**Mycobacterium ulcerans** infections cause progressive muscle atrophy and dysfunction, and mycolactone impairs satellite cell proliferation

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Submitted 17 June 2010; accepted in final form 3 January 2011

Houngbédi GM, Bouchard P, Frenette J. Mycobacterium ulcerans infections cause progressive muscle atrophy and dysfunction, and mycolactone impairs satellite cell proliferation. Am J Physiol Regul Integr Comp Physiol 300: R724–R732, 2011. First published January 5, 2010; doi:10.1152/ajpregu.00393.2010.—Clinical observations from Buruli ulcer (BU) patients in West Africa suggest that severe *Mycobacterium ulcerans* infections can cause skeletal muscle contracture and atrophy leading to significant impairment in function. In the present study, male mice C57BL/6 were subcutaneously injected with *M. ulcerans* in proximity to the right biceps muscle, avoiding direct physical contact between the infectious agent and the skeletal muscle. The histological, morphological, and functional properties of the muscles were assessed at different times after the injection. On day 42 postinjection, the isometric tetanic force and the cross-sectional area of the myofibers were reduced by 31% and 29%, respectively, in the proximate-infected muscles relative to the control muscles. The necrotic areas of the proximate-infected muscles had spread to 7% of the total area by day 42 postinjection. However, the number of central nucleated fibers and myogenic regulatory factors (MyoD and myogenin) remained stable and low. Furthermore, Pax-7 expression did not increase significantly in mycolactone-infected muscles, indicating that the satellite cell proliferation is abrogated by the toxin. In addition, the fibrotic area increased progressively during the infection. Lastly, muscle-specific RING finger protein 1 (MuRF-1) and atrogin-1/muscle atrophy F-box protein (atrogin-1/MAFbx), two muscle-specific E3 ubiquitin ligases, were upregulated in the presence of *M. ulcerans*. These findings confirmed that skeletal muscle is affected in our model of subcutaneous infection with *M. ulcerans* and that a better understanding of muscle contractures and weakness is essential to develop a therapy to minimize loss of function and promote the autonomy of BU patients.

skeletal muscle force; degeneration; edema; fibrosis; muscle regeneration

**BURULI ULCER (BU)** IS AN INFECTIOUS disease caused by *Mycobacterium ulcerans*. BU is currently the third most common mycobacterial disease in the world, after tuberculosis and leprosy. This neglected infectious disease, which affects mostly the poorest people in developing countries, can cause very severe chronic skin lesions if diagnosis and treatment are delayed. In West Africa, most BU patients experience disabilities resulting from cutaneous fibrosis and soft tissue contracture, which limit the range of motion of the affected joint (7, 25). While skeletal muscles play a key role in joint mobility and are located just under the necrotic skin infected by *M. ulcerans*, they have received little attention in BU studies. Moreover, clinical investigations of BU patients in West Africa have revealed that over half of the patients suffer from range-of-motion limitations and functional restrictions (7). Furthermore, anecdotal and empirical observations suggest that BU patients generally suffer from skeletal muscle necrosis, atrophy, and contracture. In our mice model of BU, the subcutaneous injection of *M. ulcerans* has been shown to cause necrosis, chronic inflammatory response, fibrosis, and skeletal muscle stiffness (17). Clinical and experimental findings, thus, indicate that *M. ulcerans* infections play a major role in the loss of skeletal muscle integrity.

Repair and maintenance of skeletal muscle are attributed to the satellite cells. In response to several stimuli, including muscle injury, quiescent satellite cells activate, proliferate, and differentiate to repair damaged skeletal muscle (4). Pax-7 is expressed and transcriptionally active in quiescent satellite cells (26). Following activation, satellite cells proliferate and coexpress Pax-7 and MyoD (26). Pax-7 is also implicated in the regulation of satellite cell self-renewal, and Pax-7-null mice present impairment in muscle regeneration combined with a progressive loss in satellite cell number due to cell cycle defect and increased apoptosis (19).

