

Ecotoxicological applications of Dynamic Energy

Budget theory *

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Abstract

The Dynamic Energy Budget (DEB) theory for metabolic organisation specifies quantitatively the processes of uptake of substrate by organisms and its use for the purpose of maintenance, growth, maturation and reproduction. It applies to all organisms. Animals are special because they typically feed on other organisms. This couples the uptake of the different required substrates, and their energetics can, therefore, be captured realistically with a single reserve and a single structure compartment in biomass. Effects of chemical compounds (e.g. toxicants) are included by linking parameter values to internal concentrations. This involves a toxico-kinetic module that is linked to the DEB, in terms of uptake, elimination and (metabolic) transformation of the compounds. The core of the kinetic module is the simple one-compartment model, but extensions and modifications are required to link it to DEBs. We discuss how these extensions relate to each other and how they can be organised in a coherent framework that deals with effects of compounds with varying concentrations and with mixtures of chemicals. For the one-compartment model and its extensions, as well as for the standard DEB model for individual organisms, theory is available for the co-variation of parameter values among different applications, which facilitates model applications and extrapolations.

Keywords: Dynamic Energy Budgets, effects on processes, kinetics, metabolism, transformation

1 Introduction

The societal interest in ecotoxicology is in the effects of chemical compounds on organisms, especially at the population and ecosystem level. Sometimes these effects are intentional, but more typically they concern adverse side-effects of other (industrial) activities. The scientific interest in effects of chemical compounds is in perturbing the metabolic system, which can reveal its organisation. This approach supplements ideas originating from molecular biology, but now applied at the individual level; the sheer complexity of biochemical organisation hampers reliable predictions of the performance of individuals. Understanding the metabolic organisation from basic physical and chemical principles is the target of Dynamic Energy Budget (DEB) theory [1, 2]. In reverse, this theory can be used to quantify effects of chemical compounds, i.e. *changes* of the metabolic performance of individuals. This chapter describes how DEB theory quantifies toxicity as a process.

The effects can usually be linked to the concentration of compounds inside the organism or inside certain tissues or organs of an organism. This makes that toxicokinetics is basic to effect studies. The physiological state of an organism, such as its size, its lipid content, and the importance of the various uptake and elimination routes (feeding, reproduction, excretion) interacts actively with toxicokinetics, so a more elaborate analysis of toxicoki-

netics should be linked to the metabolic organisation of the organism [3]. In the section on toxicokinetics, we start with familiar classic models that hardly include the physiology of the organism, and stepwise include modules that do make this link in terms of logical extensions of the classic models.

Three ranges of concentrations of any compound in an organism can be delineated: too-little, enough and too-much. The definition of the enough-range is that variations of concentrations within this range do not translate into variations of the physiological performance of the individual. Some of the ranges can have size zero; such as the too-little-range for cadmium. Effects are quantified in the context of DEB theory as changes of (metabolic) parameters as linked to changes in internal concentrations. These parameters can be the hazard rate (for lethal effects), the specific food uptake rate, the specific maintenance costs, etc. Changes in a single parameter can have many physiological consequences for the individual. DEB theory is used not only to specify the possible modes of action of a chemical compound, but also how the various physiological processes interact. An increase in the maintenance costs by some compound, for instance, reduces growth, and since food uptake is linked to body size, it indirectly reduces food uptake, and so affects reproduction (and development). The existence of the ranges too-little, enough and too-much of concentrations of any compound directly follows from a consistency argument, where no

classification of compounds is accepted (e.g. toxic and non-toxic compounds), and many compounds (namely those that make up reserve) exist that do change in concentration in the organism, without affecting parameter values. An important consequence is the existence of an internal No Effect Concentration (NEC), as will be discussed below.

Effects at the population level are evaluated from those at the individual level, by considering populations as a set of interacting individuals [4-8]. Although DEB-based population dynamics can be complex, particular aspects of population performance, such as the population growth rate at constant environmental conditions, are not very complex. Moreover particular simplifying approximations are possible. The focus of toxicology is typically at time scales that are short relative to the life span of the individual. That of ecotoxicology, however, are much longer, involving the whole life history of organisms; effects on feeding, growth, reproduction and survival are essential and typically outside the scope of toxicology. This has strong implications for the best design of models. While pharmacokinetics models frequently have many variables and parameters, such complex models are of little use for applications at the population level, where a strong need is felt for relatively simple models, but then applied to many species and in complex situations. DEB theory is especially designed for this task.

Chemical transformations are basic to metabolism, and transformations of toxicants are no exception. Compounds that dissociate in water should be considered as a mixture of ionic and molecular forms, and the pH affects that mixture. This makes that the effect of a single pure compound is of rather academic interest; we have to think in terms of the dynamic mixtures of compounds. Because of its strong links with chemical and physical principles, DEB theory has straightforward ways to deal with effects of mixtures. Some of this theory rests on the covariation of parameter values across species of organism and across chemical compounds. Theory on this covariation is implied by DEB theory, and is one of its most powerful aspects.

We first introduce some notions of DEB theory, and then discuss toxicokinetics and effects in the context of DEB theory.

2 The standard DEB model in a nutshell

[Figure 1 about here.]

The standard DEB model concerns an isomorph. i.e. an organism that does not change in shape during growth, that feeds on a single food source (of constant chemical composition), and has a single reserve a single structure and three life stages: embryo (which does not feed), juvenile (which does not allocate to reproduction) and adult (which allocates

to reproduction, but not to maturation). This is in some respects the simplest model in the context of DEB theory, which is thought to be appropriate for most animals. Food is converted to reserve, and reserve to structure. Reserve does not require maintenance, but structure does, mainly to fuel its turnover, see Figure 1. Reserve can have active metabolic functions and serves the role of representing metabolic memory. Reserve and structure do not change in chemical composition (strong homeostasis). At constant food availability, reserve and structure increase in harmony, i.e. the ratio of their amounts, the reserve density, remains constant (weak homeostasis).

The shape (and so the change of shape) is important because food uptake is proportional to surface area, and maintenance mostly to (structural) volume. The handling time of food (including digestion and metabolic processing) is proportional to the mass of food “particles”, during which food acquisition is ceased. The mobilisation rate of reserve to fuel the metabolic needs follows from the weak and strong homeostasis assumptions [9]; a mechanism is presented by Kooijman and Troost [10]. Allocation to growth and somatic maintenance (so to the soma) comprises a fixed fraction of mobilised reserve, the remaining fraction is allocated to maturation (or reproduction) and maturity maintenance.

