Muscle—spring dynamics in time-limited, elastic movements

M. V. Rosario1,†, G. P. Sutton2, S. N. Patek1 and G. S. Sawicki3

1 Department of Biology, Duke University, Durham, NC 27708, USA
2 School of Biological Sciences, University of Bristol, Bristol BS8 1TH, UK
3 Joint Department of Biomedical Engineering, North Carolina State University and University of North Carolina at Chapel Hill, Raleigh, NC 27514, USA

Muscle contractions that load in-series springs with slow speed over a long duration do maximal work and store the most elastic energy. However, time constraints, such as those experienced during escape and predation behaviours, may prevent animals from achieving maximal force capacity from their muscles during spring-loading. Here, we ask whether animals that have limited time for elastic energy storage operate with springs that are tuned to submaximal force production. To answer this question, we used a dynamic model of a muscle–spring system undergoing a fixed-end contraction, with parameters from a time-limited spring-loader (bullfrog: *Lithobates catesbeiana*) and a non-time-limited spring-loader (grasshopper: *Schistocerca gregaria*). We found that when muscles have less time to contract, stored elastic energy is maximized with lower spring stiffness (quantified as spring constant). The spring stiffness measured in bullfrog tendons permitted less elastic energy storage than was predicted by a modelled, maximal muscle contraction. However, when muscle contractions were modelled using biologically relevant loading times for bullfrog jumps (50 ms), tendon stiffness actually maximized elastic energy storage. In contrast, grasshoppers, which are not time limited, exhibited spring stiffness that maximized elastic energy storage when modelled with a maximal muscle contraction. These findings demonstrate the significance of evolutionary variation in tendon and apodeme properties to realistic jumping contexts as well as the importance of considering the effect of muscle dynamics and behavioural constraints on energy storage in muscle–spring systems.

1. Introduction

In most cases, muscle contractile force is transmitted to skeletal structures through elastic structures, inextricably coupling muscle and spring dynamics. Many animals use muscles to temporarily store energy in their springs, such as tendons, and the stored energy can be recovered later to help power movement. The time available for muscles to load in-series springs is important, because stored elastic energy is proportional to force, and muscle force declines with contraction velocity [1]; therefore, the force capacity, and consequently the energy storage capacity, of the system is limited by muscle velocity and activation dynamics. Some animals store elastic energy over long time periods prior to movement [2–4], whereas others use power amplification systems with time-limited storage phases [5–7].

Given that the time available for spring-loading varies across animals and movement types, the relationship between spring properties such as mechanical spring stiffness (defined as spring constant and referred to simply as ‘stiffness’ in this study) and muscle-loading dynamics may impact performance (figure 1). For example, in situations where rapid spring-loading is beneficial (e.g. escape jumps and predatory ambushes), organisms may not have enough time to fully load their springs before the onset of movement. Although these organisms are not generating maximal muscle force, it is possible that their muscle–spring properties maximize elastic energy storage for submaximal force production. Few...
studies have examined the evolutionary variation of spring properties \[8,9\], yet the diversity of elastic systems suggests a range of mechanical, functional and behavioural influences on their form and function.

Here, we test whether and how springs are tuned differently to permit maximal energy storage for time-limited, submaximal force production versus non-time-limited, maximal muscle contractions. We developed a dynamic muscle–spring simulation of a fixed-end contraction (figure 1) and used it to compare time-limited (bullfrog: *Lithobates catesbeiana*) and non-time limited (grasshopper: *Schistocerca gregaria*) jumping systems. Both frogs and grasshoppers require elastic elements to achieve their high-power jumping performance \[10–13\]. Bullfrogs exhibit time-limited jumps in which a fast response is necessary, whereas grasshoppers perform longer-term muscle contractions in advance of movement and thus are less impacted by time limitations. We used existing, published data from these muscle–spring systems \[10,14,15\] to simulate spring-loading over a range of allowable storage times. We addressed the following two questions. (i) How does variation in the time available for muscle contraction influence the amount of energy stored in springs with different stiffness? (ii) Do the values of spring stiffness of bullfrogs and grasshoppers maximize energy storage given the distinct loading regimes of their jumping behaviour?

