



Modeling muscle function using experimentally determined subject-specific muscle properties

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19 Abstract

20

21 Muscle models are commonly based on intrinsic properties pooled across a number of 22 individuals, often from a different species, and rarely validated against directly measured muscle forces. Here we use a rich data set of rat medial gastrocnemius muscle forces recorded during in-23 24 situ and in-vivo isometric, isotonic, and cyclic contractions to test the accuracy of forces 25 predicted using Hill-type muscle models. We identified force-length and force-velocity 26 parameters for each individual, and used either these subject-specific intrinsic properties, or 27 population-averaged properties within the models. The modeled forces for cyclic in-vivo and insitu contractions matched with measured muscle-tendon forces with r^2 between 0.70 and 0.86, 28 29 and root-mean square errors (RMSE) of 0.10 to 0.13 (values normalized to the maximum isometric force). The modeled forces were least accurate at the highest movement and cycle 30 frequencies and did not show an improvement in r^2 when subject-specific intrinsic properties 31 32 were used; however, there was a reduction in the RMSE with fewer predictions having higher errors. We additionally recorded and tested muscle models specific to proximal and distal 33 regions of the muscle and compared them to measures and models from the whole muscle belly: 34 35 there was no improvement in model performance when using data from specific anatomical regions. These results show that Hill-type muscle models can yield very good performance for 36 37 cyclic contractions typical of locomotion, with small reductions in errors when subject-specific 38 intrinsic properties are used.

39

41 Introduction

42 Ouantifying muscle forces is necessary for understanding how movements are powered 43 and controlled, yet muscle force is challenging to measure directly during voluntary movements. Some studies have recorded forces from common tendons that originate from multiple muscles 44 (Komi, 1987; Gregor et al. 1987; Hoffer et al. 1987; Biewener and Blickhan, 1988; Finni et al. 45 46 2000; Daley and Biewener, 2003); fewer studies have recorded forces directly from distinct 47 tendons (Biewener et al., 1998; Eng et al., 2019; Herzog et al., 1993; Roberts et al., 1997; 48 Walmsley et al., 1978). However, tendon and thus muscle forces for the majority of species, 49 muscles and movements have not yet been recorded; and this is particularly the case for in humans. When muscle forces cannot be directly measured, they are often estimated using 50 computational models (Zajac, 1989). One such model is the Hill-type muscle model that has seen 51 52 widespread use in more recent years for both in-situ (James et al. 1996; Sandercock and 53 Heckman, 1997; Wakeling and Johnson, 1999; Perreault et al. 2003; Wakeling et al. 2012; 54 Millard et al. 2013) and *in-vivo* studies (Hodson-Tole and Wakeling, 2009; Gerus et al. 2012; Lee et al. 2013; Dick et al. 2017). Hill-type muscle models are now embedded in many 55 musculoskeletal simulations of movement to understand muscle use during human movements in 56 57 health and disease (for review: Seth et al. 2018). Forces predicted by Hill-type muscle models are sensitive to the intrinsic muscle 58 59 properties used in the model. Muscles show individual variation in pennation angle (Wickiewicz 60 et al. 1983), muscle quality (isometric stress: Medler, 2002), passive stiffness (Azizi, 2014), intrinsic contraction speed (Sadoyama et al., 1988; Medler 2002), and excitability; with this 61

62 variation often being caused by ageing (Lauretani et al. 2003), disuse (Narici and de Boer, 2011;

63 Wisdom et al. 2014) and disease (Smith et al. 2019). Previous studies have shown how subject-

specific optimal lengths for fascicles and tendons can improve the accuracy of predicting forces 64 from muscle models (Li et al. 2009; Gerus et al. 2012). However, it is not known how subject-65 66 specific fascicle force-length and force-velocity relations would additionally affect the predicted forces, due to the difficulty in measuring these parameters for model validation. In this study we 67 investigate whether the forces predicted by Hill-type muscle models are improved when the 68 69 models use these subject-specific intrinsic properties. We test these models on *in-vivo* and *in-situ* 70 data obtained from a rat ankle plantarflexor muscle (medial gastrocnemius, MG), a muscle that 71 allows direct measurements of the muscle force from the MG tendon. We test these predictions 72 for contractions over a range of complexity: from *in-situ* isotonic, steady monotonic and cyclic work-loop contractions to unrestrained in-vivo locomotion. 73

74 Hill-type muscle models are commonly implemented with a single fascicle length and pennation that is assumed to represent the whole muscle that it is emulating (Zajac, 1989). 75 76 However, skeletal muscles have heterogeneous architecture, and show regional variations in 77 pennation angle, fascicle length, strain, fibre-type properties, and activation (eg. De Ruiter et al 1995, Ahn et al., 2018; Azizi & Deslauriers 2014; Konow et al., 2010). Previously we have 78 shown that Hill-type muscle models that have multiple contractile elements with different 79 80 intrinsic properties (but the same architecture) predict more accurate forces at high velocity (Lee 81 et al. 2013) and predict reduced metabolic costs of contraction (Lai et al. 2018). It is possible that 82 model refinements that consider the regional differences in architecture may additionally 83 improve the predicted forces. However, this has not been tested against directly measured force. 84 In this study we measure *in-vivo* excitations and length changes in the proximal and distal 85 regions of the rat MG, and test whether the forces predicted from different regions of the muscle 86 would differ, whether they differ from the forces predicted from measures of the whole muscle

87 belly, and how well these predictions match experimentally measured forces on the tendon

88 during *in-situ* and *in-vivo* contractions.

