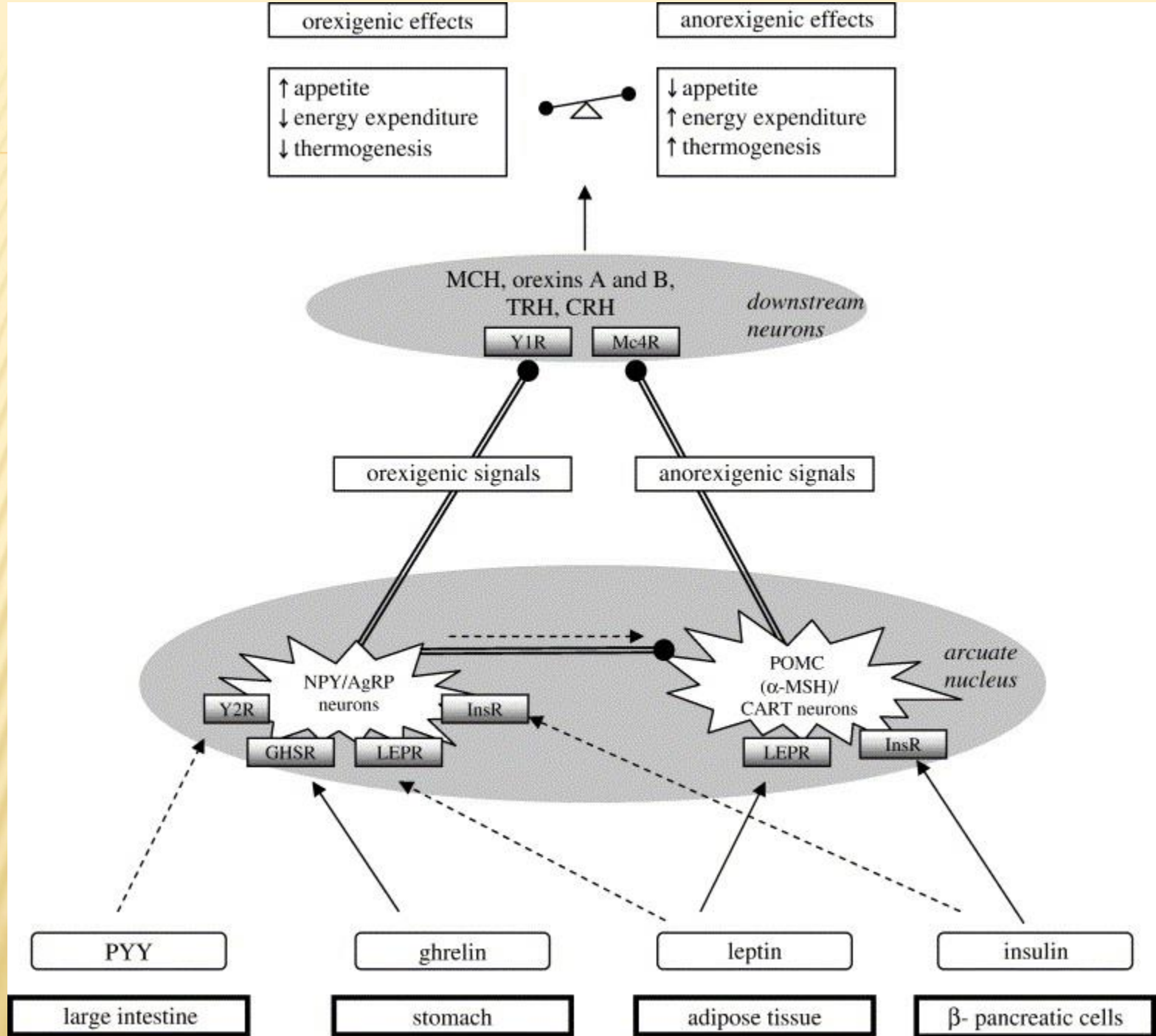
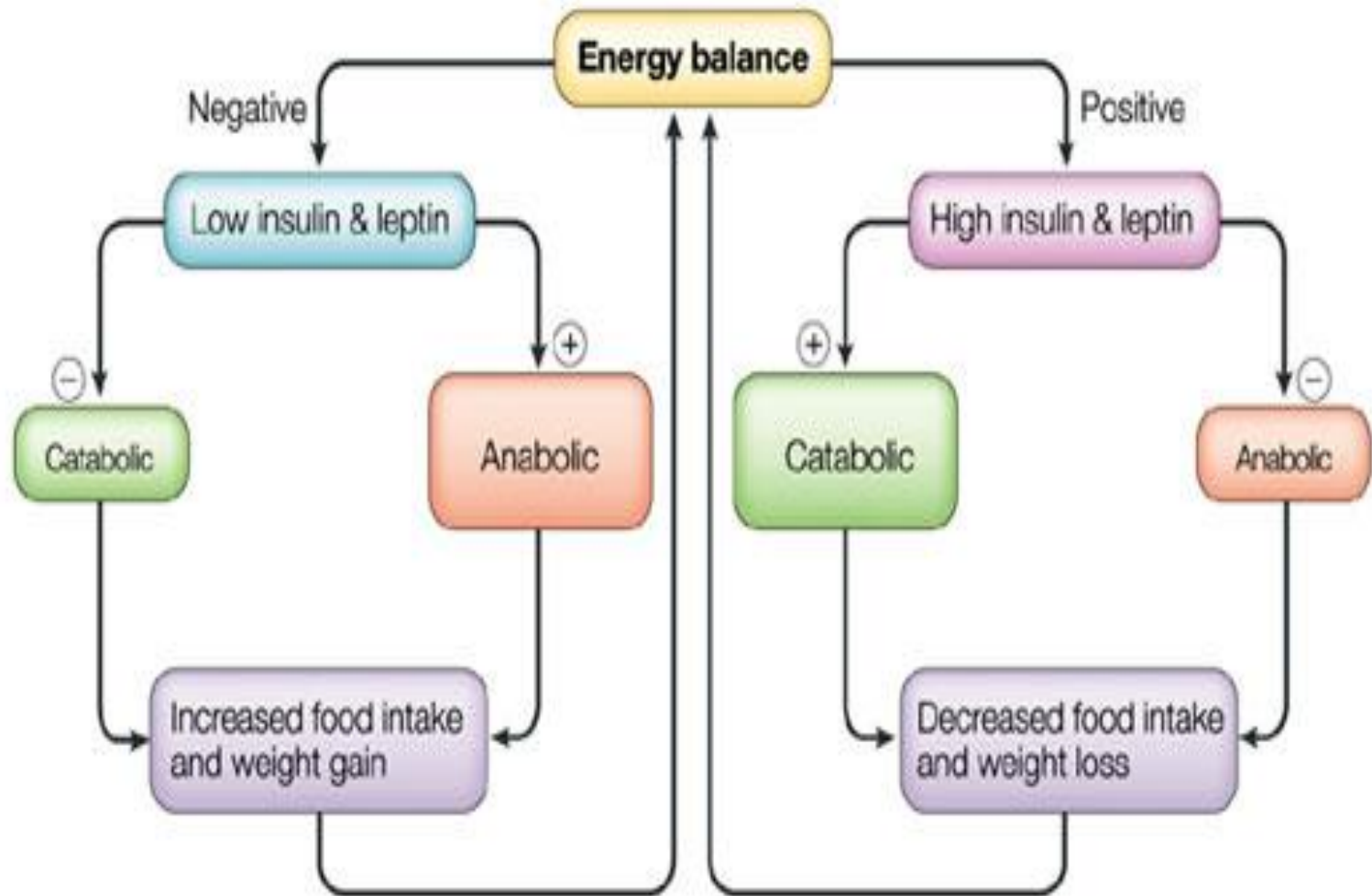