The transition from the embryo to the juvenile stage (i.e. birth) occurs by initiating assimilation when the maturity exceeds a threshold value, and from the juvenile to the

109 adult stage (i.e. puberty) by initiating allocation to reproduction and ceasing of alloca-
110 tion to maturation at another threshold value for maturity. Reserve that is allocated to
111 reproduction is first collected in a buffer, that is subjected to buffer handling rules (such
112 as spawning once per season, or convert the buffer content into an egg as soon as there is
113 enough).

114 Biomass consists of reserve and structure, and can, therefore, change in chemical com-
115 position (e.g. lipid content) in response to the nutritional condition; maturity has the status
116 of information, not that of mass or energy. Apart from some minor details, the presented
117 set of simple rules fully specify the dynamics of the individual, including all mass and
118 energy fluxes, such as the uptake of dioxygen, the production of carbon dioxide, nitrogen
119 waste and heat. It takes some time to see exactly how; Sousa *et al.* [9] gives a nice evalu-
120 ation. Aging is considered to result from a side effect of Reactive Oxygen Species (ROS),
121 and is so linked to the uptake of dioxygen [11]. A high food uptake results in a large amount
122 of reserve, so a high use of reserve, a high uptake of dioxygen, an acceleration of aging
123 and a reduction of life span. The induction of tumours is also linked to the occurrence of
124 ROS, and other reactive molecules, such as mutagenic compounds. This gives a natural
125 link between aging and tumour induction.

The standard DEB model has been extended into many directions for the various purposes. The allocation rule to the soma, for instance, can be refined to allocation to various body parts (e.g. organs), where growth of each body part is proportional to the allocated reserve flux minus what is required for maintenance of that part. Rather than using fixed fractions of the mobilised reserve, the fractions can be linked to the relative workload of the body part. This allows a dynamic adaptation of the body parts in interaction with their use. Tumours can be considered as body parts, and the “workload” of the tumour is the consumption of maintenance. This formulation produced realistic predictions of the effects of caloric restriction on tumour growth, and of the growth of tumours in young versus old hosts [12]. This approach can be extended to various types of tumours, where tumour growth is not linked to that of the whole body, but that of a particular body part. Many tumours result from destruction of local cell-to-cell communication, rather than from genotoxic effects, but these different routes have similar dynamics.

Other types of extensions of the standard DEB model concern the inclusion of variations in chemical composition of food (with consequences for the transformation of food into reserve) and size-dependent selection of different food items. For example many herbivores are carnivores when young. Animals are special because they feed on other organisms. Most other organisms take the food-compounds (energy source, carbon source, nitrate,

phosphate) that they need independently from the environment, which necessitates the inclusion of more than one reserve; see Kooijman and Troost [10] for the evolutionary perspectives. Most micro-organisms grow and divide, and don't have the three life stages delineated by the standard model. This makes that their change in shape hardly matters and that surface areas can be taken proportional to volumes, which simplifies matters considerably. The partial differential equations that are required to described the physiologically structured population dynamics of isomorphs then collapse to a small set of ordinary differential equations. Plants, on the other hand, require at least two types of structure (roots and shoots) and have a complex adaptive morphology (i.e. surface area-volume relationships); their budgets are most complex to quantify.

3 Family of toxicokinetic models

Originally (before the 1950's) the focus of toxicology, i.e. the field that gave rise to ecotoxicology, was on medical applications of compounds in a pharmacological context. Subjects where given a particular dose, and the interest is in the redistribution inside the body, and in transformation and elimination. The aim is to reach the target organ and to achieve a particular effect that restores the health or well-being of the subject. A closely related interest that developed simultaneously was health protection (disinfection and food protection

161 products), with the purpose of killing certain species of pathogenic organism (especially
162 microorganisms), or to reduce their impact.

163 After the 1950's ecotoxicology began to flourish and gradually became more independent
164 of toxicology, where the initial focus was in positive and (later) negative effects of biocides.
165 This came with extensions of the interest in the various uptake and elimination routes
166 that are of ecological relevance, and the environmental physical-chemistry of transport
167 and transformation. The aim is to kill particular species of pest organism locally (insects,
168 weeds), and to avoid effects on other, non-target, species (crop and beneficial species).

169 After the 1970's the interest further generalised to an environmental concern of avoiding
170 effects of pollutants on organisms, with an increasing attention for (bio)degradation of
171 compounds that are released into the environment, coupled to human activity [13].

172 This historic development and branching of the interest in toxico-kinetics came with a
173 narrowing of the focus on a particular aspect of toxico-kinetics in the various applications
174 that, we feel, is counter-productive from a scientific point of view. The purpose of this
175 section is an attempt to restore the coherence in the field, by emphasising the general
176 eco-physiological context and the relationship of the various models with the core model
177 for toxico-kinetics: the one-compartment model. The more subtle models account for the
178 interaction with the metabolism of the organism, which involves its metabolic organisation.

We here focus on the logical coherence of toxico-kinetics, bio-availability and metabolism, including effects (= changes in metabolism). It is not meant to be a review. For recent reviews on toxico-kinetic models, see o.a. Barber [14] and Mackay and Fraser [15]. See Table 1 for a list of frequently used symbols.

[Table 1 about here.]

3.1 One-compartment model

[Figure 2 about here.]

The core model in toxico-kinetics is the one-compartment model, see Fig. 2. It states that the uptake rate is proportional to the environmental concentration c , and the elimination rate is proportional to the internal concentration Q :

$$\frac{d}{dt}Q = k_e (P_{0d}c(t) - Q) \quad (1)$$

$$Q(t) = Q(0) \exp(-tk_e) + k_e P_{0d} \int_0^t c(s) \exp((s-t)k_e) ds \quad (2)$$

$$= Q(0) \exp(-tk_e) + P_{0d}c (1 - \exp(-tk_e)) \quad \text{for } c \text{ is constant} \quad (3)$$

where $P_{0d} = Q(\infty)/c$ is the BioConcentration Factor (BCF) and k_e the elimination rate.