| 89 | We tested the following hypotheses: |
|----|--|
| 90 | 1) That model-predictions of muscle forces will be worse at the highest contractile |
| 91 | frequencies but not the greatest magnitude of force; |
| 92 | 2) That modeled muscle forces based on subject-specific muscle properties will result in |
| 93 | significantly improved predictions of <i>in-vivo</i> force patterns compared with forces |
| 94 | modeled using muscle properties averaged across a population; and |
| 95 | 3) That region-specific fiber strain and EMG will result in significantly improved |
| 96 | predictions of <i>in-vivo</i> forces compared to measures from the whole muscle belly. |
| 97 | |
| 98 | |

99 Methods

100 Animals

All procedures were carried out under approval by the Harvard University (FAS)
Institutional Animal Care and Use Committee (protocol 20-09), according to USDA guidelines.
Seven male Sprague Dawley rats (body mass: 369±87 g; Table S1) were trained for three weeks,
using verbal encouragement and light gusts with an air-duster to locomote at three different gaits/
inclines on a motorized treadmill.

106

107 In vivo measurements

Each animal was anesthetized (isoflurane gas to effect; 1-2%, via mask). Two sets of 108 109 bipolar silver wire electromyography (EMG) electrodes and four 1-mm sonomicrometry 110 transducers were inserted into the medial gastrocnemius (MG), and a custom-fabricated 'leaf-111 spring' force transducer was attached to the MG tendon using sterile surgery procedures (Fig. 112 S1; see (Eng et al., 2019; Konow et al., 2020; Richards and Biewener, 2007). The pairs of sonomicrometry crystals were placed along proximal and distal fascicle axes (8-10mm apart, 113 114 spanning ~80-85% fascicle length). Misalignment of crystals never exceeded 5°, yielding errors 115 in fascicle strain due to misalignment <1%. The separation of the proximal and distal-most 116 crystals provided measurement of the length of the muscle belly that excluded the tendon. 117 Analgesics (Flunixin meglumine, 2 mg/kg) were administered twice over the 48 hours following 118 surgery.

119 After surgery recovery, recordings were obtained as animals moved at the steadiest 120 possible speed (walk ~ 0.3 m s^{-1} ; trot ~ 0.8 m s^{-1} ; gallop ~ 1.0 m s^{-1}) on the level and 14° upslope.

121 Electrode and transducer signals were sampled at 5 kHz (MP150 16-bit A/D converter, Biopac
122 Systems, Inc., Goleta, CA, USA).

123

124 In-situ measurements

After *in-vivo* experiments, animals were re-induced on isoflurane gas. A femur clamp 125 126 was secured to a stereotaxic frame and the foot was zip-tied to a plastic plate on the frame. The 127 MG and the sciatic nerve were exposed using blunt dissection. A cuff electrode (inter-pole 128 spacing of 5.0 mm) was secured around the sciatic nerve, and all sciatic branches distal to the 129 cuff electrode were severed, except for the one innervating MG (Tijs et al., 2014). The calcaneus was cut with the MG tendon intact and attached to the lever arm of a servomotor (Fig. S1A). The 130 131 muscle and exposed tissue were immersed in mineral oil pooled in a loosened skin flap and 132 regulated at 35+1°C.

Isometric force-length (FL) and isotonic force-velocity (FV) curves were constructed for 133 134 maximal (amplitude: 2-3 V) and submaximal (amplitude: ~ 1 V) stimulations to the nerve (1.5mA; frequency, 125Hz; pulse width, 0.2ms; train duration: 300ms) via a Grass S48 135 136 stimulator. Isotonic shortening contractions relied on servomotor force-control, with contractions 137 crossing the isometric optimal length (l_0) for both proximal and distal fascicles. As l_0 is affected by muscle activation (Holt and Azizi, 2016; Rack and Westbury, 1969; Rassier et al., 1999), we 138 139 first determined l_0 for each region independently. Three-minute rest periods between successive 140 muscle contractions were used to minimize muscle fatigue. Experimental data were only used for 141 analysis if tetanic force remained \geq 90% of the maximum isometric force F_0 during the *in-situ* 142 tests.

| 143 | In-situ forces from the MG were measured during cyclic work-loop contractions. The |
|-----|--|
| 144 | muscle length trajectories were set to match those during <i>in-vivo</i> upslope gallop strides of a |
| 145 | single rat using 3D fluoroscopy (Konow et al., 2020; peak amplitude of 1.2mm and cycle |
| 146 | frequency of 3.5Hz); with stimulus onset 60ms before peak MTU length and EMG duty cycle |
| 147 | 46%, as per in-vivo gallop data (Eng et al. 2019). This baseline work-loop condition was then |
| 148 | varied in cycle frequency (2 to 10Hz) and peak-to-peak amplitude (0.6 to 2.4mm), with some |
| 149 | conditions being low amplitude and high frequency, and others being high amplitude and low |
| 150 | frequency. All conditions were recorded for maximal and submaximal stimulations. |
| 151 | MG force and EMG signals were sampled at 4kHz; and muscle belly, proximal fascicle, |
| 152 | and distal fascicle lengths were sampled at 520Hz via the sonomicrometer. Animals were |
| 153 | euthanized after the experiment (pentobarbital sodium overdose, IP). The MG was excised post- |
| 154 | mortem, weighed and bisected in the sagittal plane to measure whole muscle belly length and |
| 155 | total length of the proximal and distal fascicles, as well as their resting pennation angle β_0 (Table |
| 156 | S1). These measurements were used to correct for the length of fascicles and muscle not spanned |
| 157 | by the sonomicrometry crystals. Lastly, tendon transducer output was calibrated to ergometer |
| 158 | force measured for a series of <i>in-situ</i> isometric tetanic contractions. |