EATING DISORDERS

ZLA 5. 4. 2018

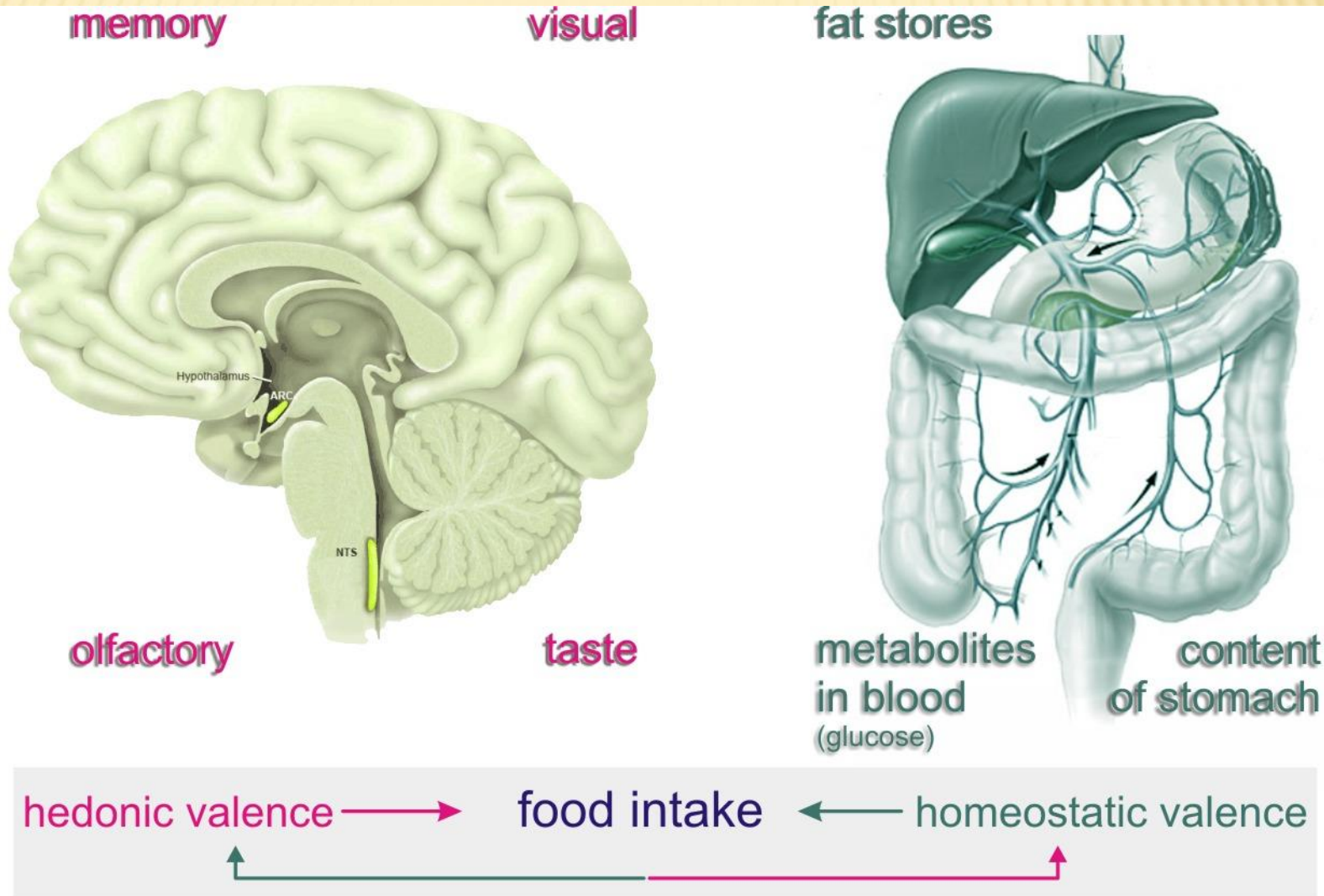


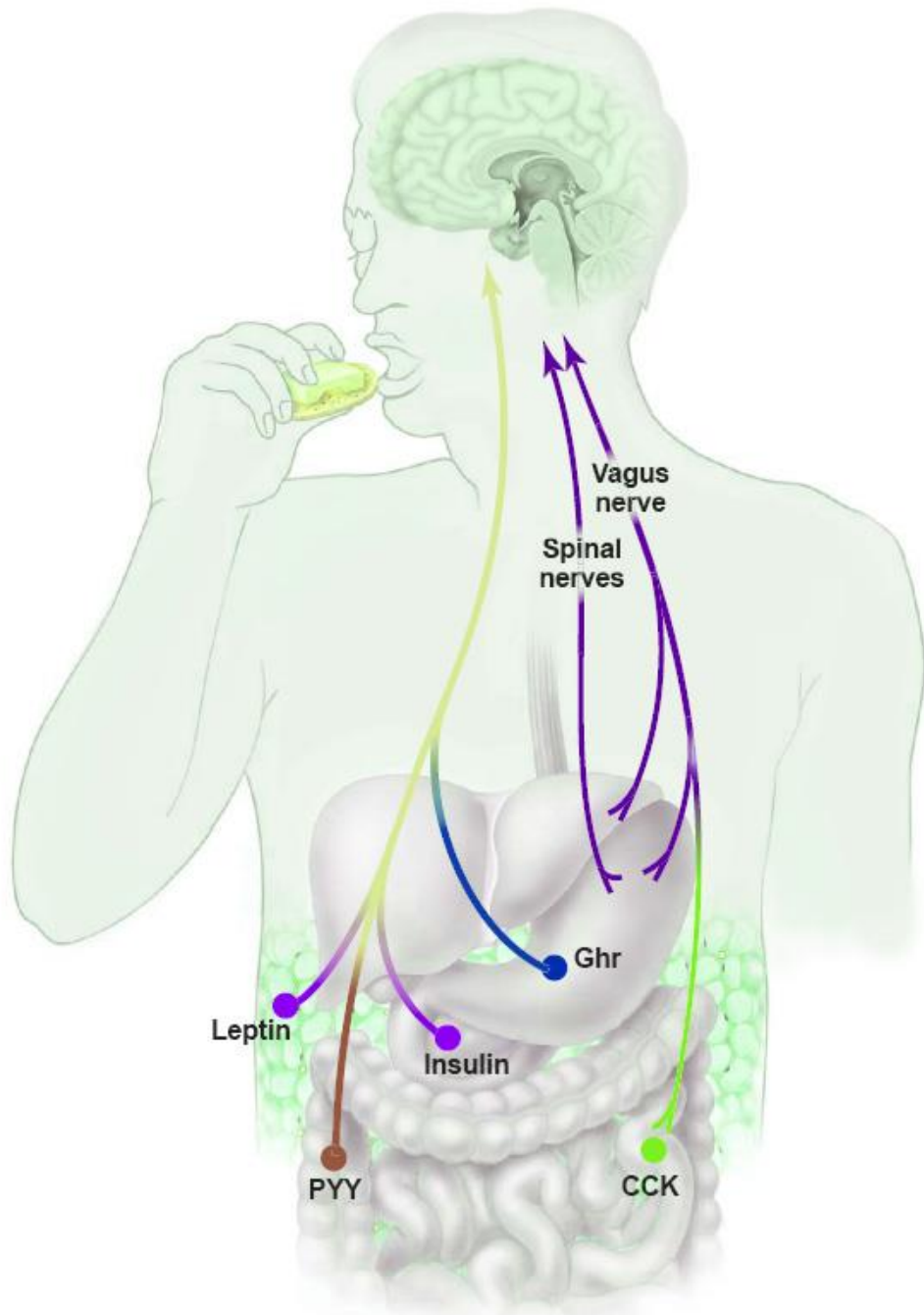


HISTORY

- ✘ Lipostatic hypothesis (Kennedy 1953) – adipose tissue products specific „lipostatic“ factor
- ✘ Glukostatic hypothesis (Mayer and Thomas 1967) – changes in glycemia lead to stimulation /inhibition of food intake (brain and liver)
- ✘ Combination of both hypotheses???

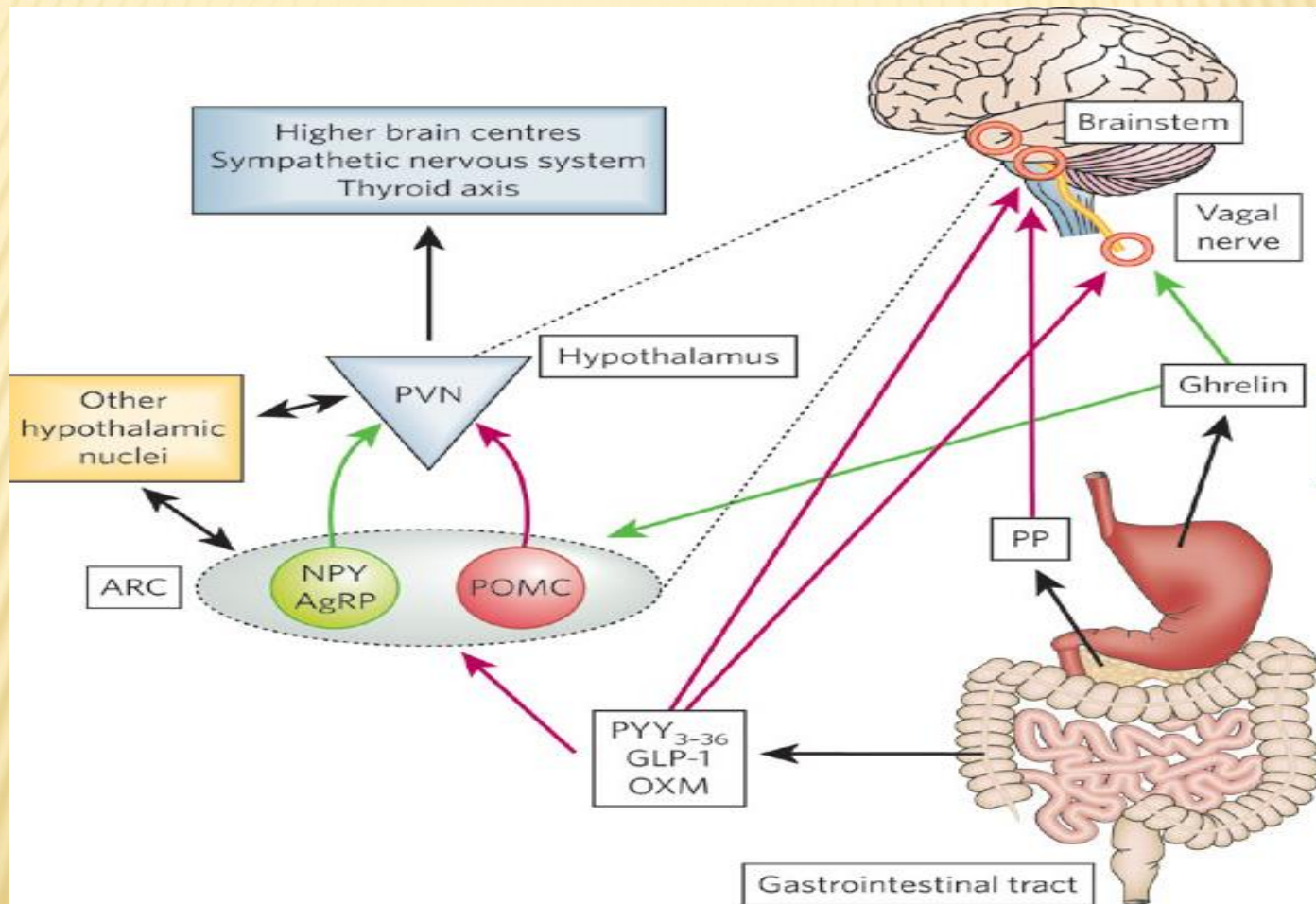
IT IS NECESSARY TO DISTINGUISH COMMON HOMEOSTATIC REGULATIONS OF FOOD INTAKE AND HEDONIC REGULATIONS





SIMPLIFIED
SCHEME OF
HOMEOSTATIC
REGULATIONS OF
FOOD INTAKE.
SUBSTANTIAL SIGNALS
ORIGINATE FROM GIT
(GHRELIN, CCK, PYY),
ADIPOSE TISSUE (LEPTIN,
OTHER ADIPOKINES) AND
PANKREAS (INSULIN,
GLUKAGON)

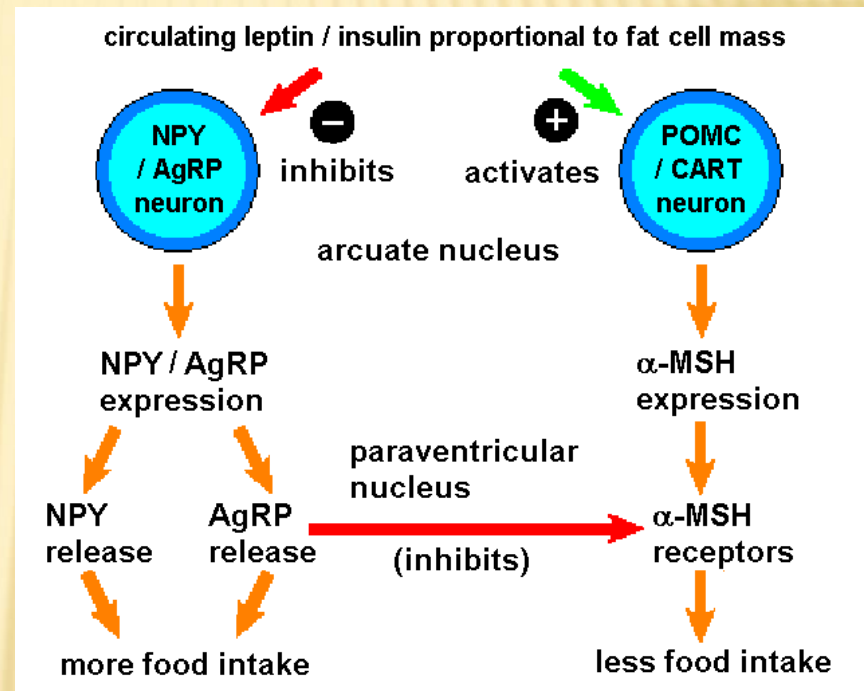
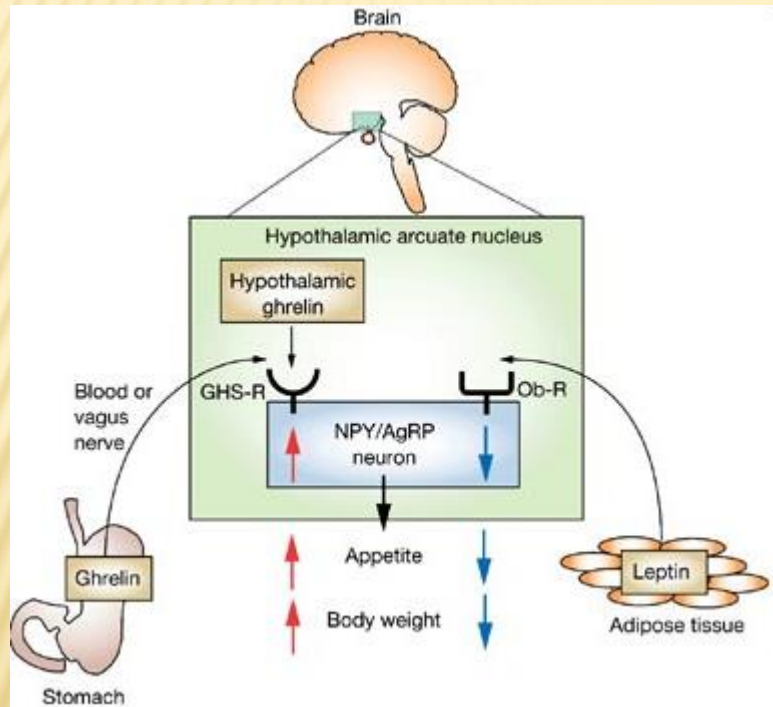
CENTRAL AND PERIPHERAL CIRCUITS IN REGULATION OF FOOD INTAKE



PYY

- ✘ Peptide YY is related to the pancreatic peptide family by having 18 of its 36 amino acids located in the same positions as pancreatic peptide. The two major forms of peptide YY are PYY₁₋₃₆ and PYY₃₋₃₆, which have PP fold structural motifs. However, the most common form of circulating PYY immunoreactivity is PYY₃₋₃₆, which binds to the Y2 receptor (Y2R) of the Y family of receptors. Peptide YY₃₋₃₆ (PYY) is a linear polypeptide consisting of 36 amino acids with structural homology to NPY and pancreatic polypeptide.
- ✘ PYY is found in L cells in the mucosa of gastrointestinal tract, especially in ileum and colon. Also, a small amount of PYY, about 1-10%, is found in the esophagus, stomach, duodenum and jejunum. PYY concentration in the circulation increases postprandially (after food ingestion) and decreases by fasting. In addition, PYY is produced by a discrete population of neurons in the brainstem, specifically localized to the gigantocellular reticular nucleus of the medulla oblongata.
- ✘ PYY exerts its action through NPY receptors; it inhibits gastric motility and increases water and electrolyte absorption in the colon. PYY may also suppress pancreatic secretion. It is secreted by the neuroendocrine cells in the ileum and colon in response to a meal, and has been shown to reduce appetite. PYY works by slowing the gastric emptying; hence, it increases efficiency of digestion and nutrient absorption after a meal.