Muscle mass is maintained by a fragile balance between protein synthesis and degradation, and muscle atrophy can occur when this balance is disturbed. The selective loss of contractile proteins relative to cellular proteins generally results in muscles with smaller myofibers (3). Chronic systemic inflammation is one of the potential conditions that can lead to significant muscle atrophy (21). For example, sepsis is characterized by chronic, systemic inflammation associated with both a reduction in protein synthesis and a breakdown of muscle proteins, in particular, the contractile proteins actin and myosin (12, 13). The atrophying muscles display increased rates of protein degradation, mainly caused by the activation of the ubiquitin proteasome system (2). In addition, the muscle-specific E3 ubiquitin ligase muscle-specific RING finger protein 1 (MuRF-1) and atrogin-1/muscle atrophy F-box proteins (atrogin-1/MAFbx) are upregulated, while myogenic regulatory factors, such as MyoD and myogenin, are usually downregulated (20, 35). Since proinflammatory cytokines possess a strong potential for mediating sepsis-induced proteolysis of myofibrillar proteins in skeletal muscles (9, 12, 21) and since a number of these cytokines are upregulated during *M. ulcerans* infections (17), we verified the impact of *M. ulcerans* infections on biceps muscle at the histological, morphological, and functional levels. We demonstrated that the presence of *M. ulcerans* and mycolactone reduces muscle force production, upregulates MuRF-1 and atrogin-1, and impairs satellite cell proliferation, all of which are associated with fibrosis and muscle atrophy and dysfunction.

**MATERIALS AND METHODS**

Animal care and feeding. Male C57BL/6 mice weighing 20–22 g (Charles River Laboratories, Saint-Constant, QC, Canada) were housed two per cage in pathogen-free conditions. All animal care and handling procedures were approved by the Université Laval Animal Protection Committee. The mice were maintained on a 12:12-h
light-dark cycle, and food and water were provided ad libitum. No differences in food and water consumption or physical activity were observed between groups.

**Experimental design.** A previously developed protocol was used to evaluate the histological, biochemical, and functional impacts of *M. ulcerans* infections on skeletal muscles (17). In the present study, the contractile properties, atrophy, fibrosis, and regeneration of biceps muscles were assessed in 72 mice randomly assigned to three groups: 1) no treatment (control, CTR, n = 24), 2) subcutaneous injection near the right biceps muscle with 30 μL of PBS containing culture media (Sham, n = 24), and 3) subcutaneous injection with *M. ulcerans* suspended in PBS [10^7 acid-fast bacilli (AFB)/30 μL] near the right biceps muscle [proximate-infected biceps (PIB); n = 24]. The biceps muscle was chosen because it is a superficial and easy muscle to dissect and it possesses two distinct tendons at its extremities. This superficial muscle allows us to perform subcutaneous injection mimicking somehow the mode of *M. ulcerans* infection. All mice from each group were killed on day 7 (n = 6), day 21 (n = 9), and day 42 (n = 9) postinjection. The infectious process did not progress extensively, and no apparent changes in physical activities were observed in infected mice.

*M. ulcerans* injection protocol and surgical procedure. The injection and surgical procedures were performed as described previously (15). Briefly, *M. ulcerans* strain 957–05 isolated from a BU patient in 2005 and identified by PCR using the insertion sequence IS2404 was grown for 6–8 wk on Lowenstein-Jensen medium at 30°C in a reduced oxygen environment (5% CO2). The mycobacteria were collected and suspended in PBS. A homogeneous aliquot (10 μL) was stained using the Ziehl-Neelsen method, and the concentration of AFB was determined at 562 nm using a Perkin-Elmer plate reader (6). The mycobacteria were suspended in PBS with 30 μL of PBS containing culture media (Sham), 70% ethanol (Sham), 1 μM CaCl2, 2 μM NaF, 8.3 μM NaVO4, 2 μM PMSF, 1 ml glycerol, 100 μl Igepal, and 10 μl of protease inhibitors in a final volume of 10 μL). The homogenate was centrifuged at 10,000 g for 5 min, and the pellet was discarded.

**RESULTS**

**Measurements of contractile properties in vitro.** To examine muscle force and function, each tendon of CTR, Sham, and PIB muscles was attached using 3–0 silk sutures. The distal tendon was tied to a rigid support between two platinum electrodes submerged in Krebs-Ringer buffer supplemented with 2 mg/mL of glucose as described previously (17). The muscle was aligned vertically, and the proximal tendon was fixed to the arm of a dual-mode servomotor (305B-LR, Aurora Scientific, Aurora, ON, Canada) controlled by Dynamic Muscle Control software. Carbogen (95% O2:5% CO2) was constantly bubbled through the solution, which was kept at 25°C. Twitch tension and isometric contractions were performed, and twitch peak tension (Pt, mN) and maximal isometric force (Pmax, mN) values were recorded, and force-frequency curves were generated as described previously (14).

