TYPES OF RECEPTORS

OUTLINE

Definition

➢ History

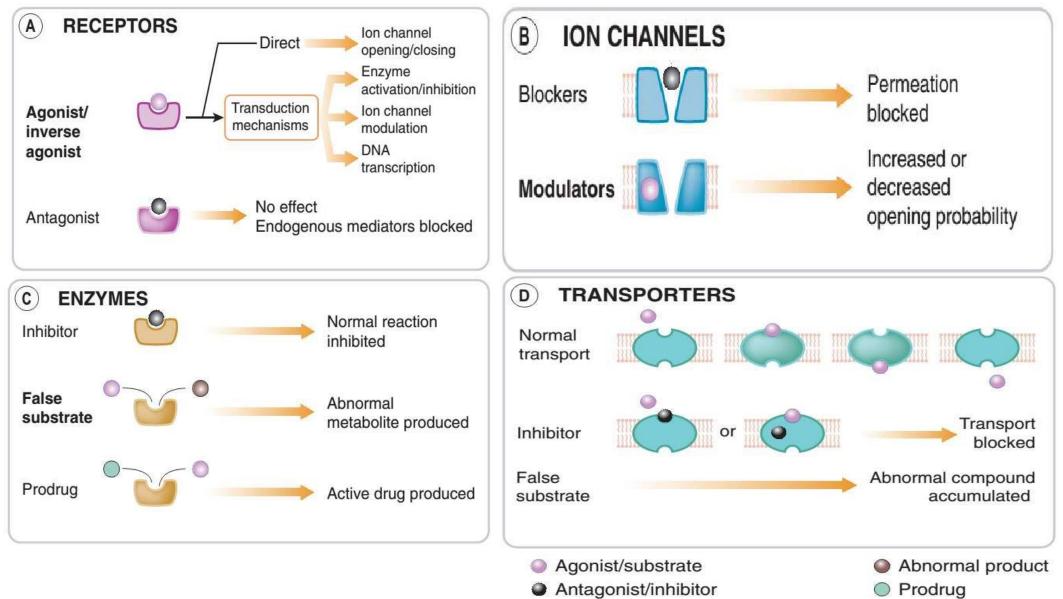
- ► Receptor Theories
- Criteria for classifying receptors

➤ Types of Receptors :

- Ligand Gated Ion Channels
- G-Protein Coupled Receptors
- Enzyme Linked Receptors
- Nuclear Receptors

➤Summary

TARGETS FOR DRUG ACTION



What are receptors?

Receptor is a cellular macromolecule, whose function is to recognize and respond to chemical signals

HISTORY

➤<u>1878</u> – John Langley proposed 'receptive substance'

1905 - Receptive substance on surface of skeletal muscle mediate drug action. Different in different species

Paul Ehrlich designated 'receptor' to be anchoring group of the protoplasmic molecule for the administered compound

Envisaged molecules extending from cells that the body could use to distinguish and mount an immune response to foreign objects

1948 - Ahlquist showed the differential action of adrenaline & demonstrated its effects on two distinct receptor populations & the theory of receptor-mediated drug interactions gained acceptance

<u>1970s</u> - Pharmacology entered a new phase following the development of receptor-labelling techniques

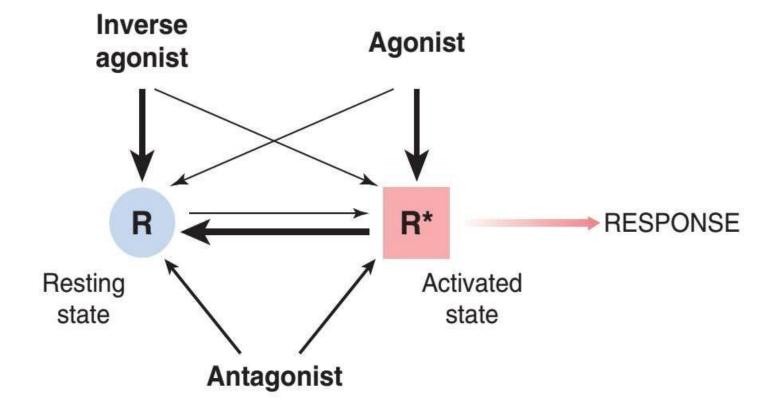
Multiple subtypes of receptors now known, which has paved the way for clinically superior drugs

CRITERIA FOR CLASSIFYING RECEPTORS

Pharmacological Criteria
 Tissue Distribution
 Ligand Binding
 Transducer Pathway
 Molecular Cloning

