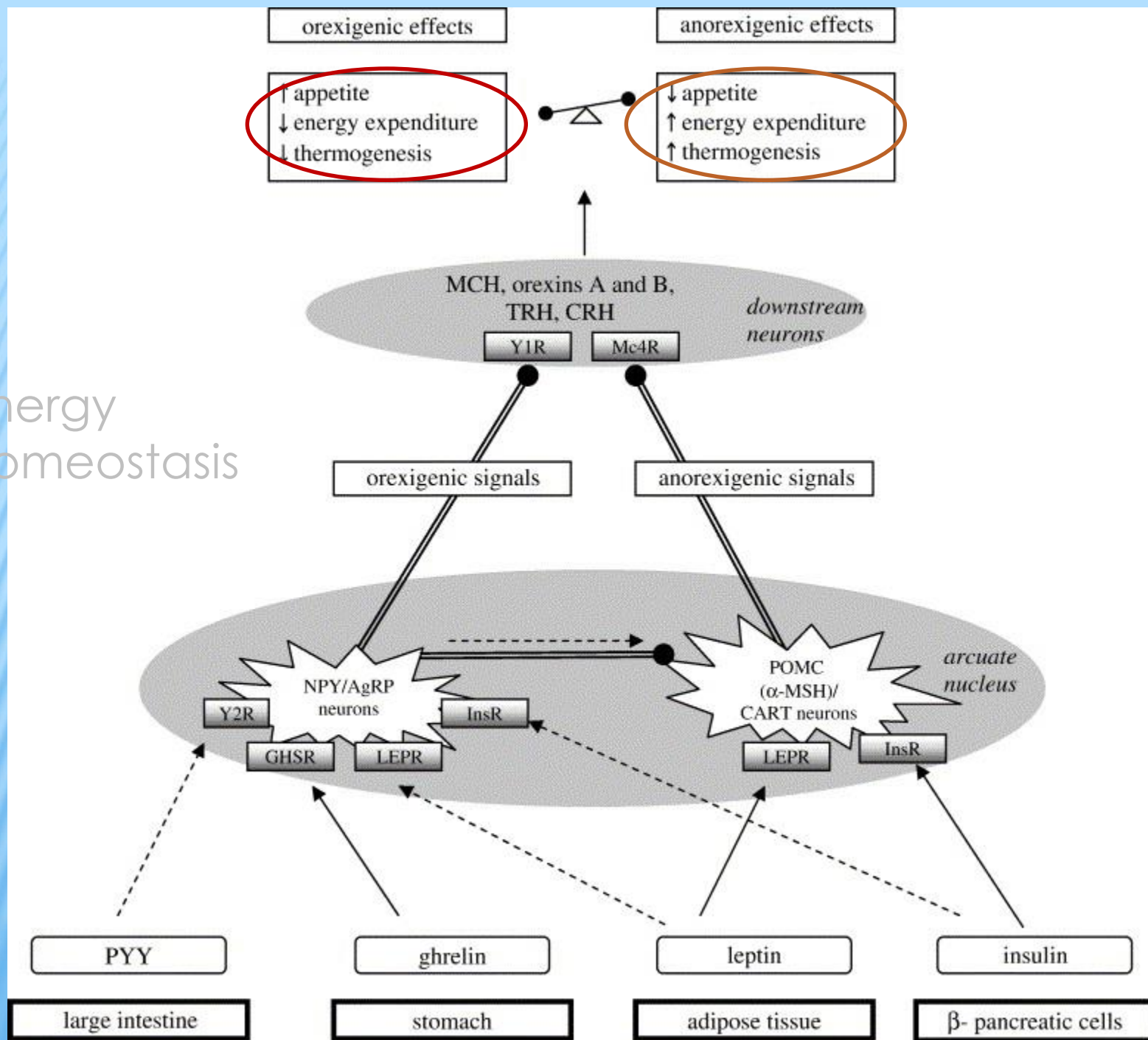


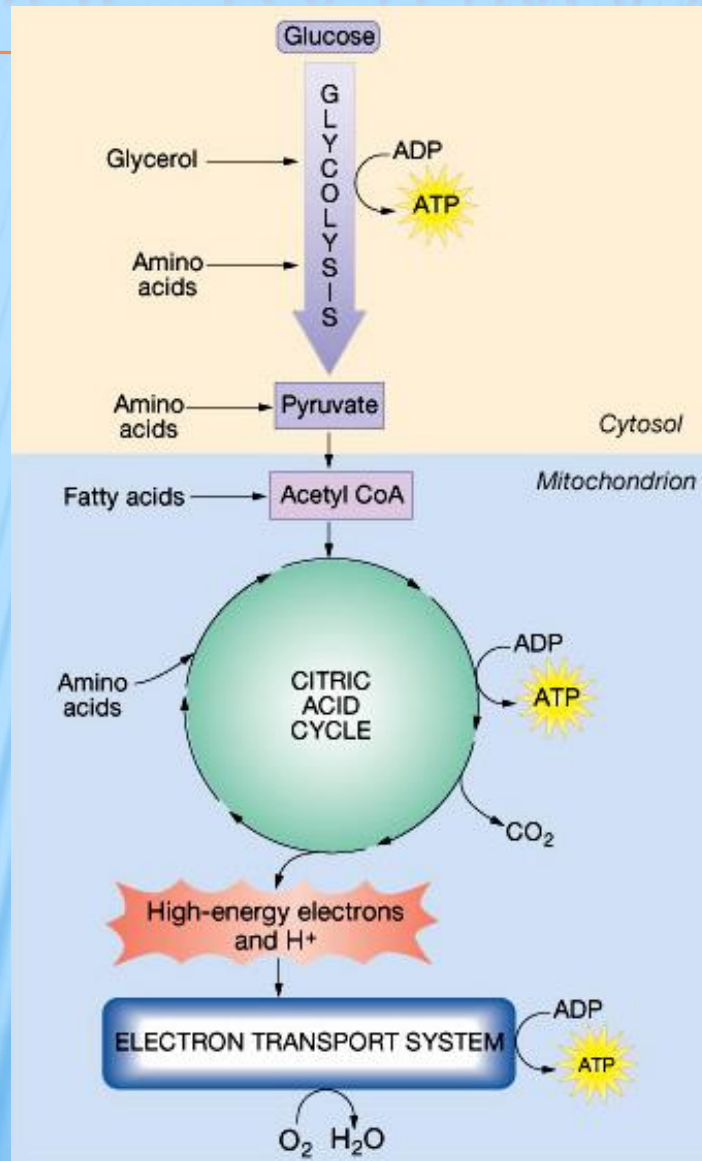
Pathophysiology of metabolism of proteins

