

Stride lengths, speed and energy costs in walking of *Australopithecus afarensis*: using evolutionary robotics to predict locomotion of early human ancestors

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This paper uses techniques from evolutionary robotics to predict the most energy-efficient upright walking gait for the early human relative *Australopithecus afarensis*, based on the proportions of the 3.2 million year old AL 288-1 ‘Lucy’ skeleton, and matches predictions against the nearly contemporaneous (3.5–3.6 million year old) Laetoli fossil footprint trails. The technique creates gaits *de novo* and uses genetic algorithm optimization to search for the most efficient patterns of simulated muscular contraction at a variety of speeds. The model was first verified by predicting gaits for living human subjects, and comparing costs, stride lengths and speeds to experimentally determined values for the same subjects. Subsequent simulations for *A. afarensis* yield estimates of the range of walking speeds from 0.6 to 1.3 m s⁻¹ at a cost of 7.0 J kg⁻¹ m⁻¹ for the lowest speeds, falling to 5.8 J kg⁻¹ m⁻¹ at 1.0 m s⁻¹, and rising to 6.2 J kg⁻¹ m⁻¹ at the maximum speed achieved. Speeds previously estimated for the makers of the Laetoli footprint trails (0.56 or 0.64 m s⁻¹ for Trail 1, 0.72 or 0.75 m s⁻¹ for Trail 2/3) may have been underestimated, substantially so for Trail 2/3, with true values in excess of 0.7 and 1.0 m s⁻¹, respectively. The predictions conflict with suggestions that *A. afarensis* used a ‘shuffling’ gait, indicating rather that the species was a fully competent biped.

Keywords: biomechanics; bipedalism; gait; evolution; palaeontology; simulation

1. INTRODUCTION

The adoption of bipedalism as the preferred mode of terrestrial locomotion is a fundamental step in divergence of the human lineage from that of other African apes. Human walking is characterized by extended postures of the hip and knee joints so that gait is relatively ‘stiff’. This ‘stiff’ gait brings about out-of-phase oscillations of the kinetic and potential energies of the centre of mass (CM) which permit energy to be exchanged between the two states conserving up to 70% of the energy of one stride for use in the next. There are, however, transitory knee-flexions at heel-strike, mid-stance and toe-off, which reduce both excessive vertical displacement of the body CM and peak vertical ground reaction forces. Bipedally walking chimpanzees by contrast use ‘compliant’ gaits where knee-flexions are sufficient and long-lasting enough to bring the oscillations of kinetic and potential energies of the CM into

phase, eliminating mechanical energy exchange, and thus increasing metabolic costs. When walking bipedally the orang-utan does maintain extended hip and knee joints (Crompton *et al.* 2003) and up to 52% energy exchange occurs, but no ape appears to be able to combine extended hindlimb posture with the long strides and high stride frequencies that are required for fast bipedal walking. Human foot function also differs from that of other apes in that thrust is imparted to the ground by the fore-foot, maximizing leverage of triceps surae, the major muscle powering heel-off and toe-off. Other apes achieve peak plantar pressures and deliver thrust in the mid-foot, reducing leverage about the ankle joint. Thus the important locomotor innovation characterizing the human lineage is likely to be the development of a long, forceful and yet efficient stride rather than upright walking *per se*. When did this occur? There are unfortunately only a limited number of feet in the fossil record of early human ancestors: the most complete is the *Homo habilis* foot OH 8, dating to about 1.8 million years old. This is contemporaneous

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with a very modern looking, skeleton, WT 15 000, of *Homo ergaster*. WT 15 000 was clearly a human-like striding biped but, while otherwise nearly complete, lacks feet.

Recent analyses of the OH 8 foot suggested it is 'ape-like' on the medial side, lacking an arch and having a mobile midfoot and opposable big toe (Kidd 1995) but 'human-like' with a stabilized midfoot, on the lateral side. Older material exists for the early human ancestor *Australopithecus afarensis*, dating to about 2–3.2 million years old. The partial feet of skeletons AL 288-1 and AL 333 also appear to lack a medial arch, and the *Australopithecus* Sts 573 foot specimen again shows an apparent 'mix' of 'human-like' and 'ape-like' features (Harcourt-Smith & Aiello 2004).

Direct evidence of foot function in human ancestors at the time of Sts 573 and *A. afarensis* is provided by the Laetoli footprint trails (3.5–3.8 million years old) made in wet volcanic ashfall deposits (Hay 1987). While Trail A, with short, broad, and poorly defined prints, and stride length resembling that of a 2.5 year old human child, may not represent a human ancestor, but instead have been made by the extinct ursid (bear ancestor) *Agriotherium africanus* (Robbins 1987), Trail G1 represents at least four complete cycles of bipedal walking by a small individual of a human ancestor, most likely a child. Alongside it run two partly superimposed trails G2 and G3, with very similar morphology but wider spacing, which are likely to represent adults of the same species. White (1980) regarded the Laetoli G prints as consistent with authorship by *A. afarensis* on the basis of the documented relationship of adult human skeletal stature to adult human footprint length. Jungers (1982) argued, however, that the short legs of *A. afarensis*, as represented by the famous 'Lucy' skeleton AL 288-1 implied short stride lengths like those in bipedal walking of the living non-human apes. Recently discovered but as yet undescribed postcranial material of *Australopithecus* from about four million years ago, and apparently of considerably greater stature, may yet prove to have been responsible for the Laetoli prints, but at present *A. afarensis* remains the best contemporaneous size-match for the Laetoli prints and the most likely maker. Stern & Sussman (1983) suggested that *A. afarensis* would have walked 'bent-hip, bent-knee' (BHBK), rather like a living chimpanzee, because of the number of skeletal features in the *A. afarensis* skeleton which are functionally associated with arboreality in living apes (e.g. curved phalanges, long trunk but short legs, etc.). Hunt (1994) has even proposed that the most efficient behaviour for *A. afarensis* would have been 'bipedal posture augmented with bipedal shuffling' as a consequence of anatomical compromise between the needs of terrestrial bipedality and arboreal climbing. However, other authors (e.g. Latimer 1991) stress characteristics associated in humans with habitual bipedalism (such as a femur where the shaft is angulated to bring the knees under the centre of gravity of the body) and conclude that this species was an habitual, erect biped. This view is now becoming the general consensus (see e.g. Ward 2002).

