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# Maximum relative speeds of living organisms: why do bacteria perform as fast as ostriches?

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## Abstract.

Self-locomotion is central to animal behaviour and survival. It is generally analysed by focusing on preferred speeds and gaits under particular biological and physical constraints. In the present paper we focus instead on the maximum speed and we study its order-of-magnitude scaling with body size, from bacteria to the largest terrestrial and aquatic organisms. Using data for about 460 species of various taxonomic groups, we find a maximum relative speed of the order of magnitude of ten body lengths per second over a  $10^{20}$ -fold mass range of running and swimming animals. This result implies a locomotor time scale of the order of one tenth of second, virtually independent on body size, anatomy and locomotion style, whose ubiquity requires an explanation building on basic properties of motile organisms. From first-principle estimates, we relate this generic time scale to other basic biological properties, using in particular the recent generalisation of the muscle specific tension to molecular motors. Finally, we go a step further by relating this time scale to still more basic quantities, as environmental conditions at Earth in addition to fundamental physical and chemical constants.

*Keywords:* biological scaling, locomotion, molecular motors, metabolic energy, Earth-based life

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## 1. Introduction

Autonomous locomotion is a fundamental property of animals, which involves an amazing diversity of mechanisms and has important consequences for ecosystems and evolution [1]. Biological scaling generally focuses on the preferred speed  $V$ , which minimizes the power or maximises the range and is primarily determined by dynamical constraints [2–5]. Because of the major role played by the gravitational acceleration  $g$ , most scalings with size  $L$  can be related to the ratio of kinetic to gravitational energy - the so-called Froude number  $Fr = V^2/Lg$ . For animals sharing similar locomotor mechanisms, this yields  $V \propto (Lg)^{1/2} \propto M^{1/6}g^{1/2}$  if the body mass  $M \propto L^3$  [6, 7].

The present study focuses instead on the maximum speed reached by animals when they are driven by circumstances to move faster than their natural speed, as for catching preys or escaping. Although such a speed has different consequences on behaviour from the preferred speed, it may be central to survival. Various scalings have been found over relatively narrow mass ranges or restricted animal groups and gaits, with allometries often compatible with a constant ratio of kinetic to gravitational energy, albeit with a value of  $Fr$  variable and higher than for the preferred speed [8–12]. However, on a very large scale and in order of magnitude, the maximum speed has been shown by Bonner [2, 13] to be roughly proportional to body length  $L$  over nearly the  $10^{20}$ -fold mass range of running and swimming organisms - a fundamental and unexplained property which generalises a result observed for fishes [14–16]. If most aquatic and terrestrial species have a similar maximum relative speed  $V_{\max}/L$ , this implies a locomotor time scale  $(L/V_{\max})$ , whose near independence on size contrasts with the general increase of physiological time scales with body mass - smaller animals having a faster pace of life [17]. Furthermore such a ubiquitous time scale - holding from bacteria to large fishes and from small mites to ostriches, requires an order-of-magnitude explanation ignoring details and based on first principles. To the best of our knowledge, this has not been previously analysed nor interpreted in the biological literature.

Interpretations should build on basic properties common to motile organisms. Chemistry yields a huge range of time scales because of the variations in reaction rates with concentration, temperature, and catalyst. The next structural level is that of proteins, whose folding time scales span a large range, from about  $10^{-4}$  to  $10^2$  s for typical numbers of amino-acids  $N \sim 10^2 - 10^3$  [18]. A larger scale is the cell size, close to  $10 \mu\text{m}$  for most organisms [19], so that the diffusion time scale of proteins in cells is  $10^{-3} - 10$  s [20]; however in the cellular complex structure, diffusion is expected to be strongly modified and overwhelmed by non-thermal processes such as active transport mediated by molecular motors [19]. A different viewpoint is to consider macroscopic biological quantities which may limit the relative speed - the mass density, the specific tension of biological motors [22], and the maximum specific metabolic rate [23], as proposed by Meyer-Vernet and Rospars [24] using dimensional analysis in a preliminary paper written in the context of physics teaching.

The present paper is organised as follows. Section 2 studies the empirical data

(supplementing, analysing and referencing the preliminary set plotted in Ref [24]) to deduce the maximum speed as a function of body’s mass and the resulting time scale. Section 3 deals with an interpretation. Since the result is approximate and nearly independent on size and locomotion style, holding both from bacteria to large fishes and for all running taxa, we make first-principle order-of-magnitude estimates. We build in particular on the universality of the specific tension [22] used by organisms for their locomotion (section 3.1), based on a recent generalisation of the muscle tension to molecular motors [21, 22]. Section 4 examines the compatibility of the results with other empirical scalings, and tries to relate them to more basic properties. Indeed, from the physicist’s perspective, proposing to explain the order of magnitude of a time scale from known biological quantities as in a Fermi’s problem is not sufficient, unless we are able to calculate these biological quantities themselves from basic numbers. Since the organisms studied evolved on Earth, these basic numbers are expected to involve environmental conditions at Earth, in addition to fundamental physical or chemical constants. In particular, as Feynman once said [25], “all the things that we see that are moving, are moving because the sun is shining”; so, can we relate this scale to the amount of solar energy reaching Earth?

## 2. Methods and empirical results

Values of the maximum speed as a function of mass were taken from tables of values published in the literature listed in subsection 2.3 for a variety of running or swimming animals of different taxonomic groups, ranging in mass from an eubacteria ( $M \simeq 2 \times 10^{-16}$  kg) to the blue whale ( $M \simeq 1.5 \times 10^5$  kg).

We only consider the velocity of organisms as a whole (i.e. the speed of the centre of mass). Hence, we include neither plants and fungi nor sessile animals. We do not consider either the motions resulting from sudden release of stored elastic energy as fleas’ jumps or spasmoneme motion [26], nor motion on inclined supports, or adhesive locomotion used for example by gastropods [27] and pili [28], nor cells’ crawling [29], which is based on a completely different mechanism from that considered here [30]. For microorganisms, we select the data for which the recorded speed cannot be confused with diffusion resulting from random directional changes, or with a drift due to gravity or chemical gradients [31].

In order to study the relative speed as a function of mass, we need empirical values of the three quantities  $V_{\max}$ ,  $M$  and  $L$ . They are determined as follows.

