

Protein Binding

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| Plasma Protein Binding | <ul style="list-style-type: none"> • Almost all drugs are bound to plasma proteins to some extent • Drug-protein complexes are large & do not readily cross membranes • ONLY unbound/free drug can diffuse through capillary walls, produce pharmacologic effect, be metabolized & excreted • Plasma protein binding profile of drug determines extent of its drug distribution & elimination rate • Drugs which compete for same plasma protein binding site may displace each other (drug-drug interaction) | | |
| Reversible & Irreversible Binding | <ul style="list-style-type: none"> • Most drugs reversibly bind to proteins via electrostatic forces Free Drug + Free Protein \rightleftharpoons Drug-Protein Complex • To maintain equilibrium between free & bound drug, if a free component goes away, a bound component will be released • Irreversible binding is less common & results in drug inactivation (e.g. cisplatin) | | |
| Examples of Plasma Proteins | Albumin (quantitatively most important) | <ul style="list-style-type: none"> • Generally acidic drugs bind more avidly to albumin • Low albumin levels (called hypoalbuminemia) = \uparrowFree Fraction of Drug in body | |
| | Main Drug Binding Sites on Albumin | Site I | <ul style="list-style-type: none"> • Phenytoin, • Sulphonamides • NSAIDs • Valproate |
| | | Site II | <ul style="list-style-type: none"> • Penicillins • Benzodiazepines • Probenecid |
| | α_1-acid-glycoprotein (reactive protein) | <ul style="list-style-type: none"> • Binding to this protein is quantitatively less important since conc. are typically 100X less than that of albumin • Reactive protein that may increase several-fold in presence of acute inflammation/stress (e.g. myocardial infarction) • Mostly binds basic drugs (e.g. lignocaine) • Elevated conc. of α_1-acid-glycoprotein = \downarrowFree Fraction of Drug in body | |
| Lipoproteins | | | |
| Specific protein carriers (e.g. thyroxine binding globulin, cortisol binding globulin) | | | |
| Free Conc. & Free Fraction | <ul style="list-style-type: none"> • In most cases, drug conc. at therapeutic doses are well below those of binding proteins & free drug fraction is constant | | |
| | Total Drug Conc. (C_t) = $C_f + C_b$ Free Drug Conc. (C_f) = $C_t * (1 - \%Protein\ Bound/100)$ Free Drug Fraction = C_f/C_t | | |
| | E.g. Drug A 80% Protein Bound | 100mg Dose Total drug conc. = 10 mg/L Free drug conc. = 2 mg/L Free drug fraction = 0.20 | 200mg Dose Total drug conc. = 20 mg/L Free drug conc. = 4 mg/L Free drug fraction = 0.20 |

Situations that can bring about a change in free drug fraction

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| Change in no. of plasma protein binding sites | ↓ Plasma Proteins | <ul style="list-style-type: none"> • ↓ No. of Binding Sites = ↑ Free Fraction of Drug |
| | Causes of ↓ Plasma Proteins | <ul style="list-style-type: none"> • ↓ Production (e.g. due to liver failure) • ↓ Intake (e.g. due to malnutrition, cachexia) • ↑ Elimination (loss via kidneys, e.g. nephrotic syndrome) • Redistribution (trauma) |
| | ↑ Plasma Proteins | <ul style="list-style-type: none"> • ↑ No. of Binding Sites = ↓ Free Fraction of Drug • e.g. Elevated α_1-acid-glycoprotein during an acute stressor |
| Δ in 'apparent' affinity of drug for plasma protein | ↓ Apparent Binding Affinity | <ul style="list-style-type: none"> • ↑ Free Fraction of Drug (e.g. Drug A in below example) • e.g. Due to reversible competitive drug interactions (Drug B displaces drug A from binding sites) |
| Development of saturable protein binding (at higher drug doses) | | <ul style="list-style-type: none"> • For a few drugs, clinically used dose may be sufficiently large to saturate protein binding sites for that drug (Corticosteroids (e.g. prednisolone), Valproate, Cefazolin) • At saturating doses, free fraction of that drug will be much greater than expected (Free drug conc. increases linearly with dose, but there is a less than linear increase in total drug conc. as saturation occurs, causing free drug fraction to increase) |

Clinical Significance of Changes in Drug Binding to Plasma Protein

- **No clinical significance** at all except for a few rare theoretical exceptions to this
- Changes in drug binding to plasma protein will **alter free fraction of a drug & total drug conc.** but **once steady state has been re-established, free drug conc. will return to as before** (with **no lasting change in drug effect**)
- There is **no significance of alterations** to plasma protein binding/plasma protein binding interactions **unless measuring total drug conc. & adjusting drug dosage accordingly** (e.g. **performing TDM**)

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| Reasons why it is not Clinically Significant In nearly all cases where a clinically important protein binding interaction has been postulated, other mechanisms, have since been shown to be in vivo cause of increased drug effect (e.g. Due to metabolic inhibition rather than a change in free drug conc.) | If ↓ Protein Binding | <ul style="list-style-type: none"> • Momentarily ↑ Free Drug Conc. = ↑ Free Drug available for distribution (V_d) & clearance (CL) processes = ↑ V_d & ↑ CL = ↓ Free Drug Conc. (back to starting conc.) • Result: ↑ Free Fraction of Drug & ↓ Total Drug Conc. but no change in free drug conc. when steady state is re-established • At re-establishment of steady state, free drug conc. is the same as originally = Same drug therapeutic effect |
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| | | <ul style="list-style-type: none"> At re-establishment of steady state, free drug conc. is the same = Same drug therapeutic effect |
| Theoretical Exceptions | <ul style="list-style-type: none"> Transient change in free drug conc. before steady-state is re-established = Excess drug effect/SE/inefficacy (very rare) (e.g. anaesthetics: thiopentone, diazoxide) <ul style="list-style-type: none"> May be a consideration in determining a loading dose but will not affect chronic dosing | |
| | Drugs with very High CL (CL >30 L/h) <ul style="list-style-type: none"> E.g. IV lignocaine (Often drug dose is titrated to effect anyway) Some drugs are cleared so rapidly that elimination process actually “drags” drug from its plasma protein binding site as it goes passed in blood Drug CL is dependent only on blood flow to eliminating organ & is not affected by extent of protein binding of drug Drug CL may not rise/fall to compensate for a change in free drug conc, since elimination organ is bypassed by drug Very few clinical examples of this | |