Many of the disputes which surround the evolution of human bipedalism arise because there is no other living analogue for early bipeds but modern humans, while the body proportions of *A. afarensis*, for example, are so different from our own (short legs and long trunk versus our own long legs and short trunk) that the mechanics of motion must have been very different (e.g. Alexander 2004). Rather than trying to interpret the behaviour of such species by a combination of analogies to humans in certain anatomical regions with analogies to other apes elsewhere, it seems sensible to adopt a reverse-engineering approach and determine what kind of locomotion a particular set of body proportions were best 'designed' to perform. Since the locomotor system is concerned primarily with the application of external force by the body, simulation techniques drawn from mechanical engineering are a potential means of predicting the significance of differences in proportions for the motion and force characteristics of bipedal locomotion.

There are two main techniques for investigating locomotor mechanics: inverse dynamics where forces are calculated based on observed movement and forward dynamics where movement is calculated based on applied forces (for review see van den Bogert 1994). In an earlier study we showed using inverse dynamic modelling that the skeletal proportions of *A. afarensis* (AL 288-1) were compatible with stable and efficient upright bipedalism (Crompton *et al.* 1998) but that mechanical energy costs of locomotion were near-doubled in BHBK gait. (Carey & Crompton 2005 have since confirmed for humans that physiological costs are indeed much higher in BHBK gait). Kramer (1999) and Kramer & Eck (2000) used similar inverse-dynamic techniques and also concluded that this species could have been an efficient upright biped. While inverse dynamics is computationally economical and has usefully been applied to assessing alternative hypotheses as to the gait of early human ancestors (Crompton *et al.* 1998; Kramer 1999; Kramer & Eck 2000; Wang *et al.* 2004) it requires that given sets of joint motions are applied to given sets of body proportions. This is not entirely satisfactory as while we often have fossil evidence of body proportions joint motion is of course not directly preserved in the fossil record. Moreover inverse dynamics can only be used for direct estimates of mechanical work. The relationship between mechanical work and metabolic energy cost is complex. For example muscles require energy to generate force even when no mechanical work is being performed or even when doing so-called 'negative work' (Winter 1990). However, recent advances in our ability to model muscle energetics (Minetti & Alexander 1997; Umberger *et al.* 2003) together with the greater availability of facilities for computationally intensive modelling have together facilitated adoption of forwards dynamic modelling. This approach does not require imposition of given sets of joint kinematics but rather predicts them from body proportions and muscle activation patterns. Musculoskeletal model based forwards dynamics simulations can calculate metabolic energy costs directly, and have been shown to produce reliable results in human walking (Sellers *et al.* 2003).

Results have also been reported for *A. afarensis* (Sellers *et al.* 2004; Nagano *et al.* 2005). Since they use a finite-state controller to activate the muscles in the model and hence to drive bipedal locomotion, the approach adopted by the both groups need make no assumptions concerning the pattern of muscle activation.

Sellers and colleagues (2003, 2004) set the search pattern to maximize locomotor efficiency; Nagano *et al.* (2005), however, sought to match a predefined kinematic configuration at the start and end of a step whilst simultaneously minimizing the metabolic energy consumed per unit distance travelled during the step. There are merits to both approaches. In the former the model is free to select any kinematic pattern which can produce forward motion and therefore makes no assumptions about the correct kinematics for the fossil animal. In the latter there are some limited assumptions about the required kinematics but these can considerably reduce the size of the search space and may improve gait cycle repeatability, allowing a much more complex model to be constructed. However, since the model constrains certain kinematic features such as speed and stride length it cannot easily be used to investigate the preferred speeds and stride lengths that suit the mechanics of the fossil (Nagano *et al.* 2005).

Our optimization method is drawn from evolutionary robotics, a technique for the automatic creation of autonomous robots (Nolfi & Floreano 2000). Whilst primarily concentrating on navigation and learning issues, evolutionary robotics also covers the spontaneous generation of locomotion in legged robots. It is therefore a tool with which we can investigate the locomotion of fossil animals. In this case values for segment proportions, joint mobilities, and muscle arrangements are either derived directly from the fossil evidence or represent anatomically likely values. A gait controller is used to activate the muscles to produce the required locomotion. Building the simulator is relatively straightforward computationally but finding an appropriate pattern to produce stable gait is difficult because the number of possible patterns is far too large for them all to be tested individually. However, such problems can be addressed effectively using genetic algorithms.

Evolutionary processes have led to highly optimized solutions to complex problems: thus, since the 1950s evolutionary theory has been used by computer scientists as a source of inspiration for optimization and machine learning algorithms. Evolutionary strategies (Rechenberg 1965) are the most developed of the early techniques. They encode the problem under consideration as a sequence of real numbers and then randomly mutate these numbers. Each time a mutation is created it is compared with the previous solution and if it performs better according to some metric criterion it replaces the original solution; otherwise it is discarded. By using a Gaussian mutation (adding a random value selected from a Gaussian probability distribution with a mean of zero) it is possible to generate any sequence, although the new sequences are more likely to be similar to the previous sequence than otherwise. This approach takes advantage of the fact that sequences close to the 'best' solution are likely to

be similar to that solution. Genetic Algorithms were invented by John Holland in the 1960s (Holland 1975). The genetic algorithm uses a population of solutions. Members of this population (called chromosomes) are allowed to contribute to the next generation by crossover, whereby two chromosomes exchange subsequences to create two new chromosomes. The selection of parent chromosomes is done randomly but is influenced by their fitness which is calculated in some way by a fitness function. The genetic algorithm *sensu strictu* uses a fixed length sequence of bits (zeros or ones) as its chromosome. However, the genetic algorithm has had such a large impact on the field of evolutionary computation that concepts such as populations and crossover have been incorporated into other techniques and the term is now used to cover almost any population-based evolutionary search technique (for a more thorough introduction to genetic algorithms see Davis 1991). These evolutionary techniques have been widely used for difficult computational search problems and are ideal for finding sets of parameters in gait controllers that produce high quality gaits. An additional advantage is that because they work on populations of solutions they are ideal for parallel implementation, and thus allow extremely high performance (van Soest & Casius 2003).

