



Grand Challenges
African Drug Discovery
Accelerator

Drug Discovery and Development Course

Introduction to Drug Discovery and Development
Jeremy Burrows

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Learning objectives

- Understand the overview of the drug discovery and development process
- Understand importance of working on the right biology
- Understand those properties that impact the chemical quality of a compound series

Drug Discovery Philosophy

- **Be curious – the best question is “why?”**
- **You will not understand everything outside your domains of expertise – ask questions!**
 - Do not pretend to be an expert in areas where you are not
 - Be responsible for understanding and delivering expertise in areas where you are expert – explain your thinking and interpretation of data to others
 - Ask, listen and learn from those who are experts outside your area
- **Focus on the goal of delivering a candidate drug**
 - Drug Discovery is not the same as doing basic science
 - Don't do something because you can do it, rather focus on answering the right questions
 - Gather data, define the issues, ruthlessly focus on solving the issues with chemistry, biology and pharmacokinetics
- **Collaboration and team-work are key**
 - It is not about “me” but the project and the project team
 - All need to learn the multi-disciplinary language of Drug Discovery

Drug Discovery – where to start?



- **Drug discovery is a misnomer**

- “a wrong or inaccurate use of a name or term”
- Except for some natural products – drugs are novel chemical entities that did not exist before they were made
 - Derived from imperfect starting compounds that have been extensively changed and optimized across many parameters

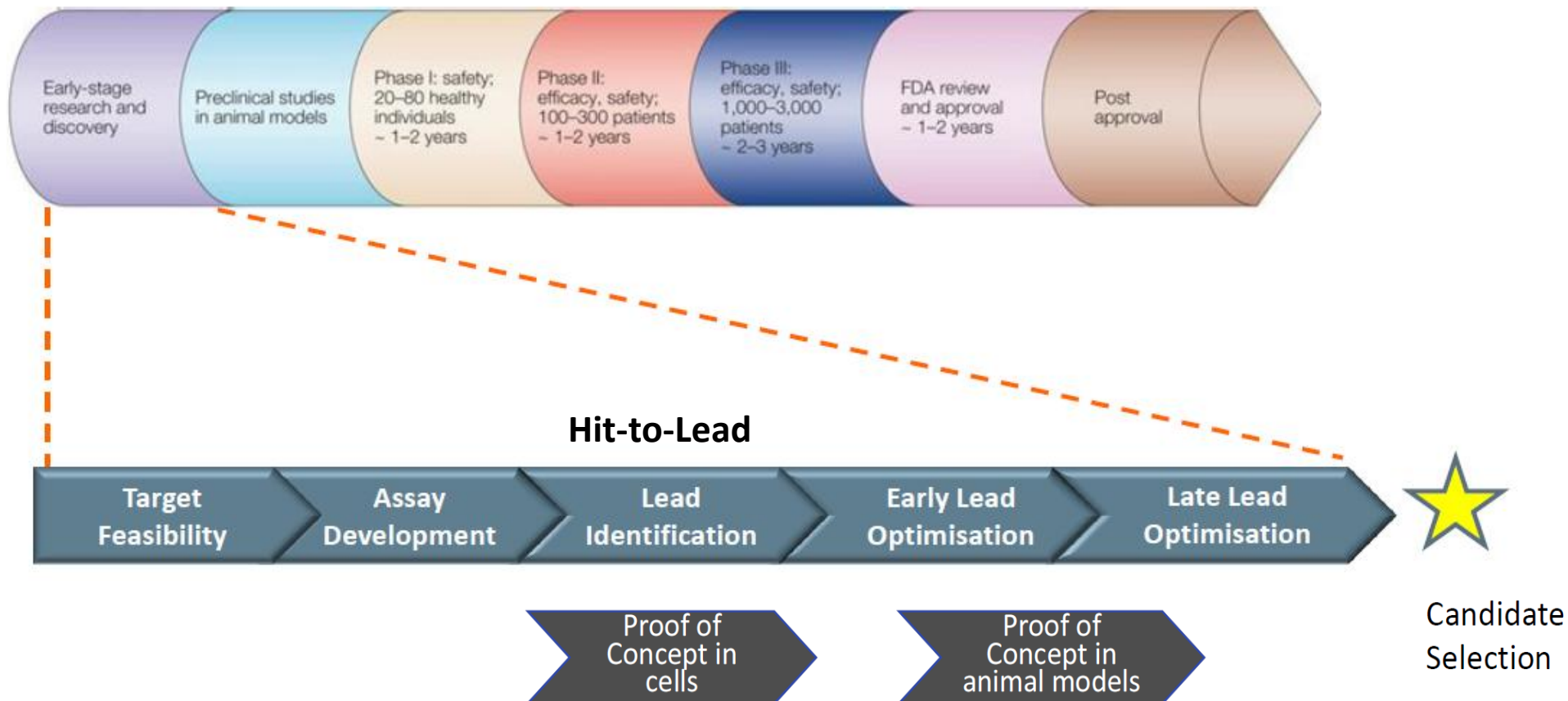
- **Drug discovery is a journey with various options**

- New biology – need to find a disease indication
- Defined disease – need to find the best approach

- **It is better to start from the end and define a route to get there!**

- Target Product Profile (TPP)
- TPP describes the profile of the medicine necessary to meet the goals of patients, care givers and payers

Drug Discovery Phases



Proof of Concept: Demonstration that the target/ compound series gives the desired response before studying in humans

What biological target to work on?