159

160 *Modelling isometric and isotonic muscle force*

We took the approach of initially determining phenomological relations to describe the intrinsic force-length and force-velocity characteristics of the muscle. We then used these intrinsic properties, combined with measures of the time-varying excitation, length and velocity of the muscle during *in vivo* and *in situ* contractions to predict what the muscle forces would be during these contractions, and compared these predicted forces to independently measured

166 forces. We used models to predict forces from the muscle belly (using the length and velocity of 167 the muscle belly, and the mean EMG-intensity from both muscle regions), and from either the 168 proximal or distal region of the muscle (using fascicle length and velocity and EMG-intensity 169 from that region).

The peak active force from the FL tests gave the maximum isometric force F_0 and the optimal length l_0 for the proximal and distal fascicles and the muscle belly. The proximal fascicle, distal fascicle and muscle belly lengths were normalized \hat{l} to their respective optimal lengths for each rat.

174 The FL data were fit to an active $\hat{F}_a(\hat{l})$ and a passive $\hat{F}_p(\hat{l})$ model (normalized to F_0) 175 using least squares minimization:

176
$$\widehat{F}_a(\widehat{l}) = e^{-\left|\frac{|\widehat{l}|^w - 1}{s}\right|^r}$$
 (equation 1)

where *r* determines roundness, *w* determines skewness and *s* determines width of the forcelength relationship (Otten, 1985; Hodson-Tole and Wakeling, 2009), and

179
$$\hat{F}_p(\hat{l}) = e^{(c_1 + c_2 \hat{l})}$$
 (equation 2)

180 where c_1 and c_2 are empirical coefficients.

181 The FV data from the concentric $(\frac{d\hat{l}}{dt} < 0)$ isotonic contractions were fit to a piecewise

182 function $\hat{F}_{\nu}\left(\frac{d\hat{l}}{dt}\right)$, normalized to F_0 , using least squares minimization (Lee et al. 2013):

183 $\hat{F}_{\nu}\left(\frac{d\hat{l}}{dt}\right) = 0$ for $\frac{d\hat{l}}{dt} < -1$ (equation 3)

184
$$\hat{F}_{\nu}\left(\frac{d\hat{l}}{dt}\right) = \frac{1 + \left(\frac{d\hat{l}}{dt}\right)\frac{1}{\nu_{0}}}{1 - \left(\frac{d\hat{l}}{dt}\right)\frac{1}{k\nu_{0}}} \qquad \text{for} \qquad -1 \le \frac{d\hat{l}}{dt} < 0$$

185
$$\hat{F}_{\nu}\left(\frac{d\hat{l}}{dt}\right) = 1.5 - \frac{0.5\left[1 - \left(\frac{d\hat{l}}{dt}\right)\frac{1}{\nu_0}\right]}{1 + 7.56\left(\frac{d\hat{l}}{dt}\right)\frac{1}{k\nu_0}} \quad \text{for} \quad \frac{d\hat{l}}{dt} \ge 0$$

where k is the curvature of the force-velocity relationship, and v_0 is the maximum unloaded 186 contraction velocity. For the eccentric portion of the FV curve $(\frac{d\hat{l}}{dt} \ge 0)$ curve parameters were 187 taken from Otten (1987): note these parameters were not specific to the individual rats. 188 189 An intensity envelope was fit to the EMG signals using an EMG-specific wavelet 190 analysis (Lee et al. 2011). The muscle excitation was taken as the square-root of this EMGintensity and normalized to the maximum excitation that occurred across all conditions for each 191 recording channel. The excitation was converted to an activation signal using three coupled 192 193 differential equations as a transfer function (Lee et al. 2011), with rat-specific time-constants 194 calculated from MG twitches recorded in the rat: these were general time-constants fit to pooled data from both proximal and distal EMG-intensities. Muscle activations $\hat{a}(t)$ as a function of 195 time t were calculated separately for the EMG from the proximal and distal locations of the 196 197 muscle, and the mean of these activations was used to model the forces for the muscle belly. Muscle forces F_m for both the *in-situ* work-loop and the *in-vivo* locomotion data were 198 199 calculated as follows:

$$F_m = F_0 \left[\hat{a}(t) \, \hat{F}_a(\hat{l}) \hat{F}_v(\frac{d\hat{l}}{dt}) + \hat{F}_p(\hat{l}) \right] \hat{F}_\beta$$

We assumed that the thickness of the muscle belly was constant during the experiment (Lee et al. 2013) and calculated it from the post-mortem fiber length and pennation β_0 . The timevarying pennation angle β_i was calculated from the fibre length during contraction and belly thickness. Given that the FL properties were measured using the fiber-lengths, but these fibers were already at a pennation angle relative to the line-of-action of the muscle belly, the effect of the time-varying pennation angle β_i during contraction was factored in by \hat{F}_{β} , where:

207
$$\hat{F}_{\beta} = \frac{\cos(\beta_i)}{\cos(\beta_0)}$$

208 for the force predictions from the proximal and distal fascicles, and

209
$$\hat{F}_{\beta} = \frac{\cos(\beta_0)}{\cos(\beta_i)}$$

210 for the force predictions from the muscle belly length.

211 We separated the *in-situ* data into work-loop cycles, and *in-vivo* data by stride cycles

212 from continuous locomotion bouts that were \sim 5 seconds long.