The product $k_e P_{0d}$ is known as the uptake rate, and is frequently indicated with k_u , which

is misleading because its units are $\text{d}^{-1} \text{m}^3 \text{C-mol}^{-1}$. Even if we would work with kg rather than C-mol, and the specific density of the organism equals 1 kg dm^{-3} , the BCF is not dimensionless [1]. It is typically more convenient to work with molalities in soils (mol kg^{-1}), and with molarities (mol l^{-1}) in water. Molalities give the uptake rate the units $\text{d}^{-1} \text{g C-mol}^{-1}$. Many workers use gram rather than mole to quantify the compound, but this choice is less practical to compare the toxicity of different compounds. The elimination rate k_e has dimension ‘per time’ and is independent of how the compound is quantified; contrary to the uptake rate, the elimination rate can be extracted from effect data and determines how fast effects build up in time, relative to the long-term effect level.

The concentrations c and Q must obviously exist, meaning that the environment and the organism are taken to be homogeneous. This condition can be relaxed without making the model more complex by allowing a spatial structure (such as organs), and an exchange between the parts that is fast relative to the exchange between the organism and the environment. Other implicit assumptions of the one-compartment model are that the organism does not change in size or in chemical composition, so changes in food availability must be negligible. The (bio)availability of the compound remains constant. So transformations can be excluded, and the environment is large relative to the organism and well-mixed. Sometimes, e.g. in the case of a large fish in a small aquarium, this is not true and the

209 dynamics of the concentration in the organism and the environment should be considered
 210 simultaneously: the 1-1 compartment model [16]. These restrictions will be removed below.

211 Since rates generally depend on temperature, and temperature typically changes in
 212 time, the elimination rate k_e can change in time as well [17]. In the sequel we will dis-
 213 cuss rates of metabolism, and like all rates, these also depend on temperature, frequently
 214 according to the Arrhenius relationship [1].

215 3.2 Multi-compartment models

216 If transport inside the organism is not fast, relative to the exchange with the environment,
 217 multiple-compartment models should be considered [18, 19]. If exchange with the environ-
 218 ment is only via compartment number 0, the change in the concentrations in the nested
 219 compartments number 0 and 1 is

$$\frac{d}{dt}Q_0 = k_e (P_{0d}c - Q_0) + k_{10}Q_1 - k_{01}Q_0; \quad \frac{d}{dt}Q_1 = k_{01}Q_0 - k_{10}Q_1 \quad (4)$$

220 where k_{01} and k_{10} are the exchange rates between the compartments, see Fig. 2. The
 221 partition coefficient between the compartments equals $P_{10} \equiv Q_1(\infty)/Q_0(\infty) = k_{01}/k_{10}$,
 222 while $P_{0d} \equiv Q_0(\infty)/c$ remains unchanged. In many cases whole body measurements are
 223 used. If M_0 and M_1 denote the masses of compartment 0 and 1, the whole body partition
 224 coefficient with the environment amounts to $P_{+d} = (M_0 + M_1P_{10})P_{0d}/M_+$, with $M_+ =$

225 $M_0 + M_1$ is the whole body mass. This rather complex behaviour of the whole body partition
226 coefficient can be a source of problems in fitting models to data. In many practical cases
227 it is not possible to identify the compartments and to measure the concentrations in these
228 compartments directly. The use of multi-compartment models cannot be recommended in
229 such cases.

230 This extension still classifies as a transport model, so in a clean environment ($c = 0$), the
231 organism will loose all its load ($Q_0(\infty) = Q_1(\infty) = 0$). Quite a few data sets on the
232 kinetics of (“heavy”) metals in organisms show that once loaded, an organism never fully
233 loses its load [20-22]. Such behaviour cannot be captured by multi-compartment models
234 because this involves an extension to transformation of compounds (technically speaking,
235 sequestered compounds belong to a different chemical species).

236 If $k_{01}, k_{10} \gg k_e$, we can assume that $Q_1(t) \simeq P_{10} Q_0(t)$, and the kinetics Eq 4 reduces to
237 Eq 1, with Q replaced by Q_0 . This situation is called time-scale separation.

238 The interpretation of the compartments can be a special tissue or organ, or, more abstract,
239 reserve (compartment 1) and structure (compartment 0). The latter makes sense in the
240 context of DEB theory, where both compartments are assumed to have a constant, but
241 different, chemical composition, while reserve is relatively rich in lipids in many animal
242 taxa. While Eq 4 assumes that the size of all compartments does not change, we will relax

on this below, when we allow more interactions with metabolism and energetics.

We can obviously include more compartments, and more complex interactions with the environment, but the number of parameters rapidly increases this way. In practice multi-compartments are used if the one-compartment model fits data badly. The introduction of more parameters generally improves the fit, but not necessarily for the right reasons. As a rule of thumb it is only advisable to use more compartment models if data on the concentrations inside the compartments are available. If lack of fit of the one-compartment model is the only motivation, alternatives should be considered that are discussed below.

3.3 Film models

Film models are conceptually related to multi-compartment models because both are extensions of the one-compartment model that include more detail in transport (so in physical factors), though in different but complementary ways. Film models are especially popular in environmental chemistry for following the transport of compounds from one environmental compartment (such as water) to another (such as air). Both compartments are assumed to be well-mixed, except for a narrow film at the interface of both compartments, where transport is by diffusion.

