

Signaling Pathways

Properties and approaches to intracellular signaling

Properties of signaling enzymes & molecules

Structure of G proteins, kinases

Example of analysis using long-term potentiation and MAP kinase

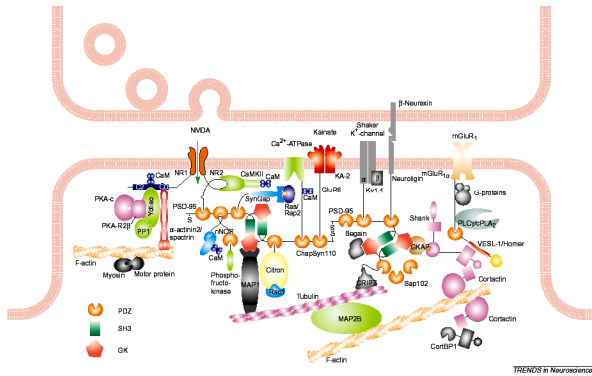
Example of analysis using Leptin receptor and Jak-Stat signaling

Approaches to molecular signaling

Example of CREB and c-Fos as molecular signals

1

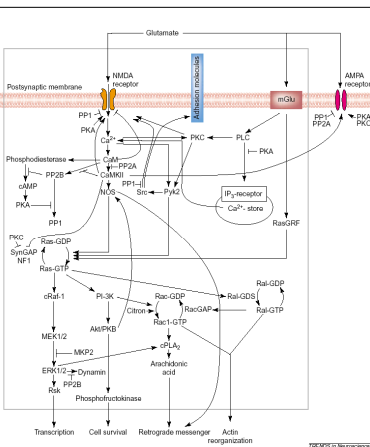
Complexity of the postsynaptic density



Interactome: the defined interactions between all proteins in the genome/proteome

Grant 2001

2



Grant 2001

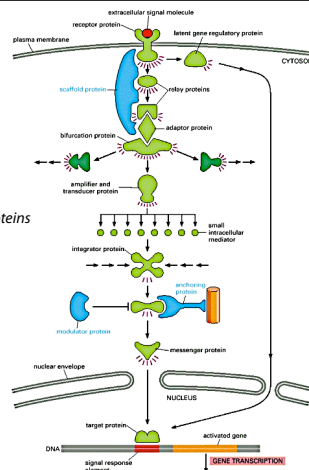
Collins 2006

Table 1 Summary of PSD protein classifications

	Total PSD
Channels and receptors	80
MAGUKs/adaptors/scaffolders	54
Serine/threonine kinases	46
Tyrosine kinase	3
Protein phosphatases	18
G proteins and modulators	77
Signalling molecules and enzymes	278
Transcription and translation	119
Cytoskeletal and cell adhesion molecules	153
Synaptic vesicles and protein transport	159
Novel	107
Other	30
Summary	1124

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1. Relay proteins
2. Messenger proteins
3. Adaptor proteins
4. Amplifier proteins
5. Transducer proteins
6. Bifurcation proteins
7. Integrator proteins
8. Latent gene regulatory proteins



General Principles of Cell Communication

<http://www.ncbi.nlm.nih.gov/books/NBK26813/>

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1. *Relay proteins* simply pass the message to the next signaling component in the chain.
2. *Messenger proteins* carry the signal from one part of the cell to another, such as from the **cytosol** to the **nucleus**.
3. *Adaptor proteins* link one signaling **protein** to another, without themselves conveying a signal.
4. *Amplifier proteins*, which are usually either enzymes or **ion** channels, greatly increase the signal they receive, either by producing large amounts of small intracellular mediators or by activating large numbers of downstream intracellular signaling proteins. When there are multiple amplification steps in a relay chain, the chain is often referred to as a **signaling cascade**.
5. *Transducer proteins* convert the signal into a different form. The **enzyme** that makes cyclic AMP is an example: it both converts the signal and amplifies it, thus acting as both a transducer and an amplifier.
6. *Bifurcation proteins* spread the signal from one signaling pathway to another.
7. *Integrator proteins* receive signals from two or more signaling pathways and integrate them before relaying a signal onward.
8. *Latent gene regulatory proteins* are activated at the cell surface by activated receptors and then migrate to the **nucleus** to stimulate gene transcription.

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Properties of Intracellular Signaling Networks

Common Intracellular Signaling Pathways



- AC = adenylyl cyclase
 PDE = phosphodiesterase
 PKA = protein kinase A
 pCREB = phospho cAMP Response Element Binding protein
 PP1 = protein phosphatase 1

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Common Intracellular Signaling Pathways



Ca⁺⁺/Calmodulin

PLC/DAG/IP3 pathway

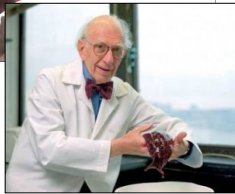
Receptor Tyrosine Kinases
e.g. Neurotrophins, JAK, STAT

MAPK (mitogen activating protein) pathway



8

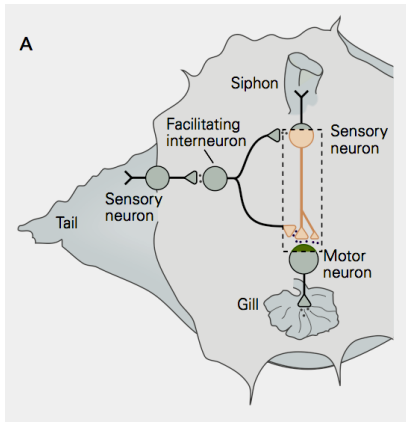
Intracellular signaling
in Aplysia learning



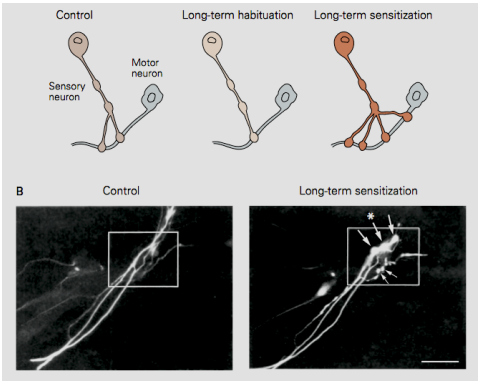


9

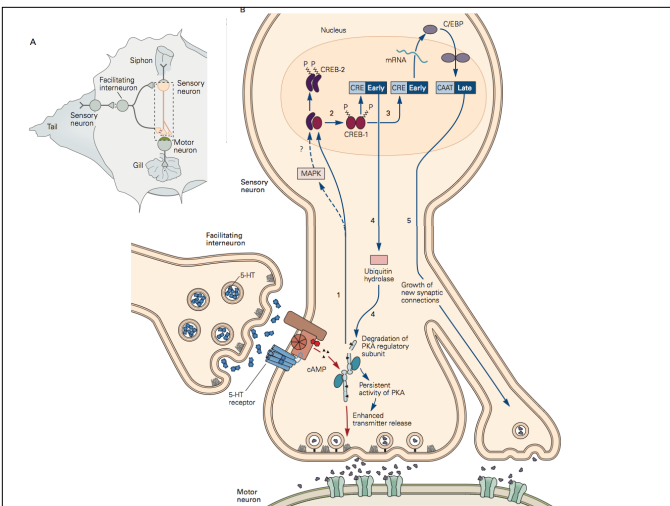
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Principles of Intracellular Signaling

Mediated by factors and enzymes (not ions)