27. 3. 2018

Energy homeostasis



METABOLIC PATHWAYS



Intermedial metabolism

METABOLISM STAGES

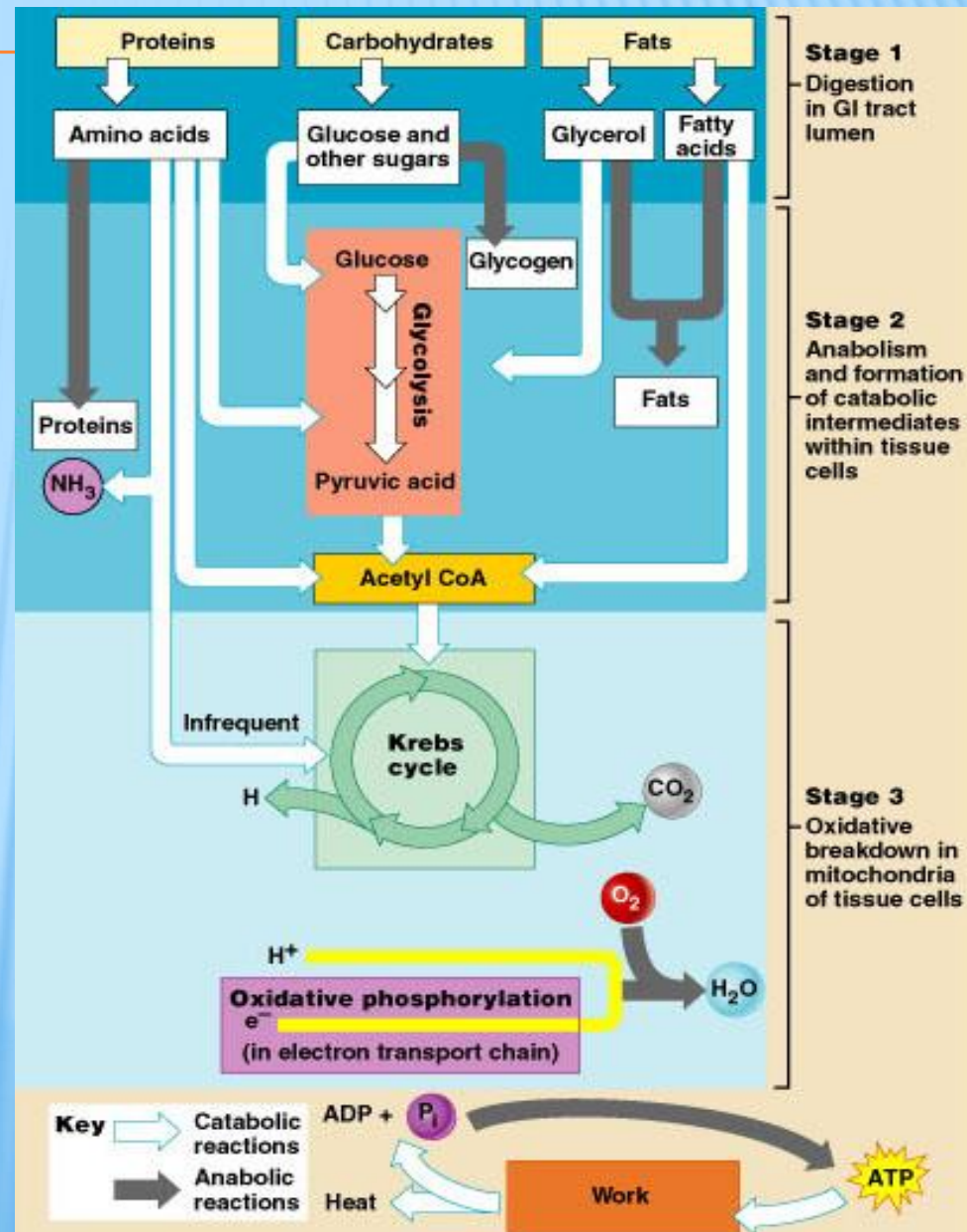
• Nutrients containing energy are processed in three stages:

1. **Digestion** – food processing; nutrients are transported to the tissue

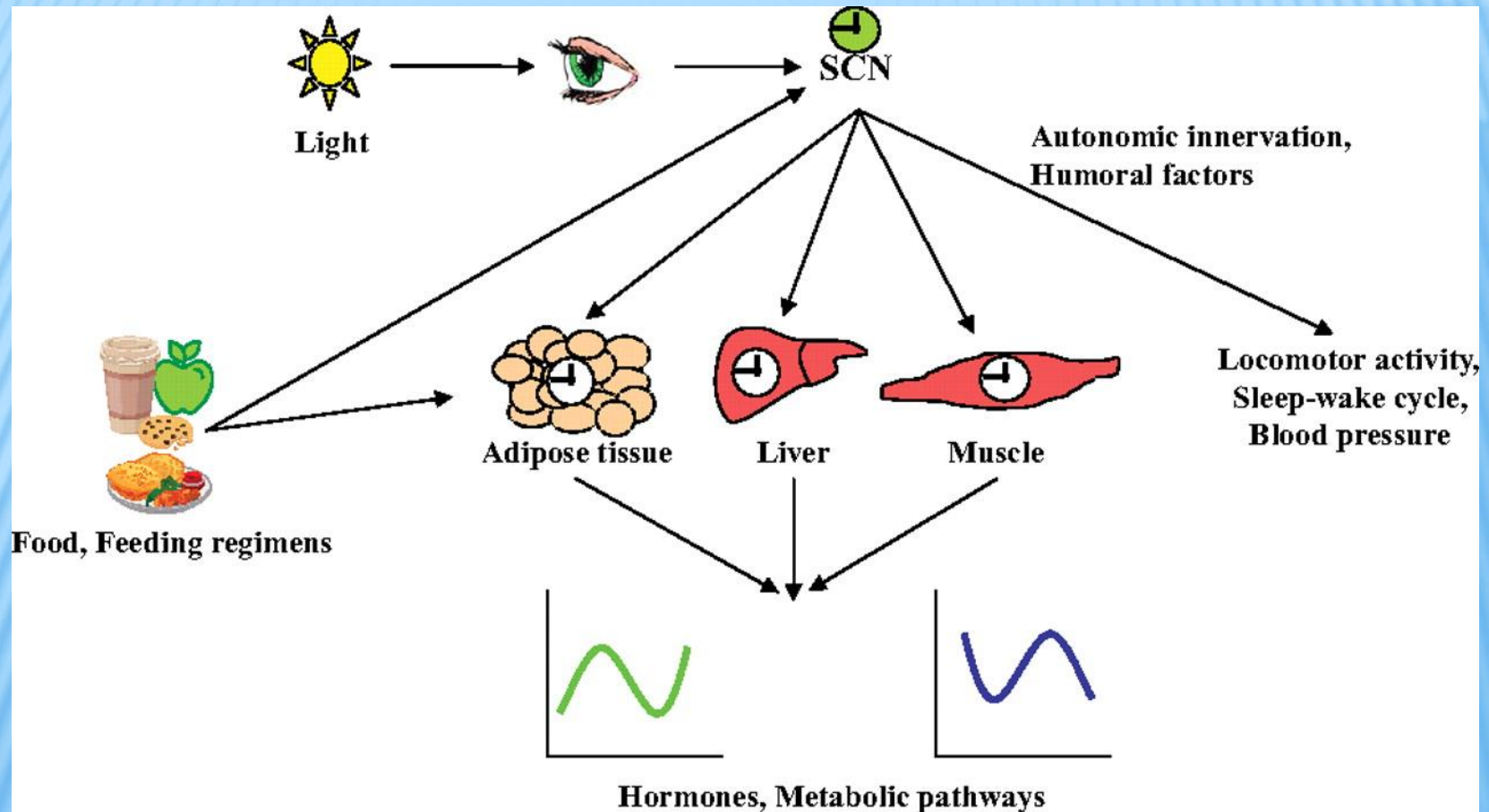
2. **Anabolism and catabolic intermedial products forming**, where nutrients are:

