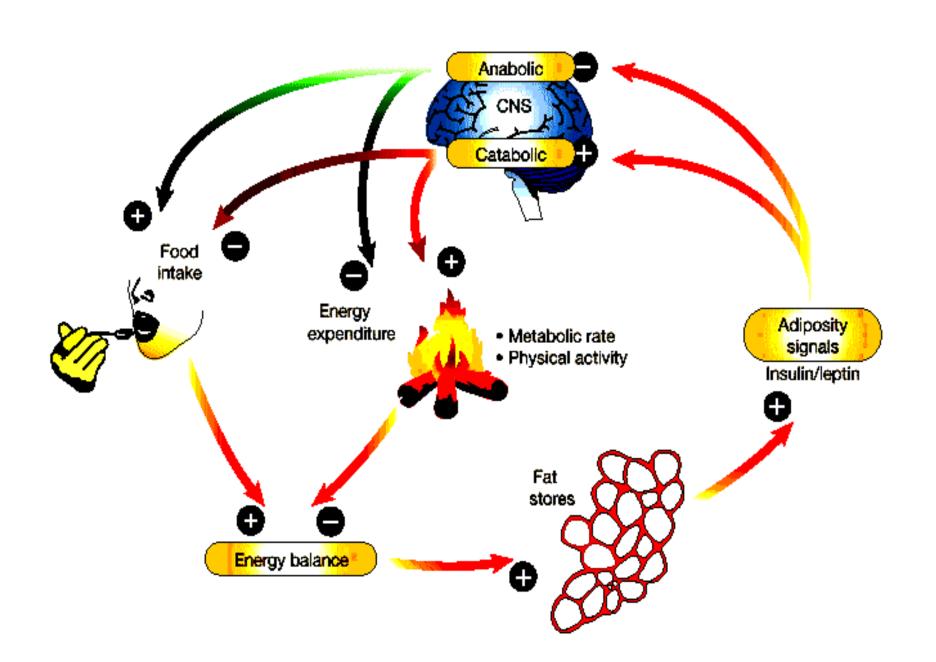
- ➤ Quantitative evaluation (energetic)
- ➤ Qualitative evaluation (sufficient and suitable proportion of proteins, lipids and sugars)
- ➤ Anabolism
- ➤ Catabolism