Diffusion of factors and enzymes takes milliseconds to seconds

Enzymes make covalent modifications

e.g. phosphorylation

Enzymatic reactions take milliseconds

Enzymatic reactions are substrate-specific

Reaction products can persist from milliseconds to weeks

Factors, Enzymes, and their substrates and compartmentalized

Members of a signaling pathways are often regulated by each other

Components of Signaling Pathway are Compartmentalized within the Neuron

membrane bound/associated

localized to PSD or dendrite

sequestered in membrane compartment

e.g. *Ca⁺⁺ in sarcoplasmic reticulum*

localized to nucleus

translocation to other compartments

e.g. *dendrite to nucleus (PKA, MAPK)*

e.g. *axon terminal to nucleus (Trks)*

Intracellular Signaling mediated by Messengers & Enzymes

Messengers:

- a generated/released small factor
- a phosphorylated protein

Enzymes:

- generate/release small factors
- phosphorylate proteins (kinases)
- dephosphorylate proteins (phosphatases)
- modify cellular targets (usually by phosphorylation)

Examples of Signaling Messengers

Properties of Intracellular Signaling Networks:

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Generated/released small factors, e.g.

DAG

membrane-bound factor derived from phospholipids.
can activate membrane-associated enzymes

cAMP

cytoplasmic factor derived from ATP by adenylate cyclase.
can activate cytoplasmic enzymes such as protein kinase A (PKA)

NO

a gaseous factor generated from amino acids by NO synthase.
can diffuse to surrounding cells and that can activate enzymes,
esp. activate guanylate cyclase → cGMP

Endocannabinoids

derived from membrane phospholipids
bind retrogradely to presynaptic cannabinoid receptors

Phosphorylated proteins, e.g.

Inhibitor I

blocks protein phosphatase I

Signaling Enzymes can:

Properties of Intracellular Signaling Networks

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Generate Small Factors

Adenylate Cyclase

generates a small factor, cAMP

Phosphorylate

Protein Kinase A

when activated by cAMP, adds $P_{O_4}^-$ to serine/threonine residues of proteins

Dephosphorylate

Protein Phosphatase 1

removes the $P_{O_4}^-$ from ser/thr residues

Modify Cellular Targets

Substrates of PKA and PP1 include

NMDA glutamate channel (regulates excitability)

CREB transcription factor (regulates gene expression)

Detection of second messengers & signaling pathways

1. Chemical assay

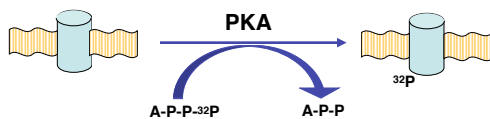
e.g. HPLC to detect small amounts of a simple compound

2. Visualization

e.g. Ca^{++} sensitive dyes that fluoresce in high Ca^{++}
e.g. antibodies, such as anti-cAMP

3. Incorporation of radio-tracer

e.g. GTP with ^{32}P for G-Proteins
ATP with ^{32}P for kinases



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Detection of second messengers & signaling pathways

Immunohistochemistry for protein or phosphorylated protein

Western blot of protein

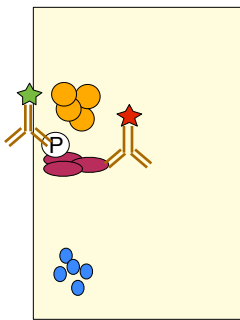
e.g. with ³²P labeling of substrate
size shift from protein (smaller) to phosphoprotein (bigger)

Western blot for phosphorylated protein

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Western Blot

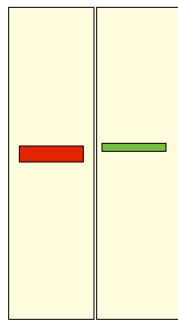
5. Blot proteins onto nylon membrane



6. Visualize proteins using labeled antibodies

7. Visualize phosphorylated proteins using phospho-specific antibodies

Developed Blot

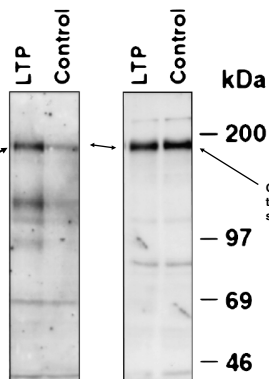


protein phospho protein

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α PY = antibody against phosphotyrosine residue

α PY α NR2B



stimulated hippocampus shows general increase in phosphotyrosine proteins.

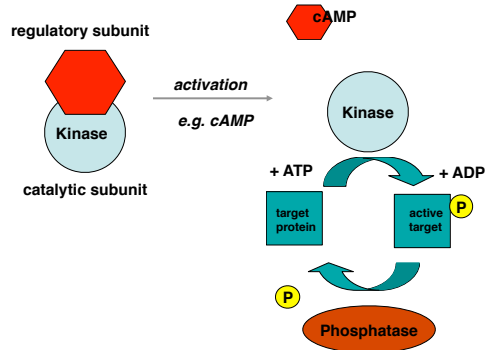
One of those proteins is the NMDA receptor 2B subunit.

Rosenblum 1996

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Kinases and Phosphatases

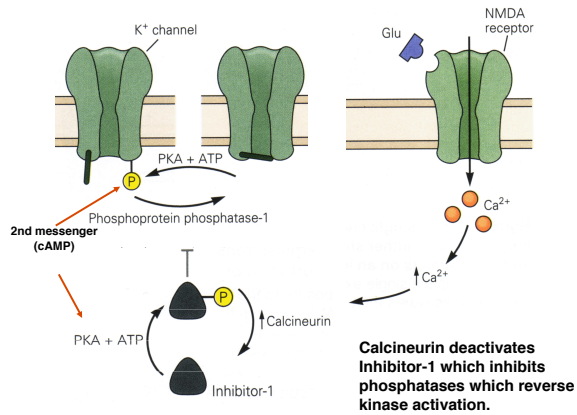
22



Squire 10.10

Don't forget to regulate the phosphatases...

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Properties of Intracellular Signaling Networks

Implications of Signaling Properties

Slower

time course of seconds to hours

Persistent

substrates remain modified for minutes to weeks

Spatial Resolution

can localize to specific compartments

Visualization

easier to detect activity and modifications of specific large molecules and proteins

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Protein Domains:

e.g., G-Protein coupled receptors:

7-transmembrane domains

ligand-binding domains

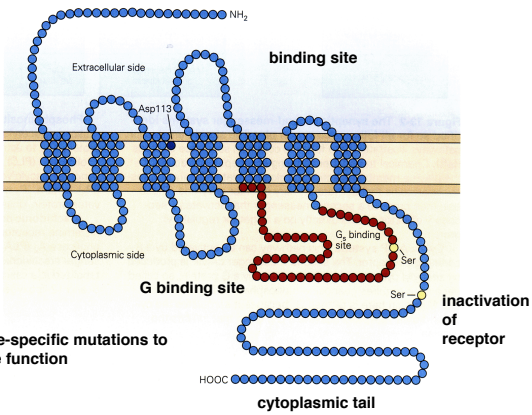
G-protein binding domains

Kinases

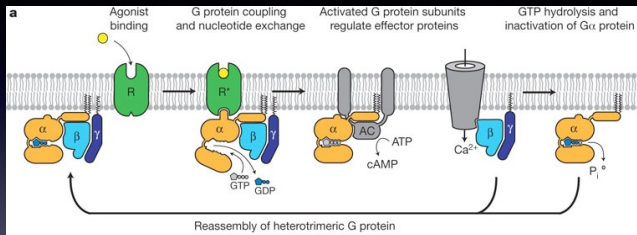
Phosphatases

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G-protein coupled receptors



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Useful toxins to block Signaling Pathways