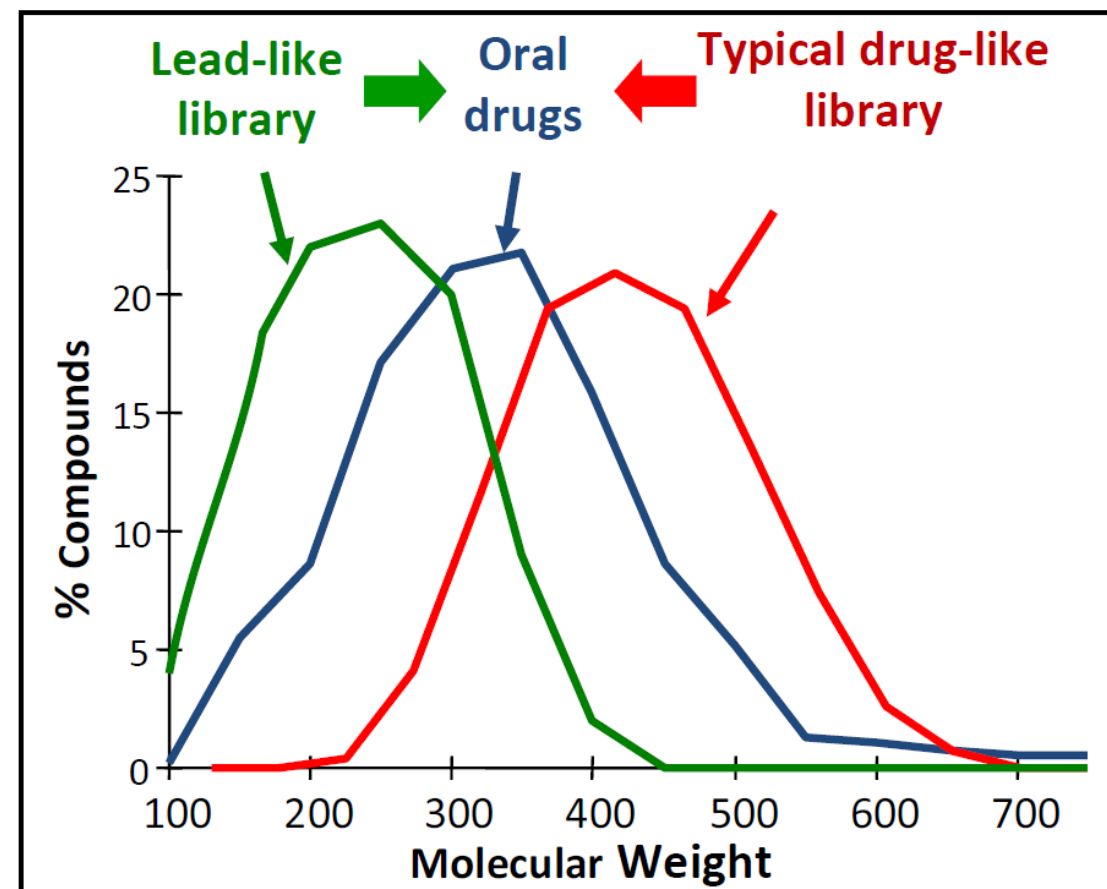
- **Critical to work on compounds having biology relevant for disease**
- **If not, even the best project team delivering the best candidate drug will deliver failure over years costing USD millions**
- **So biological target must be validated for the disease**
 - Many ways to confirm validation
 - Best to have evidence across several areas – genetic, phenotypic, disease model, chemical
- **Test cascade is the sequence of assays run to support optimisation**
- **Test cascade must also be validated and relevant for the disease**
 - *In vitro* biochemical, cellular phenotypic and *in vivo* efficacy assays are robust, validated with controls and deliver data that translate to patients
- **If not, even if working on the best target for the disease, the test cascade will lead the project team in the wrong direction**
- **Right biology and right assays!**

Hit identification options

- **Target Based**
 - High Throughput Screening (HTS); often large compound numbers in low volume (384 / 1536 well)
- **Focused Screen (biased!)**
 - Subset of molecules based on target class / structural knowledge
- **Fragment Screen**
 - High concentration screen of small molecules - biophysical methods (MWt <250)
 - Crystallography often required for optimisation / confirmation of binding
- **Virtual Screen**
 - *In-silico* docking compound library into protein model (X-ray / AlphaFold)
- **Structure based design**
 - Exploit structural knowledge to design molecules
- **Phenotypic screen**
 - Seeking to see a desired response in cellular systems
 - Screening many targets
- **Literature – work from a known ligand, could be a back-up**

What happens during optimisation?

- Optimization often results in increasing Mol Wt and modulating lipophilicity (cLogP) to deliver a lead and candidate drug
- If a library is screened where the molecules have high Mol Wt then high potency compounds may be found – but they are hard to optimize
- Small Mol Wt “lead-like” molecules have a better chance of binding to a receptor than larger drug-like molecules and may have weaker potency – but they are easier to optimize
- Focus on a screening library or hits that are “Lead-like” will give a greater chance of success



What makes a good hit?

- **Potency and selectivity**
- **Drug-like properties including low Molecular Weight (MW) and low lipophilicity (LogP)**
- **Synthetic accessibility and versatility**
- **No undesirable groups**
- **Freedom to Operate (FTO) and Intellectual Property (IP) space**

- **Potency is the measure of compound activity and reflects the thermodynamics of a compound binding to a biological target**
- **Governed by $\Delta G = \Delta H - T\Delta S$**
 - ΔH enthalpy – bond breaking (i.e. desolvation) and bond forming (how well the drug binds to the target)
 - ΔS entropy – overall impact on disorder of the system
- **Most, but not all, drugs in the nM range**
 - Weaker compounds require higher concentrations to efficiently interact with target leading to increased chances of off-target effects
 - Potency impacts dose size
- **Potency is NOT the only game in town!**

Ligand Efficiency (LE) – A measure of “Fit”