- bound to lipids, proteins and glycogen, or:
- cleaved in metabolic pathways to pyruvate and acetyl CoA.

3. **Oxidative phosphorylation** – nutrients are catabolized to CO₂, water and ATP

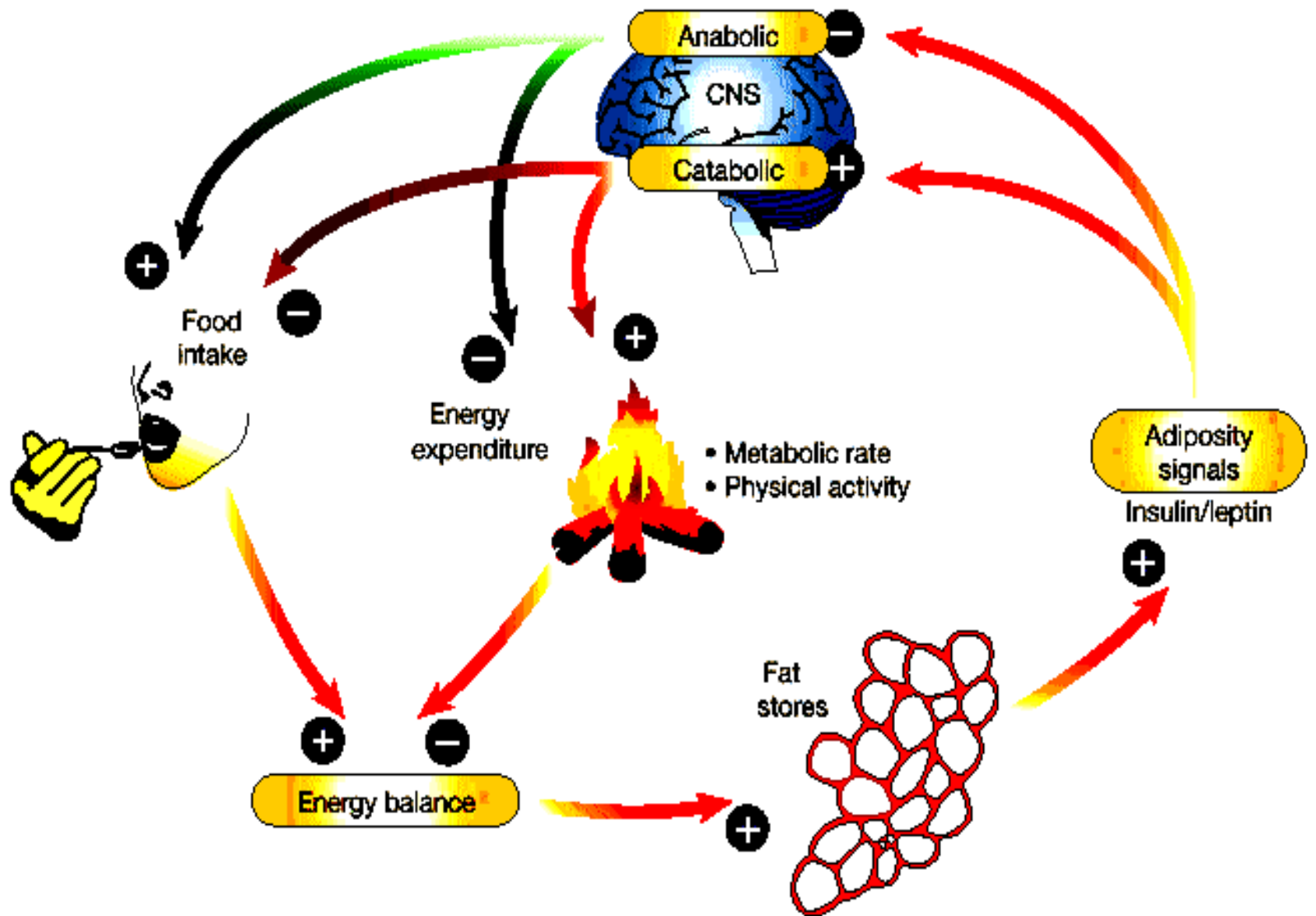


Resetting signals of the central and peripheral clocks.