259 To follow the dynamics for the densities of the compound n (mol/m), we need to define a
 260 spatial axis perpendicular to the interface and choose the origin at the boundary between
 261 the bulk and the film (on each side of the interface). Let L_i be the depth of the film, d_i
 262 the diffusivity of the compound in the film, and v_{ij} the exchange velocity of the compound
 263 between the two media. As discussed in Kooijman *et al.* [16], the dynamics of the densities
 264 is given by partial differential equations (pde's) for medium $i = 1 - j$ and $j = 0$ or 1

$$0 = \frac{\partial}{\partial t} n_i(L, t) - d_i \frac{\partial^2}{\partial L^2} n_i(L, t) \quad \text{for } L \in (0, L_i) \quad (5)$$

265 with boundary conditions at $L = 0$ (i.e. the boundary between the film and the well-mixed
 266 medium) for $v_i = d_i/L_i$

$$0 = \frac{\partial}{\partial t} n_i(0, t) - v_i \frac{\partial}{\partial L} n_i(0, t) \quad (6)$$

267 and boundary conditions at $L = L_i$ (i.e. the interface between the media where the two
 268 films meet)

$$0 = v_{ji} n_j(L_j, t) - v_{ij} n_i(L_i, t) + d_i \frac{\partial}{\partial L} n_i(L_i, t) \quad (7)$$

269 The boundary condition at $L = 0$, and the diffusion process in the film is rather standard,
 270 but we believe that the boundary condition at $L = L_i$ is presented for the first time in
 271 Kooijman *et al.* [16]. Users of the popular film models typically skip the formulation of the
 272 pde and directly focus at steady state situations; they typically use the concentration jump

273 across the interface that belongs to the situation when there is no net transport across the
 274 interface. As long as there is transport, however, the concentration jump differs from this
 275 equilibrium value.

276 The depth of the films is typically assumed to be small and the transport in the films in
 277 steady state, which makes that the density profiles in the films are linear. This leads to
 278 the 1-1-compartment kinetics for the bulk densities

$$\frac{d}{dt}n_i(0) = k_{ij}(P_{ij}n_j(0) - n_i(0)); \quad k_{ij} = \frac{v_i/\mathcal{L}_i}{1 + P_{ij}v_i/v_j - v_i/v_{ij}} \quad (8)$$

279 where $\mathcal{L}_i$ is the depth of the medium. This approximation only applies if $v_iv_j < v_{ij}v_j + v_{ji}v_i$
 280 and the transport in the film is rate limiting. The 1-1-compartment kinetics also results,
 281 however, if the film depths reduce to zero and if the diffusivities are high. The rate from
 282 i to j then reduces to $k_{ij} = v_i\mathcal{L}_i^{-1}(1 - v_i/v_{ij})^{-1}$. In these two situations, transport in the
 283 film is no longer rate limiting.

284 The applicability of film models to toxico-kinetics in organisms is still an open question.
 285 It can be argued that a stagnant water film sticks to aquatic or soil organisms (and air to
 286 a terrestrial organism), and that the skin (or cuticula) is not well served by the internal
 287 redistribution system (blood) of the organism. If toxico-kinetics is fully limited by transport
 288 in the film, and if it is not limited by the film, one-compartment kinetics results; only in the
 289 intermediary situation we can expect some deviations. Yet the discussion is not completely

academic, since these details matter for how the elimination rate depends on the partition coefficient [16, 23].

3.4 Uptake and elimination routes

We now consider extensions of the one-compartment model due to biological factors by accounting for various uptake and elimination routes. These routes depend on the type of organism, its habitat and properties of the compound. Animals that live in (wet) soil are in intense contact with the water film around soil particles, and their situation has similarities with that of aquatic animals. Direct transport through the skin can be important, which involves the surface area of an organism. Some parts of the skin are more permeable, especially that used by the respiratory system for dioxygen uptake and carbon dioxide excretion. The uptake rate might be linked to the respiratory rate, which depends on the energetics of the organism. Generally, the respiration rate scales with a weighted sum of surface area and volume, but the proportionality constants depend on the nutritional conditions of the organism [1]. For terrestrial animals, uptake via the lungs from air and via skin contact with the soil must be considered. Sometimes uptake is via drinking; the DEB theory quantifies drinking via the water balance for the individual and involves a.o. metabolism and transpiration. The details can be found in Kooijman [1].

A second important uptake route is via food and the gut epithelium. The feeding rate depends on food availability, food quality, and the surface area of the organism [1]. The elimination can follow the same routes as uptake, but there are several additional routes to consider, namely via products of organisms. The first possibility is the route that excreted nitrogen waste follows (urination). Reproductive products (mostly eggs and sperm) can also be an important elimination route. Moulting (e.g. ecdysozoans, including the rejection of gut epithelium, e.g. collembolans) or the production of mucus (e.g. lophotrochozoans) or milk (e.g. female mammals) are other possible excretion routes. The DEB theory quantifies reproductive and other products as functions of the amounts of reserve and structure of the individual. In the standard DEB model they work out to be cubic polynomials in body length, but the coefficients depend on the nutritional conditions (amount of reserves per structure) [1].

3.5 Changes in body size and composition

The body size of an organism matters in the context of toxico-kinetics for several reasons [24]. As exchange is via surface area, and is proportional to concentration, surface area-volume interactions are involved. This problem also applies to compartment and film models, but gets a new dimension if we consider changes in body size, which are linked to

the nutritional condition of organisms (lipid content), and so to (changes in) body composition. Small changes in size can have a substantial effect on the shape of accumulation curves.

If an organism does not change in shape during growth (so it remains isomorphic), surface area is proportional to volume^{2/3}, or to squared length. Moreover, dilution by growth should be taken into account, even at low growth rates. This modifies Eq 1 to

$$\frac{d}{dt}Q = (P_{0d}c(t) - Q) v_e/L - Qr \quad \text{with} \quad r = \frac{d}{dt} \ln L^3 \quad (9)$$

where L is the length, and $v_e = L_m k_e$ is the elimination velocity for maximum length L_m . The last term, Qr , represents the dilution by growth. If it equals zero, we can replace v_e/L by the constant elimination rate k'_e , but its meaning still matters if we compare the kinetics in two organisms of different size. DEB theory specifies how the change in (cubed) length depends on the amount of reserve and structure of the organism, and how the change in reserve depends on these state variables and food availability. Food intake and maintenance play an important role in growth and together they control the maximum size an organism can reach, since food intake is proportional to a surface area and maintenance to structural volume. Wallace had this insight in 1865 already [25].

The DEB theory allows for particular changes in body composition, because reserve and structure can change in relative amounts and both have a constant composition. Food

341 (substrate) is first transformed into reserve, and reserve is used for metabolic purposes,
342 such as somatic and maturity maintenance, growth, maturation and reproduction. The
343 change in reserve density for metabolic use is proportional to the reserve density per length,
344 which makes that high growth rates come with high reserve densities, i.e. the ratio of the
345 amounts of reserve and structure.

