

Mechanisms and Metabolic Implications of Regional Differences among Fat Depots

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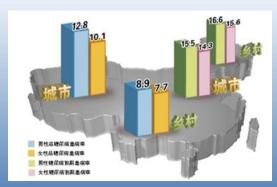
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Introduction



The widespread of diabetes

现实严峻



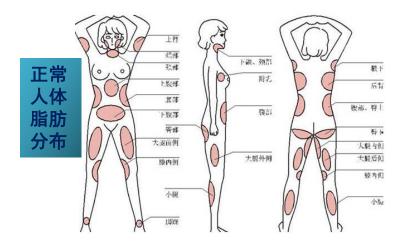
中国糖尿病患者持续增加



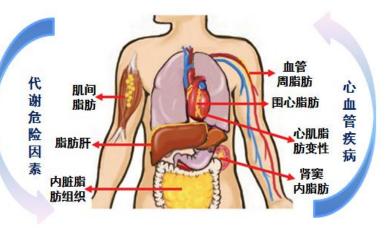
男性糖尿病比 女性风险高20%



肥胖和糖尿病主要 是机体脂代谢紊乱造成



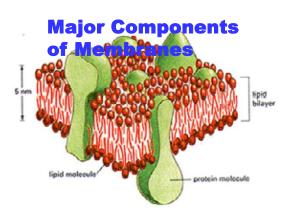
异 脏 好 防 活 者 全 局 交 数



Fat-Tissue Function

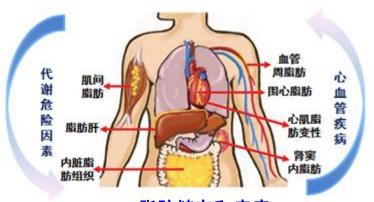








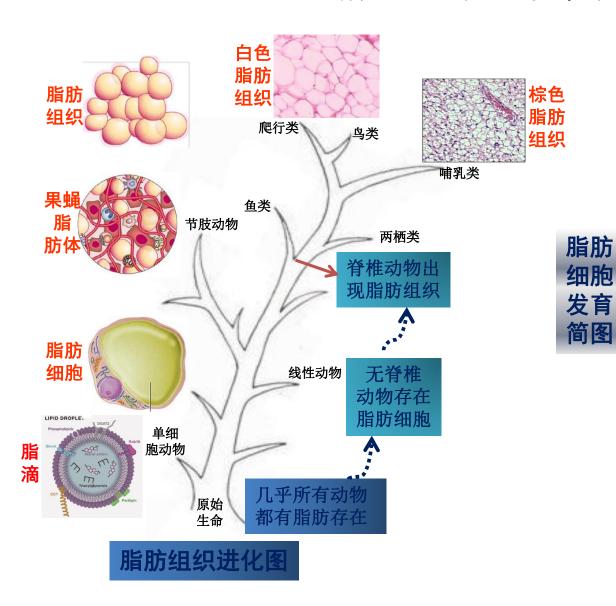


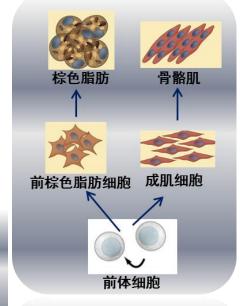


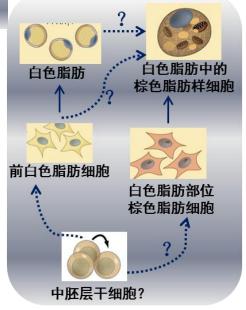
脂肪储存和疾病

脂类功能

脂肪组织的演化