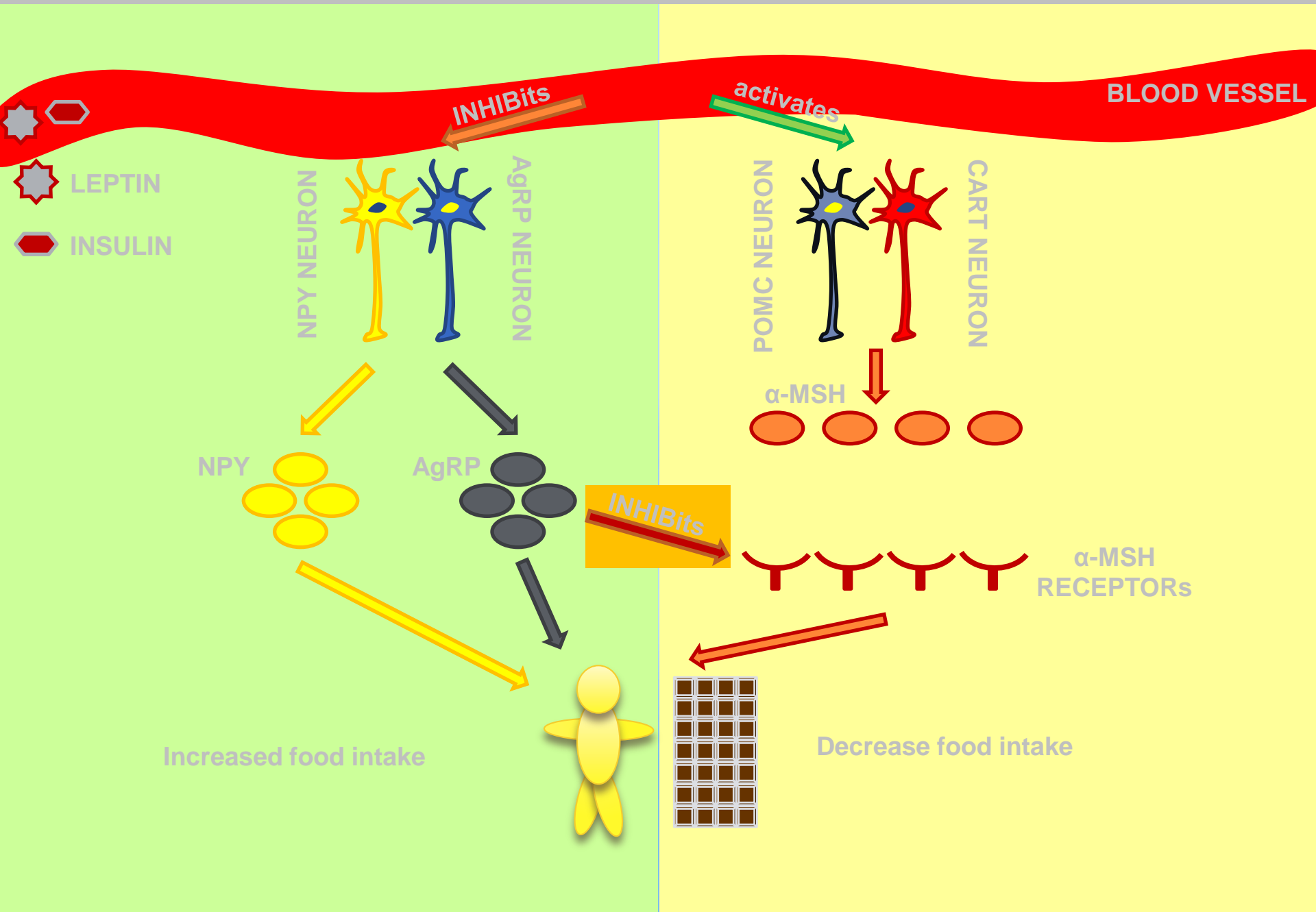


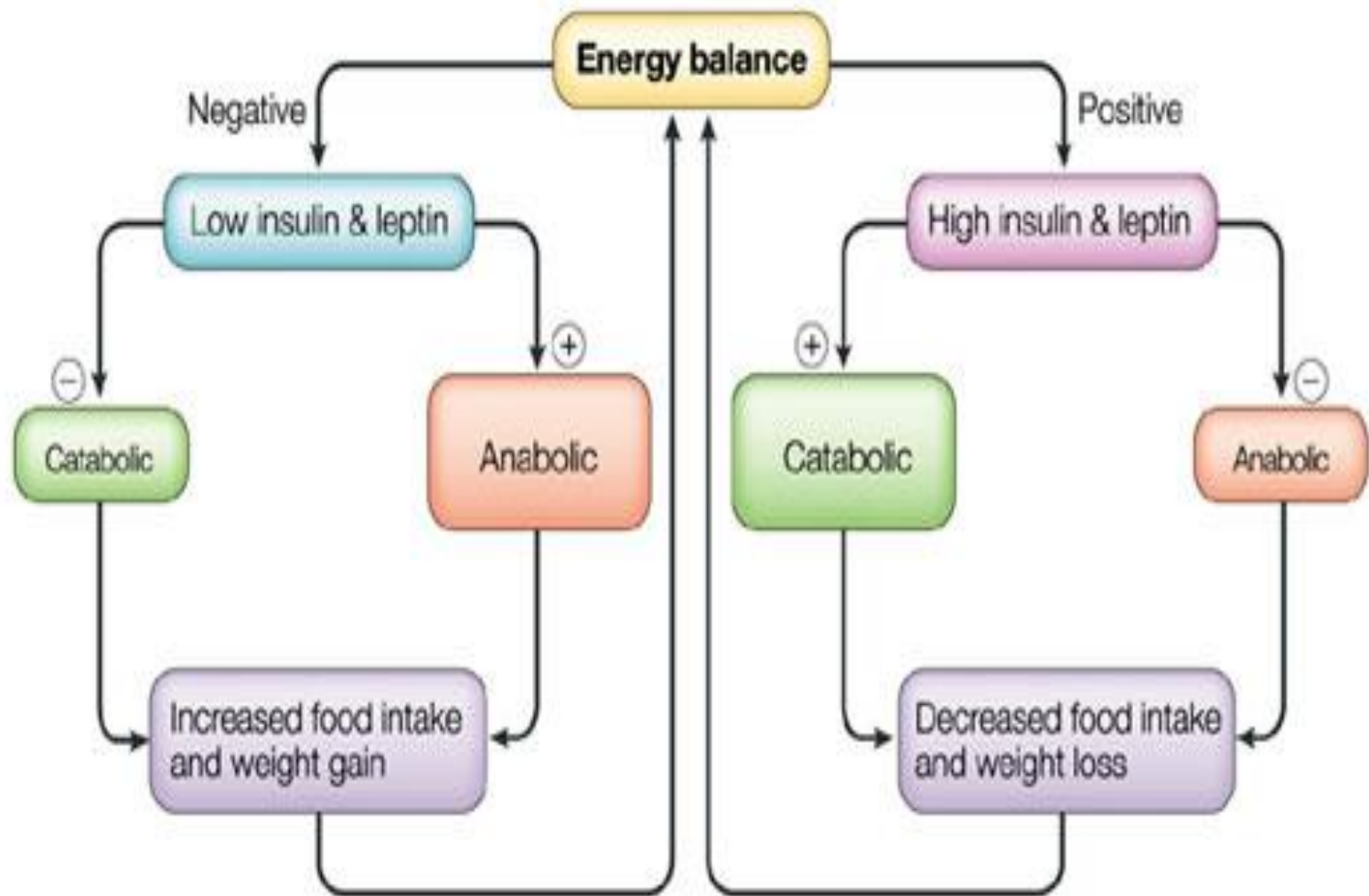
Froy O Endocrine Reviews 2010;31:1-24

ENDOCRINE
REVIEWS

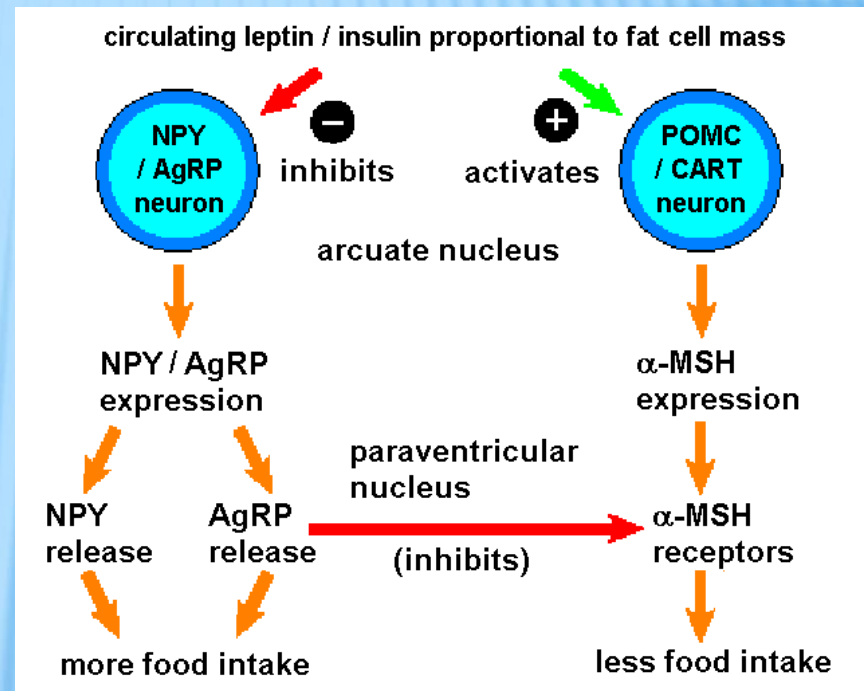
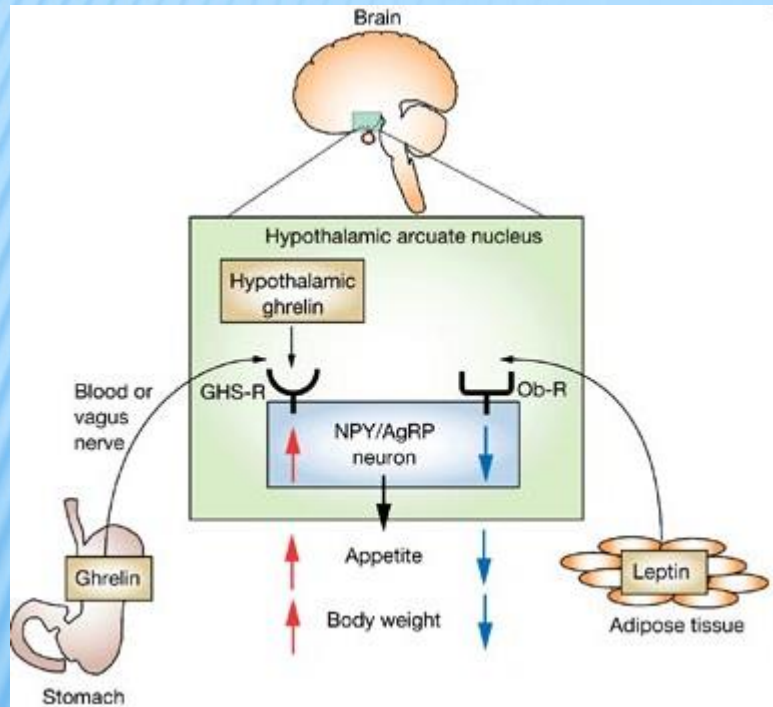


OREXIGENIC-ANOREXIGENIC PATHWAYS





Regulation of food appetite



Drug Insight: the functions of ghrelin and its potential as a multitherapeutic hormone

Masayasu Kojima and Kanji Kangawa

Nature Clinical Practice Endocrinology & Metabolism (2006) 2, 80-88

INTERNAL CHANGES OF NUTRITION MOLECULES

- × **Glycogenesis**
- × **Lipogenesis**
- × **Glycogenolysis**
- × **Gluconeogenesis**

Nutritional Epigenetics

Epigenetic effects of nutrition

Methyl donors

Vitamin B12
Folate
Choline
Betaine
Methionine
Serine
Glycine

Fatty acids

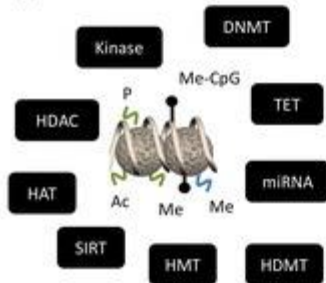
Butyrate
Arachidonic acid
Docosahexaenoic acid
Eicosapentaenoic acid

Vitamins

Retinol
Tocopherols
Vitamin C

Phytochemicals

Genistein
Soy isoflavones
Curcumin
Resveratrol
Sulphoraphane
Polyphenols



Epigenetic control gene expression Epimutations - EpiSNPs

Disease

Specific

Genes:

involved in
Metabolic
Syndrome &
Inflammaging

ADME Genes:

Phase I enzymes
Phase II
Transporters
Metabolisation
DNA repair

For example

PITX2, BRCA1, GPX3, MGMT, PLK2, TFAP2E, OSCP1, SFRP5, RASSF1A, MPO, CFTR, ...
CYP members, GSTM family
GSTP / GSTA variants
UGT / SLC22 variants
SULT2 / SULF variants
ABCA / ABCG variants
ABCB / GPX variants
ALDH variants, etc.

For example

LEP, NPY, POMC, MC4R, IRS1, INS, ADIPOQ, UCP1, TNF, FTO, GLUT4, IGF2, CEBP1, FASN, MHTFR, HIF1A, SOD2, SOD3, IFNG, PPARA, NR3C1

Disease risk
Diagnosis
Prognosis

Metabolisation
Adverse effects
Strong/weak
response

Personalized Epigenetic Biomarkers
Cancer, CVD, CNS, Inflammaging

Personalized Nutrition

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?