This study therefore applies forwards dynamic analysis and genetic algorithm optimization to reconstruct the metabolic costs associated with different stride lengths and speeds for *A. afarensis*, on the assumption that this species walked upright, following Nagano *et al.* (2005) and go on to apply our results to interpretation of the Laetoli footprint trails, assuming again that *A. afarensis* was the maker.

2. METHOD

It is extremely important that simulation studies are validated against experimental data wherever possible. Therefore, the first step in this study was to obtain such data experimentally for human subjects and to produce simulations based on those particular subjects' morphology to allow direct comparison between experimental and simulated values. This will then allow us to evaluate the likely accuracy of any subsequent *A. afarensis* simulation.

To obtain the required experimental data five healthy young adult male human subjects were required to walk on a treadmill at speeds from 0.5 to 2.0 m s⁻¹, in 0.25 m s⁻¹ increments covering the subjects' full range of walking speeds. The subjects were asked to walk at each speed for several minutes to allow the oxygen consumption rate to reach equilibrium. Then the subjects' expired air was collected using a Douglas bag and analysed to estimate metabolic power using the Weir (1949) formula. Simultaneously, the subjects' stride length was calculated by measuring the time taken for 30 strides. In addition the subjects' metabolic rate was also measured standing (i.e. progressing at 0 m s⁻¹) to allow the calculation of net locomotor costs (Heglund & Schepens 2003). The subjects were then weighed and the distances between neighbouring joint centres (first metatarsal head, ankle,

Table 1. The values for the inertial properties of the segments used in the computer simulation models.

(HAT length is from hip joint to shoulder joint, thigh from hip joint to knee joint, leg from knee joint to ankle joint, foot from ankle joint to first metatarsal head. CM position is the distance along the length of the segment (distally for the leg segments and cranially for the HAT segment). Moments of inertia are about the local origin of each segment which is the proximal joint except for the HAT which is about the hip joint.)

		subject 1	subject 2	subject 3	subject 4	subject 5	AL 288-1
HAT	mass (kg)	50.7144	56.7147	60.2403	38.9850	53.8671	22.3740
	length (m)	0.5490	0.5020	0.5190	0.4850	0.4920	0.3516
	MOI (kg m ²)	9.7338	9.1014	10.3330	5.8396	8.3035	1.7610
	CM (m)	0.3437	0.3143	0.3249	0.3036	0.3080	0.2201
thigh	mass (kg)	7.4800	8.3650	8.8850	5.7500	7.9450	3.3000
	length (m)	0.4550	0.3940	0.4470	0.4100	0.4760	0.2520
	MOI (kg m ²)	0.4516	0.3787	0.5177	0.2819	0.5249	0.0611
	CM (m)	0.1970	0.1706	0.1936	0.1775	0.2061	0.1091
leg	mass (kg)	3.4782	3.8897	4.1315	2.6738	3.6944	1.5350
	length (m)	0.4520	0.3980	0.4370	0.3150	0.4790	0.2650
	MOI (kg m ²)	0.1981	0.1718	0.2200	0.0740	0.2363	0.0300
	CM (m)	0.1957	0.1723	0.1892	0.1364	0.2074	0.1150
foot	mass (kg)	1.0846	1.2129	1.2883	0.8338	1.1520	0.4790
	length (m)	0.1690	0.1340	0.1480	0.1290	0.1530	0.0800
	MOI (kg m ²)	0.0147	0.0104	0.0134	0.0066	0.0128	0.0015
	CM (m)	0.0845	0.0670	0.0740	0.0645	0.0765	0.0400

knee and hip and also the shoulder joint) were measured to allow calculation of the segment inertial properties using the formulae recommended by Winter (Winter 1990).

Five separate computer simulations were then constructed to match the morphology of the subjects. Due to the non-deterministic nature of the modelling process they were each run multiple times. The basic design was the same as we have used previously (Sellers *et al.* 2003, 2004) with three segment left and right hindlimbs (foot, leg and thigh) and a unified head, arms and trunk (HAT) segment. The segment lengths were customized to match those of the human subjects, and assigned the calculated inertial properties (table 1).

Since this is a two-dimensional simulation the joints were all defined as hinge joints with appropriate ranges of motion (Delp *et al.* 1990). Muscles were grouped as primary flexors and extensors around the hip, knee and ankle joints; and muscle physiological cross section areas and fibre lengths of the groups were assigned by identifying the actions of the muscle and calculating a weighted average based on the physiological cross sectional area (PCSA) for each muscle contributing to the group based on data for individual muscles (Pierrynowski 1995). Attachment points were assigned based on the estimated centroid of the attachment area for a muscle group. For the knee extensor and flexors a wrapping operator was used to maintain the moment arm around the joint at an estimated 6 and 3 cm. The tendon lengths were calculated so that the muscle fibre length plus the tendon length was equal to the total path length in the anatomical position. The muscle model used was that of Umberger and colleagues (2003), based on FORTRAN code kindly supplied by Umberger. This model also requires an estimate of serial and parallel elastic properties: these were calculated assuming a 6% strain in the tendon, and a 60% strain in the passive connective tissue component

of the muscle at its maximum isometric contraction, since these values appear to be typical of vertebrate skeletal muscle (Biewener 2003). The other values required for the model were set based on mean values for lower-limb muscles (Nagano & Gerritsen 2001; Umberger *et al.* 2003). The values used are shown in tables 2 and 3.