**Assessment of area, fiber regeneration, edema, and fibrosis of skeletal muscles.** Because the histological and contractile results indicated that there was no difference between Sham and CTR muscles, muscle atrophy, repair, and fibrosis were assessed on CTR and PIB muscles only. Following the evaluation of the functional properties, the CTR and PIB muscles were trimmed of visible fat, tendons, and other connective tissues. The biceps were then weighed and stretched to near resting length, embedded in tissue freezing medium (Triangle Biomedical Sciences, Durham, NC, USA), and frozen in isopentane that was cooled in liquid nitrogen. Muscle tissue sections (10 μm thick) were stained with hematoxylin-and-eosin, and images were acquired at ×200 magnification. To measure muscle atrophy, the CSA of 200 randomly selected myofibers per muscle section were determined using ImageJ software. To assess the necrosis and repairing processes of skeletal muscles, the necrotic area and the number of central nucleated myofibers were determined and expressed per CSA. The necrotic area was estimated in each muscle section by subtracting the summed area of normal and regenerating fibers from the total area of each field. (18). Other tissue sections from CTR and PIB muscles were stained with Masson’s trichrome to color collagen blue. The surface areas of blue-stained tissues were measured and expressed as percentages of the total CSAs of the muscles. In another set of experiments, CTR, Sham, and PIB muscle samples were weighed and dehydrated for 18 h in a SpeedVac concentrator. Wet and dry muscle masses were subtracted to assess water content and edema.

**Protein sample preparation and Western blot analysis.** CTR, Sham, and PIB muscles were homogenized in 200 μL of lysis buffer (20 mM Tris, pH 7.5, 140 mM NaCl, 10 μM MgCl2, 1 μM CaCl2, 2 μM NaF, 8.3 μM NaVO4, 2 μM PMSF, 1 ml glycerol, 100 μl Igepal, and 10 μl of protease inhibitors in a final volume of 10 μL). The homogenate was centrifuged at 10,000 g for 5 min, and the pellet was discarded. The protein content of the homogenates was measured using BCA protein assay kits (EMD Chemical, Darmstadt, Germany). The protein concentrations were compared with a standard curve and determined at 562 nm using a Perkin-Elmer plate reader (6). The protein suspensions (50 μg) were separated by SDS-PAGE. The proteins were transferred to a polyvinylidene difluoride membrane (Bio-Rad, Hercules, CA) and incubated overnight at 4°C with primary antibodies (atrogen-1 rabbit polyclonal; ECM Biosciences, Versailles, KY), MuRF-1 goat polyclonal (Novus Biological, Littleton, CO), MyoD rabbit polyclonal (Santa Cruz Biotechnology, Santa Cruz, CA), and myogenin rabbit polyclonal (Santa Cruz Biotechnology) diluted 1:1000 in 5% dry milk. α-Tubulin (Santa Cruz Biotechnology) was used as an endogenous reference to monitor protein loading. Protein bands were detected using an enhanced chemiluminescence detection substrate (Western Lightning Plus ECL; Perkin Elmer, Waltham, MA). The chemiluminescent signal was acquired on Kodak film (X-OMAT; LS Kodak scientific imaging film) and relative signal bands were quantified from scans (TIFF images) by densitometric analysis of integrated gray levels corrected for local background using Quantity One software (Bio-Rad).

**Pax-7 expression in injured skeletal muscles.** To evaluate myogenic potential in various conditions, skeletal muscles were injected and chemically damaged with 10 μl of PBS (Sham), 70% ethanol (Sham), mycolactone (25 μg), or bupivacaine (50 μg). Bupivacaine is a very well known myotoxin molecule that produces very reproducible induction of muscle damage and repair. Because bupivacaine destroys myofibers but spares satellite cells, leaving the basal laminae and microcirculation intact, it can be used as a positive control for muscle regeneration and recovery. After 7 days postinjury, control and experimental muscles were then dissected, homogenized, and immunolabeled with Pax-7 (Abcam) on Western blot, as previously described to assess satellite cell survival and proliferation.

**Statistical analysis.** All data are reported as means ± SE. The data were compared with a one-way ANOVA using JMP software (JMP 7; SAS Institute, Cary, NC) and Tukey’s a posteriori test. P values <0.05 were considered significant.