# Internal changes of nutrition molecules







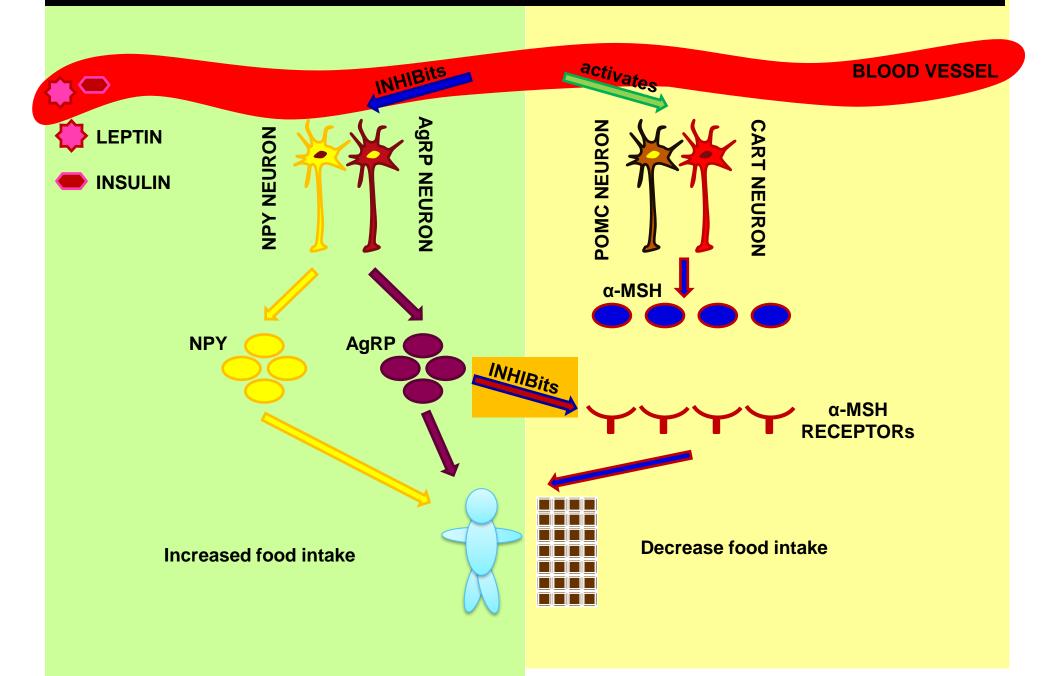


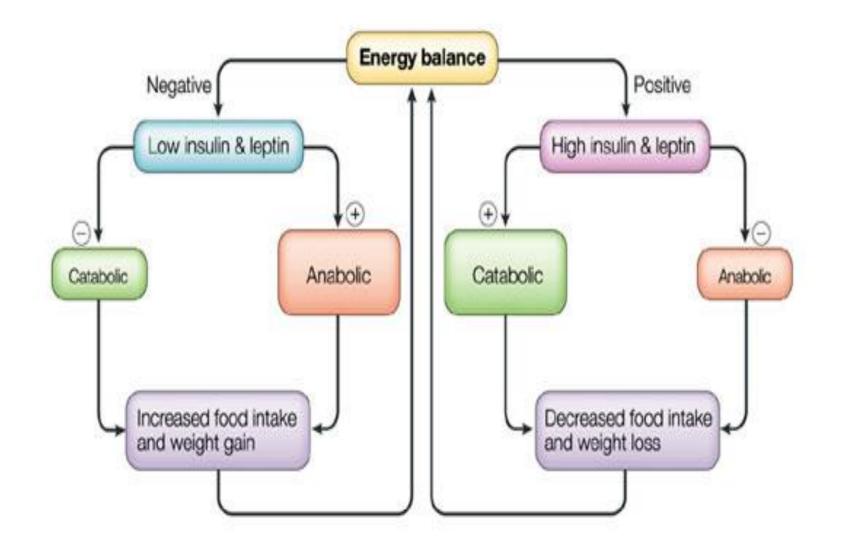
### Metabolism of substance:

- -anabolism = more complicated products are synthetized from simple absorbed compounds (s.c. assimilation) – energy is used (endergonic reaction)
- --catabolism = portion of absorbed compounds is cleaved to simpler ones (dissimilation) – energy is released (exergonic reaction)

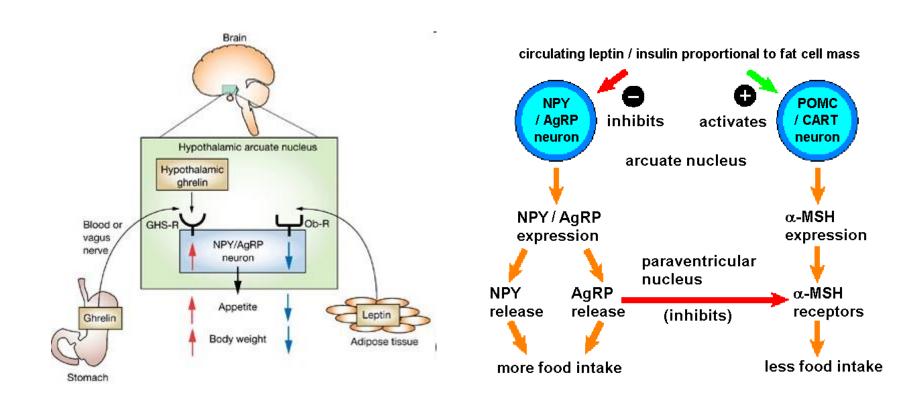


### OREXIGENIC-ANOREXIGENIC PATHWAYS

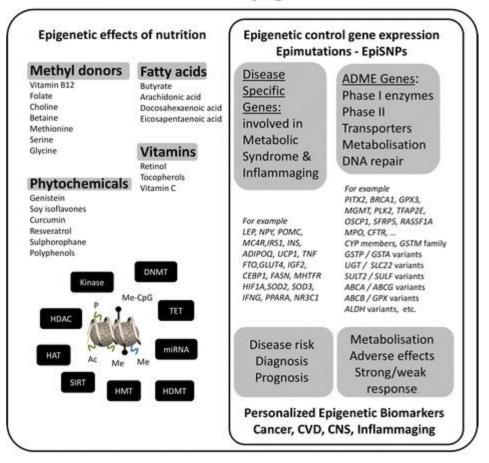




## Regulation of food appetite



#### **Nutritional Epigenetics**



2015 Mar 25;7(1):33. doi: 10.1186/s13148-015-0068-2. eCollection 2015. From inflammaging to healthy aging by dietary lifestyle choices: is epigenetics the key to personalized nutrition?