REGULATION OF FOOD INTAKE

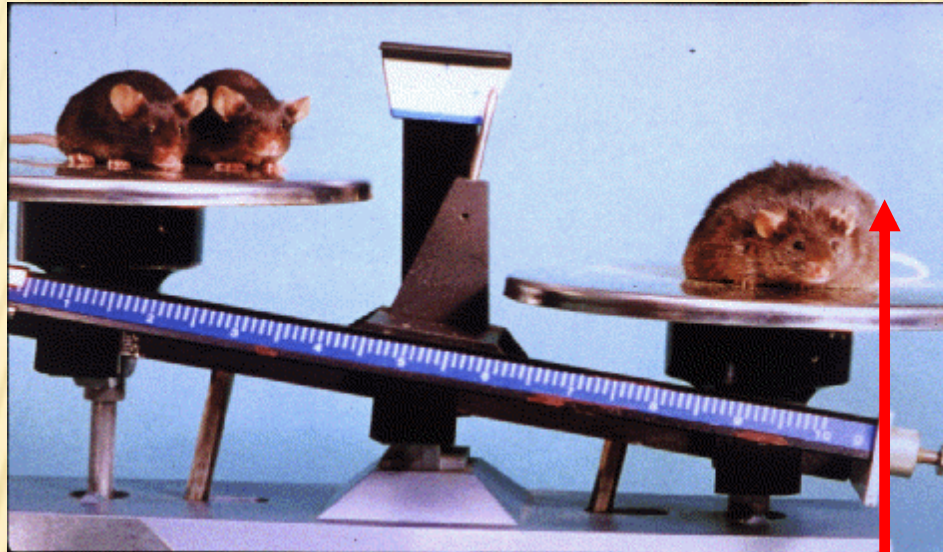


Drug Insight: the functions of ghrelin and its potential as a multitherapeutic hormone

Masayasu Kojima and Kenji Kangawa

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

1994 – ADIPOSE TISSUE IS ABLE TO PRODUCE HORMONS ITSELF – LEPTIN DISCOVERY



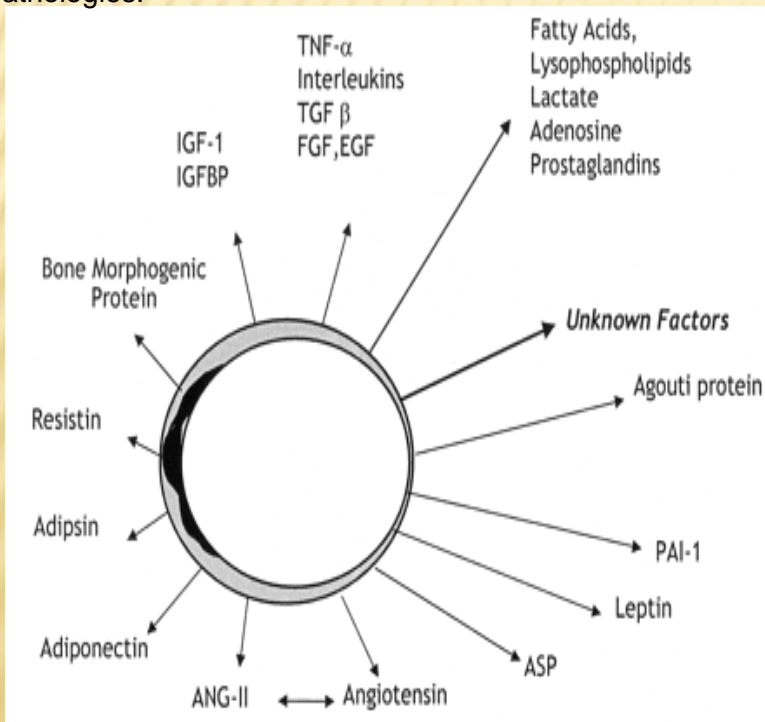
Mutation of *ob* gene coding for leptin is leading to significant obesity in mouse

Leptin therapy led to normalized body weight and/or fertility of these mutated mice

WAT produces adipokines

These factors are produced by adipocytes, but also by macrophages, fibroblasts, endothelial cells and other cells in adipose tissue

To date, a lot of adipose tissue-derived factors has been described. These factors with pleiotropic functions in many processes including regulation of energy metabolism, inflammation, food intake, insulin sensitivity etc. Markedly contribute to metabolic regulations and its pathologies.



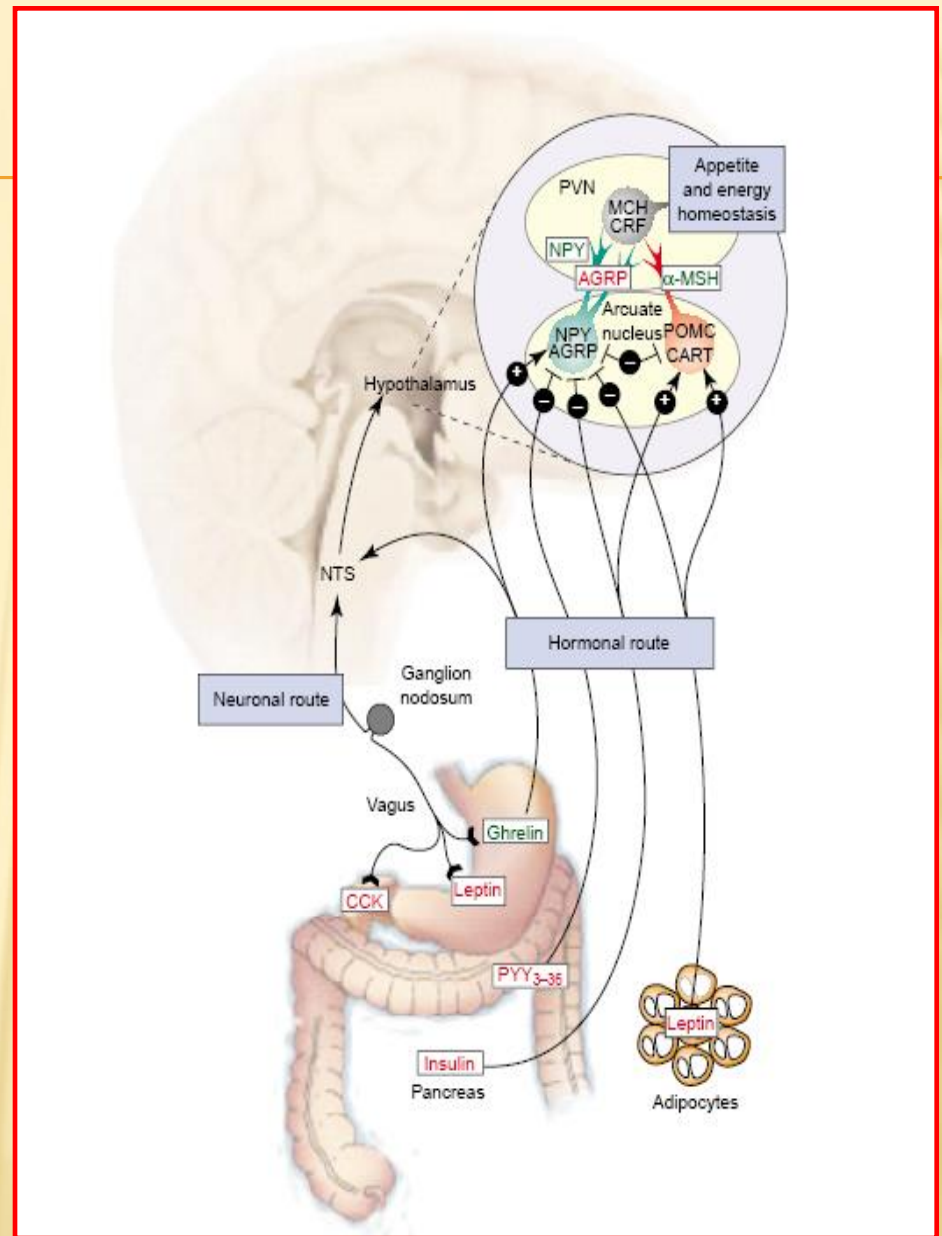
They are usually:

1. Proinflammatory (TNF- α , IL-6, resistin)
2. Anti-inflammatory (adiponektin)

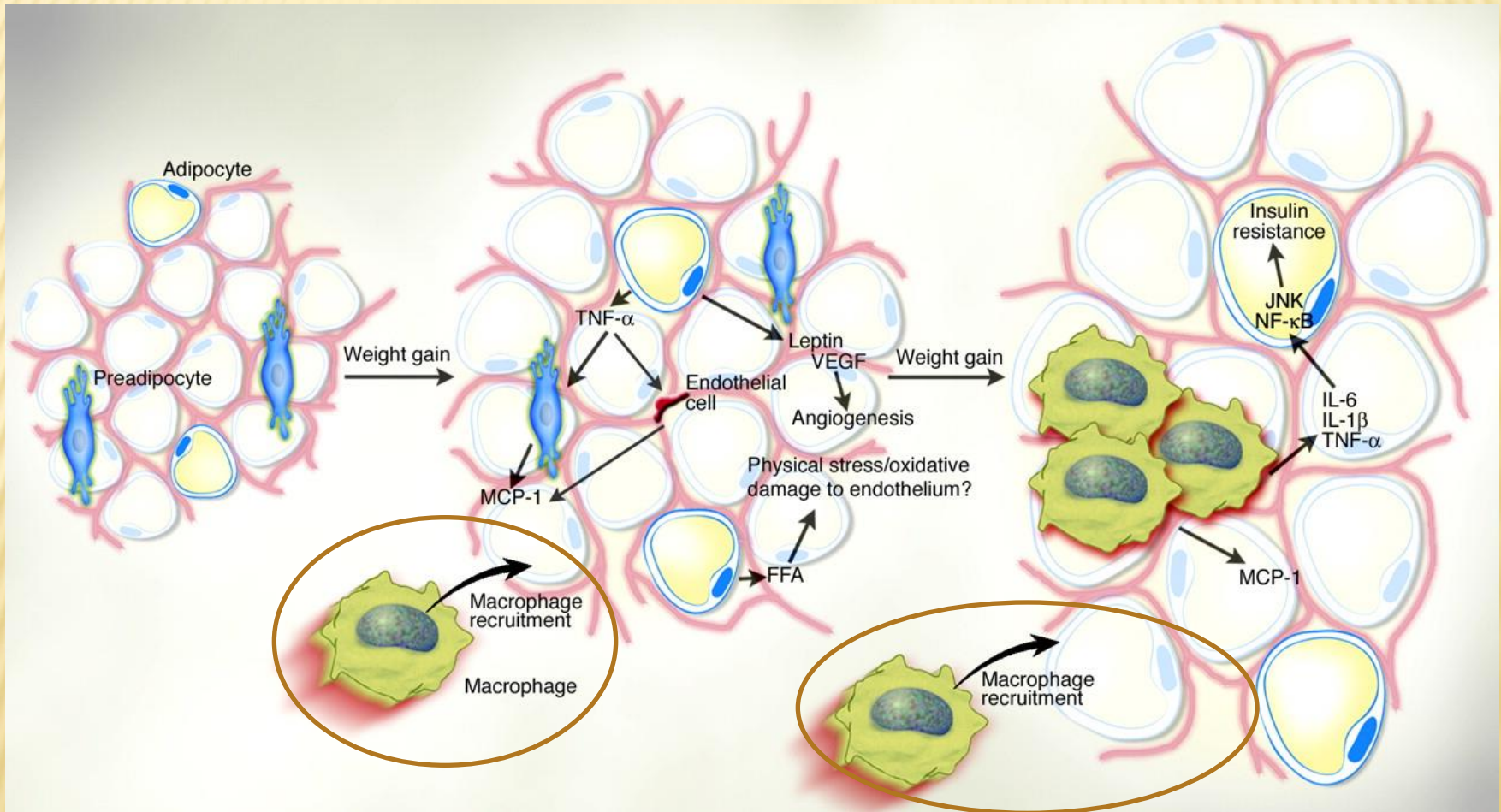
They are very important in metabolic regulations