RECEPTOR THEORIES

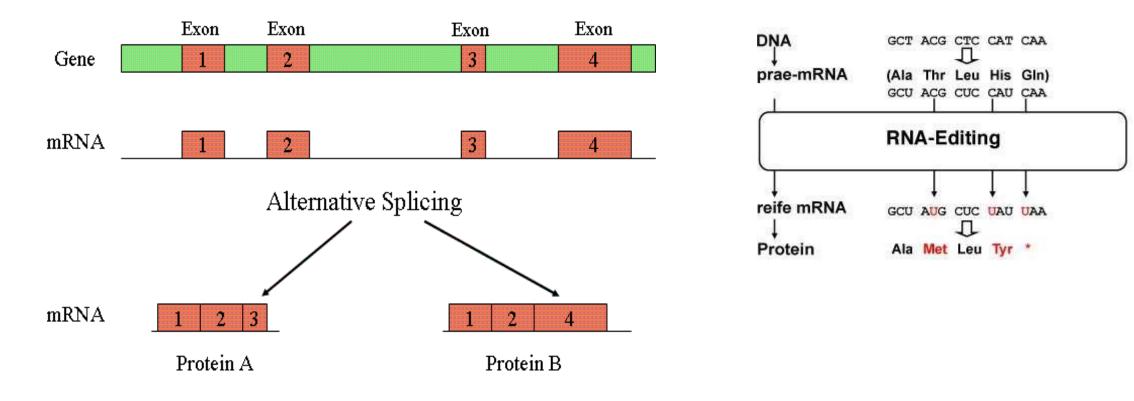
- Occupation theory (1937)
- Rate theory (1961)
- Two State Receptor Theory (1983)



TYPES OF RECEPTORS

- Ligand Gated Ion Channels
- ► G-Protein Coupled receptors
- Enzyme Linked receptors
- > Nuclear receptors

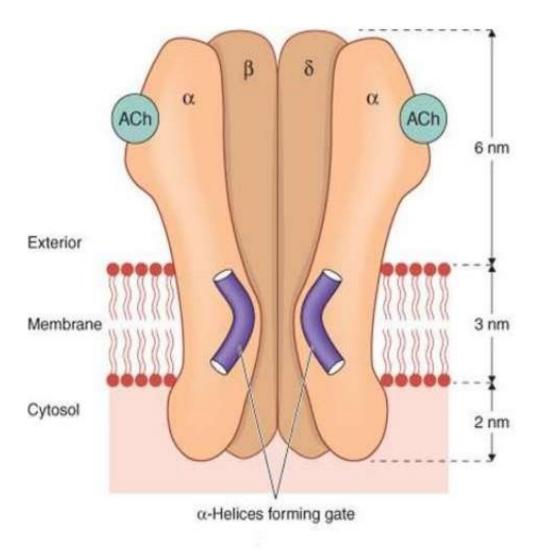
RECEPTOR SUBTYPES & NOMENCLATURE

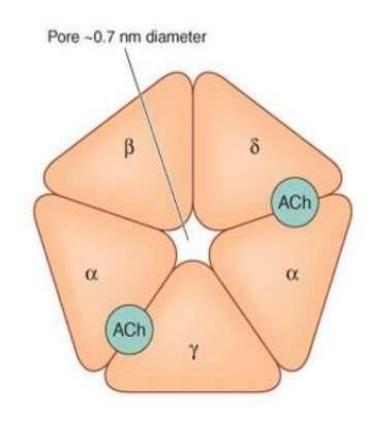


<u>IUPHAR-</u> International Union of Basic and Applied Pharmacology

TYPE 1 : LIGAND GATED ION CHANNELS

- Ionotropic Receptors
- Typically receptors on which neurotransmitters act
- Timescale : Milliseconds
- Localization : Membrane
- Effector : Ion Channel
- Coupling : Direct
- Examples : Nicotinic Ach Receptor, GABAA Receptor, Glutamate Receptor, Glycine receptor, 5 Hydroxytryptamine type 3 (5 – HT₃)





MOLECULAR STRUCTURE

➢ Nicotinic Ach receptor studied in great detail

 \geq Pentameric Assembly of 4 types of subunits α , β , γ and δ

>4 membrane spanning α -helices inserted into membrane

➤2 Ach binding sites, both must bind Ach molecules for receptor activation

Lining of central transmembrane pore formed by helical segments of each subunit (negatively charged AA). 5 helices sharply kinked inwards halfway, forming a constriction

GATING MECHANISM

Ach molecules bind, twists the α subunits, kinked helices either straighten out or swing out of the way, opening channel pore

Conductance produced by different Ach like agonists is the same whereas mean channel lifetime varies

Mutation of a critical residue in helix changes channel from being cation selective (excitatory) to being anion selective (typical of receptors for inhibitory transmitters like GABA) > Most excitatory neurotransmitters cause

Increase in Na+ and K+ permeability

net inward current carried mainly by Na+

Depolarization of the membrane (probability to generate action potential)