### 2.1. Maximum speed

How is the empirical “maximum speed” defined? Does it mean a maximum speed sustainable for long periods, or a top speed briefly attainable in a sprint? The top speed requires the additional contribution of transient (anaerobic) processes leading to fatigue, whereas animals in prolonged (aerobic) locomotion must achieve oxygen balance, so that

the speed at which the maximal rate of oxygen consumption is attained is usually called the maximum aerobic speed.

Most data on mammals concern the transient maximum (top) speed, which has been found to be in average 2-3 times higher than the maximum sustainable speed [8, 32]. For ectothermic species, which typically move in short bursts of exercise followed by periods of recovery, most recorded data concern the “burst” speed, expected to be close to the top speed. For fishes, the average top speed (called “escape” speed) is known to be 3–4 times greater than the average sustained speed [15, 35]. This order-of-magnitude similarity of the transient top speed and the maximum sustained speed is in line with the observed similarity in order of magnitude of the maximum anaerobic (transient) and aerobic (sustainable) metabolisms [36] (see section 3.2). Since we seek order-of-magnitude values over a wide range of species and masses, we therefore consider the top speed, which represents the majority of the available maximum speed data.

Note that since both original and secondary sources were consulted, with experimental conditions not always available, the results may not be homogeneous in this respect. Indeed, the motivation to run and swim as fast as possible depends on the species and the experimental procedure; convincing a mite or an ostrich to run fast does not require the same tools. Therefore, since the diversity of animal sizes and gaits necessarily produces large differences in experimental set-ups, the data have variable degrees of accuracy - a problem inherent to considering organisms over a 20-fold mass range.

Another question concerns the temperature, which affects reaction rates and therefore speed. As far as possible, we selected the data corresponding to the mean body temperature of animals active in the field. For ectothermic organisms, the range of such temperatures is expected to produce a significant range in speeds; we shall discuss this point in section 4.3. We did not make corrections for the sake of consistency, because no such corrections were applied neither for deducing the constancy of the specific tension nor the constancy of the maximum specific metabolism (see e.g. discussion in Ref. [23]), which are used in our interpretation.

Similarly, we did not correct either for the phylogenetic context [33, 34]. Although phylogenetic considerations might affect the regressions (see e.g. [37, 38]), they should not invalidate our main result that the motional time scale has the same order of magnitude over the entire mass range.

## 2.2. Mass and length

When only one of the two parameters  $M$  and  $L$  was available we determined the other one from the documented relation between them. Since organisms of different sizes must compensate for varying dynamical and physiological conditions, they cannot be precisely geometrically similar. As size increases - implying larger dynamic forces, whereas the elasticity of materials remains constant, the length-to-diameter ratio  $L/D$  may have to decrease in order to prevent excessive deformations, yielding the relation  $D^2 \propto L^3$ ,

whence  $M \propto L^4$  [2, 39]; this so-called elastic similarity holds for some properties in limited size ranges [2]. Another relevant constraint is the resistance of tendons and skeletons, which leads to stress similarity within some taxa [40]. However, recent observations show that for large-scale variations expressed in order of magnitude - which is what is needed here - geometric similarity is generally more adequate, i.e.  $M \propto \rho L^3$ ,  $\rho$  being the mass density, with a constant of proportionality depending on shape as summarized below.

First, for all terrestrial non-flying mammals, mass and length can be approximated by  $M \simeq (3L)^3$  (in SI units) over most of their size range [41] (although elastic similarity tends to hold for very large mammals [42]). By comparing with empirical allometries holding for various taxa [12, 43, 44], we have verified that extending this relation to non-mammals produces errors that do not affect significantly the results. An exception is the case of very elongated species as snakes and worms; hence we did not include limbless terrestrial taxa; we shall return to this point in section 4.2.

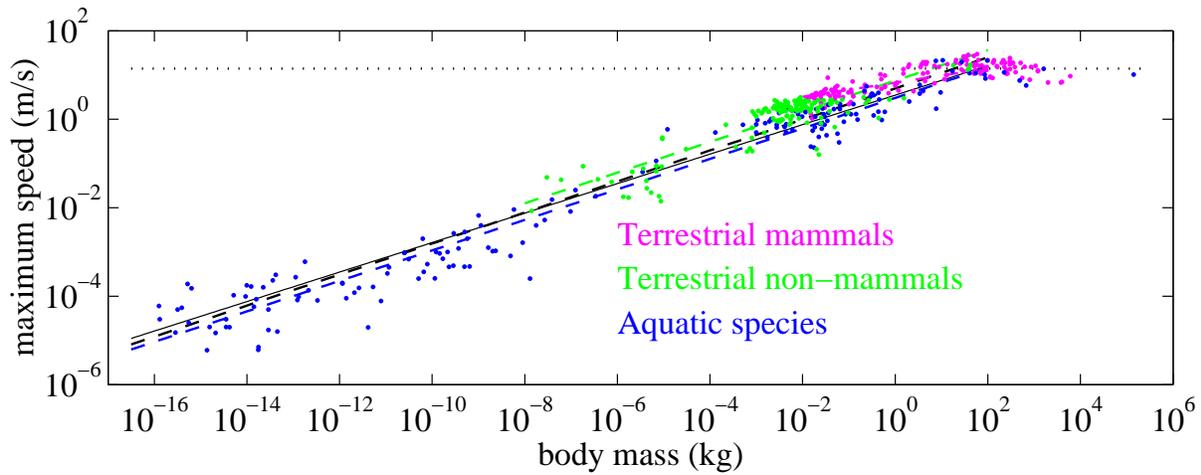
Second, swimming species (except microorganisms) tend to have a stream-lined shape with  $M \simeq (2L)^3$  (in SI units) [45].

Third, motile microorganisms have a much greater diversity of shapes (with a median at  $L/D \simeq 3$ ) [16, 46, 47], the elongation being expected to facilitate their orientation by increasing the resistance to rotation induced by Brownian motion [16]; because of this diversity, we keep only microorganism data for which both  $L$  and  $M$  or  $L$  and  $D$  are available (contrary to the preliminary data plotted in Ref. [24]). When only  $L$  and  $D$  are available, we estimate  $M$  using the prolate spheroid approximation  $M = (\pi/6)\rho L^3(D/L)^2$ . For our microorganism data set, the mean of  $L/D \simeq 4.8$  ( $M \simeq (1.6L)^3$ ) with a standard deviation  $\simeq 6.4$  and a median of 2.4 ( $M \simeq (3.3L)^3$ ).