G-Protein Toxins:

1. Cholera Toxin

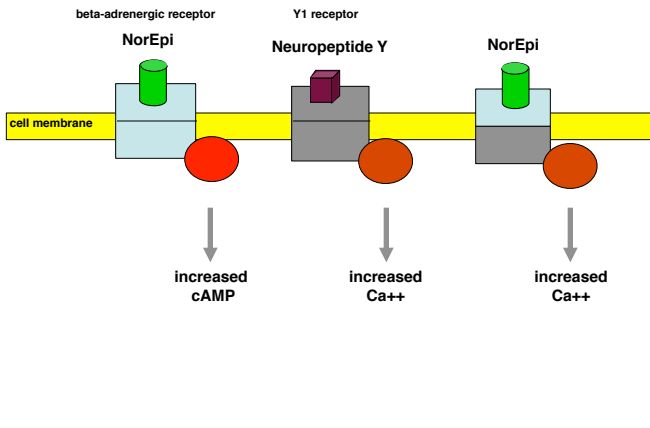
allows binding of GTP, but prevents hydrolysis.

Causes overproduction of cAMP, leading to loss of electrolytes and water from intestinal cells.

2. Pertussis Toxin

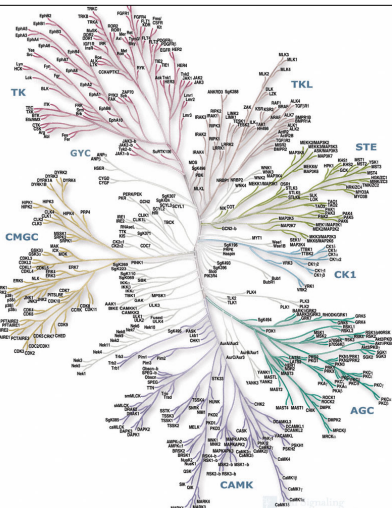
blocks release of GDP from alpha subunit, so G-protein locked in the inactive state.

Chimeric receptors

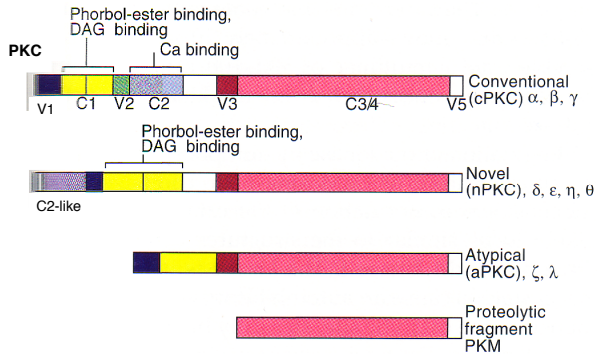


Complexity of the kinase families

Human Protein Kinases



Kinases have domains that can be mixed and matched



Alternative splicing/processing of PKC enzyme Squire 10.12

Kinases phosphorylate specific sites in targets

TABLE 10.4 Consensus Phosphorylation Sites of Some Protein Kinases

Protein kinase	Consensus phosphorylation site
PKA	R-R / K-X-S* / T*
PKG	R / K _{2,3} -X-S* / T*
cPKC	(R / K _{1,3} -X _{2,0})-S* / T*-(X _{2,0} -R / K _{1,3})
CaM kinase II	R-X-X-S* / T*
MLCK (smooth muscle)	(K / R ₂ -X)-X _{1,2} -K / R ₃ -X _{2,3} -R-X ₂ -S*-N-V-F

Source: Adapted from Kennelly and Krebs.⁵⁴
 R, Arg; K, Lys; S*, phospho-Ser; T*, phospho-Thr; X, polar amino acid; N, Asn; V, Val; F, Phe.

- 1) can use sequence to predict phosphorylation site
- 2) can raise antibody against phosphorylated consensus site

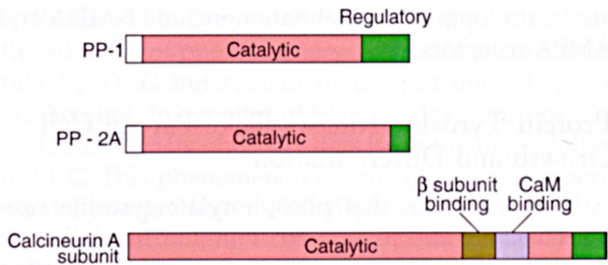


FIGURE 10.14 Domain structure of the catalytic subunits of some Ser/Thr phosphatases. The three major phosphoprotein phosphatases, PP-1, PP-2A, and calcineurin, have homologous catalytic domains but differ in their regulatory properties.

The NEW Scansite 3

Database Search Results

Home News Search Input Scansite Features Scan Protein for Motifs Search a Sequence Database for Motifs Find Sequence Match Calc. MolWeight and pI Calc. Amino Acid Composition Databases and Motifs FAQ Tutorial About Collaborator and Administrator Login

Motifs: PKA_Rin
Database: SwissProt
Species restriction: Organism Class: Mammals
Keyword restriction:
Sequence restriction:
Number of Phosphorylation Sites: 0
Isoelectric Point: from 0
Molecular Weight: from 0

Search Results
Total Number of Proteins in Database: 533049
Number of Proteins Matching Restrictions: 65559 (these proteins have been scored using the given motif(s))
Number of Predicted Sites Found: 179736
Median of Scores: 0.555
Median Absolute Deviation of Scores: 0.03960

Predicted Motif Sites
Please allow popups in your browser settings to make links in the table work properly!
Displaying up to 50 predicted motif sites. You can download the complete list of results in the section below

Scan this Protein!	Score	Accession	Protein Annotations	Site [Protein Kinase A]	Sequence [Protein Kinase A]	M	W
Scan!	0.104	KCNK9_CANVD	Description: RecName: Full=Potassium channel subfamily K member 9; AltName: Full=Acid-sensitive potassium channel protein TASK-3; AltName: Full=TWIK-related acid-sensitive K(+) channel 3.; Keywords: Potassium channel, Glycoprotein, Ion transport, Potassium transport, Ionic channel, Transport, Transmembrane helix, Potassium, Membrane.	8364	RRLMLRRAKSY*****	40	

40

The NEW Scansite 3

21 matches

Scan!	0.221	HNLD0_HUMAN	Description: RecName: Full=Protein Wnt-10a; Flags: Precursor.; Keywords: Hypotrichosis, Signal, Wnt signaling pathway, Polymorphism, Developmental protein, Palmoplantar keratoderma, Glycoprotein, Secreted, Ectodermal dysplasia, Extracellular matrix, Disease mutation, Reference proteome, Complete proteome; Accessions: Q9H7S8, Q96TAT, Q9QZT5;	8332	PGPRRRAaPADLVTF	46450	
Scan!	0.221	CREB1_HUMAN	Description: RecName: Full=Cyclic AMP-responsive element-binding protein 1; Short=CREB-1; Short=CREB; Keywords: Ubi conjugation, Chromosomal rearrangement, Transcription, Alternative splicing, Host-virus interaction, DNA-binding, Isopeptide bond, Phosphoprotein, Transcription regulation, Activator, Reference proteome, Complete proteome, Nucleus; Accessions: P21934, Q9UMA7, P16220;	8133	EILSRAPAYRKIIND	36692	

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Cyclic AMP-responsive element-binding protein 1 - Homo sapiens (Human)

UniProtKB

P16220 (CREB1_HUMAN) Reviewed, UniProtKB/Swiss-Prot

Last modified September 18, 2013. Version 168. History...