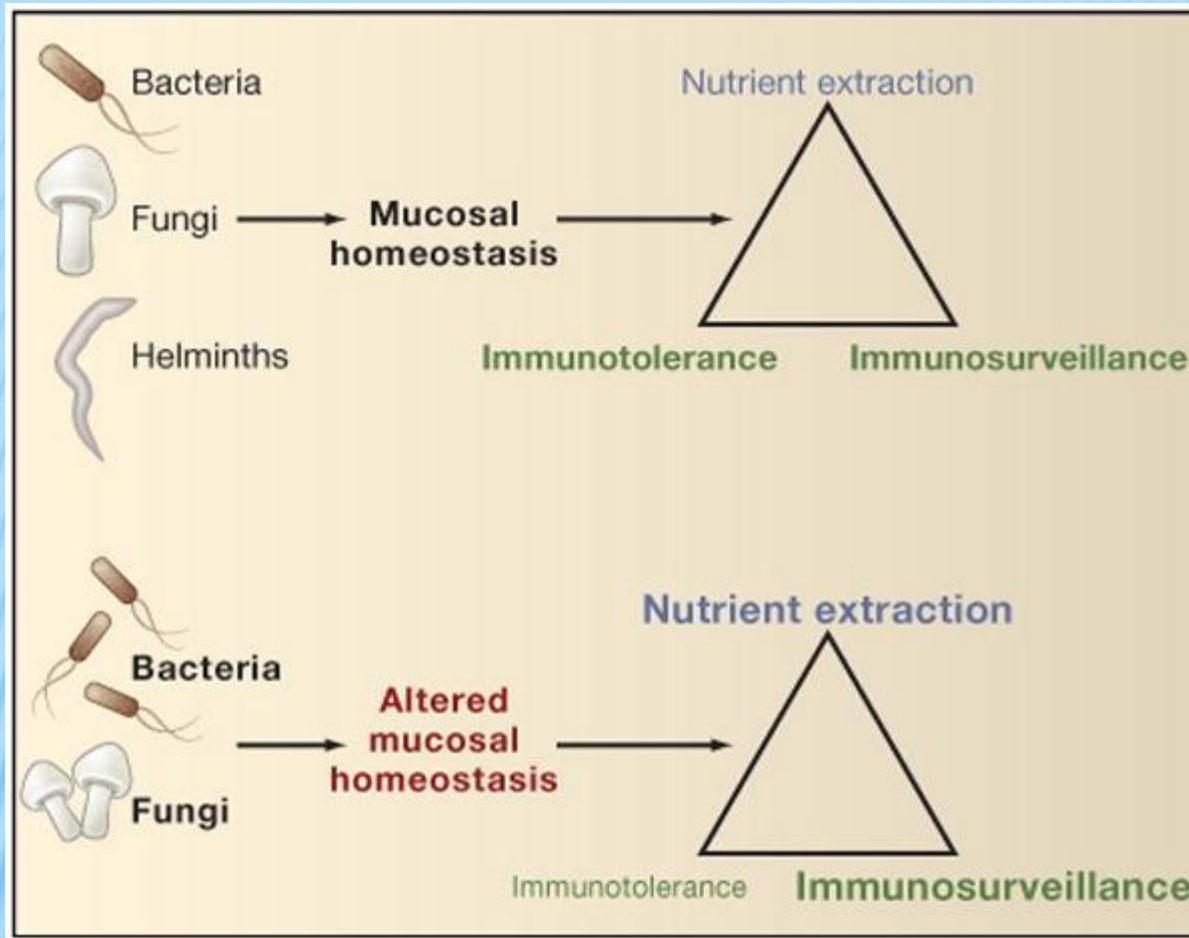
[Vel Szic KS¹,](#)

[Declerck K¹,](#)

[Vidaković M²,](#)

[Vanden Berghe W¹.](#)

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.

METABOLISM

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

- Anabolism
- Catabolism

METABOLISM OF SUBSTANCE:

- ✘ -anabolism = more complicated products are synthesized 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**)

Nutrient	Compound	Transcription factor
<i>Macronutrients</i>		
Fats	Fatty acids Cholesterol	PPARs, SREBPs, LXR, HNF4, ChREBP SREBPs, LXRs, FXR
Carbohydrates	Glucose	USFs, SREBPs, ChREBP
Proteins	Amino acids	C/EBPs
<i>Micronutrients</i>		
Vitamins	Vitamin A Vitamin D Vitamin E	RAR, RXR VDR PXR
Minerals	Calcium Iron Zinc	Calcineurin/NF-ATs IRP1, IRP2 MTF1
<i>Other food components</i>		
	Flavonoids Xenobiotics	ER, NFκB, AP1 CAR, PXR



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

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

PROTEIN METABOLISM

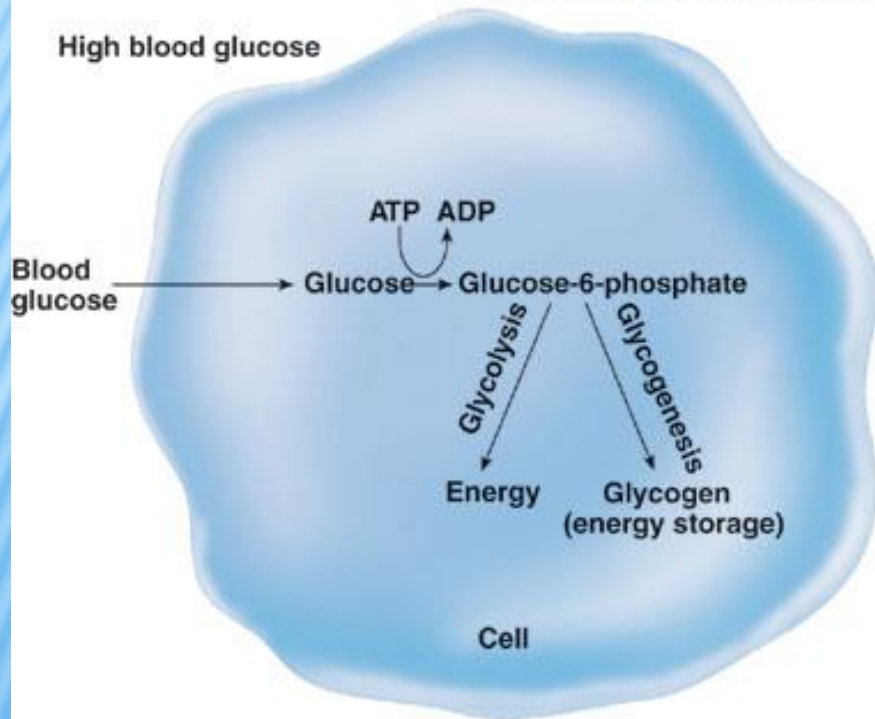
- ✘ Non essential amino acids can be formed by **transamination** (= transfer of amino acid group to keto acid).
- ✘ When using for energy formation, they undergo **oxidative deamination**. NH_3 and keto acids are side products of the reaction. NH_3 is changed to urea and excreted by urine.
- ✘ Amino acids are not stored in the body.