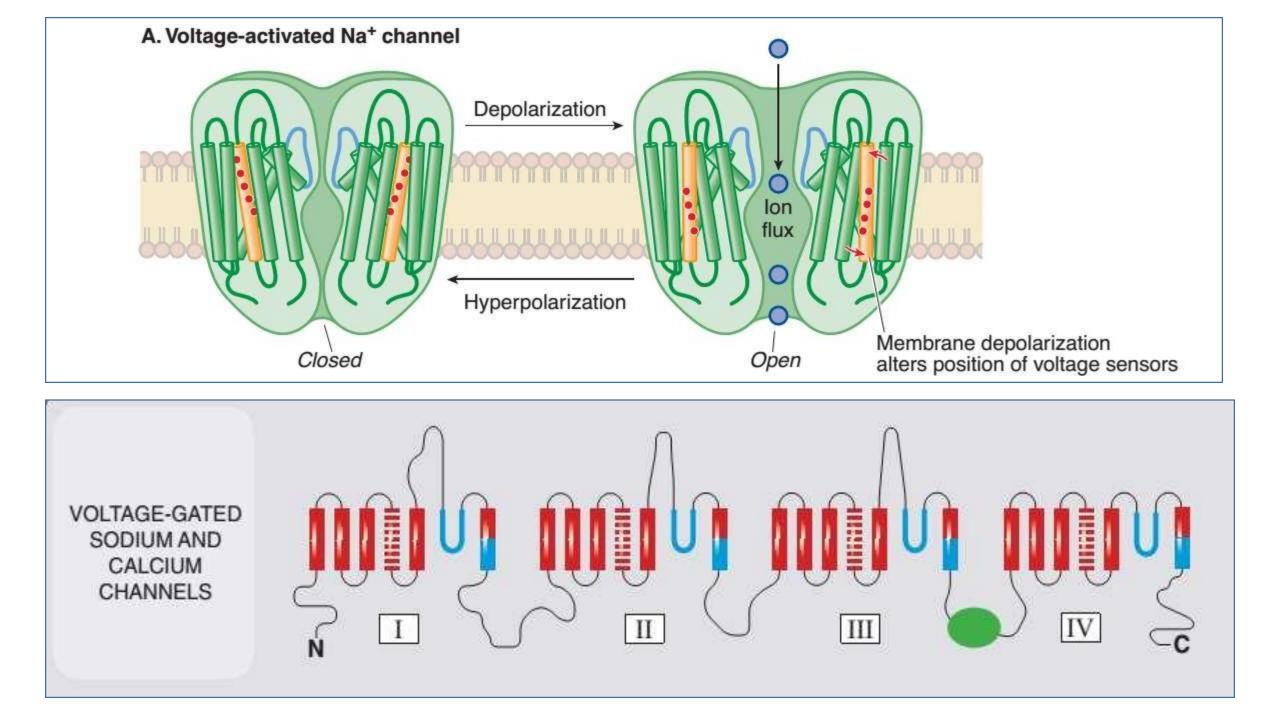
➢Speed of this response implies that coupling between the receptor and ionic channel is a direct one (no intermediate biochemical steps involved)

VOLTAGE OPERATED CHANNELS

These channels open when the cell membrane is depolarised. They underlie the mechanism of membrane excitability

Activation induced by membrane depolarisation is short lasting, even if the depolarisation is maintained

The most important channels in this group are selective sodium, potassium or calcium channels

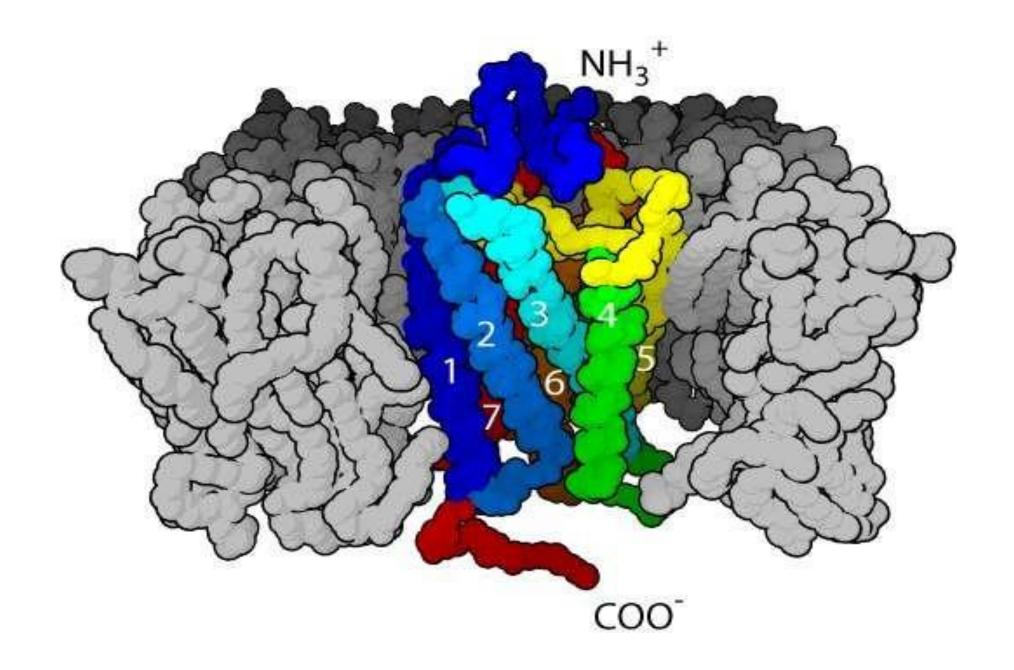


Ligand Gated receptors v/s VOCs

- ROCs appear to assume only two states whereas VOCs undergo a third state called refractory (inactivated) state.
- Voltage gated channels have no major endogenous modulator (like Ach)

TYPE 2 : G – PROTEIN – COUPLED RECEPTORS

- Alfred Goodman Gilman & Martin Rodbell (1994)
- Metabotropic or 7 Transmembrane/ Heptahelical receptors
- Largest family
- Timescale : Seconds
- Location : Membrane
- Effector : Channel or Enzyme
- Coupling : G- Protein
- Examples : adrenoceptors, Muscarinic Ach, histamine, serotonin, opioid, cannabinoid, amine, peptide, prostanoid receptors