Vel Szic KS<sup>1</sup>,

Vel Szic KS<sup>1</sup>,

Declerck K<sup>1</sup>,

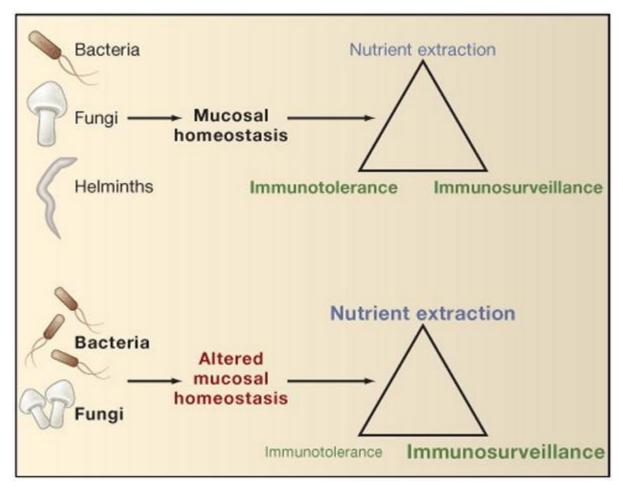
Vidaković M<sup>2</sup>,

Vanden Berghe W<sup>1</sup>.

#### **Personalized Nutrition**

#### Overview of the mechanisms and consequences of epigenetic regulation by nutritional compounds.

Modulation of different classes of chromatin writers-erasers by phytochemicals (left panel). Genes encoding absorption, distribution, metabolism, and excretion (ADME) proteins can be epigenetically regulated and thereby determine individual nutritional responses. Epigenetic modification of disease-related genes can contribute to diagnosis (biomarker) as well as disease prevention or progression (right panel).



The loss of universal helminth infection as occurred in earlier human evolution may alter the numbers or types of bacterial and fungal commensals and thus affect normal mucosal tissue homeostasis. In susceptible or highly exposed individuals, such alterations might alter the balance between immunotolerance, immunosurveillance and nutrient extraction. This imbalance may contribute to the appearance of inflammatory systemic dysregulation at mucosal surfaces, resulting in increases in asthma and allergic diseases, particularly in the setting of environmental changes that have increased exposure to indoor allergens and pollutants, and even to increases in obesity, which can be a risk factor for severe asthma.

### Catabolic states

Are induced by dysregulation of metabolic events by inflammation( cytokines), stress (A, GKs), chronic immobilization.

- Acute severe diseases (adaptation on starvation decreases, protein malnutrition can develop).
- Cancer cachexia (cytokines TNF, IL-1 and IL-6).
- Trauma, burns, fever, painful states, AIDS (wasting syndrome).



### Organ changes in protein and energy deficits

- Loss of body weight (drop of 40% is leading to death).
- ➤ECT volume is nor changing or relative expansion of ECT against ICT. When oncotic pressure was decreasing, edema could develop.
- ➤ Decreased cardiac output.
- ➤ Decreased function of respiratory system due to decreased contractility of respiratory muscles.
- > Decreased stomach motility and stomach production of HCL
- ➤ Decreased exocrine function of pancreas
- ➤ Decrease liver mass with decrease proteins, lipids and glycogen in secondary malnutrition
- In primary malnutrition liver mass increased due to lipids infiltration and increased production of glycogen.



### Organ changes in protein and energy deficits

- Kidney weight decreases.
- ➤Increased ECT volume due to decreased osmotic gradient in the kidney.
- Increased secretion in endocrine system.
- ➤Immunodeficiency state (decrease of non specific immune functions).
- Atrophy of the skin end GIT epithelia with functional lesions of barriers against external environment.
- Decreased wound healing in severe protein malnutrition.



## The human starvation

- response is unique among animals in that human brains do not require the ingestion of glucose to function. During starvation, less than half the energy used by the brain comes from metabolized glucose. Because the human brain can use ketone bodies as major fuel sources, the body is not forced to break down skeletal muscles at a high rate, thereby maintaining both cognitive function and mobility for up to several weeks.

