Cell Biology Lecture 9

Cell Signaling - principles Sesilja Aranko 22.11.2023 Alberts • Johnson • Lewis • Morgan • Raff • Roberts • Walter

Molecular Biology of the Cell

Sixth Edition

Chapter 15 Cell Signaling, Part I Pages: 813-831

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Course overview – Tentative schedule

Date	Lecture		Chapters & Topi	cs		Assignments
25.10.	1		Course overview, DNA, Chromosomes, Genome,	<u>Ch</u> . 4		
27.10.	2 -G	art 1	Replication, Repair, Recombinat	ion, Ch. 5		
1.11.	3	Ĩ	From DNA to protein, Ch. 6			
3.11.	4		Control of gene expression, Ch. 7			
8.11.	5		Membrane structures, Ch. 10 Membrane transport, Ch. 11			Assignment I (Essay) Draft I (8.11.)
10.11.	6 -G	2	Intracellular compartments and r Ch. 12	orotein sortin	IQ.	
15.11.	7	Part	Intracellular compartments and p Ch. 12 Susanna Mäkinen, Solar Food	orotein sortin s	ıg,	Assignment II – Draft I (15.11.)
17.11.	8		Membrane Traffic, Ch. 13 iGEM team 2023			+iGEM intro
22.11.	9		Cell signalling, Ch. 15			Assignment II – Peer review (22.11.)
24.11.	10 -G	<u>ب</u>	Cell signalling, Ch. 15			Assignment I (Essay) Draft II (24.11.)
29.11.	11	Part	Cell cycle. Ch. 17 Jere Weltner, Folkhälsan			
1.12.	12		Apoptosis, Ch. 18			Assignment II – final version (1.12.)
7.12.	EXAM		December 7th			
8.12.	Final version essay		December 8th			Assignment I (Essay) Final version (8.12.) Aim at finishing before exam date. Use last days for polishing.

LEARNING OUTCOMES

 Can describe the principles of cell signaling and apply these to evaluate realistic cases of signaling

CONTENTS

- Principles of cell signaling
- Signaling through G-protein-coupled receptors
- Signaling through enzyme-coupled receptors

- Communication between cells in multicellular organisms
- \rightarrow signaling pathways:
- Activated by an extracellular signal molecule, typically binding to a receptor protein that is embedded in the plasma membrane of the target cell

Signal molecule can be protein or peptide, but also amino acid, nucleotide, steroid etc., even dissolved gas such as carbon monoxide



- Communication between cells in multicellular organisms
- \rightarrow signaling pathways:
- Activated by an extracellular signal molecule, typically binding to a receptor protein that is embedded in the plasma membrane of the target cell
- The receptor activates one or more intracellular signaling pathways, involving a series of signaling proteins and small chemical messengers
- Finally, one or more of the intracellular signaling molecules alters the activity of effector proteins and thereby the behavior of the cell



• Extracellular signals can act over short or long distances



• Receptors can be on the cell surface or intracellular



• In both cases, high *specificity* and *affinity*

 Each cell is programmed to respond to specific combinations of extracellular signals



- Combination of signals allows having less different receptors, yet multiple different responses
- To add up complexity, also inhibitory pathways

Various responses induced by the neurotransmitter acetylcholine



- Same type of acetylcholine receptor (a Gprotein-coupled receptor)
- Intracellular signals produced are interpreted
- Receptor protein is different (an ion-channelcoupled receptor)

THREE CLASSES OF CELL-SURFACE RECEPTOR PROTEINS



- Synaptic signaling between nerve cells
- Signal molecules (neurotransmitters) transiently open ion channel

THREE CLASSES OF CELL-SURFACE RECEPTOR PROTEINS



THREE CLASSES OF CELL-SURFACE **RECEPTOR PROTEINS**



Synaptic signaling between nerve cells Signal molecules (neurotransmitters) transiently open ion channel

- **G**-protein mediates signal from receptor to target protein
 - Receptor is an enzyme that act directly or by activating an associated enzyme

- Cell-surface receptors relay signals via *intracellular signaling molecules* ("second messengers")
- Intracellular signaling proteins act as molecular switches



- 30-50% proteins in human phosphorylated
- Ser/Thr kinases (majority)
- Tyr kinases
- Kinase cascades

- Cell-surface receptors relay signals via *intracellular signaling molecules* ("second messengers")
- Intracellular signaling proteins act as molecular switches



- G-proteins
- Monomeric GTPases

- G-proteins
- Monomeric GTPases



- GAP= GTPase activating protein
- *GEF=guanine nucleotide exchange factor*
- GAPs inactivate the protein by stimulating it to hydrolyze its bound GTP to GDP
 - GDP remains tightly bound to the inactivated GTPase
- **GEFs activate** the inactive protein by stimulating it to release its GDP
 - The concentration of GTP in the cytosol is 10 times greater than the concentration of GDP → the protein rapidly binds GTP and is thereby activated.