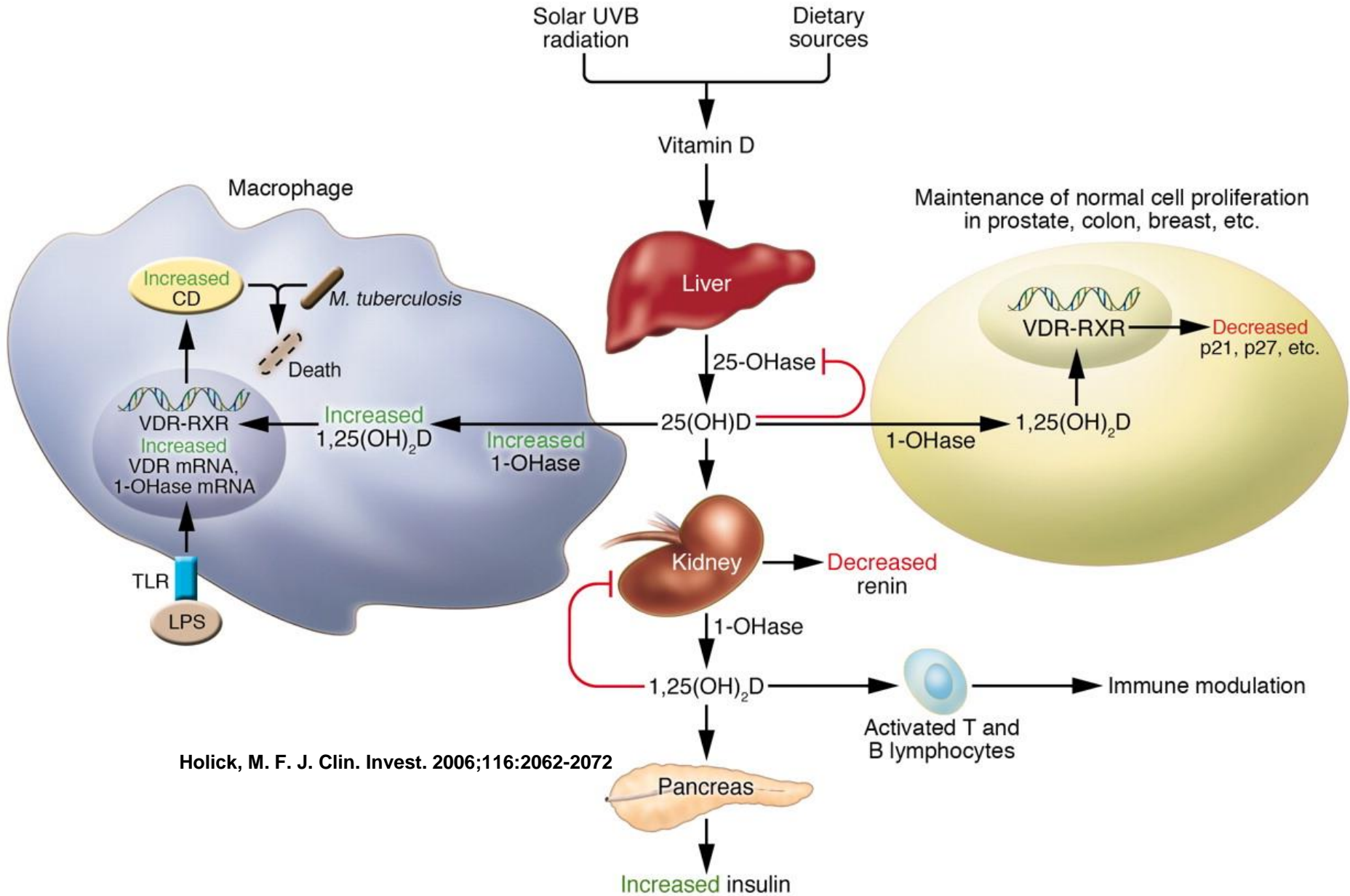
Food Intake and GIT...

In food intake and especially in postprandial suppression of appetite, many substances produced in GIT play substantial roles.

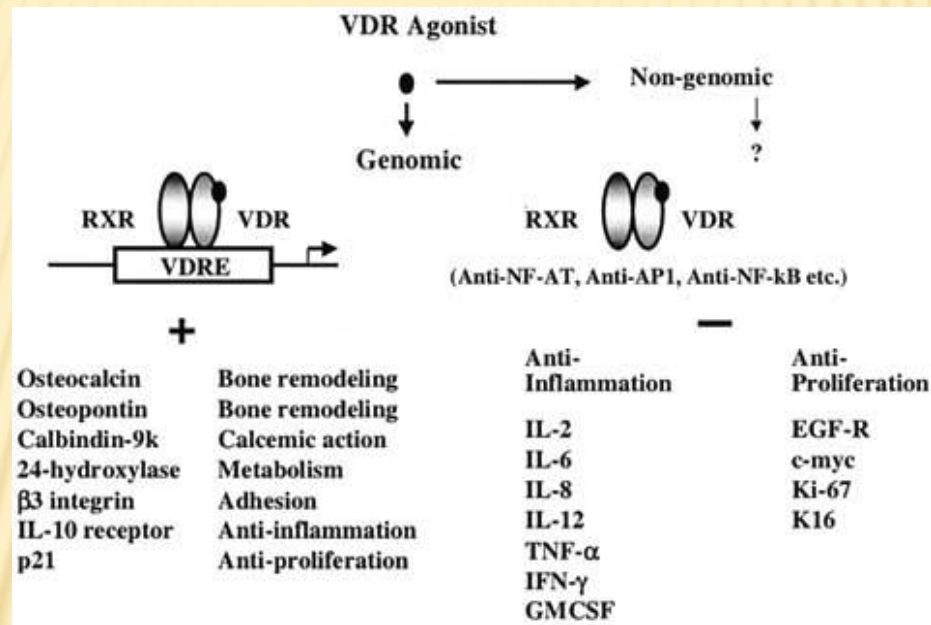


OBESITY IS ASSOCIATED WITH LOCAL INFLAMMATORY RESPONSE IN FAT



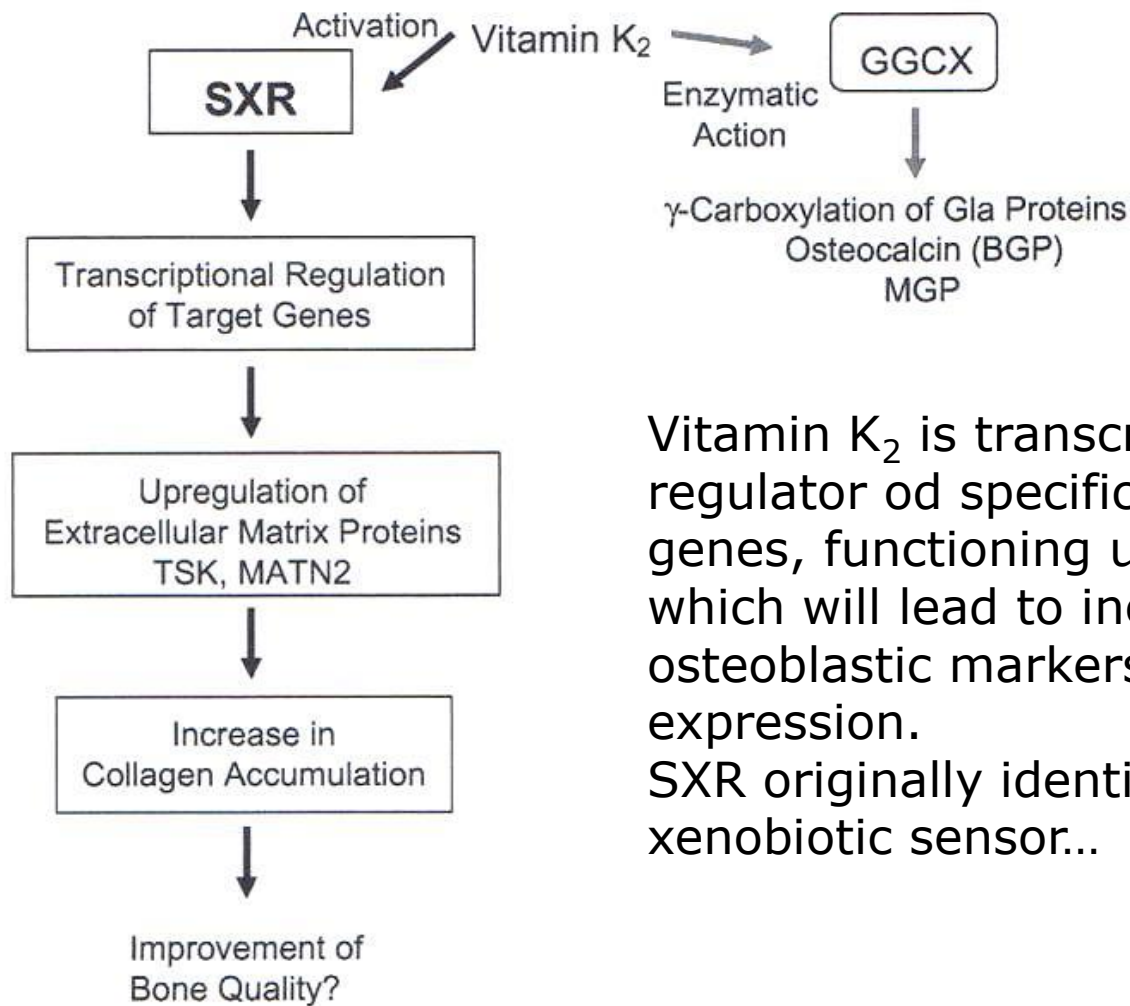


Regulation of gene expression by VDR



Vitamin K and bones

- Vitamin K₂ is substantial cofactor for γ -carboxylase, enzyme which catalyses conversion of specific residuals of glutamic acid to Gla residuals
- Vitamin K₂ is necessary for γ -carboxylation of proteins of bone matrix which contain Gla as **MGP (= matrix Gla protein) a osteocalcin.**
- Uncompleted γ -carboxylation of osteocalcin and MGP during vitamin K decrease lead to osteoporosis and high risk of fractures. Vitamin K₂ stimulates synthesis of osteoblastic markers and bone deposition.
- Vitamin K₂ decreases bone reabsorption by inhibition of osteoclasts formation and by decrease of their resorption activity.
- Vitamin K₂ treatment induces osteoclast apoptosis, but inhibits osteoblasts apoptosis which is leading to increased bone formation.
- Vitamin K₂ supports osteocalcin expression (increases its mRNA) which can be further modulated by 1, 25-(OH)₂ vitamin D₃.



Vitamin K₂ is transcription regulator of specific bone genes, functioning using SXR which will lead to increase of osteoblastic markers expression. SXR originally identifies as xenobiotic sensor...

Fig. 3. SXR- and vitamin K₂-dependent regulatory mechanisms of bone metabolism in osteoblastic cells. SXR promotes collagen accumulation in osteoblastic cells by regulating the transcription of its target genes including those encode extracellular matrix proteins. Vitamin K₂ plays a role in the posttranslational modification of Gla proteins by functioning as a coenzyme of γ -glutamyl carboxylase (GGCX) and also acts as a potent SXR ligand in bone metabolism

OSTEOPOROSIS

- ✘ is a skeletal disease characterised by low bone mass and microarchitectural deterioration with a resulting increase in bone fragility and hence susceptibility to fracture.
- ✘ It is an important public health issue because of the potentially devastating results and high cumulative rate of fractures; in white populations, about 50% of women and 20% of men older than 50 years will have a fragility fracture in their remaining lifetime.