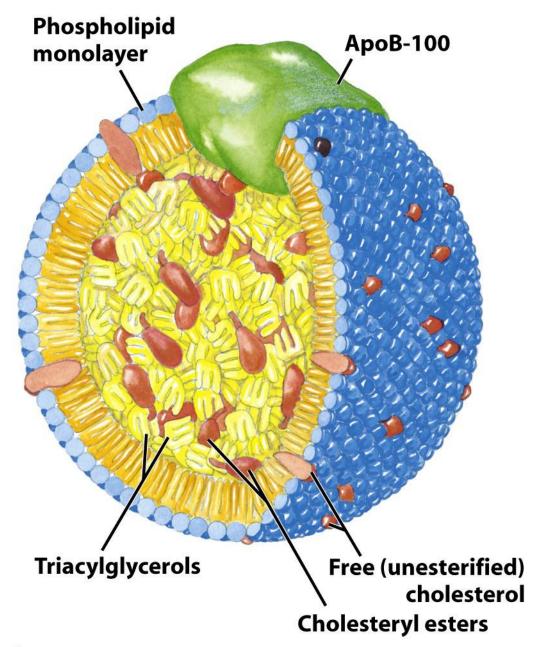
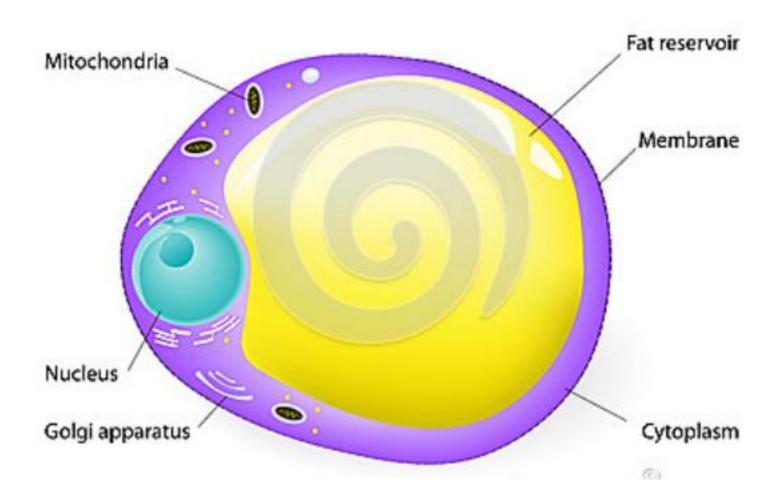


Figure 21-39a
Lehninger Principles of Biochemistry, Fifth Edition
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ADIPOCYTE



脂肪组织分布的多样性

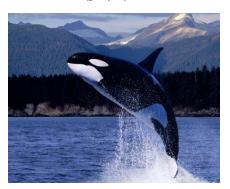
不同动物脂肪储存部位不一样

Gesta et al. Cell. 2007

Speci	es	Caenorhabditis elegans	Drosophila melanogaster	Carcharodon carcharias	Cyprinus carpio	Xenopus laevis	Gallus gallus domesticus	Mus musculus	Homo sapiens
Fat sto	rage	Stored in intestinal cells	Stored in the "fat body"	Stored in liver	Stored in WAT	Intra-abdominal WAT (no sub- cutaneous WAT)	Subcutaneous and internal WAT	Subcutaneous and internal WAT	Subcutaneous and internal WAT

