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Risk Assessment: does it need redefinition?

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IRAS

Risk assessment of chemicals in relation to human health

- hazard identification
- hazard characterisation (dose-response)
- exposure assessment
- risk characterisation



toxicological risk assessment

“Classical:”

animal experiments: quantification (LD50, NOEL)

extrapolation to human situation (safety factors)

threshold vs. non-threshold extrapolation approaches

establish safety standards for human exposure

uncertainties

mechanisms of action (in vitro methods)



BASIS OF TOXICITY

- interaction of chemical with biological system leading to adverse effects
- molecular target:
“toxicological receptor”



BASIS OF TOXICITY

interactions are determined by:

- structural properties of compound

AND

- structural properties of target molecule

nature of physico-chemical reactions:

primary mechanism of toxic action



BASIS OF TOXICITY

following the primary interaction:

pathophysiological changes;

finally resulting in clinically manifest toxicity

complete process: *mode of action*



PARADIGM:

toxicity is determined by:

the *critical*

concentration and time of exposure (dose metric) to

the *critical compound* (metabolite?) on

the *critical site of action*



in vitro toxicology

“Classical” *in vitro* toxicology:
finding concentrations (not dose)
need to extrapolate to intact organism
lack of biotransformation/ kinetics
concentrated on cytotoxicity, rather than on
mechanisms of importance *in vivo*



basal cytotoxicity

disturbance of house-hold functions of cells:

- * integrity of cellular membranes
- * intactness of energy supplies
- * maintenance of cellular compartments



cell-specific functional disturbance

differentiated functions of cells:

e.g.

neuronal activity

albumin production

barrier function / specific transport

etc.

etc.



in vitro toxicology

Mechanisms of action:

- may be studied in detail in *in vitro* systems
- may give the intrinsic ability of a chemical to cause a certain toxic action



use of *in vitro*-derived data in hazard and risk assessments

e.g. reactivity + basal cytotoxicity:

answers towards: corrosivity, irritancy

reactivity towards DNA: genotoxicity



developments in cellular biology

general cellular biology

differentiation, de-differentiation, apoptosis

cell-cell interactions

receptor-mediated processes

signaling processes

metabolic changes

genomics, proteomics, metabonomics

adaptation processes



developments in cellular toxicology

biomarkers for cellular injury:

- oxidative stress
- disturbances in cellular membranes
- disturbances in intercellular communication
- cytokine profiles
- DNA damage
- etc.

cell- or tissue-specific effects



use *in vitro* assays in toxicodynamics

to determine (aspecific) cytotoxicity
to determine tissue-specific toxicity
to find relevant concentrations for
functional cellular disorders (e.g. EC₂₀
values):

‘response surrogates’



developments in structure-activity relationships

chemical-biological interactions:

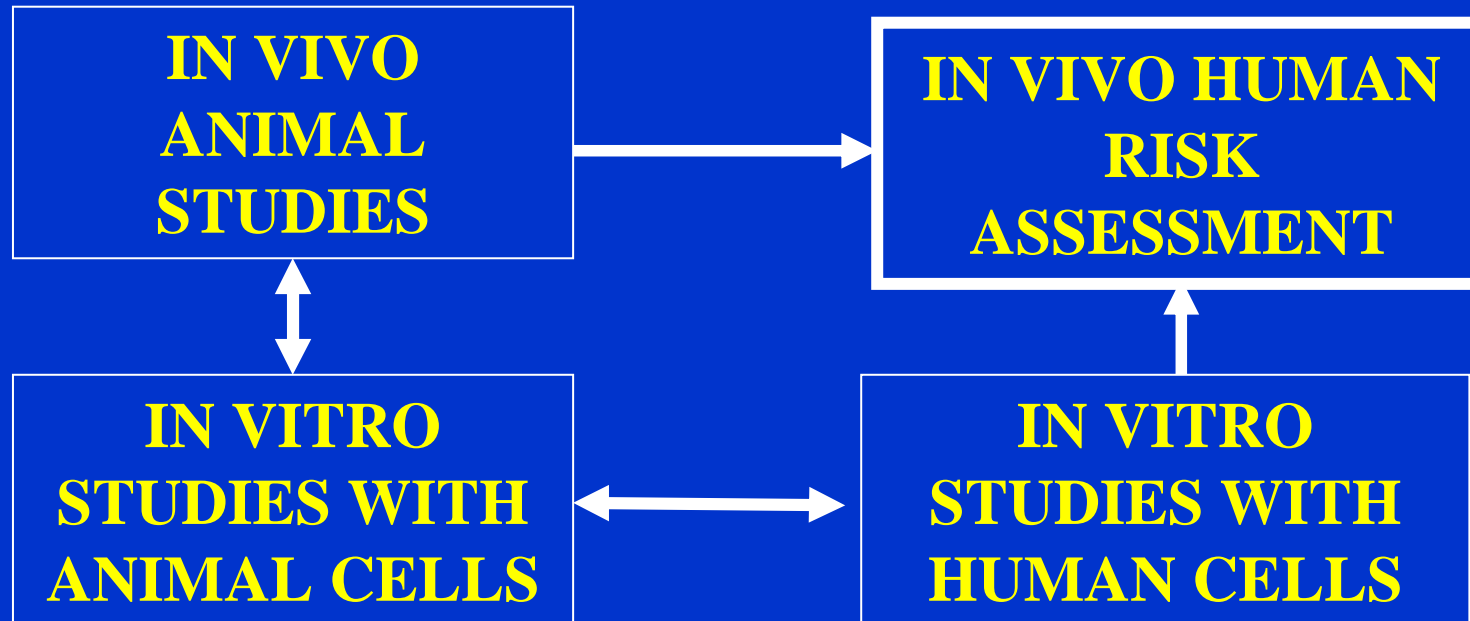
- covalent binding
- receptor binding
- hydrophobic/hydrophilic interactions

knowledge-based computer systems

QSARs



THE PARALLELOGRAM APPROACH



clinical effects

possibility of chemical to reach targets

- in appropriate amount or concentration
- in appropriate form (bound, metabolite?)

