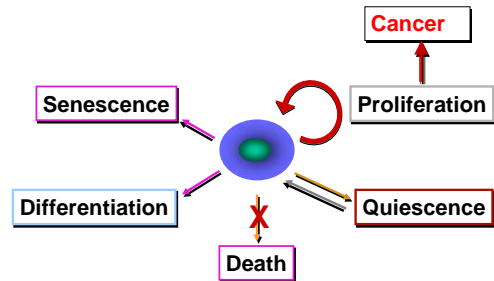


细胞信号转导 (Cell Signaling)

陈晔光
清华大学生命科学学院
ygchen@tsinghua.edu.cn

1

Fates of a cell



2

General introduction

Common features of signal transduction
Cell surface signal transducers, receptors

Ion channels

Secondary messengers

cAMP

cGMP

Lipids

Calcium

G proteins

Trimeric G proteins

Monomeric G proteins, Tyr kinase/MAP kinase

Protein modules

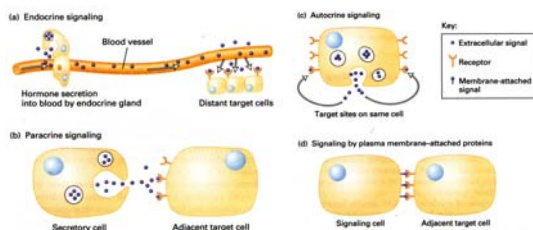
3

What is "Signal Transduction"?

This expression first made its mark in the biological literature around 1974. Physical scientists and electronic engineers had earlier used the term to describe the conversion of energy or information from one form into another. Signal transduction at the cellular level refers to the movement of signals from outside the cell to inside; cascade of information from the plasma membrane to the nucleus in response to an extracellular stimulus in living organisms.

4

General schemes of intercellular signaling

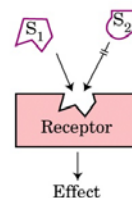


5

Four basic features of signal-transducing systems

(a) Specificity

Signal molecule fits binding site on its complementary receptor; other signals do not fit.

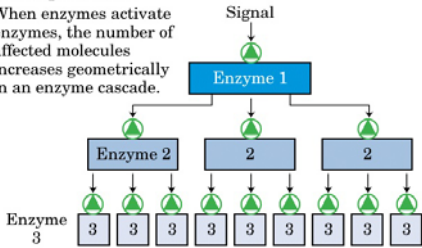


6

Four basic features of signal-transducing systems

(b) Amplification

When enzymes activate enzymes, the number of affected molecules increases geometrically in an enzyme cascade.

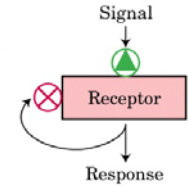


7

Four basic features of signal-transducing systems

(c) Desensitization/Adaptation

Receptor activation triggers a feedback circuit that shuts off the receptor or removes it from the cell surface.

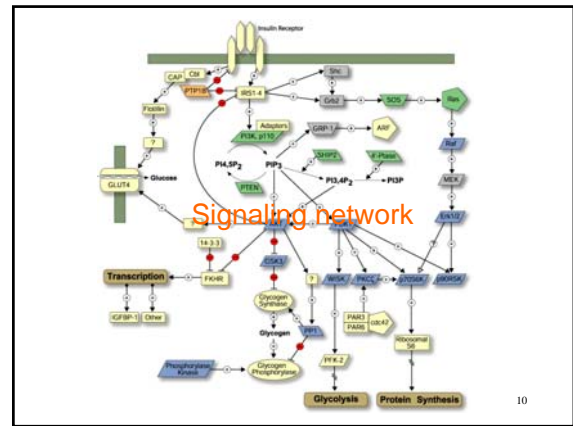
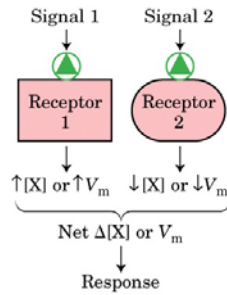


8

Four basic features of signal-transducing systems

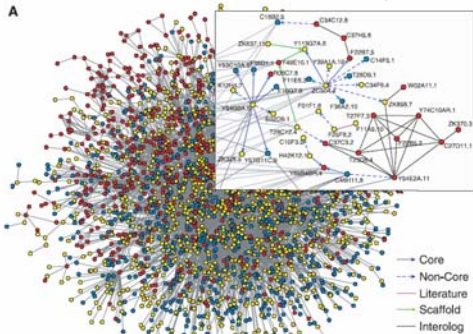
(d) Integration

When two signals have opposite effects on a metabolic characteristic such as the concentration of a second messenger X, or the membrane potential V_m , the regulatory outcome results from the integrated input from both receptors.



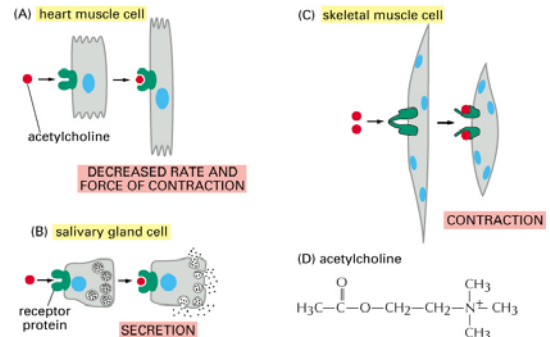
10

Protein interactome network in *C. elegans*

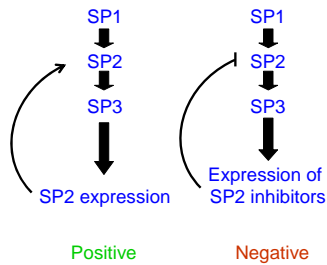


Li, et al., 2004, Science, 303:540

Context-specificity of cell signaling: Different cells respond differently to the same extracellular signal

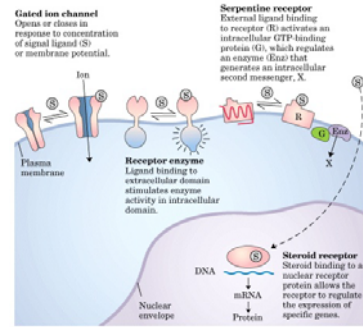


Positive and negative feedback mechanisms



13

Four general types of signal transducers



14

Ligand-activated cell-surface receptors

- Ion-channel receptors: acetylcholine
- G-protein-coupled receptors: epinephrine, glucagon, serotonin
- Tyrosine kinase-linked receptors: interferons
- Tyrosine kinase receptors: EGF, PDGF, insulin
- Tyrosine phosphatase receptors: CD45
- Serine/threonine kinase receptors: TGF β , BMP
- Guanylate cyclase receptor: atrial natriuretic factor