346 Reserves are in many animal taxa relatively enriched in lipids, which might have a strong
347 influence on the kinetics of hydrophobic compounds. The body burden of eel in a ditch that
348 is polluted with mercury or PCB might greatly exceed that of other fish partly because eel
349 is relative rich in lipids. This illustrates the importance of lipid dynamics. Freshly laid eggs
350 consist almost exclusively of reserve, which makes egg production a potentially important
351 elimination route for lipophyllic compounds. The reserve allocation to reproduction is via
352 a buffer that comes with species-specific buffer handling rules. Many aquatic species spawn
353 once a year only (e.g. most bivalves and fish), which implies that the buffer size gradually
354 increases between two spawning events and makes a sharp jump down at spawning. The
355 body burden can also make a jump at spawning (up or down, depending on the properties
356 of the compound).

357 The difference in lipid content between reserve and structure invites for the application of a
358 nested two-compartment model, where the exchange with the environment is via structure.

This links up nicely with food uptake, because reserve does not play a role in it, and food uptake is also proportional to squared structural length. An important difference with the nested two-compartment model is, however, that the size of the compartments typically changes in time, especially the reserve. When redistribution of the compound between the compartments reserve and structure is relatively fast, and the nested two-compartment model for the compound reduces to a one-compartment one, reserve dynamics still affects toxico-kinetics, because the lipid content is changing in time. The resulting dynamics for active uptake from food amounts to

$$\frac{d}{dt}Q = (P_{Vd}c_d + P_{VX}fc_X)\frac{v_e}{L} - Q(P_{VW}\frac{v_e}{L} + r) \quad \text{with} \quad P_{VW} = 1 + P_{EV}(m_E + m_{ER}) \quad (10)$$

where c_d and c_X are the concentrations of the compound in the environment and in food, f is the scaled functional response, P_{Vd} and P_{VX} are the partition coefficients of the compound in structure and environment or food. The reserve density m_E and the reproduction buffer density m_{ER} now modify the partition coefficient between structure and biomass (i.e. reserve plus structure), via the partition coefficient between reserve and structure P_{EV} . DEB theory specifies how structural length L , the reserve density m_E and the reproduction buffer density m_{ER} change in time.

The reproduction buffer is not of importance in all species, and not always in males. If food density is constant, the reserve density m_E becomes constant. In those situations

the structure-biomass partition coefficient P_{VW} is constant as well. If also the dilution by growth can be neglected, i.e. $r = 0$ and L is constant, Eq 10 still reduces to the one-compartment model Eq 1.

Many accumulation-elimination experiments are done under starvation conditions; e.g. it is hardly feasible to feed mussels adequately in the laboratory. The reserve density decreases during the experiment, so the chemical composition is changing, which can affect the toxico-kinetics [26].

Some situations require more advanced modelling of the uptake and eliminations route, where e.g. gut contents exchanges with the body in more complex ways, and defecation might be an elimination route.

3.6 Metabolism and transformation

Both uptake and elimination can depend on the metabolic activity [27]. Respiration is frequently used as a quantifier for metabolic activity. This explains the popularity of body size scaling relationships for respiration [28], and the many attempts to relate many other quantities to respiration. In the context of DEB theory, however, and that of indirect calorimetry, respiration is a rather ambiguous term, because it can stand for the use of dioxygen, or the production of carbon dioxide or heat. These are not all proportional

to each other, however. Moreover all these three fluxes have contributions from various processes, such as assimilation, maintenance, growth etc. Since the use of reserves fuels all non-assimilatory activities, this is an obvious quantifier to link to the rate at which compounds are transformed or taken up. For uptake of compounds via the respiratory system, the use of dioxygen might be a better quantity to link to uptake under aerobic conditions.

Respiration rates turn out to be cubic polynomials in structural length in DEB theory, which resemble the popular allometric functions numerically in great detail. The coefficients depend on the nutritional conditions in particular ways. Since elimination rates are inversely proportional to length because of surface area-volume interactions, as has been discussed, and the specific metabolic rate is very close to this relationship, it is by no means easy to evaluate the role of metabolism in direct uptake and elimination in undisturbed subjects.

The role of metabolism is easier to access for elimination via products and if the elimination rate is not proportional to the internal concentration, but has a maximum capacity. The classic example is the elimination of alcohol in human blood [29]. This type of kinetics can be described as

$$\frac{d}{dt}Q = k_e P_{0d}c - k_e Q / (K_Q + Q) \quad (11)$$

where K is a half saturation constant for the elimination process. It reduces to the one-compartment model Eq 1 for small internal concentrations, relative to the half saturation constant, $K_Q \gg Q$. The elimination rate can now be linked to metabolic activity, and so to body size. If particular organs are involved, such as the liver in the case of alcohol, the DEB theory can be used to study adaptation processes to particular metabolic functions. In the case of alcohol, the uptake term should obviously be replaced by a more appropriate one that applies to the particular subject. Many toxicants are metabolically modified. This especially applies to lipophilic compounds, which are typically transformed into more hydrophilic ones, which are more easily excreted but also metabolically more active. The rate of transformation can be linked to the metabolic rate, and so depends on body size and nutritional conditions. These metabolic products can be more toxic than the original lipophilic compound.

4 Bio-availability

Compounds are not only transformed in the organism, but also in the environment which affects their availability. Many have an ionic and a molecular form, which are taken up at different rates; the ionic species requiring counter ions, which complicates their uptake. Speciation depends on the concentration of compound and environmental properties, such

427 as the pH. It can vary in time and also occurs inside the organism, but the internal pH
428 usually varies within a narrow band only. Models for mixtures of chemicals can be used in
429 this case (see section on effects). Internal concentration gradients could develop if transport
430 inside the organism is slow; film models should be used in this case. A nice example of a case
431 where concentration gradients result from transformation in combination with transport is
432 the fluke *Fasciola* which has an aerobic metabolism near its surface with the micro-aerobic
433 environment inside its host, but an anaerobic metabolism in the core of its body [30].

434 Another problem, which occurs especially in soils, is that the transport through the medium
435 can be slow enough for concentration gradients to develop around the organism. Film
436 models should then be used again.

