

An Overview of Pharmacokinetic- Pharmacodynamic modelling

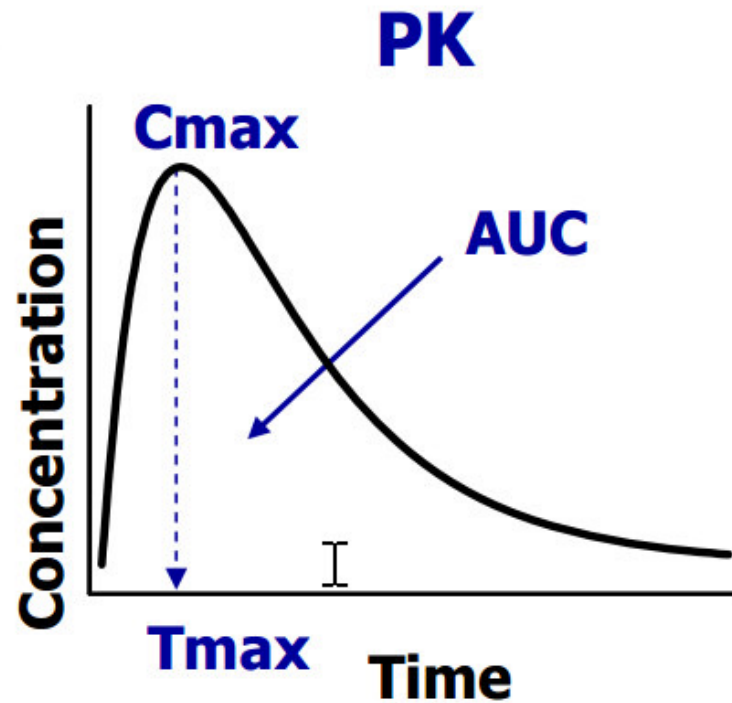
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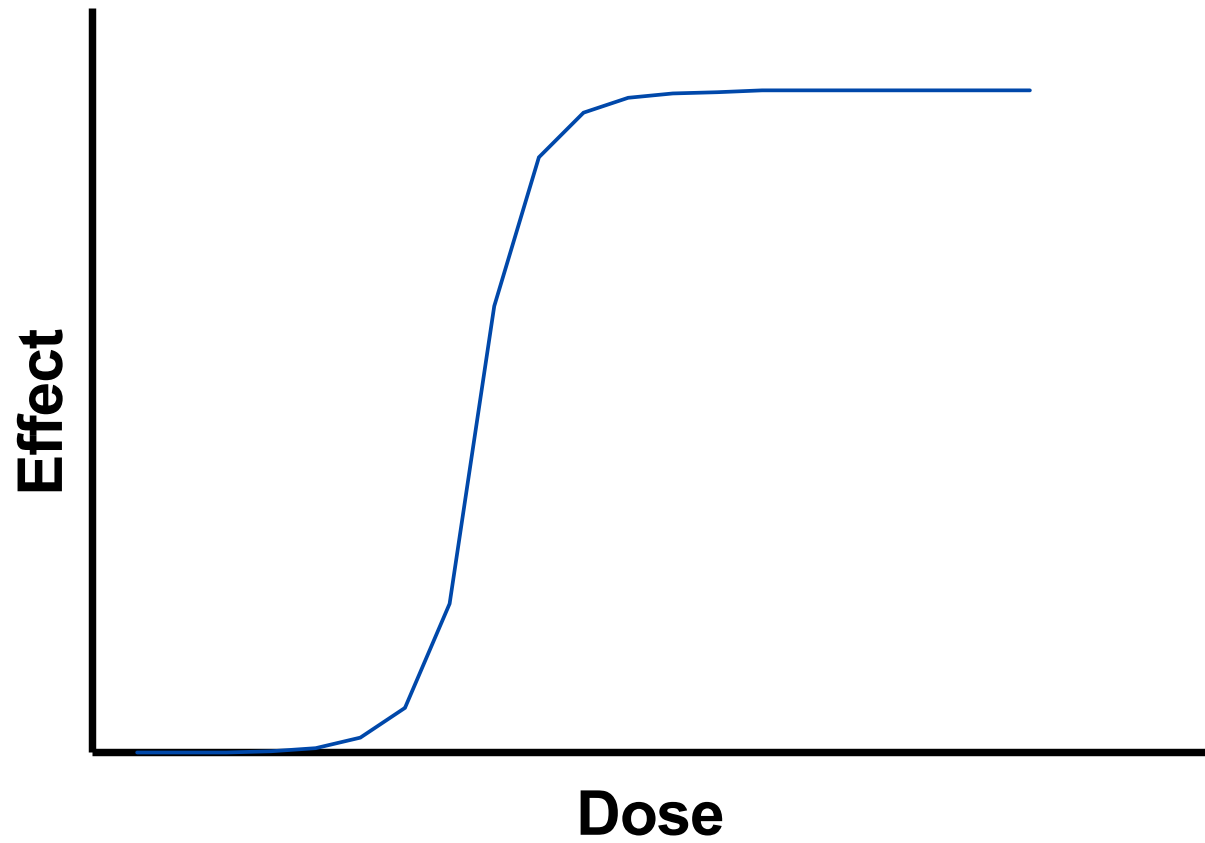
A few basic definitions