#### Human starvation

- This response is extremely important in <u>human evolution</u> and allowed for humans to continue to find food effectively even in the face of prolonged starvation.
- Initially, the level of <u>insulin</u> in circulation drops and the levels of <u>glucagon</u>,
   <u>epinephrine</u> and <u>norepinephrine</u> rise. At this time, there is an up-regulation of <u>glycogenolysis</u>, <u>gluconeogenesis</u>, <u>lipolysis</u>, and <u>ketogenesis</u>.
- The body's glycogen stores are consumed in about 24 hours. In a normal 70 kg adult, only about 8,000 kilojoules of glycogen are stored in the body (mostly in the striated muscles).

## Human starvation

- The body also engages in gluconeogenesis to convert glycerol and glucogenic amino acids into glucose for metabolism.
- Another adaptation is the <u>Cori cycle</u>, which involves shuttling lipid-derived energy in glucose to peripheral glycolytic tissues, which in turn send the <u>lactate</u> back to the <u>liver</u> for resynthesis to glucose.
   Because of these processes, blood glucose levels remain relatively stable during prolonged starvation.



#### Human starvation

 However, the main source of energy during prolonged starvation is derived from triglycerides. Triglycerides are broken down to fatty acids via lipolysis. Epinephrine precipitates lipolysis by activating protein kinase A, which phosphorylates hormone sensitive lipase (HSL) and perilipin. These enzymes, along with CGI-58 and adipose triglyceride lipase (ATGL), complex at the surface of lipid droplets. The concerted action of ATGL and HSL liberates the first two fatty acids. Cellular monoacylglycerol lipase (MGL), liberates the final fatty acid. The remaining glycerol enters gluconeogenesis.

#### Malnutrition

Malnutrition is a disorder in body composition in which inadequate macronutrient (protein, carbohydrate, and fat) or micronutrient (vitamins, minerals, and trace elements) intake results in decreased body mass, reduced organ mass, and most importantly, decreased organ function.



## Nutrition: proteins

– Adequate intake of protein is one of the key nutritional factors to maintain independence, predominantly by preventing loss of muscle mass and strength (sarcopenia), frailty and associated comorbidities in later life.



## Sarcopenia

– A gradual decline in muscle mass is observed from the third decade of life with a 30–50% decrease reported between the ages of 40 and 80. Muscle strength is correlated with muscle mass and rapidly declines after the age of 50. The beginning of the fourth decade of life might therefore be interpreted as the time when muscle ageing process begins and for this reason it is the optimal time for implementing appropriate dietary changes, to prevent or delay the onset of sarcopenia.



## Type D behaviour

- is characterized by the joint tendency to experience negative emotions and to inhibit these emotions while avoiding social contacts with others. The observation that cardiac patients with type D personality are at increased risk for cardiovascular morbidity and mortality underlines the importance of examining both acute (e.g. major depression) and chronic (e.g. certain personality features) factors in patients at risk for coronary events.
- Both type D dimensions (negative affectivity and social inhibition) are associated with greater cortisol reactivity to stress. Elevated cortisol may be a mediating factor in the association between type D personality and the increased risk for coronary heart disease and, possibly, other medical disorders.



## Personality type D ("distressed" personality)

- Type-D denotes the synergistic effect of negative affectivity (tendency to experience negative emotions) and social inhibition (tendency to inhibit self-expression).
- As a result, type-D patients experience more feelings of anxiety, depression, and anger, but inhibit self-expression in order to avoid disapproval by others.
- Type-D is associated with a four- to fivefold increased risk of death or myocardial infarction in cardiac patients.
- Type-D personality was positively associated with the cortisol-awakening response, independently of age, sex, and body mass.



## Proteinuria as a cause of a loss of proteins

#### Table 1 | Types of proteinuria

Types	Characteristics
Glomerular	Most common form, up to 90%
	Feature of chronic kidney disease
	Loss of albumin and higher molecular weight proteins
Tubular	Low molecular weight proteins, such as
	β2-microglobulin
Overflow	Increased production, that is, light chains in multiple
	myeloma
Post-exercise	Transient benign
	Can be up to 10 g/day
Post-prandial	Transient physiological proteinuria
	Possibly through insulin action in podocytes
Infection- associated	Physiological response
	Mediated by toll-receptors
	Possibly involved in clearing pathogens from the
	circulation

Table 1. The principle genes involved in congenital nephrotic syndrome and in associated syndromes