MOLECULAR STRUCTURE

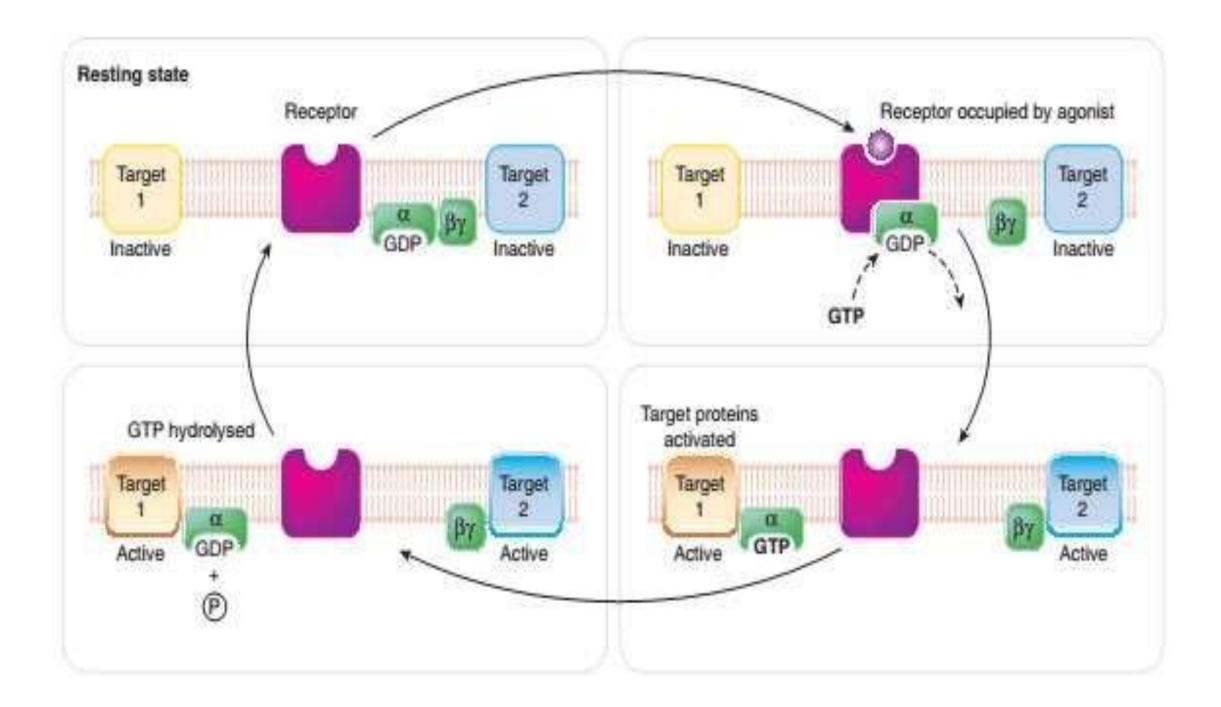
- Single polypeptide chain 1100 residues. 7 Transmembrane α -helices, an extracellular N-terminal domain and intracellular C-terminal domain
- 3rd cytoplasmic loop couples to the G- Protein

Family	Receptors ^b	Structural features
A: rhodopsin family	The largest group. Receptors for most amine neurotransmitters, many neuropeptides, purines, prostanoids, cannabinoids, etc.	Short extracellular (N terminal) tail. Ligand binds to transmembrane helices (amines) or to extracellula loops (peptides)
B: secretin/ glucagon receptor family	Receptors for peptide hormones, including secretin, glucagon, calcitonin	Intermediate extracellular tail incorporating ligand-binding domain
C: metabotropic glutamate receptor/ calcium sensor family	Small group Metabotropic glutamate receptors, GABA _B receptors, Ca ²⁺ -sensing receptors	Long extracellula tail incorporating ligand-binding domain

G-PROTEINS AND THEIR ROLE

- Family of membrane-resident proteins whose function is to recognize activated GPCRs and pass on the message to the effector systems that generate a cellular response
- Function of G-Protein:
- 3 subunits (α , β , γ) are anchored to the membrane through attached lipid residues

Coupling of the α subunit to an agonist-occupied receptor causes bound GDP to exchange with intracellular GTP; α –GTP complex dissociates from receptor and from $\beta\gamma$ complex



 Amplification : A single agonist—receptor complex can activate several Gprotein molecules, each of these can remain associated with the effector enzyme for long enough to produce many molecules of product (Second messenger)

• Four main classes of G-protein (Gs, Gi, Go and Gq) show selectivity with respect to both the receptors and the effectors with which they couple