15

General introduction

- Common features of signal transduction
- Cell surface signal transducers, receptors

Ion channels

Secondary messengers

- cAMP
- cGMP
- Lipids
- Calcium

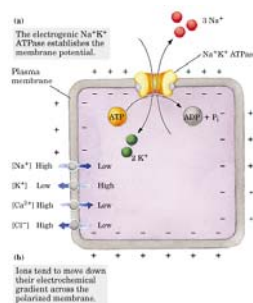
G proteins

- Trimeric G proteins
- Monomeric G proteins, Tyr kinase/MAP kinase

Protein modules

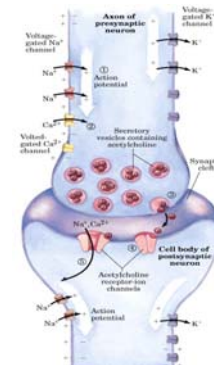
16

Transmembrane electrical potential



17

Role of voltage-gated and ligand-gated ion channels in neural transmission

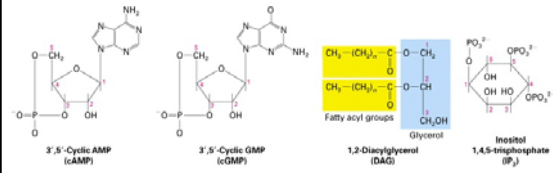


18

General introduction
 Common features of signal transduction
 Cell surface signal transducers, receptors
 Ion channels
 Secondary messengers
 cAMP
 cGMP
 Lipids
 Calcium
 G proteins
 Trimeric G proteins
 Monomeric G proteins, Tyr kinase/MAP kinase
 Protein modules

19

Second messengers

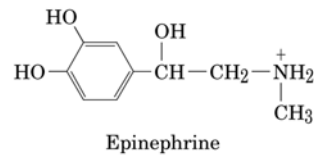


20

Secondary messengers:

cAMP
 cGMP
 Lipids
 Calcium

21

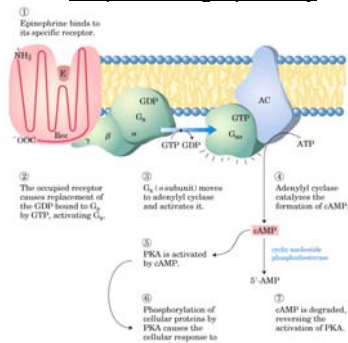


Epinephrine

Produced by Chromaffin cells in adrenal medulla
 Functions:
 Increase heart rate, blood pressure,
 sweating, rate of respiration
 Stimulate conversion of glycogen to glucose

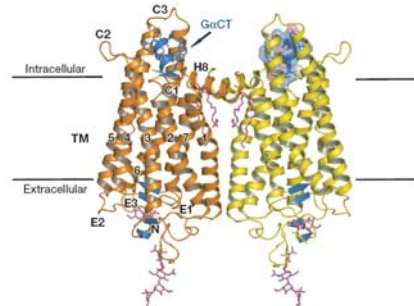
22

Transduction of the epinephrine signal:
 the β -adrenergic pathway



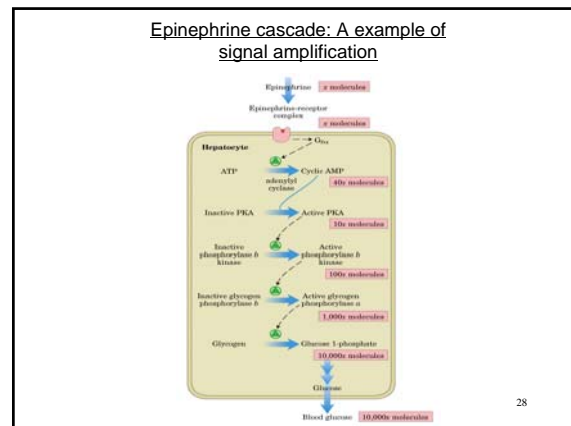
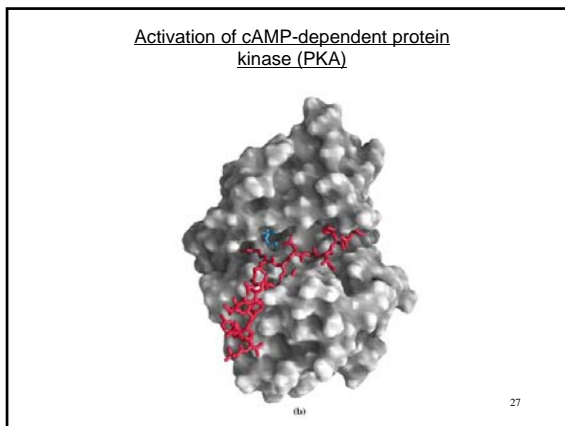
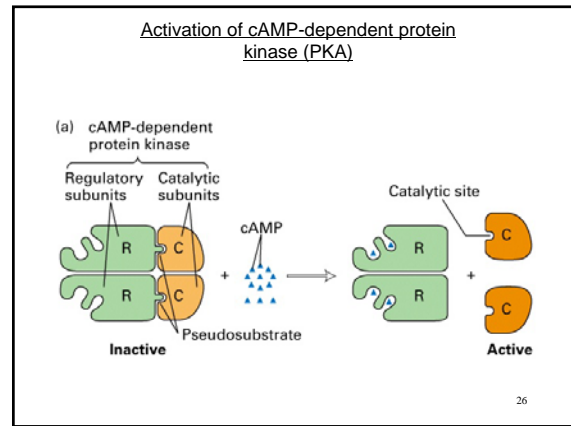
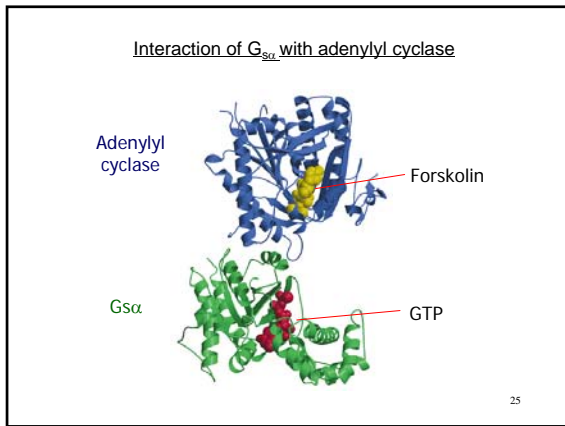
23