**RESULTS**

*M. ulcerans* causes a significant loss of muscle force. To evaluate the functional impact of *M. ulcerans* on the biceps muscles, twitch tension (P, mN) and maximal isometric force (absolute P0, mN) values were recorded in CTR, Sham, and PIB muscles on days 7, 21, and 42 postinfection. The Pt values...
for the PIB muscles were 12.8% and 25% lower than the Sham muscles on days 21 and 42, respectively (Fig. 1A). The absolute $P_0$ values were 18% and 31% lower for the PIB muscles than for the Sham muscles on days 21 and 42, respectively (Fig. 1B). The decline in muscle force was time dependent in PIB muscles, and significant differences were observed between PIB muscles at 7, 21, and 42 days. These decrements were nearly constant since the force-frequency curves for the PIB muscles were significantly lower than the Sham and CTR muscles (Fig. 1C). However, *M. ulcerans* had no significant influence on speed-related contractile parameters, since the time-to-peak tension (TPT; ms) and half-relaxation time (RT½; ms) values were not significantly different from those of the

![Figure 1A](image1.png)

**Figure 1A.** Isometric contractile properties of biceps muscles. Maximum twitch tension $P_t$ (A), maximum tetanic force $P_0$ (B), and force-frequency curves (C) are shown for control (CTR), sham (Sham), and proximate-infected biceps (PIB) muscles on days 7, 21, and 42 postinfection. Isometric force decreased progressively during the infection. All values are means ± SE; $n = 6$ to 9. No significant differences were observed between CTR and Sham muscles. *Significantly different from the matched CTR muscle on day 7.* #Significantly different from the matched PIB muscle on day 7. ‡Significantly different from the matched Sham muscle on day 21. ‡‡Significantly different from the matched PIB muscle on day 21; ($P < 0.05$).

![Figure 1B](image2.png)

![Figure 1C](image3.png)

**Table 1.** The morphological characteristics were obtained from control, Sham, and proximate-infected biceps muscles on days 7, 21, and 42 postinjection.

<table>
<thead>
<tr>
<th>Day 7</th>
<th>Day 21</th>
<th>Day 42</th>
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</thead>
<tbody>
<tr>
<td>CTR</td>
<td>Sham</td>
<td>PIB</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Body weight, g</td>
<td>25.1 ± 0.57</td>
<td>25.04 ± 0.39</td>
</tr>
<tr>
<td>Biceps-dry weight, mg</td>
<td>3.89 ± 0.12</td>
<td>3.90 ± 0.11</td>
</tr>
<tr>
<td>Biceps-water content, mg</td>
<td>12.90 ± 0.20</td>
<td>13.24 ± 0.12</td>
</tr>
</tbody>
</table>

All values are means ± SE; $n = 6$ to 9. CTR, control; PIB, proximate-infected biceps. *Significantly different from the matched Sham muscle on day 7.* †Significantly different from the matched PIB muscle on day 7; ($P < 0.05$).
M. ulcerans and mycolactone cause muscle edema, damage, atrophy, and fibrosis with impairment in satellite cell proliferation. The M. ulcerans infections resulted in significant edema in the biceps muscles, as indicated by the wet weight

Sham muscles at all time points (data not shown). Although M. ulcerans causes myofiber atrophy, edema, and the accumulation of connective tissue, which can all change the size of skeletal muscles, we also found a significant decrease in muscle-specific force (sP₀, N/cm²) at 21 and 42 days postinjection relative to sham muscles (data not shown).

Fig. 2. Histological staining of cross-sectional biceps muscles. A and B: CTR and PIB muscles on day 42 were stained with Masson’s trichrome. B: Masson’s trichrome-stained fibrotic tissues are indicated in brackets. The fibrotic muscle area (C) was quantified for CTR and PIB muscles on days 7, 21, and 42 postinjection. All values are expressed as means ± SE; n = 6 to 9. *Significantly different from the matched CTR muscle. #Significantly different from the matched PIB muscle on day 7 (P ≤ 0.05). Scale bar = 100 μm.