• Cholera toxin and pertussis toxin

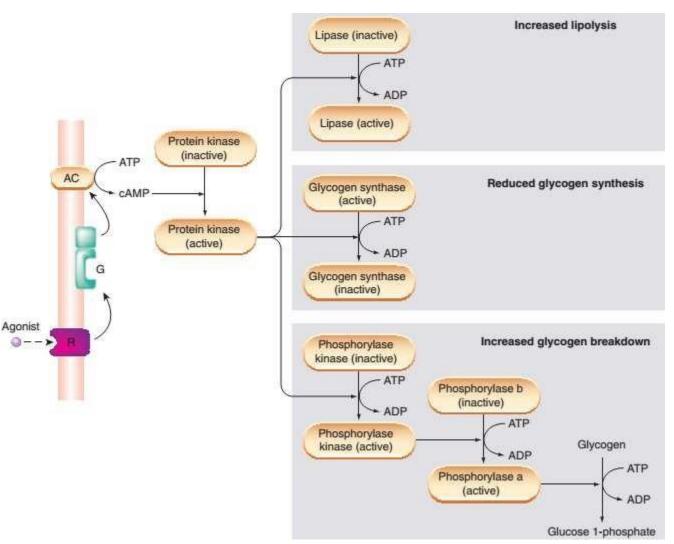
Subtypes	Associated receptors	Main effectors			
Ga subunits					
Gαs	Many amine and other receptors (e.g. catecholamines, histamine, serotonin)	Stimulates adenylyl cyclase, causing increased cAMP formation			
Gα _i	As for Gα _s , also opioid, cannabinoid receptors	Inhibits adenylyl cyclase, decreasing cAMP formation			
Gα₀	As for Gα _s , also opioid, cannabinoid receptors	?Limited effects of α subunit (effects mainly due to βγ subunits)			
Gαq	Amine, peptide and prostanoid receptors	Activates phospholipase C, increasing production of second messengers inositol trisphosphate and diacylglycerol			
Gβγ subunits	All GPCRs	 As for Gα subunits (see above). Also: activate potassium channels inhibit voltage-gated calcium channels activate GPCR kinases (GRKs, activate mitogen-activated protein kinase cascade 			

Targets for G-Proteins

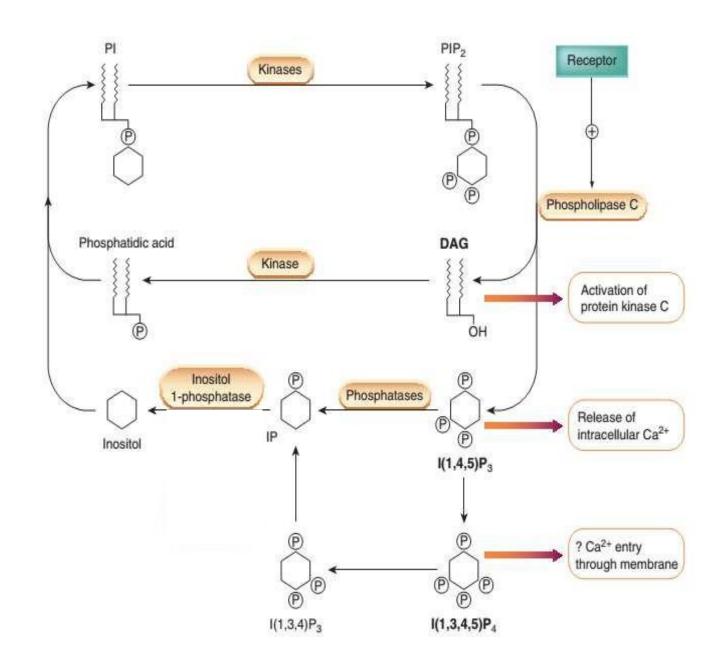
Adenylate cyclase

 catalyses formation of the intracellular messenger cAMP

 cAMP activates various protein kinases that control cell function in many different ways by causing phosphorylation of various enzymes, carriers & other proteins



- Phospholipase C/IP3/DAG
 - catalyzes the formation of IP3 and DAG from membrane phospholipid
 - IP3 increases free cytosolic Ca2+ (releasing Ca2+ from intracellular compartments) which initiates many events
 - DAG activates protein kinase C, which controls many cellular functions by phosphorylating proteins



> Ion Channels appears to be general mechanism for controlling K⁺and Ca⁺⁺ channels by direct interaction between the $\beta\gamma$ -subunit of G₀ and the channel

➢ Phospholipase A₂ (formation of arachidonic acid and eicosanoids)

The Rho/Rho kinase system G_{12/13} type G-protein. α subunit interacts with guanosine nucleotide exchange factor, which facilitates GDP–GTP exchange at another GTPase, Rho. On exchange Rho activated & activates Rho kinase phosphorylates substrate proteins

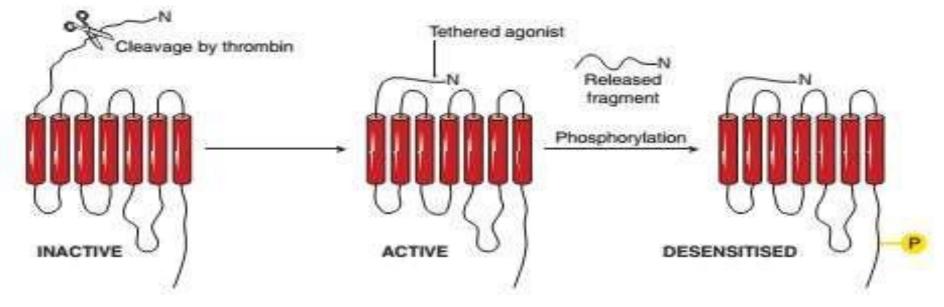
The MAP kinase system activated by cytokines and growth factors acting on kinase-linked receptors and by GPCR ligands. Controls processes involved in cell division, apoptosis and tissue regeneration

