

Donald Voet • Judith G. Voet • Charlotte W. Pratt

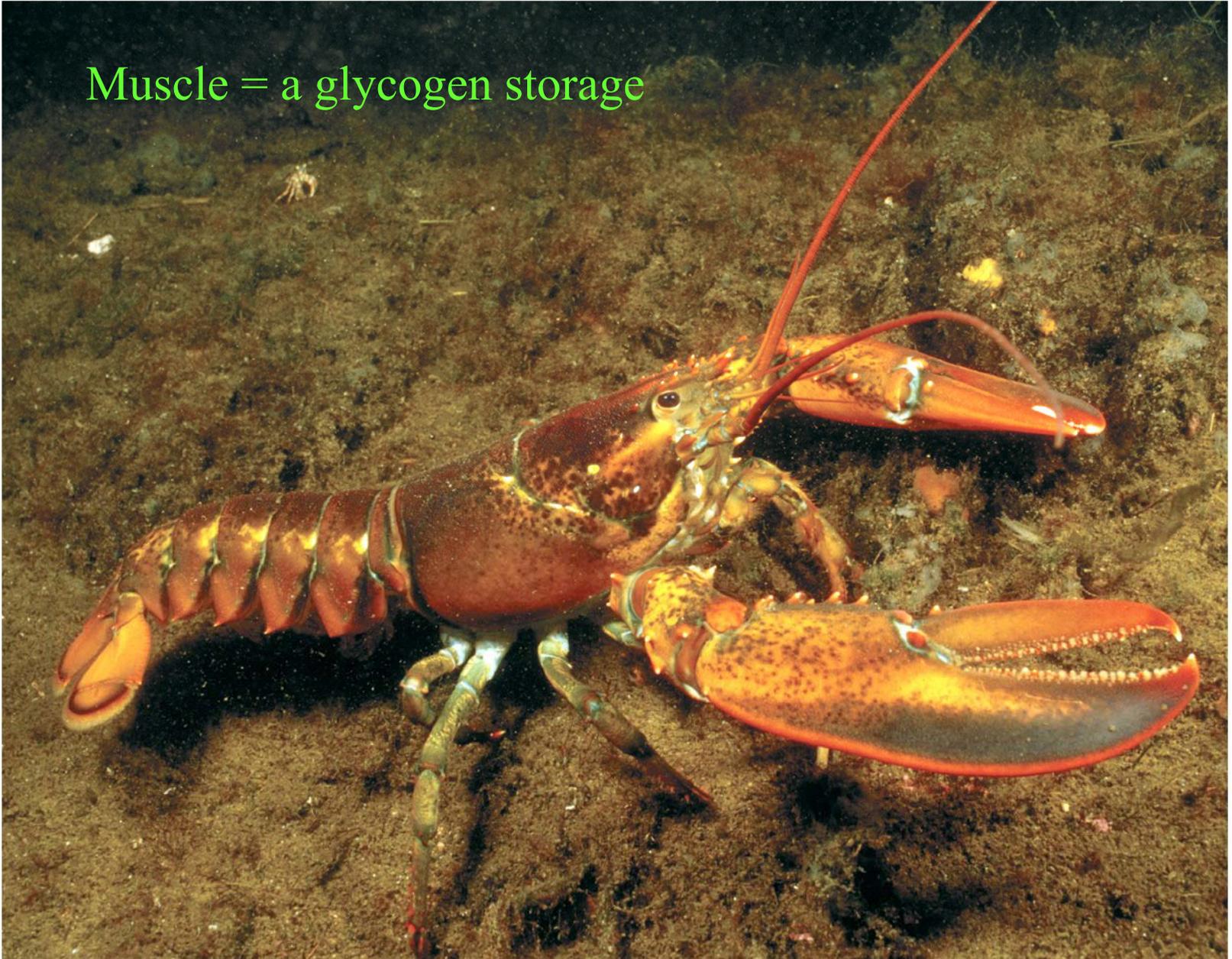
Fundamentals of Biochemistry

Second Edition

Chapter 15:

Glycogen Metabolism and Gluconeogenesis

Muscle = a glycogen storage



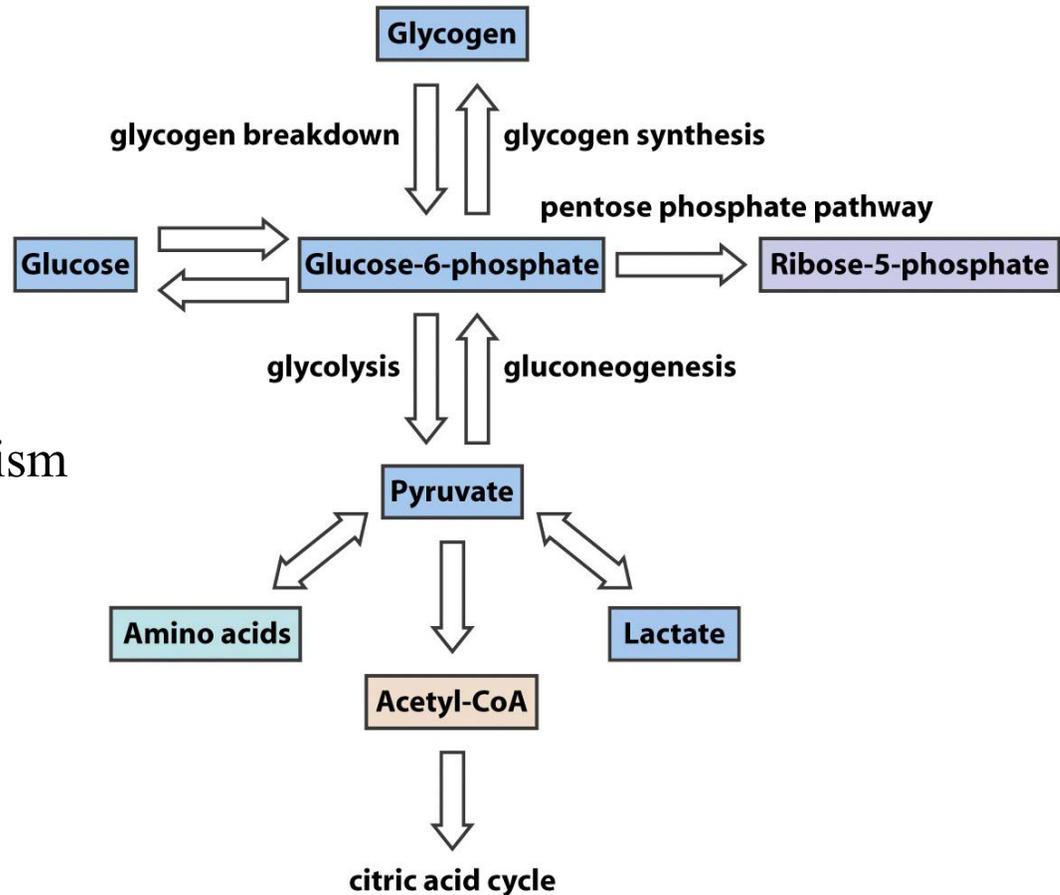
Glucose storage

Glycogen: animals, fungi, and bacteria

Starch: plants

Constant supply of glucose in animals: ~5 mM in blood

Maintenance of the concentration



Overview of glucose metabolism

Glycogen breakdown (glycogenolysis)

$\alpha(1-4)$ -linked D-glucose with $\alpha(1-6)$ -linked branches every 8-14 residues

Intracellular granules of 100~400 Å diameter

Abundant in muscle (up to 1-2%) and liver (up to 10%)

A complex with enzymes catalyzing synthesis and breakdown

Glycogen phosphorylase

Glycogen debranching enzyme

phosphoglucomutase

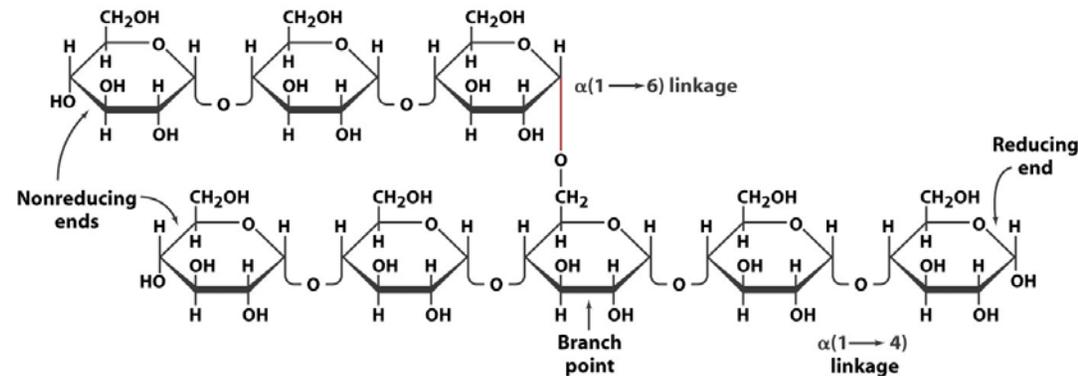


Figure 15-2a Fundamentals of Biochemistry, 2/e
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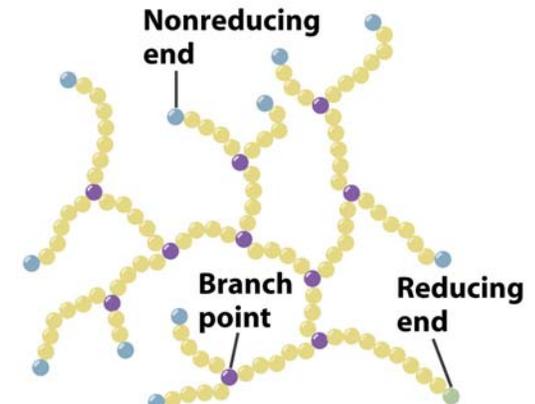


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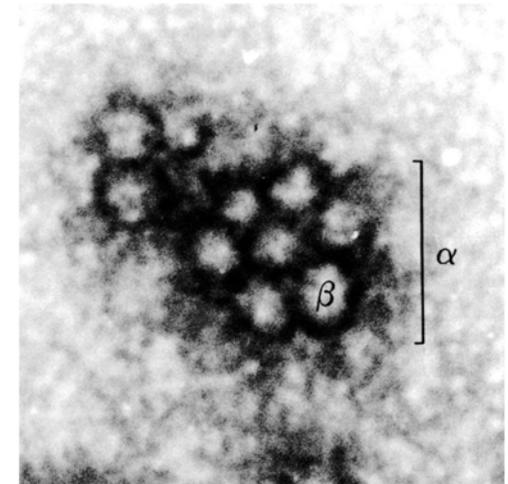
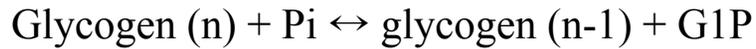


Figure 15-2c Fundamentals of Biochemistry, 2/e

Glycogen phosphorylase



Dimer of 97 kD subunits

Allosteric regulation

Inhibitors: ATP, G6P, glucose

Activator: AMP

Covalent modification (ser-14)

Phosphorylase a (phosphorylated)

Phosphorylase b (dephosphorylated)

