Impaired neuromuscular transmission and skeletal muscle fiber necrosis in mice lacking Na/Ca exchanger 3

Sophie Sokolow,1,2 Mario Manto,3 Philippe Gailly,4 Jordi Molgó,5 Clarisse Vandebrouck,4 Jean-Marie Vanderwinden,6 André Herchuelz,2 and Stéphane Schurmans1

1Institut de Recherches en Biologie Humaine et Moléculaire–Institut de Biologie et de Médecine Moléculaires (IRIBHM-IBMM), Université Libre de Bruxelles, Gosselies, Belgium
2Laboratory of Pharmacology and Therapeutics and Laboratory of Experimental Neurology, Université Libre de Bruxelles, Brussels, Belgium
3Laboratoire de Neurobiologie Cellulaire et Moléculaire, Unité Propre de Recherche 9040, Centre National de la Recherche Scientifique, Gif-sur-Yvette, France
4Laboratory of Neurophysiology, Université Libre de Bruxelles, Brussels, Belgium
5Institut Fédératif de Neurobiologie Alfred Fessard, Laboratoire de Neurobiologie Cellulaire et Moléculaire, Unité de Recherche 9040, Centre National de la Recherche Scientifique, Gif-sur-Yvette, France
6Laboratory of Experimental Neurology, Université Libre de Bruxelles, Brussels, Belgium

We produced and analyzed mice deficient for Na/Ca exchanger 3 (NCX3), a protein that mediates cellular Ca2+ efflux (forward mode) or Ca2+ influx (reverse mode) and thus controls intracellular Ca2+ concentration. NCX3-deficient mice (Ncx3−/−) present a skeletal muscle fiber necrosis and a defective neuromuscular transmission, reflecting the absence of NCX3 in the sarcolemma of the muscle fibers and at the neuromuscular junction. The defective neuromuscular transmission is characterized by the presence of electromyographic abnormalities, including low compound muscle action potential amplitude, a decremental response at low-frequency nerve stimulation, an incremental response, and a prominent postexercise facilitation at high-frequency nerve stimulation, as well as neuromuscular blocks. The analysis of quantal transmitter release in Ncx3−/− neuromuscular junctions revealed an important facilitation superimposed on the depression of synaptic responses and an elevated delayed release during high-frequency nerve stimulation. It is suggested that Ca2+ entering nerve terminals is cleared relatively slowly in the absence of NCX3, thereby enhancing residual Ca2+ and evoked and delayed quantal transmitter release during repetitive nerve stimulation. Our findings indicate that NCX3 plays an important role in vivo in the control of Ca2+ concentrations in the skeletal muscle fibers and at the neuromuscular junction.


Introduction
In order to avoid prolonged elevation of cytosolic free calcium concentration ([Ca2+]i), cells use multiple channels, pumps, exchangers, and chelators (1). The Na/Ca exchanger (NCX) is a plasma membrane protein responsible for Ca2+ extrusion from cells. Indeed, in its forward mode, NCX extrudes one Ca2+ against three entering Na+, using the Na+ gradient across the plasma membrane as the energy source (2, 3). In view of its high exchange capacity, NCX is well suited for rapid recovery from high [Ca2+]i and may play an important role in maintaining Ca2+ homeostasis and protecting cells from Ca2+ overload and death (2, 4). In its reverse mode, NCX allows Ca2+ entry in cells (2).

Three isoforms of the exchanger (NCX1, NCX2, and NCX3) have been cloned and are encoded by distinct genes (4–6). NCX1 is most abundant in heart but is widely distributed in most cells (5, 7). NCX1 is required for Na+-dependent Ca2+ extrusion and for protection against apoptosis in embryonic cardiomyocytes (8). NCX2 expression is restricted to the CNS, brain, and spinal cord (9). Evidence for an essential role of NCX2 in the clearance of residual Ca2+ in presynaptic terminals during hippocampal short- and long-term plasticity was recently reported (9). The expression of the NCX3 gene is restricted to brain and skeletal muscle (5–8).