The *A. afarensis* model was produced from the human model by scaling the latter to the proportions of the AL 288-1 fossil skeleton (Johanson *et al.* 1982). A number of similar models have been produced previously (Crompton *et al.* 1998; Kramer 1999; Kramer & Eck 2000; Sellers *et al.* 2004; Wang *et al.* 2004; Nagano *et al.* 2005) although all differ in the numerical values chosen for various parameters. Segment lengths (the distance between joint centres rather than the lengths of the bones themselves) were those adopted by Kramer (1999) except for the HAT length (defined as the distance from the hip to the shoulder joint centre) which was estimated from the Lovejoy reconstruction (Weaver *et al.* 1985). There is considerable disagreement in literature concerning estimated total body weights: values vary from 12.3 to 38.9 kg (Aiello & Dean 1990). The simulations listed above all use values close to 30 kg; we chose 33 kg following Kramer (1999). In our previous work we reconstructed the inertial parameters of the segments based on both modern human and modern chimpanzee values. However, we found very little effect of mass distribution on results (Sellers *et al.* 2004) so in this paper we used human values reconstructed according to Winter's method (Winter 1990). Muscle values were obtained directly by scaling from the human values since we have no way of estimating these soft tissue parameters. The muscle length was scaled based on the segment-length scaling from the human to the AL 288-1 models, while muscle PCSA was scaled so that relative muscle volume was maintained (i.e. area scale = mass scale/length scale).

Table 2. The values used for the muscles in the human and *A. afarensis* (AL 288-1) model.

(Note that the origins and insertions are in segment local coordinates with the first value being the distance distally along the segment and the second value being distance perpendicular to the segment axis.)

		human		AL 288-1	
hip extensor	origin (m)	0.052	-0.076	0.037	-0.055
	insertion (m)	0.3	0.005	0.191	0.003
	PCA (m ²)	0.0137	—	0.0068	—
	fibre length (m)	0.13	—	0.082	—
hip flexor	origin (m)	-0.081	0.071	-0.059	0.052
	insertion (m)	0.16	0.029	0.102	0.018
	PCA (m ²)	0.0075	—	0.0037	—
	fibre length (m)	0.15	—	0.095	—
knee extensor	origin (m)	0.25	0.029	0.159	0.018
	insertion (m)	0.076	0.043	0.047	0.027
	PCA (m ²)	0.0203	—	0.0102	—
	fibre length (m)	0.106	—	0.067	—
knee flexor	origin (m)	0.25	0	0.159	0
	insertion (m)	0.08	-0.01	0.05	-0.007
	PCA (m ²)	0.0159	—	0.008	—
	fibre length (m)	0.104	—	0.066	—
ankle dorsiflexor	origin (m)	0.14	0.033	0.087	0.02
	insertion (m)	0	0.057	0	0.018
	PCA (m ²)	0.0051	—	0.004	—
	fibre length (m)	0.089	—	0.055	—
ankle plantarflexor	origin (m)	0.057	-0.029	0.036	-0.018
	insertion (m)	0.038	-0.057	0.023	-0.035
	PCA (m ²)	0.0264	—	0.0208	—
	fibre length (m)	0.055	—	0.034	—

Table 3. The general modelling parameters used in the model.

(Note that the joint range angles are measured from the anatomical position and are measured in an anticlockwise direction assuming the subject is facing in the direction of the *X* axis with the *Y* axis pointing vertically.)

general muscle parameters	force per unit area (N m ⁻²)	250 000	—
	relative V_{\max} (s ⁻¹)	12	—
	width of force-length curve	0.97	—
	fast twitch proportion	0.43	—
	density (kg m ⁻³)	1059.7	—
joint range (°)	hip	-10	120
	knee	-90	0
	ankle	-40	33

Knee moment arms were reduced to 4 and 2 cm. The values used are shown in tables 1 and 2.

The biomechanical simulation is performed using custom-written software written in C++ using the open-source Dynamechs simulation library (McMillan *et al.* 1995). This simulator is based on the highly efficient Featherstone algorithm (Featherstone 1987) so that joint constraints are solved analytically rather than by using the more commonly applied 'penalty' methods. The simulator is implemented as a set of C++ classes which are incorporated into a user-written program. The parameters of the model are defined using an XML file which includes instructions that specify the muscle activation pattern during the simulation. The simulator solves the equations of motion and produces movements based on the external and internal forces generated by the model. Ground interaction was achieved using standard Dynamechs point contact elements attached to the front and back

of the foot. These incorporate both static and dynamic friction using the standard linear Coulomb model and using damped elastic penetration to generate ground reaction forces. Elastic parameters were adjusted to restrict interpenetration to a few centimetres whilst still allowing a reasonable integration step-size. Too high a ground stiffness rapidly produced numerical instability in the model. Joint limits were similarly implemented as relatively stiff, damped torsional springs that activated when the joint reached its anatomical limit.

To obtain stable bipedal walking there needs to be a suitable activation pattern. This is produced by a separate program that implements the genetic algorithm optimization. The separation of the genetic algorithm from the simulator allows extremely flexible deployment with versions of the software running on Unix, Windows and MacOSX and running at multiple sites connected via the Internet. This allows multiple

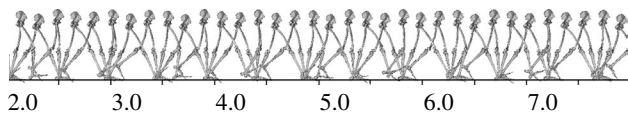


Figure 1. Overlay image of animation output of walking simulator for the *A. afarensis* model. The scale is in metres and the individual images are taken at 0.2 s intervals. The movie file from which this animation was produced is available in the electronic supplementary material.