Fig. 3. Histological and morphological analyses of biceps muscles. A and B: CTR and PIB muscles on day 42 were stained with hematoxylin and eosin. C: The cross-sectional area of muscle fibers was quantified for CTR and PIB muscles on days 7, 21, and 42 postinjection. The arrow indicates a central nucleated fiber. All values are means ± SE; n = 6 to 9. *Significantly different from the matched CTR muscle. #Significantly different from the matched PIB muscle on day 7 (P ≤ 0.05). Scale bar = 100 μm.
and dry/wet weight ratios of PIB muscles compared with the sham muscles. The edema progressed steadily, with the water content of the PIB muscles increasing from 2.6% to 12% between days 7 and 21 relative to the sham muscles (Table 1). Interestingly, the edema decreased thereafter, with no significant differences between the CTR, Sham, and PIB muscles on day 42. To assess muscle fibrosis, CTR and PIB muscle sections were stained with Masson’s trichrome. As expected, there was more collagen in the PIB muscles than in the CTR muscles (Figs. 2, A and B). The area of fibrotic tissue increased over time and reached 4.5% and 5.3% of the muscle CSA on days 21 and 42, respectively (Fig. 2C). The histological observations with hematoxylin and eosin suggested that the edema was mainly concentrated between muscle fascicles (Figs. 3, A and B). Furthermore, the M. ulcerans infections caused a significant reduction in myofiber CSA, with the mean values for the PIB muscles dropping by 17% and 29% on days 21 and 42, respectively, relative to the CTR muscles (Fig. 3C). The number of central nucleated myofibers was also determined to estimate the number of regenerating fibers. The CTR muscles did not display any fiber damage or central nucleated myofibers while the number of regenerating fibers remained stable at ~6–7 myofibers/mm² from days 7 to 42 in infected skeletal muscles (Fig. 4A). Consistent with this observation, the levels of expression of myogenic regulatory factors MyoD and myogenin remained relatively low and stable at all experimental time points (Figs. 4, B and C).

Histological observations also showed that muscle injected with 70% ethanol (Sham) had minor inflammatory reaction (Fig. 5A), whereas skeletal muscles injected with PBS (Sham) presented no sign of histological damage (Fig. 5B). The injection of mycolactone (25 μg) or bupivacaine (50 μg) induced extensive damage with significant cell infiltration at 7 days postmuscle injury (Fig. 5, C and D). The extent of damage was particularly important following mycolactone injection (Fig. 5C). More importantly, the level of expression of transcription factor Pax-7, expressed on quiescent and activated satellite cells, increased by more than 12-fold in bupivacaine-injected muscles but remained similar to Sham in mycolactone-injected muscles at 7 days postinjury (Fig. 5E). These results clearly indicated for the first time that satellite cell proliferation is abrogated in the presence of mycolactone (Fig. 5E). As expected, the PIB muscles displayed significant areas of necrosis on days 21 and 42 compared with day 7 (Fig. 6A). The area of necrotic tissue increased and reached 7% of the muscle CSA on day 42. This catabolic state was associated with an upregulation of the muscle-specific ubiquitin ligases (E3) atrogin-1/MAFbx (muscle atrophy F-box protein) and MuRF-1 (musclespecific RING-finger 1). These two muscle-specific genes for
atrophy rose 2.4- and 1.5-fold, respectively, at day 42 compared with CTR muscles (Figs. 6, B and C).

DISCUSSION

The loss of contractile proteins and normal sarcomeric organization associated with muscle wasting is the hallmark of diverse catabolic conditions, such as muscle disuse, burn injury, cancer, renal failure, AIDS, chronic obstructive pulmonary disease, space flight, muscular dystrophy, and aging (14, 22, 30, 31, 34). Loss of muscle mass is also commonly seen in animal models and patients with sepsis (1, 17, 27). This catabolic state of skeletal muscles is mainly driven by polyubiquitination, which involves a cascade of proteolytic enzymes, such as ubiquitin-activating enzyme, ubiquitin-conjugated enzyme, and ubiquitin-protein ligase. Ultimately, the loss of muscle mass results in weakness and fatigue that limit ambulation and prolong rehabilitation.

The present findings showed that the M. ulcerans infections caused a 31% and 29% decrease in the force and CSA of myofibers, respectively, which was associated with an upregulation of atrogin-1 and MuRF-1, two muscle-specific ubiquitin-ligases involved in protein degradation. The ubiquitin-proteasome system is the major pathway responsible for the turnover of muscle protein, in general, and, more specifically, in sepsis (35). For example, atrogin-1 and MuRF-1 play an essential role in sepsis-induced muscle wasting. More recent data have
indicated that MuRF-1 acts by targeting myofiber proteins, such as titin, troponin, myosin light-chain-2, and nebulin (33), leading to degradation and muscle loss of mass. Another major mechanism of sepsis-induced muscle wasting is the activation of the calpain system. Calpain and caspase-3 appear to play a pivotal role in ubiquitin-proteasome-mediated muscle wasting by promoting myofibrillar proteolysis (5, 29). Both the ubiquitin-proteasome and calpain proteolytic pathways have been shown to be partly controlled by proinflammatory cytokines (9, 12, 21), such as TNF-α and IL1β, which increase during M. ulcerans infections (17). While the assessment of protein synthesis and degradation was not within the scope of the work reported here, other research on muscle atrophy and sepsis suggests that the loss of muscle mass and the reduction in the CSA of myofibers in mice with M. ulcerans infections is likely the result of an upregulation of protein degradation combined with a reduction in protein synthesis.