Structure of Opsin associated with the C-terminal Gs α



Scheerer et al. Nature, 2008

24



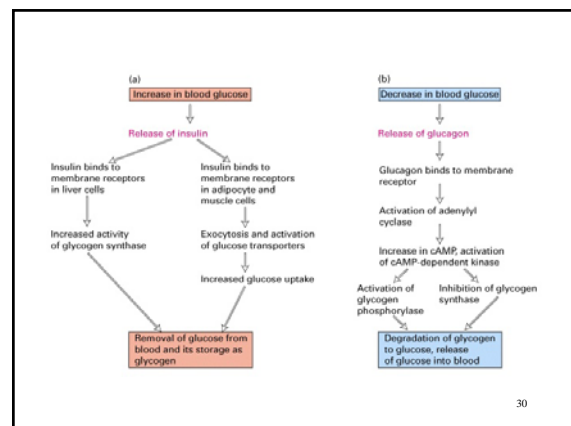
cAMP as a secondary messenger

TABLE 20-3 Metabolic Responses to Hormone-Induced Rise in cAMP in Various Tissues

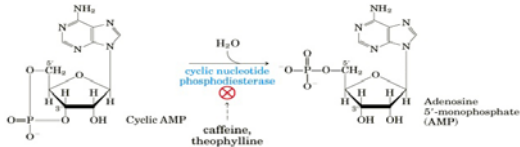
Tissue	Hormone Inducing Rise in cAMP	Metabolic Response
Adipose	Epinephrine; ACTH; glucagon	Increase in hydrolysis of triglyceride; decrease in amino acid uptake
Liver	Epinephrine; norepinephrine; glucagon	Increase in conversion of glycogen to glucose; inhibition of synthesis of glycogen; increase in amino acid uptake; increase in gluconeogenesis (synthesis of glucose from amino acids)
Ovarian follicle	FSH; LH	Increase in synthesis of estrogen, progesterone
Adrenal cortex	ACTH	Increase in synthesis of aldosterone, cortisol
Cardiac muscle cells	Epinephrine	Increase in contraction rate
Thyroid	TSH	Secretion of thyroxine
Bone cells	Parathyroid hormone	Increase in resorption of calcium from bone
Skeletal muscle	Epinephrine	Conversion of glycogen to glucose
Intestine	Epinephrine	Fluid secretion
Kidney	Vasopressin	Resorption of water
Blood platelets	Prostaglandin I	Inhibition of aggregation and secretion

SOURCE: E. W. Sutherland, 1972, *Science* 177:401.

29



Inactivation of cAMP



31

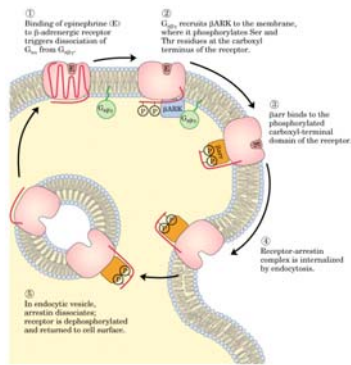
Medicine: β -adrenergic receptor

Cardiac muscle cells possess β_1 receptor, whose activation increases heart rate. Practolol (心得灵), an β_1 -selective antagonist, can slow heart contraction and is used to treat cardiac arrhythmia and angina.

Smooth muscle cells have β_2 receptors, whose activation promotes relaxation. Terbutaline (特布他林), an agonist selective for β_2 , is used in the treatment of asthma.

32

Desensitization of the β -adrenergic receptor



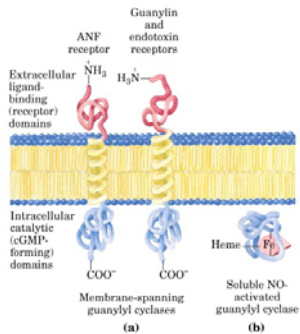
33

Secondary messengers:

- cAMP
- cGMP
- Lipids
- Calcium

34

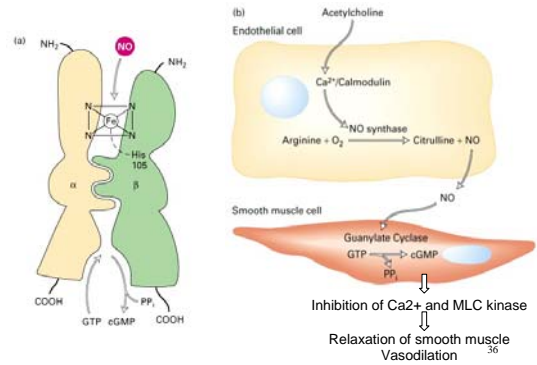
Guanylyl cyclases



Endotoxin, produced by *E. coli* and other gram-negative bacteria in intestine, stimulates cGMP production, increases Cl⁻ secretion and consequently decreases reabsorption of water by the intestinal epithelium, producing diarrhea

35

NO (Nitric Oxide) signaling



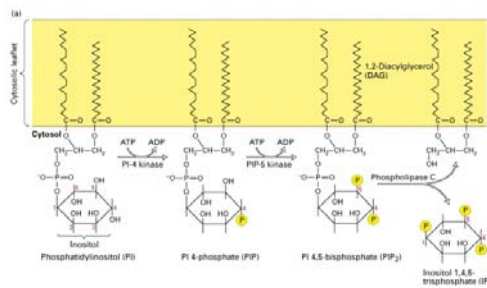
36

Secondary messengers:

- cAMP
- cGMP
- Lipids
- Calcium

37

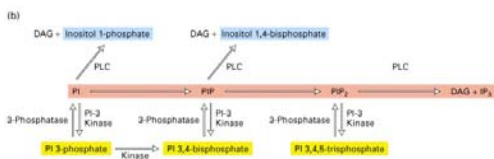
Lipids as messengers



磷脂酰肌醇

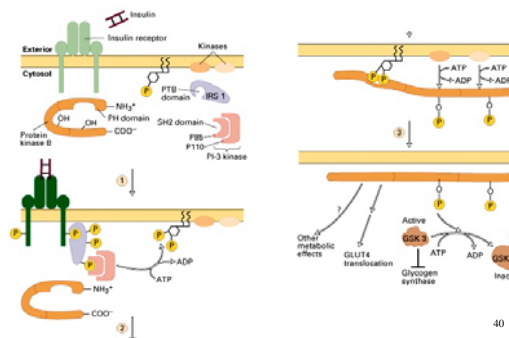
38

Lipid phosphorylation



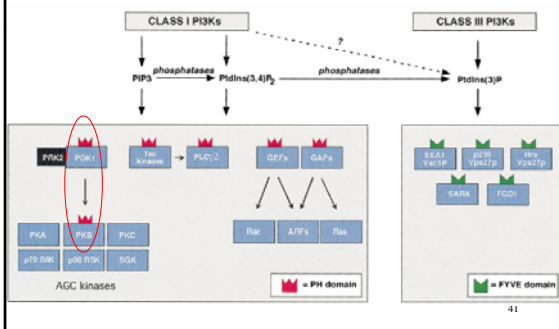
39

Insulin regulates glucose metabolism via PI3K



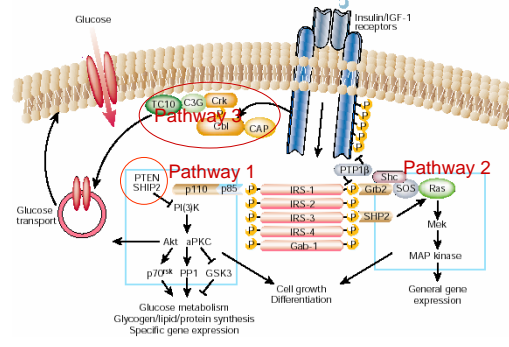
40

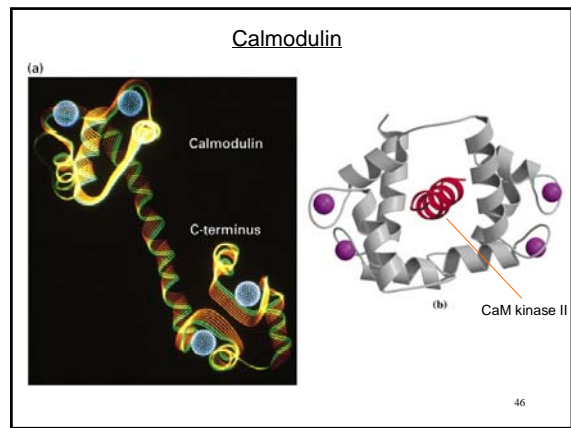
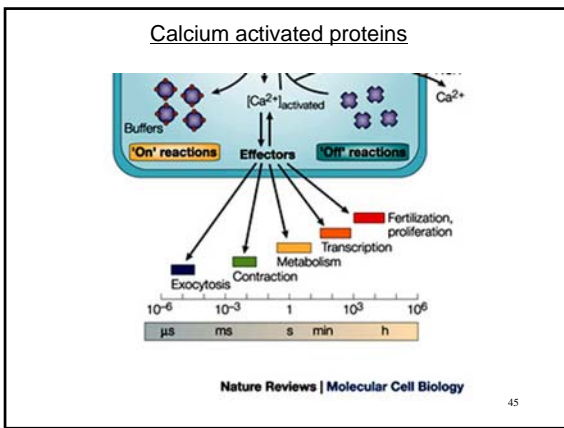
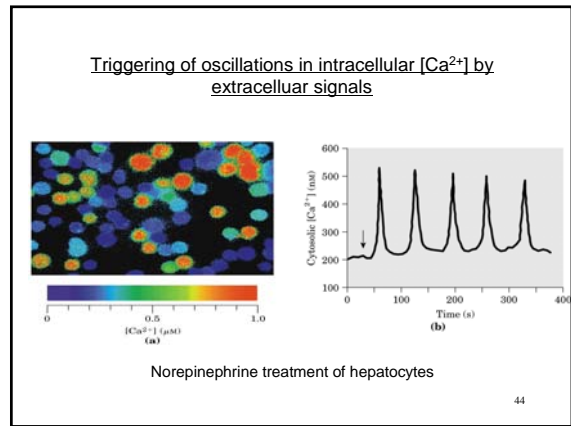
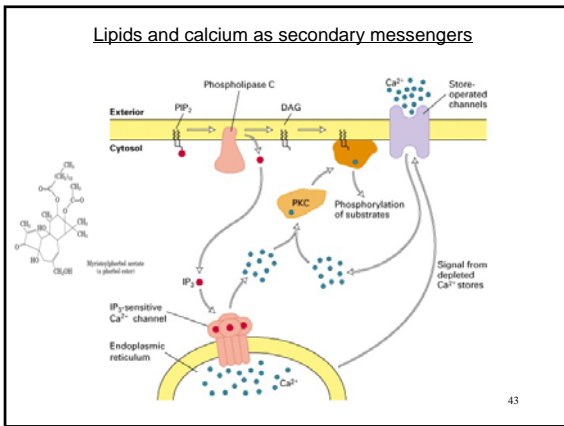
PH domain-containing proteins



41

Glucose metabolism controlled by Insulin signaling





- ### Calcium activated proteins
- Calmodulin: CaM kinase; phosphorylase b kinase (glycogen breakdown)
 - Protein kinase C
 - Calcineurin: protein phosphatase 1B
 - Troponin - muscle contraction
 - Synaptotagmin (neurotransmitter release)
 - Guanylyl cyclase
- 47

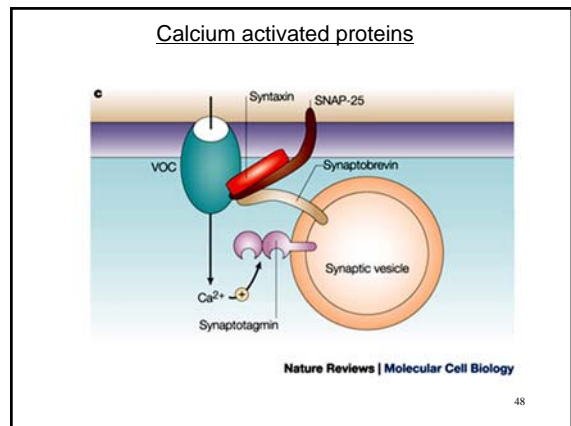


TABLE 20-4 Cellular Responses to Hormone-Induced Rise in Inositol 1,4,5-Triphosphate (IP₃) and Subsequent Rise in Cytosolic Ca²⁺ in Various Tissues