simulations to be run in parallel on separate computers rather than sequentially on a single computer which very greatly increases the overall speed. Even so it can take many thousands of simulation runs to find a good solution and it is important to keep the simulation as simple as possible and to minimize the complexity of the control system so that the search process can run as quickly as possible. The current implementation uses three activation phases per step and for each phase the model requires a phase duration as well as the muscle activation around the six joints. Thus the whole model is controlled by 21 parameters which are reused for subsequent steps by reversing the right- and left-side activations at each step. The activation pattern provides a relative activation signal for the muscular elements in the model (a value between 0 and 1 maintained for the duration of the phase). This is converted into a muscle activation using the first order model of He *et al.* (1991) as implemented in Nagano & Gerritsen (2001) and Umberger *et al.* (2003). Co-activation was not allowed in this model although limited co-contraction could occur because of the time lag inherent in the first order activation model. The ‘fitness’ of the simulator is measured by the forward progression made by the model before it uses up a fixed amount of metabolic energy or falls over; this ensures that unstable patterns have a very low fitness. However, this fitness condition alone will only generate a very limited range of speeds at the slow end of values for walking (Sellers *et al.* 2004). To force the model to walk faster a linear time-dependent penalty function was used whereby the energy available to the model decreased over time. With a larger rate of energy decrease the model adopts faster walking speeds since they are now more efficient, which forced the model to adopt a larger range of speeds whilst still favouring the most efficient gait for any given speed. The highest walking speeds were never used since the model would spontaneously adopt running or skipping gaits, since these are more energy efficient than high-speed walking gaits.

3. RESULTS

Figure 1 shows a set of overlay images produced by the simulator for the *A. afarensis* model. The simulation shows a fairly naturalistic walking style but there are artifacts due to the fact that each step only has three independent phases. This means that the activation level of all muscles in a group is constant within a particular phase, so that some aspects of the kinematics (such as the dorsiflexion of the ankle joint during the beginning of the support phase) are an unavoidable

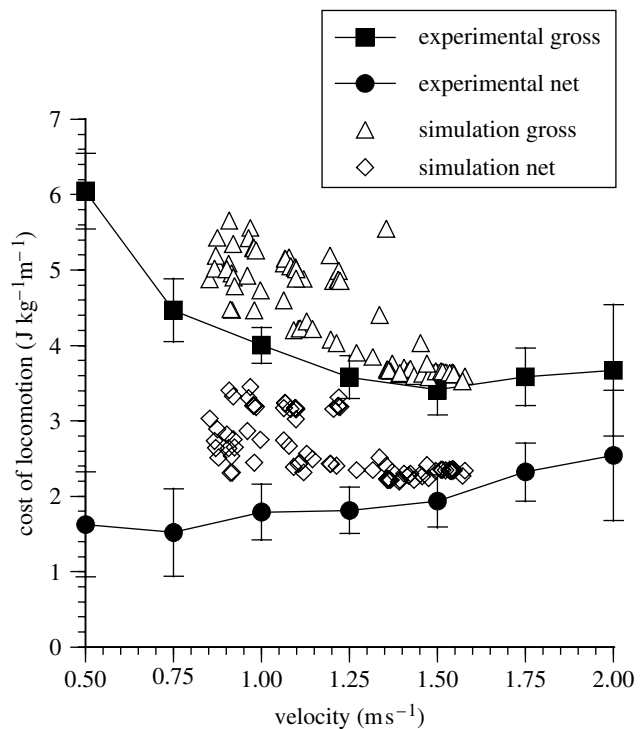


Figure 2. The cost of locomotion for the five experimental subjects and the cost of locomotion estimated by the simulator. The error bars represent the 95% confidence limits of the mean. Simulated gross values represent the cost of walking plus the cost of standing. The simulation is required to minimize costs; thus it is the *lowest* value at a given speed that represents the model’s best estimate.

compromise. These effects would be reduced by increasing the number of phases in the gait cycle but this would greatly increase the computational time required. The kinematics approximate ‘compass gait’, an extreme form of ‘stiff’ walking (Cavagna *et al.* 1977; Alexander & Jayes 1978). The ‘compass gait’ pattern derived is in part due to the model’s two-dimensionality: in human walking the pelvis is rotated forwards on the swing-leg side in a horizontal plane just before that leg makes ground contact, and then tilted in the transverse plane during mid-stance (an action emphasized during speed walking) both of which reduce the maximum vertical movement of the centre of gravity. These ‘determinants of gait’ (Saunders *et al.* 1953) obviously cannot occur in our two-dimensional simulation.

Figure 2 shows the results from the experimental measurement of cost of locomotion along with the results from the simulator. The experimental data produced the expected U-shaped relationship between gross cost of transport and walking speed. The most efficient walking occurred at about 1.5 m s^{-1} . The metabolic costs calculated by the simulator are entirely musculoskeletal costs: they do not for example include any costs of the ‘expensive tissues’ such as the brain or liver (Aiello & Wheeler 1995). Thus by subtracting the cost of standing from the cost of walking we can correct for most of this discrepancy between simulated and experimental results. The much flatter curve produced shows a reasonable match between experimental and simulated results. There is then very good agreement

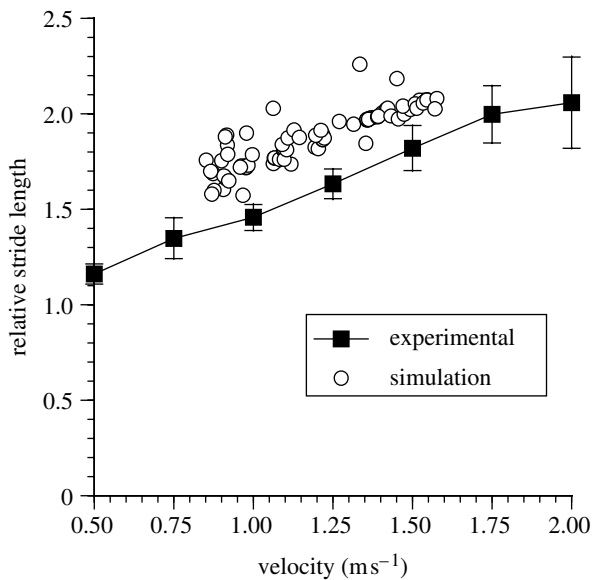


Figure 3. The relative stride length values obtained for the five experimental subjects and for the simulator. The error bars represent the 95% confidence limits of the mean. Note that the relative stride length is calculated by dividing the actual stride length by the total leg length (the sum of the thigh and leg segments).

with our experimental data over the full speed range (bearing in mind that the actual cost will always be lower than the lowest predicted value, given the limitations of the model). We can also reconstruct the gross cost for the simulator, by adding back in the cost of standing for the experimental subjects and this is also shown in the figure.