C-terminal

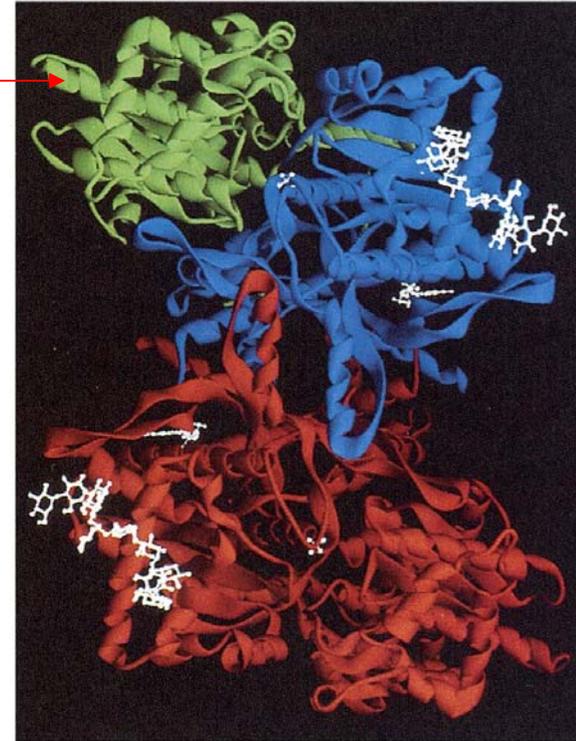


Figure 15-3a Fundamentals of Biochemistry, 2/e

Continuous feeding of substrate

Narrow crevice:
5 units away from a branching point

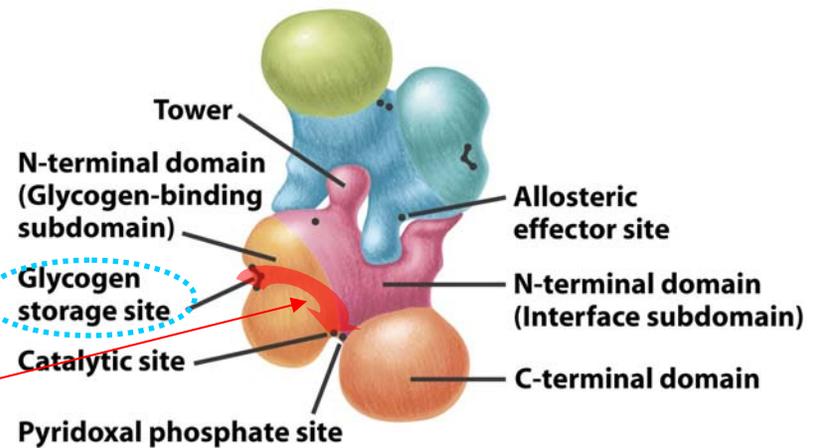


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Glycogen debranching enzyme

Two separate active sites:

Transferase: $\alpha(1-4)$ transglycosylase

$\alpha(1-4)$ glucosidase

The maximal rate is slower than phosphorylase

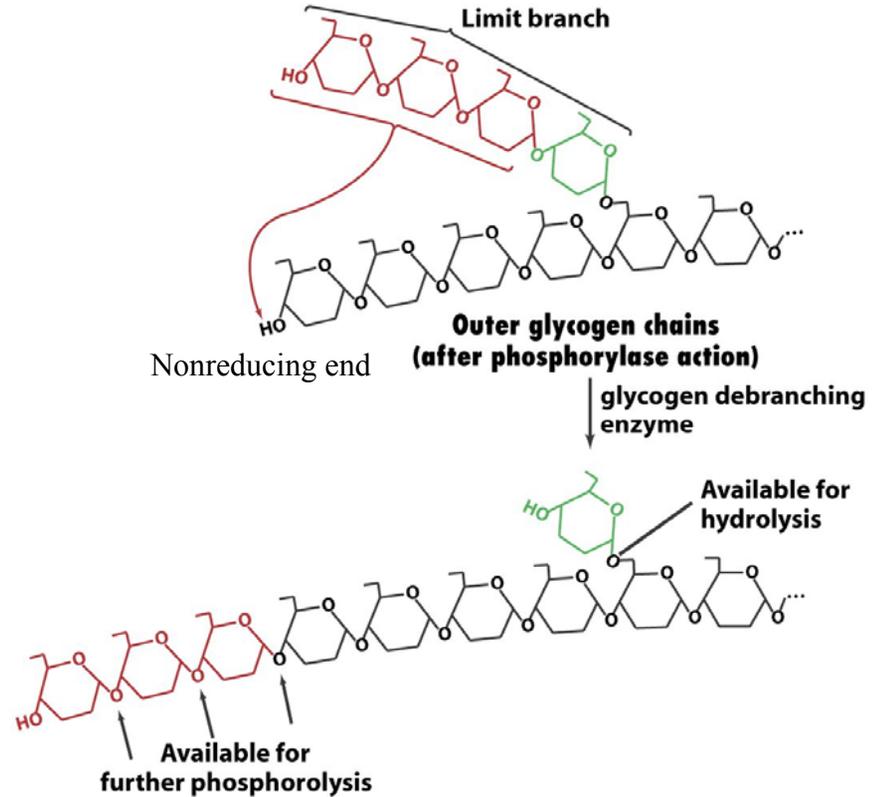


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Phosphoglucomutase

G1P to G6P via G1,6P

Hexokinase step is bypassed

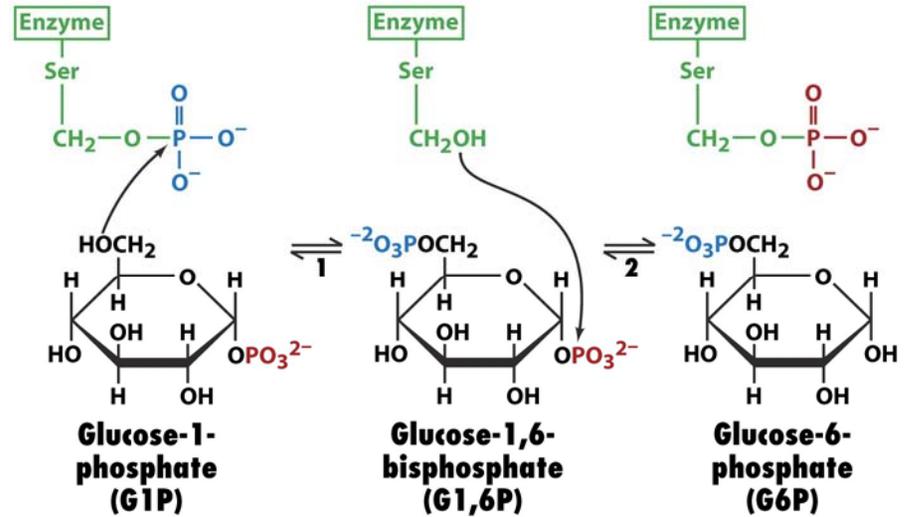


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Glucose-6-phosphatase

G6P to Glucose

Liver enzyme in ER membrane

G6P is transported to ER and hydrolyzed

Leave liver via GLUT2

Type I glycogen storage disease: G6P process defect in liver

Hereditary Glycogen Storage Diseases

Type	Enzyme Deficiency	Tissue	Common Name	Glycogen Structure
I	Glucose-6-phosphatase	Liver	von Gierke's disease	Normal
II	α -1,4-Glucosidase	All lysosomes	Pompe's disease	Normal
III	Amylo-1,6-glucosidase (debranching enzyme)	All organs	Cori's disease	Outer chains missing or very short
IV	Amylo-(1,4 \rightarrow 1,6)-transglycosylase (branching enzyme)	Liver, probably all organs	Andersen's disease	Very long unbranched chains
V	Glycogen phosphorylase	Muscle	McArdle's disease	Normal
VI	Glycogen phosphorylase	Liver	Hers' disease	Normal
VII	Phosphofructokinase	Muscle	Tarui's disease	Normal
VIII	Phosphorylase kinase	Liver	X-linked phosphorylase kinase deficiency	Normal
IX	Phosphorylase kinase	All organs		Normal
0	Glycogen synthase	Liver		Normal, deficient in quantity

Box 15-2 table 1 Fundamentals of Biochemistry, 2/e
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Incidence?