### 2.3. Maximum speed versus body mass

Figure 1 shows the measured maximum (top) speed as a function of mass for about 460 species of different taxonomic groups. The data for 155 terrestrial mammal species (taken from [9] plus some species from [8, 48] not included in [9]) are plotted in magenta. The data for 137 other running species (a set roughly three times larger than considered in Ref. [24]) are plotted in green. They include 14 ant species [2, 11, 49], 16 other arthropod species (including crabs [50], mites [2, 51], spiders [52, 53], cockroaches [54] and beetles [55]), 100 species and subspecies of lizards [12], and 7 species of terrestrial birds [56–60]. This yields a terrestrial mass range spanning nearly 12 orders of magnitude, from  $10^{-8}$  kg (clover mite) to 6000 kg (African bush elephant). The maximum speeds for 95 swimming mites, fishes, cetaceans, crustaceans, penguins, turtles, and squids [2, 5, 14, 15, 61–67] and 71 microorganisms [2, 46, 68–70] are also plotted (in blue). This yields a  $7 \times 10^{20}$ -fold range in mass, from  $2 \times 10^{-16}$  kg (eubacteria) to  $1.5 \times 10^5$  kg (blue whale).

We have superimposed (dashed) regression lines of  $\log_{10}V_{\max}$  versus  $\log_{10}M$  in magenta for running mammals (slope 0.16 with 95% confidence limits (CL) [0.14, 0.18],



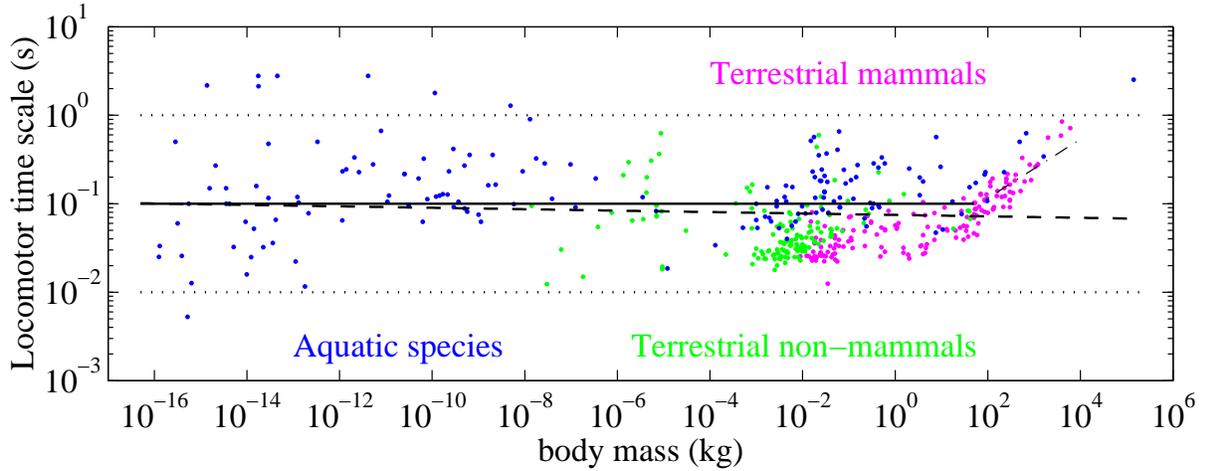
**Figure 1.** Maximum speed versus body mass for terrestrial mammals (in magenta) and non-mammals (in green), and aquatic species (in blue) from the references cited. The corresponding regression lines are shown (dashed) in similar colors, and the regression line for all data is shown in (dashed) black. Maximum running and swimming speed (black solid line, equation (7)) and upper limit at large masses (dotted) are also shown as estimated in section 3.

intercept 0.83, correlation coefficient  $r=0.80$ ,  $n = 155$ ), green for running non-mammals (slope 0.35 with 95% CL [0.31, 0.38], intercept 0.87,  $r=0.88$ ,  $n = 137$ ), and blue for swimming (slope 0.34 with 95% CL [0.33, 0.36], intercept 0.48,  $r=0.98$ ,  $n = 166$ ). The regression line for all running and swimming data is plotted in (dashed) black (slope 0.35 with 95% CL [0.34, 0.36], intercept 0.70,  $r=0.97$ ,  $n = 458$ ). In all data subsets the slope of the regression line is highly significantly different from 0 ( $P < 10^{-3}$ ). The proportion of the total variance explained by the regression is 64% for running mammals, 77% for running non-mammals, 96% for swimmers and 94% for swimmers and runners together.

Figure 1 also shows that the maximum speed tends to level-off for large organisms, which confirms a known result [2, 8, 9] and strongly affects the fitted exponent of terrestrial mammals which includes a high number of large animals; we shall return to this point later.

#### 2.4. Maximum relative speed and corresponding time scale

In figure 2 the time scale  $\tau = L/V_{\max}$  is plotted as a function of body mass for the organisms considered in subsection 2.3. The slope of the regression line of  $\log_{10}\tau$  versus  $\log_{10}M$  for all data, plotted as a dashed line (slope -0.008 with 95% CL  $[-0.017, 5 \times 10^{-4}]$ , intercept -1.127,  $r = 0.087$ ,  $n = 458$ ) is not significantly different from 0 ( $P = 0.06$ ). Terrestrial species have a mean  $\simeq 0.082$  s (SD  $\simeq 0.10$  s, median  $\simeq 0.049$  s,  $n = 292$ ), close to that of mammals only (0.091 s). Swimming species have a mean  $\simeq 0.27$  s (SD  $\simeq 0.49$  s, median  $\simeq 0.12$  s,  $n = 166$ ). Note that microorganisms ( $M \lesssim 10^{-8}$  kg, mean  $\simeq 0.37$  s, SD  $\simeq 0.66$ ) and other swimmers ( $M \gtrsim 10^{-8}$  kg, mean  $\simeq 0.20$  s, SD  $\simeq 0.28$ ) have similar medians ( $\simeq 0.12$  s) and similar logarithmic means ( $\simeq 0.14$  s) despite the



**Figure 2.** Locomotor time scale as a function of body mass for terrestrial mammals (in magenta) and non-mammals (in green), and aquatic species (in blue) from the references cited in subsection 2.3. The dashed line  $\tau = 10^{-1.127} \times M^{-0.008}$  is the regression of  $\log_{10}\tau$  versus  $\log_{10}M$  for the whole data set. The solid line (with values smaller and larger by one order of magnitude dotted) is from (6), estimated in section 3; the limit for large masses estimated in subsection 3.4 is shown dash-dotted. For clarity the scale is twice larger on the speed axis than on the mass axis.

huge difference in mass. The larger variability of microorganism data may result in part from the increased difficulty in measurement of both size and speed as size decreases. The mean of all data together is  $\simeq 0.15$  s (SD  $\simeq 0.3$  s, median  $\simeq 0.07$  s). More than 98% of the data lie in the range  $10^{-2} < \tau < 1$  s, which is remarkably narrow compared to the  $10^{20}$ -fold variation in body mass. The next section proposes an order-of-magnitude interpretation of these empirical results.