437 A major problem in the translation of laboratory toxicity tests to field situations is the
438 formation of ligands with (mainly) organic compound that are typically abundant in the
439 field, but not in the bioassay. Ligands reduce the availability substantially, and typically has
440 a rather complex dynamics. Moreover compounds can be transformed by (photo)chemical
441 transformation and by actions of (micro)organisms. This implies that the concentration
442 of available compounds changes in time, and our methodology to assess effects of chemical
443 compounds should be able to take this into account.

These processes of transformation require compound-specific modelling and this short section demonstrates that bio-availability issues interact with toxico-kinetics and effects of chemical compounds in dynamic ways, which calls for a dynamic approach to effects of chemicals [3, 31].

5 Effects at the individual level

Compound affect individuals via changes of parameter values as functions of the internal concentration [32, 33]. The parameter values are independent of the internal concentration in the “enough”-range of the compound. This implies the existence of an internal No Effect Concentrations (NECs) at either end of the “enough”-range; the upper end is typically of interest for ecotoxicological applications. Outside the “enough”- range the value of the target parameter is approximately a linear function of the internal concentration as long as the changes in parameter value are small; the inverse slope is called the tolerance concentration (a large value means that the compound is not very toxic). Small changes in parameter values do not necessarily translate into small changes of some end-point, such as the cumulative number of offspring [34] or the body size [35] at the end of some standardised exposure period. DEB theory specifies how exactly changes in parameter values translate into the performance of the individual. Typical target parameters are the specific

461 maintenance costs, or the yield of structure on reserve, or the maximum specific assimila-
462 tion rate or the yield of reserve on food or the yield of offspring on reserve. For effects on
463 survival, the hazard rate serves the role of target parameter, and the inverse “tolerance con-
464 centration” for the hazard rate is called the killing rate. Mutagenic compounds can induce
465 tumours [36], but also accelerate ageing by enhancing the effects of ROS. The partitioning
466 fraction for mobilised reserve can be the target parameter for endocrine disruptors.

467 Using sound theory for how effects depend on internal concentrations, DEB-based theory
468 can handle varying concentrations of toxicant [37-40], even pulse exposures [41]. DEB
469 theory applies to all organisms, including bacteria that decompose organic pollutants.

470 A proper description of this process should account for adaptation [42], co-metabolism
471 [43] and the fact most bacteria occur in flocculated form in nature [44], which affect the
472 availability of the compound.

473 The model of linear effects of internal concentrations on parameter values has been ex-
474 tended into several directions, such as adaptation to the compound, inclusion of the recent
475 exposure history via receptor dynamics [45] and attempts to include particular molecular
476 mechanisms [46].

5.1 Mixtures and NECs

Mixtures of compounds affect parameters values via addition of the effects of single compounds, plus an interaction term which is proportional to the products of the internal concentrations of the compound [47]. This interaction term can be positive or negative; a construct that is the core of the analysis of variance (ANOVA) model and rests on a simple Taylor series approximation of a general non-linear multivariate function, which only applies for small changes of parameter values; the non-linearities of the effects should be taken into account for larger changes. These non-linearities might well be specific for the compound and the species and, therefore, lack generality. Notice that linear effects on parameter values translate into non-linear effects of the performance of the individual because the DEB models are non-linear. Also notice that each DEB parameter has a NEC value for any compound; the lowest value among all parameters might be considered as *the* NEC of a compound for the organism, but its estimation requires to study effects on all 13 parameters of the standard DEB model, in principle. Since this study can be demanding, it is in practice essential to talk about the NEC of a compound for an organism *for a particular DEB parameter*.

The NEC reflects the ability of the individual to avoid changes of performance. From a statistical point of view, this robust parameter has very nice properties [48-50]. The NEC

is *not* meant to imply that some molecules of a compound don't have an effect, while other molecules do. The removal of a kidney in a healthy person can illustrate the NEC concept: the removal implies an effect at the sub-organismic level, but this effect generally does not translate into an effect at the individual level. The NEC, therefore, depends on the level of observation. We can delineate three cases of how compounds in the mixture combine for the NEC

- the presence of other compounds is of no relevance to the NEC of any particular compound
- the various compounds add, like they do for effects, and at the moment effects show up, the amounts of the compounds that show no effects remain constant
- the various compounds add, like they do for effects, and the amounts of the compounds that show no effects continue changing with the internal concentration of the compounds; if a compound continues accumulation more than other compounds, its NEC increases while that of the other compounds in the mixture decreases

The third case is possibly most realistic, but also computationally the most complex. In many practical situations the results are very similar to the second case, which can be used as an approximation. If compounds in a mixture are equally toxic, and so all have

512 the same NEC, the second case is formally identical to this special case [51]. This way
513 of described effects of mixtures turns out to fit well with experimental data [47] and each
514 pair of compounds have a single interaction parameter, which does not change in time. If
515 there are k compounds in a mixture, there are $k(k-1)/2$ interaction parameters, just like
516 in ANOVA.

517 A further reflection on the NEC might clarify the concept. Any compound affects (in
518 principle) all DEB parameters (including the hazard rate), but the NEC for the various
519 parameters differ. This makes that, if the internal concentration increases, the parameter
520 with the lowest NEC first starts to change, but other parameters follow later. In a mixture
521 of compounds, this can readily lead to a rather complex situation where in a narrow range
522 of (internal) concentrations of compounds in a mixture several parameters start changing
523 [51]. Even in absence of the above-mentioned chemical interactions of compounds on a
524 single parameter, interactions via the energy budget occur, which are hard to distinguish
525 from the chemical interactions on a single parameter. Chemical interactions are typically
526 rare, but interactions via the budget always occur.

5.2 Hormesis

Hormesis, the phenomenon that low concentrations of a toxicant seem to have a stimulating rather than an inhibiting effect on some endpoint, can result from interactions of the compound with a secondary stress, such as resulting from very high levels of food availability. If a compound decreases the yield of structure on reserve, it reduces growth and delays birth (if an embryo is exposed) and puberty (in the case of juveniles), but also reduces the size at birth. A reduction of growth indirectly reduces reproduction, because food uptake is linked to size. Since it also reduces size at birth, the overall effect can be a hormesis effect on reproduction (in terms of number of offspring per time) [52]. Indirect effects on reproduction differ from direct effects by not only reducing, but also delaying reproduction. This has important population dynamical consequences.