depends on:

- **exposure**
- **biokinetic behaviour:** absorption, distribution, metabolism, elimination



biokinetics

physicochemical properties

- of the chemical
as well as those
- of the tissues: **PARTITIONING**

AND



biokinetics

the physiology of the organism

e.g.

structure of the gut, nose, lungs, skin

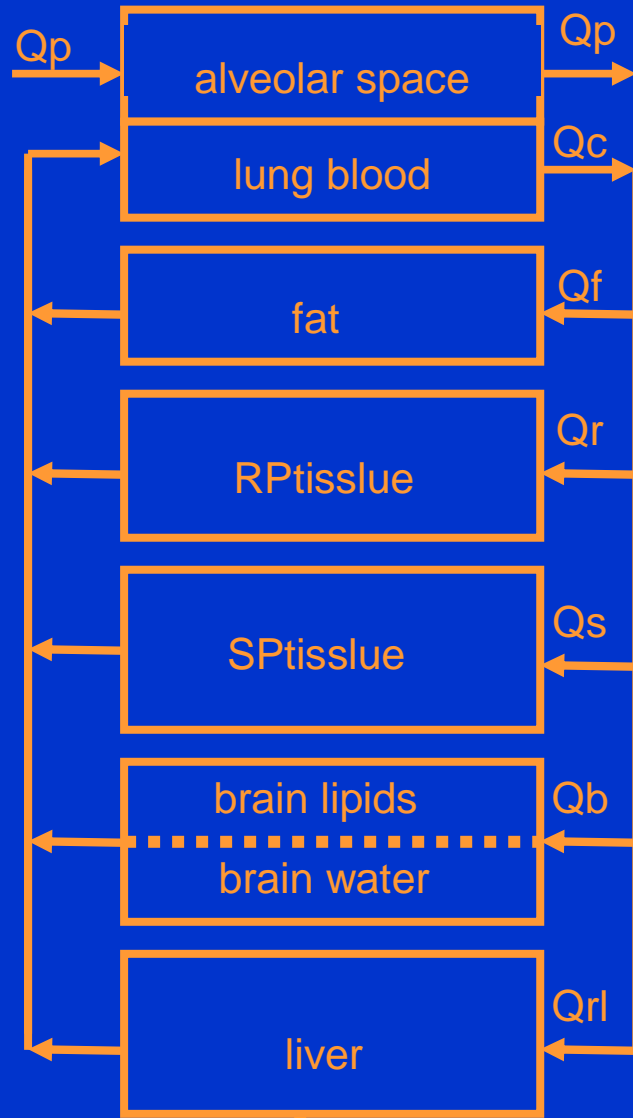
blood flows

metabolising systems

kidney function

etc.





V_{max}
 K_m



alternative methods

Many possibilities for studying mechanisms on the cellular and molecular level

However: use in risk assessment:

- very limited possibilities for a one-to-one replacement
- need for using information from different sources: **integration**



