كلية الطب البيطري – جامعة تكريت ماجستير أدوية

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Advanced Pharmacology

Pharmacokinetic Principles

PHARMACODYNAMICS د جسام الدين سالم النجار

How Drugs Act Targets for drug Action

Protein Targets For Drug Binding:

- 1. Regulatory Proteins
- 2. Structural Proteins





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Regulatory Proteins:

A. Receptors:

Are macromolecular proteins act as recognition sites for drugs (agonist or antagonist). They are functionally silent in the absence of the drug.

- B. Enzymes:
- **1. Competitive inhibitor:**
- **a. Reversible**: neostigmine inhibits acetylcholinesterase, carbidopa inhibits dopa decarboxylase
- b. Irreversible: aspirin inhibits COX
- **2. False substrate** → abnormal product
 (fluorouracil)
- **3. Prodrug**: A parent compound lacks activity & needs enzymatic degradation to convert into the active form (cortisone & enalapril).

C. Carriers:

Transport of ions & organic molecules across cell membrane requires carriers.

- Loop diuretics block Na/K/2Cl co-transporter
- TCA & cocaine block N.A carrier (uptake1)
- cardiac glycosides block Na+ / K+ pump
- Omeprazole blocks proton pump.

D. Ion channels:

- 1. Ligand gated ion channel: gating is controlled by ligand binding
- 2. Voltage-gated ion channel: controlled by membrane potential.
- Drug-channel binding:
- Direct: either <u>Blockers</u> (e.g local anesthetics block voltage-gated Na+ channel) or <u>Modulators</u> where the drug binds to an accessory site of the channel affecting gating (e.g. Ca²⁺ channel blockers inhibit opening of Ca²⁺ channel)
- **b.** Indirect: involving G-protein

- 2. Structural proteins:
- Colchicine interacts with tubulin.
- Ciclosporin acts on immunophilins.
- Therapeutic antibodies act against cytokines ,e.g. infliximab (anti-TNF-α antibody)
- **Exceptions:**
- Chemotherapeutic drugs: antimicrobial agents anticancers (interact directly with DNA).
- Some drugs produce their effect without binding to any cellular components, e.g. antacids, chelating drugs, osmotic diuretics & bulk laxatives.

Drug-Receptor interaction Full & Partial agonist & antagonist

- Drug-Receptor binding may or may not result in receptor activation → response.
- Occupation → affinity (tendency of the drug to bind the receptor)
- Activation → efficacy (ability of a drug, once bound, to initiate changes → effect.



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- Agonist: a drug that binds to the receptor" affinity" → activation of the receptor" efficacy".
- a. Full agonist: possesses ↑ affinity & efficacy (+1). Large % of receptors reside in (R*) → maximal tissue response
- b. Partial agonist: possesses ↑ affinity & intermediate efficacy (0-1) i.e.↓ no. of receptors are activated even at 100% occupancy → submaximal tissue response.

They have low intrinsic activity (act as an agonist, if no full agonist is present, or as antagonist if full agonist is present, e.g. Pindolol

 Antagonist: a drug that binds to the receptor " affinity" without causing activation " zero efficacy" i.e. equal affinity for (R) & (R*).

Receptor Families & Signaling Mechanisms

- Receptors link to a variety of cellular components (enzymes, ion channels...etc).
- The operation of this linkage is known as transduction mechanism.
- Four families of receptors are distinguished on the basis of their signaling mechanisms:



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- 1. Ion channel-linked receptor (inotropic):
- Transmembrane receptor coupled directly to an ion channel, involved in fast neurotransmission.
- Transduction: millisecond.
- Ex. nAch, GABA_A, 5HT3 & glutamate receptors.
- nAch receptor: consists of 5 protein subunits $(2\alpha,\beta,\Upsilon,\delta)$.
- Each subunit consists of polypeptide chain crosses the membrane 4 times (4 transmembrane helices).
- Gating mechanism: requires 2 molecules of Ach, conformational change, transient opening of the channel → Na+ influx.



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- **2.** G-protein coupled receptor (GPCR) (metabotropic):
- Transmembrane receptor coupled to intracellular effector systems via G-protein.
- Transduction: seconds. e.g. mAch, adrenergic, opiates, chemokines & dopamine receptors also odorant & visual receptors.
- Structure: A single polypeptide chain, comprises 7 transmembrane spanning segments (serpentine or heptahelical receptor).

STRUCTURE OF G-PROTEIN LINKED RECEPTOR



- **G-protein:** consists of $(\alpha, \beta, \Upsilon)$ trimer, with GDP or GTP binds α subunit. It interact with the third intracellular loop of the receptor.
- The family of G-proteins contain several subfamilies (with 个 selectivity for a particular set of receptors and specific group of effectors) e.g. Gs, Gi, Go & Gq.

TABLE 2–1G proteins and their receptors and effectors.

G Protein	Receptors for	Effector/Signaling Pathway
G _s	β -Adrenergic amines, glucagon, histamine, serotonin, and many other hormones	↑ Adenylyl cyclase \rightarrow ↑ cAMP
G _{i1} , G _{i2} , G _{i3}	α_2 - Adrenergic amines, acetylcholine (muscarinic), opioids, serotonin, and many others	Several, including: \downarrow Adenylyl cyclase $\rightarrow \downarrow$ cAMP Open cardiac K ⁺ channels $\rightarrow \downarrow$ heart rate
G _{olf}	Odorants (olfactory epithelium)	↑ Adenylyl cyclase \rightarrow ↑ cAMP
Go	Neurotransmitters in brain (not yet specifically identified)	Not yet clear
Gq	Acetylcholine (muscarinic), bombesin, serotonin (5-HT _{1C}), and many others	\uparrow Phospholipase C \rightarrow \uparrow IP_3, diacylglycerol, cytoplasmic Ca^{2+}
G_{t1}, G_{t2}	Photons (rhodopsin and color opsins in retinal rod and cone cells)	↑ cGMP phosphodiesterase $\rightarrow \downarrow$ cGMP (phototransduction)

cAMP, cyclic adenosine monophosphate; cGMP, cyclic guanosine monophosphate.