### 3. Physical interpretations

The locomotor time scale of living organisms is expected to depend on at least three basic macroscopic properties, whatever their size and locomotion style: their mass density  $\rho$ , which is close to that of water

$$\rho \simeq 10^3 \text{ kg m}^{-3} \quad (1)$$

the maximum force per cross-section area  $\sigma$  (specific tension) that they can exert, and the maximum power per mass unit  $b_M$  (maximum specific metabolic rate) that they can use for moving. Let us consider in turn the two latter factors.

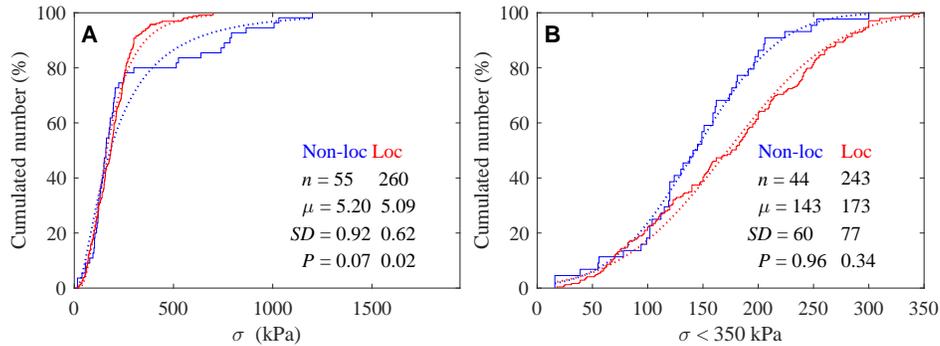
#### 3.1. Specific tension

The force per cross-sectional area of skeletal muscles at constant length (the so-called isometric specific tension) is known to be remarkably constant whatever the muscle and body mass, because of the similar construction of skeletal muscles based on the myosin molecule [71, 72]. This notion of specific tension has been formally generalised

to individual molecules [21, 22] and to propulsion organelles and individual molecular motors propelling microorganisms [22], showing that their maximum specific tension has a similar statistical distribution as that of conventional muscles, and that the maximum applied force per unit cross-section area is about

$$\sigma \sim 2 \times 10^5 \text{ N m}^{-2} \quad (2)$$

over about 19 orders of magnitude in body mass, from microorganisms to the largest animals [22].

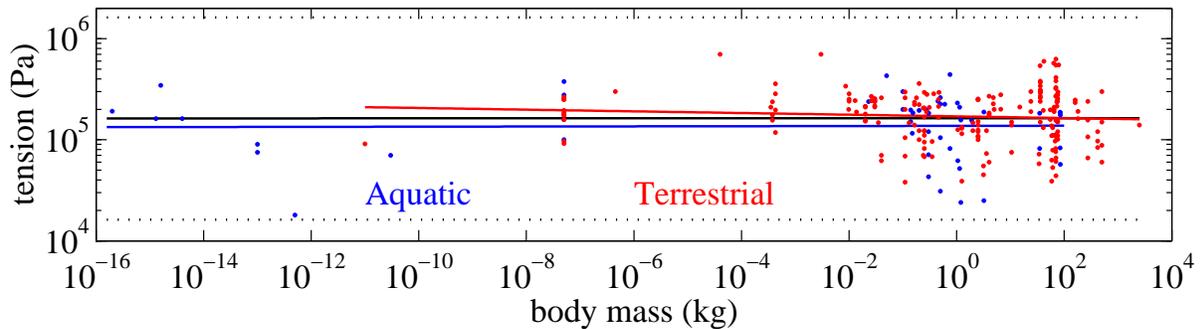


**Figure 3.** Empirical cumulative distribution functions (CDFs) of the specific tension  $\sigma$  used for locomotion (red) and other functions (blue) in aquatic and terrestrial species. Panel A shows empirical CDFs of all  $\sigma$  values (solid) fitted to lognormal CDFs of mean  $\mu$  and SD indicated (dotted). Panel B shows empirical CDFs of values  $\sigma < 3.5 \times 10^5$  Pa fitted to Gaussians (dotted) of mean and SD indicated. CDFs of locomotors and non-locomotors are not significantly different at level 5% ( $P = 0.13$ ) in A and significantly different ( $P = 0.01$ ) in B. All comparisons based on Kolmogorov-Smirnov tests. (References of the data in Rospars and Meyer-Vernet [22]).

We are interested here in the specific tension used for locomotion. Figure 3 compares the cumulative distributions of the values of  $\sigma$  used (in red) or not (in blue) for locomotion. The values smaller than  $3.5 \times 10^5$  Pa (corresponding to the inflection point of the distributions in figure 3A), which can be fitted to Gaussians, are plotted in the right-hand-side panel.

Figure 4 shows the specific tension used for locomotion as a function of cell or body mass for 260 terrestrial and aquatic species. The slope of the regression lines ( $-0.008$  with 95% CL  $[-0.022, 0.005]$ ,  $n = 207$ ,  $P = 0.22$  for terrestrial species,  $0.0008$  with 95% CL  $[-0.017, 0.018]$ ,  $n = 53$ ,  $P = 0.93$ , for aquatic species) and  $6 \times 10^{-5}$  with 95% CL  $[-0.0096, 0.0097]$ ,  $n = 260$ ,  $P = 0.99$  for all data together) are not significantly different from 0. The mean of all values of  $\sigma$  is  $1.84 \times 10^5$  (SD  $1.1 \times 10^5$ , median  $1.74 \times 10^5$ )  $\text{N m}^{-2}$ .