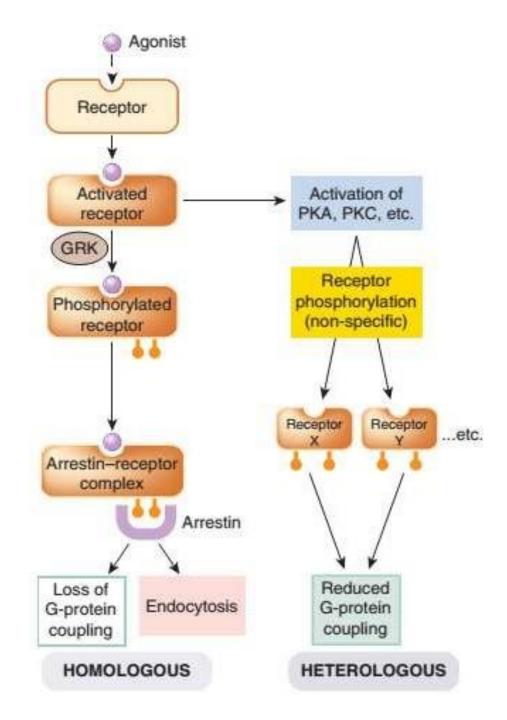
Protease Activated Receptors

End of extracellular N-terminal tail of the receptor snipped off to expose 5-6 N-terminal residues that bind to receptor domains in extracellular loops, functioning as 'tethered agonist'

A PAR molecule can be activated only once

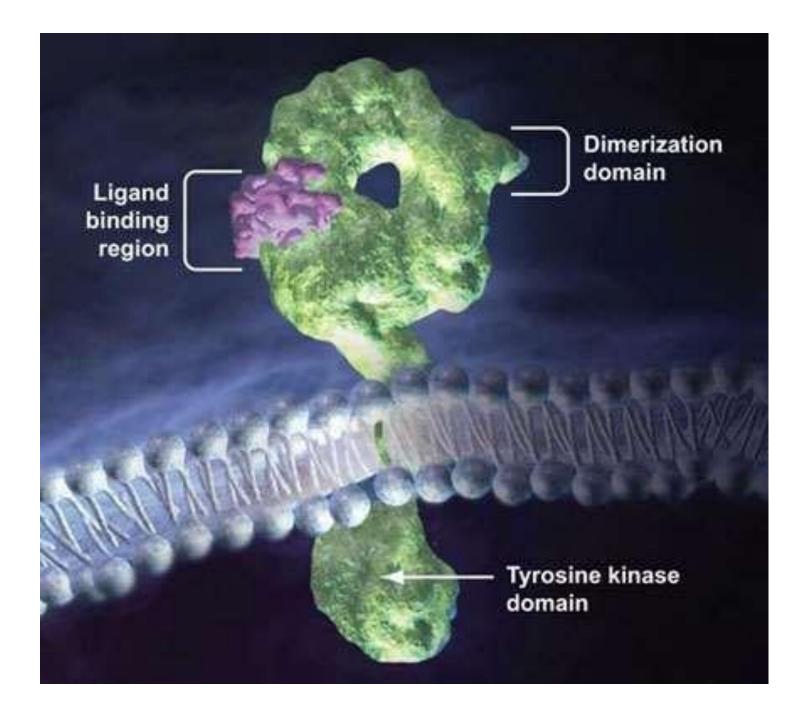
Inactivation by further proteolytic cleavage that frees tethered ligand, or by desensitization





TYPE 3 : KINASE LINKED AND RELATED RECEPTORS

- Large, heterogenous group responding mainly to protein mediators.
- Timescale : Hours
- Location : Membrane
- Effector : Protein Kinases
- Coupling : Direct
- Examples : Insulin, Growth Factors, Cytokine, ANF receptors



MOLECULAR STRUCTURE AND TYPES

Large proteins - single chain ~ 1000 residues, single membrane-spanning helical region, with a large extracellular ligand-binding domain, and an intracellular domain of variable size and function

Types –

- <u>Receptor tyrosine kinases (RTKs</u>)
- <u>Serine/threonine kinases</u>

• <u>Cytokine receptors</u>. lack intrinsic enzyme activity. When occupied, they associate with and activate, a cytosolic tyrosine kinase, such as Jak (the Janus kinase)

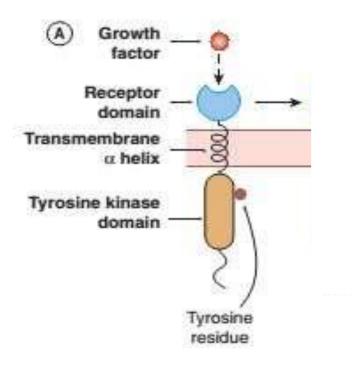
SIGNAL TRANSDUCTION

Generally involves dimerization autophosphorylation of tyrosine residues, act as acceptors for the SH2 domains of intracellular proteins

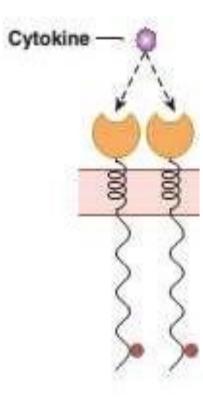
Involved mainly in events controlling cell growth and differentiation, and act indirectly by regulating gene transcription

> Two important pathways are:

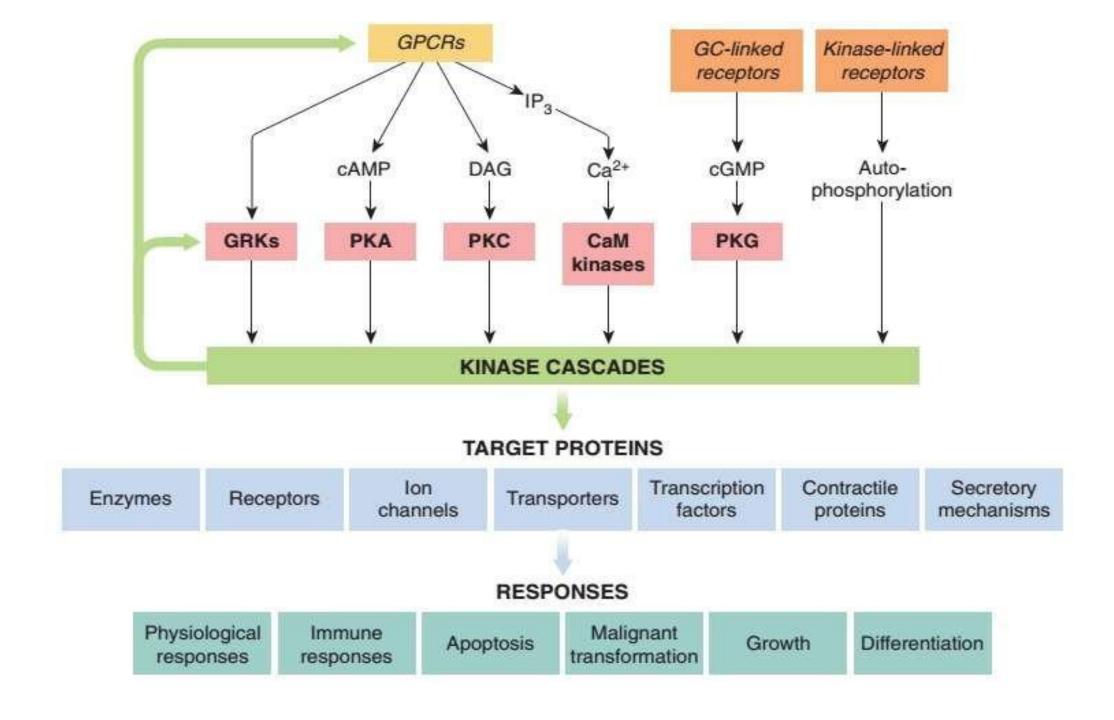
the Ras/Raf/ MAP kinase pathway - cell division, growth and differentiation.
 the Jak/Stat pathway activated by many cytokines - controls the synthesis and release of many inflammatory mediators







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TYPE 4 : NUCLEAR RECEPTORS

- Regulate gene transcription. ?Misnomer
- Timescale : Hours
- Location : Intracellular
- Effector : Gene transcription
- Coupling : Via DNA
- Examples : Steroid Hormones, Thyroid Hormones, Retinoic acid and Vitamin D receptors
- This family includes a great many (40%) orphan receptors

MOLECULAR STRUCTURE

AF1			AF2	16
N-terminus AF1 Co-activator region	Core DNA binding domain	Hinge region	Ligand–binding domain AF2 Co-activator region HSP binding	C-terminal extension

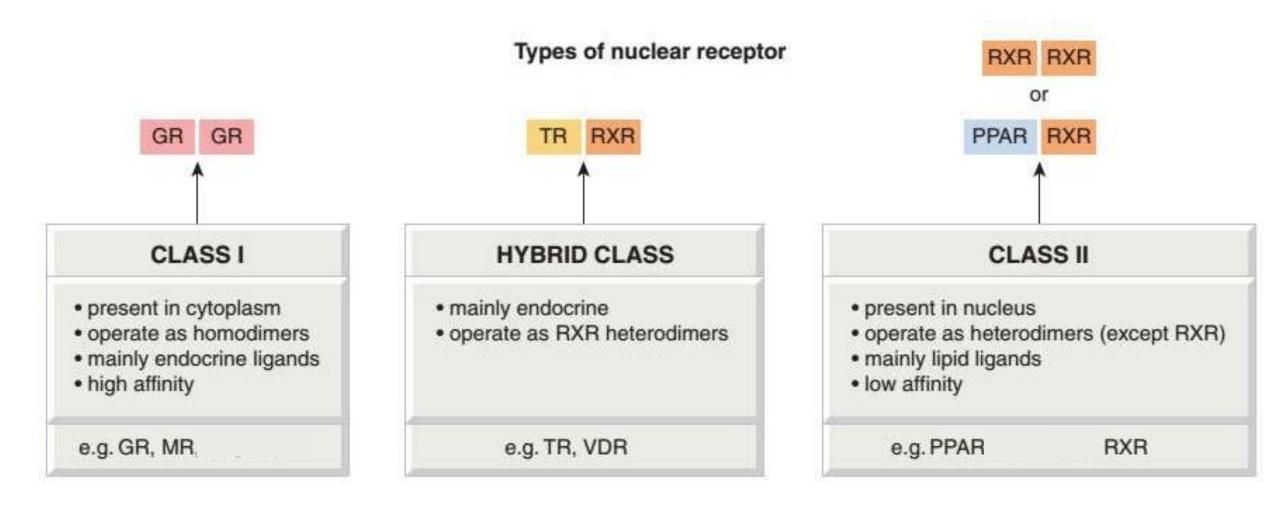
The N-terminal domain harbours the AF1 site that binds to other cell specific transcription factors and modifies the binding or activity of the receptor itself