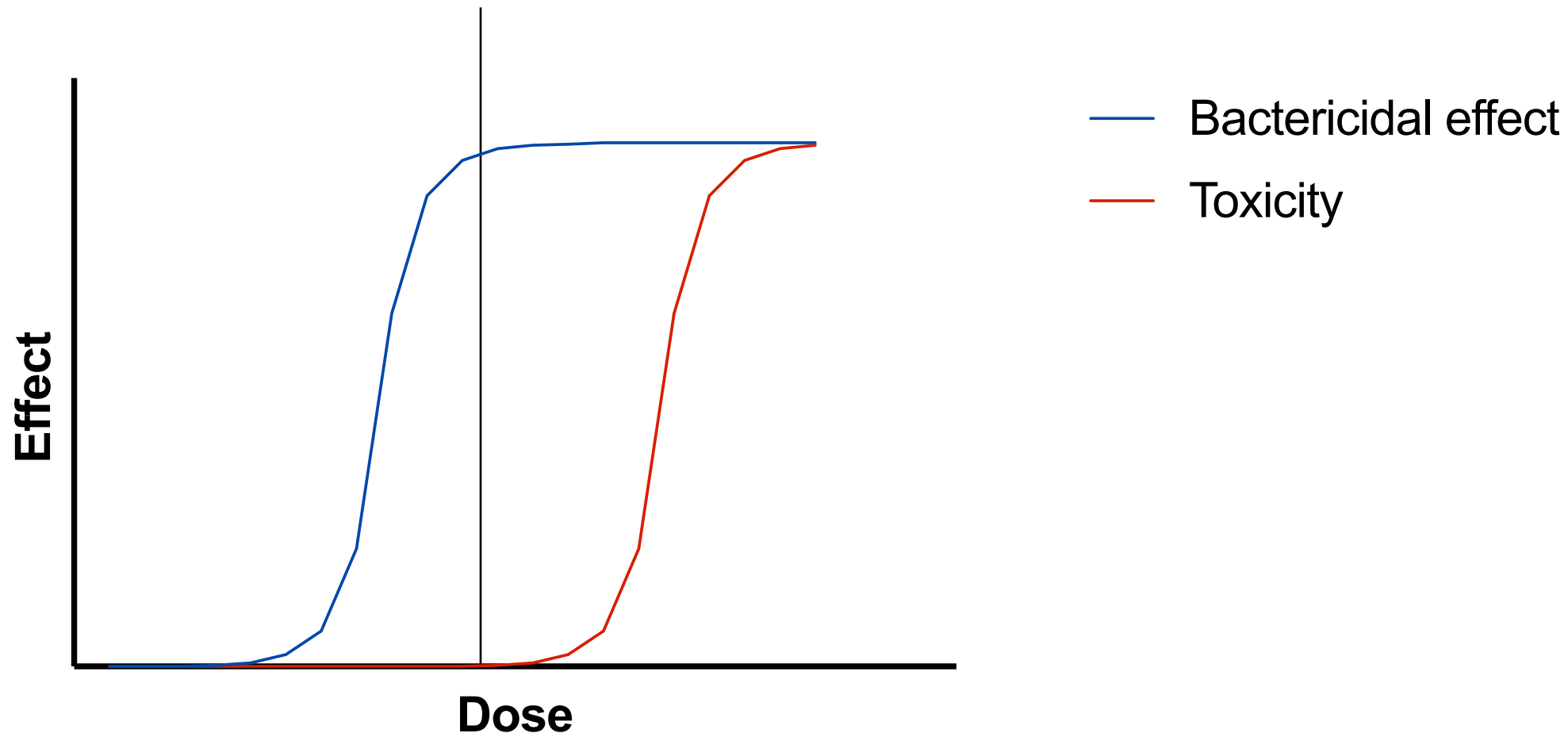
Pharmacokinetics – What a human (or animal) body does to a drug

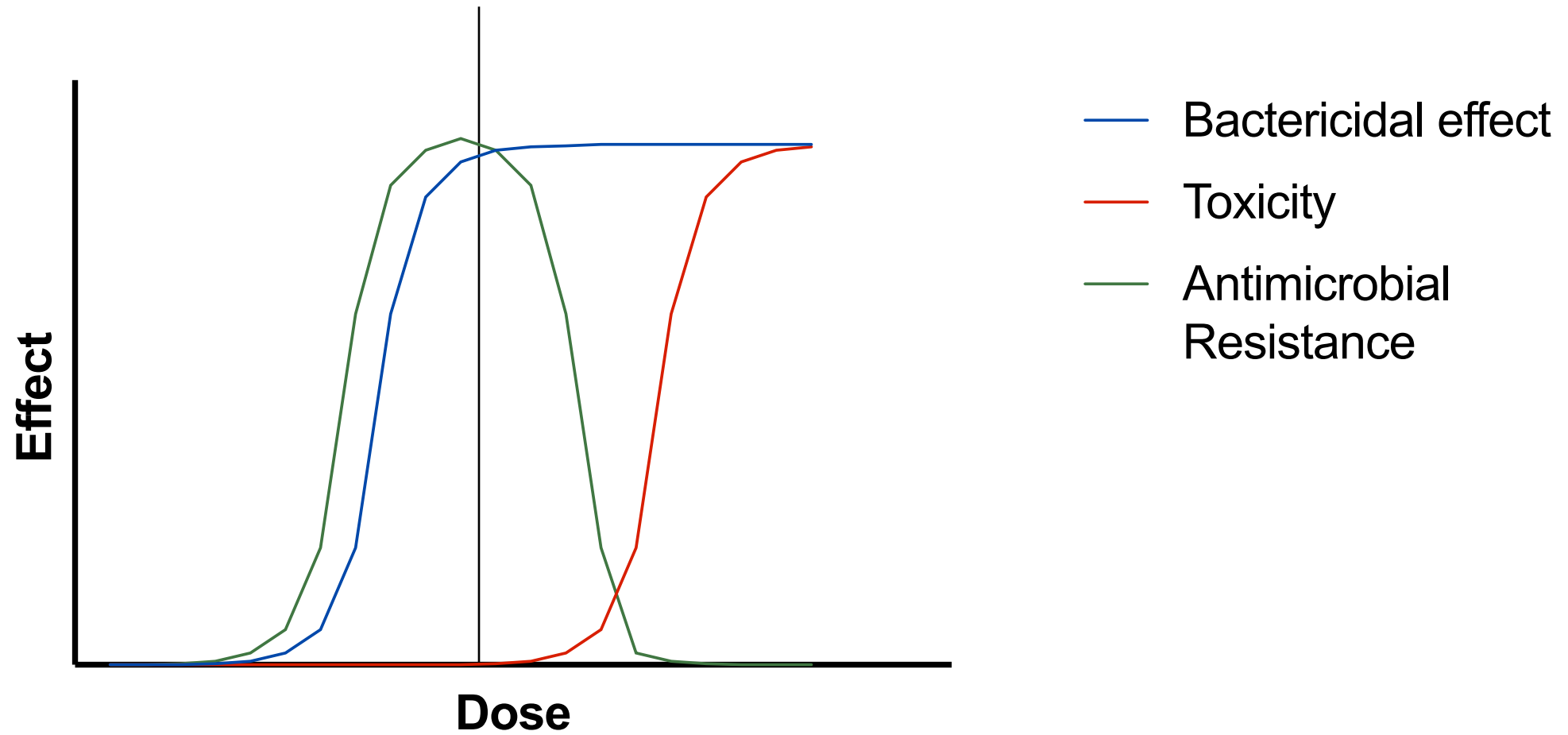


A few basic definitions

Pharmacodynamics – What a drug does to the ‘body’ (whether the host or the pathogen)