Clusters with 100%, 90%, 50% identity | Documents (6) | Third-party data

Names - Attributes - General annotation - Ontologies - Interactions - All products - Sequence annotation - Sequences - References - Cross-refs - Entry Info - Customize order

Names and origin

Protein names	Recommended name: Cyclic AMP-responsive element-binding protein 1 Short name=CREB-1 Short name=cAMP-responsive element-binding protein 1
Gene names	Name CREB1
Organism	Homo sapiens (Human) [Reference proteome]
Taxonomic identifier	9606 [NCBI]
Taxonomic lineage	Eukaryota · Metazoa · Chordata · Craniata · Vertebrata · Euteleostomi · Mammalia · Eutheria · Euarchontoglires · Primates · Haplorhini · Hominoidea · Homo

Protein attributes

Sequence length	341 AA.
Sequence status	Complete.
Protein existence	Evidence at protein level

General annotation (Comments)

42

Cyclic AMP-responsive element-binding protein 1 - Homo sapiens (Human)

www.uniprot.org/uniprot/P16220

Names | Attributes | General annotation | Ontologies | Interactions | All products | Sequence annotation | Sequences | References | Cross-refs | Entry info

Gene Ontology (GO)

Biological process

- Fc epsilon receptor signaling pathway
- Traceable author statement. Source: Reactome
- MyD88-dependent toll-like receptor signaling pathway
- Traceable author statement. Source: Reactome
- Notch signaling pathway
- Traceable author statement. Source: Reactome
- TRIF-dependent toll-like receptor signaling pathway
- Traceable author statement. Source: Reactome
- Type 1 pneumocyte differentiation
- Inferred from electronic annotation. Source: Compara
- activation of phospholipase C activity
- Traceable author statement. Source: Reactome
- axon guidance
- Traceable author statement. Source: Reactome
- epidermal growth factor receptor signaling pathway
- Traceable author statement. Source: Reactome
- fibroblast growth factor receptor signaling pathway
- Traceable author statement. Source: Reactome
- innate immune response
- Traceable author statement. Source: Reactome
- lactation
- Inferred from electronic annotation. Source: Compara
- lung sacculle development
- Inferred from electronic annotation. Source: Compara
- negative regulation of transcription by competitive promoter binding
- Inferred from direct assay (C34943, I181932). Source: BHF-UCL
- neutrophil TRP receptor signaling pathway
- Traceable author statement. Source: Reactome
- phosphatidylinositol-mediated signaling
- Traceable author statement. Source: Reactome
- pituitary gland development
- Inferred from electronic annotation. Source: Compara

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www.ebi.ac.uk/QuickGO/GProteinTerm=P16220

Accession P16220

Gene CREB1

Taxonomy Homo sapiens

Description Cyclic AMP-responsive element-binding protein 1

Annotation

Displaying annotations 1 to 111 of 111 for 1 proteins

Database	Gene Product ID	Symbol	Qualifier	GO Identifier	GO Term Name	Aspect	Evidence	Reference	With	Tax
UniProtKB	P16220	CREB1		GO:0002224	toll-like receptor signaling pathway	P	TAS	Reactome REACT_6966		9606
UniProtKB	P16220	CREB1		GO:0002224	toll-like receptor signaling pathway	P	TAS	Reactome REACT_9020		9606
UniProtKB	P16220	CREB1		GO:0002255	MyD88-dependent toll-like receptor signaling pathway	P	TAS	Reactome REACT_27215		9606
UniProtKB	P16220	CREB1		GO:0002252	MyD88-dependent toll-like receptor signaling pathway	P	TAS	Reactome REACT_6788		9606
UniProtKB	P16220	CREB1		GO:0002255	MyD88-independent toll-like receptor signaling pathway	P	TAS	Reactome REACT_6782		9606
UniProtKB	P16220	CREB1		GO:0002256	MyD88-independent toll-like receptor signaling pathway	P	TAS	Reactome REACT_6809		9606
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UniProtKB	P16220	CREB1		GO:006355	regulation of transcription, DNA-dependent	P	IEA	InterPro:IPRO01630/InterPro:IPRO03102/InterPro:IPRO04827	9606	9606
UniProtKB	P16220	CREB1		GO:006355	regulation of transcription, DNA-dependent	P	IEA	Ensembl/Compara	Ensembl/ENSMUSP0000059973	9606
UniProtKB	P16220	CREB1		GO:006355	regulation of transcription, DNA-dependent	P	IEA	UniProt KeywordsGO (UniProtKB/Swiss-Prot)	UniProtKB-KW-KW-0805	9606

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PhosphoSitePlus: a resource for protein phosphorylation and other post-translational modifications

www.phosphosite.org/homeAction.do?sessionId=854D1864D1C43FC98991BAF1B1DB9

PhosphoSitePlus: a resource... CREB1 Homo sapiens P16220

PhosphoSitePlus

with grant support from

Cell Signaling TECHNOLOGIES

Home

ABOUT PHOSPHOSITE | USING PHOSPHOSITE | CURATION PROCESS | CONTACT

PhosphoSitePlus® (PSP) is an online systems biology resource providing comprehensive information and tools for the study of protein post-translational modifications (PTMs). See About PhosphoSite above for more information.

Please cite the following reference for this resource: Hornbeck PV, et al. (2012) *Nucleic Acids Res.* 2012;40(D4):78 (Epub ahead of print).

A PROTEIN MODIFICATION RESOURCE

PROTEIN OR SUBSTRATE SEARCH

Protein Name: CREB1

SEARCH

ADVANCED SEARCH AND BROWSE OPTIONS

- Protein, Sequence, or Reference Search
- Site Search
- Comparative Site Search
- Browse MS2 Data By Disease
- Browse MS2 Data By Cell Line
- Browse MS2 Data By Tissue

WHAT'S NEW

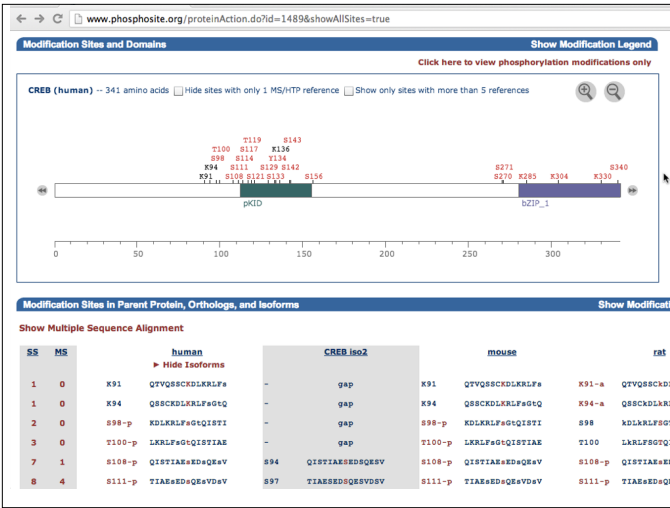
- Jul 2012 Download Datasets of Regulatory or Disease-Associated Sites.
- Dec 2011 Download "PhosphoSitePlus: a comprehensive resource..." in January 2012 issue of *Nucleic Acids Research*.
- Jul 2011 Multiple Sequence Alignment (MSA) added to the Protein Page.
- Jul 2011 Download PyMOL & Chimera Scripts from the Structure Viewer window.
- Jul 2011 New Tutorial for Navigating PhosphoSitePlus®.
- Apr 2011 Name Term Indicates links to protein-specific reagents from CST™