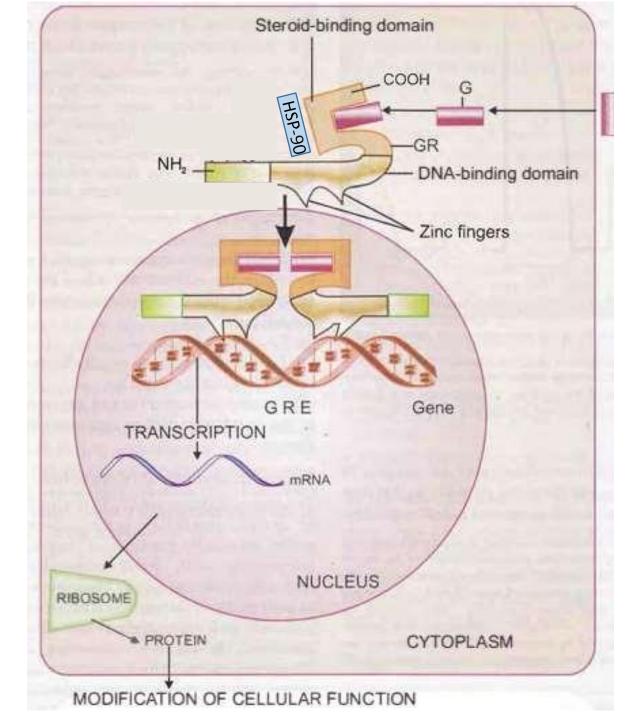
➢ The Core domain consists of the structure responsible for DNA recognition and binding. They bind to the hormone response elements located in genes to regulate them

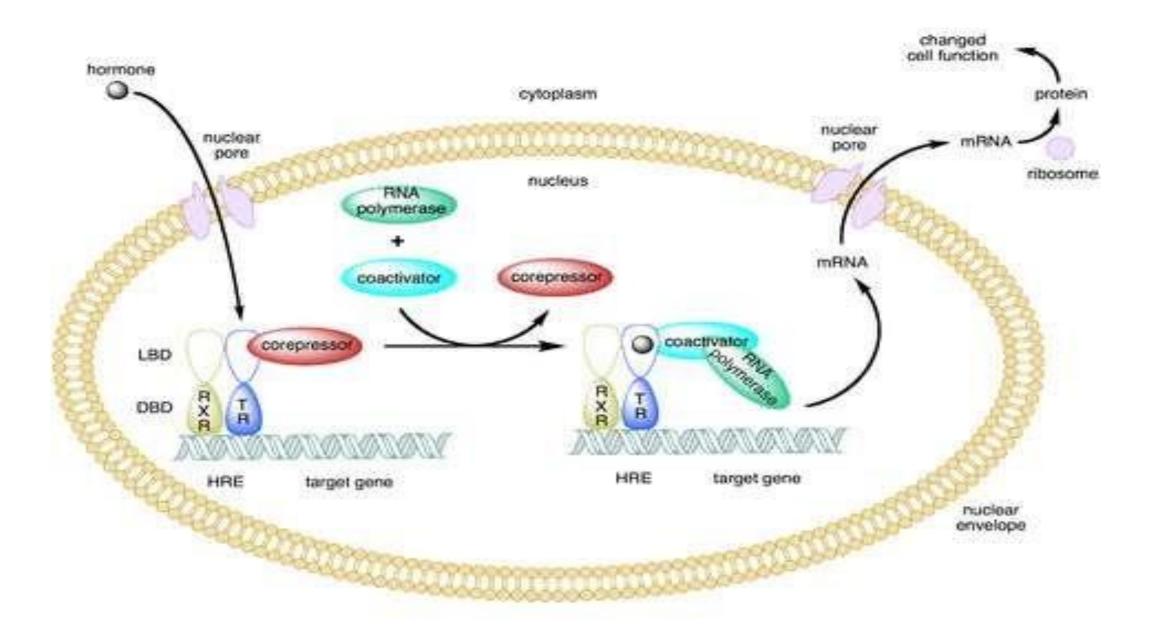
Hinge region in the molecule allows it to dimerise with other NRs and also to exhibit DNA binding

C-terminal domain contains the ligand binding module and is specific to each class of receptor

CLASSIFICATION OF NUCLEAR RECEPTORS



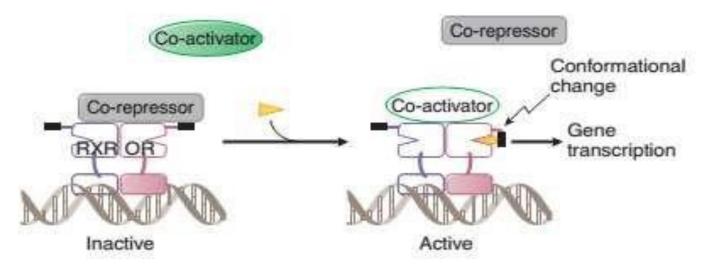




CONTROL OF GENE TRANSCRIPTION

Hormone Response Elements : Short (4-5 bp) sequences of DNA to which the NRs bind to modify gene transcription, present symmetrically in pairs.

Recruits co-activators or co-repressors to modify



This receptor family responsible for the pharmacology of approximately 10% prescription drugs

SUMMARY