Osteoporosis



Normal



Osteoporotic bone

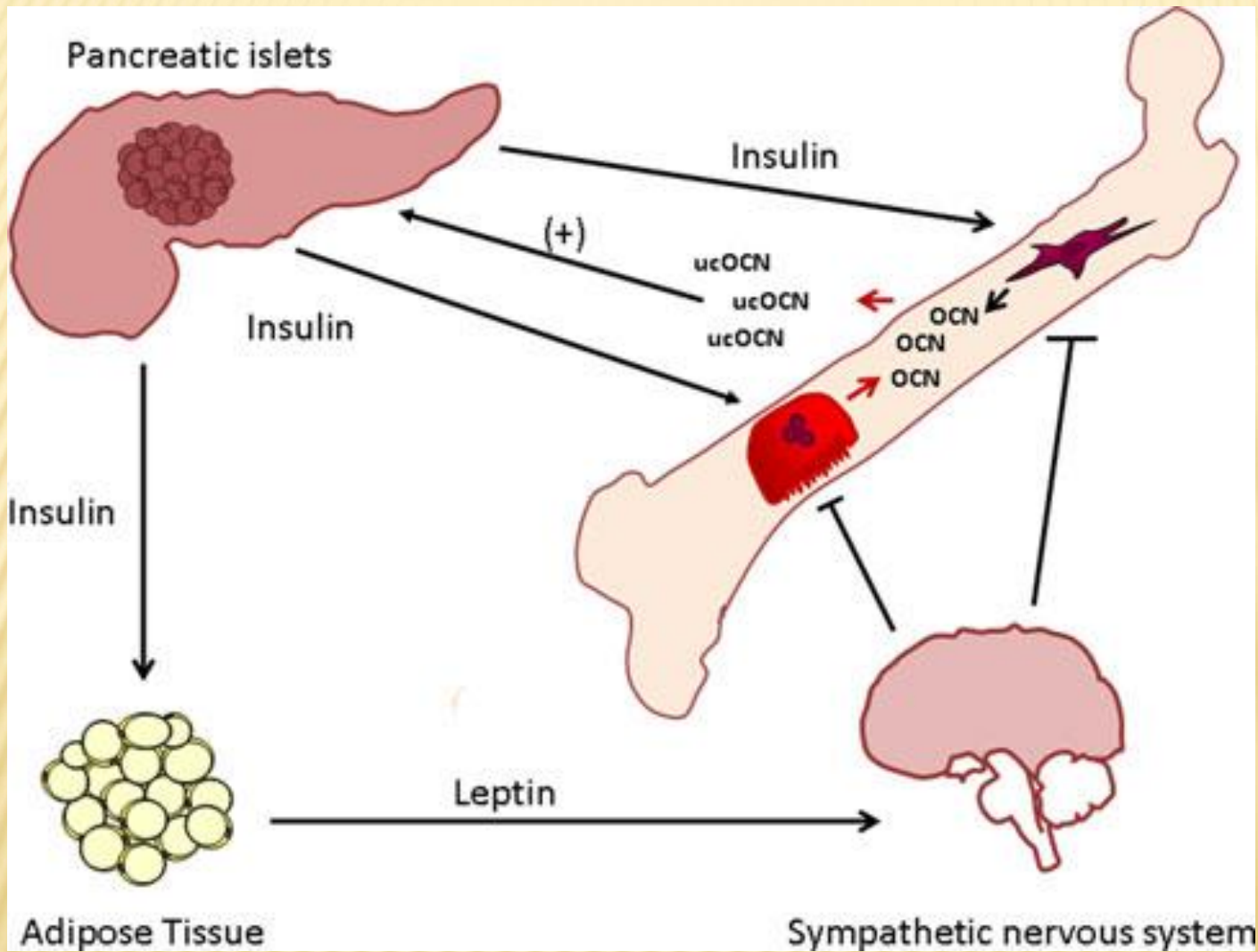
© Steiner 2004

OSTEOPOROSIS

- ✘ Oestrogen has a central role in normal physiological remodelling, and oestrogen deficiency after the menopause results in a remodelling imbalance with a substantial increase in bone turnover.
- ✘ This imbalance leads to a progressive loss of trabecular bone, partly because of increased osteoclastogenesis.
- ✘ Enhanced formation of functional osteoclasts seems to be the result of increased elaboration of osteoclastogenic proinflammatory cytokines such as interleukin-1 and tumour necrosis factor, which are negatively regulated by oestrogen.
- ✘ A direct effect of oestrogen in accelerating osteoclast apoptosis has also been attributed to increased production of transforming growth factor β .

OSTEOPOROSIS - CAUSES

- ✘ Glucocorticoids excess
- ✘ Estrogene deficiency
- ✘ Vitamin K2 deficiency?

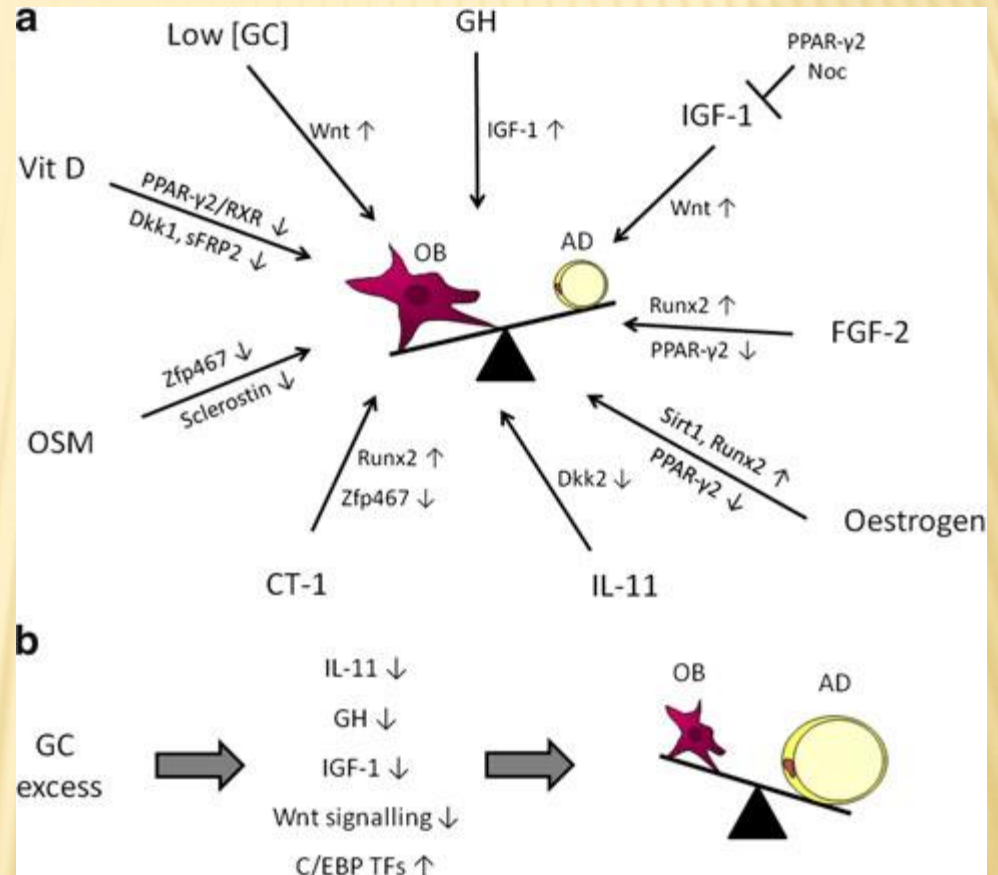


Cortisol generally antagonizes insulin ...

Equilibrium between osteoblastogenesis (OB) and adipogenesis (AD)

- a) Several endogenous factors support osteoblastogenesis against adipogenesis.
- b) High levels of cortisol prefer adipogenesis to osteoblastogenesis

Low [GC] low (physiological) concentrations of glucocorticoids, GH- growth hormone, IGF-1 insulin-like growth factor-1, FGF-2 fibroblast growth factor-2, IL-11 interleukin-11, CT-1 cardiotrophin-1, OSM oncostatin M, OB osteoblast, AD-adipocyte



COMMON ADVERSE EFFECTS OF GLUCOCORTICOID THERAPY

- ✘ There is substantial and accelerated decreases in bone mineral density (BMD) with oral glucocorticoid therapy, most pronounced in the first year, with trabecular bone more quickly affected than cortical bone.
- ✘ Even after only 2 months of high-dose glucocorticoids, studies show markedly decreased BMD at the lumbar spine, femoral neck and whole body, with the greatest loss in the trabecular lumbar vertebrae.
- ✘ In light of the high incidence of glucocorticoid-induced osteoporosis and associated fractures, screening and treatment rates for glucocorticoid-induced osteoporosis has come under substantial scrutiny.
- ✘ Less than 50% of patients receiving long-term glucocorticoids have been evaluated for osteoporosis, and less than 25% have been treated. There is great variability among clinicians in both the awareness of glucocorticoid-induced osteoporosis and the importance of prevention and treatment as the standard of care.
- ✘ The use of antiosteoporotic medication was observed to be most common among postmenopausal women, where it approached **50%**.

COMMON ADVERSE EFFECTS OF GLUCOCORTICOID THERAPY- **GLUCOCORTICOID-INDUCED OSTEOPOROSIS**

- ✘ Glucocorticoid-induced osteoporosis is the most common type of iatrogenic osteoporosis and a frequent cause of secondary osteoporosis.
- ✘ An estimated 50% of patients taking glucocorticoids for longer than 6 months will develop secondary osteoporosis .
- ✘ The absolute risk for glucocorticoid-induced osteoporosis is higher in patients aged 65 years or older given their baseline age-related fracture risk, although the relative risk of fracture related to glucocorticoid use may be even higher in patients under 65 .

PATHWAYS OF TRANSCRIPTION FACTORS PARTICIPATING IN NUTRITION BASED INTERACTIONS

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



SIZE AND ENDOCRINE PROFILE OF ADIPOCYTES IS CORRELATED WITH THE WHOLE ADIPOSITIVITY



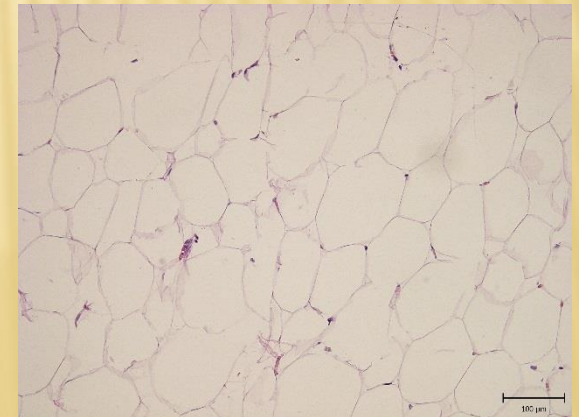
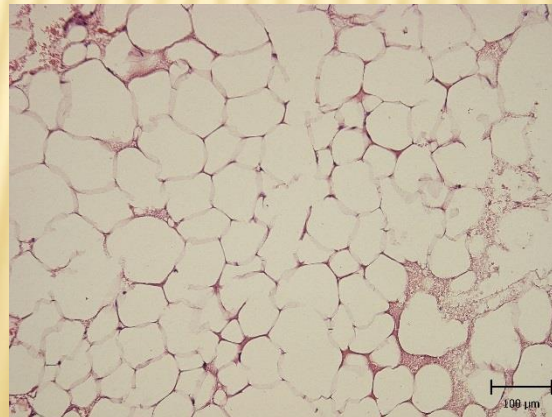
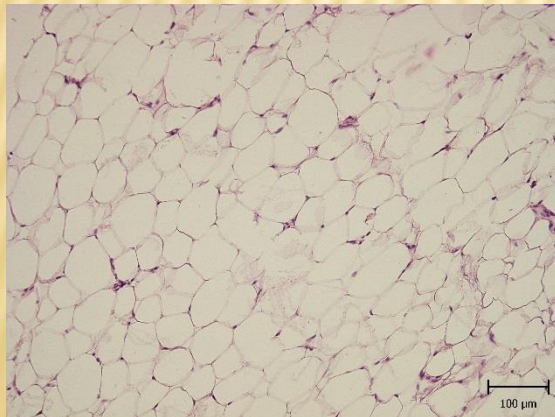
Malnutrition
(Anorexia nervosa)