5.3 Co-variation of parameter values

A very powerful property of the standard DEB model and the one-compartment model is that they imply rules for how parameter values co-vary among species and compounds [53, 23, 9]; this variation directly translates into how expected effects vary. These expectations can be used to fill gaps in knowledge about parameter values, but cannot replace the need for this knowledge. Evolutionary adaptations and differences in mode of ac-

tion of compounds can cause deviation from expected parameter values for species, and
 compounds, respectively.

The reasoning behind the scaling relationships for the standard model rests on the as-
 sumption that parameters that relate to the local biochemical environment in an organism
 are independent of the maximum body size of a species, but parameters that relate to
 the physical design of an organism depends on the maximum size. Strange enough, this
 simple assumption fully species the covariation of parameter values. The application is
 best illustrated with the maximum length L_m an endotherm can reach in the standard
 DEB model. This length is a simple function of three parameters, $L_m = \kappa\{p_{Am}\}/[p_M]$,
 where κ is the fraction of mobilised reserve that is allocated to the soma, $\{p_{Am}\}$ is the
 surface-area specific maximum assimilation energy flux and $[p_M]$ is the volume-specific so-
 matic maintenance cost. Since κ and $[p_M]$ depend on the local biochemical environment,
 they are independent of maximum length, which implies that $\{p_{Am}\}$ must be proportional
 to maximum length. All other parameters can be converted in simple ways to quantities
 that depend on the local biochemical environment; these transformations then defined how
 they depend on maximum length. When we divide the maturity at birth and puberty by
 the cubed maximum length, we arrive at a maturity-density, which reflects the local bio-
 chemical environment and should not depend on maximum length. So the maturity at

birth and puberty are proportional to the cubed maximum length. Many quantities, such as the use of dioxygen by an individual, can be written as functions of parameter values and amounts of reserve and structure. So the maximum respiration rate of a species is a function of parameter values, while we know of each parameter how it depends on maximum length. It can be show that maximum respiration rate scales between a squared and a cubed maximum length, and the weight-specific respiration with weight to the power $-1/4$, a well-known result since Kleiber [54].

The reasoning behind the scaling relationships for the one-compartment model rests on the assumption that transport to and from the compartment is skewly symmetric [16]. The ratio of the concentrations in the compartment and the environment at equilibrium is a ratio of uptake and the elimination rates, just like the maximum length of the individual in the standard DEB model. This implies, see [16], that the uptake rate is proportional to the square root of the partition coefficient, and the elimination rate is inversely proportional to the square root of the partition coefficient. Film models are extensions of the one-compartment model, that behave at the interface between the environments basically in the same way as an one-compartment model; only around this interface they differ because film models account for concentration gradients. This deviation can be taken into account with the result that elimination rates are (almost) independent of the partition coefficient

for low values of the partition coefficient and inversely proportional to it at high values.

Effects parameters can be included into the scaling reasoning in similar ways, with the result that the NEC is inversely proportional to the partition coefficient, and the tolerance concentration or the killing rate is proportional to the partition coefficient.

The octanol-water partition coefficient is frequently taken as a substitute for the body-water partition coefficient, with the advantage that reliable computational methods exist to evaluate this partition coefficient from the chemical structure of the molecule. In practice, however, octanol is not an ideal chemical model for organisms which change the chemical composition of their bodies. This makes that the co-variation of NECs, elimination and killing rates show less scatter and can be expected on the basis of the variation between each of these three parameters and the octanol-water partition coefficient [23].

We are unaware of any descriptive model for toxico-kinetics and/or metabolic organisation for which theory on the co-variation of parameter values is available, and we doubt that it is even possible to derive such theory for descriptive models. Co-variation theory is not available for the so-called net-production models [55], for instance, where maintenance needs are first subtracted from assimilation before allocation to storage, growth or reproduction.

The fact that the predicted relationships of now over thirty eco-physiological quantities, such as length of the embryonic and juvenile periods, maximum reproduction rate, maxi-

598 mum growth rate, maximum population growth rate, vary with the maximum body size of
599 species in ways that match empirical patterns provides strong support of the DEB theory.

600 The one-compartment and standard DEB models share the property that the independent
601 variable (the partition coefficient in the case of toxicokinetics and the maximum length in
602 the case of budgets) can be written as a ratio of an incoming flux (of toxicant and food,
603 respectively) and an outgoing flux (excretion and maintenance, respectively). This shared
604 property seems to be crucial for the core theory.

605 One of the many possible applications of the scaling relationships is in the effects of mix-
606 tures of compounds with similar modes of action, such as the poly-chlorinated hydro-
607 carbons. Suppose we know the concentrations and the partition coefficients of the com-
608 pounds in the mixture. We then link the elimination rates, the NECs and the tolerance
609 concentrations to the partition coefficients in the way described, and estimate the three
610 proportionality constants for the results of a bioassays with the mixture.

611 The sound theoretical basis for effects of toxicants in combination with rules for the co-
612 variation of parameter values offers the possibility for extrapolation, from one individual
613 to another, from one species of organism to another, and, sometimes, from one type of
614 compound to another [23]. These crosslinks partly reduce the need for a huge experimental
615 effort that should be invested in more advanced forms of environmental risk assessment,

616 such as discussed in Brack *et al.* [13]. Moreover, the theory simplifies to parameter poor
617 models under particular conditions. It has been demonstrated that many popular empirical
618 models turn out to be special cases of the general theory [1]. This might help in particular
619 applications.

620 **6 Effects at the population and ecosystem level**

621 At high food levels, organisms grow and reproduce fast and the maintenance costs com-
622 prise only a tiny fraction of the budget of the individuals. If a compound increases the
623 maintenance cost for individuals, say by a factor two, these effects are hardly felt by a
624 fast-growing population. Fast growth never lasts long in nature, due to the depletion of
625 food resources. At carrying capacity, where the generation of food resources just matches
626 the maintenance needs of a population (this is the maintenance needs of the collection of
627 individuals plus a low reproductive output that cancels the mortality), maintenance costs
628 comprise the dominant factor of the budget of individuals. If a toxic compounds now in-
629 creases the maintenance cost by a factor two, it in fact reduces the carrying capacity by
630 a factor two. This simple argument shows that the effects of toxicants on populations is
631 dynamic, even if the concentration of the compound would be constant [56]. It also shows
632 that no single quantifier for toxicity can exist at the population level.