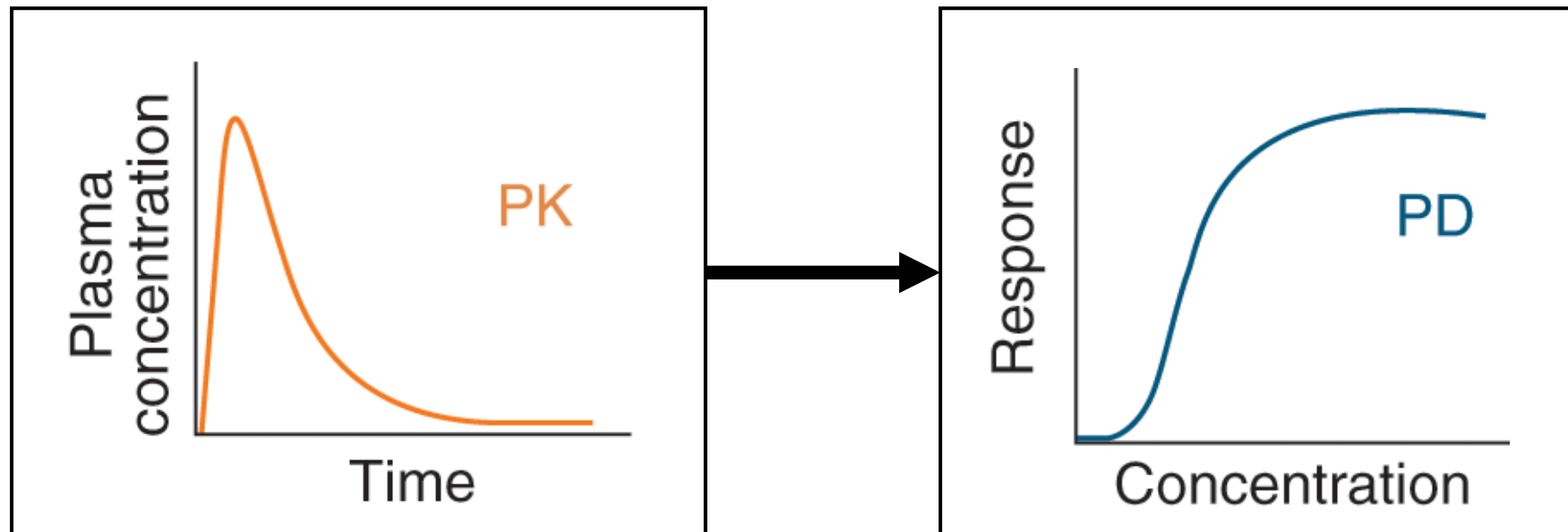






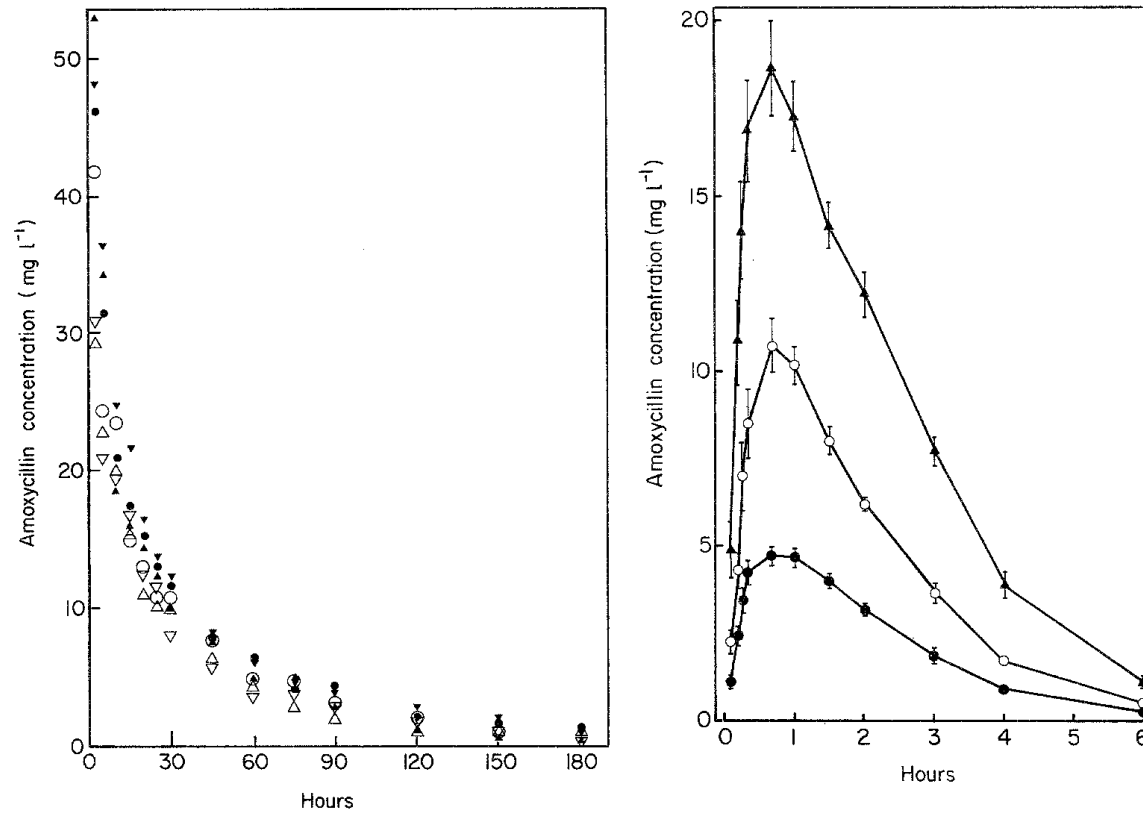
A few basic definitions

PK-PD modelling – A structural model and mathematical equations that describe the dynamic concentration of a drug in a given compartment and the effects of the drug in that compartment.