Glycogen synthesis

UDP-glucose pyrophosphorylase

Glycogen synthase

Glycogen branching enzyme

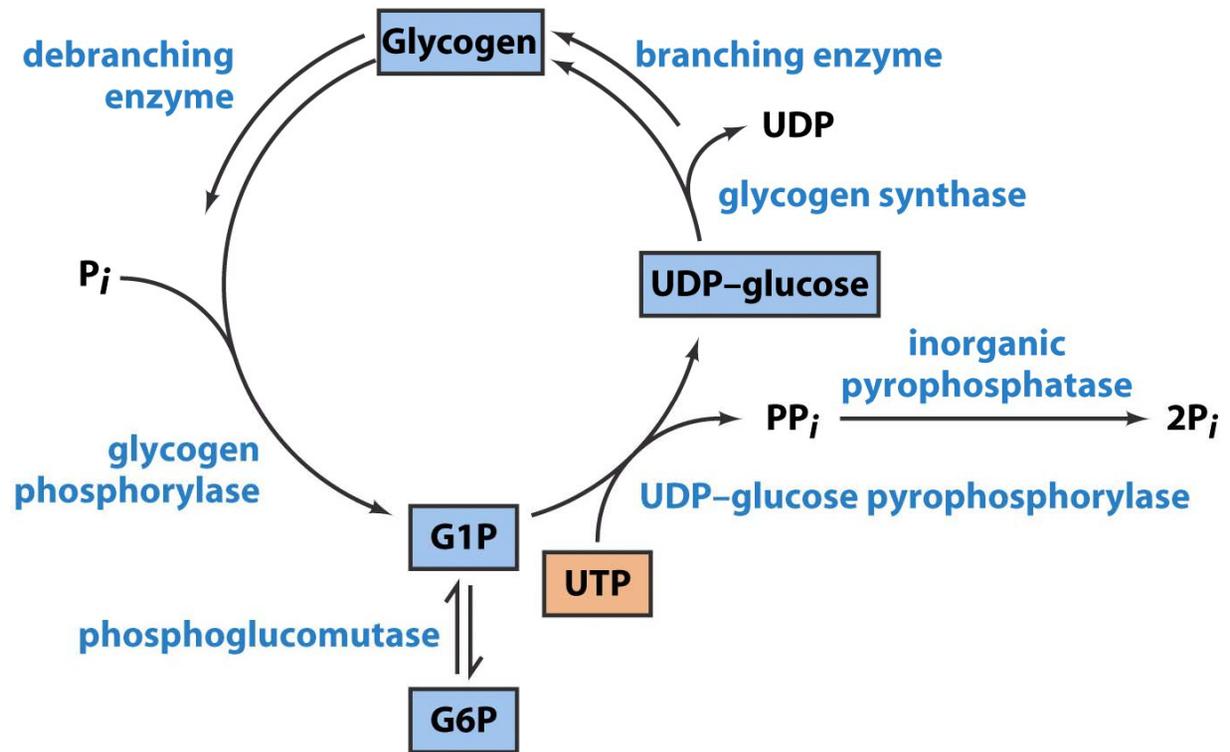
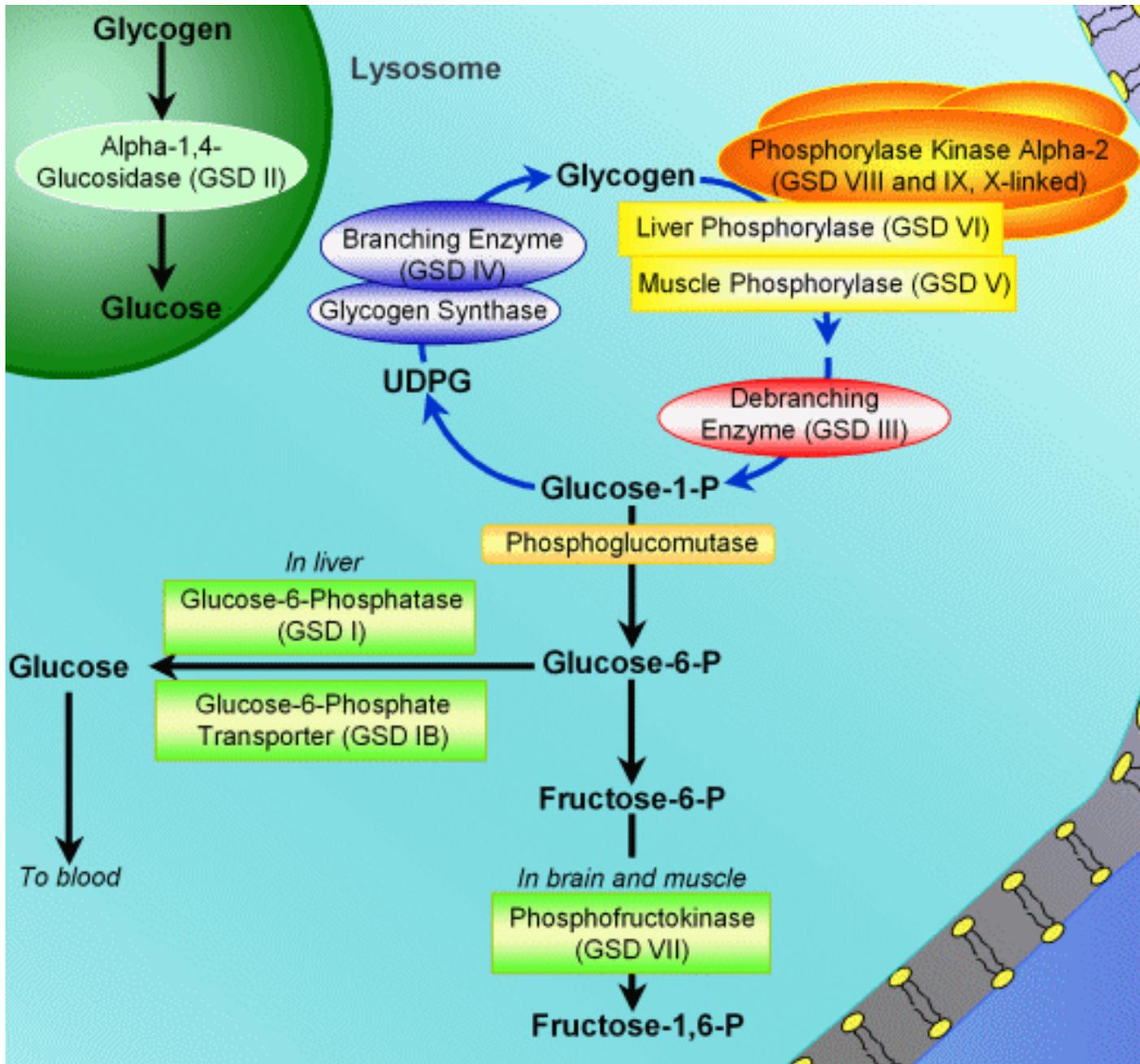


Figure 15-8 Fundamentals of Biochemistry, 2/e
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A. UDP-glucose pyrophosphorylase

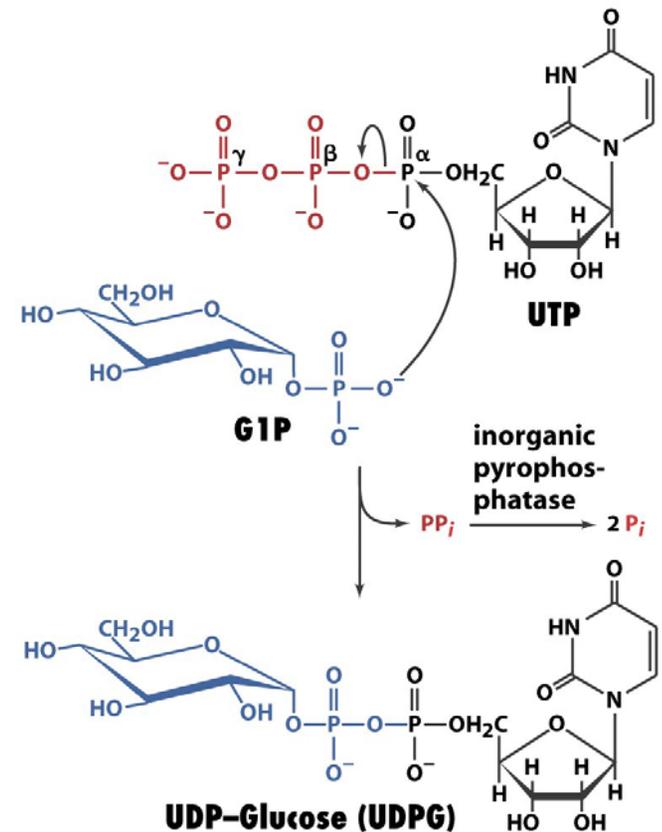


Figure 15-9 Fundamentals of Biochemistry, 2/e
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B. Glycogen synthase

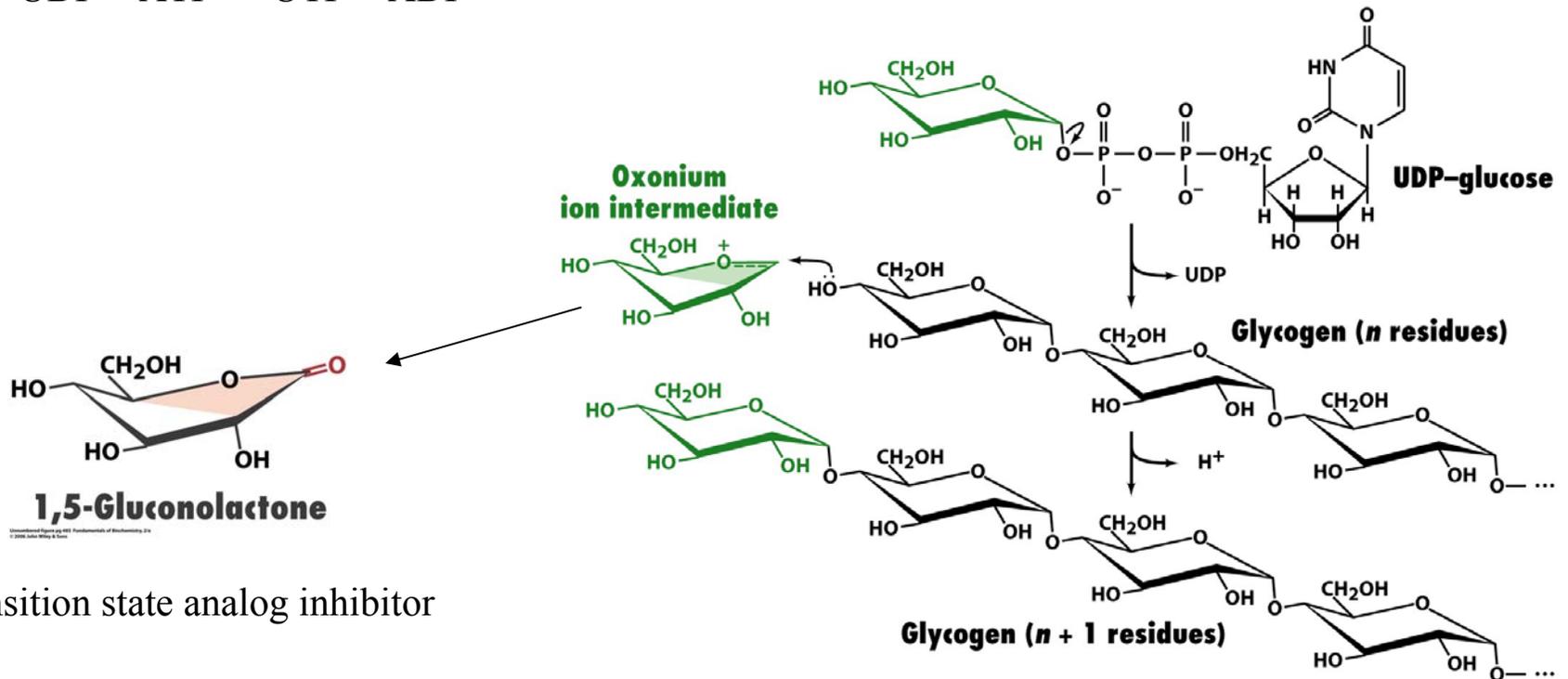
Catalyzes $\alpha(1-4)$ linkage to the **existing** glucan chain

UDPG + glycogen (n) \rightarrow UDP + glycogen (n+1)