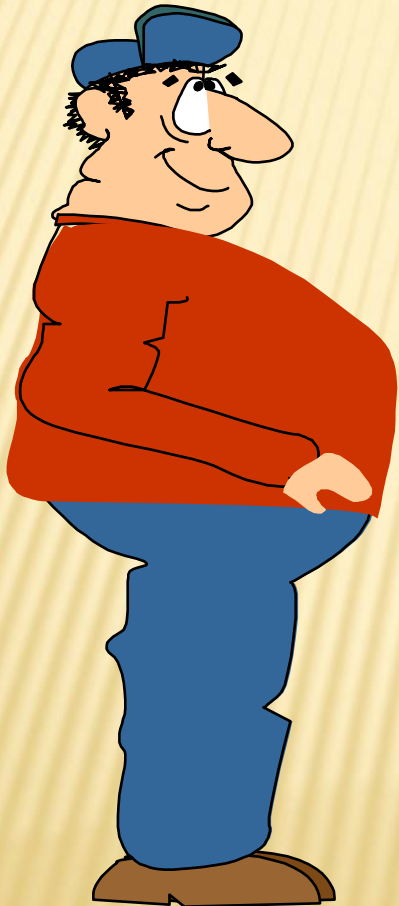
Normal state (slight overweight))



Obesity



DIAGNOSTIC CRITERIA OF OBESITY (BMI)



$$\text{BMI} = \frac{\text{Weight (kg)}}{\text{Height (m}^2\text{)}}$$

Classification	BMI (kg/m ²)	metabolic rate
Normal body weight	18.5–24.9	average
Overweight	25–29.9	increased
Obesity I	30.0–34.9	middle
Obesity II	35.0–39.9	high
Obesity III	≥40.0	very high

WAIST SIZE SEEMS TO BE THE BEST INDICATOR OF VISCERAL OBESITY

Women



Men

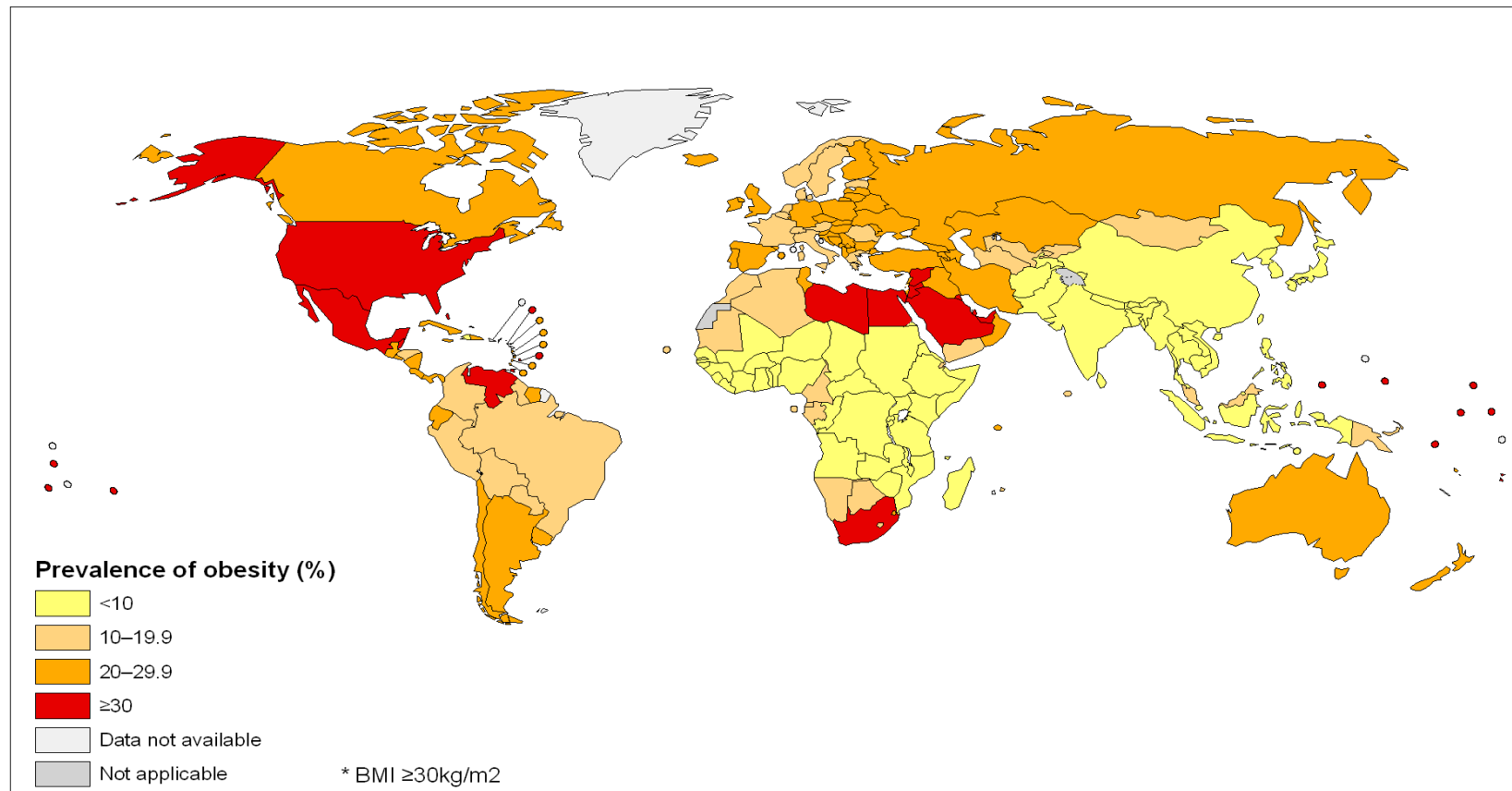
>88 cm = highly increase risk¹
>80 cm = higher risk¹

>102 cm = highly increased risk¹
>94 cm = higher risk¹

¹Lean MEJ, et al. Lancet;1998;351:853–6

STANDARDISED PREVALENCE OF OBESITY 2008

Prevalence of obesity*, ages 20+, age standardized Both sexes, 2008



The boundaries and names shown and the designations used on this map do not imply the expression of any opinion whatsoever on the part of the World Health Organization concerning the legal status of any country, territory, city or area or of its authorities, or concerning the delimitation of its frontiers or boundaries. Dotted lines on maps represent approximate border lines for which there may not yet be full agreement.

Data Source: World Health Organization
Map Production: Public Health Information
and Geographic Information Systems (GIS)
World Health Organization



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OBESITY

Has obesity genetic background?

Height in Feet	18	20	21	23	24	26	27	29	30	32	34	35	37	38
5'8	18	20	21	23	24	26	27	29	30	32	34	35	37	38
5'10	17	19	20	22	23	24	26	27	29	30	32	33	35	36
6'0	16	18	19	20	22	23	24	26	27	28	30	31	33	34
6'2	15	17	18	19	21	22	23	24	26	27	28	30	31	32
6'4	15	16	17	18	20	21	22	23	24	26	27	28	29	30
6'6	14	15	16	17	19	20	21	22	23	24	25	27	28	29
6'8	13	14	15	17	18	19	20	21	22	23	24	25	26	28

■ Healthy Weight ■ Overweight ■ Obese

GENETICS OF OBESITY

✘ Argumenst why yes:

HERITABILITY of OBESITY

Family studies	30-50%
Adoption studies	10-30%
Twin studies	50-90%

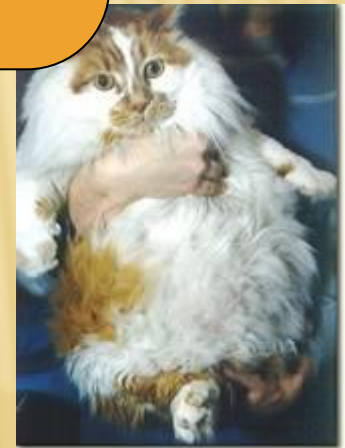
compared to DZ twins)

+ Large family studies

GENETICS OF OBESITY – ARGUMENTS WHY NOT



IS OBESITY DONE BY
CULTURAL EATING HABITS?

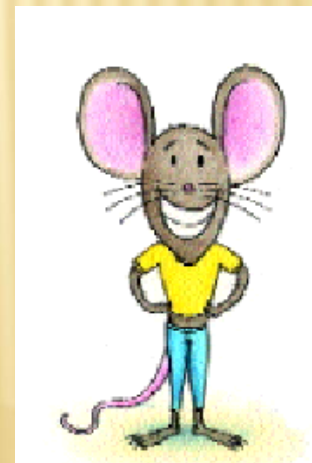


CLASSIFICATION OF OBESITY SYNDROMES

pleiotropic syndromes with obesity

„monogenic“ syndromes with obesity

polygenic (complex) syndromes



PLEIOTROPIC SYNDROMES WITH OBESITY

- ✘ About 30 syndromes, in which obesity represents constant component



MONOGENIC SYNDROMES WITH OBESITY

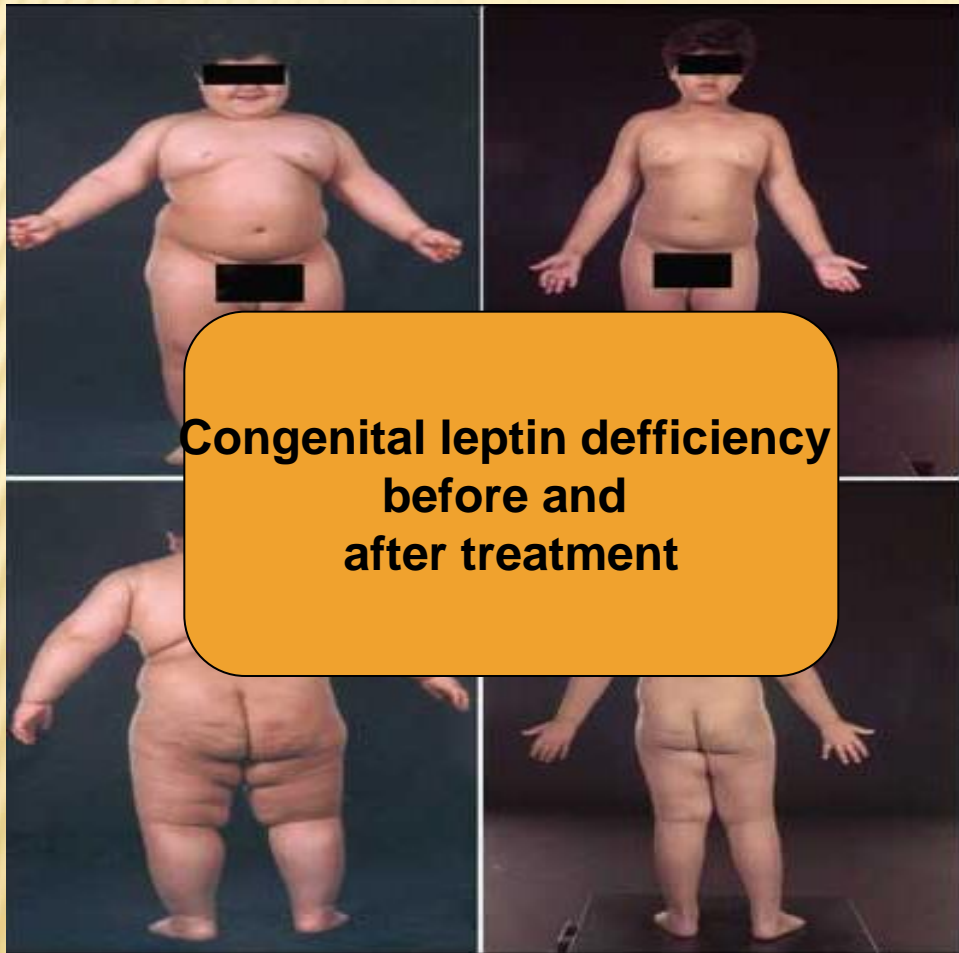
Genes Implicated in Monogenic Obesity and the Traits Found To Be Associated with Them in Genome-Wide Association Studies (GWAS)