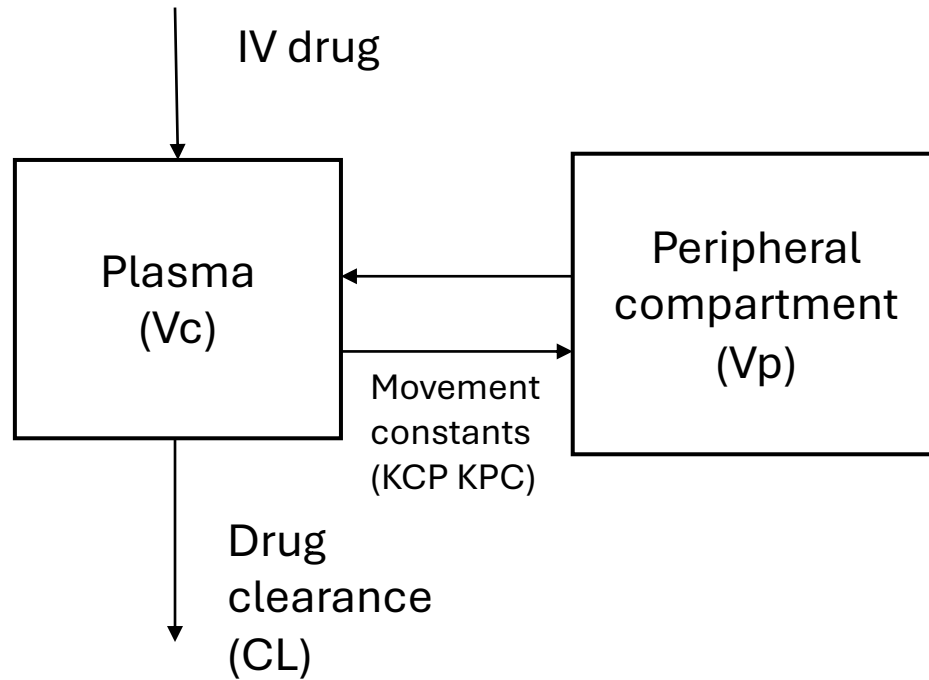


Pharmacokinetics

Tradition antimicrobial pharmacokinetics



Population Pharmacokinetic modelling



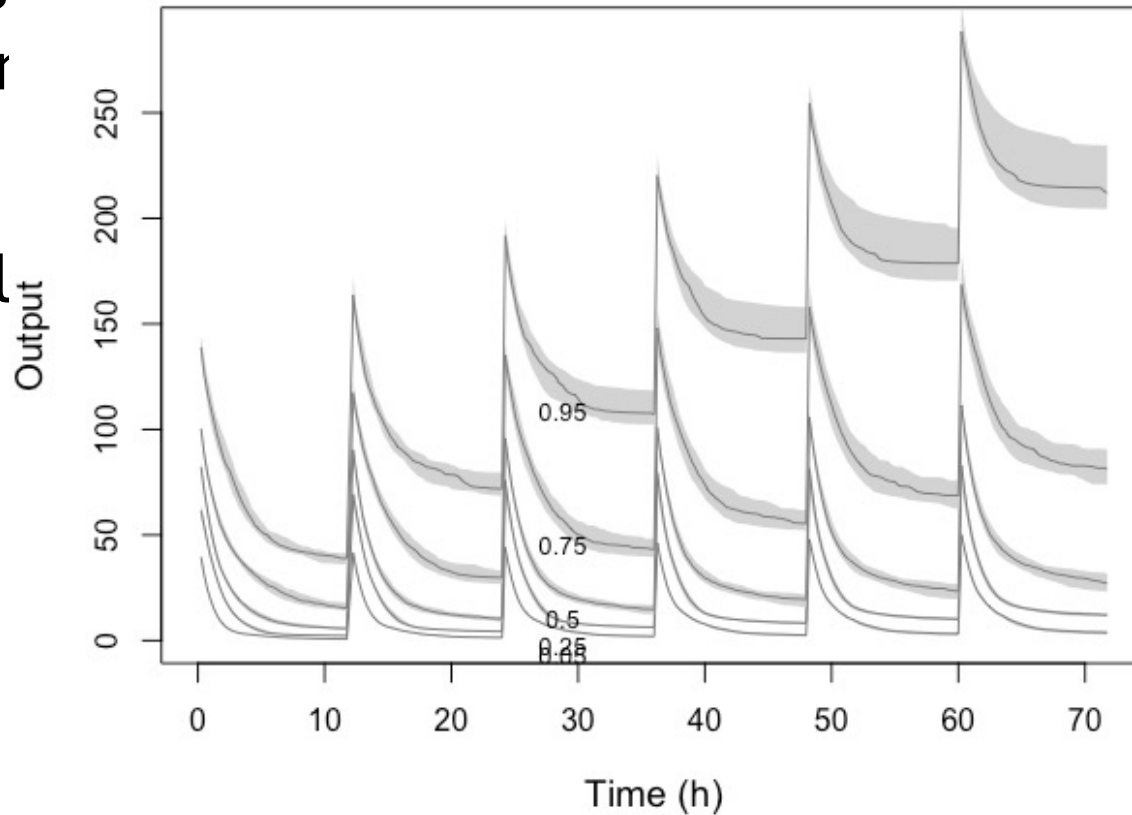
$$V_c = x * (y * \text{body weight}) * (z * \text{age})$$

$$CL = x * (y * \text{renal function}) * (z * \text{liver function})$$

PopPK modelling

Allows inc
drug pharr

Can simul



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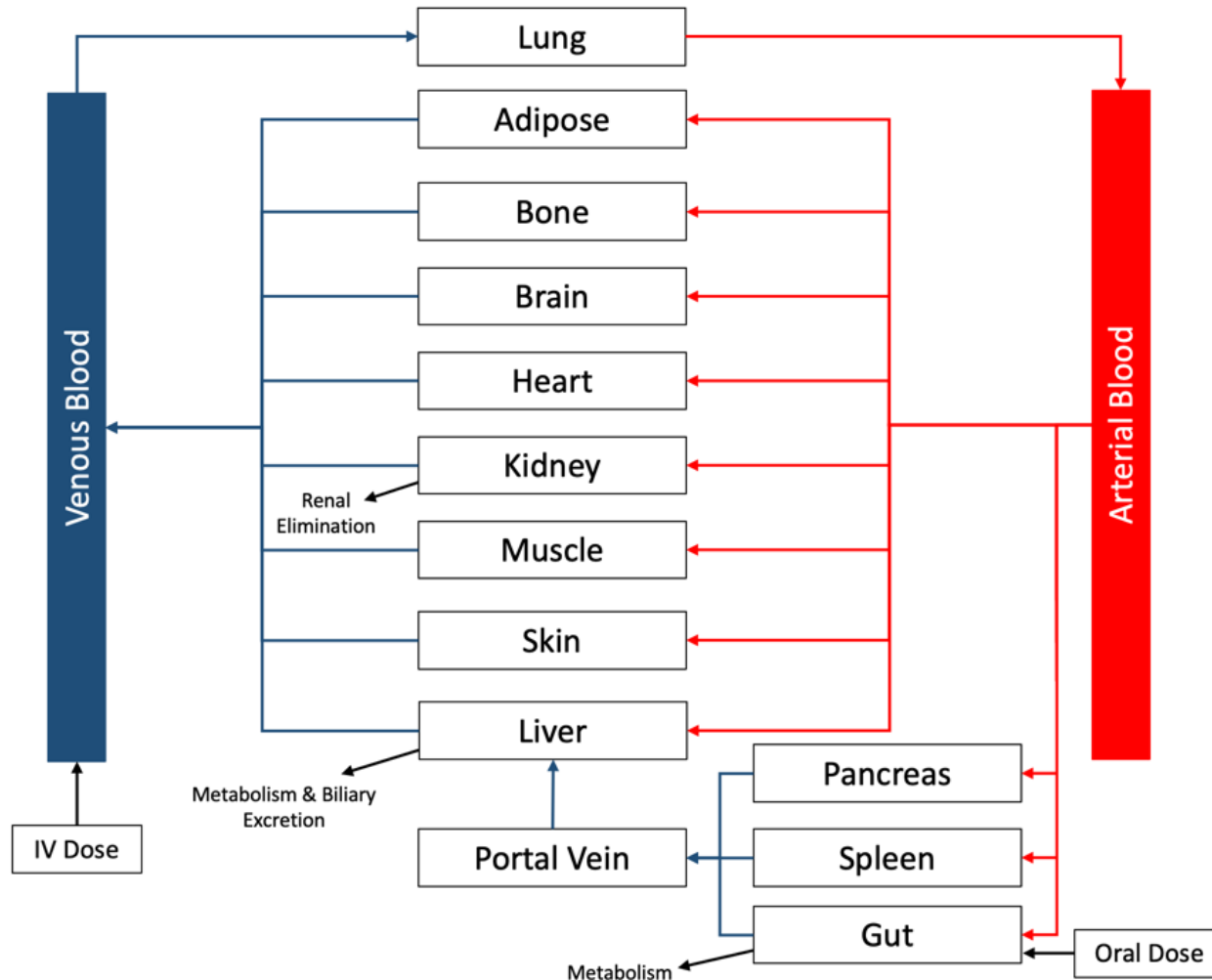
However...