Figure 3 shows the results from the experimental measurement of stride length along with the results from the simulator. Grieve & Gear (1966), Charteris *et al.* (1982) and Alexander (1984) normalize stride length by stature. But since stride length is necessarily strongly influenced by leg length (see e.g. Jungers 1982) and since any flexed trunk posture may reduce biomechanically effective 'stature' (Alexander 1984), normalization against leg length is probably more appropriate, and is adopted here. Recalling that stride length was not itself one of the criteria optimized there is good agreement between the experimental and simulation data over the full range of simulated speeds, although the simulator consistently overestimates the stride length for a given speed (or conversely underestimates the speed for a given stride length).

Figure 4 shows the results from the *A. afarensis* simulation, and includes for comparison cost of locomotion data for young children (of similar body size to AL 288-1) taken from the literature (Heglund & Schepens 2003). Once again the simulator produces values which are reasonably faithful estimates of the experimental net cost of locomotion. The simulated speed range 0.6–1.3 m s⁻¹ is certainly consistent with allometrically predicted values for walking in animals between 9 and 25 kg. (cf. Heglund & Taylor 1988). In order to estimate the gross cost of locomotion we need to estimate the cost of standing in *A. afarensis*. The power cost of standing with straight knees for 5–6 year

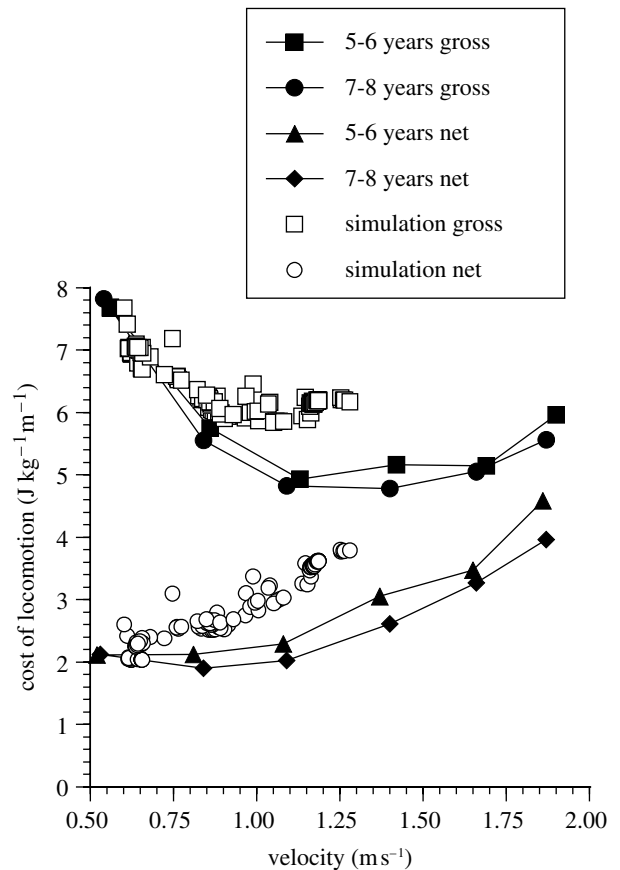


Figure 4. The cost of locomotion for the *A. afarensis* simulator. For comparison the graph also shows the experimentally derived cost of locomotion for modern human children of similar body mass and stature to *A. afarensis* (redrawn using data from Heglund & Schepens 2003).

old children is 3.08 W kg⁻¹ and for 7–8 year olds is 3.02 W kg⁻¹ (Heglund & Schepens 2003) so a value of 3.05 W kg⁻¹ was used to calculate the gross cost of locomotion from the simulator.

Figure 5 shows the relationship between walking speed and stride length for the simulator. We have added relative stride length information which may pertain directly to *A. afarensis*, derived from the 3.5 to 3.8 million year old footprint trails at Laetoli, Tanzania, plotted against velocities estimated by Charteris *et al.* (1982) and Alexander (1984). We have not included the site A trail because of its uncertain affinity (Robbins 1987). The simulator predicts walking speeds of approximately 0.65–0.7 m s⁻¹ for trail G1, and 0.95–1.0 m s⁻¹ for trail G2/3. Since for our human subjects the simulator consistently underestimated speeds in comparison to speeds experimentally recorded for given stride lengths, it is quite possible that the *A. afarensis* simulation also underestimates speed for given stride lengths: actual speeds might have been still higher.