Integrated testing schemes: elements

physico-chemical properties

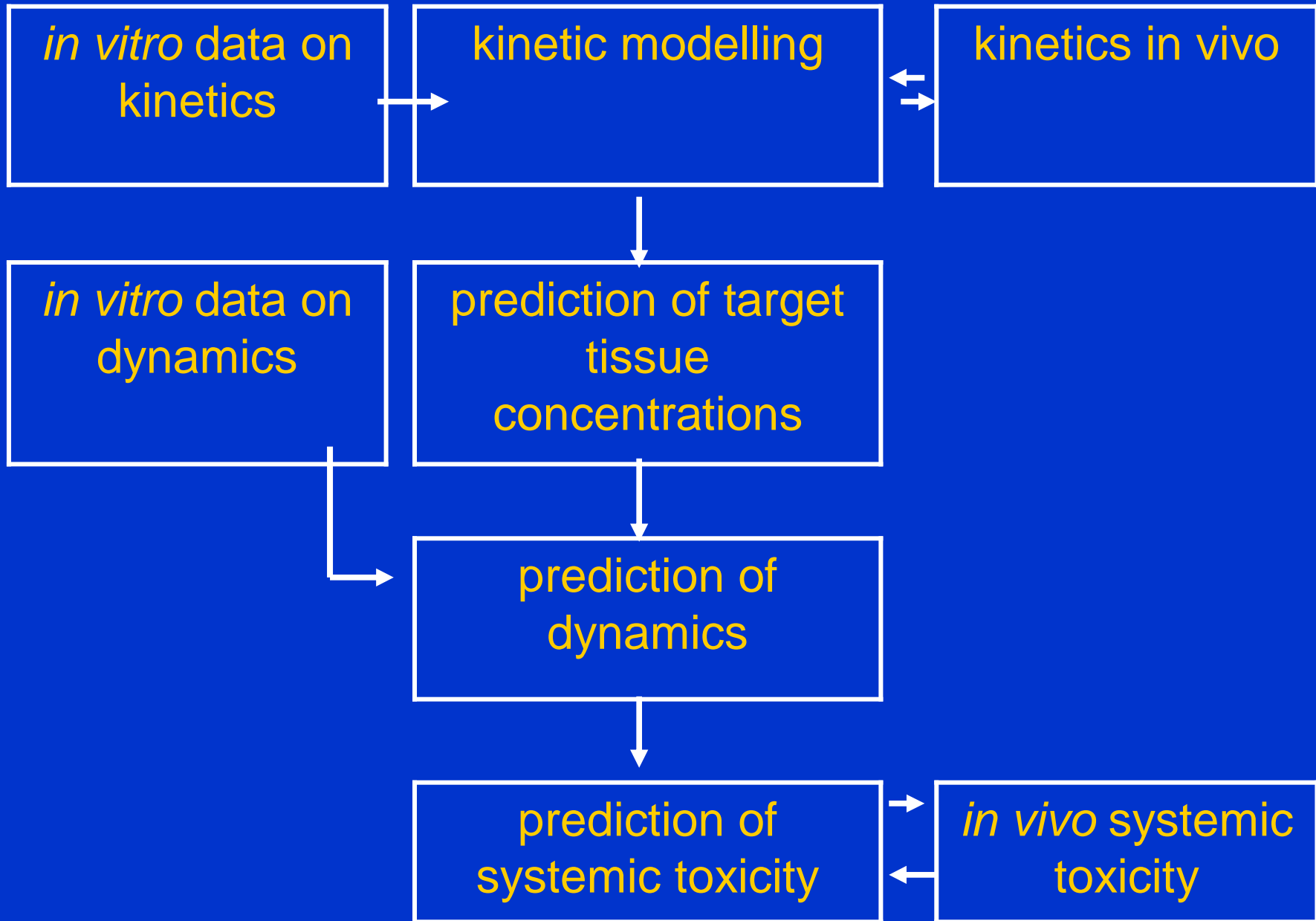
QSARs

in vitro data

kinetic and dynamic modelling

evaluation against *in vivo* data





the case of acrylamide

In vitro data on toxicity

Kinetic model to predict dose causing toxic effects



**determine basal
cytotoxicity
(e.g. Nutral Red test):**

determine EC₅₀ and EC₂₀

**determine tissue-specific
alterations, e.g. in neuronal cells:**