哺乳动物脂肪储存部位差异很大

皮下



驼峰



全身



全身/浅皮 下脂肪



脂肪组织分布的多样性

相近、同种动物脂肪储存的主要部位不一样





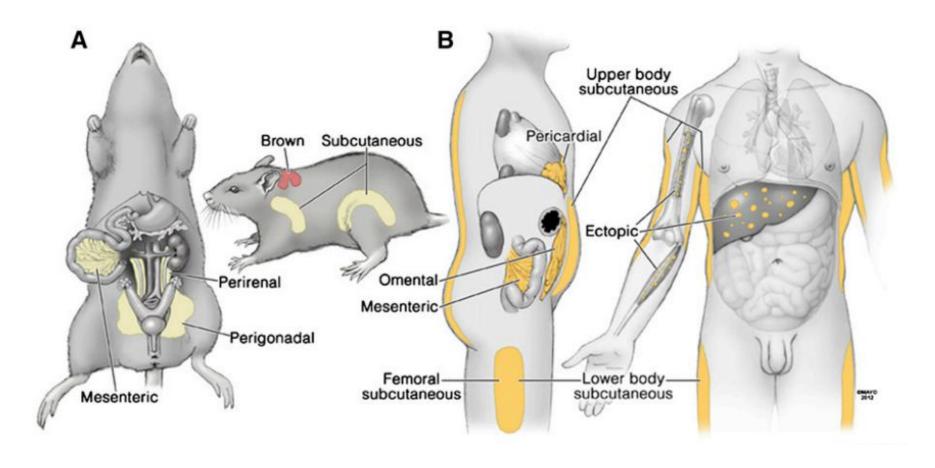








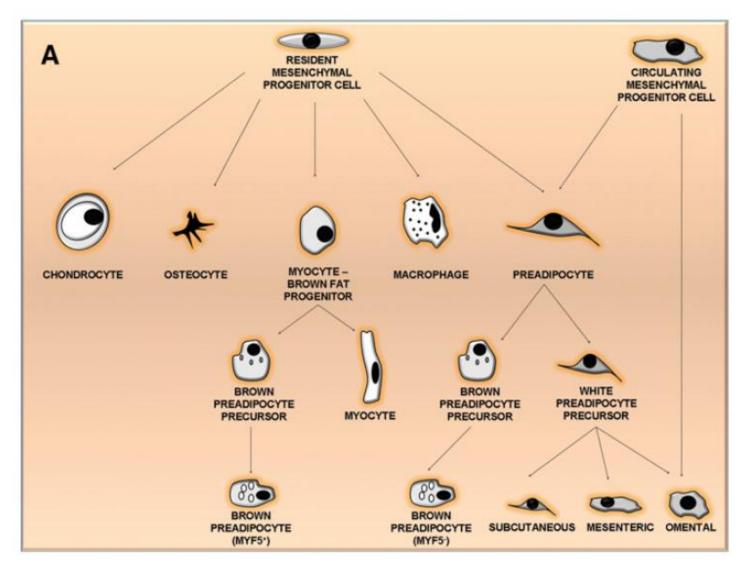
冬眠的动物和迁徙的鸟类异<u>位脂</u> 肪存储不会导致相应的疾病



Anatomy of Major Fat Depots in Rodents and Humans

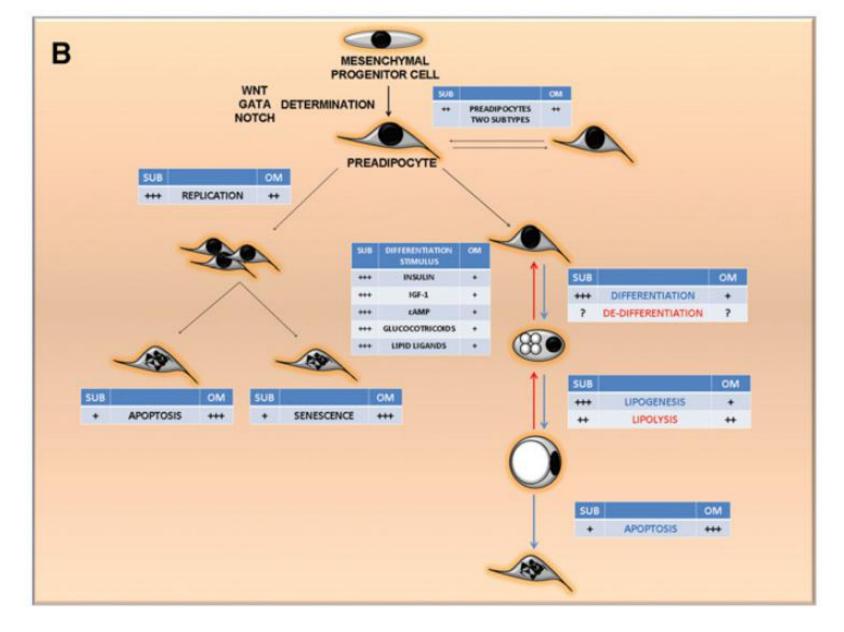
intra-abdominal:omental and mesenteric depots, also termed visceral fat lower-body:gluteal fat, subcutaneous leg fat, and intramuscular fat upper-body: subcutaneous fat