2. Methods

We ran simulations of bullfrog and grasshopper muscle–spring systems with varying spring stiffness \(k_{\text{simulation}}\) and determined which \(k_{\text{simulation}}\) resulted in maximal energy storage \(k_{\text{maxE}}\). We focused on spring stiffness, because this single value determined the relationship between force and spring stretch. Additionally, because spring stiffness is a mechanical property, it allowed us to compare the mechanical behaviour of springs that are composed of different materials. We omitted the duration of muscle contraction using static models and included the duration of muscle contraction using dynamic models. After all simulations were run, we compared published results of spring stiffness from bullfrog tendons \(K_{\text{bullfrog}}\) and grasshopper apodeme-cuticular springs \(K_{\text{grasshopper}}\) with the results of the simulations. Below, we outline how the simulations predicted energy storage in muscle–spring systems as a function of \(k_{\text{simulation}}\).

Two factors, spring stretch \(\Delta x_s\) and spring stiffness \(k_{\text{simulation}}\), were required in order to calculate stored energy:

\[
\text{energy} = \frac{1}{2} k_{\text{simulation}} \Delta x_s^2.
\]

Determining \(\Delta x_s\) was complicated by the interaction between muscle and spring. For example, an increase in \(k_{\text{simulation}}\) suggested higher energy storage (equation (2.1)); but springs with higher values of \(k_{\text{simulation}}\) stretch less for a given muscle force. Consequently, it was possible to increase \(k_{\text{simulation}}\) such that the resulting decrease in \(\Delta x_s\) reduced stored energy. Therefore, to account for the interactions between muscle and spring, we developed the following muscle–spring model.

(a) Muscle–spring model

We simulated dynamics within muscle–spring systems by connecting, in series, a model of a muscle to a model of a spring (figure 1). Specifically, we connected a Hill-type muscle to a Hookean spring \[1,5,16\]. We kept constant the muscle properties across all simulations while varying the spring stiffness, \(k_{\text{simulation}}\). Muscle and spring models were mathematically connected and implemented in R \(v.\ 3.2.1\), Vienna, Austria). In the following
sections, we explain how different instances of the model were used to predict force and elastic energy storage over a range of contraction scenarios.

(i) Hill-type muscle model
We used a Hill-type muscle model to predict muscle force as a function of three factors: muscle length, muscle velocity and muscle activation [5]. The relative contributions to muscle force by these three functions are described by equations (2.2)–(2.4):

\[ F_{\text{length}}(\Delta x_m, L_0, a_L, b_L, s) = e^{-(v/(v_{\text{max}}))^b} \]

(2.2)

\[ F_{\text{velocity}}(v, a_v, b_v) = \frac{1 - \left(\frac{v}{v_{\text{max}}}\right)}{1 + \left(\frac{v}{v_{\text{max}}/4}\right)} \]

(2.3)

\[ F_{\text{activation}}(t_{\text{contraction}}, r_{\text{act}}) = \begin{cases} r_{\text{act}} & t_{\text{contraction}} \cdot r_{\text{act}} < 1 \\ \frac{r_{\text{act}}}{1} & t_{\text{contraction}} \geq 1 \end{cases} \]

(2.4)

where \( x_m \) is the length of the muscle with units of muscle lengths; \( a_v, b_v, a_L, b_L \) and \( s \) are phenomenological parameters that were fitted to describe the shape of the muscle’s length–tension curve; \( v \) is muscle shortening velocity (\( \Delta x_m/\Delta t \)); \( v_{\text{max}} \) is the maximum shortening velocity of muscle contraction; \( t_{\text{contraction}} \) is the time the muscle has been contracting; and \( r_{\text{act}} \) is the linear rate of muscle activation. We used each of these three functions in the Hill-type muscle model to scale maximum force production; therefore, these functions were evaluated from 0 to 1 and represented the fraction of maximum force that was produced by a single component (i.e. length, velocity or activation) independent of all others (figure 2).