Tissue	Hormone Inducing a Rise in IP ₃	Cellular Response
Pancreas (acinar cells)	Acetylcholine	Secretion of digestive enzymes, such as amylase and trypsinogen
Parotid (salivary gland)	Acetylcholine	Secretion of amylase
Pancreas (β cells of islets)	Acetylcholine	Secretion of insulin
Vascular or stomach smooth muscle	Acetylcholine	Contraction
Liver	Vasopressin	Conversion of glycogen to glucose
Blood platelets	Thrombin	Aggregation, shape change, secretion of hormones
Mast cells	Antigen	Histamine secretion
Fibroblasts	Peptide growth factors, such as bombesin and FGFβ	DNA synthesis, cell division
Sea urchin eggs	Spermatozoa	Rise of fertilization membrane

source: M. J. Berridge, 1987, *Ann. Rev. Biochem.* 56:159; M. J. Berridge and R. F. Irvine, 1984, *Nature* 312:315.

49

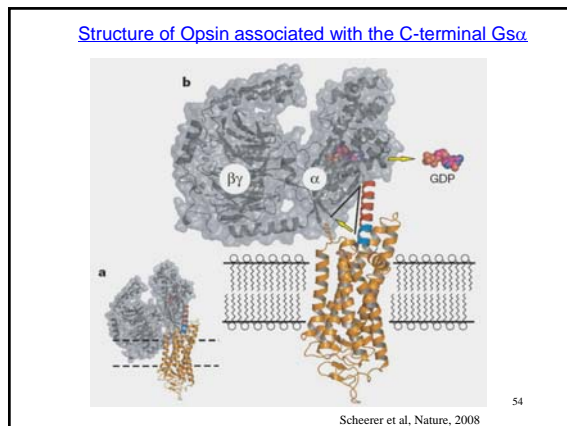
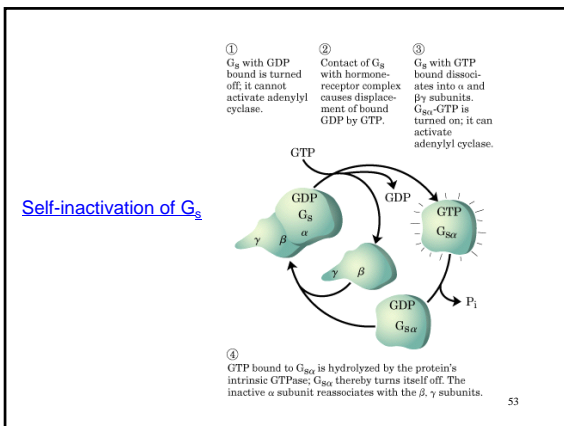
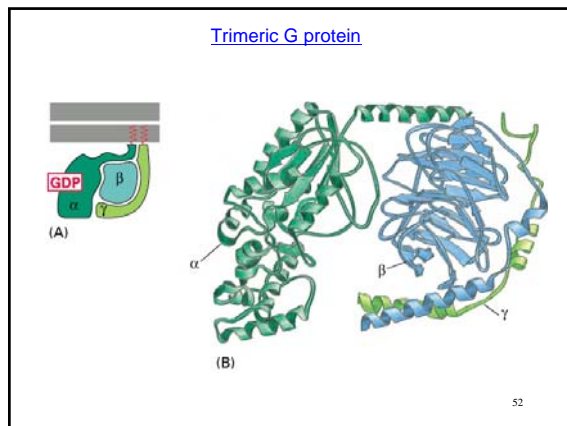
General introduction
 Common features of signal transduction
 Cell surface signal transducers, receptors
 Ion channels
 Secondary messengers
 cAMP
 cGMP
 Lipids
 Calcium
 G proteins
 Trimeric G proteins
 Monomeric G proteins, Tyr kinase/MAP kinase
 Protein modules

50

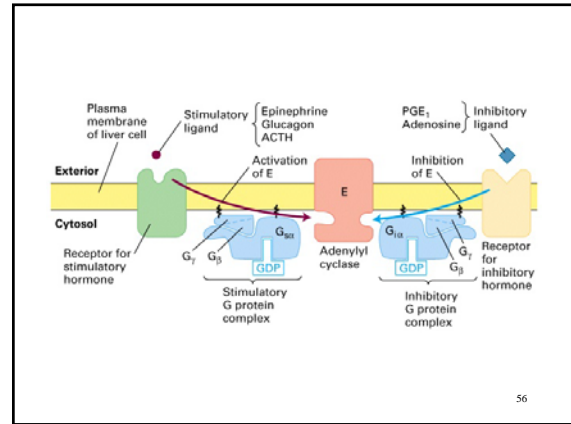
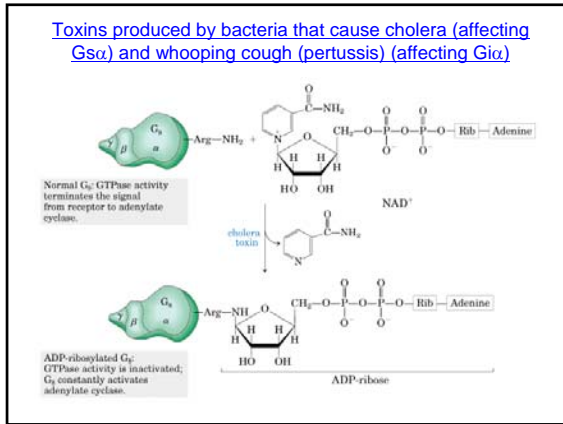
GTP-binding proteins: A superfamily

- Trimeric G proteins: G_s, G_i, G_q, G_t
- Small GTP-binding proteins
 - ✓ Ras: cell growth
 - ✓ Rac, Rho, Cdc42: cell migration
 - ✓ Rab: membrane trafficking
 - ✓ ARF: membrane trafficking
 - ✓ Ran: Nuclear transport
- Other GTP-binding proteins: Dynamin, EF-Tu