Phosphorylation Site Statistics

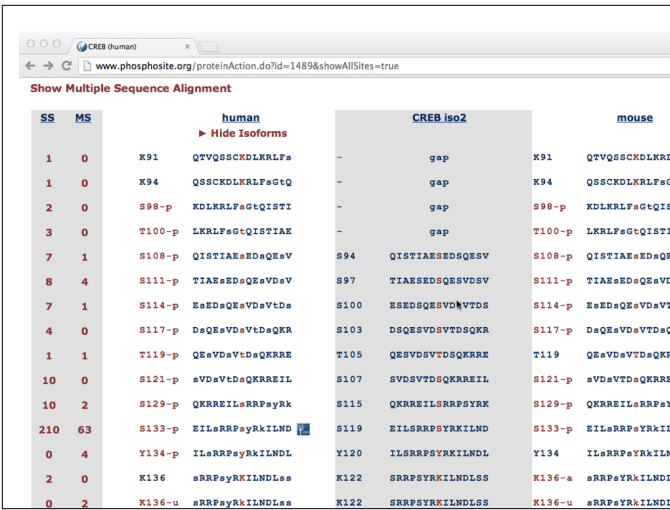
Non-redundant sites:	209,635
Non-redundant proteins:	19,846
Sites curated from literature:	109,423
All sites using site-specific (SS) methods:	12,315
All sites using discovery-mode MS (MS) methods:	100,230

© 2012 Hornbeck, Hubbs, Eick, Reed, Mende, and Roth

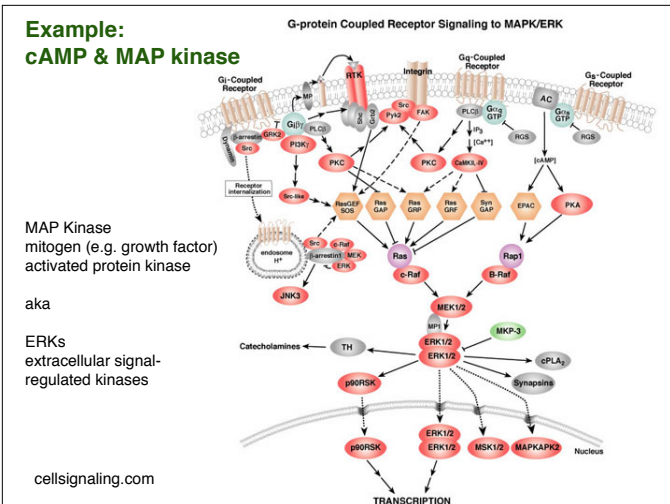
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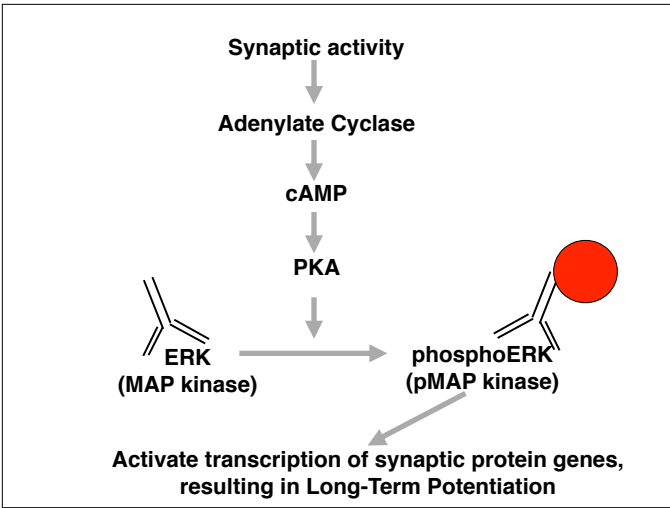


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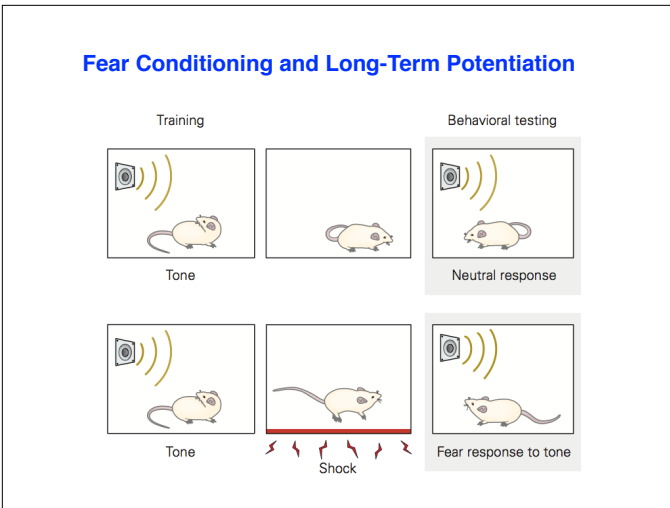


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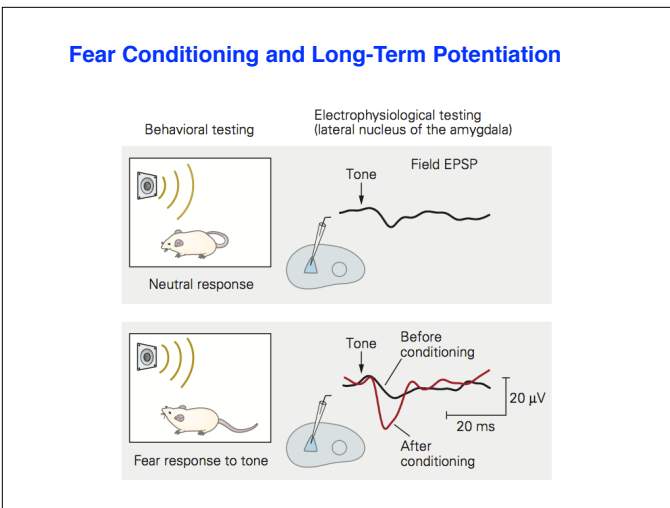
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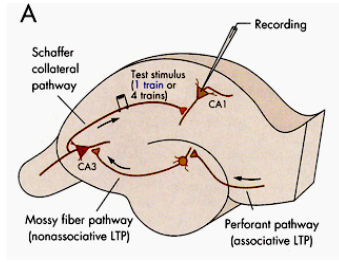


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Long-Term Potentiation (LTP) in Hippocampus

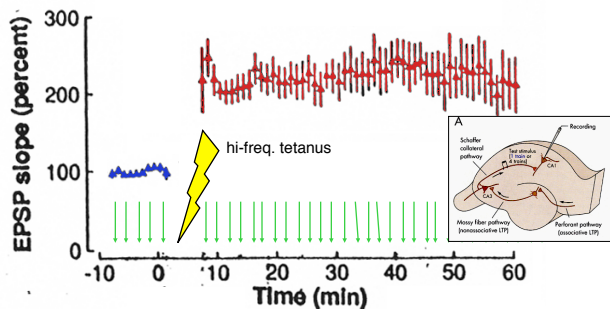
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Long-Term Potentiation (LTP) in Hippocampus

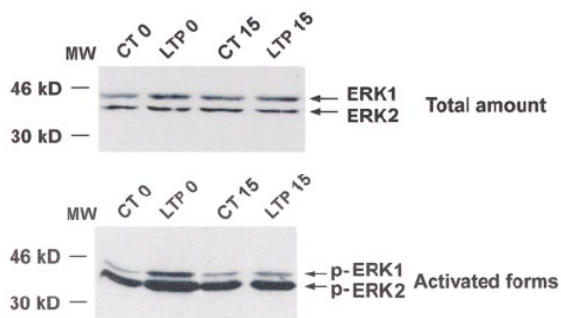
High frequency stimulus strengthens transmission at a synapse, by upregulating future transmitter release, receptor number, second messengers, etc.