Cellular Mechanisms of Fat Growth and Function

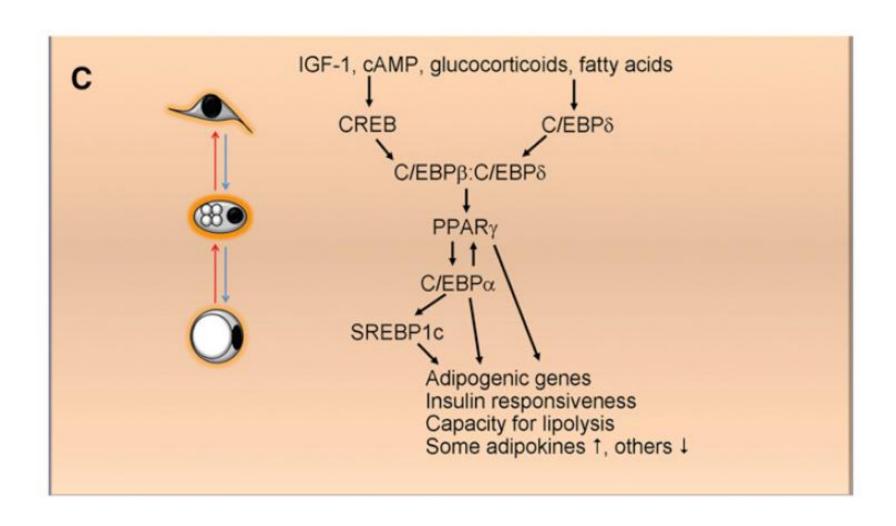


Circulating progenitors may contribute, especially to visceral fat developent

Regional variation in fat-tissue cell dynamics



preadipocytes can replicate, differentiate into adipocytes, or possibly revert into multipotent progenitors again

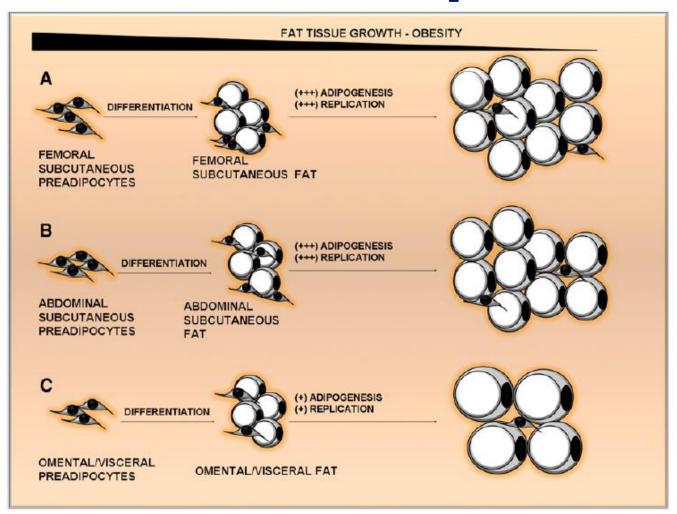


Key transcription factors involved in adipogenesis.

Adipokine or		L. S. S. S.	
Secreted Factor	Source Cells	Depot	References
Leptin*	fat cells	subcutaneous > omental	(Wiest et al., 2010)
Adiponectin* HMW	fat cells	omental > subcutaneous	(Kovacova et al., 2012)
PAI-1	preadipocytes	possibly, omental > subcutaneous	(Xu et al., 2012)
IL-6*	preadipocytes, macrophages, activated endothelial cells, large fat cells	visceral > subcutaneous	(Fontana et al., 2007)
TNF-α*	preadipocytes, macrophages, adipocytes	mesenteric > omental = subcutaneous	(Cartier et al., 2008; Xu et al., 2012)
MCP-1*	preadipocytes, macrophages	visceral > subcutaneous	(Madani et al., 2009; Miller et al., 2011)
Angiotensinogen	fat tissue	omental > subcutaneous	(Dusserre et al., 2000; van Harmelen et al., 2000)
RANTES*	stromal vascular fraction, fat cells	gastric fat pad > omental = subcutaneous	(Madani et al., 2009)
CSF-1	endothelial cells, fibroblasts	visceral > subcutaneous	(Harman-Boehm et al., 2007)
Omentin	stromal vascular cells	omental > subcutaneous	(Yang et al., 2006)
RBP4	preadipocytes, adipocytes	visceral < subcutaneous	(Kos et al., 2011)
Chimerin	unknown	visceral < subcutaneous	(Alfadda et al., 2012)
Vaspin	fat cells	visceral > subcutaneous	(Hida et al., 2005; Klöting et al., 2006)