Muscle growth and regeneration are attributed to satellite cells. Our results showed that M. ulcerans and mycolactone have negative impacts on the regeneration process since the number of centronucleated myofibers and the level of myogenic regulatory factors MyoD and myogenin and Pax-7 did not increase over time in the PIB muscles. MyoD is one of the earliest markers of myogenic commitment, which is expressed in activated satellite cells (23), while myogenin is essential for the development of functional skeletal muscles (24, 27). The lack of a significant increase in MyoD and myogenin content suggested that the presence of M. ulcerans prevented muscle regeneration. More importantly, the level of Pax-7 expression increased by 12-fold in bupivacaine-injected muscles, while the level of Pax-7 expression in mycolactone-injected muscle remains stable, indicating that satellite cell proliferation is inhibited at 7 days postinjury. These results are in agreement with previous findings demonstrating that injecting skeletal muscles with mycolactone totally abrogates muscle regeneration (16). The fact that the number of regenerating fibers did not significantly increase in these skeletal muscles may, thus, be due to an increase in satellite cell apoptosis, satellite cell inactivation, and/or the decreased unavailability of cytokines and growth factors that govern the muscle regeneration process. Furthermore, mycolactone can diffuse far from the site of infection (11, 15), penetrate membranes of other cell types, induce cytoskeleton rearrangement, and ultimately cause cell death by apoptosis (10, 28).

Alternate interpretations for muscle dysfunction are also possible. First, an in vivo study of mice footpad infections demonstrated that M. ulcerans can invade and produce nerve necrosis (8). It is, thus, possible that biceps nerves and/or neuromuscular junctions may be partly dysfunctional, leading to progressive muscle atrophy. Second, the accumulation of
leukocytes and fibrotic tissue over time indicates that infected biceps muscles are subject to an inappropriate chronic inflammatory reaction leading to scar formation instead of muscle regeneration (17). The inadequate chronic inflammatory response may lead to fibrosis caused by persistent irritant or an upregulation of fibrogenic growth factors and cytokines, which together stimulate the synthesis and deposition of extracellular matrix (36). Previous observations have shown that muscle regeneration does not occur in the presence of mycolactone and that the collagen content of skeletal muscle increases by 134%, indicating that mycolactone stimulates fibroblasts, which are a major cell type in skeletal muscles (16). These results are consistent with the increase in the fibrotic areas of the PIB muscles observed on days 21 and 42. Further in vitro and in vivo studies are needed to clarify the molecular mechanisms by which M. ulcerans, and/or mycolactone, and/or recruited leukocytes that secrete proinflammatory cytokines are involved in muscle atrophy and muscle regeneration impairment.

**Perspectives and Significance**

BU is a neglected disease. Little research has been conducted on the impact of M. ulcerans on skeletal muscles. The present study demonstrated that M. ulcerans infections lead to a progressive and significant decrease in tetanic force production and twitch tension of biceps muscles close to the site of infection. Histological, morphological, and biochemical analyses also indicated that M. ulcerans infections and mycolactone induce myofiber atrophy and fibrosis, and they inhibit satellite cell proliferation. These results confirmed the involvement of skeletal muscles at the functional level in our model of subcutaneous infection with M. ulcerans. The subcutaneous injection of the M. ulcerans was chosen because it mimics the mode of contamination of the disease from bug bites to humans. Furthermore, mice injected with M. ulcerans present a discrete nodule at 7 days postinfection. Several mice exhibit sequentially a local edema and skin ulceration at 21 and 42 days postinfection, respectively (17). The same chronological manifestation of the disease is found in humans, making this animal model useful and relevant. We trust that the present work will set the stage for an extensive research program on skeletal muscle rehabilitation directed toward a better understanding of muscle contracture and weakness that often lead to a loss of function, autonomy, and dignity in BU patients (see the following Web address for details on this disease: http://www.who.int/buruli/en).

**ACKNOWLEDGMENTS**

We thank Dr Sévérin Anagonou and his team for the M. ulcerans strain 957-05. We also gratefully acknowledge the critical comments of Patrice Bouchard. This investigation was supported by grants to J. Frenette from the Natural Sciences and Engineering Research Council of Canada and the Canadian Institutes of Health Research.

**DISCLOSURES**

No conflicts of interest, financial or otherwise, are declared by the authors.

**REFERENCES**