- **Binding contribution per non-H atom**

- “Goodness of fit”

$$LE = \frac{-RT \ln Ki}{HA}$$

HA = no. of Heavy atoms i.e. non-H atoms

R = Gas Constant ($0.001987 \text{ kcal K}^{-1} \text{ mol}^{-1}$)

T = Temperature in Kelvin

Ki = Potency in M

$$LE = pIC_{50} * 1.37 / HA \text{ [} pIC_{50} \text{ is } -\log_{10}(IC_{50}) \text{]}$$

Has units kilocalories per mole per non-H atom

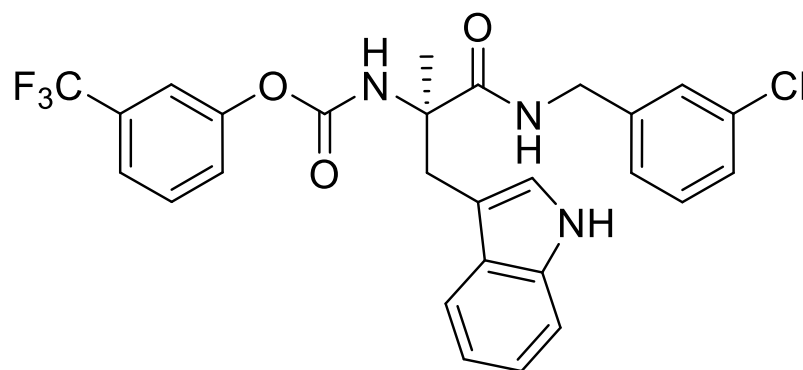
Lipophilicity Ligand
Efficiency

$$LLE = pIC_{50} - \log D$$

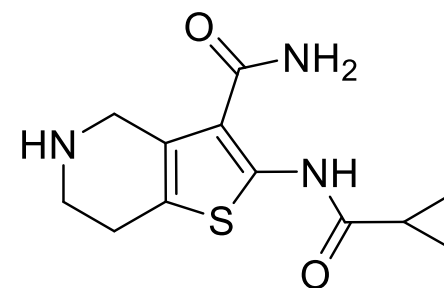
- **Optimising a low LE hit has risks**

- Poor, non-optimal fit to target; Potency gained by bulk and not specific interactions

Mentimeter: Which molecule would you rather work on?

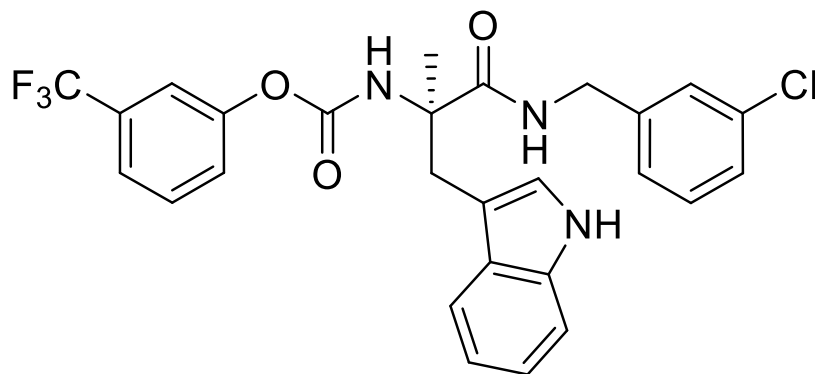


$K_i = 100\text{nM}$

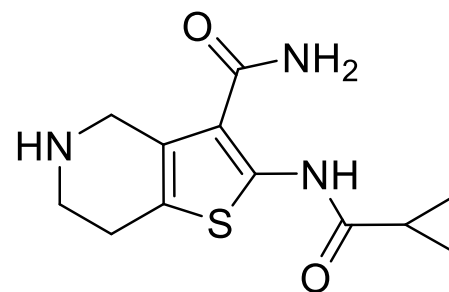


$K_i = 10,000\text{nM}$

Which molecule would you work on?



Ki = 100nM
Heavy atoms = 37
LE = 0.26



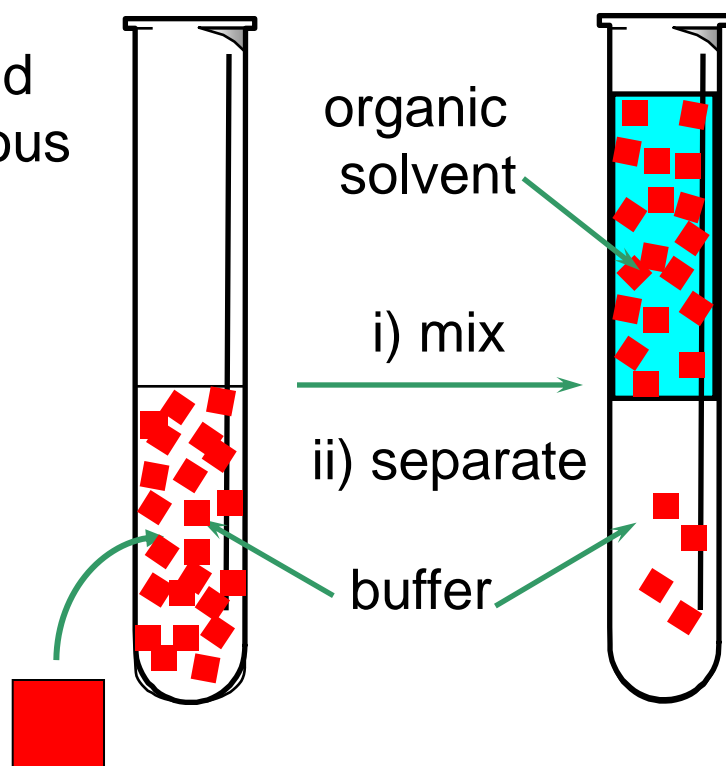
Ki = 10,000nM
Heavy atoms = 18
LE = 0.38