The order of magnitude of this ubiquitous value of  $\sigma$  can be interpreted from first principles. Molecular motors as the myosin molecule [73] move by steps via a conformational change of their 3-D structure, so that the elementary step length is of the order of magnitude of their size, which is the typical protein size  $a_0 \sim 6$  nm



**Figure 4.** Specific tension used for locomotion versus cell or body mass for terrestrial (in red) and aquatic (in blue) species, and corresponding log-log regression lines, with the regression for all data plotted in black ( $\sigma = 10^{5.21} \times M^{0.00}$ ). The dotted lines are the values smaller and larger by one order of magnitude. For clarity the scale is about twice larger on the tension axis than on the mass axis. (References of the data in Rospars and Meyer-Vernet [22]).

[74]. Each such step is mainly powered by the ATP release energy, of standard value  $W_0 \simeq 0.5 \times 10^{-19}$  J per molecule, so that the elementary force exerted by the motor is  $F_0 \sim W_0/a_0$  in order of magnitude [75], whence the force per unit motor’s cross-section [22]

$$\sigma \sim W_0/a_0^3 \quad (3)$$

Substituting the above values of  $W_0$  and  $a_0$  into (3) yields (2).

### 3.2. Mass-specific metabolic rate

Most analyses of the energy consumption rate  $B$  of living organisms concern the so-called “basal” (“resting”) or “field” metabolic rates, depending on whether the organism is at rest or is performing its normal tasks. To compare organisms of different body mass  $M$ , it is convenient to define the mass-specific metabolic rate  $b = B/M$ . Power laws of the form  $b = \alpha M^\beta$  have been fitted to various sets of data, yielding various scaling exponents, and fostering an amazing variety of models [76]. Since exchanges with the environment act through surfaces ( $\propto M^{2/3}$ , yielding  $b \propto M^{-1/3}$  - although such surfaces may scale differently [77, 78]), whereas internal resources vary with volume ( $\propto M$ , yielding  $b \propto M^0$ ), one may expect  $-1/3 < \beta < 0$ , depending on the relative importance of external and internal constraints [79], if other properties as the transport system co-adjust [80]. This range covers most of the observations, including the so-called Kleiber’s law exponent  $\beta = -1/4$  which remains controversial [17, 38] and in particular has been found not to hold for prokaryotes and protists [82]. Recently,  $b$  has been found to be constant in order of magnitude across the  $10^{20}$ -fold mass range of living organisms in comparable physiological states at natural temperatures [81] - a result compatible with the variety of exponents  $\beta$  because of the variation of the intercepts  $\alpha$  for different taxa and mass ranges.

An important point is that the exponent  $\beta$  varies with the physiological state of the organisms, approaching zero at high level of activity [79, 84], which is the relevant state at maximum speed. Indeed, at high activity, the metabolism is expected to be mainly determined by the resource demand of working tissues, which scales as their mass ( $\propto M$ ), as does the oxygen and energy stored which are used during bursts of maximal activity. Hence the maximal specific metabolic rate is expected to be roughly constant in order of magnitude - at least when normalized to the mass of active tissues, as observed [23, 83–85].

This maximal specific metabolic rate  $b_M$ , ubiquitous in order of magnitude across the  $10^{20}$ -fold mass range of living organisms, may represent a biochemical limit universal for life; measurements on dividing bacteria and moving animals during peak activities yield the maximal value

$$b_M \sim 2 \times 10^3 \text{ W kg}^{-1} \quad (4)$$

per unit mass of active tissue [23], like the mitochondria [84, 86–88] at their natural temperature. For example in mammals, over 90% of the energy at maximum metabolic rate is consumed in locomotor muscles, which are densely populated of mitochondria - the energy factories of eukaryotic cells - and we have  $b_M \simeq 1600 \text{ W/kg}$  of mitochondria mass [83, 84]. The value per unit body mass is smaller by the proportion of active tissue in the body (typically 1-50%), and is typically 5-30 times higher than the resting specific metabolic rate  $b$  for both vertebrates [1] and invertebrates [89]. In particular, the maximum metabolic rate per unit body mass amounts to only about 90 and 100 W/kg body mass for respectively (20 g) European woodmouses (*Apodemus sylvaticus*) and (28 kg) pronghorns (*Antilocapra americana*) [83], whose bodies have a large proportion of mitochondria, whereas it amounts to only about 20 W/kg body mass for human [36] and insect [89] runners.

Most of these measurements concern the aerobic metabolism, measured via oxygen consumption. At (short term) peak activity, part of the energy - or even most of it for ectoterms, is supplied by transient anaerobic processes [90]. However, the total maximal energy output is found to be generally similar in order of magnitude [36], the smaller aerobic power of ectoterms being compensated by a larger anaerobic power (e.g. [91–93]).

### 3.3. Consequences on the locomotor time scale

A basic order-of-magnitude estimate of the locomotor time scale can be derived from the constraints addressed above. Consider first a micro-organism whose locomotion is powered by  $N$  molecular motors of size  $a_0$ , making it move by about its length  $L$  at the frequency  $f$  so that its speed is  $V \simeq Lf$ . Assuming that each motor (of mass  $\rho a_0^3$ ) uses the elementary energy  $W_0$  for each such step, the total power is  $NW_0f$ , which cannot exceed the maximum value  $N\rho a_0^3 \times b_M$ . This yields  $f_{\max} \sim \rho a_0^3 b_M / W_0$  [24], whence  $\tau \sim W_0 / (\rho a_0^3 b_M)$ . From the expression (3) of  $\sigma$  we deduce

$$\tau \sim \sigma / (\rho b_M) \quad (5)$$

Substituting (1), (2) and (4) into (5) yields

$$\tau = L/V_{\max} \sim 0.1 \text{ s} \quad (6)$$

Consider now an animal of length  $L$ , cross-section  $S_{\perp}$ , and mass  $M \sim \rho S_{\perp} L$ , moving via cyclic (of frequency  $f$ ) motion of appendages such as runners' legs or fishes' tail and fins. For an appendage  $i$  of length  $L_i$ , muscle section  $S_i$ , and tension  $\sigma_i$  rotating by  $\theta_i$  with a lever arm  $l_i$ , the work performed is  $W_i = \Gamma_i \theta_i$  with the torque  $\Gamma_i \sim \sigma_i S_i l_i$ , so that the total work of the appendages is  $W \sim \sum_i \sigma_i \theta_i S_i l_i$ . With the approximations  $\theta_i \sim 1$  radian,  $\sigma_i \sim \sigma$ ,  $L \sim L_i \sim l_i$  and  $S_{\perp} \sim \sum_i S_i$ , the specific work per cycle is thus  $W/M \sim \sigma/\rho$  - a gross simplification which also neglects the role of elastic elements and of gears and joints, and assumes that the applied tension is the maximal one, despite the actual variation of tension with muscle speed. If the step size is approximated by  $L$ , and both the efficiency and the proportion of muscle and active tissue are approximated by 100%, the specific power is therefore  $\sigma f/\rho$ , which cannot exceed  $b_M$ . This yields the time scale (5), i.e. (6), which is drawn as a solid black line on figure 2.