Each of the factors impacting muscle force production were combined to estimate muscle force \( F_{\text{muscle}} \) by multiplying the results of equations (2.2)–(2.4) with each other and the maximum tetanic force of the muscle \( F_{\text{max}} \).

\[ F_{\text{muscle}} = F_{\text{max}} \cdot F_{\text{length}} \cdot F_{\text{velocity}} \cdot F_{\text{activation}}. \]

(2.5)

In this model, maximum tetanic force was generated when each of the constituent effects on muscle force \( F_{\text{length}}, F_{\text{velocity}} \) and \( F_{\text{activation}} \) equaled 1.

(ii) Hookean spring model
We represented the series elastic component of the muscle–spring model with a linear, Hookean spring. Although biological springs are not Hookean, many springs, including those of bullfrogs and grasshoppers, approximate linear behaviour over a significant region of the force–displacement curve [10,17]. For this model, spring force was determined only by the displacement through which it is stretched \( (\Delta x_s) \) and the spring stiffness \( (k_{\text{simulation}}) \).

\[ F_{\text{spring}} = -k_{\text{simulation}} \Delta x_s. \]

(2.6)

(iii) Static muscle–spring model
We allowed the muscle and spring models to interact by setting two groups of variables equal: \( F_{\text{muscle}} \) equalled \( F_{\text{spring}} \), and the muscle shortening length change equaled the negative of spring stretch length change. This reduced muscle force (figure 2). We further simplified equation (2.7) to represent a static, steady-state solution by setting the dynamic components \( F_{\text{activation}} \) and \( F_{\text{velocity}} \) to 1:

\[ F_{\text{max}} \cdot F_{\text{length}}(\Delta x_m, L_0, a_L, b_L, s) \cdot F_{\text{velocity}}(v, v_{\text{max}}) \cdot F_{\text{activation}}(t_{\text{contraction}}, r_{\text{act}}) = -k_{\text{simulation}} \Delta x_s. \]

(2.7)

To simplify the model, variables that described muscle properties (i.e. variables that were only used to determine the shape of the Hill-type muscle components) were held constant for a given muscle. We used simplified equation (2.7) to represent a static, steady-state solution by setting the dynamic components \( F_{\text{activation}} \) and \( F_{\text{velocity}} \) to 1:

\[ F_{\text{max}} \cdot F_{\text{length}}(\Delta x_m, L_0, a_L, b_L, s) = -k_{\text{simulation}} \Delta x_s. \]

(2.8)

with \( \Delta x = \Delta x_m = -\Delta x_s \).

Solving for \( \Delta x \) in equation (2.8) resulted in the maximum internal stretch of that particular spring by its in-series muscle. This value was used to calculate maximum stored elastic energy in the static simulations, the case in which contraction time to store spring energy is not limited (see figure 1 for schematic).

(iv) Dynamic muscle–spring model
The dynamic muscle–spring model was identical to the static model with one exception: we did not set \( F_{\text{activation}} \) and \( F_{\text{velocity}} \) equal to 1 in equation (2.7). Holding all muscle properties constant and considering velocity as \( \Delta x_m \), and \( \Delta t \) resulted in the dynamic model

\[ F_{\text{max}} \cdot F_{\text{length}}(\Delta x_m, L_0, a_L, b_L, s) \cdot F_{\text{velocity}}(\Delta x, \Delta t) \cdot F_{\text{activation}}(t_{\text{contraction}}, r_{\text{act}}) = -k_{\text{simulation}} \Delta x_s. \]

(2.9)