PROTEINS

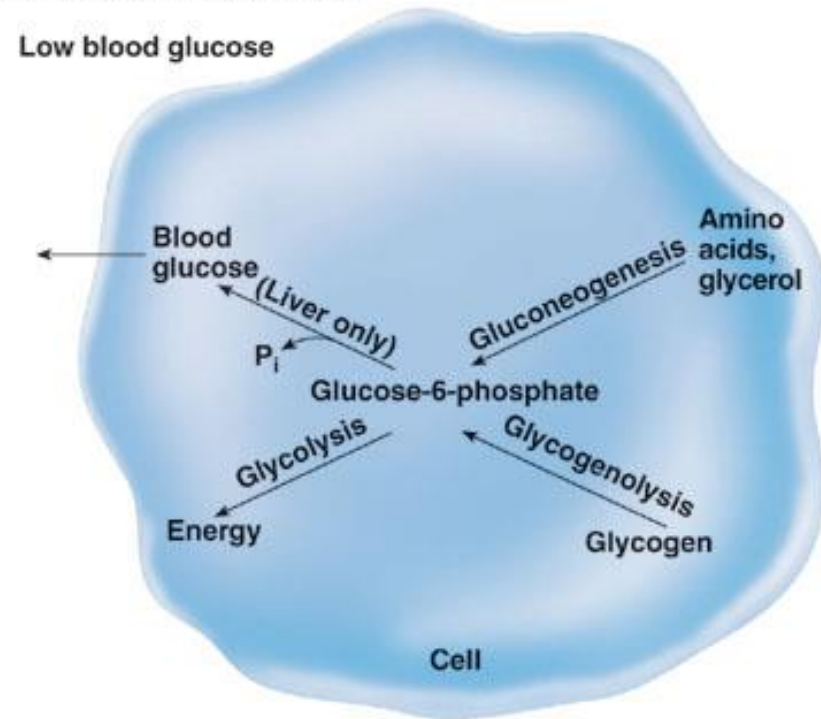
- ✘ Main building material of cells and tissues, enzymes, some hormones and compounds of pigments
- ✘ Food sources: animal (contain all AAs) and plant (not all contain essential AAs) proteins ⇒ digestion in stomach – **pepsin** (from pepsinogen, activated by HCL) and small intestine – **trypsin, aminopeptidases** (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)

INTERNAL CHANGES OF NUTRITION MOLECULES

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(a)



(b)

METABOLIC REACTION OF LIPIDS AND PROTEINS

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

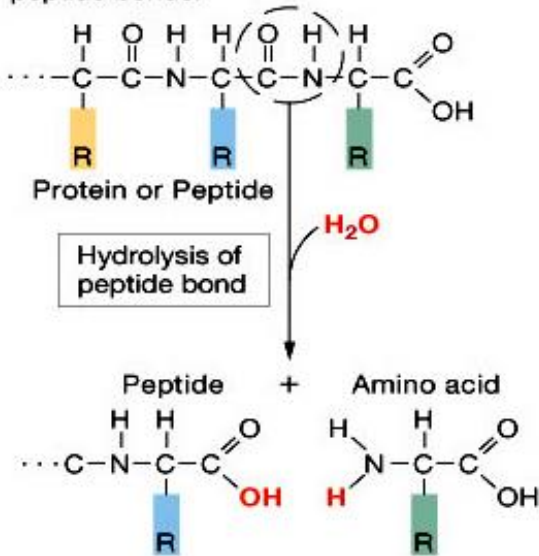
Proteins

Transamination	Transfer of an amine group from an amino acid to α -ketoglutaric acid, thereby transforming α -ketoglutaric acid to glutamic acid
Oxidative deamination	Removal of an amine group from glutamic acid as ammonia and regenerating α -ketoglutaric acid (NH_3 is converted to urea by the liver)

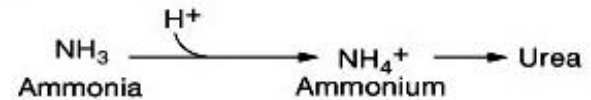
CATABOLISM OF PROTEINS

(a) Protein catabolism

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

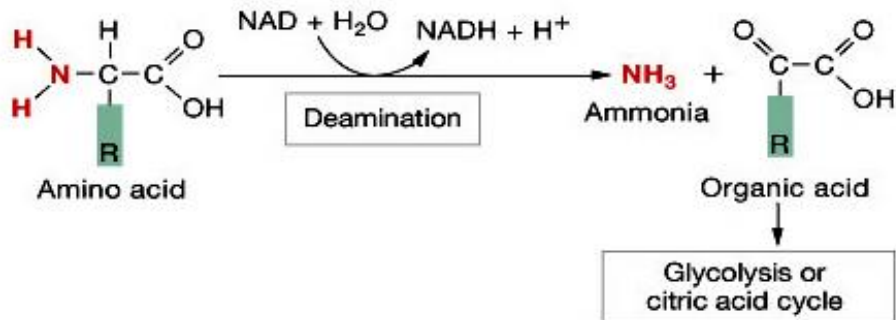


(c) Ammonia is toxic and must be converted to urea.



(b) Deamination

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



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 not 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 and 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 human evolution and allowed for humans to continue to find food effectively even in the face of prolonged starvation.
- ✘ Initially, the level of insulin in circulation drops and the levels of glucagon, epinephrine and norepinephrine rise. At this time, there is an up-regulation of glycogenolysis, gluconeogenesis, lipolysis, and ketogenesis.
- ✘ 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 Cori cycle, which involves shuttling lipid-derived energy in glucose to peripheral glycolytic tissues, which in turn send the lactate back to the liver 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.

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.

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.

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 β 2-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			
<i>WT1</i>	11P13	Wilms tumor 1	IDMS, DDS, Frasier syndrome, WAGR syndrome, ISRNS
<i>LMX1B</i>	17q11	Lim homeobox transcription factor 1- β	Nail-patella syndrome
<i>INF2</i>	14q32.33	Inverted formin-2	FSGS
<i>CD2AP</i>	6p12	CD2-associated protein	FSGS (adult)
AR			
<i>NPHS1</i>	19q13.1	Nephrin	CNF
<i>NPHS2</i>	1q25-31	Podocin	Idopathic CNS, SRNS
<i>LAMB2</i>	3p21	Laminin β 2 chain	Pierson's syndrome
<i>PLCE1</i>	10q23	Phospholipase C epsilon 1	SRNS, DMS
<i>PDSS2</i>	6q21	Decaprenyl disphosphate synthase, subunit 2	NS with Leigh syndrome
<i>ITGA3</i>	17q21.33	Integrin α 3	NS with interstitial lung disease
<i>ARHGDI1</i>	17q25.3	Rho GDP dissociation inhibitor 2	Idiopathic CNS
<i>SCARB2</i>	4q21.1	Scavenger 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.