G1P + UTP \rightarrow UDPG + 2 Pi

Glycogen (n) + G1P + UTP \rightarrow glycogen (n+1) + UDP + 2 Pi

UDP + ATP \leftrightarrow UTP + ADP



*Initiation of glycogen synthesis

Glycogenin: priming glycogen synthesis

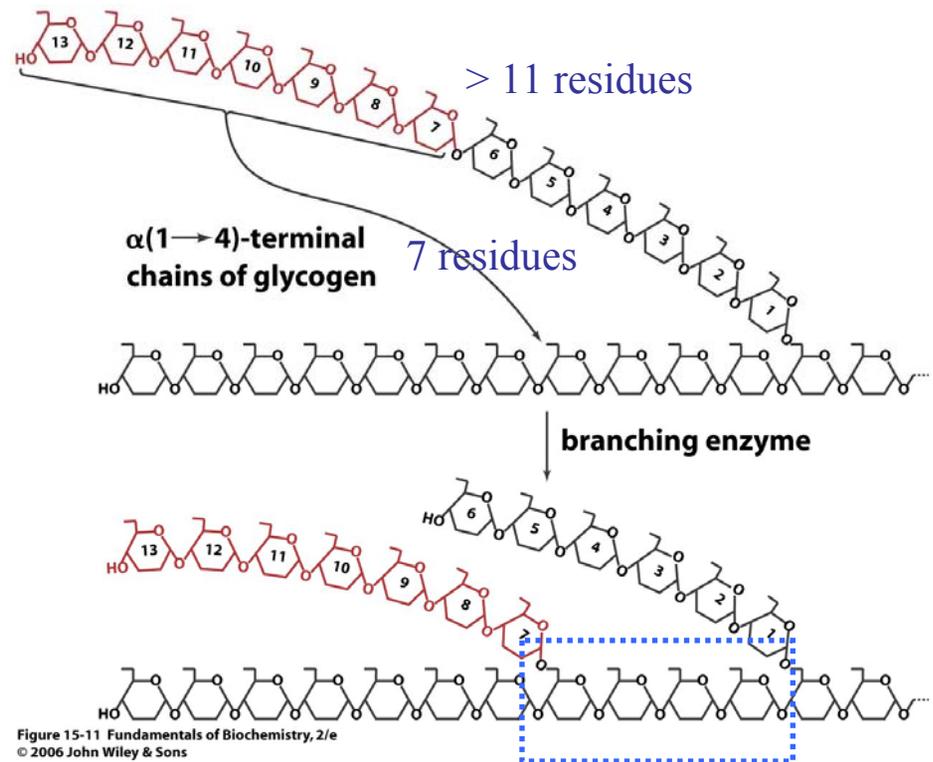
Glycosyltransferase

Attaches a glucose of UDPG to the Tyr194 and extends up to 7 additional residues

Glycogen synthase takes over the role

C. Glycogen branching enzyme

Amylo-(1,4 → 1,6)-transglycosylase

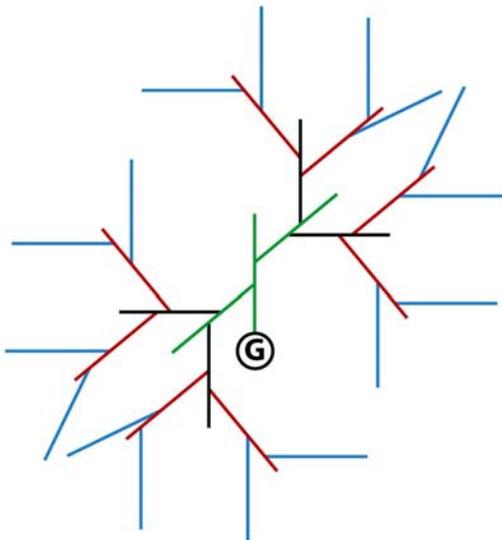


Optimizing glycogen structure

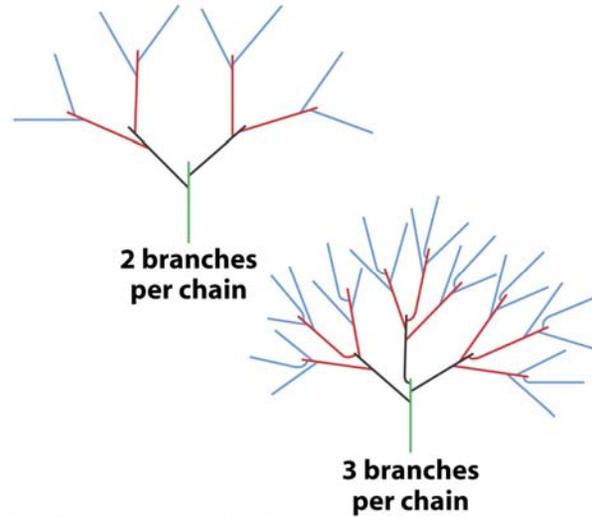
Branching & interval

The largest amount of glucose in the smallest possible volume

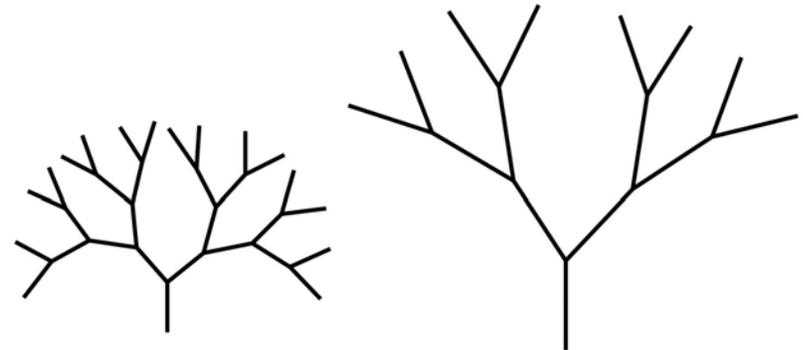
Branching frequency and length



Box 15-3 figure 1 Fundamentals of Biochemistry, 2/e
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Box 15-3 figure 2 Fundamentals of Biochemistry, 2/e
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Box 15-3 figure 3 Fundamentals of Biochemistry, 2/e
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Control of glycogen metabolism

Allosteric & covalent modification

A. Allosteric control of phosphorylase & synthase

Independent control of v_f and v_r

Effectors: ATP, G6P, AMP

Glycogen breakdown:

High demand for ATP: low [ATP] & [G6P], high [AMP]

Phosphorylase is stimulated

Synthase is inhibited

Glycogen synthesis: When [ATP] & [G6P] are high

B. Covalent modification

Hormonal control

A set of kinases and phosphatases

Reverse regulation of phosphorylase and synthase

Synthase a: dephosphorylated form
more active

Synthase b: phosphorylated form
less active
activated by G6P

Covalent modification of glycogen phosphorylase

Phosphorylase a: phosphorylated, active (even without AMP stimulation)

Phosphorylase b: dephosphorylated, inactive

3 involved enzymes

Phosphorylase kinase

Protein kinase A (PKA)

Phosphoprotein phosphatase-1

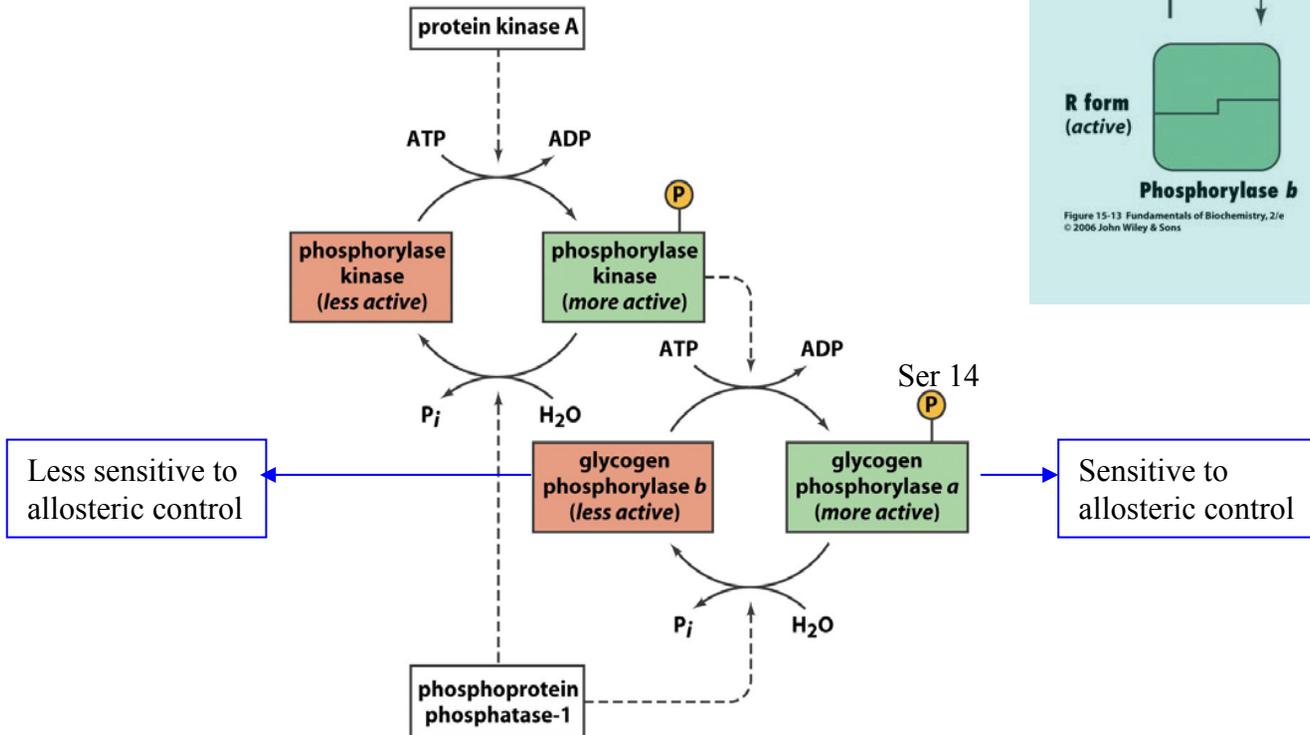


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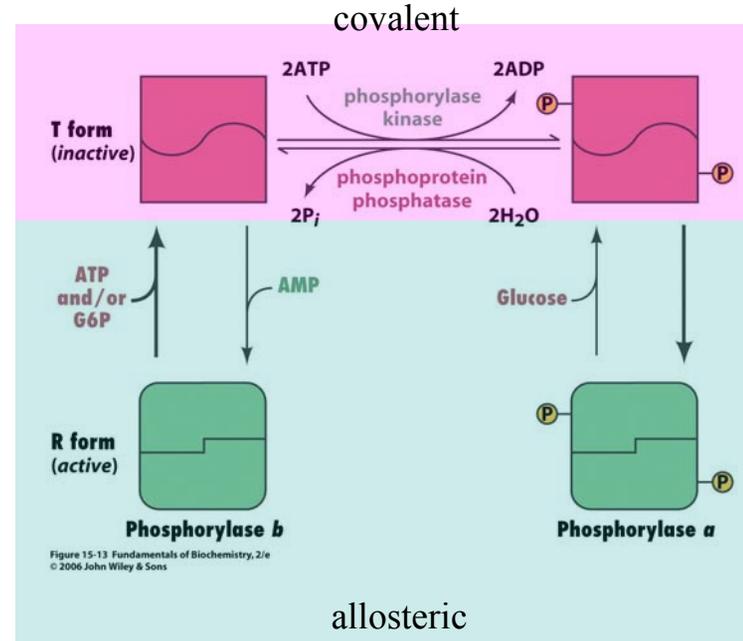
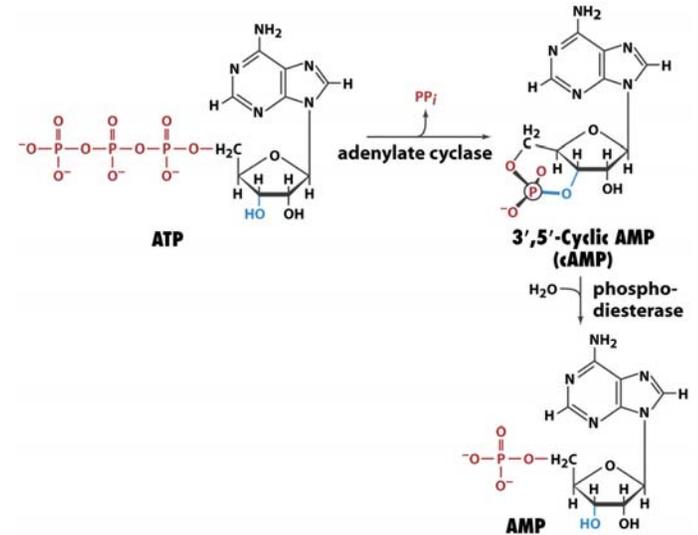