51



Toxins produced by bacteria that cause cholera (affecting $G_{\alpha s}$) and whooping cough (pertussis) (affecting $G_{\alpha i}$)



G_{α} Subclass*	Effect	Associated Effector Protein	2nd Messenger
$G_{\alpha s}$	↑	adenylyl cyclase	cAMP
	↑	Ca ²⁺ channel	Ca ²⁺
	↓	Na ⁺ channel	Change in membrane potential
$G_{\alpha i}$	↓	adenylyl cyclase	cAMP
	↑	K ⁺ channel	Change in membrane potential
	↓	Ca ²⁺ channel	Ca ²⁺
$G_{\alpha q}$	↑	Phospholipase C	IP ₃ , DAG
$G_{\alpha 12}$	↑	Phospholipase C	IP ₃ , DAG
$G_{\alpha 13}$	↓	Ca ²⁺ channel	Ca ²⁺
$G_{\alpha o}$	↑	cGMP phosphodiesterase	cGMP
$G_{\alpha 14}$	↑	Phospholipase C	IP ₃ , DAG
	↓	Adenylyl cyclase	cAMP

*A given G_{α} may be associated with more than one effector protein. To date, only one major $G_{\alpha o}$ has been identified, but multiple $G_{\alpha 12}$ and $G_{\alpha 13}$ proteins have been described. In some cases (not indicated in this table) effector proteins are regulated by coincident binding to G_{α} and $G_{\beta\gamma}$.

key: ↑ = stimulation; ↓ = inhibition; IP₃ = inositol 1,4,5-trisphosphate; DAG = 1,2-diacetylgllycerol.

source: See A. G. Dolphin, 1987, Trends Neurosci, 10:53; L. Birnbaumer, 1992, Cell 71:1069.

Family/subunit	Mass (kDa × 10 ⁻³)	% Amino acid identity ^a	Toxin ^b	Tissue distribution	Representative receptors	Effector/role	
$G_{\alpha s}$	$G_{\alpha s1}$ (27)	44.2	100	CTX	Ubiquitous	BAR ^c , glucagon	↑ Adenylyl cyclase
	$G_{\alpha s2}$	45.7	-	CTX	Ubiquitous	TSR, others	↑ Ca ²⁺ channels
	$G_{\alpha s3}$	44.7	88	CTX	Olfactory neuro-epithelium	Olfactant	↑ Na ⁺ channels
$G_{\alpha i}$	$G_{\alpha i1}$	40.3	100	PTX	Nearly ubiquitous	M ₂ Cho, α_2 AR, others	↓ Adenylyl cyclase
	$G_{\alpha i2}$	40.5	88	PTX	Ubiquitous	others	↑ Ca ²⁺ channels
	$G_{\alpha i3}$	40.5	94	PTX	Nearly ubiquitous	others	↑ Adenylyl cyclase (?)
	$G_{\alpha i4}$	40.0	73	PTX	Brain, others	Met-Enk, α_2 AR, others	↑ Phospholipase C (?)
	$G_{\alpha i5}$	40.1	73	PTX	Brain, others	others	↑ Phospholipase A ₂ (?)
$G_{\alpha q}$	$G_{\alpha q1}$	40	68	CTX, PTX	Retinal rods	Rhodopsin	↑ cAMP-specific phosphodiesterase
	$G_{\alpha q2}$	40.1	68	CTX, PTX	Retinal cones	Cone opsin	?
	$G_{\alpha q3}$	40.5	67	CTX (?)	Taste buds	Taste (?)	?
$G_{\alpha 12}$	40.9	60	-	Brain, adrenal pituitary	M ₂ Cho (?)	↓ Adenylyl cyclase (?)	
$G_{\alpha 13}$	$G_{\alpha 131}$	42	100	-	Nearly ubiquitous	M ₂ Cho, α_2 AR, others	↑ Phospholipase C ₂
	$G_{\alpha 132}$	42	88	-	Nearly ubiquitous	Lung, kidney, liver	↓ Ca ²⁺ channels
	$G_{\alpha 133}$	41.5	79	-	?	?	?
	$G_{\alpha 134}$	43	57	-	B cells, myeloid cells	?	?
$G_{\alpha 14}$	$G_{\alpha 141}$	43.5	58	-	T cells, myeloid cells	?	↑ Phospholipase C ₂
	$G_{\alpha 142}$	44	100	-	Ubiquitous	?	?
	$G_{\alpha 143}$	44	67	-	Ubiquitous	?	?

^a% Amino acid identity: comparison is with the first-listed member of each family.

^bCholera toxin (CTX) and pertussis toxin (PTX) catalyze the ADP-ribosylation of an Arg residue (CTX) and a Gln residue (PTX), respectively, of the indicated subunits.

^cSince various $G_{\alpha s}$ and $G_{\alpha i}$ short forms of $G_{\alpha s}$, $G_{\alpha i}$, long forms of $G_{\alpha s}$, $G_{\alpha i}$.

^dReceptor abbreviations: BAR, β -adrenergic; M₂Cho, M₂ muscarinic cholinergic; α_2 AR, α_2 -adrenergic; met-enk, met-enkephalin; M₂Cho, M₂ muscarinic cholinergic; M₂Cho, M₂ muscarinic cholinergic; M₂Cho, M₂ muscarinic cholinergic.

Subunit	Mass (kDa × 10 ⁻³)	% Amino acid identity ^a	Tissue distribution	Effector/role	
β	β_1	37.3	100	Ubiquitous	Required for G_{α} -receptor interaction
	β_2	37.3	90	Nearly ubiquitous	
	β_3	37.2	83	Nearly ubiquitous	
	β_4	37.2	89	Nearly ubiquitous	
γ	γ_1	8.4	100	Retina, other (?)	Support of agonist-induced receptor phosphorylation and desensitization
	γ_2	7.9	38	Brain, adrenal, other (?)	
	γ_3	8.5	36	Brain, testis	
	γ_4	(?partial)	(34)	[Kidney, retina (?)]	
	γ_5	7.3	25	Liver, other (?)	
	γ_6	7.5	35	Brain, other (?)	

^a% Amino acid identity: comparison is with the first-listed member of each family.