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Western Blot to identify phosphoprotein

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LTP 0 = immediately after stimulation
LTP 15 = 15 min after stimulation

Davis 2000

Immunohistochemistry for phosphoMAP Kinase

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CT 0



LTP 0



LTP 15

Davis 2000

Some useful analogs to stimulate 2nd Messenger systems

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Forskolin

activates Adenyl Cyclase

permeable analogs of cAMP/cGMP

activate PKA/PKG

phorbol esters

(e.g. TPA)

activate PKC by simulating DAG

A variety of new drugs...

Note: most of these drugs cannot be used systemically.

Useful toxins to block Signaling Pathways

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G-Protein Toxins:

1. Cholera Toxin

allows binding of GTP, but prevents hydrolysis.

Causes overproduction of cAMP, leading to loss of electrolytes and water from intestinal cells.

2. Pertussis Toxin

blocks release of GDP from alpha subunit, so G-protein locked in the inactive state.

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Phosphatase Inhibitors

should enhance kinase effects by blocking removal of phosphate from substrate

Okadaic acid

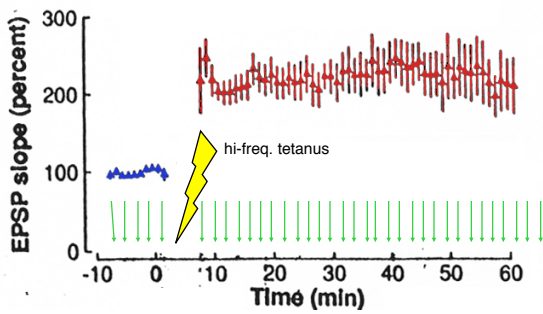
blocks PP1 and PP2a
diarrhetic shellfish toxin

Vanadate (VO_4^-)

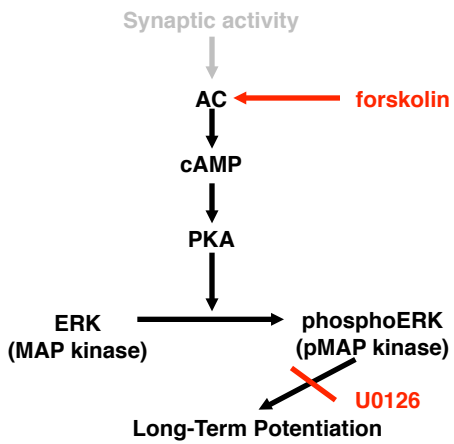
blocks tyrosine phosphatases
mimics phosphate group...

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Long-Term Potentiation (LTP) in Hippocampus



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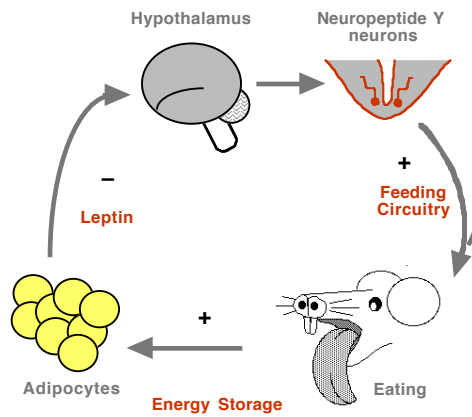
Criteria for Establishing a Signaling Pathway

- Receptor should have appropriate domains
- Kinase should be activated by the physiological stimulus
- Downstream targets of pathway should be activated by stimulus.
- Drugs that alter kinase/phosphatase activity in the pathway should alter physiological response
- Genetic manipulations should alter signaling and physiological response

Example: Leptin / Jak-Stat / SOCS pathway

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Feedback Loop: Leptin -> NPY



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Obese Mutant Mice

ob/ob obese mouse
db/db diabetic mouse
fa/fa fatty (Zucker) rat

have impaired leptin signaling

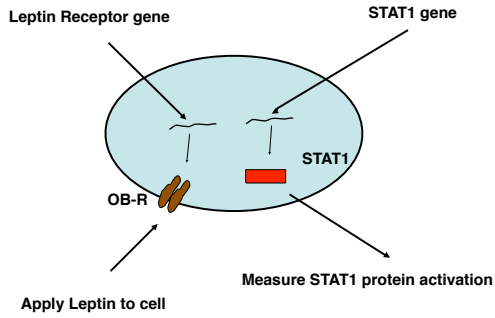
db/db mice lack functional leptin receptors



	Body mass	Adiposity
wildtype (+/+):	18 g	12 %
obese (ob/ob)	64 g	60 %

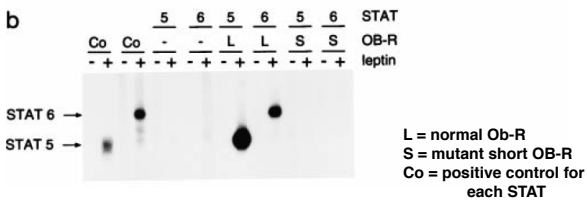
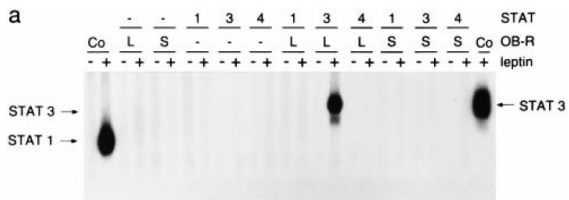
66

Activation of STAT in transfected Cell



Repeat for STAT3,4,5, and 6

Measure phosphoStat levels in presence of leptin and leptin receptor



✓ Which Stat (Stats 1-6) is coupled to Leptin Receptor?

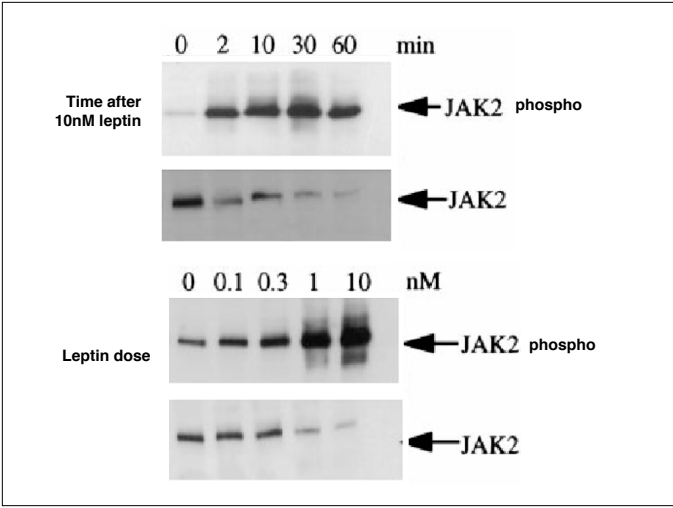
Are JAK & Stat phosphorylated?