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PKA (protein kinase A)

cAMP-dependent protein kinase (cAPK)

Ser/Thr of numerous cellular proteins



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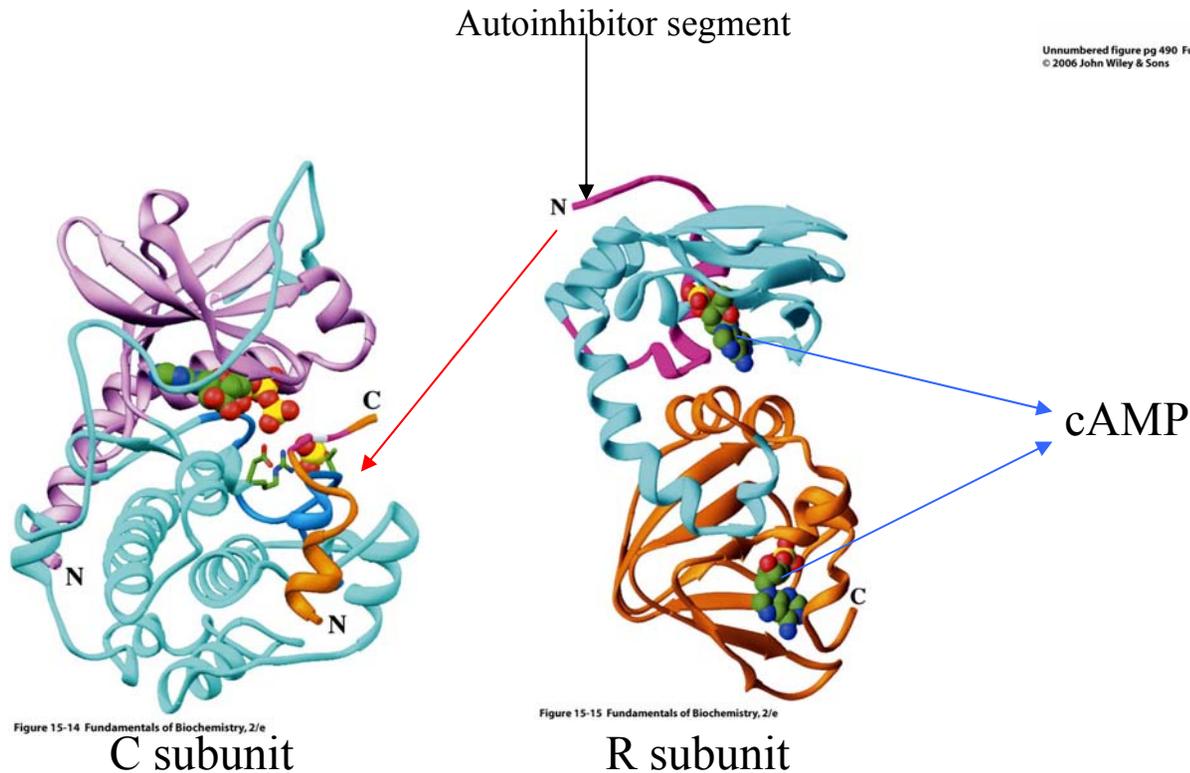


Figure 15-14 Fundamentals of Biochemistry, 2/e

Figure 15-15 Fundamentals of Biochemistry, 2/e

Phosphorylase kinase

4 subunits: $\alpha\beta\gamma\delta$ (γ , catalytic subunit; $\alpha\beta\delta$, regulatory subunits)

Catalytic subunit structure is similar to the C subunit of PKA (autoinhibition)

Activation of the catalytic subunit

Phosphorylation of $\alpha\beta$

Ca^{++} binding to δ (calmodulin)

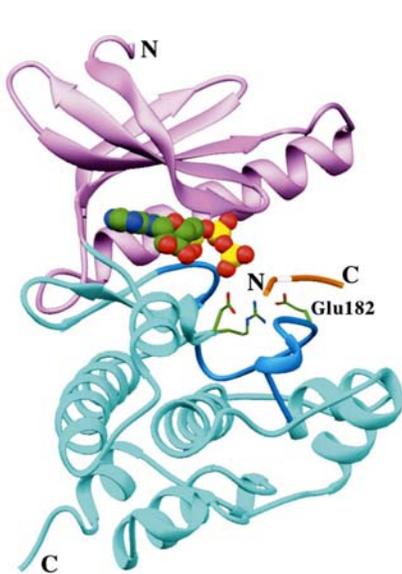


Figure 15-16 Fundamentals of Biochemistry, 2/e

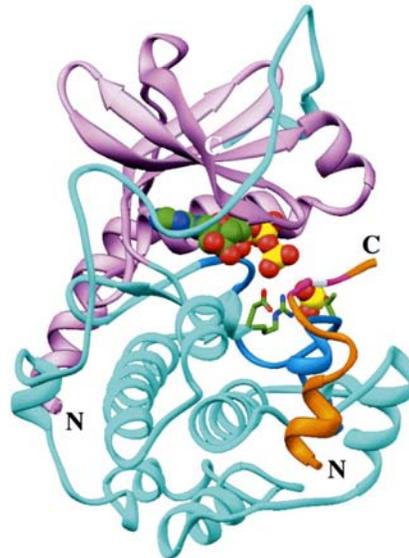


Figure 15-14 Fundamentals of Biochemistry, 2/e

Calmodulin (CaM)

Ca²⁺-binding protein

Highly conserved in eukaryotes

Helix-loop-helix motif: EF hand

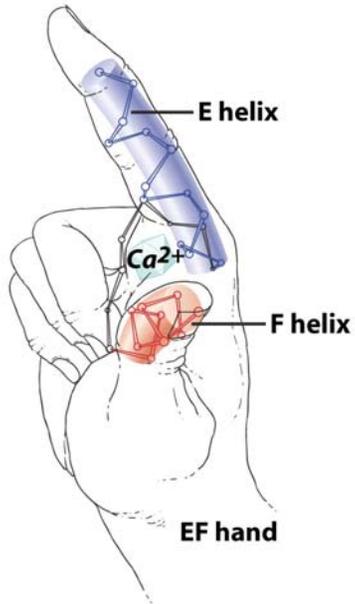


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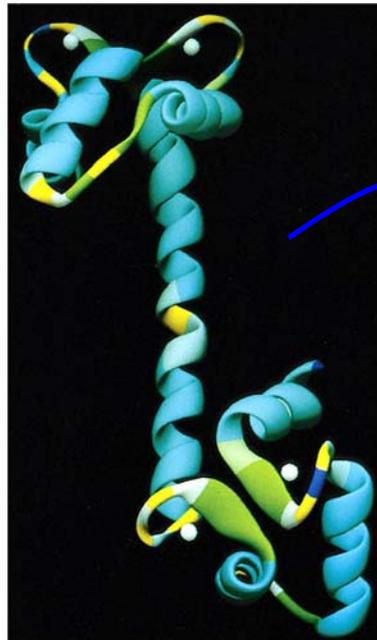


Figure 15-17 Fundamentals of Biochemistry, 2/e

Conformational change
by calcium binding

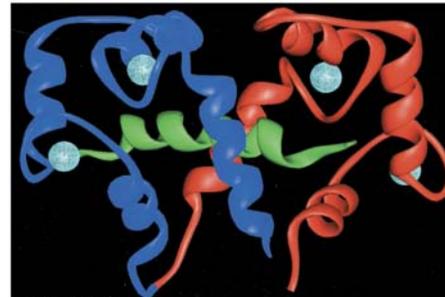


Figure 15-16a Fundamentals of Biochemistry, 2/e



Target polypeptide

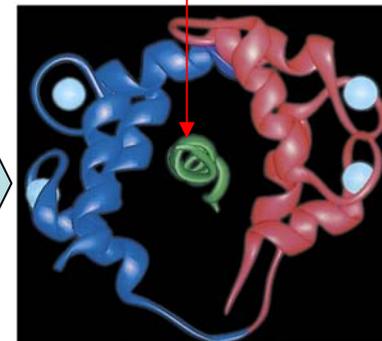


Figure 15-16b Fundamentals of Biochemistry, 2/e

Phosphoprotein phosphatase-1

Control in muscle

Catalytic subunit (PP1c) + glycogen binding subunit (G_M subunit)

Active only when bound to glycogen: regulated by the phosphorylation of G_M subunit

Further regulation by phosphoprotein phosphatase inhibitor 1 (inhibitor-1)

Dual effect of cAMP

PKA: activate phosphorylase, deactivate inhibitor-1

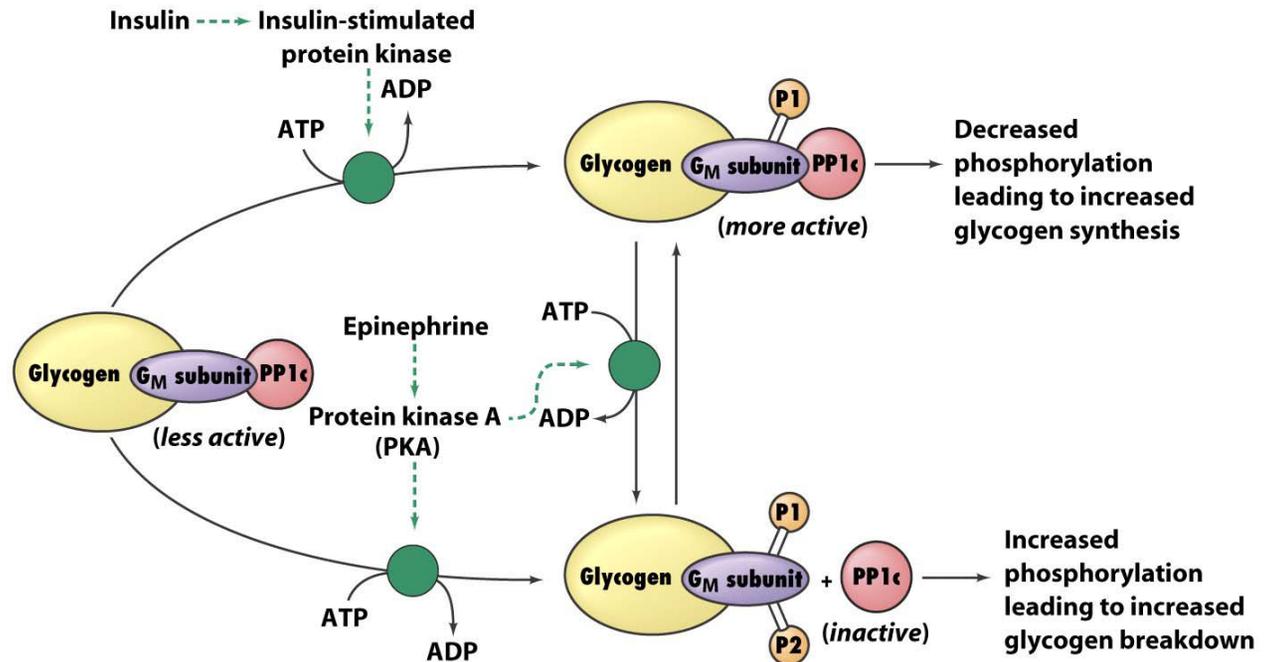
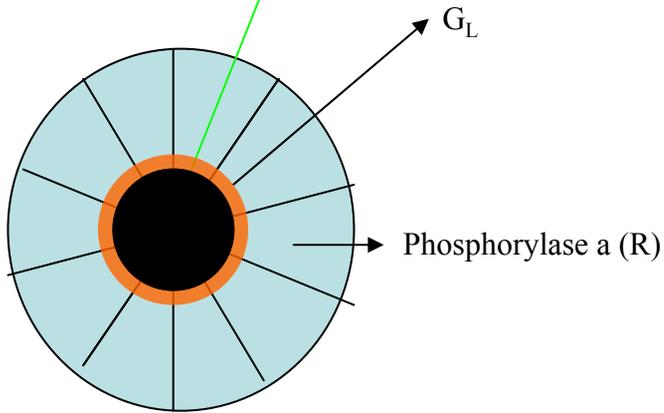