213

214 Statistical analysis:

The predicted muscle forces were evaluated by their coefficient of determination (r^2) and root-mean square error (RMSE, relative to F_0) by comparison with experimentally measured tendon forces. Values in the text are reported as mean \pm standard deviation. All statistical tests used the General Linear Model function in Systat v.12 and involved an initial full-factorial design, factoring individual, modeling method (population based vs. subject-specific), and muscle region. Model-simplifications were carried out when factors were non-significant.

To test **hypothesis 1** we used a nested factorial design to hold either the "force" or "frequency" parameter constant, while testing across conditions for the alternate parameter. Force was dictated *in-vivo* by the treadmill slope condition (Konow et al., 2020), and *in-situ* by sub- or supra-maximal muscle stimulation. Frequency was dictated *in-vivo* by gait condition, and *in-situ* by varying work-loop frequency. Tests were run separately for the *in-situ* and *in-vivo* conditions, and within each condition, for the population average (All) and subject specific (Rat) methods of modeling.

- To test **hypothesis 2** we isolated the effect of the method factor to discriminate the effect of using subject-specific (Rat) as opposed to population averages (All) muscle activation and strain-trajectories when modeling muscle force.
- To test **hypothesis 3** we used post-hoc testing with pair-wise comparisons of r^2 and RMSE for the muscle's proximal and distal regions, and the whole muscle belly.
- 234

235 Results

236 Consistent active isometric FL curves were obtained across animals for proximal and 237 distal MG regions, as well as for the whole muscle (Fig. 1A-C). The active FL curve was 238 narrower for the muscle belly compared with the proximal and distal regions. The passive FL 239 curves, however, were more variable across animals (Fig. 1D-F), in large part due to variable 240 disruption of adjacent fascial compartments needed to isolate the MG to obtain in-situ FL and 241 FV measurements. For isotonic FV relationships (Fig. 1G-H), the normalized maximum shortening velocities were significantly different between regions ($F_{2,10}=14.88$; p=0.001) without 242 effect of individual (F_{5.10}=3.32; p=0.050). Maximum shortening velocities of the muscle belly 243 $(5.06\pm0.98 \text{ s}^{-1})$ were lower than for the proximal (10.82±3.01 s⁻¹; p<0.01) and distal (9.82±3.22 s⁻¹) 244 245 ¹; p < 0.05) region, while the difference between the proximal and the distal region was not 246 significant (p>0.05) These FL and FV relations were determined before, and independently of, 247 the evaluation of the Hill-type model force predictions of the cyclic in situ contractions and the 248 in vivo behaviours.

In general, the time-varying length and activation resulted in excellent model predictions of MG force for both the *in-vivo* locomotor conditions (Fig. 2A-C) and the *in-situ* work-loops (Fig. 2D). However, these model predictions were not as good as for the steady FL and FV data

252 (Fig. 3). When averaged across the proximal, distal and muscle belly rat-specific models, the mean r^2 for the model predictions was 0.996±0.004, 0.857±0.100 and 0.699±0.207 for the 253 254 steady, cyclic *in-situ*, and *in-vivo* conditions, respectively, and the mean RMSE for the model 255 predictions was 0.016 ± 0.013 , 0.101 ± 0.050 and 0.133 ± 0.063 and for the steady, cyclic *in-situ*, 256 and *in-vivo* conditions, respectively. The prediction-accuracy of forces became worse with each increase in *in-vivo* stride frequency (walk to gallop in Figs. 4 and S3) (GLM: F_{2.1179} = 107.99; 257 p<0.001) and *in-situ* work-loop frequencies (Figs. S2 and S3) (F_{5,693} = 35.98; p<0.001), but 258 generally exhibited similar or lower r^2 , and greater RMSE with increased *in-vivo* force as the 259 260 locomotor grade changed from level to upslope, and increased *in-situ* stimulation intensity. The quality of model-predictions of forces, as shown by their r^2 , did not substantially 261 262 change between models with subject-specific muscle parameters compared to the population 263 means (Fig. 5A). Indeed, the statistical tests for in-vivo showed no significant main effect of the model type (subject-specific or population-averaged parameters) on the r^2 values (F_{1.1175} = 0.01; 264 p>0.05), but there was a statistically significant decrease of the RMSE values ($F_{1,1179} = 12.98$; 265 p<0.001) for the predicted forces when using subject-specific data (Fig. S3. For *in-situ* work 266 loops, the subject-specific model yielded significantly higher r^2 (F_{1.689} = 57.62; p<0.001). In 267 268 general, we found that models using the population-averaged parameters predicted forces with 269 higher RMSE errors more often than models that used subject-specific muscle parameters (Fig. 270 5B).

The model predictions of *in-vivo* force did not improve when using region-specific fiber strain and EMG as compared to the data for the whole muscle belly (Fig. 3). In our populationaverage (All) modeling approach, averaging proximal and distal activation and using muscle belly length to model the muscle belly forces, in fact, yielded significantly higher r^2 than using

regional data collected from distal or proximal compartments ($F_{1,666} = 18.39$; p<0.001). The posthoc test remained significant for the individual-level comparison (p<0.001) despite significant individual variation across the regions. None of the three corresponding RMSE comparisons were statistically significant ($F_{1,666} = 0.007$; p>0.05).