Does Stat translocate to the nucleus?

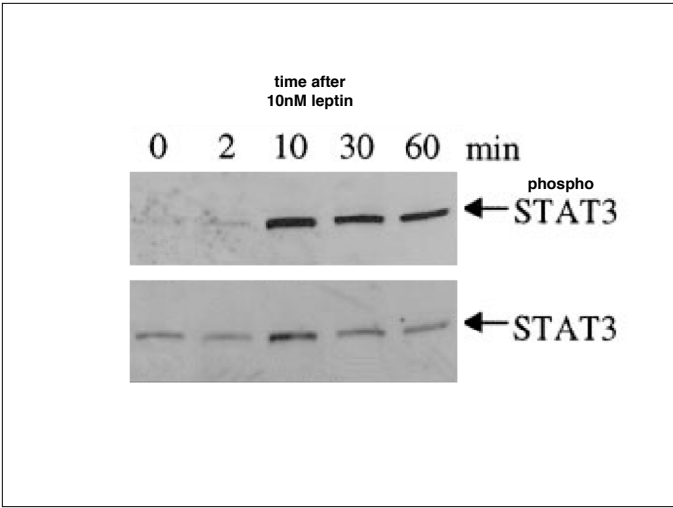
Is SOCS induced? Which SOCS (SOCS1-3)?

Does tyrosine phosphatase downregulate Leptin Receptor activity?

Is this signaling pathway absent in receptor KO mutant?



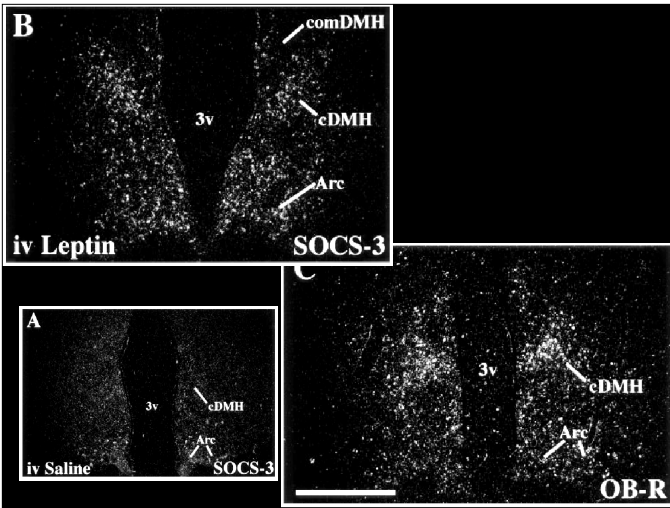
73



74

- ✓ Which Stat (Stats 1-6) is coupled to Leptin Receptor?
- ✓ Are JAK & Stat phosphorylated?
- Does Stat translocate to the nucleus?
- Is SOCS induced? Which SOCS (SOCS1-3)?
- Does tyrosine phosphatase downregulate Leptin Receptor activity?
- Is this signaling pathway absent in receptor KO mutant?

75



79

Which Stat (Stats 1-6) is coupled to Leptin Receptor?
 Are JAK & Stat phosphorylated?
 Does Stat translocate to the nucleus?
 Is SOCS induced? Which SOCS (SOCS1-3)?
 Does tyrosine phosphatase downregulate Leptin Receptor activity?
 Is this signaling pathway absent in receptor KO mutant?

80

Blockade of Tyrosine Phosphatases with Vanadate enhances leptin effect

orthovanadate	(-)	1	10	10 ²	10 ⁴	(μ M)
100 nM OB	(-)	(+)				
p-STAT3						
STAT3						

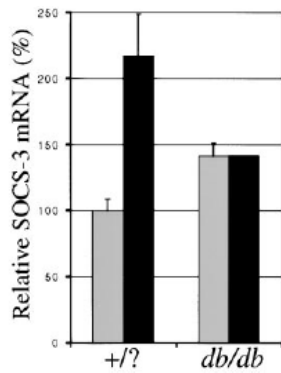
81

82

- ✓ Which Stat (Stats 1-6) is coupled to Leptin Receptor?
- ✓ Are JAK & Stat phosphorylated?
- ✓ Does Stat translocate to the nucleus?
- ✓ Is SOCS induced? Which SOCS (SOCS1-3)?
- ✓ Does tyrosine phosphatase downregulate Leptin Receptor activity?
- Is this signaling pathway absent in receptor KO mutant?

83

SOCS3 not induced in Receptor KO mutant.

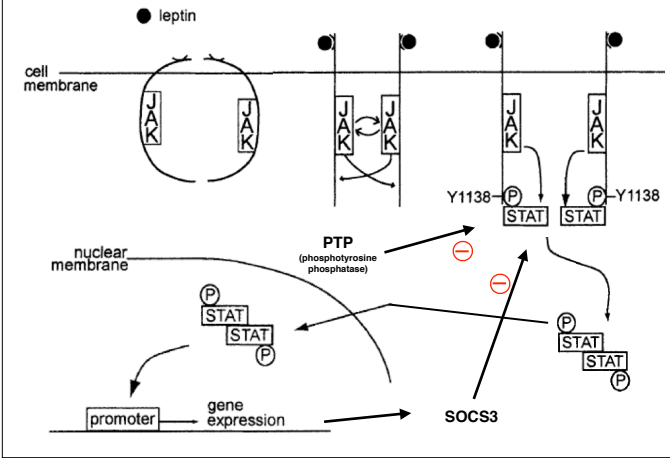


84

- ✓ Which Stat (Stats 1-6) is coupled to Leptin Receptor?
- ✓ Are JAK & Stat phosphorylated?
- ✓ Does Stat translocate to the nucleus?
- ✓ Is SOCS induced? Which SOCS (SOCS1-3)?
- ✓ Does tyrosine phosphatase downregulate Leptin Receptor activity?
- Is this signaling pathway absent in receptor KO mutant?

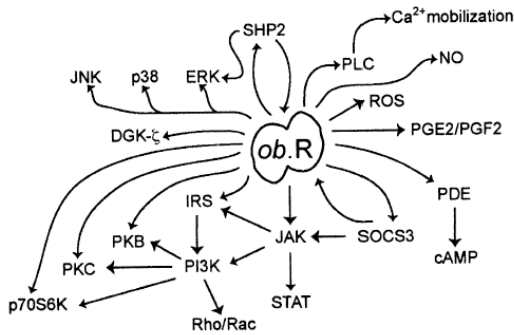
Leptin receptor (Ob-R) signaling pathway

85



Signaling pathways regulated by leptin.