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Phosphoprotein phosphatase-1

Control in liver

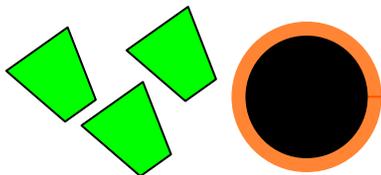


High glucose

Phosphorylase a (T): exposed Ser 14

Dephosphorylation

Phosphorylase b



Phosphorylase a is a glucose sensor in liver

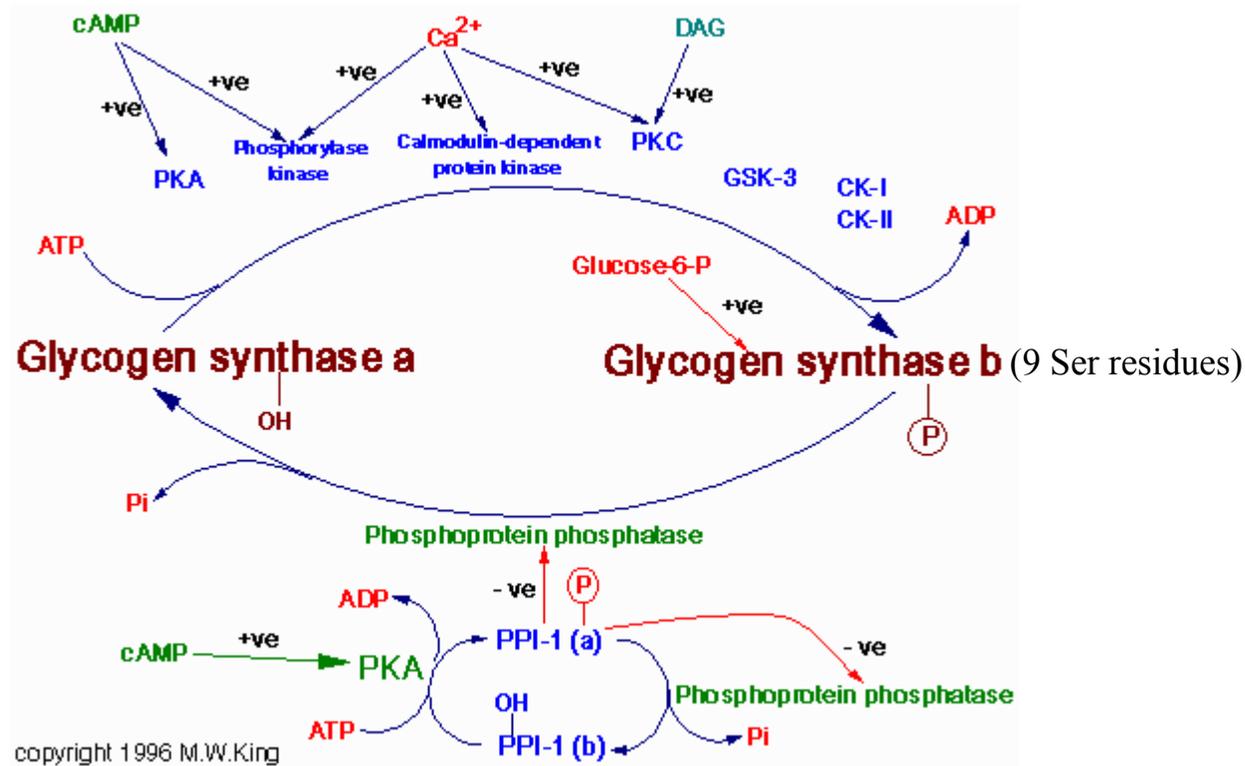
Phosphoprotein phosphatase-1 is bound to glycogen through glycogen binding subunit (G_L)

G_L is not subject to control via phosphorylation

Controlled by binding to phosphorylase a

간에서는 phosphorylase a에 결합된 상태로 조절된다 (보통 10개의 phosphorylase a에 1개꼴로 존재). T와 R 모두에 결합할 수 있다. 하지만 R form의 경우는 Ser14 잔기가 감추어져 있기 때문에 탈인산화가 일어나지 않는다. glucose의 농도가 증가하면서 T form으로 전환되면 Ser14가 노출되면서 탈인산화가 일어나 phosphorylase b로 전환된다. 이것은 phosphatase-1와의 친화력이 떨어지기 때문에 phosphatase-1은 떨어져 나온다. 그렇지만 phosphatase-1은 보통 10개의 phosphorylase a에 1개꼴로 존재하기 때문에 phosphorylase의 90% 이상이 phosphorylase b로 바뀌기 전에는 완전히 떨어지지 않는다. 따라서 glycogen synthase를 활성화시키지 못한다. 이것은 phosphorylase와 glycogen synthase가 동시에 활성화되는 것을 억제한다.

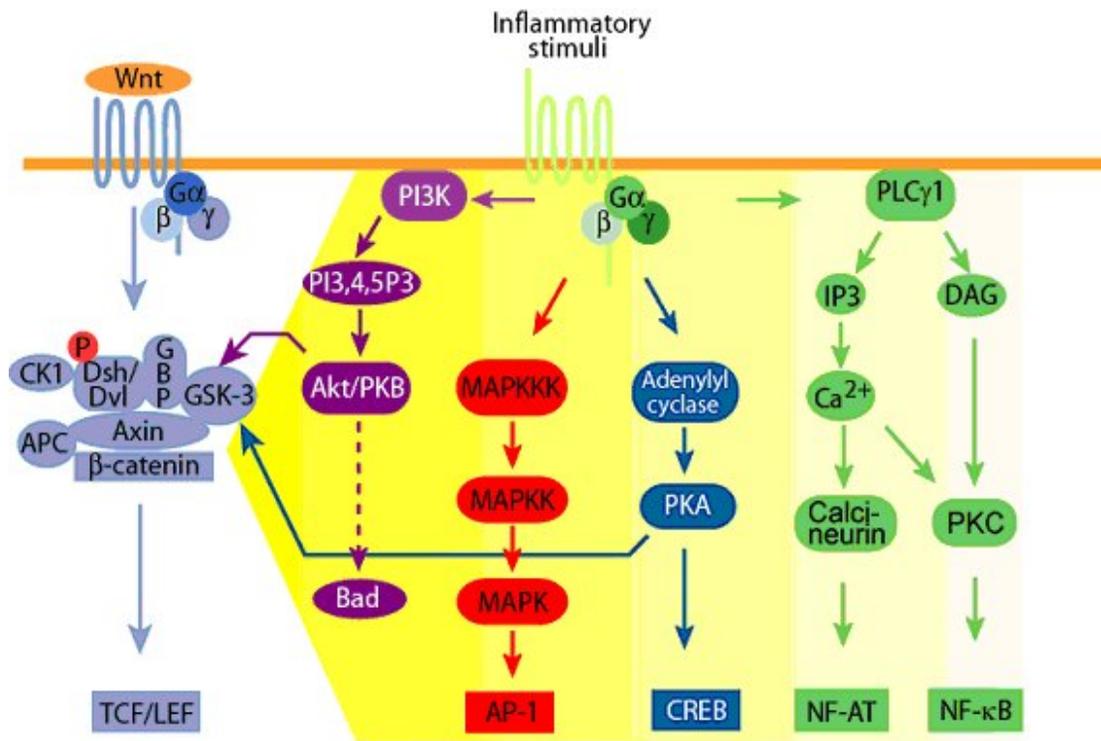
Glycogen synthase



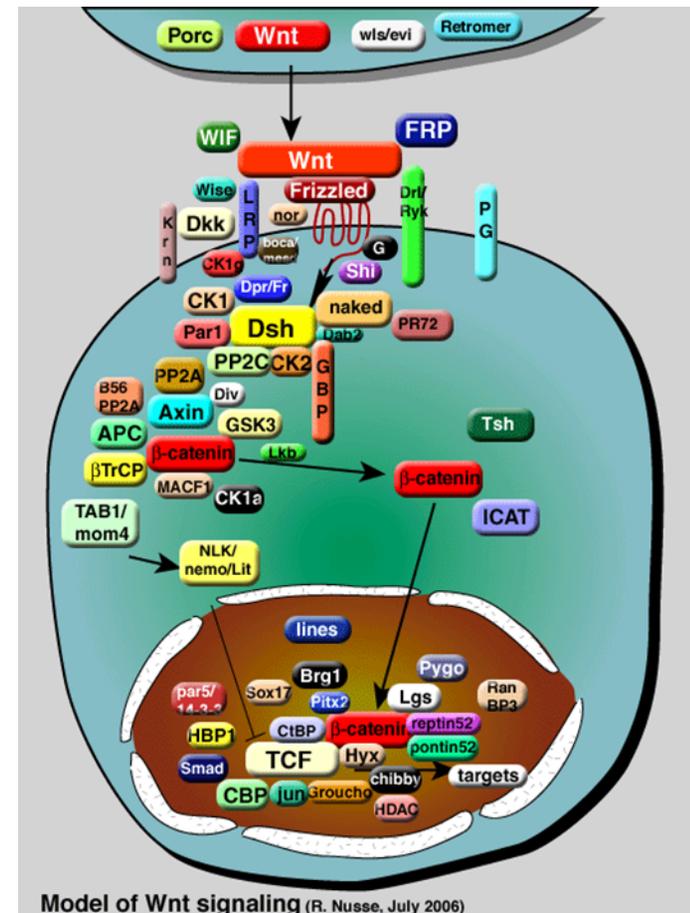
Pathways involved in the regulation of glycogen synthase. See the text for details of the regulatory mechanisms. PKA is cAMP-dependent protein kinase. PPI-1 is phosphoprotein phosphatase-1 inhibitor. Whether a factor has positive (+ve) or negative (-ve) effects on any enzyme is indicated. Briefly, glycogen synthase a is phosphorylated, and rendered much less active and requires glucose-6-phosphate to have any activity at all. Phosphorylation of glycogen synthase is accomplished by several different enzymes. The most important is synthase-phosphorylase kinase the same enzyme responsible for phosphorylation (and activation) of glycogen phosphorylase. PKA (itself activated through receptor mediated mechanisms) also phosphorylates glycogen synthase directly. The effects of PKA on PPI-1 are the same as those described above for the regulation of glycogen phosphorylase. The other enzymes shown to directly phosphorylate glycogen synthase are protein kinase C (PKC), calmodulin-dependent protein kinase, glycogen synthase kinase-3 (GSK-3) and two forms of casein kinase (CK-I and CK-II). The enzyme PKC is activated by Ca²⁺ ions and phospholipids, primarily diacylglycerol, DAG. DAG is formed by receptor-mediated hydrolysis of membrane phosphatidylinositol bisphosphate (PIP₂). (web.indstate.edu/thcme/mwking/glycogen.html)

GSK3 & Wnt signalling

Wnt proteins form a family of highly conserved secreted signaling molecules that regulate cell-to-cell interactions during embryogenesis. Wnt genes and Wnt signaling are also implicated in cancer.