Substituting into (6) the average relation  $M \simeq (2.8L)^3$  deduced from the (logarithmic) average of  $M/L^3$  for the plotted data, we find the maximum speed

$$V_{\max} \sim 3.5 M^{1/3} \quad (7)$$

which is drawn as a solid black line in figure 1 and is very close to the experimental dashed black line.

It may seem surprising that these order-of-magnitude estimates, which are little more than dimensional analysis estimates and make so drastic assumptions as to the actual mechanisms of locomotion [4, 86] agree with observation. It appears that the large number of factors that have been approximated by unity turn out to compensate each other in order of magnitude.

### 3.4. Upper limit to the maximum speed

Figure 1 suggests that the speed reaches a maximum value  $\sim 15$  m/s for masses exceeding  $\sim 50$  kg ( $L \simeq 1.5$  m), which confirms known results [2, 8, 9]. Since this upper limit holds for both running and swimming (with a Reynolds number  $Re$  varying from  $10^{-5}$  for bacteria to more than  $10^8$  for large fishes), it is not expected to result from constraints related to the locomotion style [40, 94] or to the drag, which depend on the ambient density and on  $Re$ . Such a ubiquitous constraint may derive instead from an argument inspired by Hill [95]. Consider again the periodic motion of an appendage of length  $L_i$  and mass  $M_i \sim \rho S_i L_i$ . This motion is expected to be more limited by muscle tension than by the (higher) mechanic strength. With the torque  $\Gamma_i \sim \sigma S_i L_i$  and the angular momentum  $I_i \sim M_i L_i^2$ , the maximum angular acceleration is  $d^2\theta/dt^2 \sim \Gamma_i/I_i \sim \sigma/\rho L_i^2$ . Setting  $L_i \sim L$ , this yields the minimum time for the appendage to reach a significant angle  $\theta \sim 1$  radian

$$\tau_m \sim L(\rho/\sigma)^{1/2} \quad (8)$$

We deduce the upper limit of the top speed  $V_M \sim L/\tau_m \sim (\sigma/\rho)^{1/2} \sim 14$  m/s from (1) and (2), which is plotted (dotted) in figure 1. In figure 2 we have plotted (dash-dotted) the time scale (8) evaluated as a function of  $M$  by using the (logarithmic) average of  $M/L^3$  for the plotted data, which yields  $\tau_m \sim 0.025 \times M^{1/3}$ .

## 4. Discussion

### 4.1. Comparison with previous work

Let us verify that the present results are compatible with empirical scalings previously derived for narrower size ranges and/or restricted taxa. Allometries in living organisms are difficult to determine, mainly because the available data cannot be randomly drawn, truly independent and exactly comparable (see e.g. discussion in [89]), and generally depend on the range of taxa and sizes sampled. Hence we focus on the previously measured ranges of values of  $\tau$  rather than on allometry, and compare them to the range shown in Figure 2.

The maximum relative running speed of mammals has been shown to vary with body mass as  $V_{\max}/L = 21.5 \times M^{-0.17}$  in the range [0.015–6000] kg [9]. Not surprisingly since the set considered in the present paper is nearly the same, this scaling is equivalent to the regression between  $V_{\max}$  and  $M$  found in section 2.3 for mammals with the relevant relation between  $M$  and  $L$ . This scaling yields  $\tau = L/V_{\max} \simeq 0.07$  s in the (logarithmic) middle of that range, and makes  $\tau$  vary by less than a factor of three from this value in the whole studied mass range. For invertebrates, the maximum frequencies found by Full [89] for runners and swimmers yield  $\tau \sim 0.1 - 1$  s in the mass range [ $2 \times 10^{-5} - 0.5$ ] kg. For still smaller masses, the various published relationships for running ants [11] can be converted into  $\tau \sim 0.07 - 0.3$  s at 28 °C, in the mass range [ $3 \times 10^{-7} - 4 \times 10^{-5}$ ] kg. Note also that the running stride frequency has been found to scale with body mass as  $f = 4.32 \times M^{-0.148} \text{ s}^{-1}$  in the range [ $3 \times 10^{-8} - 10^3$ ] kg [51]; this allometry yields  $\tau = 1/f \sim 0.11$  s in the middle of the range with a variation by less than a factor of ten around this value in the whole range. Consider now still smaller masses, i.e. microorganisms. The relationship between the upper bound of swimming speeds of various microorganisms as a function of size published in Ref. [96] yields  $\tau \simeq 1.6 \times d^{0.3}$  s (in SI units; note that the given size  $d$  is an equivalent spherical value), which makes  $\tau$  vary by less than a factor of three around 0.07 s in the mass range [ $2 \times 10^{-15} - 3 \times 10^{-8}$ ] kg. Consider finally fishes and cetaceans. The published relationship [10] can be converted into  $V_{\max}/L = 5.7 \times M^{-0.17}$  in the range [ $10^{-3} - 10^5$ ] kg. This yields  $\tau \simeq 0.26$  s in the (logarithmic) middle of the range, with a variation of less than a factor of 5 around this value in the whole range. Note that the scalings for running mammals and for fishes and cetaceans both correspond to a constant Froude number in a restricted mass range (with a larger value for running mammals), but in both cases the speed levels off at body lengths exceeding about 1 m, and are compatible with our results.

How does the maximum relative speed of organisms compare with the maximum

shortening speed of muscles  $V_m/L_m$  (expressed in units of muscle length  $L_m$ )? For a large variety of species of different taxonomic groups covering a body mass range  $[1.9 \times 10^{-6} - 500]$  kg, Medler [72] found a significant relationship  $V_m/L_m \simeq 3.3 \times M^{-0.17}$  (in S.I. units). However, all the values of  $L_m/V_m$  lie in the range  $[0.026 - 3]$  s, i.e. a time scale for muscles about 3 times larger than the value for the whole animal.