- **At Hit stage it is most critical to understand the selectivity over the most relevant off-targets**
 - Cellular selectivity >10 fold over cytotoxicity in hit
 - If biological target known, selectivity to closest human targets of concern, plus targets known to cause toxicity such as hERG
 - Helps to define the scale of the challenge for Hit-to-Lead
 - Selectivity can be altered with chemistry – so criteria increase with later phases
- **What selectivity window is needed ultimately?**
 - Depends on impact of non-selectivity!
 - Need to define what assays are used to define selectivity
 - At least 100-1000 fold selectivity typical goal for a candidate
 - hERG – 30 fold margin based on IC_{20} vs free C_{max} at therapeutic dose

Lipophilicity (LogP)

- Partition coefficient is called LogP
- Measure of lipophilicity or degree to which a compound will partition into 'fatty' tissue vs water

Add
Compound
to Aqueous



Measure analyte in aqueous phase before and after shaking with organic solvent.

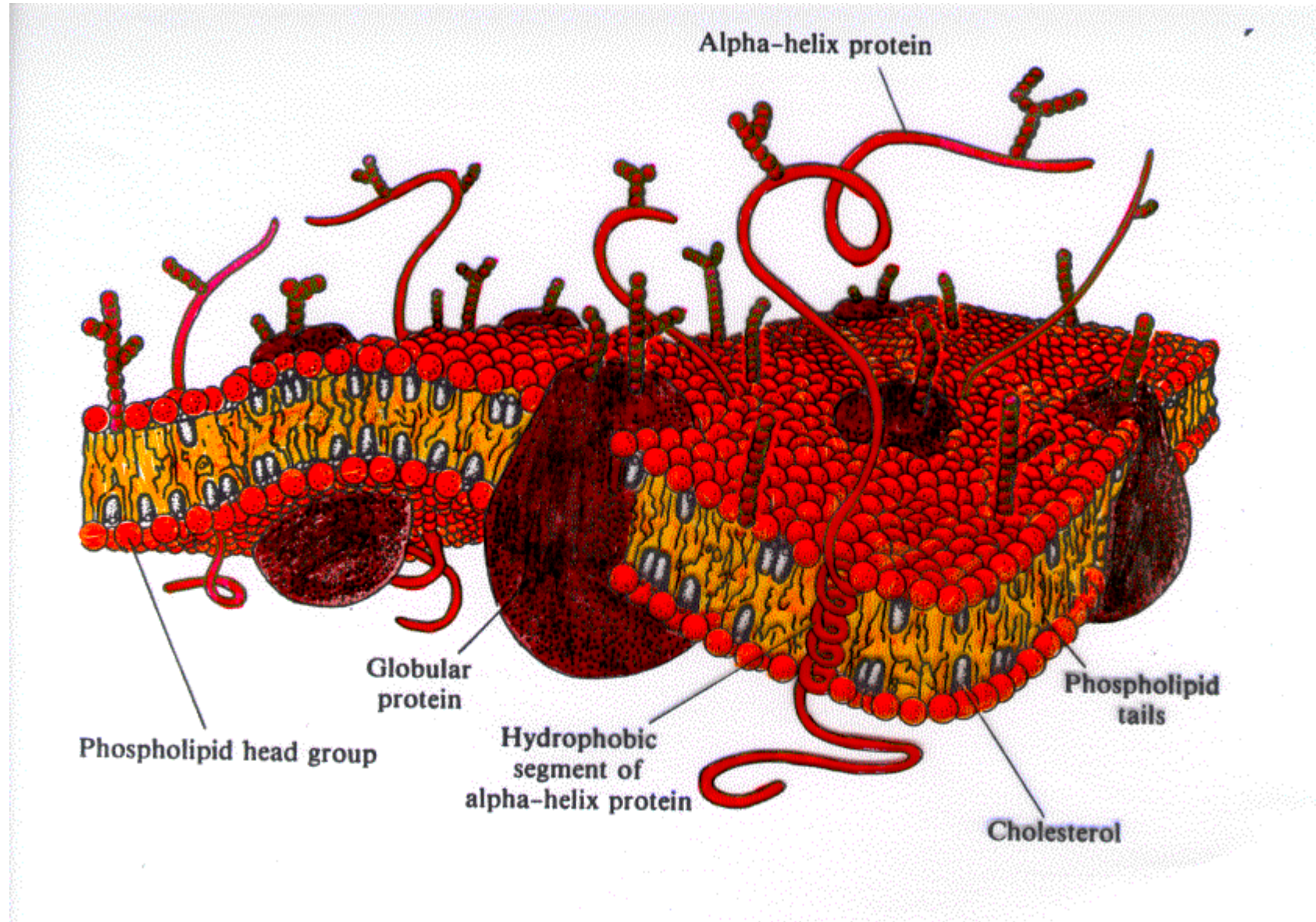
$$P = \frac{\text{conc in org}}{\text{conc in aqu}} = \frac{18}{2} = 9$$

$$\text{LogP} = \text{Log}_{10}P = 0.954$$

clogP = calculated LogP

- **Why is solubility important?**
 - Drugs must be in solution to be absorbed
- **What solubility do I need?**
 - Depends upon route of administration – e.g. oral versus IV
 - Related to dose size
 - Higher potency means lower solubility tolerated
- **Thermodynamic solubility measure is best, but kinetic OK for trends**
 - Remember that compound that resembled brick dust? Forget it!
- **A compound is more soluble as its partitioning into water increases i.e. lower logP the better**
 - If a compound is ionized to an acid or base – more soluble
 - LogD is distribution coefficient and corrects LogP for ionization
 - High melting point lowers solubility