4. DISCUSSION

The predicted metabolic energy costs for the adult humans match the experimental data surprisingly well, although a perfect match has not yet been achieved. It has to be remembered that the predictions made by the

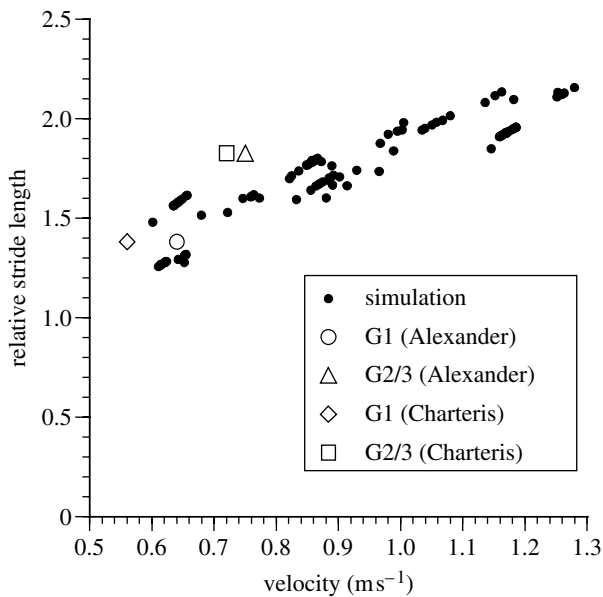


Figure 5. Open markers: estimated velocities plotted against Laetoli stride lengths for trails G1 and G2/3 as calculated by Alexander (1984) and Charteris *et al.* (1982). Solid circles: stride length and speed predictions from the *A. afarensis* simulator. The trend of the points, whilst largely in agreement with Alexander's G1 value clearly predicts higher velocities for the relative stride length of the G2/3 trails.

simulator do not exactly correspond to either the net or the gross cost of locomotion as measured experimentally. The experimentally determined gross cost of locomotion also includes energy costs not associated with the musculoskeletal system (such as by the liver, brain, etc.) and will therefore always be considerably greater than the costs estimated by the simulator. Calculation of net cost does correct for this while at rest. However, some of non-musculoskeletal costs are dependent on locomotor speed (those of the circulatory and respiratory systems for example) so that the correction becomes inaccurate at higher speeds. In addition the net cost of locomotion also subtracts the purely postural costs of the musculoskeletal system. Whilst this is mostly desirable since the model does not include the muscles acting in a primarily postural role, such as the axial musculature, some of the muscles it does include also have a postural cost at rest. There is currently no experimental way of estimating what the simulation actually predicts: the metabolic cost associated with the pattern of activation of the locomotor muscles alone.

In addition the model does not include many of the energy-saving features of the human locomotor system, in part because it is limited to two dimensions. There are no two-joint muscles to help energy transfer, and the spring mechanisms are limited to the tendons and the passive elasticity of the muscles. Storage of spring energy in ligaments is not included, and the choice of spring constant for the existing elements has not been optimized to maximize energy return. It is therefore not surprising that the actual metabolic costs of locomotion are somewhat lower than the values produced by the simulator. In addition the model is two-dimensional, so the costs and benefits inherent in the extra dimension

are ignored. These would become extremely important when investigating the value of particular anatomical structures and innovations since with a three-dimensional model we could, for example, investigate the effects of pelvis width independently of other morphological variation.

The predicted costs of locomotion: $7.0 \text{ J kg}^{-1} \text{ m}^{-1}$ ($2.0 \text{ J kg}^{-1} \text{ m}^{-1}$ net) for the lowest speeds, falling to $5.8 \text{ J kg}^{-1} \text{ m}^{-1}$ at 1.0 m s^{-1} ($2.9 \text{ J kg}^{-1} \text{ m}^{-1}$ net) and rising to $6.2 \text{ J kg}^{-1} \text{ m}^{-1}$ ($3.8 \text{ J kg}^{-1} \text{ m}^{-1}$ net) at the maximum walking speeds achieved, are consistent with those we have previously published for single speeds (Sellers *et al.* 2004) even though we are using a completely different muscle model in this simulation. They are also consistent with the metabolic cost predicted by the only fully three-dimensional simulation (Nagano *et al.* 2005): $7.7 \text{ J kg}^{-1} \text{ m}^{-1}$. Nagano and colleagues' value includes an estimate for non-locomotor metabolic costs, and so directly represents the gross metabolic costs of locomotion. However, it should be noted that their cost estimate is for a speed of only 0.615 m s^{-1} , at the bottom end of our range of speeds. Our gross cost is also in good agreement with experimental values for children (figure 4).

All other things being equal, our data suggest that 1.0 m s^{-1} was the optimal walking speed for *A. afarensis*. However, the optimal speed depends on the degree of time pressure on the animal—an animal with plenty of free time will actually use less energy over an extended time period by travelling at speeds which incur costs approaching the minimum of the net costs of locomotion (Sellers *et al.* 2004). Of course it is seldom the case that an animal is under no time constraint: higher speeds will often be adopted.

Our model predicts walking speeds for the two Laetoli trackways which were unequivocally made by human relatives, G1 and G2/3, of approximately 0.7 and 1.0 m s^{-1} respectively. These estimates substantially exceed the estimate of Charteris *et al.* (1982), based on relationships between stride length and speed in human adults and children (Grieve & Gear 1966): 0.56 m s^{-1} for trail G1, and 0.72 m s^{-1} for trail G2/3. They also exceed, to a lesser degree, Alexander's (1984) estimates, using dynamic scaling, of 0.64 m s^{-1} for G1 and 0.75 m s^{-1} for G2/3. G-2/3 consists of overlapping trails most likely made by two adults of *A. afarensis*. When compared to our predictions for the relationship of stride length, speed and metabolic costs, the predicted speed for G2/3 suggests that the makers were walking at, or near, their energetically optimum speed. Further, they were walking well within the range of predicted speeds for an animal of equivalent body size (cf. Heglund & Taylor 1988). Modern humans commonly adopt walking speeds between 1.0 and 1.7 m s^{-1} , depending on the situation (Bornstein & Bornstein 1976). Wirtz & Ries (1992) note, however, that young adults most commonly choose to walk at 1.5 m s^{-1} ; this is near the energetically optimum speed we recorded for our young adult human subjects. Our estimate for *A. afarensis* is nevertheless, although slow compared to a young adult human, within the range of absolute values for humans, despite a short stature of about 1.1 m . Thus, our model offers further evidence

that *A. afarensis* was a fully competent biped. This contrasts dramatically with Hunt's (1994) suggestion that *A. afarensis*' gait would have been 'bipedal shuffling'. It also necessarily casts doubt on his conclusion that *A. afarensis* was a 'postural biped' with a bipedal gait compromised by the competing needs of terrestrial bipedality and arboreal climbing.