Gene symbol	Gene name	Phenotype	Associated traits
<i>BDNF</i>	Brain-derived neurotrophic factor	Wilms tumor, aniridia, genitourinary anomalies, mental retardation, and obesity (WAGRO) syndrome	Obesity, BMI, weight
<i>CART</i>	Cocaine- and amphetamine-regulated transcript	Severe obesity	
<i>LEP</i>	Leptin	Morbid obesity due to leptin deficiency	
<i>LEPR</i>	Leptin receptor	Severe obesity due to leptin receptor deficiency	Serum level of C-reactive protein, serum level of leptin receptor
<i>MC4R</i>	Melanocortin-4 receptor	Early-onset severe obesity	Obesity, BMI, waist circumference, height, serum level of HDL cholesterol
<i>NTRK2</i>	Neurotrophic tyrosine kinase, receptor, type 2	Early-onset severe obesity, hyperphagia, developmental delay	
<i>PCSK1</i>	Proprotein convertase subtilisin/kexin type 1 gene, or prohormone convertase 1	Early-onset severe obesity	BMI, serum proinsulin level, fasting serum glucose level (interaction with BMI)
<i>POMC</i>	Proopiomelanocortin	Early-onset severe obesity, adrenal insufficiency, red hair	Obesity, height
<i>PPARG</i>	Peroxisome proliferator-activated receptor gamma	Severe obesity, insulin resistance, lipodystrophy	Type 2 diabetes, fasting serum insulin level (interaction with BMI), plasma level of plasminogen activator inhibitor type 1
<i>SIMI</i>	Single-minded homolog 1 (<i>Drosophila</i>)	Early-onset severe obesity, Prader-Willi syndrome	

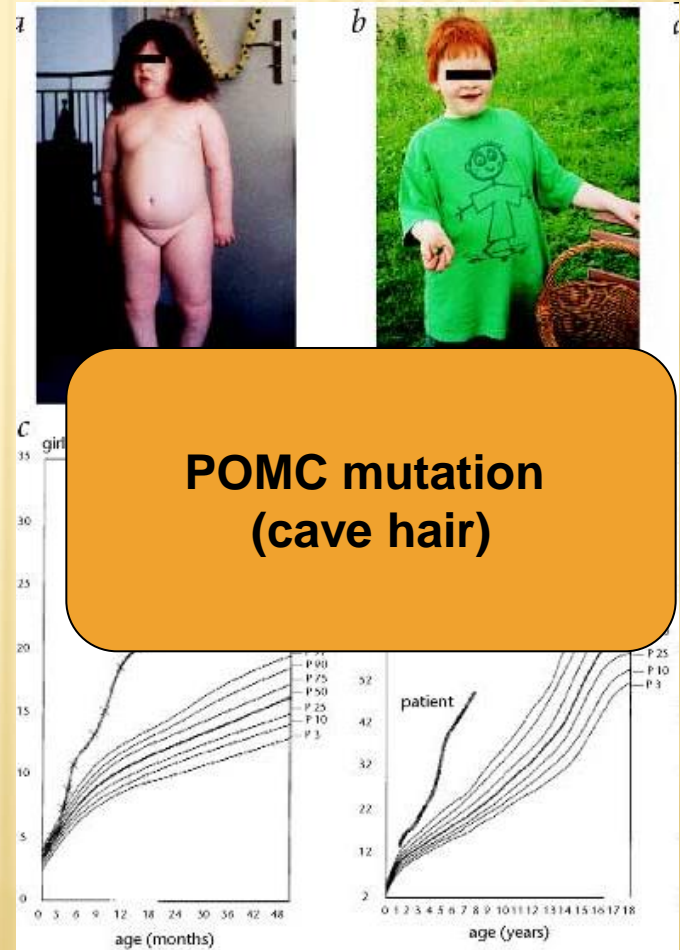
Note. BMI, body mass index; HDL, high-density lipoprotein.

N C Med J. 2013; 74(6):530-533

MONOGENIC OBESITY SYNDROMES



**Congenital leptin deficiency
before and
after treatment**



**POMC mutation
(cave hair)**

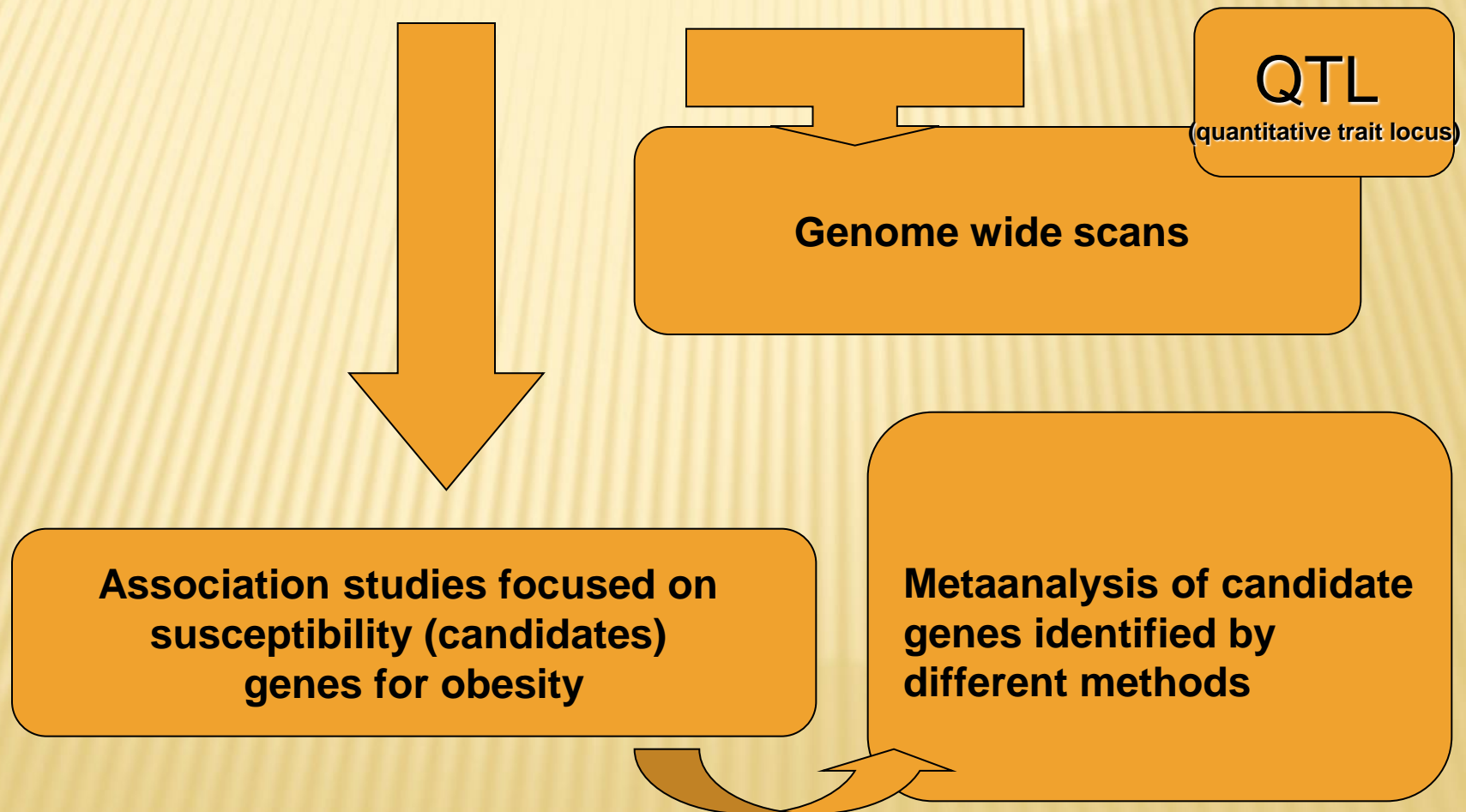
SY PRADER- WILLI AS A CLINICAL EXAMPLE

- ✘ Hypotonic children, mental retardation, small figure, behavioral complications (hyperphagy as a result of incontrollable appetite – one of the most common causes of children obesity)
- ✘ Loss of expression of paternally imprinted genes at 15q11.2-q13 chromosome as a result of microdeletion in the region.



OBESITY AS A COMMON (COMPLEX) DISEASE

✘ Susceptibility genes



FUTURE

- ✘ [BMC Med.](#) 2017; 15: 50.
- ✘ Published online 2017 Mar 7. doi: [10.1186/s12916-017-0800-1](#)
- ✘ PMID: PMC5340003
- ✘ **Developmental pathways to adiposity begin before birth and are influenced by genotype, prenatal environment and epigenome**
- ✘ [Xinyi Lin](#),^{#1} [Ives Yubin Lim](#),^{#1,2} [Yonghui Wu](#),¹ [Ai Ling Teh](#),¹ [Li Chen](#),¹ [Izzuddin M. Aris](#),¹ [Shu E. Soh](#),^{1,3} [Mya Thway Tint](#),^{2,3} [Julia L. Maclsaac](#),⁴ [Alexander M. Morin](#),⁴ [Fabian Yap](#),⁵ [Kok Hian Tan](#),⁵ [Seang Mei Saw](#),^{6,7,8} [Michael S. Kobar](#),⁴ [Michael J. Meaney](#),^{1,9} [Keith M. Godfrey](#),¹⁰ [Yap Seng Chong](#),^{1,2} [Joanna D. Holbrook](#),¹ [Yung Seng Lee](#),^{1,3,11} [Peter D. Gluckman](#),^{1,12} [Neerja Karnani](#),^{1,13} and on behalf of the GUSTO study group

OBESITY AS A COMPLEX DISEASE

- ✘ Genetic, epigenetic and prenatal environmental factors are linked to offspring size and adiposity at birth and in early childhood.
- ✘ Individual prenatal environmental influences on birth weight was identified; some of these prenatal environment variables [maternal ppBMI, GWG („Gestational Weight Gain“) and glucose levels] continued to associate with offspring size and adiposity in early childhood.

OBESITY AS A COMPLEX DISEASE

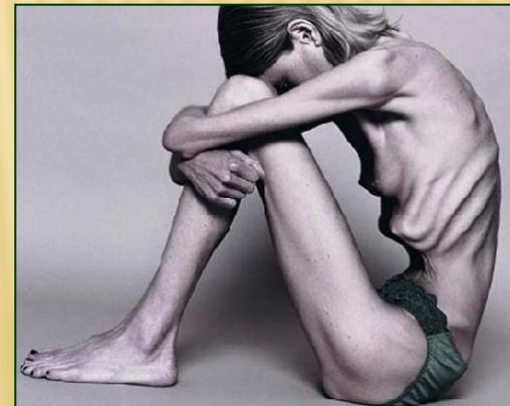
- ✘ Genetic variation, as captured by PRS („Polygenic Risk Score“), not only influenced birth weight, but also child size and adiposity up to 48 months of age, independent of birth weight.
- ✘ The PRS was constructed using adiposity-linked genetic risk variants previously reported in an adult population. The association of adult adiposity risk score with size and adiposity in pediatric population indicates that the effects of genetic risk variants can be detected as early as birth.