Proliferation Differentiation Cell survival



Model of Wnt signaling (R. Nusse, July 2006)

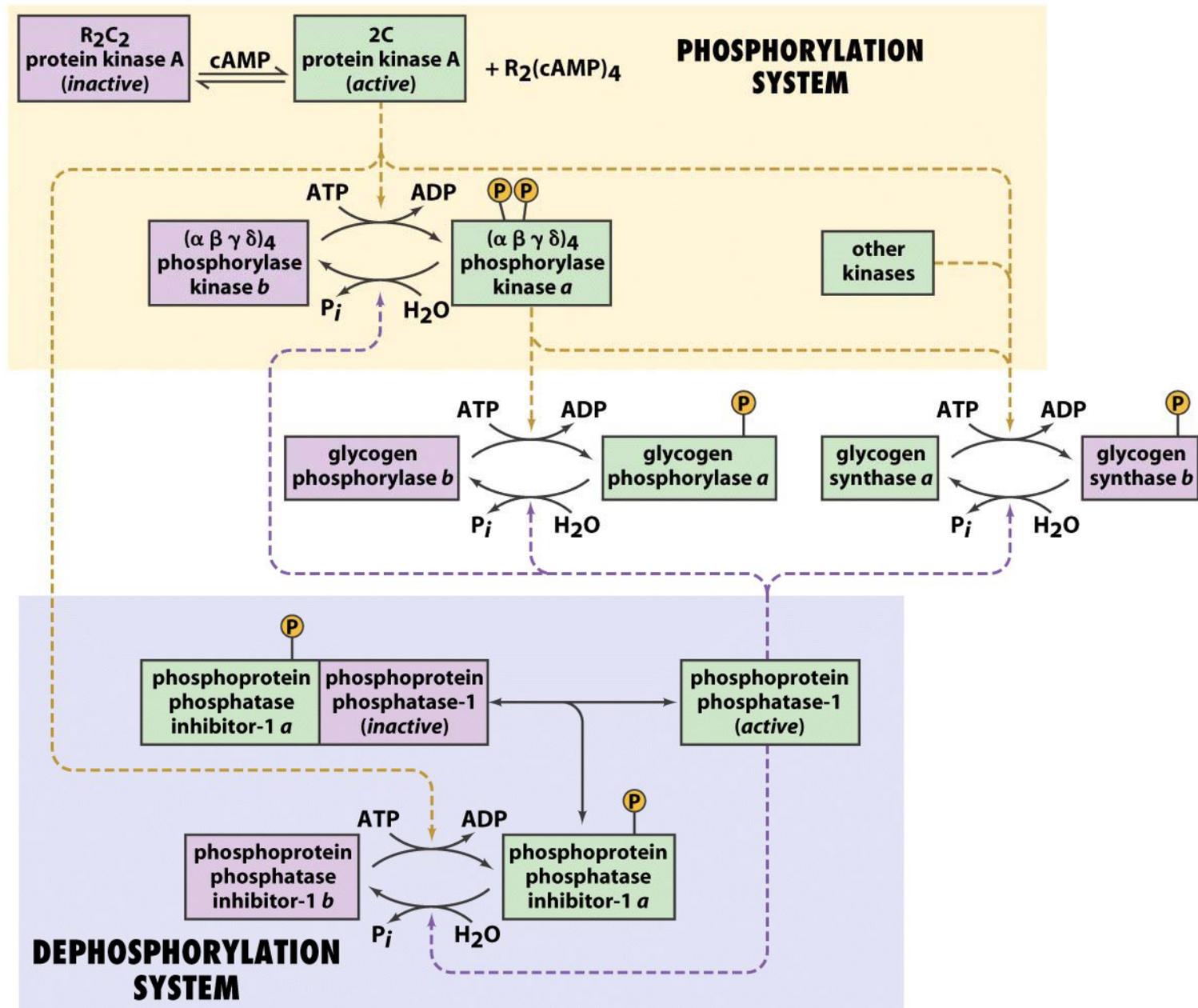
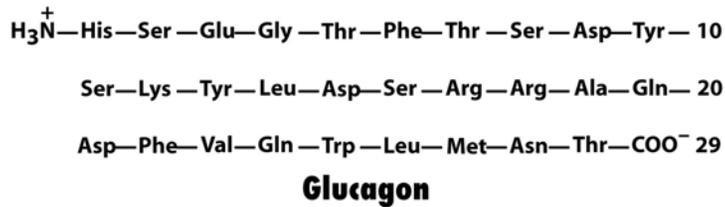


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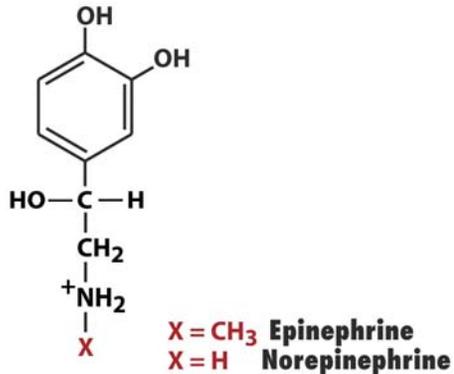
Hormonal effects on glycogen metabolism

Insulin & glucagon: control blood glucose

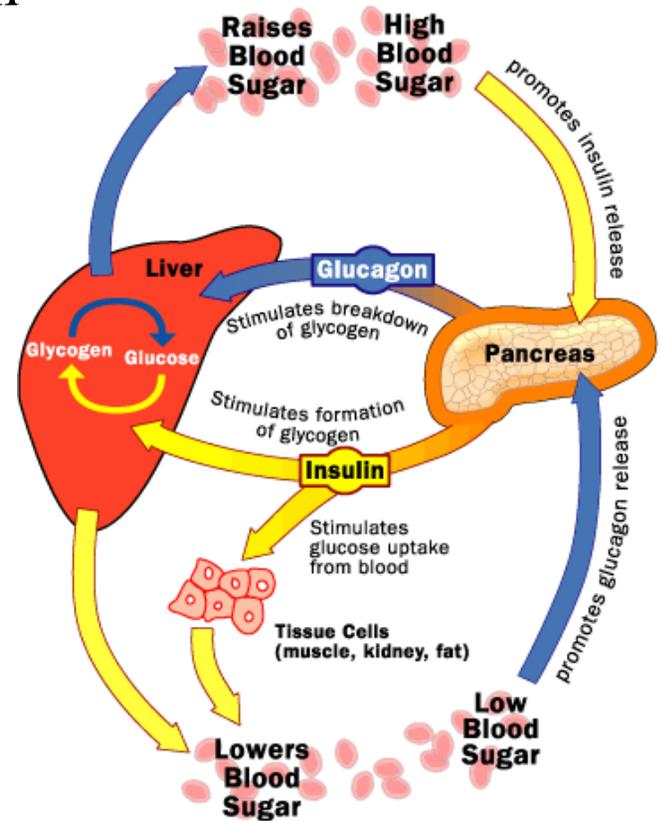
Epinephrine & norepinephrine (in response to stress)



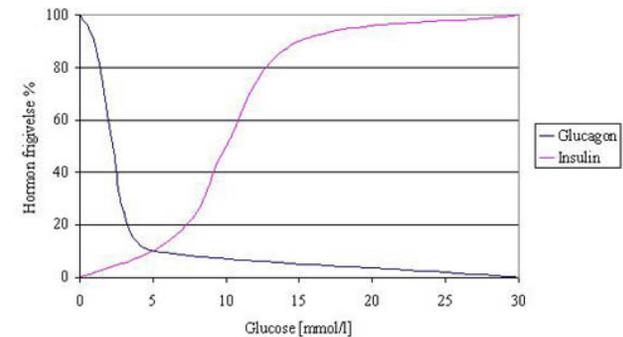
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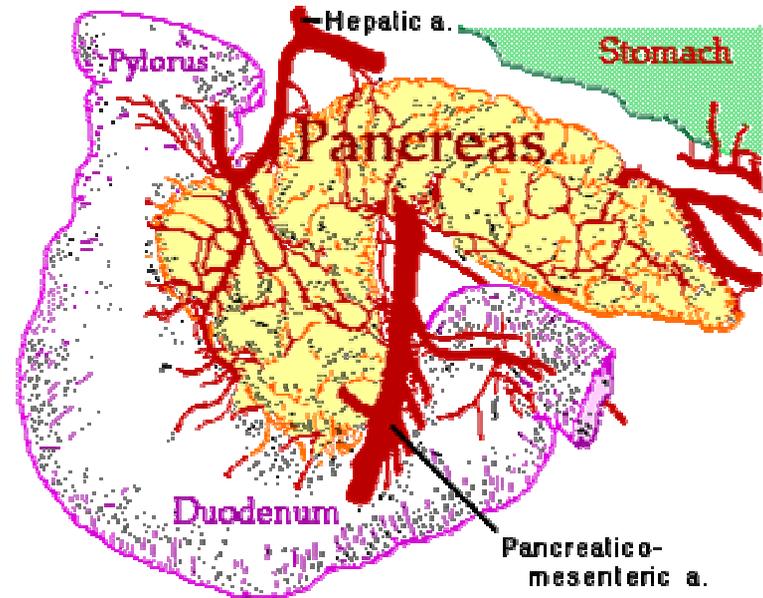
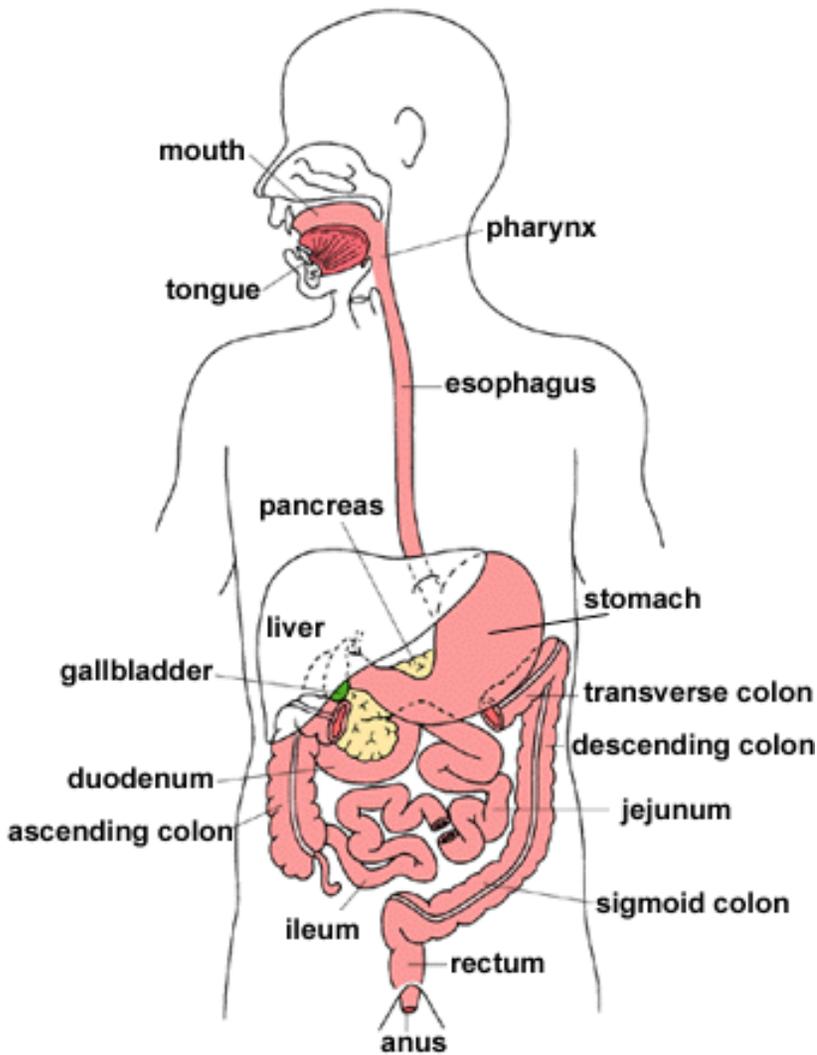