#### 4.2. Some limitations and extensions

As noted in section 2, our result has several limitations since we have excluded locomotory styles which cannot be approximated by the elementary models of section 3, even in order of magnitude. In particular, it is not surprising that the rough constancy of  $\tau$  does not hold for flyers [2], because our proposed interpretation is not expected to hold. Indeed, since they move in air, their speed must be high enough to ensure weight support. Hence their top speed is expected to be so large that it should be determined by air (pressure) drag [97] rather than by muscular forces. Writing that the drag power loss  $\propto S_{\perp} V^3$  ( $S_{\perp}$  denoting the body frontal area) should not exceed the maximum available power  $b_M M$  with  $M \propto S_{\perp} L$  yields  $V_{\max} \propto L^{1/3}$ , which is close to the empirical mass allometry of the maximum flying speed [2].

However, one may consider extensions to very elongated species that were not included in our data (except for the microorganisms for which both mass and length were known), as worms and snakes. It is not easy to persuade a worm [98] to move at its maximal speed, but it is easier for terrestrial snakes. The latter have been found to move at  $V_{\max} \simeq 1.2 M^{1/3}$  for 140 species of snakes over a  $[10^{-3} - 1]$  kg mass range with  $M \simeq (0.6 L)^3$  [99], for slithering by lateral undulations or helical side winding based on travelling waves [100]. This yields  $\tau \simeq 1.4$  s, whereas the red racer - a snake speed champion - can even reach  $\tau \simeq 0.8$  s, near the upper values of the speed time scale found here. Another kind of limbless locomotion based on undulatory travelling waves is that of the sandfish lizard, who can swim in sand at  $V \simeq 0.35 L f$  with  $f_{\max} \simeq 4$  Hz for  $M \simeq 1.6 \times 10^{-2}$  kg and  $L \simeq 8.3 \times 10^{-2}$  m [101]. This yields  $\tau \simeq 0.7$  s, also close to the upper values of the speed time scale found here. The fact that wavelike locomotion yields a time scale in the range of our results - albeit close to the upper values - is not surprising since the order-of-magnitude estimate made in section 3.3 might possibly be generalised to wavelike locomotion by summing over the undulations (of wavelength  $\lambda$ ) along  $L$  (with  $S_i \simeq S_{\perp}$ ), rather than summing on limb sections as  $S_{\perp} \simeq \sum S_i$  (with  $L_i \simeq L$ ). Since the step size is expected to be of the order of  $\lambda$  instead of  $L$ , this should yield  $\tau$  greater than the value given by Eq.(5) by the factor  $L/\lambda$ .

#### 4.3. Temperature effects

The maximum speed is expected to increase with body temperature, as do the reaction rates, and this is indeed observed in ectothermic species. For example the equivalent activation energy  $W_a \simeq 8 \times 10^{-20}$  J for ants [11] - of the same order of magnitude as  $W_0$ , produces a variation in speed of a factor of  $\exp[W_a \Delta T / T^2] \simeq 3.6$  (with  $W_a \simeq 0.5$

eV and the temperatures in eV) over their typical field temperature range of 15 - 35 °C. However, some organisms use compensating strategies to maintain their level of performance. For example, some fishes use different muscle fibres to adapt to changes in temperature [102] or can even change the temperature of their muscles [103]. This is the reason why we have as far as possible selected the organisms' maximum speeds at their typical temperatures, as was also the case for the determination of the values (2) and (4) of the specific tension and of the maximum specific metabolic rate. The data plotted in figures 1 and 2 correspond to body temperatures ranging from about 10 to 40 °C; assuming an equivalent activation energy  $\simeq W_a$ , this should produce a variation in speed of a factor of about 12, which represents a major contribution to the dispersion of the data.

#### 4.4. What basic quantities determine $\tau$ ?

Can we relate  $\rho$ ,  $\sigma$  and  $b_M$ , which determine  $\tau$  according to (5), to more basic quantities? Since these three properties concern Earth-based life, the basic quantities are expected to be ambient conditions at Earth in addition to fundamental physical and chemical constants.

Consider first the mass density  $\rho$ . Life is based on liquid water at Earth, of surface temperature  $T_{\oplus} \simeq 300$  K, and of present atmospheric pressure  $10^5$  Pa near the middle of the range required for liquid water (from the triple point pressure  $\simeq 6 \times 10^2$  Pa to the critical point pressure  $\simeq 2 \times 10^7$  Pa). The density  $\rho$  of living matter is close to that of water, of order of magnitude  $\rho \sim 18 \times m_p/a_{\text{H}_2\text{O}}^3$ , where  $m_p$  is the proton's mass and  $a_{\text{H}_2\text{O}} \simeq 0.3$  nm is the size of the  $\text{H}_2\text{O}$  molecule.

Second, consider the tension  $\sigma$  of muscles and molecular motors, whose order of magnitude is mainly determined by the elementary energy  $W_0$  and size  $a_0$  of life molecular motors from (3). The quantity  $W_0$  - the standard free energy involved in biological molecules (of the same order as the energies associated with H (hydrogen)-bonds [104]), can be expressed in terms of the temperature  $T_{\oplus}$  as

$$W_0 \simeq 12 kT_{\oplus} \tag{9}$$

which is high enough compared to  $kT_{\oplus}$  for not being destroyed by thermal agitation, but small enough for not being too far from thermal equilibrium. A similar reasoning might determine  $a_0$  since, as Schrödinger first noted [105], biological structures must be relatively large in order to limit the relative importance of fluctuations, which in the case of proteins, requires a number  $N$  of amino-acids satisfying  $N^{-1/2} \ll 1$  to enable robust folding by limiting structural fluctuations (see e.g. [106]). More precisely, one may argue that the motor force  $W_0/a_0$  should be weak enough for not breaking the motor's 3-D structure held by weak forces such as H-bonds [107]. Writing that the motor force should not exceed the thermal energy  $kT_{\oplus}$  divided by the distance over which H-bonds operate, i.e. the size  $a_{\text{H}_2\text{O}}$  of the  $\text{H}_2\text{O}$  molecule, we get [22]

$$a_0 \gtrsim a_{\text{H}_2\text{O}} \times (W_0/kT_{\oplus}) \simeq 4 \text{ nm} \tag{10}$$

which suggests that  $\sigma$  is determined via (3) by fundamental constants, in addition to the temperature and the existence of liquid water.