Lipid modification of trimeric G proteins

Subunit	Sequence	Modification
α_s	M G [] L G N S K T E D Q R N E -	P
α_{i1}	M [] T L S A E D K A A V E R -	M, P
α_t	M [] A G A S A E E K H S R E L -	M
α_q	M T L E S I M A [] L S E E A K A R R I N -	P

Prenylation

Subunit	Sequence	Modification
γ_1	- K G I P E D K N P F K E L K G G C ⁺ V I S	F
γ_2	- T P V P A S E N P F R E K K F F C ⁺ A I L	GG

P: palmitoylation; M: myristoylation; F: farnesylation; GG: geranylgeranylation

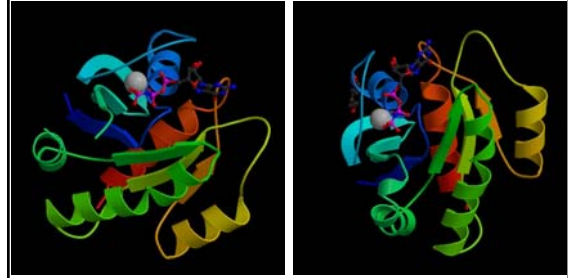
Small GTP-binding proteins

- Ras: cell growth
- Rac, Rho, Cdc42: cell migration
- Rab: membrane trafficking
- ARF: membrane trafficking
- Ran: Nuclear transport

61

Ras

⚠ Oncogenes discovered from murine sarcoma viruses (Harvey virus - H-ras, Kirsten virus - K-ras) and N-ras;
 ⚠ Activated in 10-50% of human tumors (G12V, Q61L, both of which are resistant to GAP).



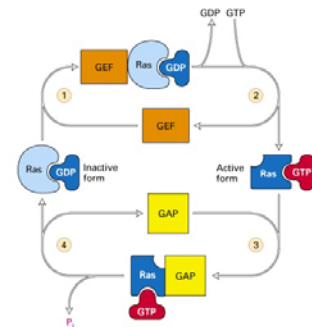
Conserved motifs in Ras-like proteins

N-terminus 4-31aa	Guanine nucleotide binding domain 160 aa					Extension 13-49 aa	C-terminus
	PM1	G1	PM2	PM3	G2	G3	Caax xCC CxC
	GxxxxGKs	F	T	DxxG	nKxD	ExSA	

PM: phosphate/Mg²⁺ binding regions; G: guanine base binding regions;
 Caax: a=aliphatic, x= any residue

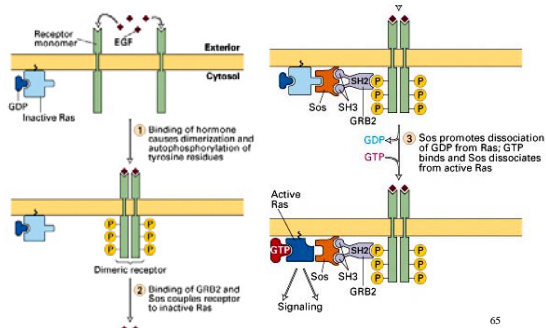
63

The GTP/GDP cycle of small GTP-binding proteins



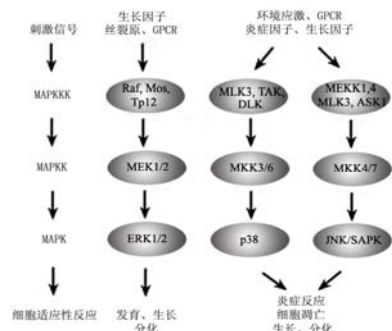
64

Ras mediates signaling from Receptor Tyrosine Kinase (RTK)

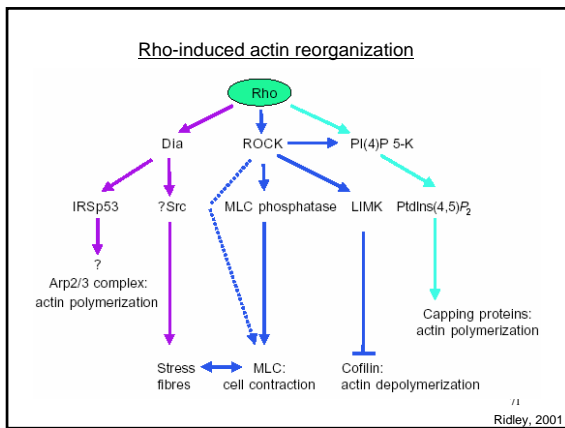
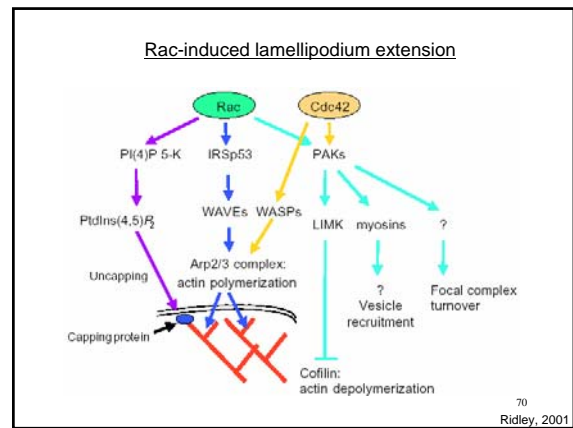
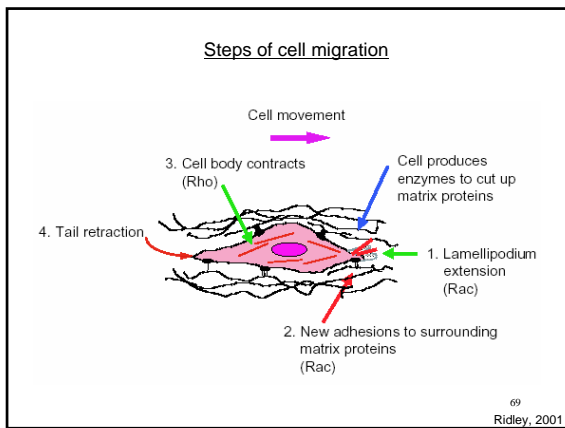
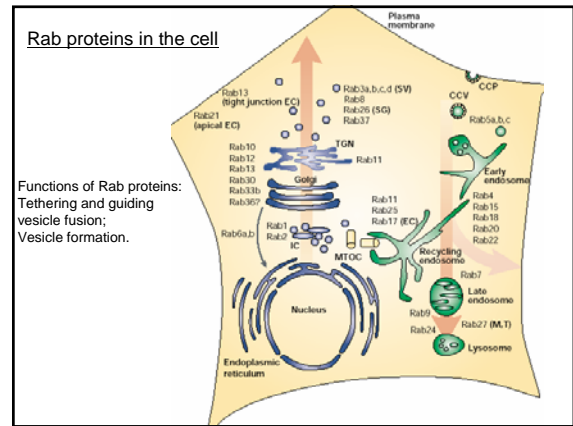
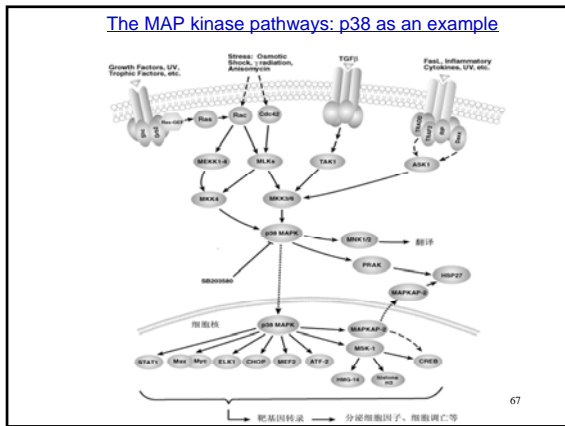