If a toxic compound increases the cost of growth or reproduction, the effects hardly depend on the growth rate of the population, so on the food level, which shows that the mode of action is important for how effects on individuals translate to those on populations. It might be difficult to tell the various modes of action apart on the basis of the results from a standardised toxicity bioassay. The reason why the mode of action is still important is in the biological significance of the observed effect, which must be found at the population level. Details in the reproduction strategy of populations turn out to be important for how effects on reproduction translate to the population level [57].

Although bioassays with meso-cosms have the charm of being close to the actual interest of effects of toxicants to be avoided, the experimental control is extremely weak which results in a huge scatter of trajectories of experimental meso-cosms. The result is that the effects have to be huge to recognise them as effects [58]. Moreover the expected long term behaviour of chemically perturbed ecosystems is very complex, as shown by bifurcation analysis [59].

The specific population growth rate integrates the various performances of individuals naturally, and can rather easily be evaluated [60]. A delay of the onset of reproduction can be at least as important as a reduction of the reproduction for the fate of the population.

7 Concluding remarks

We argued that models for effects of chemical compounds should have three modules:

- dynamic energy budgets for how organisms generally deal with resource uptake and allocation
- toxico-kinetics for how organisms acquire the compounds
- changes of budget parameters as function of the internal concentrations

We discussed the basics for each of these modules: the standard DEB model, the one-compartment model and the linear change in target parameters. We also indicated where and how these models can be extended, from simple to more complex, to include particular phenomena. We discussed how budgets affect both the kinetics and the effects and, therefore, have a central role in effects models. Practice teaches that the restriction of realistic modelling is not in the model formulation as such, but in the useful application of these models to data. More complex models have more parameters and many of these parameters are by no means easy to extract from available experimental data. They require knowledge of physiological and ecological processes that are typically outside the scope of typical (eco)toxicological research. Kooijman *et al.* [61] discusses why any particular application of DEB theory requires only a limited set of parameter values, and how these

values can be obtained from simple observation on growth and reproduction at several levels of food availability.

The practical need to fill in gaps in knowledge about parameter values is the reason why due attention has been given to theory for the co-variation of parameter values; this theory naturally follows from the logical structure of the one-compartment model and the standard DEB model. Extensions of both models can modify the co-variation of parameters, as has been discussed.

Contrary to descriptive models, models with strong links with underlying processes can be used for a variety of extrapolation purposes, from acute to chronic exposure, from one species to another, from one compound to another, from individuals to populations, from laboratory to field situations [31]. Such extrapolations are typically required in environmental risk assesment, where NECs should play a key role [62]. The use of models to predict exposure in the environment is frequent, but to predict effects is still rare. The complexity of the response of organisms to changes in their chemical environment doubtlessly contributed to this. Yet we think that thirty years of applications of DEB theory to quantify effects of compounds on organisms have demonstrated that the theory is both effective and realistic. Many of the computations behind the models in this chapter can be done with the freely downloadable software package DEBtool: <http://www.bio.vu.nl/thb/deb/deblab/>

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819 List of Figures

820	1	The standard DEB model with fluxes (moles per time) and pools (moles).	
821		Assimilation is zero during the embryo stage and becomes positive at the	
822		transition to the juvenile stage (birth) if food is available. Age is zero at the	
823		start of the embryo stage. Reproduction is zero during the juvenile stage	
824		and becomes positive at the transition to the adult stage (puberty), when	
825		further investment into maturation is ceased.	50
826	2	The scheme of the one- and two-compartment models. The factor P_{0d} con-	
827		verts an external concentration into an internal one; all rates labelled k have	
828		dimension ‘per time’. In the two-compartment model P_{0d} does not have the	
829		interpretation of the bioconcentration factor.	51

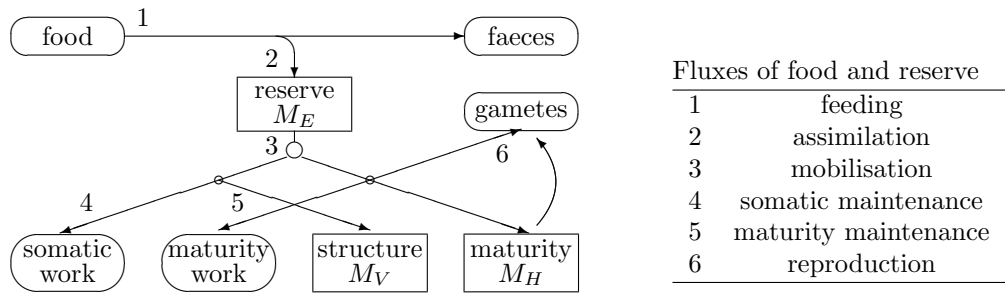


Figure 1:

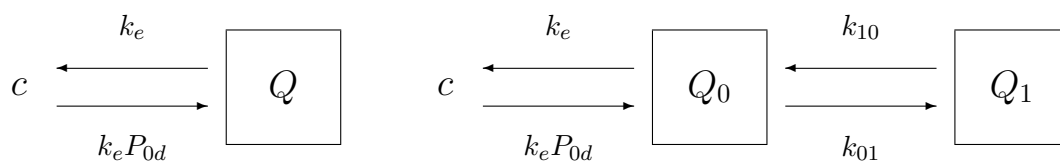


Figure 2:

830 List of Tables

831	1	List of frequently used symbols, with units and interpretation. The unit	
832		C-mol represents the number of C-atoms in a organism as multiple of the	
833		number of Avogadro.	53

Table 1:		
symbol	units	interpretation
t	d	time
c	M	concentration of compound in the environment
n_i	mol m^{-1}	density of compound
Q	mol C-mol^{-1}	concentration of compound in an organism
r	d^{-1}	specific growth rate of structure
k_e	d^{-1}	elimination rate
k_{01}, k_{10}	d^{-1}	exchange rates between compartments
P_{0d}	$\text{mol C-mol}^{-1} \text{M}^{-1}$	BioConcentration Factor (BCF)
v_{ij}, v_e	m d^{-1}	velocity, elimination -
d_i	$\text{m}^2 \text{d}^{-1}$	diffusivity
m_E, m_{ER}	C-mol C-mol^{-1}	reserve density, reproduction buffer density