Solving for \( \Delta x \) at each time step was complicated by the relationship between muscle length and contraction velocity, because \( \Delta x_m \) affected muscle force in two ways. First, \( \Delta x_m \) affected \( F_{\text{length}} \) directly; values of \( \Delta x_m/L_0 \) smaller than 1 (as a result of muscle shortening contraction) decreased muscle force (figure 2). Second, for a given \( \Delta t \), greater values of \( \Delta x_m \) resulted in greater contraction velocities. This reduced muscle force.
The amounts of energy stored with springs (15.37 N mm$^{-1}$) by rearranging the standard equation for two springs in series, which resulted in the equation

$$K_{\text{grasshopper}} = \frac{K_{\text{apodeme}} \cdot K_{\text{SLP}}}{K_{\text{apodeme}} + K_{\text{SLP}}}, \quad (2.10)$$

where $K_{\text{apodeme}}$ and $K_{\text{SLP}}$ are the stiffness values of the apodeme (31.4 N mm$^{-1}$) and semilunar process (30 N mm$^{-1}$), respectively [10]. The values of $k_{\text{simulation}}$ were uniformly generated from 5 to 350 N L$^{-1}$ increments of 0.1 N L$^{-1}$.

**Simulation parameters.** We simulated all muscle contractions with time steps of 0.001 s. The total number of steps was not determined before simulation. Instead, simulations terminated when change in muscle length reduced to less than 0.001 L$_0$ between time steps.

**Identification of $k_{\text{maxE}}$.** The determination of the spring stiffness that permitted maximal energy storage in the static simulations was straightforward. The value of $\Delta x$ in equation (2.8) was solved for many values of $k_{\text{simulation}}$. The stored energy for each simulation was calculated using equation (2.1). The value of $k_{\text{simulation}}$ that resulted in the greatest stored energy was recorded as $k_{\text{maxE}}$.

Obtaining $k_{\text{maxE}}$ from the dynamic muscle–spring model followed a similar process; however, the data required an additional pre-processing step. For each time step, $\Delta x$ was calculated for various values of $k_{\text{simulation}}$ via equation (2.9).

To test the effect of muscle contraction duration ($t_{\text{contraction}}$), we ran simulations with truncated duration to exclude time steps that were greater than $t_{\text{contraction}}$. From the truncated data-set, $k_{\text{maxE}}$ was determined using the methods above for the static and dynamic muscle–spring models.

### 3. Results

#### (a) Static simulation

For both the bullfrog and the grasshopper, the amount of stored elastic energy was maximized for an intermediate spring stiffness (figure 4). As $k_{\text{simulation}}$ increased, stored energy rose, levelled off and declined. In the bullfrog, the spring stiffness that permitted maximal energy storage ($k_{\text{maxE}}$ dotted lines in figure 4) equalled 20.98 N mm$^{-1}$, more than double the measured value of bullfrog tendon. The amounts of energy stored with $k_{\text{maxE}}$ and $K_{\text{apodeme}}$ were 20.43 and 14.13 mJ, respectively (table 2). In the grasshopper, $k_{\text{maxE}}$ equalled 18.0 N mm$^{-1}$ and the amounts of energy...
Figure 3. Regardless of activation rate, the spring stiffness that permits maximal energy storage ($k_{\text{maxE}}$) is dependent on the duration of muscle contraction ($t_{\text{contraction}}$). For example, $k_{\text{maxE}}$ at 300 ms (approximating the static, steady state) is higher than $k_{\text{maxE}}$ measured at 50 and 100 ms. For fast activations, $k_{\text{maxE}}$ is more sensitive at smaller durations of muscle contraction, demonstrated by the steep slope. The fast, intermediate and slow activations reach maximum activation within 100, 200 and 300 ms, respectively. Data shown are from the bullfrog model.

stored with $k_{\text{grasshopper}}$ were 2.24 and 2.21 mJ, respectively (table 2).