279

280 Discussion

We show here that generally good, to excellent, fits of muscle force can be achieved by muscle models that incorporate subject-specific muscle properties, as well as when properties are pooled across individuals. Rat-specific MG models yielded fits with r^2 that ranged from 0.699– 0.857 for *in-vivo* locomotor and *in-situ* work-loop contractions to 0.996 for steady FL and FV contractions (Fig. 3), and the RMSE for these rat-specific muscle models ranged from 0.101– 0.132 during dynamic *in-vivo* and *in-situ* contractions, respectively, to as low as 0.016 for steady monotonic contractions (relative to F_{0} ; Fig. 3).

An important aspect of our modeling approach is that the models were not optimized 288 across all input parameters of the model to achieve a best fit. Instead, we relied on rat-specific 289 290 values of measured intrinsic muscle properties (active and passive isometric FL, l_0 , isotonic FV) 291 and fascicle architecture, which were then driven by measured activations (from indwelling 292 EMG) of the muscle during both steady and dynamic *in-vivo* and *in-situ* contractions. Using this 293 approach, we achieved the best predictive fits from Hill-type muscle models to measured muscle 294 forces that have been reported to date. In comparison, prior animal (cat (Perreault et al., 2003; Sandercock and Heckman, 1997); goat (Lee et al., 2013) and human (Dick et al., 2017)) muscle 295 models have achieved fits against measured muscle forces, with r^2 ranging from 0.40–0.63 and 296 297 RMSE ranging from 0.10–0.23 (average ranges when they were reported relative to F_0). It

should be noted that the increases in the mean r^2 and decreases in mean RMSE were larger than their median values, due to the skewed nature of the distributions of r^2 and RMSE. Not only did the mean and median measures improve with the rat-specific intrinsic properties (Fig. 5), but there was a reduction in the skewness of the distributions, with a reduction in the number of cases of poor model fits when compared to models that used population-averaged intrinsic properties.

304 A large source of variability, and likely error, arising from our rat MG model was the 305 considerable experimental variation in l_0 and passive FL properties measured for the MG across 306 individual rats (Fig. 1D-F, Table 2). The resting length for elastic tissues, including tendons, is 307 often assumed to occur at muscle l_0 . Passive muscle stiffness strongly influences the time-308 varying force predicted by a muscle model in relation to activation and changes in muscle length 309 (Thelen, 2003; Li et al. 2009; Gerus et al. 2012, 2015). Nevertheless, despite the considerable 310 variation in resting length that we observed across individual muscle preparations, which likely 311 reflected the unavoidable disruption of fascial compartments to obtain *in-situ* measurements of 312 muscle properties (Tijs et al. 2020), the models predicted good fits (Fig. 2) for dynamic changes 313 in force during *in-situ* work-loop and *in-vivo* locomotion contractions.

The predicted muscle forces showed worse fits (*r*²) and greater errors (RMSE) for the higher cycle and stride frequencies, in a manner consistent with previous reports for cyclic contractions (James et al. 1996; Wakeling and Johnston 1999, Dick et al. 2017). This speedsensitivity may reflect the increasing importance of time-sensitive processes such as deactivation rates (Askew and Marsh, 1998) as cycle durations become shorter. Whilst errors appear greater for upslope versus level *in-vivo* gait, and maximal versus submaximal *in-situ* stimulations, it should be noted that these errors are all relative to the maximal isometric force: the errors

321 relative to the peak force during each contraction cycle were less for cycles that generated 322 greater force. The greater r^2 values for level *versus* upslope locomotion may reflect the more 323 complex patterns of muscle fascicle strain during upslope locomotion, where the muscle 324 experiences periods of both active lengthening and active shortening (Fig. 2; Eng et al. 2019; Konow et al., 2020). Active lengthening of a muscle likely incurs history-dependent effects that 325 326 are not captured by the steady FL and FV properties used in Hill-type models. When a muscle 327 develops force with prior active lengthening its force may be elevated compared to its isometric 328 force at the same length (Abbott and Aubert 1952; Cavagna and Citterio 1974; Edman et al. 329 1982; Herzog and Leonard 2002; Hisey et al. 2009), and force may be reduced when the muscle 330 reaches that length with prior active shortening (Abbott and Aubert 1952; Marechal and Plaghki 331 1979; Meijer et al. 1998; Herzog et al. 2000). These history-dependent effects may be included 332 in models of muscle force (Rode et al. 2009; Herzog et al. 2012; Herzog 2014; Nishikawa et al. 333 2012, McGowan et al. 2013; Ross et al. 2018); however, most current Hill-type models do not 334 consider contraction history and so there is still scope for increasing model accuracy. Whilst accounting for active lengthening is important for eccentric contractions, the amount of active 335 336 lengthening was small for these typical gaits in the rat and thus the scope for such model 337 improvement is limited in this case.

Our modelled MG forces did not improve when region-specific length and EMG data were used, rather than muscle belly length and mean EMG intensity across the muscle. This possibly reflects the complexity and heterogeneity of the architecture and recruitment within the rat MG. Additionally, the supramaximal stimulation of the sciatic nerve for the *in situ* contractions will have induced the same recruitment for the two regions, reducing the scope for region-specific differences in force. It would seem that the whole muscle output cannot

necessarily be predicted from one isolated region, but rather will emerge from the outputs andinteractions between the different muscle regions.

346 We were surprised to find, therefore, that modelled forces did not statistically improve when subject-specific intrinsic properties were used compared to the models with more generic 347 population-averaged parameters: this may partly reflect the very good model fits that were 348 349 already achieved with the generic models. However, it should be noted that the muscle models 350 with subject-specific models were more likely to achieve lower errors than the generic model. 351 Consequently, for clinical assessments there may be considerable merit in using patient-specific 352 parameters for muscle models, which may deviate more substantially from the distribution of 353 values across a healthy population.