-morphology

-changes in cell physiology

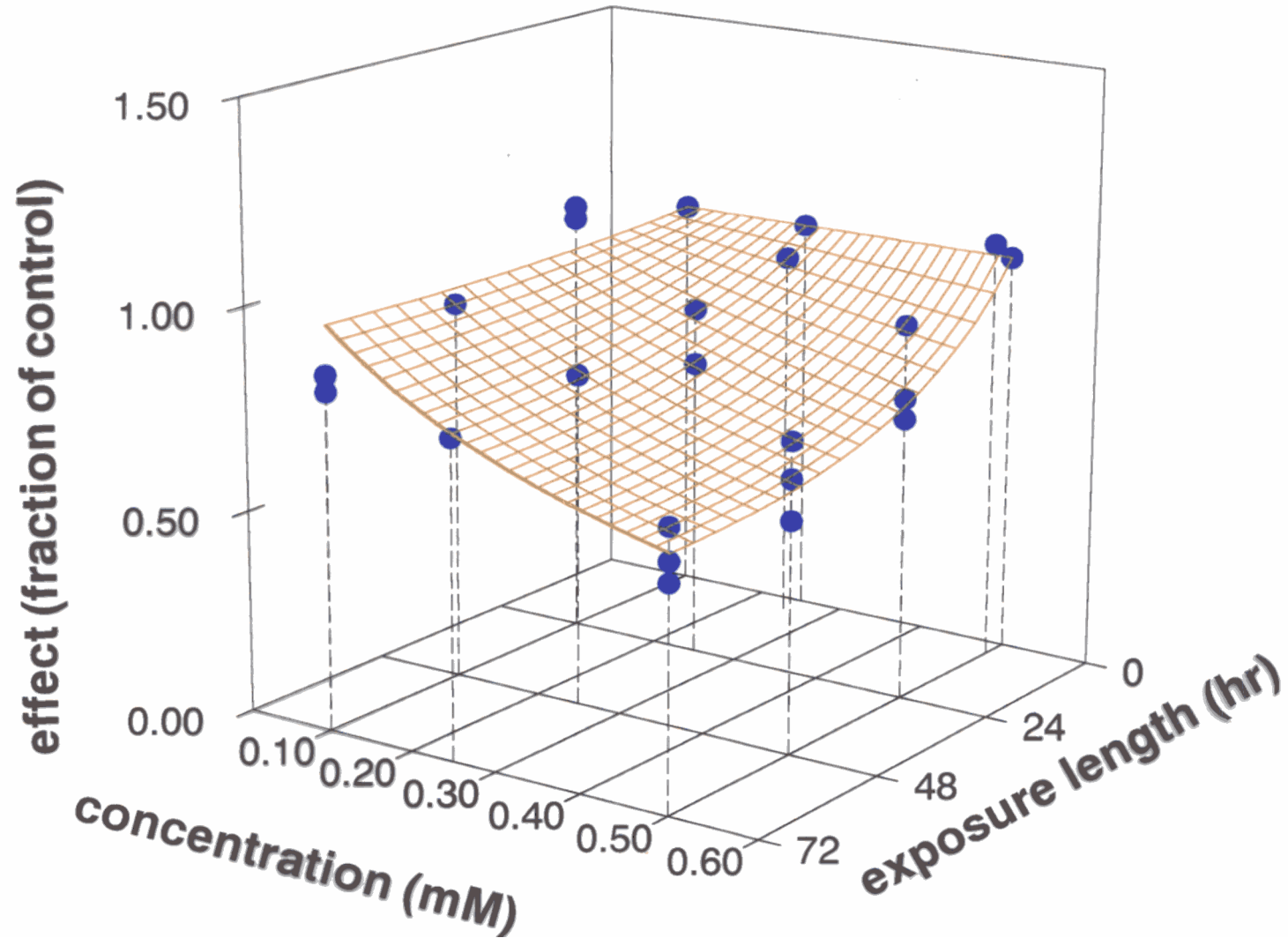
-neurochemical alterations

determine relevant EC₅₀ and EC₂₀

**-determine ratios of EC₅₀s for basal/neurotoxic effects;
-decide on specific neurotoxicity action
-use EC₂₀-neurotoxicity as a basis for a LOAL calculation,
making use of the integrated model described before.**



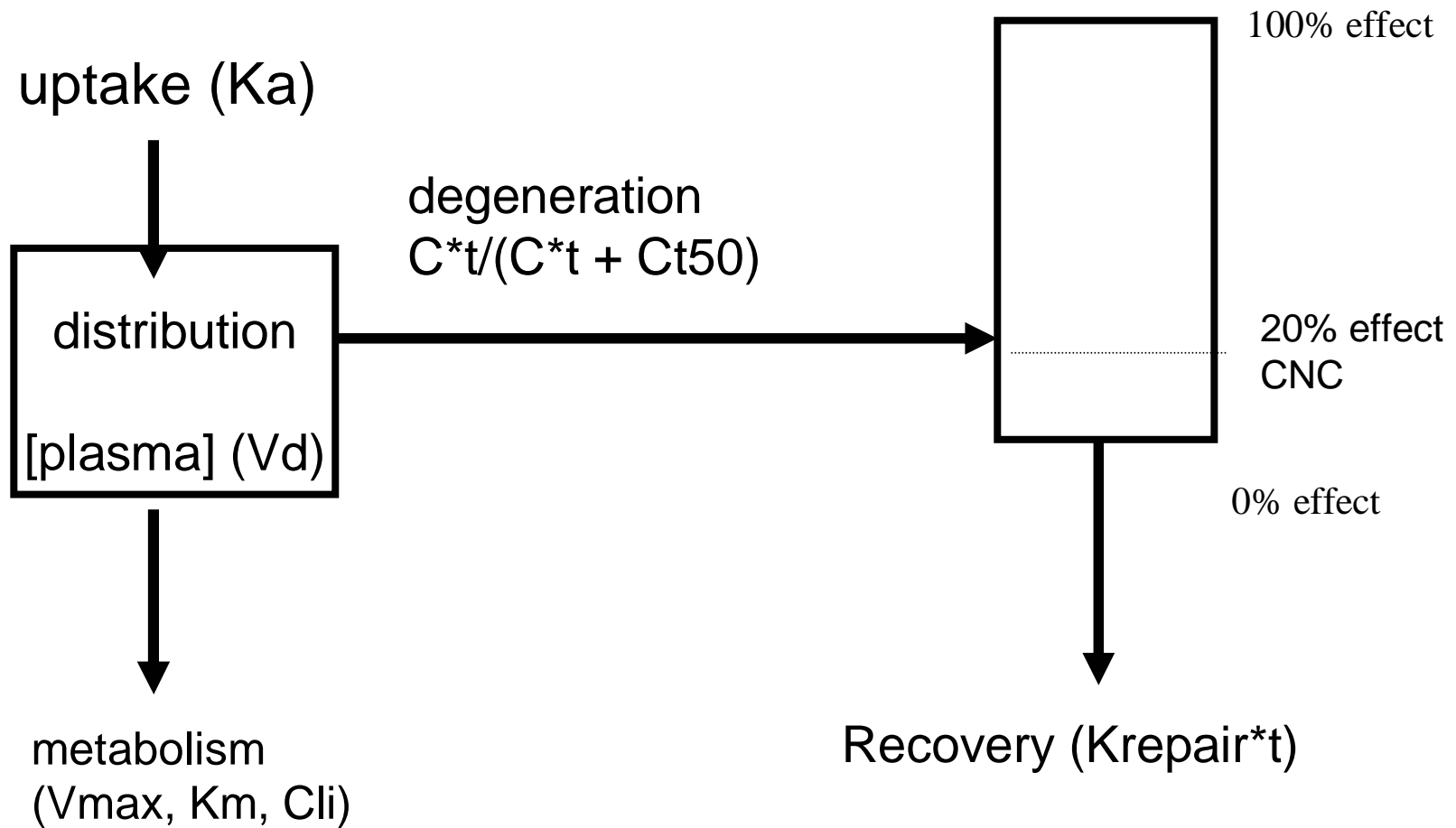
acrylamide's effects on # of neurites/cell