(b) Dynamic simulations

As $t_{\text{contraction}}$ increased, more elastic energy was able to be stored. For example, maximal elastic energy values stored at 50, 100 and 300 ms were 5.09, 14.46 and 20.43 mJ, respectively, for the bullfrog and 0.03, 0.14 and 1.08 mJ for the grasshopper (table 2). In addition, all values resulting from the 5000 ms dynamic simulation matched those of the static simulation; therefore, because we reached the static steady-state solution by 5000 ms, we did not simulate muscle contraction past this time step.

Similar to the static simulation, dynamic simulations also revealed that an intermediate spring stiffness resulted in maximal stored energy; however, the value of $k_{\text{maxE}}$ was dependent on $t_{\text{contraction}}$ (figures 3 and 4). Our simulation of the bullfrog muscle–spring system also showed that $k_{\text{maxE}}$ was higher for faster rates of contraction (figure 3); therefore, as a point of comparison between the bullfrog and the grasshopper, unless otherwise stated, all reported results were taken from simulations at the highest $t_{\text{act}}$ tested, resulting in tetanic contractions occurring in 100 ms.

In the bullfrog, $k_{\text{maxE}}$ for a realistically time-limited contraction (50 ms) was 7.14 N mm$^{-1}$, less than half that predicted by the static solution (20.98 N mm$^{-1}$). This difference was a direct consequence of the force–velocity property of the frog muscle. Additionally, $K_{\text{bullfrog}}$ (6.69 N mm$^{-1}$) was much closer to $k_{\text{maxE}}$ at 50 ms (7.14 N mm$^{-1}$) than to the $k_{\text{maxE}}$ of the static simulation (20.98 N mm$^{-1}$). Alternatively, $k_{\text{maxE}}$ in the grasshopper for a biologically relevant contraction duration (300 ms) was 12.0 N mm$^{-1}$, which matched the result predicted by the static solution (table 2). Regardless of simulation, as $t_{\text{duration}}$ increased, so did $k_{\text{maxE}}$ until the solution generated by the static solution was reached. This was shown by the rightward shift of the dotted line in figure 4 as time increased. When considering the highest value of $t_{\text{act}}$ used in the simulations, $k_{\text{maxE}}$ did not level out until 150 ms (figure 3). Additionally, the peak values of energy storage all occurred at the highest rates of activation (see the electronic supplementary material).

4. Discussion

By simulating the dynamic interaction between muscle and spring during a fixed-end contraction, we asked two questions: (i) Does reducing the time available for spring-loading affect which springs store the most energy? (ii) Do the values of spring stiffness in bullfrogs and grasshoppers permit maximum energy storage based on their contrasting loading regimes? For both the bullfrog and the grasshopper, the time available for muscle contraction determined which spring stiffness permitted maximal energy storage. As time restriction increased (i.e. as less time was available for muscle contraction), the values of spring stiffness that permitted maximal stored energy decreased (figure 4). Although the greatest amounts of elastic energy were predicted using the static solution, this solution was not reached until 5000 ms in the grasshopper, a duration of muscle contraction that is much greater than what occurred in other experiments (table 2). Consequently, static simulations may be insufficient to model muscle–spring systems in some cases. The static solution, however, offered an upper bound of $k_{\text{maxE}}$ and maximum stored energy in biological systems.

In both the bullfrog and the grasshopper, empirically measured values of spring stiffness approximately matched $k_{\text{maxE}}$ when taking time-limited loading into account. In the bullfrog, dynamic simulation revealed that when the duration of muscle contraction was restricted to biologically relevant contraction durations (50 ms), $k_{\text{maxE}}$ and $K_{\text{bullfrog}}$ were similar (7.14 and 6.69 N mm$^{-1}$, respectively). Therefore, the incorporation of muscle dynamics into the simulation not only allowed the muscle–spring model to behave in a more realistic way, but it also countered the results of the static simulation and suggested that bullfrog tendons maximize energy storage at short time scales. Conversely, results from the dynamic simulation of the grasshopper muscle–spring system suggested that the grasshopper spring system maximized energy at relatively long time scales. In the case of grasshoppers, which load their springs with longer durations than bullfrogs (300 and 50 ms, respectively), the static simulation provided reasonable estimates of $k_{\text{maxE}}$ and maximal stored energy. It is important to note that biological springs can be tuned over evolutionary time to perform a multitude of mechanical behaviours over a wide range of loading regimes; therefore, there may be functional reasons that explain mismatch between our predictions of optimal stiffness and the stiffness with which organisms operate. Regardless, these two cases of dynamic fixed-end contractions demonstrate that muscle–spring system performance depends on the interaction between storage time available and muscle–spring properties.