Some caution must be expressed, since uncertainties with respect to dimensions of *A. afarensis* apply to the present study as much as to previous studies, and our predicted velocities must therefore be regarded as very approximate values. Nevertheless, any error may actually be in underestimation, as the model of our human subjects predicted velocities consistently lower than experimentally recorded.

Although chimpanzees may possibly not have particularly short strides compared to *Homo* or *Australopithecus* (Alexander 1984) it is extremely doubtful that in bipedal gait they could sustain a similar velocity to either genus. The single extant study of metabolic costs in bipedalism (Taylor & Rowntree 1973) shows an intercept and slope much greater even than that of humans imitating chimpanzee-like (BHBK) gait (Carey & Crompton 2005). Even the normal quadrupedal locomotion of chimpanzees is expensive compared that of other mammals (Taylor & Rowntree 1973). Assuming that chimpanzees tend to walk quadrupedally at their most efficient speed, not even male chimpanzees, which tend to walk at about 0.88 m s^{-1} (Pontzer & Wrangham 2004), match the optimum performance predicted for bipedalism of *A. afarensis*: an even more marked gap exists if we consider instead female chimpanzees (0.78 m s^{-1}).

Thus, within the limits of our model, and assuming that Taylor & Rowntree's (1973) data are reliable, the bipedal performance of *Australopithecus afarensis*, as predicted by our model is not only much closer to that of modern humans than to that of bipedally walking great apes, but at normal walking speeds, shows a clear speed/cost advantage over chimpanzee quadrupedalism. Climbing remains significantly more energetically expensive than terrestrial quadrupedalism for chimpanzees, despite their musculoskeletal adaptations (Pontzer & Wrangham 2004). Pontzer & Wrangham (2004) have shown that the costs of locomotion in chimpanzees are nevertheless dominated by terrestrial walking, because of very high daily travel distances, *versus* only limited use of climbing. If we make the major (and quite likely incorrect) assumption that the African apes existing at the time of *A. afarensis* were ecologically, morphologically and physiologically similar to modern common chimpanzees, then our data would tend to support Rodman & McHenry's (1980) argument that the adoption of bipedalism offered energetic advantages to early human ancestors.

However, the advantage to chimpanzees of increased capabilities in arboreal climbing must be taken into consideration, and chimpanzees can certainly achieve speeds in quadrupedal locomotion well in excess of the maximum speed achieved by our simulation of bipedalism in *A. afarensis*. Unfortunately, voluntary bipedalism is a rare event in other great apes, and the measures needed to assess its physiology usually

impossible if not unethical, so we are unlikely to be able to make a direct assessment of the physiological reasons for the gulf between bipedal performance of living apes, and that of humans, and, which is more to the point, their fossil relatives. A modelling approach similar to the present one is probably our best available tool.

Adoption of habitual, efficient bipedalism would be expected to exert an immediate influence on ranging behaviour and time budgets. Studies of modern human bipedalism and bipedalism in non-human primates can be used to quantify its consequences for time and energy budgets (for a detailed review see Schmitt 2003). But extension of these data to early human ancestors and their relatives is problematical. As we have seen the earliest unquestionable bipeds (e.g. *A. afarensis*) were little more than half the mass of modern human adults. Current approaches (e.g. Pontzer & Wrangham 2004) therefore use allometric scaling equations to assign costs to locomotor activities, but even these are subject to errors relating to differences in body proportions which outside genus *Homo* may be quite large. However, if the nature and scale of these differences could be understood, simulation approaches using agent-based modelling, a technique from computer science which focuses on behaviour of the individual, and variation within a population, as agents move through simulated environments are very promising future possibilities.

5. CONCLUSION

Assuming that the early human relative *A. afarensis* was the maker of the Laetoli footprint trails G1 and G2/3, our study suggests that, by 3.5 million years ago, some at least of our early relatives, despite small stature, could sustain efficient bipedal walking at absolute speeds within the range shown by modern humans.

The predicted velocity of walking of the adults which made Trail G2/3, 1.0 m s^{-1} , is close to the predicted energetic optimum for this species, but less than the energetic optimum for our human subjects (*ca.* 1.5 m s^{-1}). It is also less than the maximum velocity achieved by our *A. afarensis* simulations (*ca.* 1.2 m s^{-1}). It is however greater than the normal quadrupedal walking speed of common chimpanzees, and almost certainly represents a lower cost of transport over the same distance.

An evolutionary robotics approach can thus provide a relatively complete assessment of walking behaviour of fossil species such as *A. afarensis*, and may suggest its possible advantages over gaits of living non-human primates. However, our model still overestimates costs, in part because its two-dimensional nature limits it to a relatively rigid 'compass gait', and tends to underestimate speed. Our present model does not encompass the detail of foot anatomy which would be necessary to investigate running and cannot accurately mimic the force production capabilities of the human forefoot. This may be why the model tends to switch to running at relatively slow speeds (1.3 m s^{-1} (AL 288-1 model) or 1.5 m s^{-1} (human model)—human subjects both in

our experiments and as found by others tend to prefer to run at speeds in excess of 2.0 m s^{-1} (e.g. Thorstensson & Robertsson 1987; Minetti *et al.* 1994).

Computational costs are still such that models such as the present one which can *sui generis* produce stable walking are still limited in sophistication with respect to detailed functional anatomical questions, while more biorealistic models (such as that of Nagano *et al.* 2005) have not achieved stability over multiple cycles, and cannot be easily used to predict vital gait parameters such as speed and stride length. The next generation of simulators, however, should be able to cope both with individual muscle recruitment and multiple bone/soft tissue interactions, and the elastic storage of energy, which will greatly aid our understanding of the evolution of the human locomotor system, particularly at or near the walk/run transition.

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