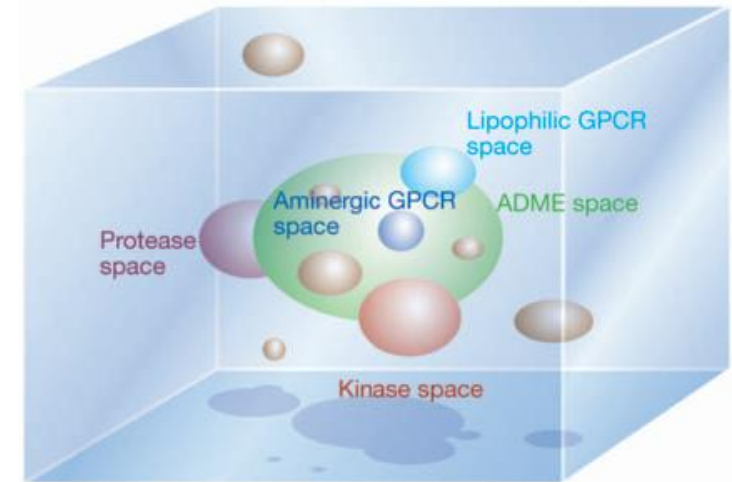
Permeability



tPSA =
Topological
Surface
Area

Lipinski's Rule of 5

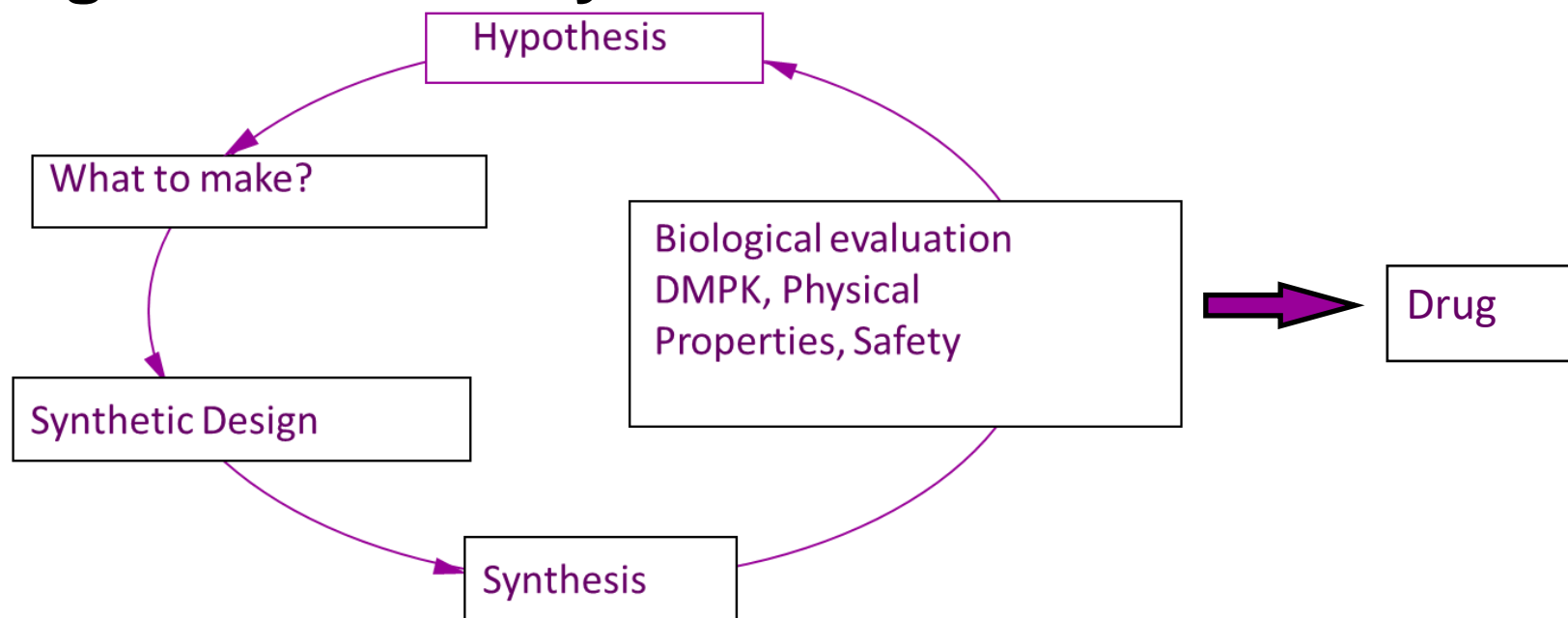
- Empirical rule, based on observation (marketed drugs) of physicochemical Assessment of orally bio-available drugs
- Poor absorption is likely when molecule has
 - >5 Hydrogen-bond donors (NH, OH)
 - >10 H-bond acceptors (Heteroatoms)
 - cLogP >5
 - MW >500
- Oral drugs need to be ADME compliant
 - Absorption, Distribution, Metabolism, Elimination
 - Also called DMPK – Drug Metabolism, Pharmacokinetics



C. Lipinski and A. Hopkins,
Nature, **432**, 855-861, (2004)

Synthetic accessibility and versatility

- The central engine of a drug discovery project is the Hypothesis-design-make-test cycle



- If a chemical series' synthesis is inaccessible the cycle cannot complete or is drastically slowed down
- Feasibility to modify the chemical structure at different points – synthetic versatility – important to explore scope of series

Undesirable groups and toxicophores

- **Includes groups known to:**
 - Be reactive
 - Be unstable
 - Be Toxic
 - Give rise to toxic species on metabolism
 - Lead to rapid metabolism and excretion
 - Have caused clinical candidates to fail
- **Such chemistry should be avoided or optimised out in Hit Generation**