The dynamic simulation of the bullfrog also demonstrated the importance of dynamics for all rates of muscle activation. At the fastest muscle activations, $k_{\text{maxE}}$ did not level out until 150 ms (figure 3); therefore, bullfrog muscle–spring systems that complete energy storage within 150 ms are more sensitive to muscle dynamics than those that do not. Given that maximal in vitro activation of bullfrog muscle is reached in 100 ms [15], this further demonstrated the importance of muscle dynamics in bullfrog spring systems.
The duration of muscle contraction \( t_{\text{contraction}} \) determines whether the spring stiffness of bullfrog tendon \( K_{\text{bullfrog}} \) (indicated by arrow) permits maximal energy storage. For example, in the static simulation, \( K_{\text{bullfrog}} \) does not coincide with the peak of the curve (indicated by dotted line). Results from the static simulation suggest that \( K_{\text{bullfrog}} \) does not permit maximal energy storage. Conversely, during time-limited muscle contractions (50 ms), \( K_{\text{bullfrog}} \) is closer to the peak of the curve. The leftward shift of the peak as \( t_{\text{contraction}} \) is reduced suggests that muscle–spring dynamics become increasingly important with shorter durations of muscle contraction. Conversely, in grasshoppers, the static solution is a close approximation of the results from the biologically relevant loading time (300 ms). Results from simulations that occur at biologically relevant loading times are boxed. Note that the scale of the y-axis is different in each panel.

Table 2. As the duration of muscle contraction approaches biologically relevant durations (italicized values), \( k_{\text{max}} \) approaches the measured spring stiffness \((K_{\text{bullfrog}} \text{ and } K_{\text{grasshopper}})\). Static simulations accurately model systems that exhibit relatively long loading times, such as the grasshopper (grey-shaded values). Dynamic simulations are necessary for systems that exhibit time-limited contraction, such as the bullfrog (unshaded).

<table>
<thead>
<tr>
<th>( t_{\text{contraction}} ) (ms)</th>
<th>( k_{\text{max}} ) (N mm(^{-1}))</th>
<th>( K_{\text{bullfrog}} ) (N mm(^{-1}))</th>
<th>( E_{\text{max}} ) (mJ)</th>
<th>( E_{\text{bullfrog}} ) (mJ)</th>
<th>( k_{\text{max}} ) (N mm(^{-1}))</th>
<th>( K_{\text{grasshopper}} ) (N mm(^{-1}))</th>
<th>( E_{\text{max}} ) (mJ)</th>
<th>( E_{\text{grasshopper}} ) (mJ)</th>
</tr>
</thead>
<tbody>
<tr>
<td>50</td>
<td>7.14</td>
<td>6.69</td>
<td>5.09</td>
<td>5.09</td>
<td>6.50</td>
<td>15.37</td>
<td>0.04</td>
<td>0.03</td>
</tr>
<tr>
<td>100</td>
<td>15.17</td>
<td>6.69</td>
<td>14.46</td>
<td>12.19</td>
<td>6.75</td>
<td>15.37</td>
<td>0.14</td>
<td>0.13</td>
</tr>
<tr>
<td>200</td>
<td>20.98</td>
<td>6.69</td>
<td>20.25</td>
<td>14.12</td>
<td>8.75</td>
<td>15.37</td>
<td>0.53</td>
<td>0.51</td>
</tr>
<tr>
<td>300</td>
<td>20.98</td>
<td>6.69</td>
<td>20.43</td>
<td>14.13</td>
<td>12.00</td>
<td>15.37</td>
<td>1.08</td>
<td>1.07</td>
</tr>
<tr>
<td>5000</td>
<td>20.98</td>
<td>6.69</td>
<td>20.43</td>
<td>14.13</td>
<td>18.00</td>
<td>15.37</td>
<td>2.24</td>
<td>2.21</td>
</tr>
<tr>
<td>static</td>
<td>20.98</td>
<td>6.69</td>
<td>20.43</td>
<td>14.13</td>
<td>18.00</td>
<td>15.37</td>
<td>2.24</td>
<td>2.21</td>
</tr>
</tbody>
</table>