PopPK models only as good as the available PK data, and most PK data is from healthy (usually male) Caucasian volunteers

Limitations in extrapolation of prediction to

- Certain demographics (ethnicities, age)
- Co-morbidities (e.g. Renal and liver disease)
- Certain medical contexts (e.g. Pregnancy, critical illness)
- Body sites beyond that sampled (e.g. bone, CNS)

Physiology-based PK modelling



Uses physiological, biochemical and anatomical insights of the body

with

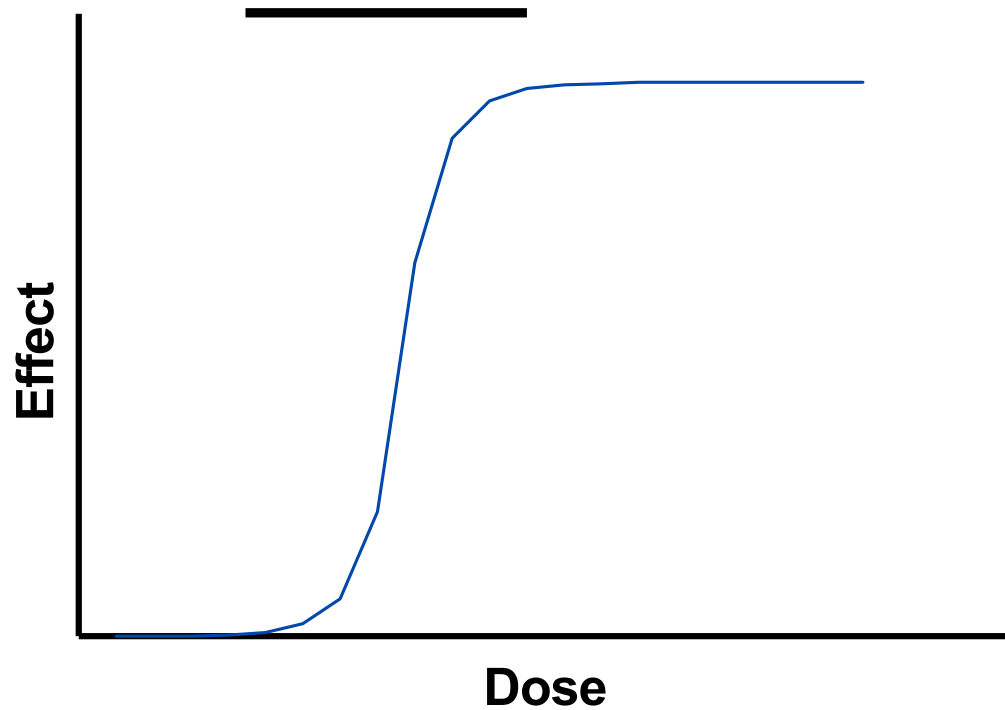
Phys-chemical and pharmacological properties of drug to predict pharmacokinetics

Can predict:

1. Drug concentrations in particular tissues
2. Variations of pharmacokinetics with different special populations
3. Interactions of drugs not tested in combination
4. Predict human pharmacokinetics for molecules in early drug development)

Pharmacodynamics

Pharmacodynamic modelling



$$\frac{dX}{dt} = X * \left(Kg * \left(1 - \frac{X}{POPmax} \right) - \frac{Kk * \left(\frac{X}{E50} \right)^H}{\left(1 + \left(\frac{X}{E50} \right)^H \right)} \right)$$