- **Important to know if a hit series has been published**
- **Chemical substructure searches of literature**
 - Provide information on novelty and known pharmacology of series
 - if not novel – information can help with synthesis, other biological activity and competition
- **If the project plan is to ultimately file a patent application, then the compounds patented must be novel and inventive**
 - No issue if series is novel from the outset – just need to keep confidential
 - If not novel, then chemistry will need to factor this in as an additional constraint
 - Must not publish or disclose novel chemical structures and data before patent application filed

- Phase to drive project towards target product profile, addressing key issues along the way
- Aim is to validate chemical series as being able to deliver a compound meeting the profile by showing the series is:
 - Potent and Selective
 - Drug-like (physicochemical)
 - Drug-like (DMPK/ADME)
 - Potential for acceptable safety
- At **lead** stage the optimal profile does not have to exist in one compound, but each component should exist within the series
- Scope to deliver a **candidate** is believed to exist

- **DMPK = Drug Metabolism and Pharmacokinetics**
- **ADME = Absorption, Distribution, Metabolism and Elimination**
- **PD = Pharmacodynamics**

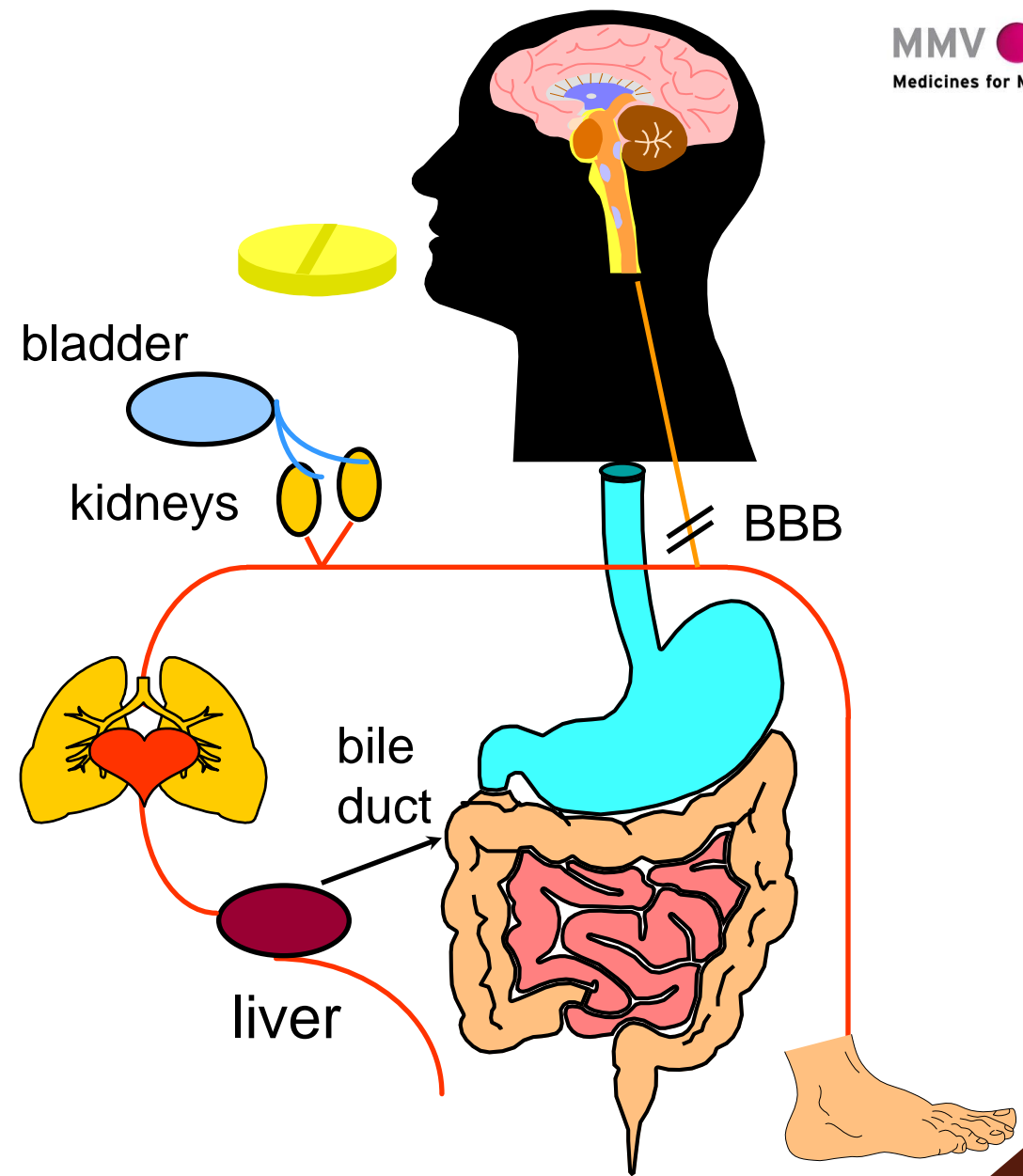
- **Pharmacokinetics is what the body does to a drug**
 - Want appropriate PK in animals (for efficacy and safety studies) and then to optimise, based on predicted human PK, to a candidate drug

- **Pharmacodynamics is what the drug does to the body**
 - Want to predict what drug concentration/ exposure results in efficacy
 - Demonstrate efficacy at those concentrations in disease model
 - Predicted human efficacious concentration defines human dose target

DMPK: Oral Dosing

To be active, oral drugs must:

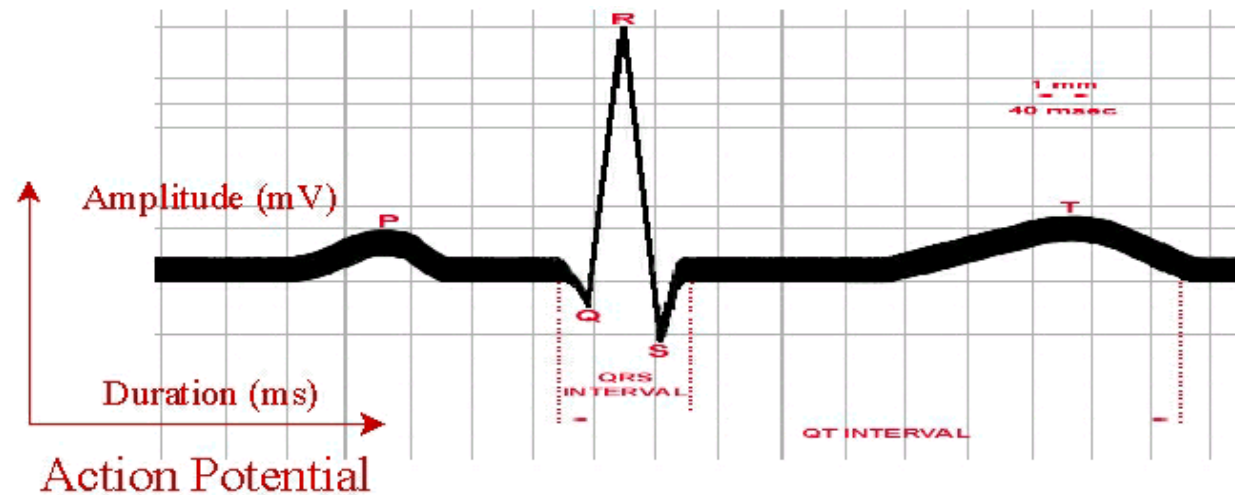
- dissolve
- survive range of pH (1.5-8)
- survive intestinal flora/ fauna
- cross membranes
- survive liver (oxidation and conjugation)
- avoid active transport to bile
- avoid excretion by kidneys
- partition into appropriate place(s)



- **This phase fine tunes properties of the series such that the entire package is present in one or more compounds**
 - Optimisation of all properties, using the same or similar assays as used in Hit-to-Lead
 - Often requires synthesis and in-depth evaluation of a large number of related analogues
 - Can be expensive owing to the large number of complex studies required
 - These are usually built in at the latter stages owing to low through-put or high cost of such tests which limits numbers of compounds that can be tested
- **Candidate Criteria should be clearly defined**

Additional studies: QT effects & Genotoxicity

- **30% of post-marketing withdrawal of drugs has been attributed to QT prolongation**
 - Majority inhibit the I_{Kr} current mediated by hERG channels
 - I_{Kr} - 'rapid' delayed rectifier current that conducts potassium (K^+) ions out of the muscle cells of the heart



- **Ames test used to assess the mutagenic potential of chemical compounds**
 - Uses well characterised bacterial strains

Additional studies: CMC (Chemistry Manufacturing and Control)

- **Cost of goods**
 - Criterion will depend on dose and therapy area
- **Chemical stability**
 - Check for decomposition in a range of situations to mimic the intestine
- **Shelf-life stability**
 - Compound subjected to humid conditions at elevated temperature to mimic accelerated shelf storage
- **Crystal form**
 - Salt selection (for acidic or basic drugs) or co-crystal
 - Potential look at polymorphic forms to find the most stable polymorph
 - Solubility and F% then measured with correct final form

Drug discovery into drug development

- **A candidate drug is an experimental “drug”**
- **Still a lot to do before it is a drug**
 - No direct evidence it is safe in animals let alone humans
 - No direct evidence it works in real patients
- **Clinical trials**
 - Long, expensive and littered with failure
- **Main causes of clinical failures are**
 - Efficacy – it doesn’t work! (~55%)
 - Safety – unexpected toxicity (~30%)
- **Safety and efficacy – risk/ benefit – assessed by independent stringent regulatory authority**

- **Preclinical development – 12-24 months**
 - GLP rodent and non-rodent and other safety studies, GMP scale-up
- **Phase I – 3-9 months**
 - Healthy volunteers – safety, PK and maximum dose (20-80 male volunteers)
- **Phase IIa – 6-12 months**
 - Patients – safety and efficacy (100-200 patients)
- **Phase IIb – 12-24 months**
 - Patients – safety and efficacy (200-500 patients)
- **Phase III – 2-4 years**
 - Patients – safety efficacy (up to 3,000 patients)
- **Phase IV – Post approval studies to monitor for safety**

- **Drug Discovery is all about finding the ‘sweet spot’ where all properties necessary for a drug come together**
- **Candidate drug and target must be consistent with the TPP**
- **The principles and art of drug discovery are consistent regardless of biological target or disease**
 - Specific disease biology knowledge necessary
- **Regulators or patients do not care whether a drug was discovered in Pharma or Academia**
 - The data packages, compound/ science quality have to be the same
- **Drug discovery is multi-disciplinary and highly collaborative in nature**
 - Asking for help and advice is not a weakness

- Pharmacokinetics and metabolism in drug design – D.A.Smith, H. van de Waterbeemd and D. K. Walker. Wiley-VCH, ISBN 3-527-30197-6
- The organic chemistry of drug design and drug action. R.B.Silverman. Academic Press, ISBN 978-0126437324
- Goodman and Gilman's The pharmacological basis of therapeutics. L. Brunton, J. Lazo, K. Parker. McGraw-Hill Medical, ISBN 978-0071422802
- Drug Discovery handbook – S. C. Gad (Ed). Wiley (2005), ISBN ISBN 0-471-21384-5
- Principles of early drug discovery – S. Rees et al, Br J Pharmacol. 2011 Mar; 162(6): 1239–1249
- Drug discovery process - <https://www.youtube.com/watch?v=DhxD6sVQEYc>



Grand Challenges
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Think with the end in mind: TPP