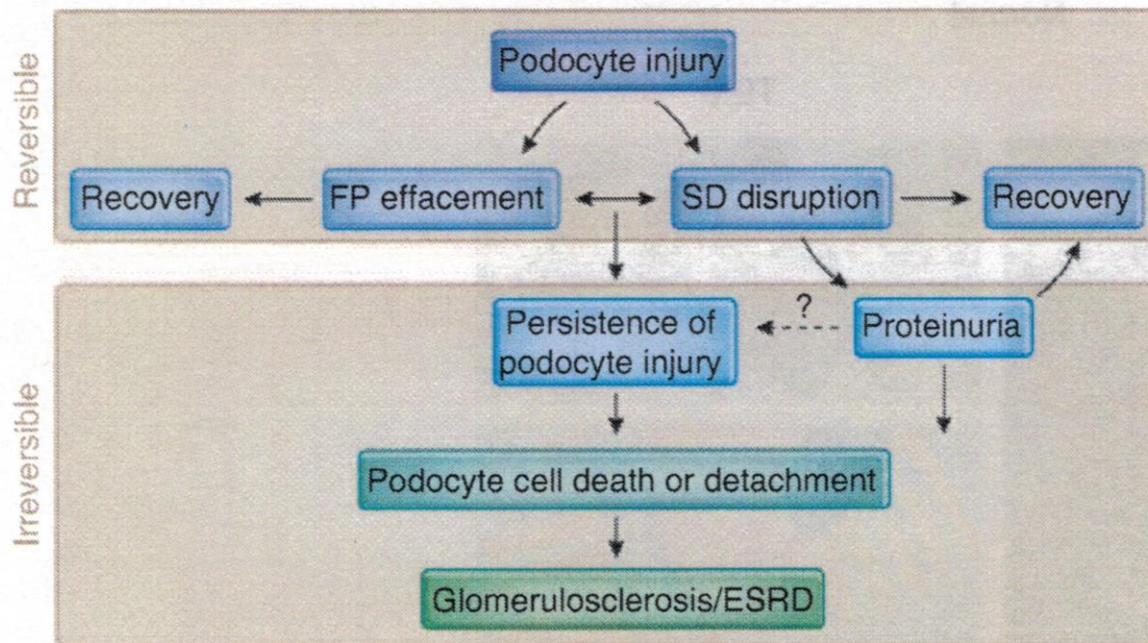


Figure 2 | Consequences of podocyte injury. Podocytes can be injured in many human and experimental glomerular diseases, leading to structural changes, such as foot processes (FP) effacement and slit diaphragm (SD) disruption that are reversible.¹²⁴ Persistence of podocyte injury can cause cell death or detachment of podocytes from the glomerular basement membrane (GBM).⁷² The resulting loss of podocyte will ultimately lead to irreversible glomerulosclerosis and end-stage renal failure (ESRD).⁷³ 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.

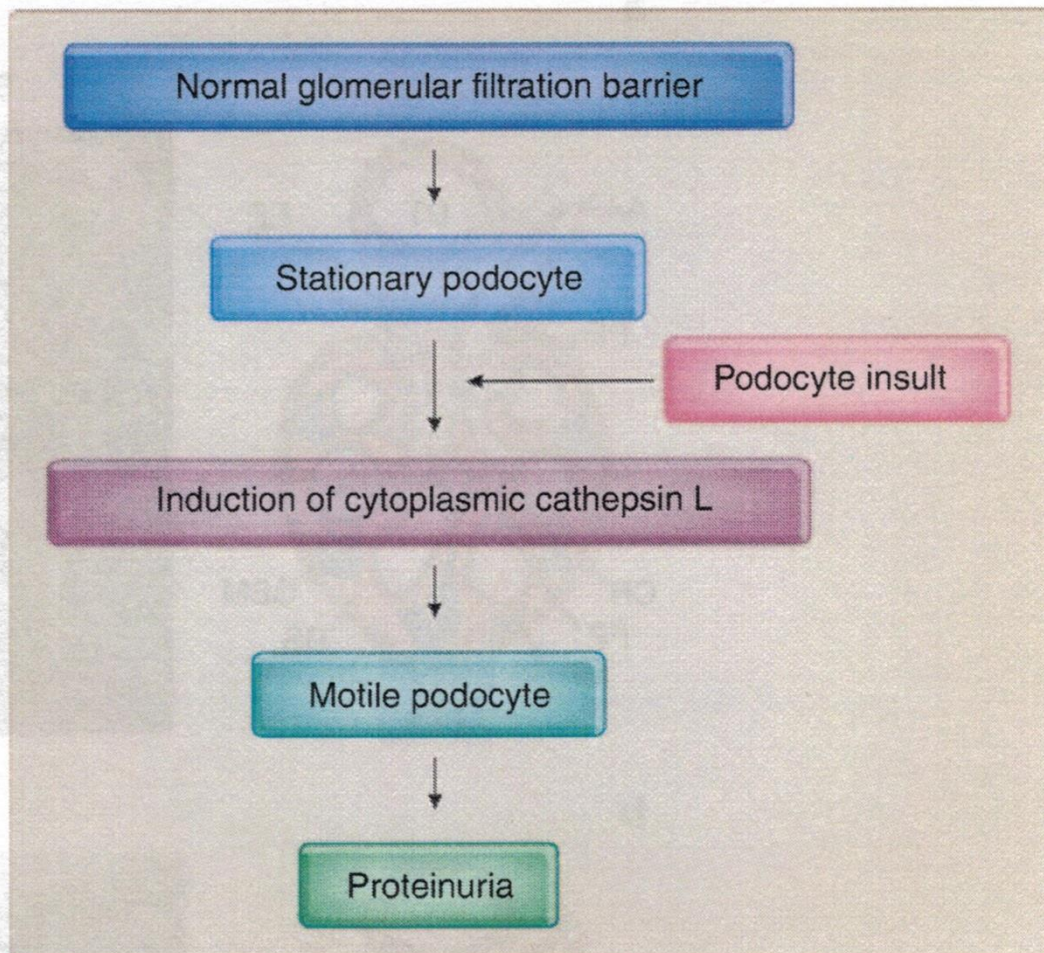


Figure 3 | Induction of cathepsin L in podocytes precedes FP effacement and proteinuria. Upon an insult, stationary podocytes upregulate cytoplasmic cathepsin L expression and activity and develop motile podocyte foot processes (FPs). This migratory response leads to FP effacement, slit diaphragm remodeling,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

DĚKUJI VÁM ZA POZORNOST