Genes	Locus	Protein	Phenotype
AD			
WT1	11P13	Wilms tumor 1	IDMS, DDS, Frasier syndrome, WAGR syndrome, ISRNS
LMX1B	17q11	Lim homeobox transcription factor $1-\beta$	Nail-patella syndrome
INF2	14q32.33	Inverted formin-2	FSGS
CD2AP	6p12	CD2-associated protein	FSGS (adult)
AR			
NPHS1	19q13.1	Nephrin	CNF
NPHS2	1q25-31	Podocin	Idopathic CNS, SRNS
LAMB2	3p21	Laminin $\beta$ 2 chain	Pierson's syndrome
PLCE1	10q23	Phospholipse C epsilon 1	SRNS, DMS
PDSS2	6q21	Decaprenyl disphosphate synthase, subunit 2	NS with Leigh syndrome
ITGA3	17q21.33	Integrin $\alpha 3$	NS with intestitial lung disease
ARHGDIA	17q25.3	Rho GDP dissociation in inhibitor 2	Idiopathic CNS
SCARB2	4q21.1	Scuvenger receptor class B, number 2	Action myoclonus-renal failure syndrome
Unknown			Galloway-Mowat syndrome

AD: autosomal dominant; AR: autosomal recessive; IDMS: idiopathic diffuse mesangial sclerosis; DDS: Denys-Drash syndrome; WAGR: Wilm's tumor, aniridia, genitourinary abnormalities and mental retardation; FSGS: focal segmental glomerulosclerosis; CNS: congenital nephrotic syndrome; CNF: CNS of the Finnish type; SRNS: steroid-resistant nephrotic syndrome; DMS: Denys-Drash syndrome.

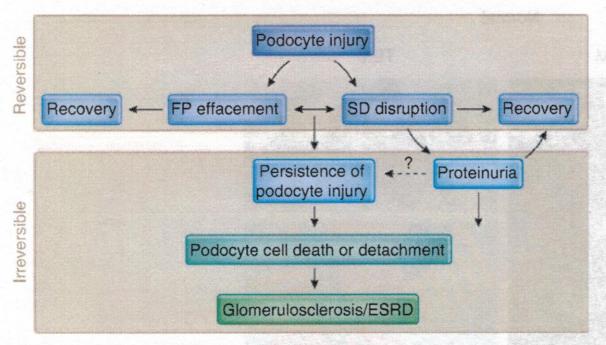


Figure 2 | Consequences of podocyte injury. Podocytes can be injured in many human and experimental glomerular diseases, leading to structural changes, such as foot processes (FP) effacement and slit diaphragm (SD) disruption that are reversible. Persistence of podocyte injury can cause cell death or detachment of podocytes from the glomerular basement membrane (GBM). The resulting loss of podocyte will ultimately lead to irreversible glomerulosclerosis and end-stage renal failure (ESRD). The role of proteinuria in the progression of ESRD is a matter of debate. In some patients, nephrotic-range proteinuria can persist over years without progression to ESRD.

**Mundel P**, Reiser J. **Kidney Int**. 2010 Apr;77(7):571-80.

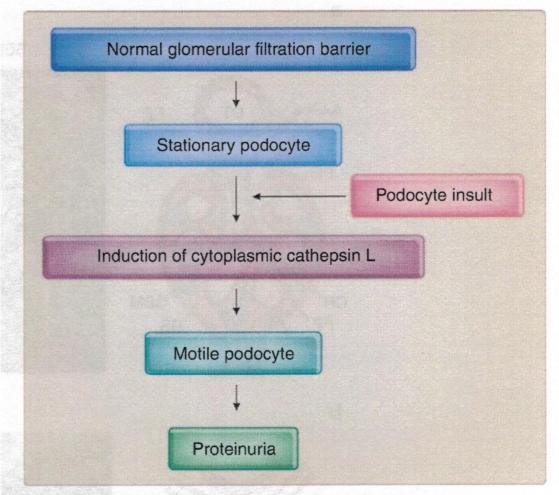


Figure 3 | Induction of cathepsin L in podocytes precedes FP effacement and proteinuria. Upon an insult, stationary podocytes upregulate cytoplasm cytosolic cathepsin L expression and activity and develop motile podocyte foot processes (FPs). This migratory response leads to FP effacement, slit diaphragm remodeling, and proteinuria.<sup>58</sup>

## Other causes of hypoalbuminimia

- Chronic severe diseases of liver
- Enteropathy with loss of proteins
- Malnutrition



### Děkuji vám za pozornost





#