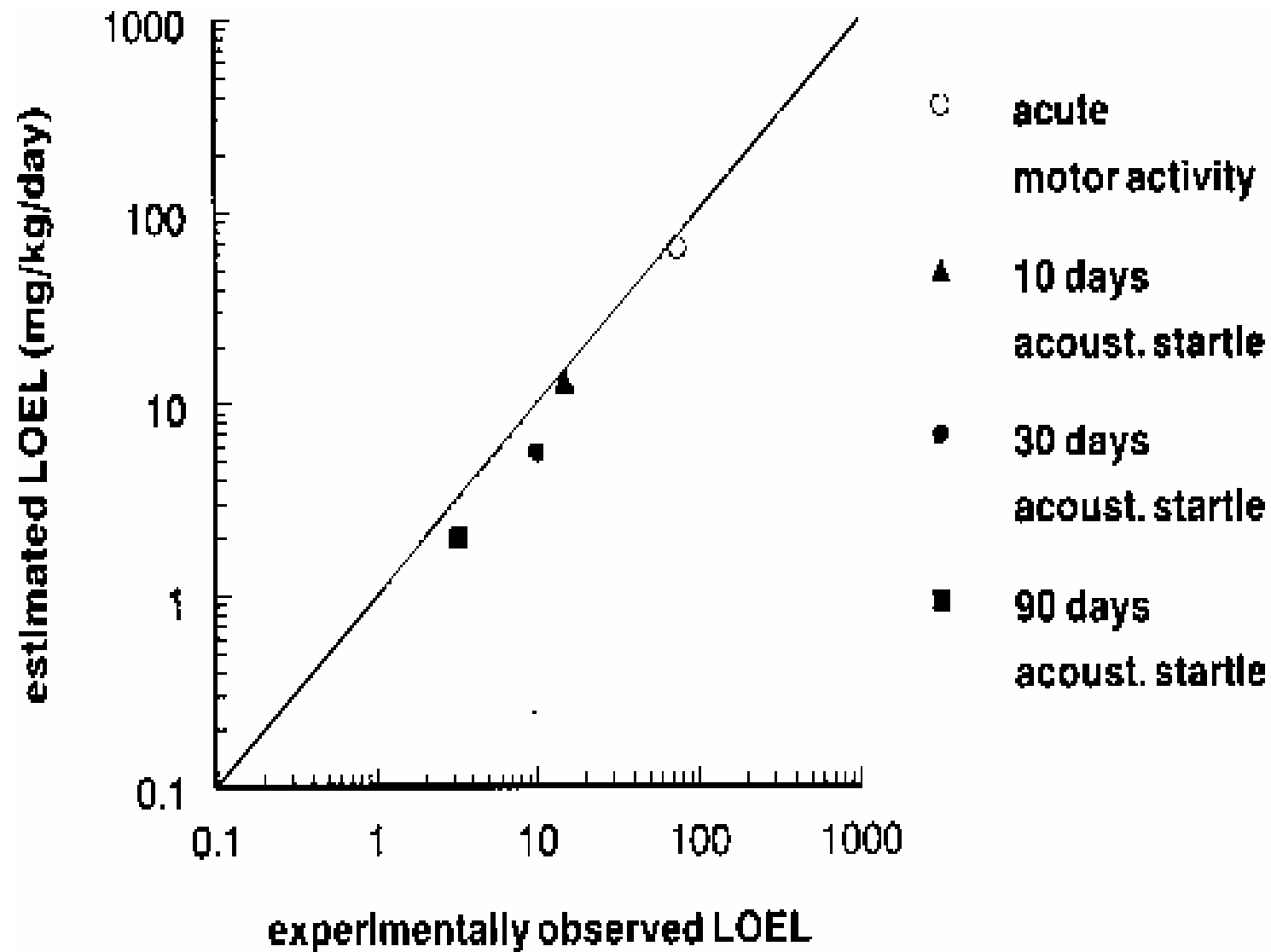


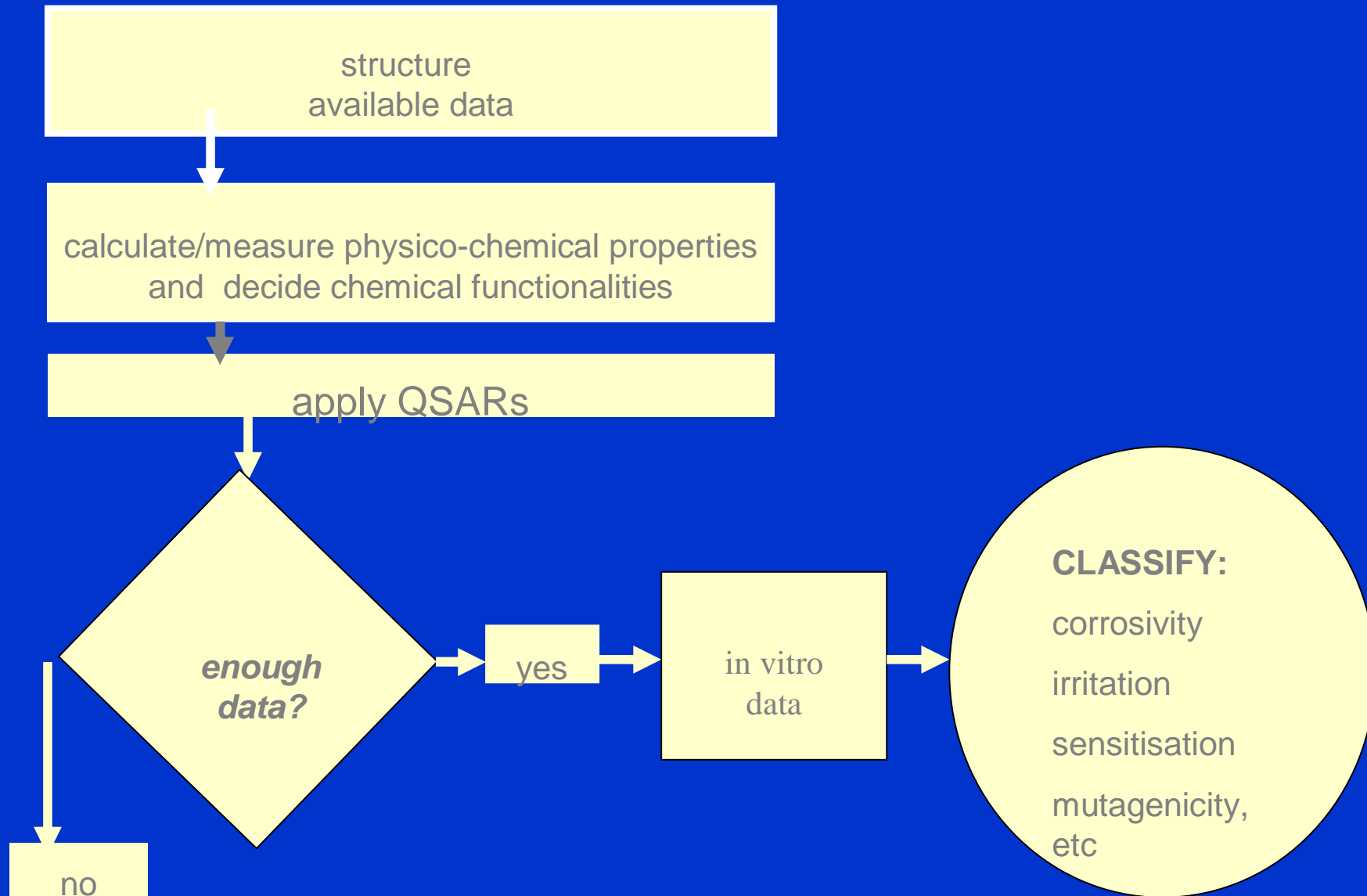
KINETIC-DYNAMIC MODELLING

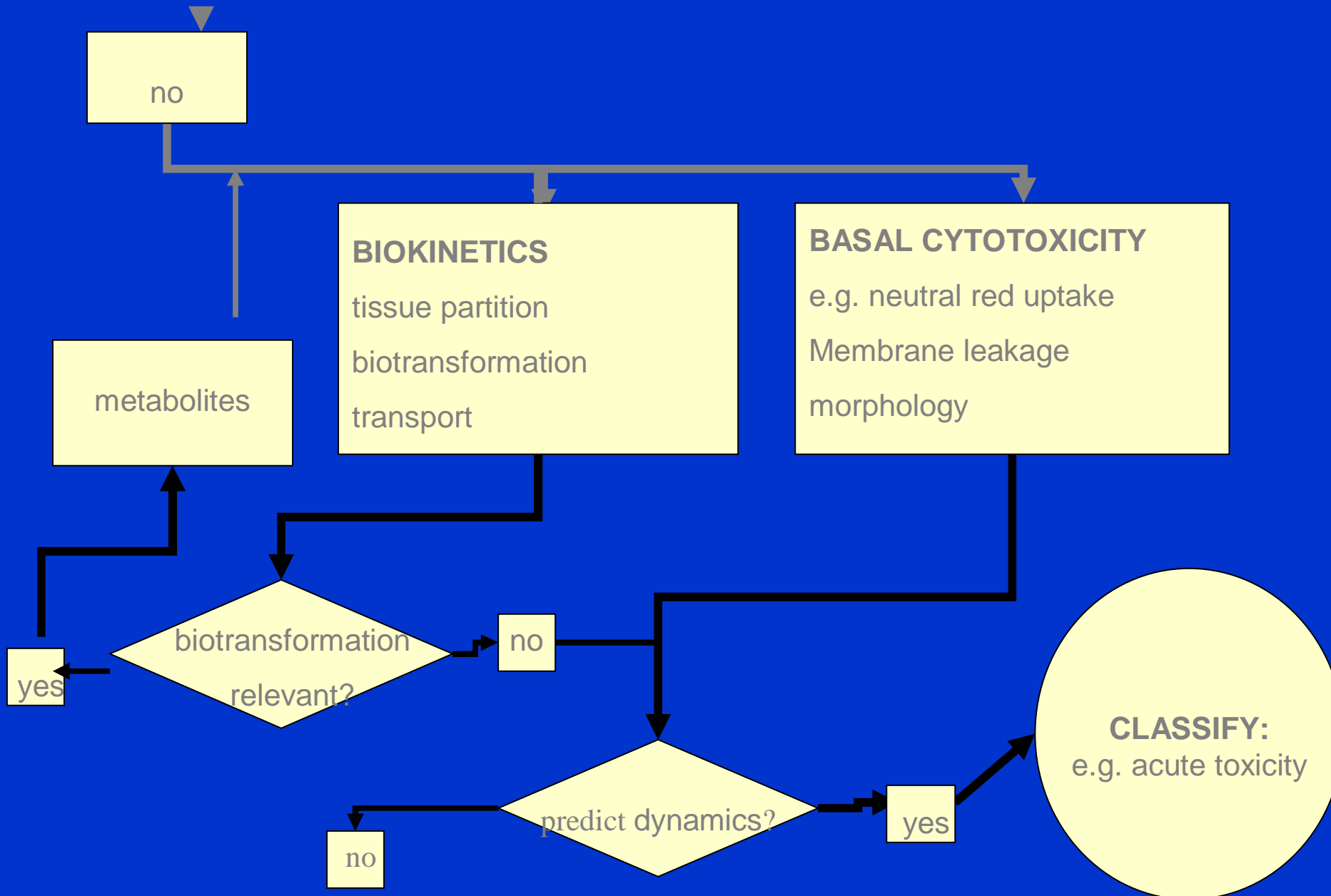
KINETICS

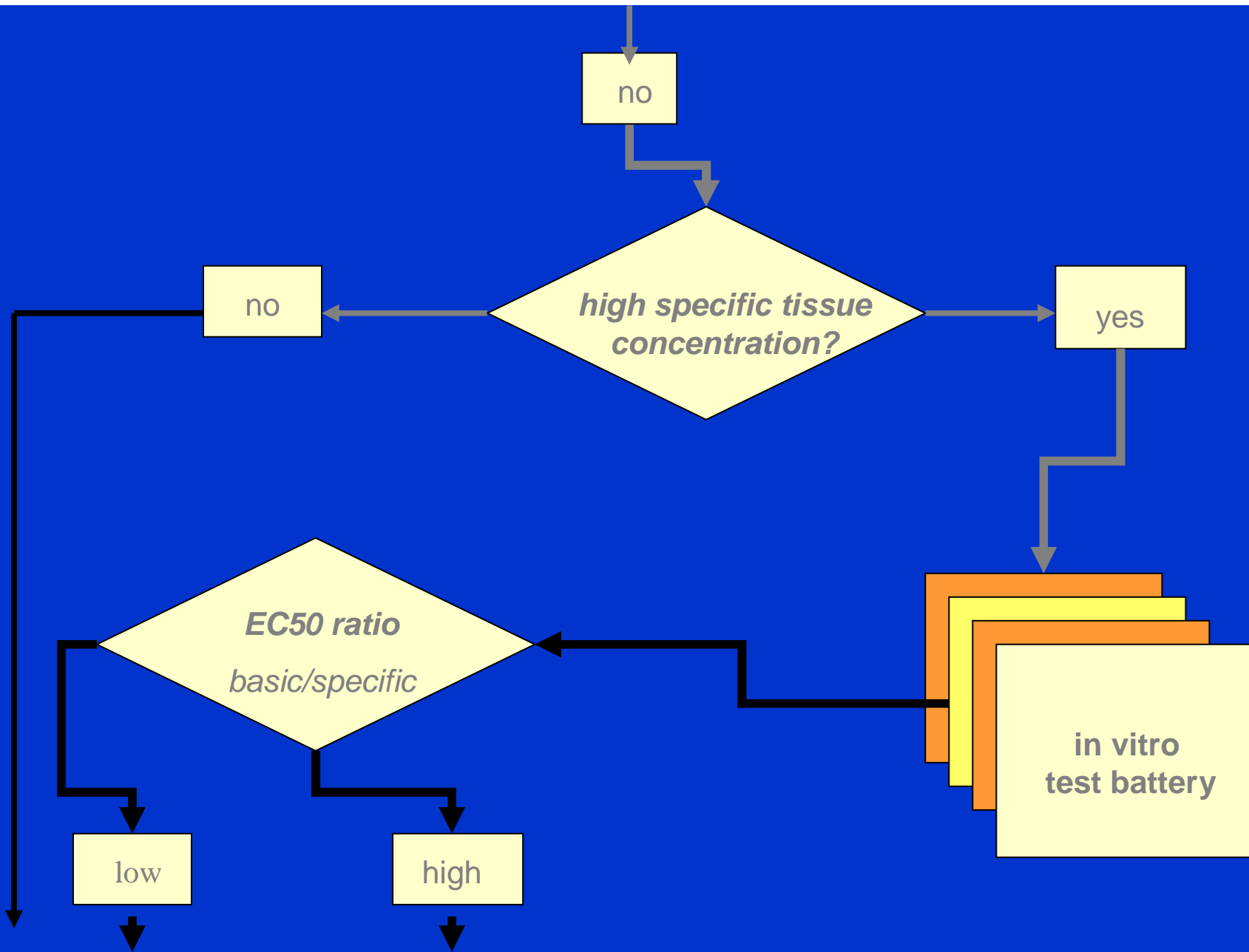
DYNAMICS

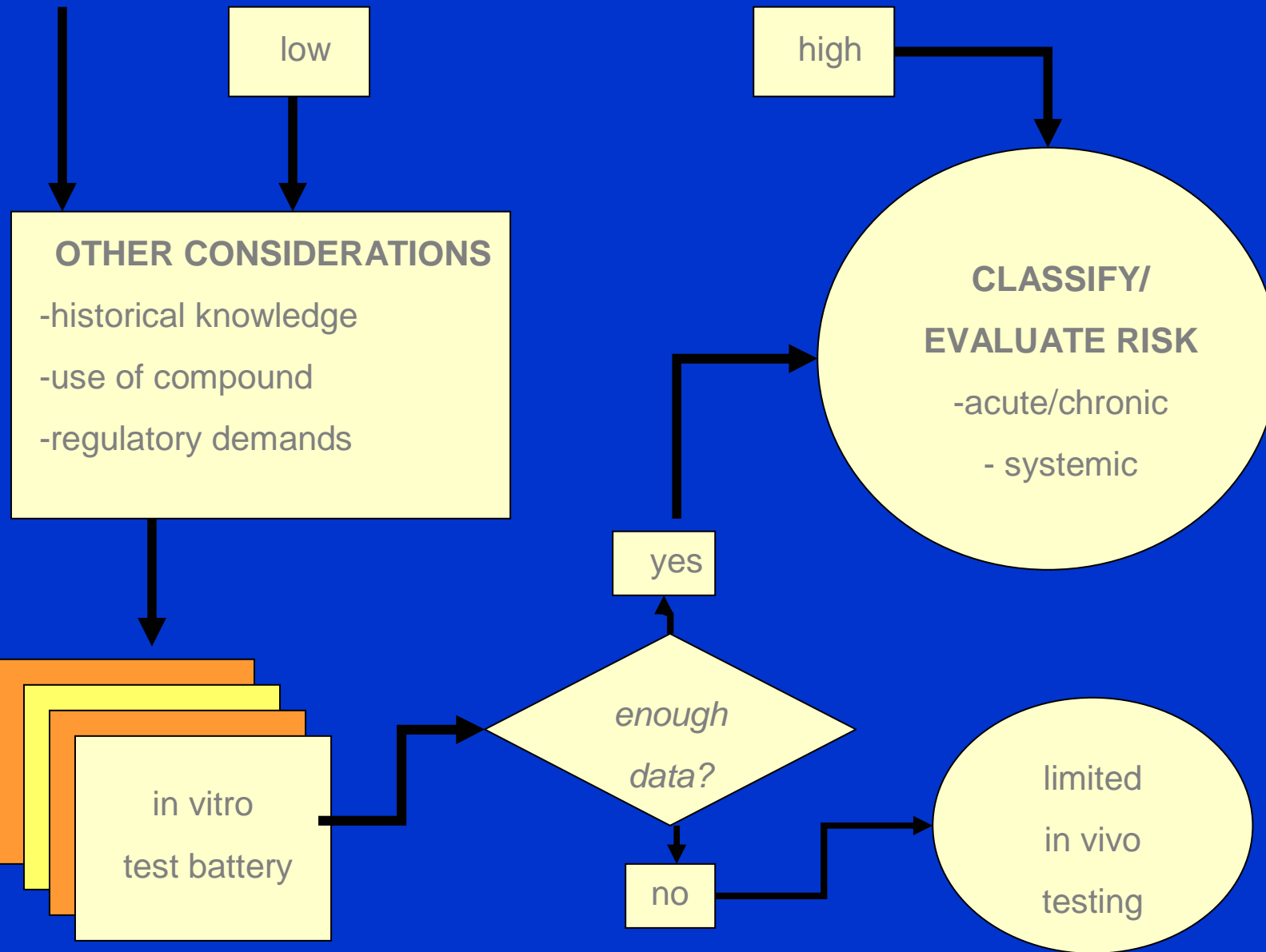












Basic data set

EXTERNAL DOSE

TOXIC RESPONSE

Physiologically-based kinetic parameters

EXTERNAL DOSE

INTERNAL DOSE

TOXIC RESPONSE

Physiologically-based kinetic model

EXTERNAL DOSE

INTERNAL DOSE

TARGET ORGAN DOSE

TOXIC RESPONSE

Physiologically-based kinetic model plus local target organ metabolism

EXTERNAL DOSE

INTERNAL DOSE

TARGET ORGAN DOSE

TARGET ORGAN METABOLISM

TOXIC RESPONSE

Biologically-based dose-response model

EXTERNAL DOSE

INTERNAL DOSE

TARGET ORGAN DOSE

TARGET ORGAN METABOLISM

TARGET ORGAN RESPONSES

TOXIC RESPONSE

in vitro data in risk assessment: perspectives

use as stand-alone methods

use of in vitro data in integrated schemes

challenges: choice of the most relevant
battery of tests

can we predict mode of action/target organ
on the basis of structure



CONCLUSIONS

integration of *in vitro* data in risk evaluation is possible, provided that:
biokinetics are taken into account
(absolute necessity)

integration of all available data in a stepwise (hierarchical) approach will improve the transparency and efficacy of the risk assessment process

there are many possibilities for the use of non-animal data in this process