65

MAP kinase pathways



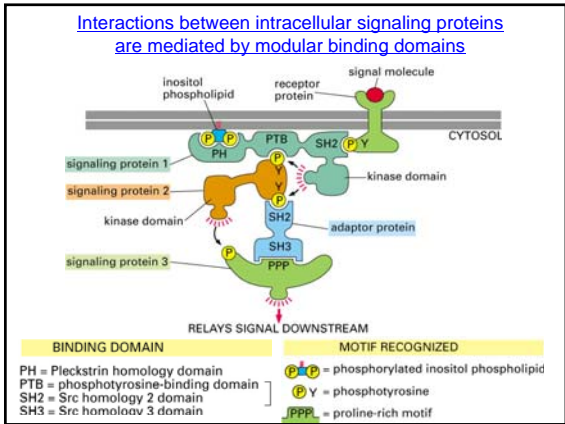
66



- General introduction
 - Common features of signal transduction
 - Cell surface signal transducers, receptors
 - Ion channels
 - Secondary messengers
 - cAMP
 - cGMP
 - Lipids
 - Calcium
 - G proteins
 - Trimeric G proteins
 - Monomeric G proteins, Tyr kinase/MAP kinase
 - Protein modules
- 72

Protein Modules: protein building blocks

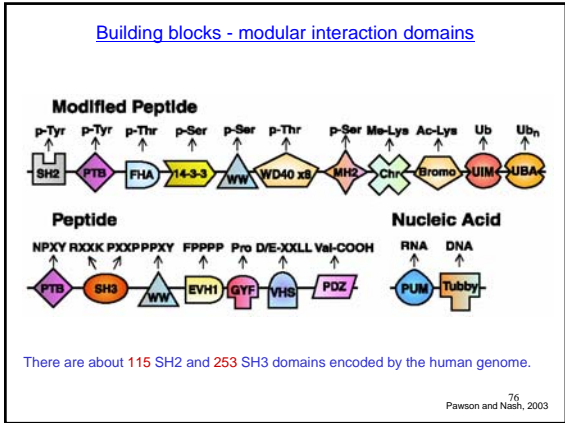
73



Protein modules

PTB domains: ~ 100-150 aa, bind to NPXY motifs: Shc, IRS-1
 PDZ domains: ~80-90 aa, recognize short peptide motifs (4-5 residues) at the C-terminus of membrane proteins, usually containing a hydrophobic residue at the very end; protein-protein interaction: Dishevelled, FAP
 SH2 (src homology): ~100 aa, binds to phosphotyrosine residues: Src, Grb2, Shc, STAT
 SH3: binds to proline-rich sequences (PXXP): Src, Nck
 WW domains: bind to Pro-rich sequences (XPPXY): Nedd4 (E3 ubiquitin ligase), Smurf, Dystrophin
 Death domains: Fas
 LIM domains: recognize turn-based motifs
 PH (Pleckstrin-homology) domains: associate with phosphoinositides (PI_{3,4}P₂; PI_{4,5}P₂; PI_{3,4,5}P₃), target proteins to the plasma membrane: Akt, SOS
 FYVE domains: associate with phosphoinositides (PI₃P), target proteins to endosomes: EEA1, SARA

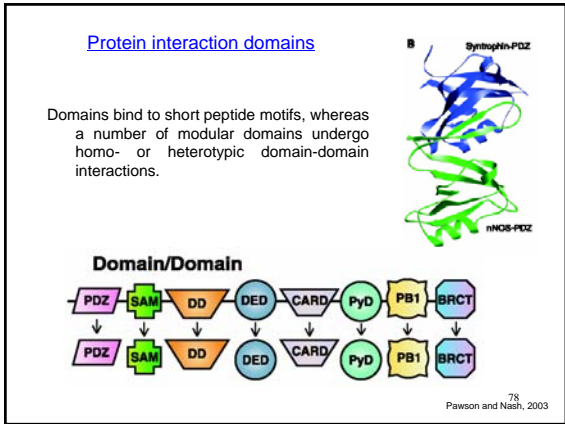
75



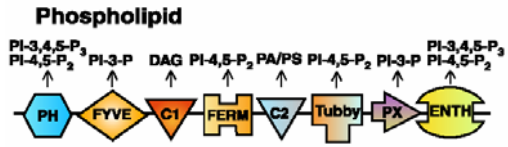
Repeated motifs

Some interaction domains are assembled from repeated motifs (up to 50 copies): HEAT, TPR, Arm, ankyrin, leucine-rich, Pumilio repeat.

77
Pawson and Nash, 2003

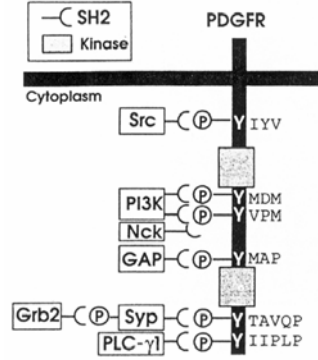


Protein-lipid interaction domains



79
Pawson and Nash, 2003

Specificity of SH2 binding to phosphotyrosine



80
Pawson, 1995

谢谢!

81