- G-proteins
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• Activated G-proteins act as GEFs

- Intracellular signals must be specific and robust in a noisy cytoplasm. Ways to ensure specificity:
- 1. Spesificity of catalytic site towards specific residue
- 2. Selection of **consensus sequence** near the active site
- **3. Docking sites** that provide enforced proximity and allosteric regulation of a correct kinase–substrate pair
- **4. Localization**, increases local concentrations. May be achieved by **condensation**
- **5. Scaffolds**, form dynamic ternary complexes with kinases and substrates
- 6. Competition between correct and wrong residues
- 7. Multisite phosphorylation and kinetic proofreading, minimize the harm by wrong phosphorylations

- Intracellular signaling complexes form at activated cellsurface receptors:
 - 1. Preformed signaling complexes on a scaffold protein
 - 2. Assembly of signaling complex on an activated receptor
 - 3. Assembly of signaling complex on phosphoinositide binding site
- Helps to enhance specificity

1. Preformed signaling complexes on a scaffold protein



A receptor and some of the intracellular signaling proteins it activates in sequence are **preassembled into a signaling complex** on the inactive receptor by a large *scaffold protein*

2. Assembly of signaling complex on an activated receptor



- A signaling complex assembles *transiently* on a receptor only after the **binding** of an extracellular signal molecule has activated the receptor
- Typically, activated receptor phosphorylates itself at multiple sites, which then act as docking sites for intracellular signaling proteins

3. Assembly of signaling complex on phosphoinositide binding site



- Activation of a receptor leads to the increased phosphorylation of specific phospholipids (phosphoinositides) in the adjacent plasma membrane
- **PIPs serve as docking sites** for specific intracellular signaling proteins, which can now interact with each other

 Modular interaction domains mediate interactions between intracellular signaling proteins



Modular interaction domains guide the formation of a specific signaling complex



example based on the insulin receptor

Modular interaction domains guide the formation of a specific signaling complex



Formation of large receptor clusters by multivalent interactions among signaling proteins



receptor tails

extracellular

two adaptor proteins:

- One contains one SH2 domain, which binds phosphorylated tyrosines on the receptors, and two SH3 domains.
- The other contains three proline-rich regions that can bind to SH3 domains, plus a protein kinase domain.

Formation of large receptor clusters by multivalent interactions among signaling proteins

Numerous multivalent binding interactions can occur among the three components in this system, generating a crosslinked protein matrix or condensate in which the protein kinases of the receptor and adaptor protein are concentrated, potentially providing a more effective signal output.

extracellular

The cross-linking of the matrix can be enhanced further by including adaptor proteins with domains that interact with modified phospholipids in the membrane.

Question

A kinase cascade organized by a scaffold protein or composed of freely diffusing components

Compare:

- Amplification
- Speed
- Crosstalk between pathways



PROTEIN PHOSPHORYLATION AND KINASES

- Protein phosphorylation is the most common post-translational modification
- Affects almost every basic cellular process



Peptidyl serine



Peptidyl phosphoserine

- Ser/Thr kinases (majority)
 - cAMP/cGMP, diacylglycerol, and Ca²⁺/calmodulin
- Tyr kinases
 - Typically, receptor kinases or receptor associated kinases



https://www.nature.com/articles/nrm2203

Protein kinases phosphorylate typically 1-~hundreds of residues out of ~700,000 potential target residues \rightarrow a range of mechanisms to ensure specificity