86

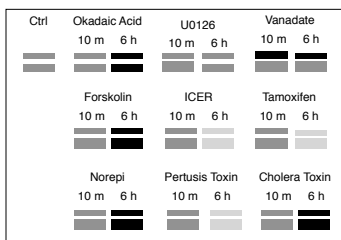


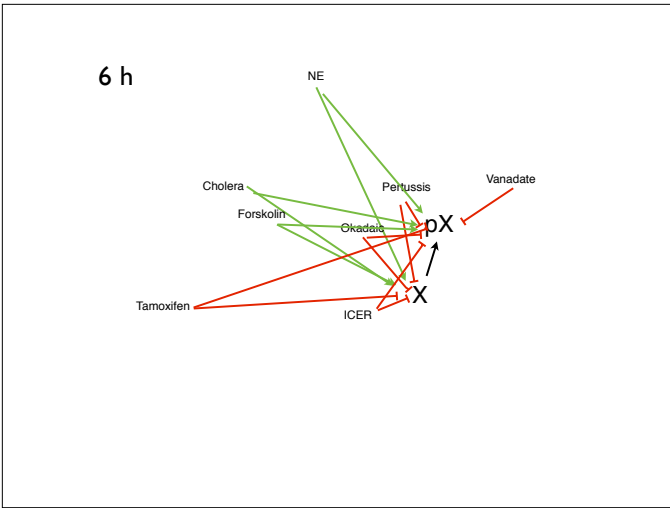
"Note that arrows (a) are intended to depict regulation rather than imply activation and (b) do not imply a direct association as in some cases multiple potential intermediates are as yet uncharacterized or excluded."

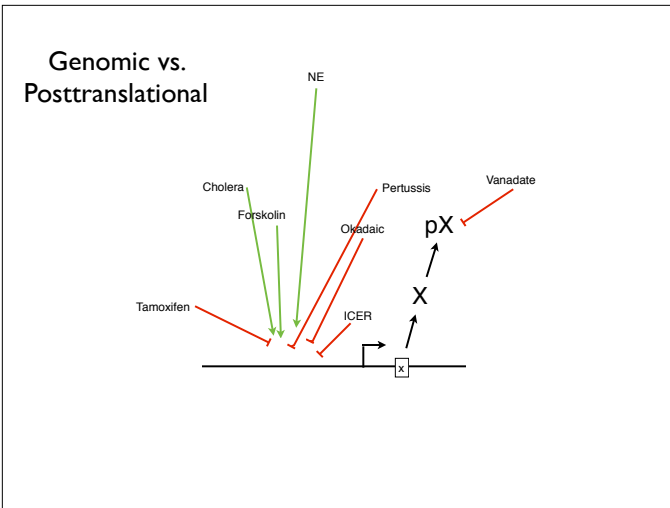
Figure 3. shows a series of Western Blots for a specific protein extracted from treated cells. The protein is visualized with a specific antibody that yields 2 bands: a major band and a slightly larger molecular weight band that contains a subset of the total protein. For each blot, the cells were treated for either 10 min or 6 h with the indicated compound. The compounds decreased or increased the intensity of one or both bands as indicated.

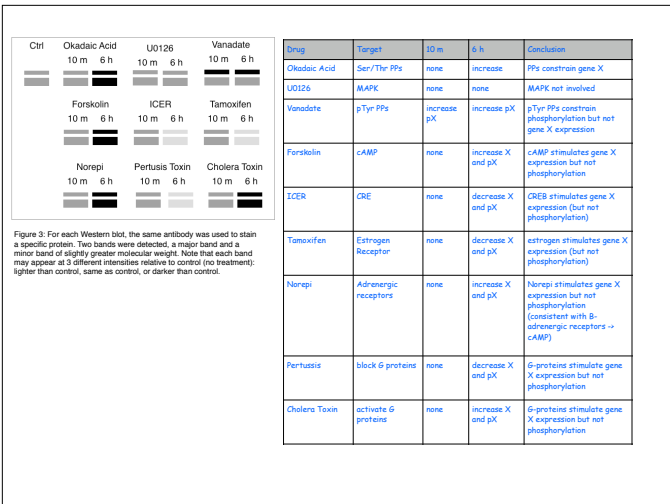
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Outline the regulation of the protein's gene expression and phosphorylation (i.e. arrows connecting receptors, enzymes, transcription factors).



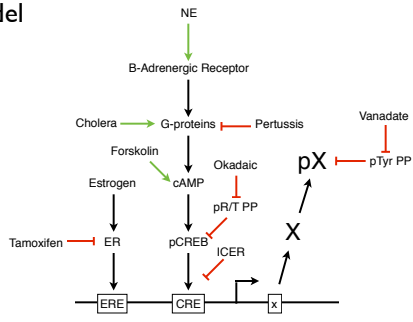






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Model

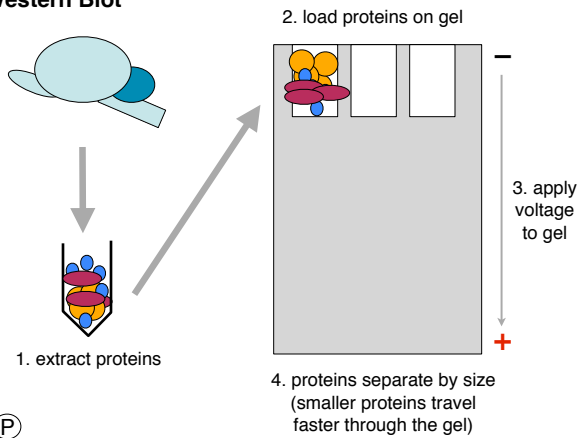


95

end

96

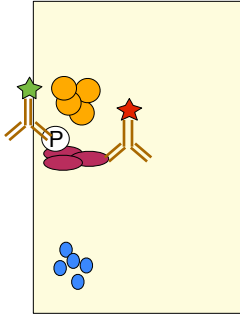
Western Blot



Western Blot

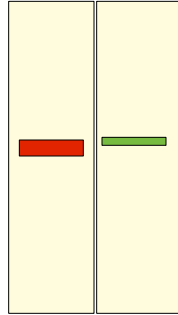
5. Blot proteins onto nylon membrane

Developed Blot



6. Visualize proteins using labeled antibodies

7. Visualize phosphorylated proteins using phospho-specific antibodies



protein phospho protein

2D Gel of Postsynaptic Density

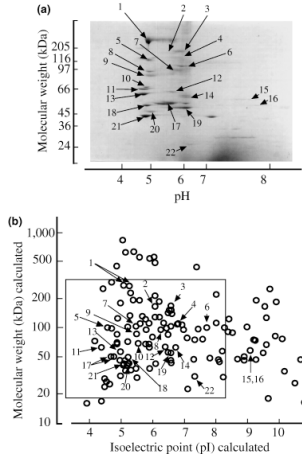


Figure 1. Proteins in the PSD fraction revealed by two-dimensional (2D)-gel electrophoresis and 2D-display. (a) 2D-gel electrophoresis of the PSD fraction. PSD protein (200 µg) was subjected to 2D-gel electrophoresis and stained with Coomassie blue. Some of the major Coomassie-stained protein spots were identified; 1, α and β fodrin; 2, NMDA-receptor 2B subunit; 3, densin 180; 4, SAPAP; 5, neurofilament M protein; 6, GluR1; 7, SAP-97; 8, α actinin; 9, PSD-95/SAP-90; 10, HSP70; 11, neurofilament L protein; 12, TOAD-64; 13, α internexin; 14, β CaM kinase II; 15, IRSp58; 16, IRSp53; 17, α and β tubulin; 18, GFAP; 19, α CaM kinase II; 20, homer 1b; 21, actin; 22, VDAC. (b) Theoretical 2D-gel representation of data assuming that identified peptides correspond to full-length proteins. PSD proteins with an identification number greater than 5 are shown. Their predicted molecular weight and pI values are indicated. The region in the box indicates the region measured in 2D-gel electrophoresis. Proteins are numbered the same as in (a).