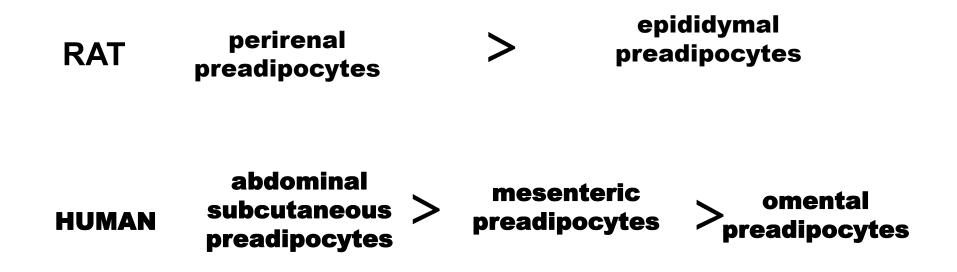
Preadipocytes generate adipokines, paracrine factors, hormones, and metabolic signals in a manner distinct from fat cells. Several adipokines are secreted in a fat-depot-dependent fashion

Regional Differences in Preadipocyte Characteristics Replication



Mechanisms of Fat-Tissue Growth during the Progression of Obesity Vary among Depots

Differences in cell-dynamic properties of preadipocytes are partly cell autonomous



replication

Adipogenesis

- abdominal subcutaneous preadipocytes had a greater capacity for adipogenesis than omental cells
- PPAR γ and C/EBPa expression is higher in human abdominal subcutaneous than in omental differentiating preadipocytes

adipogenesis appears to vary among depots

Apoptosis and Senescence

- Apoptotic fat cells are more abundant in human omental than abdominal subcutaneous fat, which may induced by serum deprivation or TNF-a
- Cellular senescence may contribute to regional variation in age- and obesity-related increases in fat-tissue inflammation, glucose intolerance, loss of capacity to store fat in subcutaneous depots, and lipotoxicity.

Inherent Differences

- Genome-wide expression profiles of primary preadipocytes cultured in parallel from different depots of mice and humans are highly distinct. Expression of >500 genes varies significantly among human abdominal subcutaneous, omental, and mesenteric preadipocytes and fat tissue.
- Whether developmental regulators directly cause regional differences in preadipocyte and fat-depot characteristics is being actively in vestigated.
- Most developmental genes are subject to epigenetic regulation

A Special Role for Leg Fat?

- leg fat appears to serve an important role in disposing of excess dietary fat in women
- Adipose tissue lipolysis in leg fat is normally exquisitely sensitive to insulin this makes lower-body depots an ideal place to store fat when it is ingested in excess of short-term energy needs.
- Greater amounts of leg fat signal a lesser metabolic risk and a more normal fatty-acid profile, although whether leg fat plays a protective role or (opposite of visceral fat) signals generally normal function of subcutaneous fat remains to be determined.

Mesenteric Fat: An Underappreciated Role?

- Mesenteric fat may make an important contribution to metabolic dysfunction, acting in a manner distinct from omental fat.
- The cellular and gene-expression properties of human mesenteric fat are cell autonomously distinct from omental fat.

Metabolically Protective Role of Subcutaneous Fat

- Certain subcutaneous fat regions appear to be metabolically, immunologically, and mechanically protective.
- The high capacity for certain subcutaneous regions to generate new fat cells may prevent visceral fat enlargement and systemic lipotoxicity.

Intra-abdominal Fat: Cause or Indicator?

- Central fat is associated with elevated risk for diabetes, hypertension, atherosclerosis, dyslipidemia, and cancers.
- In normal-weight or moderately overweight people, visceral obesity is strongly associated with insulin resistance, but in severe obesity, it is a weak independent predictor.

Conclusions

- Fat distribution is closely linked to metabolic disease risk.
- Distribution varies with sex, genetic background, disease state, certain drugs and hormones, development, and aging.
- Preadipocyte replication and differentiation, developmental gene expression, susceptibility to apoptosis and cellular senescence, and adipokine secretion vary among depots.
- How interdepot differences in these molecular, cellular, and pathophysiological properties are related is incompletely understood.
- Whether fat redistribution causes metabolic disease or whether it is a marker of underlying processes that are primarily responsible is an open question.

thanks