The results of the simulations hinted at the relationship between compliant springs and energy storage. As muscle contraction duration decreases, the total amount of elastic energy that can be stored decreased (figure 4). Additionally, the sensitivity of \( k_{\text{max}} \) to muscle dynamics increased as the duration of muscle contraction decreased (figure 3). Therefore, in muscle–spring systems that are time-limited, reducing spring stiffness could help maximize energy in situations in which stored energy is decreased owing to short contraction durations. In short, when muscle dynamics became important, optimal spring stiffness decreased.

Given that the results from the simulation were generated by connecting a Hill-type muscle model to a Hookean spring model, it is important to note the limitations of these constituent models in the context of this study. The Hill muscle model has been shown to accurately represent general trends in the relationship between the dynamics of muscle activation and force production [19–21]. This relationship, however, was highly dependent on activation dynamics [22,23] and may not have been accurately represented in this study. Instead of focusing on the intricacies of neuronal firing, we simplified muscle activation as a linear ramp to test, in general, whether muscle activation rate affected time-limited energy storage. To that effect, the model demonstrated that muscle dynamics played a part in determining which spring stiffness permitted maximal energy storage.

Additionally, our muscle–spring model does not incorporate inertial effects of muscle mass on contraction velocity [24] or activation-dependent shifts in the muscle length–tension relationship [25]. As a first approximation of how these effects may affect our interpretation of our results, we conducted a sensitivity analysis of our model by perturbing each parameter \( \pm 20\% \) (including \( V_{\text{max}} \) and starting length to represent inertial and activation effects, respectively). We found that while increases in \( V_{\text{max}} \) led to increases in \( E_{\text{max}} \), predictions of \( k_{\text{max}} \) were relatively insensitive (see the electronic supplementary material); therefore, inertial effects had little effect on our predictions of optimal
sensitive to variation in muscle activation rate, reported values of muscle stiffness. Conversely, the most sensitive parameter in the shape of the length–tension curve in regions that span the full excursion that muscles experience during fixed-end contractions. Data regarding activation-dependent length–tension curves should be incorporated in future simulations as they become available.

The results assumed that muscle contracts at rates such that maximal activation occurs within 100 ms. In reality, different jumps from the same animal could vary in muscle activation rate, thereby affecting the amount of spring stretch. The simulations show that bullfrog muscles that took longer than 100 ms to reach maximal activation stored less elastic energy for $k_{\text{maxE}}$ (figure 3). Because the simulations were sensitive to variation in muscle activation rate, reported values of $E_{\text{max}}$ and $k_{\text{maxE}}$ should not be interpreted as exact predictions of optimal bullfrog performance. Nonetheless, these values do provide a qualitative view of how muscle and spring parameters interact during time-limited energy storage.

Another simplification of the model involved the use of a linear spring. Most biological structures, including bullfrog tendons, exhibit a toe region of low spring stiffness early in the force–displacement curve followed by a linear region of higher spring stiffness. Many studies remedy this by measuring spring stiffness in the linear region of the force–displacement curve. In addition, it is important to note that the simulation only predicted the amount of energy stored, but other factors such as mass, material properties and morphological lever systems directly impact the unloading of energy [26,27].