354

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358

359 Declaration of Competing Interest

360 The authors declare that they have no known competing financial interests or personal

361 relationships that could have appeared to influence the work reported in this paper.

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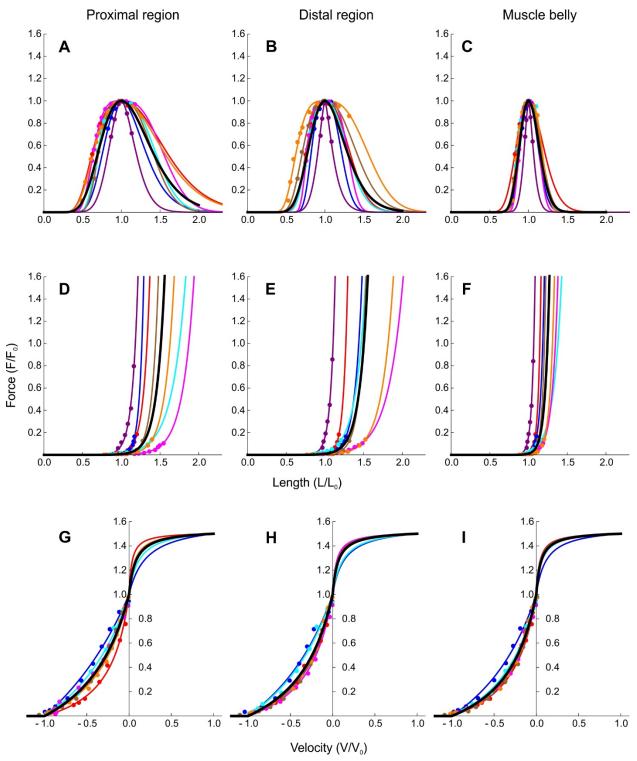
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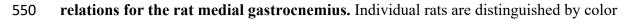
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549 Fig. 1. Isometric force-length (active A-C; passive (D-F), and isotonic force-velocity (G-I)



551 (experimental data: points and fitted models: lines). The pooled FL and FL curves (black lines)

552 were fit to all the data across the different individuals.

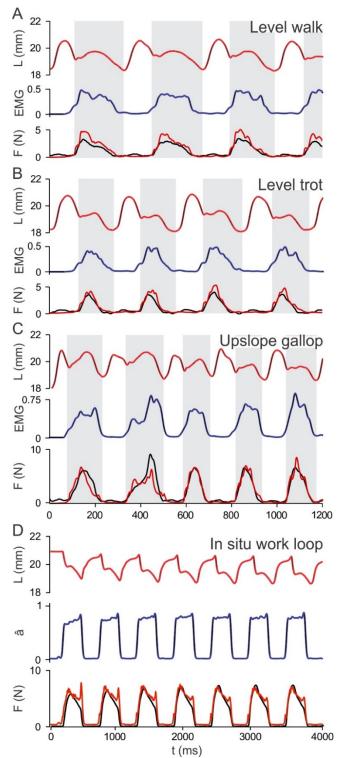




Fig. 2. Rat MG length, EMG and forces for *in-vivo* locomotion (A-C) and *in-situ* work-loop
recordings. Length is from the sonomicrometry measures of the muscle belly. EMG-intensity
(*in-vivo*) or activation (*in-situ*) is pooled between the proximal and distal sites. Forces were
measured at the tendon (black) or predicted from the muscle models (red lines).

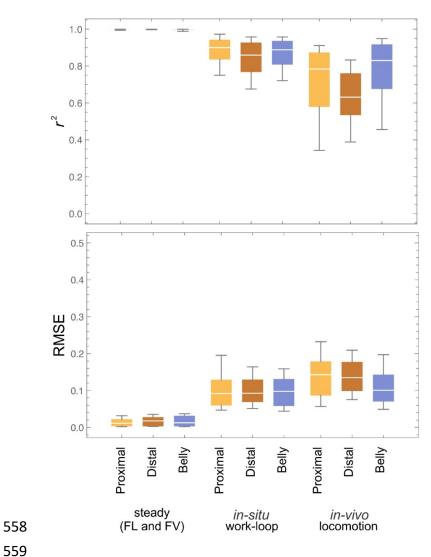
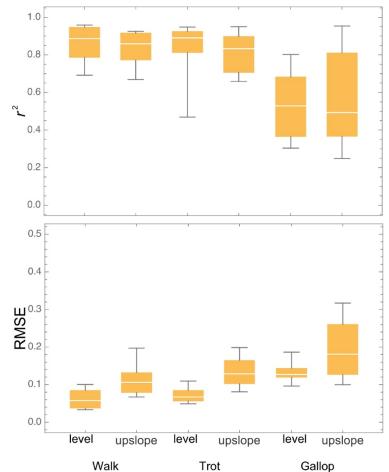


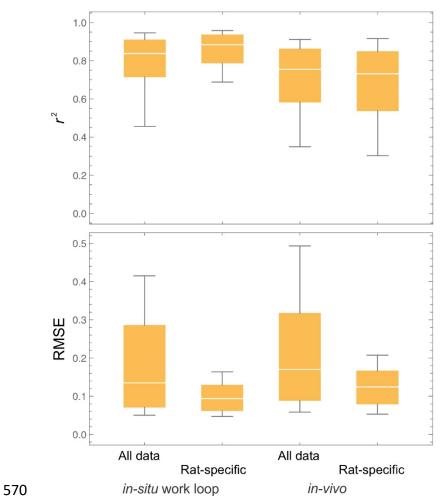
Fig. 3. Summary box plots (r² and RMSE) comparing quality of modelled forces when 560

561 using rat-specific intrinsic properties in the models. Box plots show median values with 25/75

562 quartiles, and whiskers delineating the 90% quartile ranges.