Work on the right biology

Many options to find chemistry – validation of series is key

Optimisation is not just about potency

DMPK and safety too determine whether you have a drug

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Practical lessons from a lifelong quest to discover medicines

- You need a thick skin: the science will beat you with a stick every day
- Stay humble
- And yet: you are constantly exposed to magic; the science is continually mind-blowing
- Helping make drugs that save people's lives: indescribable
- Stay positive
- Keep learning, try new things, think big, take chances
- Biology is what usually gets you
- Work hard, sweat the details, avoid unforced errors
- Structure can help you—and so can cellular phenotype
- Integrate everything: technologies, disciplines, hypotheses, mindsets, literature data, you name it
- Teams make drugs. Great teams behave in predictable ways and thrive in supportive environments. But teams are complex—and the interpersonal stuff is at least as important as the science
- Luck is a big factor in all our successes. Did I mention that you need to stay humble?
- Good fortune in our lives enables all we accomplish—pay it forward

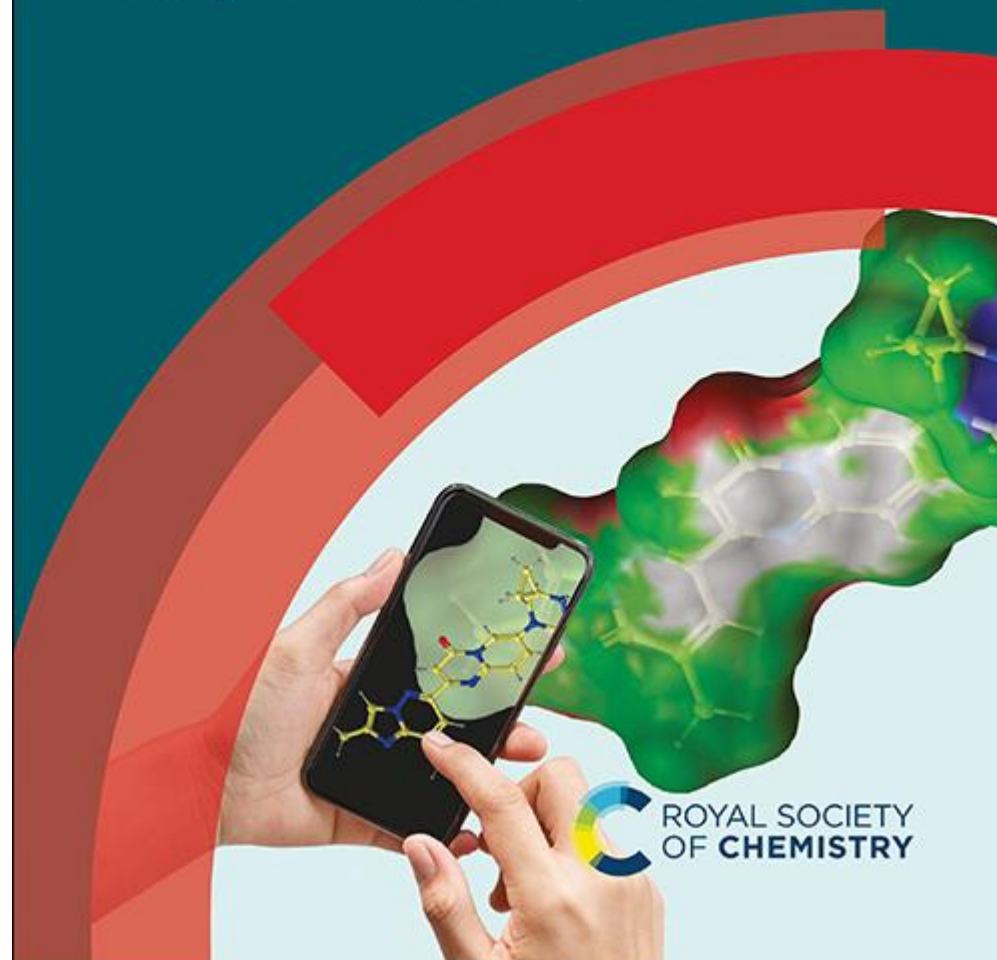


2nd Edition

The Handbook of Medicinal Chemistry

Principles and Practice

Edited by Andrew M. Davis and Simon E. Ward



Abbreviations

- **Target:** biological molecule, usually a protein, whose function is to be modulated
- **Hit:** small molecule with promising activity in a primary screening assay
- **Lead:** molecule with suitable properties for optimization to give a drug candidate
- **Candidate:** molecule selected for development – no further optimisation
- **Asset:** a molecule or project with value associated with its stage in the pipeline
- **Pipeline:** an organisation's ensemble of projects at various stages
- **TPP:** Target Product Profile – description of the required drug
- **KO:** Genetic Knockout (usually mouse)
- **FTIH:** First Time in Human (aka Phase I Clinical Trial) – safety assessment in healthy volunteers
- **PoC:** Proof of Concept (aka Phase II Clinical Trial) – compound/mechanism gives desired clinical response
- **NOAEL:** No Observed Adverse Effect Level
- **MTD:** Maximum Tolerated Dose
- **MW/MWt:** Molecular Weight
- **MED:** Minimum Effective Dose
- **HTS:** High Throughput Screen
- **SAR:** Structure-Activity Relationship – relationship between the compound's structure and its activity
- **DMPK:** Drug Metabolism & Pharmacokinetics – how the drug is handled by the body
- **ADMET:** Absorption, Distribution, Metabolism, Elimination or Excretion, Toxicology
- **PD:** Pharmacodynamics – the activity of the drug in an animal or patient
- **DDI:** Drug-Drug Interaction – the effect of one drug on another, e.g. blocking or increasing metabolism
- **LO:** Lead Optimisation