Third, consider the energy requirements. Where do the number given in Eq.(4) come from? Life is sustained by a continuous flow of ordered energy [105], mainly acquired at Earth through solar radiation and chemical energy contained in food. Solar radiation (of effective temperature  $T_{\odot} = 5700$  K) provides at the top of the Earth's atmosphere an average energy flux  $S_{\odot}/4$ , where  $S_{\odot} = 1.36 \times 10^3$  W m<sup>-2</sup> is the solar constant, and in the absence of atmospheric greenhouse effect would yield an average temperature at the Earth surface

$$T = (W_{\text{ph}}/\sigma_{SB})^{1/4} \quad (11)$$

where  $W_{\text{ph}} = (1 - A)S_{\odot}/4 \simeq 240$  W m<sup>-2</sup>,  $A \simeq 0.3$  is the mean albedo and  $\sigma_{SB}$  is the Stefan-Boltzmann constant. The actual temperature  $T_{\oplus}$  is higher, while remaining of the same order of magnitude and below the run-away greenhouse limit [108].

Since living organisms acquire energy through their body surface  $S$ , but spend it within their body mass  $M$ , the way to become larger during evolution (thereby decreasing the ratio  $S/M$ ) while keeping a constant specific metabolic rate, is to devise more efficient means of delivering energy: becoming multicellular, increasing internal temperature and using sophisticated distribution networks [17].

Therefore we must consider the basic constraints to energy inputs at the smallest size level, that of a micro-organism or a cell deriving energy from photosynthesis (assuming that complementary resources are available). Photosynthesis is presently used essentially by non-motile organisms as plants, except for a minority of bacteria having photosynthetic capabilities [109] but the situation was different earlier in the evolution of life [110]. Note that with  $T_{\odot} \simeq 20 T_{\oplus}$  the typical solar photon energy  $h\nu$  exceeds the standard free energy (9) involved in biological molecules. Assuming for simplicity a spherical organism of diameter  $L$  (of transverse surface-to-mass ratio  $1.5/(\rho L)$ ) fully exposed to solar radiation and able to convert it, the maximum specific metabolic rate is then

$$b_{M\text{ph}} \sim 1.5\beta_{\text{ph}}W_{\text{ph}}/(\rho L) \quad (12)$$

where  $\beta_{\text{ph}} \sim 0.1$  is the maximum yield of photosynthesis [111]. Substituting  $W_{\text{ph}}$ ,  $\beta_{\text{ph}}$ , and  $\rho$  yields

$$b_{M\text{ph}} \sim 3 \times 10^{-2}/L \quad (13)$$

Comparing (13) with the empirical value (4), it follows that the maximum metabolic rate can be sustained by photosynthesis (which requires  $b_M \lesssim b_{M\text{ph}}$ ) if  $L \lesssim 15$   $\mu\text{m}$ . It is remarkable that this crude estimate is of the same order of magnitude as the average size of an animal cell [19]. Larger motile organisms have indeed different shapes or organisation, or derive their energy from other sources. Hence, if the cell size  $L$  is determined [19, 20, 112], the solar radiation  $W_{\text{ph}}$  might determine  $b_M$  via (12) and therefore the time scale  $\tau$  via (5), since  $\rho$  and  $\sigma$  are determined by fundamental constants and the temperature - itself determined by solar radiation.

## 5. Concluding remarks

A few remarks are in order. First of all, the constancy of the locomotor time scale is rough and approximate, as is the case for the specific tension and the maximal specific metabolic rate which determine it according to (5). Yet this approximate constancy - with 98% of the organisms considered here having a time scale between  $10^{-2}$  and 1 second, holds throughout a  $10^{20}$ -fold mass range of aquatic and terrestrial species, with in particular bacteria, ants, fishes and mammals performing roughly as fast as ostriches. For motion in water, it is remarkable that the time scale remains approximatively the same at Reynolds numbers varying from about  $10^{-5}$  for bacteria (for which the drag force  $\propto V$  is determined by viscosity whereas inertia is negligible) to more than  $10^8$  for large fishes (for which the drag force  $\propto V^2$  is determined by dynamic pressure).

As noted in section 4.1, this large scale order-of-magnitude constancy does not preclude - and is indeed compatible with - various scalings valid in narrower ranges of size and taxa. When considering narrow ranges, the detailed properties of the organisms and their locomotion style must be taken into account, as for example the proportion of active tissue or the role of elastic elements; these properties are expected to change the scaling, and it is only on a very large scale that these variations can be transcended. Similarly, as noted in section 2.1, even the large-scale scalings may be criticized because our data are not truly random, independent and comparable. However, the constancy of the order of magnitude of the motional time scale over the entire mass range is a more robust result, supported by our interpretation (Eq. 5) which neglects factors able to affect the scaling with  $M$  without changing the order of magnitude of  $\tau$ .

It is worth noting that the special case of wheel-like locomotion is also accounted for. Indeed, consider the stomatopod somersaulters, who invented the (macroscopic) wheel [113, 114], whereas swimming bacteria use (microscopic) wheels when moving their flagella with rotary motors [115]. These stomatopods propel themselves on beaches, taking the shape of a wheel, in a motion akin to that of a one-legged animal [114] at a maximum speed of 5.6 cm/s for a 2 cm length [116], i.e. a time scale of about 0.4 s, well within the range shown in Fig. 2. This is not surprising, given the similarity in energetics and dynamics of locomotion whatever the number of legs [117].

It is important to keep in mind that, as noted in the introduction, alternative interpretations of the time scale of order of magnitude 0.1 s might be considered since this value lies in the middle of the range of other relevant biological time scales [118]. In particular, it equals the time scale corresponding to the median maximal turn-over rate of enzymes catalyzing biological reactions [119]. Other constraints or scalings might also be found.

Finally, the discussion leads us to conjecture that the values of  $\rho$ ,  $\sigma$  and  $b_M$ , which determine the ubiquitous motional time scale according to our proposal, are themselves determined by the available solar energy, in addition to fundamental physical and chemical constants. This contrasts with the preferred speed (mainly determined by gravity [120]), which may have consequences for putative life on other planets [121].

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