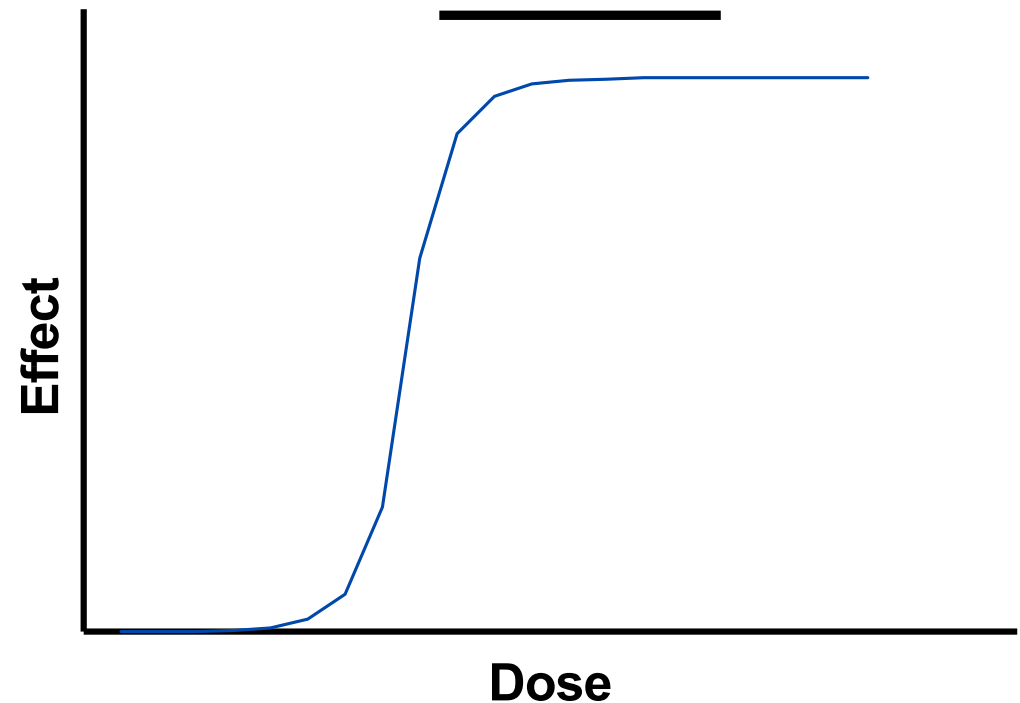
Where X = bacterial population

Pharmacodynamic data from Humans

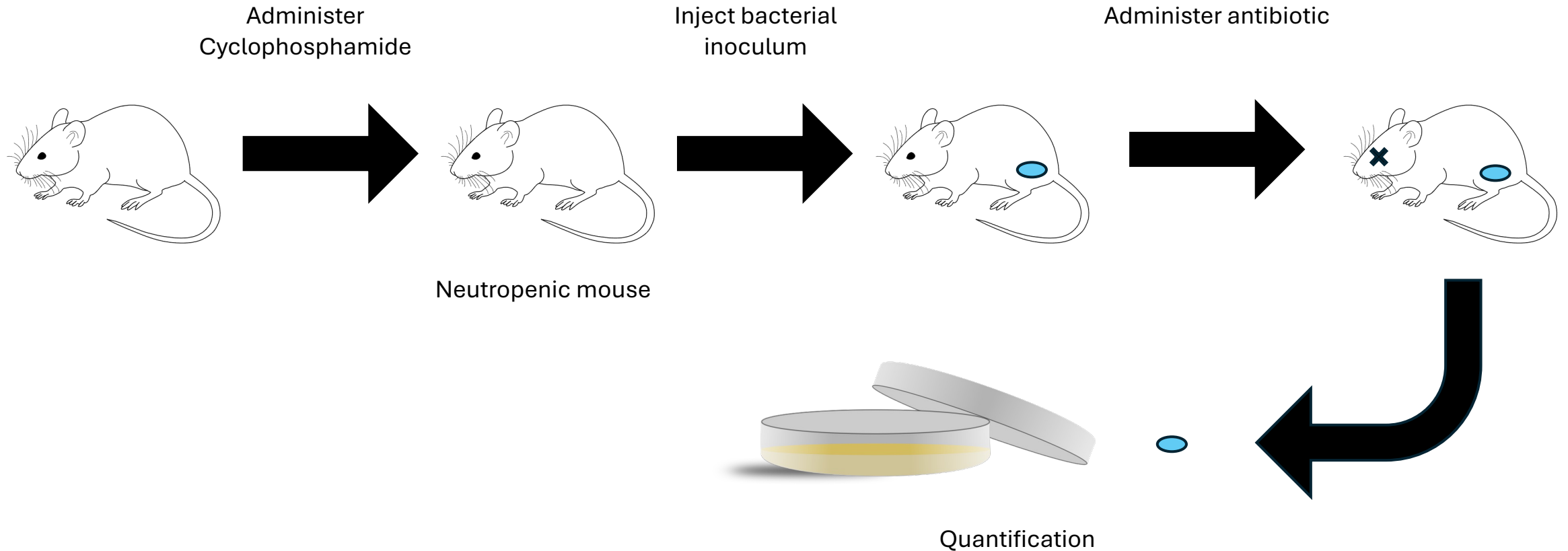
In theory, most ideal situation

However

- Data usually either categorical (e.g. mortality, microbiology success) or use of surrogates (e.g. inflammatory markers)
- Cannot measure effect for sub-maximal effect (ethically, at least)



Neutropenic mouse model



Neutropenic mouse model

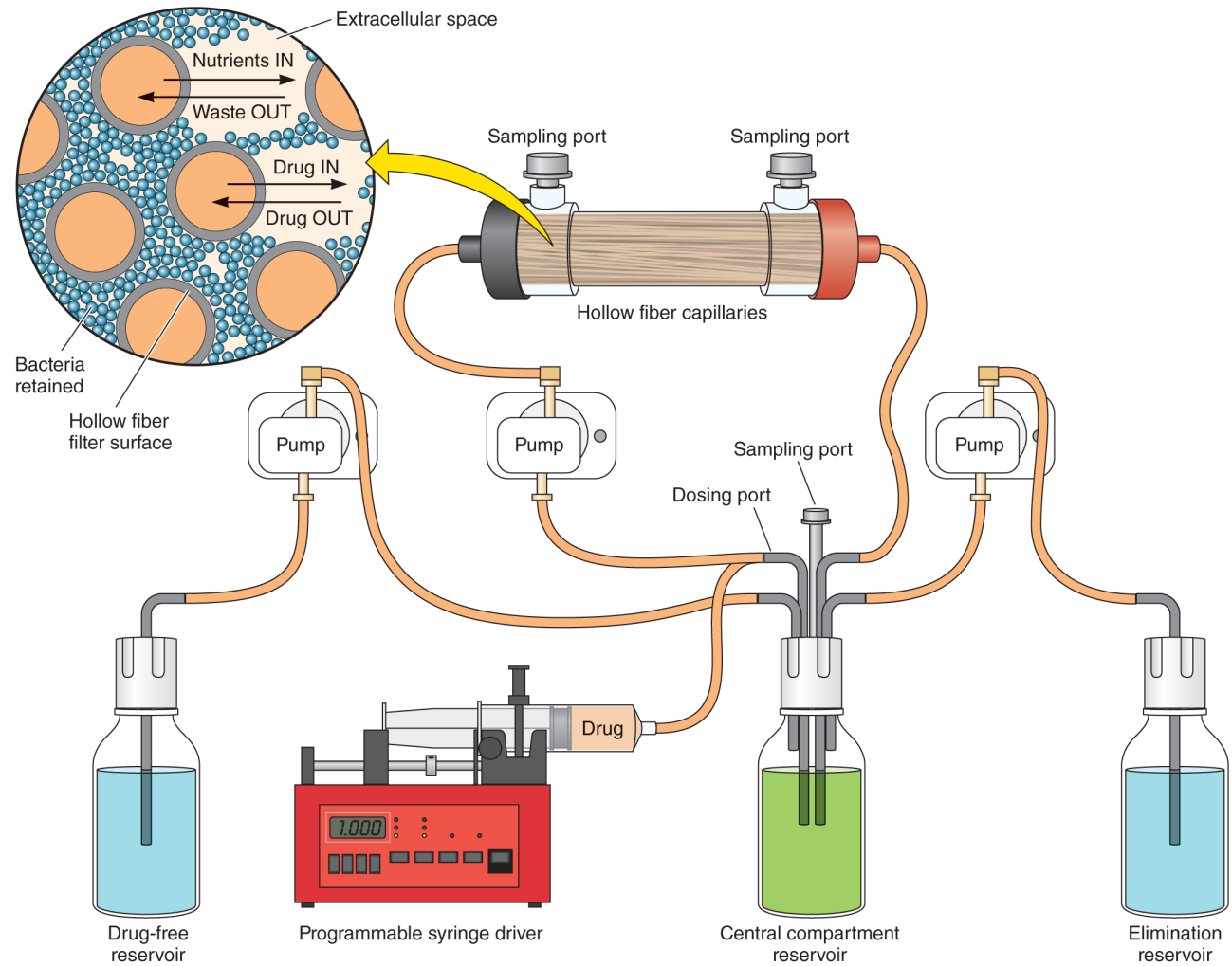
Benefits

- Standardised and reproducible
- Lots of experience with the model
- Conclusions readily translate to humans
- Cornerstone of PK/PD target definition

Limitations

- Resistance is almost never measured
- Mice usually cannot survive for >24-48h
- Mice are small, and therefore small inoculum
- Differences in animal PK and physiology