Depth of the kinase catalytic cleft
 → tyrosine or serine/threonin e residues

IRK: deep cleft to accommodate Tyr









https://www.nature.com/articles/nrm2203

2. Local interactions near the phosphorylation site \rightarrow consensus sequences

Table 1 Consensus phosphorylation sites of some protein kinases								
Kinase	Full name	Consensus phosphorylation site	Refs					
РКА	Protein kinase A or cAMP- dependent protein kinase	R-R-X-S/T-Φ	5,39					
CDK	Cyclin-dependent kinase	S/T-P-X-K/R	5,39					
ERK2	Extracellular-regulated kinase-2	P-X-S/T-P	5,136					
CK1*	Casein kinase-1	pS-X-X-S/T	5,137					
CK2 [‡]	Casein kinase-2	S/T-D/E-X-E/D	5,138					
GSK3	Glycogen synthase kinase-3	S-X-X-PS	5,139					
CaMK2	Calmodulin-dependent protein kinase-2	R-X-X-S/T	5,136					
ABL	Abelson murine leukaemia virus tyrosine kinase	I/V/L-Y-X-X-P/F	5,140					
EGFR	Epidermal growth factor receptor	E-E-E-Y-F	5,141					
Src	Rous sarcoma virus tyrosine kinase	E-E-I-Y-E/G-X-F	5,141					
IRK	Insulin receptor tyrosine kinase	Y-M-M-M	5,141					
PKB/AKT	Protein kinase B	R-X-R-X-X-S/T	142					
PKD	Protein kinase D	L/I-X-R-X-X-S/T	40					
PIM1-3	Proviral integration site kinases 1–3	R-X-R-X-X-S/T	40,143					

*CK1 is a well-conserved Ser/Thr-specific protein kinase, the regulation and function of which are incompletely understood. *CK2 is also a well-conserved Ser/Thr kinase that is unrelated to CK1 and is implicated in the regulation of diverse biological phenomena. pS, phosphorylated Ser; X, any residue; Φ, hydrophobic residue.

nttps://www.nature.com/articles/nrm2203

3. Docking sites are separated from the catalytic site of the kinase and the phosphorylation site of the substrate, can provide enforced proximity and allosteric regulation of a correct kinase–substrate pair

4. Localization restricts kinases to a subset of substrates and increases local kinase concentrations.

5. Scaffolds, which form dynamic ternary complexes with kinases and substrates and might contribute to kinase specificity in several



https://www.nature.com/articles/nrm2203

https://www.cell.com/trends/cell-biology/fulltext/S0962-8924(22)00260-4

6. Competition ensures that the correct residues are targeted for phosphorylation. This mechanism can suppress the phosphorylation of off-target substrates and can add thresholds and temporal ordering to phosphorylation responses.



7. Multisite phosphorylation and kinetic proofreading, minimize the consequences of aberrant phosphorylations

https://www.nature.com/articles/nrm2203

- The relationship between signal and response varies in different signaling pathways
 - Timing (speed of response)
 - Sensitivity
 - Dynamic range
 - Persistence
 - Signal processing

- The relationship between signal and response varies in different signaling pathways
 - Timing (speed of response)
 - Sensitivity
 - Dynamic range
 - Persistence
 - Signal processing
 - Coordination of multiple responses for one signal
 - Integration (of multiple signals)
 - Including AND gates



THE SPEED OF A RESPONSE

• The **speed** of a response depends on the **nature** of the signaling molecules



THE SPEED OF A RESPONSE AND TURNOVER

- The **speed** of a response depends on the **turnover** of signaling molecules
- The synthesis rates are
 (A) decreased or (B)
 increased by a factor of 10
- The concentrations of molecules that are degraded rapidly (*red lines*) change quickly, the concentrations of those that are degraded slowly (*green lines*) change proportionally more slowly



been increased by a factor of 10

been *decreased* by a factor of 10

- Cells can respond abruptly to a gradually increasing signal
- Response increases gradually as the concentration of extracellular signal molecule increases
- Eventually reaching a plateau (signaling pathway saturated)
- *=Hyperbolic* response curve
- E.g. hormones



concentration of signal molecule -----

• Cells can respond abruptly to a gradually increasing signal

- Response is switchlike
- The cell switches completely between a low and high response
- Typical to control cell states



concentration of signal molecule -----

- Cells can respond abruptly to a gradually increasing signal
- The signaling system reduces the response at low signal concentrations
- Produces a steeper response at some intermediate signal concentration
- = Sigmoidal response curve
- Examples, allostreric regulation (next slide) or simultanous activation and inhibition of opposite reactions



concentration of signal molecule -----

Activation curves for an allosteric protein as a function of effector molecule concentration

- The curves show how the sharpness of the activation response increases with an increase in the number of effector molecules that must be bound simultaneously to activate the target protein
- An example of sigmoidal response
- E.g. phosphorylation of multiple sites or binding of multiple signalling molecules required