Our dynamic simulations revealed a phenomenon that potentially affects all spring systems that are transiently loaded by muscle. That is, muscles that cannot develop isometric force because of time restriction can achieve significant amounts of elastic energy storage when coupled with springs of lower stiffness than would be predicted in the static case. For example, because bullfrogs lack morphological latches, they are not able to load their springs with peak isometric force. Instead, the bullfrog uses a dynamic catch mechanism, which temporarily resists force via inertial loads and mechanical advantage about moving joints [28]. The dynamic catch is able to resist some muscle contraction to permit spring-loading, but not long enough for isometric contractions to develop. With the exception of salamanders and chameleons, which probably contain anatomical latches [29,30], and toads, which have been hypothesized to store energy via co-contraction of antagonistic muscle [31], it is likely that vertebrates are inherently subject to time-limited energy storage, and potentially benefit from springs less stiff than expected.

Conversely, we predict that some invertebrate systems with anatomical latches may operate with relatively higher spring stiffness that can permit maximal energy storage over long storage times. Systems that have anatomical latches, and those that use rigid connections of body parts to resist muscle contraction, can develop isometric contractions during spring-loading. For example, snapping shrimp (Alpheus californiensis), trap-jaw ants and frog-hoppers (Phillea spumarius) contain body parts that lock together to form a latch and have springs that are connected to slow, forceful muscles that contract isometrically for up to several seconds [2–4,32–34]. Given the amount of power amplification observed in these systems, it is likely that these muscle–spring systems are operating with spring stiffness that permits maximal energy storage.

In some systems, the determination of an optimal spring stiffness can be complicated by an active latch, in which an antagonist muscle contracts to keep a system latched. Active latches may permit variation in the amount of stored energy prior to movement [35–37]. For example, the bush cricket can use changes in both joint angle and activation of the latching muscle to determine how much force holds the latch in place [36]. Meanwhile, a larger muscle can load the spring until it exceeds the force of the latch, thereby initiating movement. Given that the bush cricket can control the amount of energy stored, it is possible that it operates with a spring stiffness that results in the most stored energy for a wide range of situations. Although this idea is speculative, this study provides the tools necessary to test this hypothesis in other active latch systems in future work.

5. Conclusion

When testing for maximal energy storage, it is important to consider the dynamic interaction of muscle and spring. Our simulations revealed that within the realm of biologically relevant time scales, the more time available for loading by muscle, the stiffer the series spring required for maximum elastic energy storage. Muscles that load in-series springs over shorter time scales benefit from less stiff springs. At short time scales, muscle force is small owing to low activation and high velocity, and less stiff springs allow the spring to stretch more for a given amount of force. Thus, it is necessary to determine the effect, if any, of muscle dynamics on energy storage before concluding whether or not muscle–spring systems maximize energy storage.

Ethics. All animal information was based on previously published datasets.

Data accessibility. All R code used for simulations can be found at http://dx.doi.org/10.17605/OSF.IO/Z385A. Data generated from the simulation have been uploaded to Dryad (http://dx.doi.org/10.5061/dryad.089cr).

Authors’ contributions. M.V.R. implemented the mathematical model, ran all dynamic simulations, analysed the data and prepared the manuscript. G.P.S. conceived the original idea of the static model and guided the study to include grasshoppers, S.N.P. helped prepare the manuscript and identified key points/questions of the study. G.S.S. helped develop the dynamic model and simulations, provided data from bullfrogs, offered insight on data analysis/interpretation and helped edit the manuscript.

Competing interests. We have no competing interests.

Funding. This research was supported by grants awarded to M.V.R. (DOE FG02-97ER25308) and S.N.P. (NSF IOS-1439850).

Acknowledgements. The authors thank W. M. Kier, K. K. Smith, D. M. Boyer and E. Azizi for providing feedback on the manuscript. We also appreciate comments and contributions from P. Green, P. S. L. Anderson, K. Kagaya, R. Crane. We thank the DOE CSGF for training provided for high-speed computing.
References