Hollow Fibre infection model



Hollow Fibre infection model

Benefits

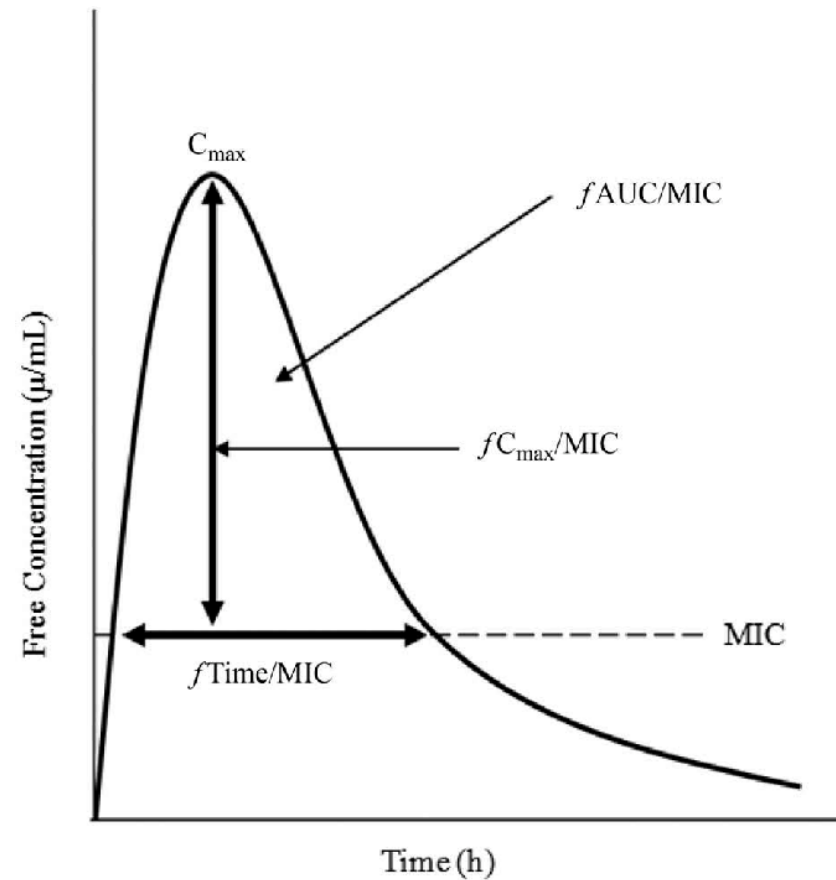
- Can examine resistance
- Can use high inoculum infections over a long period of time
- No use of animals
- Can use whichever PK profile you like

Limitations

- Expensive
- Cannot replicate holistic tissue partitioning
- Lacks physiological context (protein binding, immune effectors etc.)
- Lack of track record with translation to humans

PK/PD model outcomes

1. Define the relevant pharmacodynamic index



PK/PD model outcomes

2. Define the pharmacodynamic target index for e.g. bacterial stasis, 1-log bacterial kill or suppression of emergence of resistance

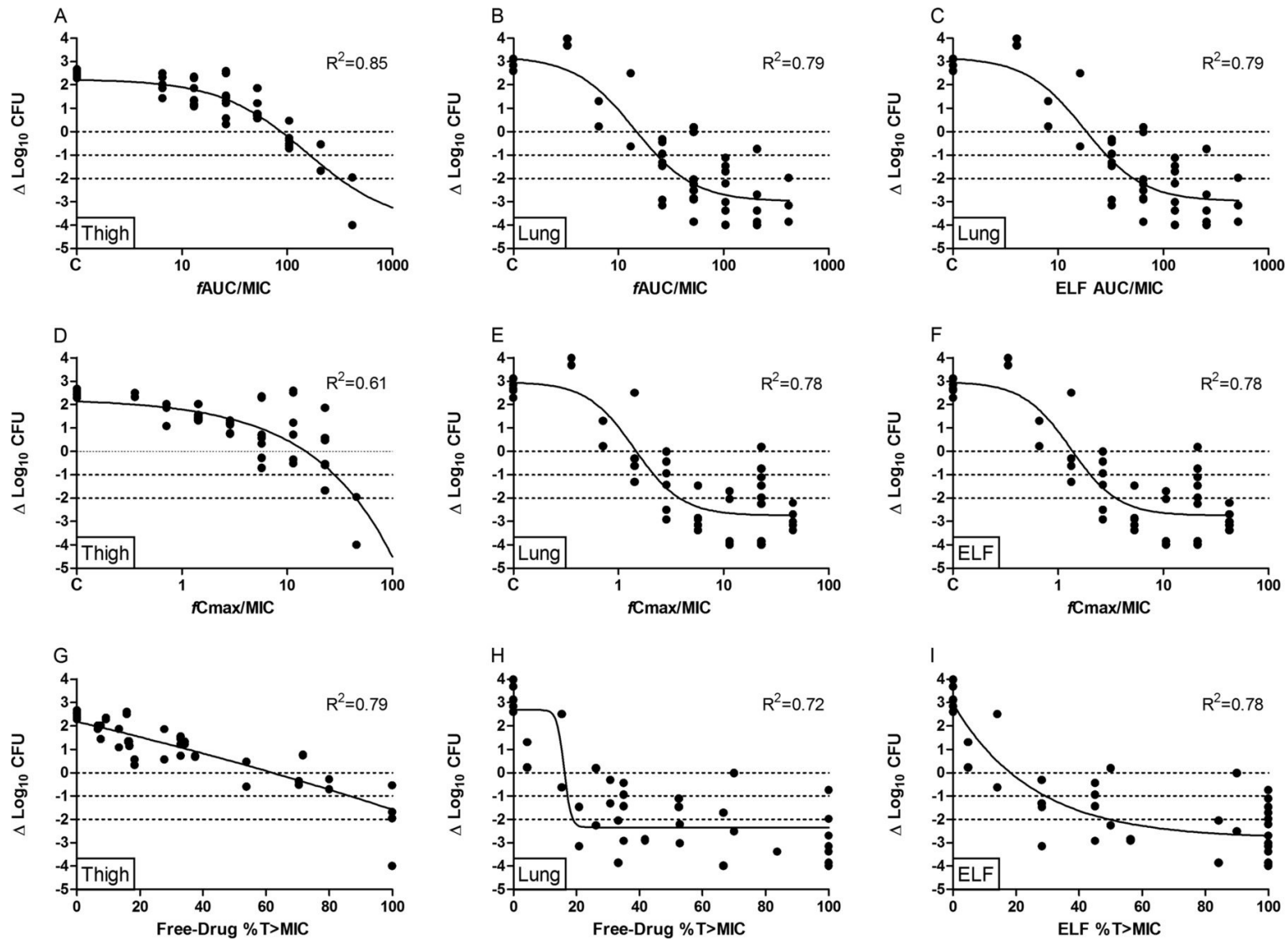
e.g.