POSITIVE AND NEGATIVE FEEDBACK

- Positive feedback can generate an all-or-none response
- Negative feedback is a common feature of intracellular signaling systems



POSITIVE FEEDBACK

- Positive feedback can generate an all-or-none response
- May be bistable exists in on-state or off-state that *persists* after original signal level drops
- E.g. cell memory in the differentiation of cells permanent change without a change in DNA



POSITIVE FEEDBACK

- Positive feedback can generate an all-or-none response
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- The input signal is an activated protein kinase (S) that phosphorylates and thereby activates another protein kinase (E)
- (A protein phosphatase (I) dephosphorylates and inactivates the activated E kinase)
- Without feedback, the activity of the E kinase is proportional to the level of stimulation by the S kinase
- With the positive feedback loop, the transient stimulation by S kinase switches the system from an "off" state to an "on" state, which then persists after the stimulus has been removed



time during which the input signal (activated S kinase) is present

- The input signal is an activated protein kinase (S) that phosphorylates and thereby activates another protein kinase (E)
- A protein phosphatase (I) dephosphorylates and inactivates the activated E kinase



(C)

- Without feedback, the activity of the E kinase is proportional to the level of stimulation by the S kinase
- Short delay: the system shows a response when the signal is first increased, but the feedback quickly dampens the response—which then declines to some intermediate level at which the input signal and feedback are balanced



- Without feedback, the activity of the E kinase is proportional to the level of stimulation by the S kinase
- Long delay: the response rises unopposed at first, allowing kinase activity to reach maximum levels before it feeds back to shut itself off. Then the sudden drop in activity removes the negative feedback, unleashing another pulse of kinase activity
- If conditions are right, the result is sustained oscillations for as long as the stimulus is present



- Without feedback, the activity of the E kinase is proportional to the level of stimulation by the S kinase
- Short delay: the system shows a response when the signal is first increased, but the feedback quickly dampens the response—which then declines to some intermediate level at which the input signal and feedback are balanced
- Short delay adaptation, response to change



ADAPTATION AND DESENSIZATION

- Cells can adjust their sensitivity to a signal adaptation and desensization
- Negative feedback with a short delay dampens the initial response to receptor activation



ADAPTATION AND DESENSIZATION

- Cells can adjust their sensitivity to a signal adaptation and desensization
- In some cases, the activated receptor rapidly activates a stimulatory pathway while also initiating a slower inhibitory pathway—resulting in a transient output response. This is called a delayed feed-forward loop.
- Various mechanisms can inactivate a cell-surface receptor



TRANSPORT INTO THE CELL FROM THE PLASMA MEMBRANE: ENDOCYTOSIS

• Specific proteins are retrieved from early endosomes and returned to the plasma



SIGNALING THROUGH G-PROTEIN-COUPLED RECEPTORS

 GPCR Desensitization depends on receptor phosphorylation



A GRK phosphorylates only activated receptors because it is the activated GPCR that turns on the GRK. The binding of an arrestin to the phosphorylated receptor prevents the receptor from binding to its G protein and also directs its endocytosis

Question

Tell 3 examples of how a signal can be sharpened?

SUMMARY

- Modular interaction domains mediate interactions between intracellular signaling proteins
- The relationship between signal and response varies in different signaling pathways
- The **speed of a response** depends on the **turnover** of signaling molecules
- Cells can respond abruptly to a gradually increasing signal
- Positive feedback can generate an all-or-none response
- **Negative feedback** is a common feature of intracellular signaling systems
- Cells can adjust their sensitivity to a signal

CELL-FREE PROTEIN SYNTHESIS



Brookwell, A.; Oza, J.P.; Caschera, F. Biotechnology Applications of Cell-Free Expression Systems. Life 2021, 11, 1367. https://doi.org/10.3390/life11121367

CELL-FREE VS *IN-VIVO* PROTEIN EXPRESSION

Cell-Free Protein Synthesis Cell-Based Protein Synthesis Extract Preparation Transformation and Expression Harvest Lysis Culture Growth Transformation Culture Growth/Expression Harvest Post-lysis Processing Cell Extract Expression Capability to plug-and-play reaction components and 0 0 1 conditions 0 0 Centrifugation Supernatant 70 Protein Expression Cell Extract Applications 12 Prototyping Discovery Biomanufacturing

ADVANTAGES AND DISADVANTAGES OF EACH?

https://bioprocessintl.com/upstream-processing/expression-platforms/toward-a-roadmap-for-cell-free-synthesis-in-bioprocessin