563WalkTrotGallop564Fig. 4. Box plots (r^2 and RMSE) across *in-vivo* conditions averaged for the whole muscle565belly. Data are taken for the whole muscle belly. These models using rat-specific intrinsic566properties yielded forces with lower r^2 and higher RMSE for gallop compared with trotting and567walking conditions (see Fig. S3 for statistical evaluation). RMSE was generally higher for568upslope gait versus level gait conditions. Box plots show median values with 25/75 quartiles, and569whiskers delineating the 90% quartile ranges.



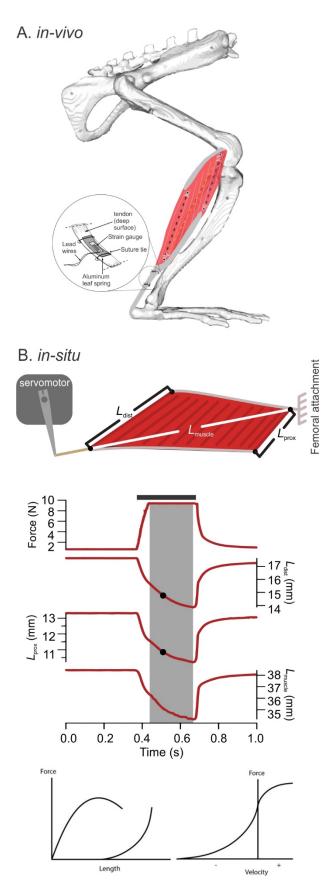
571 Fig. 5. Summary box plots (*r*² and RMSE) across steady versus dynamic *in-situ* work-loop

572 and *in-vivo* conditions. Note how the models using intrinsic properties pooled across the

573 population of rats had model fits with higher RMSE than compared to models with rat-specific

574 properties. Box plots show median values with 25/75 quartiles, and whiskers delineating the 90%

575 quartile ranges.



577 Fig. S1. Schematic of (A) *in-vivo* and (B) *in-situ* force and length measurements. (A) *In-vivo*

- 578 MG forces were recorded using a leaf-spring force transducer surgically attached to the free MG
- tendon (Richards and Biewener, 2007; Eng et al. 2019). Proximal and distal fascicle length
- 580 changes, as well as whole muscle belly length were recorded using 1.0 mm sonomicrometry
- 581 crystals, along with fine-wire EMG electrodes implanted in both regions (not shown). (B) In-situ
- 582 MG forces were measured by isolating the MG tendon and calcar attachment to a lever-based
- 583 ergometer, with the proximal origin of the muscle remaining attached to the femur, which was
- 584 fixed by a clamp. Stimulation of the muscle via the sciatic nerve elicited tetanic FL and FV
- 585 contractions under supramaximal (2-3 v) and submaximal (~1 v) conditions, allowing active and
- 586 passive FL and isotonic FV curves to be constructed.

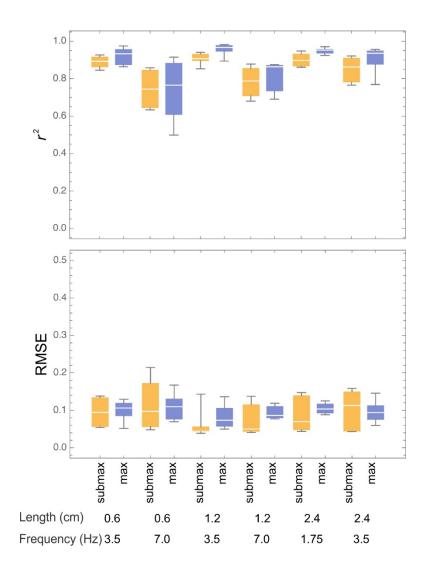


Fig. S2. Box plots (r^2 and RMSE) comparing quality of modelled forces when using rat-589 590 specific intrinsic properties in the models across *in-situ* work-loop conditions (Alength & frequency) for the whole muscle belly. Predictive fits of rat-specific muscle models to forces 591 592 measured during *in-situ* work-loops were generally similar across the range of imposed lengths 593 and frequencies, with r^2 averaging 0.88+0.10 and RMSE averaging 0.09+0.04. No significant difference in RMSE or r^2 was observed for model fits to supramaximal versus submaximal 594 595 intensity work-loops (Fig. S3). Box plots show median values with 25/75 quartiles, and whiskers 596 delineating the 90% quartile ranges.