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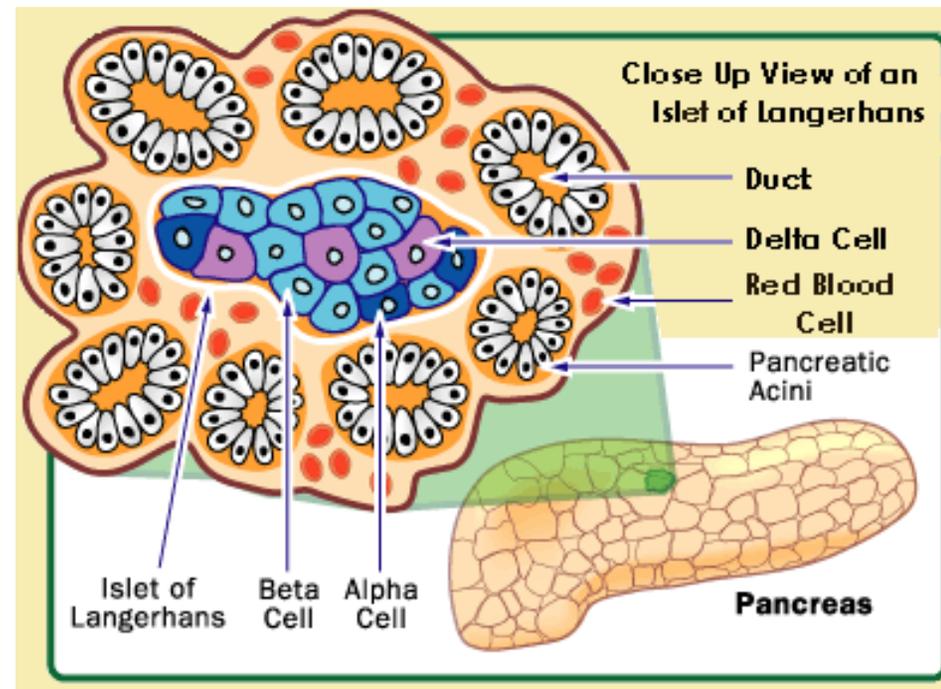


Insulin kontra glucagon





α -cells: glucagon
 β -cells: insulin



Stimulate pancreas to secrete glucagon

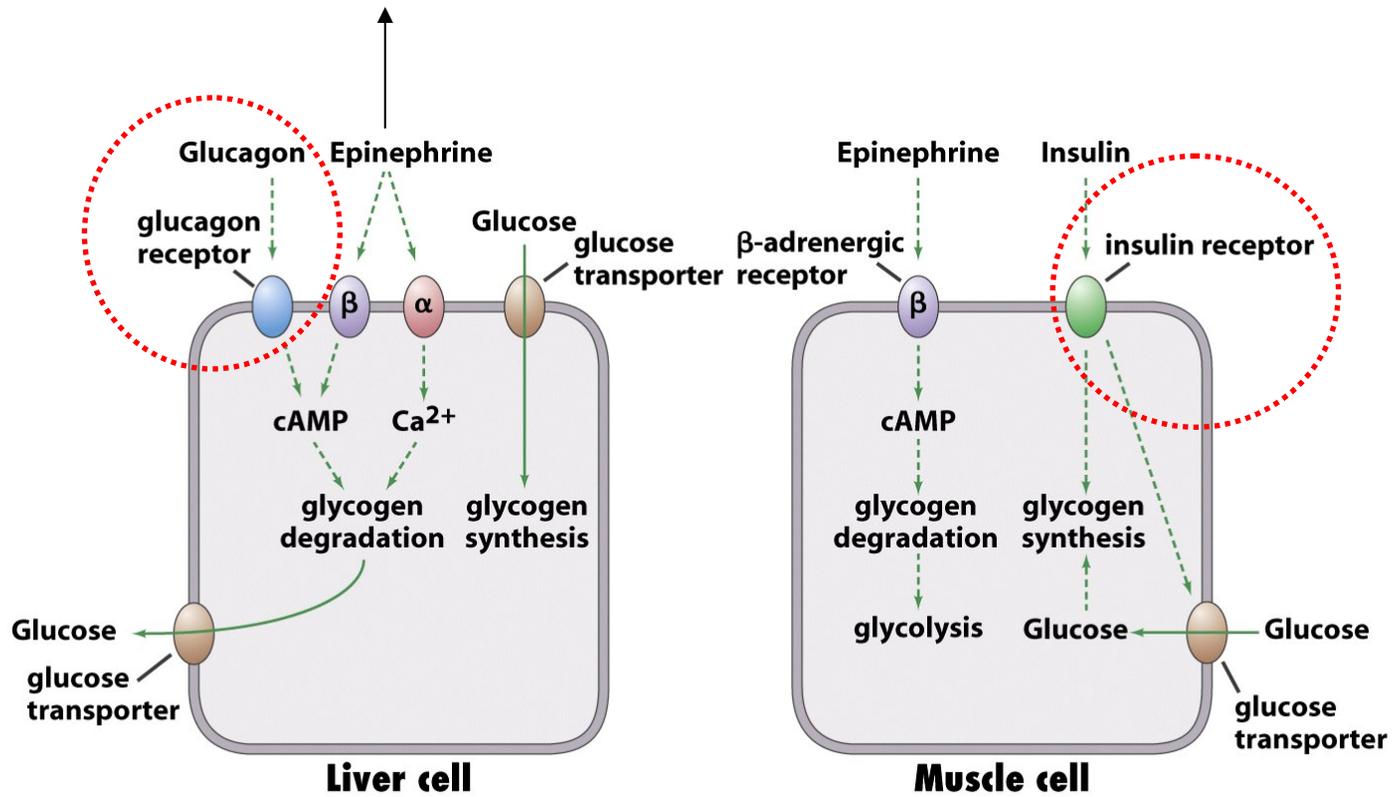
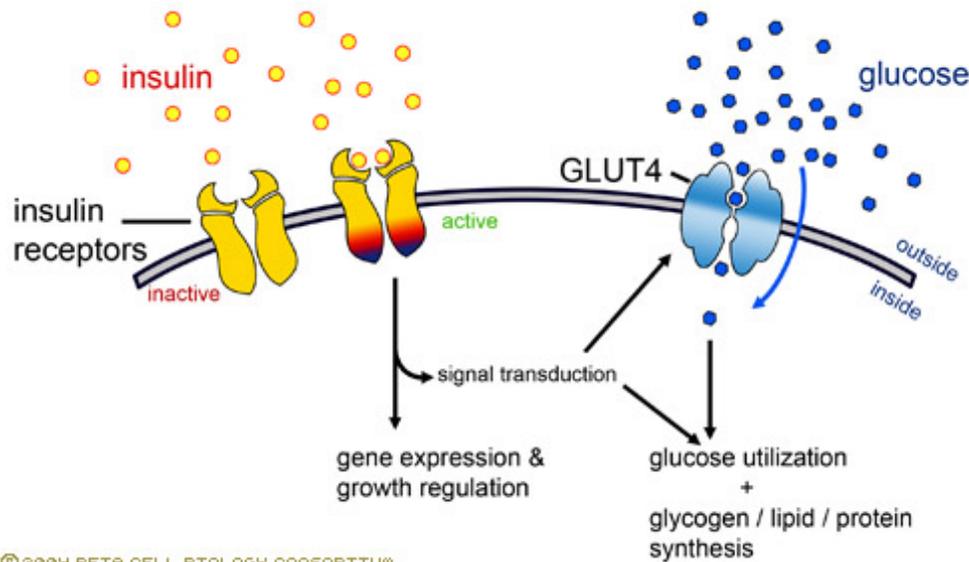


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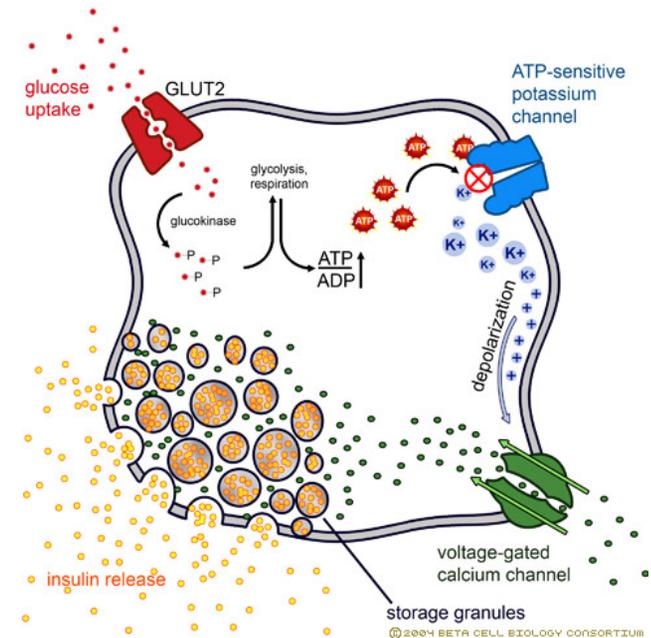
Adrenergic receptors: β -adrenergic (cAMP), α -adrenergic (calcium ion)

Insulin receptors:

- The receptors for insulin are found on most mammalian cells – action of insulin is mediated through these receptors.
- Impaired action of insulin can result from defects in the receptors or defects in post-receptor events.



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Signal transduction