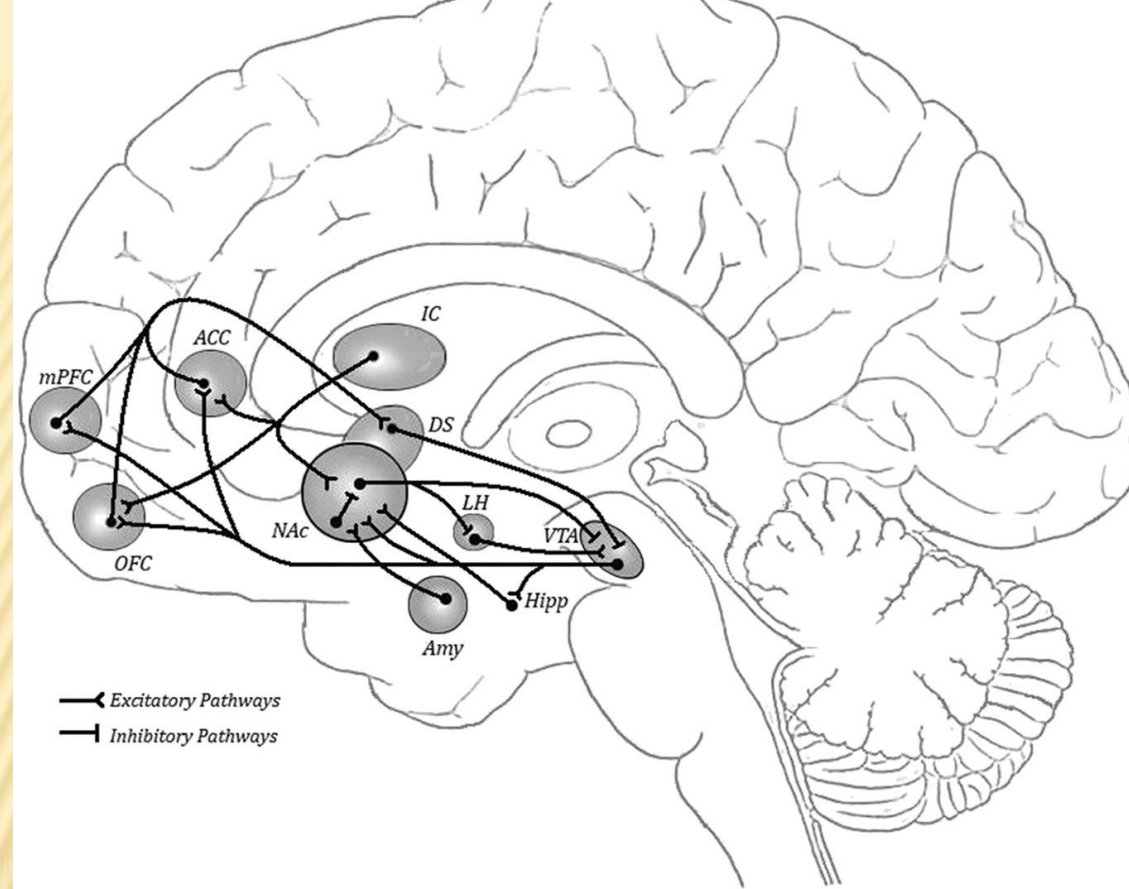
OBESITY AS A COMPLEX DISEASE ([BMC MED.](#) 2017; 15: 50)

- ✘ Neonatal methylation levels at seven loci were associated with birth weight. At six of the seven loci, there was suggestive evidence that the associations continued to persist up to 48 months of age. Among them, two of the loci (*CDKN2B/P4HA3*) also showed suggestive association with child BMI at 48 months. Even though the associations in early childhood did not survive multiple testing corrections, these CpGs still hold potential as biomarkers of adverse metabolic trajectory as the prevalence of obesity increases with age and might become more apparent later in the life-course. Lastly, methylation levels at three of seven loci associated with birth weight (*IGDCC4*, *MIRLET7BHG*, *CACNA1G*) also showed significant associations with the prenatal environment; however, similar analyses with childhood weight and adiposity measures showed suggestive associations.
- ✘ Birth weight seems to be influenced by both genetic and prenatal environment factors, possibly acting through different mechanisms, either by altering the epigenome (evidenced by CpGs that were associated with prenatal environment and/or SNPs) or independently of the epigenome (e.g. the PRS).

ANOREXIA NERVOSA (AN) AND BULIMIA NERVOSA (BN)

- ▶ are complex psychiatric disorders of great importance for public health policies, as they are associated with a high burden of morbidity and mortality due to their severe medical and psychological consequences. Etiopathogenesis of these eating disorders (EDs) continues to remain elusive, with the result that their treatment is often unsuccessful.
- ▶ AN is a severe psychiatric disorder leading to life-threatening weight and fat loss. This illness could be characterized by irrational fear of becoming fat, abnormal eating behavior, hyperactivity, GIT complications and wide variety alterations of hormonal and metabolic systems.
- ▶ The exact etiopathogenesis is unknown and the way of treatment remain limited.





Schematic representation of brain reward circuits.

ACC anterior cingulate cortex; Amy amygdala; DS dorsal striatum; Hipp hippocampus; IC insular cortex; LH lateral hypothalamus; mPFC medial prefrontal cortex; NAc nucleus accumbens; OFC orbitofrontal cortex; VTA ventral tegmental area.

The brain reward system integrates basic and emotional stimuli, such as hunger, satiety, desire, pleasure and fear, with higher order cognitive processes aimed at modulating further actions or representation of the general experience. These high order processes involve anterior cingulate cortex (ACC), orbitofrontal cortex (OFC) and ventromedial prefrontal cortex (PFC), which are necessary for identification of rewarding stimuli, inhibition of emotional responses, and promote behavioral outcomes ([Haber and Knutson, 2010](#); [Wittman et al., 2010](#); [Sripada et al., 2011](#)). Overall, the PFC provides inhibitory influences on motivation and reward-directed behavior, integrating sensory inputs, memories, goals, and physiological states with the aim to provide an adequate performance ([Miller and Cohen, 2001](#)). ACC and dorsolateral prefrontal cortex (DLPFC) may also serve to monitor potential conflict situations induced by reward stimuli ([Walton et al., 2003](#); [Vogt et al., 2005](#)). Therefore, they have a *gating role* in action selection following reward cues ([Goldstein and Volkow, 2011](#)). Indeed, by a top-down effect, both OFC and ACC provide a negative feedback to mesolimbic areas regulating reward-seeking motivation ([Goldstein and Volkow, 2011](#)).

ANOREXIA NERVOSA (AN) AND BULIMIA NERVOSA (BN)

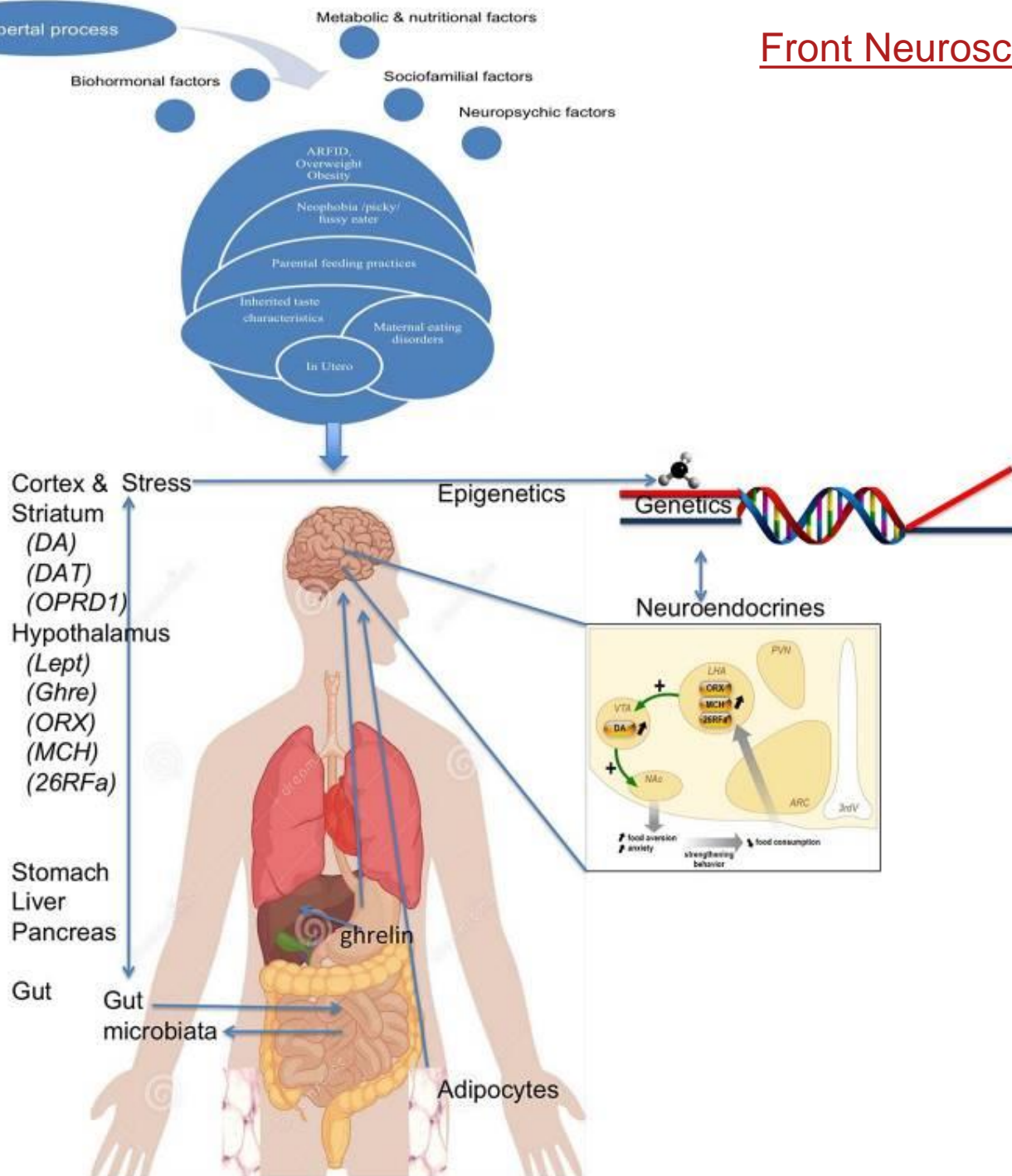
- ✘ Functional magnetic resonance imaging (fMRI) techniques have been employed to investigate the brain's processing of reward elicited by both food-related and non-food-related stimuli in AN and BN ([O'Hara et al., 2015](#)).
- ✘ It has been shown that, compared to healthy controls, AN patients exhibit **abnormal activation of different brain areas, including the parietal, the orbito-frontal, the dorso-lateral prefrontal, the anterior cingulate and the medial prefrontal cortex** ([Frank, 2015a](#) ; [Frank, 2015b](#)) after exposure to visual food cues, especially for highly palatable foods.
- ✘ Similarly, **altered insula, striatum or orbitofrontal responses to sweet stimuli have been found in recovered or symptomatic AN and BN patients.**
- ✘ These findings suggest a dysregulation of brain mechanisms involved in the processing of food-related rewarding stimuli in the pathophysiology of EDs.

PATHOPHYSIOLOGY OF AN

- ✘ An integral pathophysiological scenario that fits to the natural history of AN with the following steps can be proposed:
- ✘ (1) enhanced vulnerability to stress (genetic, epigenetic, or environmental factors);
- ✘ (2) major stressing events activating the stress-axis, increased intestinal permeability, and increased virulence of the microbiota;
- ✘ (3) bacterial proteins (e.g., ClpB) challenge the immune response and due to molecular mimicry, cause increased production of Igs cross-reactive with neuropeptides (e.g., α -MSH);
- ✘ (4) this results in altered food intake, anxiety, gastrointestinal discomfort and other consequences of altered central and peripheral melanocortin signaling;
- ✘ (5) global malnutrition and some specific macro- and micro-nutrient deficiencies contribute to the perpetuation of gut barrier and immune dysfunction as well as behavioral symptoms.

OREXIGENIC NEUROPEPTIDES

- ✘ Orexigenic neuropeptides including ghrelin, orexins and 26RFa are up-regulated in AN and it is thought that this orexigenic profile reflects an adaptive mechanism of the organism to promote food intake and thus to counteract undernutrition. However, this adaptive mechanism is ineffective in increasing food consumption leading to the **concept of a global resistance of AN patients to orexigenic signals**.
- ✘ We can speculate that a chronic increase of the activity of LHA orexigenic neurons expressing orexins, MCH, or 26RFa could reinforce dopamine-induced anxiety in the reward system of AN patients and thus the aversion to ingest food.



LOSS OF ADIPOSE TISSUE IS SUPPOSED TO BE THE MAIN FACTOR CONTRIBUTING TO THE BODY WEIGHT LOSS

	NW (n = 50)	AN (n = 30)
Weight (kg)	62.2 ± 1.54	45.8 ± 1.89*
Lean mass (kg)	39.1 ± 0.76	37.8 ± 1.01
BMI (kg/m²)	21.2 ± 0.42	15.7 ± 0.47*
Body fat content (%)	24.3 ± 0.79	7.1 ± 0.88*
Total fat skinfold (mm)	120.5 ± 12.17	42.1 ± 4.78*
Abdominal skinfold (mm)	12.5 ± 2.13	4.8 ± 1.59*
Insulin (pmol/l)	28.3 ± 4.53	14.2 ± 3.67*
Glucose (mmol/l)	4.7 ± 0.08	4.1 ± 0.11
Menstruation	Yes - regular cycle	Secondary amenorrhea

GENETIC ASSOCIATIONS???

- ✘ AN is 11x more frequent in relatives of probands compared to physiological population.
- ✘ BN is 4-5x more frequent in relative women.
- ✘ ~15% risk of eating disorders in relatives of AN and BN vs. 4% risk in healthy population.

DĚKUJI VÁM ZA POZORNOST