$C_{max}:MIC > 10$

$AUC:MIC > 200$

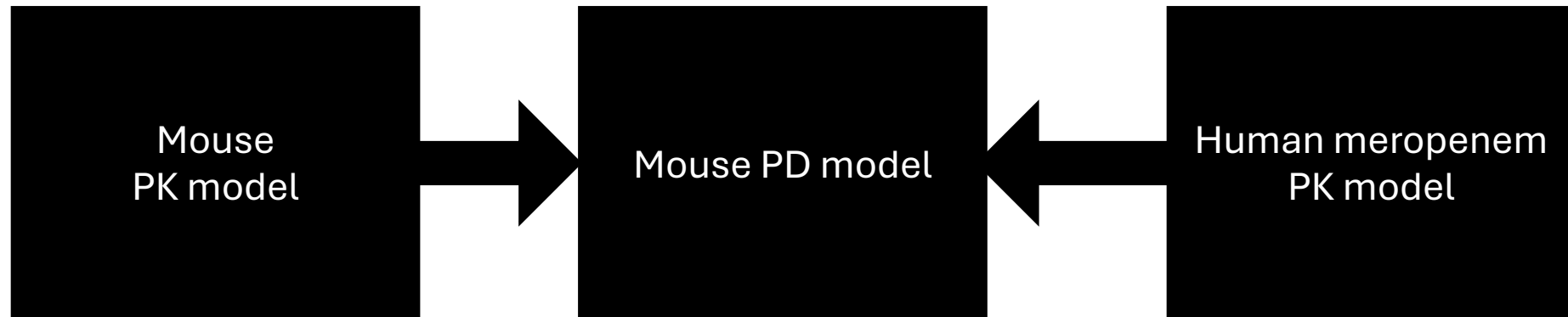
$\%Time\ above\ MIC > 50\%$

...and define regimen in humans that meets them



PK/PD model outcomes

3. Simulate pharmacodynamic outcome in humans.

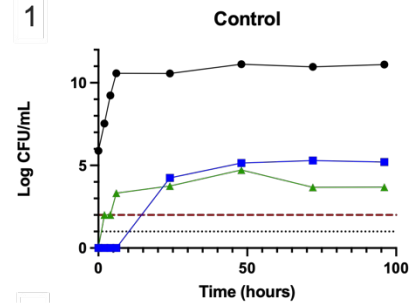


PK/PD model outcomes

4. Characterise pharmacodynamic interactions of drugs

i.e. Determine if drugs are synergistic or antagonistic for both bactericidal effect and emergence of resistance

1



● Total Counts ■ Flomoxef resistance counts ▲ Fosfomycin resistant counts Limit of detection (total counts) ---- Limit of detection (resistant counts)

In summary...

This whole process underpins optimal dose
selection of antimicrobials

Some big limitations...

1. Human pharmacokinetics is often poorly characterised and/or from a limited population

- Early antimicrobials not modelled well (if at all)
- Data largely from (usually male) Caucasian healthy volunteers
- Data often lacking from specific demographics and patient groups

2. Pharmacodynamics for bactericidal effect typically not characterised well for many antibiotics

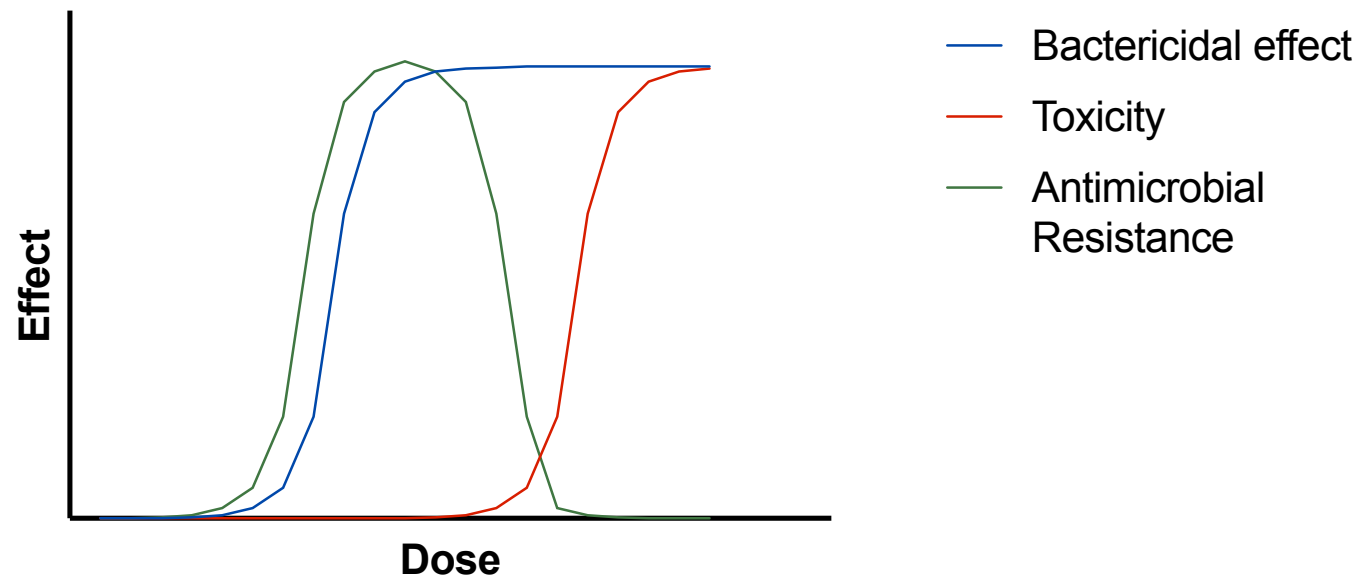
- Particularly older (and more frequently used antibiotics)

e.g. amoxicillin, flucloxacillin

- Particularly in certain infection sites

3. Pharmacodynamics for antimicrobial resistance often not characterised at all

- Drugs are not dosed appropriately to prevent emergence of resistance
- Includes newly developed drugs
- Arguably is the main reason for antimicrobial resistance





Clinical Microbiology
Reviews



Antimicrobial Chemotherapy | Review

Challenges for global antibiotic regimen planning and establishing antimicrobial resistance targets: implications for the WHO Essential Medicines List and AWaRe antibiotic book dosing

Nada Reza,^{1,2} Alessandro Gerada,^{1,2} Katharine E. Stott,^{1,2} Alex Howard,^{1,2} Mike Sharland,³ William Hope^{1,2}

4. There are lot of assumptions built in

- Translatability from *in vivo* / *in vitro* platform
- The true 'MIC'
- Extrapolation of pharmacodynamics to other strains/species

5. Resource and time intensive

- A PK/PD experimental package for a single drug-bug combination will take 6+ months and cost £100,000s
- Impossible to keep up with evolving microbiology
- Impossible to cover every drug-bug combination

Possible solutions

1. Just need to do more of everything
2. Utilise novel *in silico* options e.g. PBPK modelling
3. Greater PK data availability
4. Regulatory input (for antimicrobial resistance emergence)
5. Rethink of how we examine pharmacodynamics
 - Better ways to use human pharmacodynamic data?
 - Better measures of antimicrobial susceptibility than MIC?

Thank you