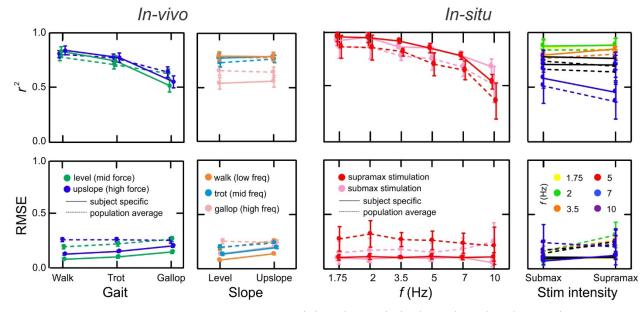


Fig. S3. Least-Square Means plots summarizing the statistical results related to testing 598 hypothesis 1; that model-predictions of muscle forces will be less accurate at the highest 599 600 contractile frequencies but not the greatest magnitude of force. For both in-vivo (Left) and in-situ (Right), we used a nested factorial design to keep either the "force" or "frequency" parameter 601 constant, while testing across conditions for the alternate parameter. Force was dictated *in-vivo* 602 by the treadmill slope condition (Konow et al., 2020), and *in-situ* by sub- or supra-maximal 603 muscle stimulation. Frequency was dictated *in-vivo* by gait condition, and *in-situ* by varying 604 605 work-loop frequency. Within each condition, tests were run separately for the population average and subject specific model inputs. Results are largely consistent across conditions, with r^2 values 606 607 being significantly reduced by increasing cycle frequency but remaining largely invariant by 608 changes in force. RMSE values are generally better for modeling based on population average 609 data. For statistical test results, please see the results section.

- 611
- 612
- 613

| Table 1. Active FL parameters. See equation 1. Rat-specific parameters are shown as mean±SD. | | | | | | | | | |
|--|--------|-------|--------|--------|-------|-------|-------------|-------------|--|
| rwS l_0 [mm] F_0 [N] r^2 RMS | | | | | | | | | |
| Proximal | Pooled | 2.37 | 0.217 | 0.475 | 8.57 | 8.52 | 0.857 | 0.128 | |
| | Rat | 2.83 | 0.206 | 0.501 | 8.74 | 9.88 | 0.998 | 0.013 | |
| | | ±0.75 | ±0.256 | ±0.691 | ±2.45 | ±6.26 | ± 0.002 | ± 0.007 | |
| Distal | Pooled | 2.29 | 0.135 | 0.417 | 9.48 | 8.54 | 0.732 | 0.209 | |
| | Rat | 2.93 | 0.091 | 0.289 | 9.14 | 9.89 | 0.994 | 0.019 | |
| | | ±0.78 | ±0.159 | ±0.592 | ±3.03 | ±6.27 | ±0.012 | ±0.012 | |
| Belly | Pooled | 2.17 | 0.096 | 0.567 | 22.0 | 8.56 | 0.846 | 0.125 | |
| | Rat | 2.57 | 0.079 | 0.477 | 21.9 | 9.90 | 0.994 | 0.019 | |
| | | ±0.50 | ±0.204 | ±1.236 | ±3.1 | ±6.26 | ±0.006 | ±0.012 | |

| Table 2. Passive FL parameters. See equation 2. Rat-specific parameters are shown as | | | | | | | | | | | |
|--|------------------------|-----------|-----------|-------------|-------------|--|--|--|--|--|--|
| mean±SD. | | | | | | | | | | | |
| | c_1 c_2 r^2 RMSE | | | | | | | | | | |
| Proximal | Pooled | -13.0 | 7.74 | 0.021 | 0.329 | | | | | | |
| | Rat | -14.3±4.2 | 9.26±4.11 | 0.997±0.003 | 0.005±0.006 | | | | | | |
| Distal | Pooled | -12.3 | 7.44 | 0.040 | 0.308 | | | | | | |
| | Rat | -14.1±4.6 | 9.26±4.51 | 0.998±0.001 | 0.006±0.009 | | | | | | |
| Belly | Pooled | -21.3 | 16.7 | 0.079 | 0.185 | | | | | | |
| | Rat | -23.8±7.0 | 19.3±6.72 | 0.995±0.007 | 0.008±0.012 | | | | | | |

| Table 3. FV parameters. See equation 3. Rat-specific parameters are shown as mean \pm SD. | | | | | | | | |
|--|--------|-------------|----------------|-------------|-------------|--|--|--|
| | | k | $v_0 [s^{-1}]$ | r^2 | RMSE | | | |
| Proximal | Pooled | 0.802 | 10.82 | 0.983 | 0.064 | | | |
| | Rat | 1.24±1.20 | 10.82±3.01 | 0.998±0.002 | 0.027±0.013 | | | |
| Distal | Pooled | 0.638 | 9.82 | 0.989 | 0.051 | | | |
| | Rat | 0.775±0.473 | 9.82±3.22 | 0.997±0.002 | 0.028±0.011 | | | |
| Belly | Pooled | 0.509 | 5.06 | 0.989 | 0.051 | | | |
| | Rat | 0.599±0.391 | 5.06±0.98 | 0.997±0.002 | 0.026±0.014 | | | |

| | Body mass | MG mass | l _{muscle} | Proximal | Distal | Proximal | Distal |
|-----|-----------|---------|---------------------|-----------------------------------|-----------------------------------|----------------------------------|----------------------------------|
| Rat | (g) | (g) | (mm) | $l_{\text{fascicle}} (\text{mm})$ | $l_{\text{fascicle}} (\text{mm})$ | θ_{fascicle} (deg) | θ_{fascicle} (deg) |
| 1 | 207 | 0.446 | 19.1 | 6.7 | 8 | 27 | 19 |
| 2 | 247 | 0.673 | 27.6 | 6.8 | 8.3 | 20 | 15 |
| 3 | 289 | 0.719 | 29.1 | 7.8 | 8.9 | 21 | 17 |
| 4 | 362 | 0.976 | 33.2 | 8.8 | 10.3 | 22 | 17 |
| 5 | 380 | 1.035 | 31.8 | 8.4 | 10.3 | 25 | 21 |
| 6 | 360 | 0.911 | 30.1 | 7.9 | 9.6 | 21 | 20 |