In skeletal muscle fibers, the presence of NCX1 and NCX3 has been detected on the sarcolemma, but the exact roles played by these exchangers are still unclear (7, 8, 10). The expression of both exchanger proteins is developmentally regulated in rat skeletal myotubes, and their different subcellular distribution might be associated with specific physiological functions (8, 10, 11).

Received for publication April 17, 2003, and accepted in revised form November 19, 2003.

Address correspondence to: Sophie Sokolow, IRIBHM-IBMM, Université Libre de Bruxelles, rue des Professeurs Jeener et Brachet 12, B-6041 Gosselies, Belgium. Phone: 32-2-650-98-24; Fax: 32-2-650-98-20; E-mail: ssokolow@ulb.ac.be.

Conflict of interest: The authors have declared that no conflict of interest exists.

Nonstandard abbreviations used: electromyography (EMG); cytosolic free calcium concentration ([Ca2+]i); Na/Ca exchanger (NCX); neuromuscular junction (NMJ); vesicular acetylcholine transporter (VACHT); plasma membrane Ca2+ ATPase type 1 (PMCA1); flerox digitorum brevis (FDB); N-methyl-d-glucamine (NMGD); Evans blue dye (EBD); compound muscle action potential (CMAP); hypoxanthine phosphoribosyltransferase (HPRT); single-fiber electromyography (SFEMG); single-fiber action potential (SFAP); mean consecutive discharge (MCD); evoked endplate potential (EPP); spontaneous miniature endplate potential (MEPP); embryonic stem (ES); NCX3 immunoreactivity (NCX3-ir); sarcoplasmic reticulum (SR); 2,5-diter(butyl)-1,4-benzoquinone (TBQ); SR Ca2+ ATPase (SERCA); repetitive nerve stimulation (RNS); low rate of repetitive stimulation (LRRS); high rate of repetitive stimulation (HRRS).
Methods

Generation of Ncx3 knockout mice. Exon 2 of the murine Ncx3 gene (GenBank, accession number AF321404), which encodes for the first two-thirds of the protein, was cloned from a 129/Sv genomic phage library. A targeting vector was constructed by inserting the neomycin resistance cassette (neo) into the unique EcoRI restriction site of that exon. Standard procedures were used to generate Ncx3−/− mice. Experimental mice were 2 to 6 months old, of both sexes, and had F2 genetic backgrounds from 129/Sv and CD1 mice. Control mice consisted of age-matched littermates with two WT alleles at the Ncx3 locus (Ncx3+/+). All animal studies were authorized by the Animal Care Use and Review Committee of the Université Libre de Bruxelles.

RNA hybridization and RT-PCR. Messenger RNA (0.5 µg, MicrofastTrack 2.0 kit, Invitrogen, Carlsbad, California, USA) was extracted from gastrocnemius muscles. The RNA probe was an α32P [UTP]-radiolabeled 450 bp in vitro–transcribed DNA fragment (MAXscript, Ambion, Austin, Texas, USA). A GAPDH riboprobe was used to normalize the amount of RNA present on the membrane. OneStep RT-PCR was performed according to the manufacturer’s protocol (Qiagen, Valencia, California, USA). The primers pair was as follows: Ncx3, 5′-CAGTGGTTGTCTGGAAGCCGG-3′ (located in exon 2) and 5′-TGGGGGTGGTCACACCAGTACT-3′ (located in exon 9); hypoxanthon phosphoribosyltransferase (HPRT) served as a standard to balance the amount of RNA used, 5′-GCTGGTTAAGGACCCTCT-3′ and 5′-CACAGGACTAGAACCACCTGC-3′.

Primary antibodies. (i) Anti-mouse Ncx3 exchanger; peptide DGNFTPLEGEVKEDS from the mouse Ncx3 protein sequence was used for rabbit immunization (Eurogentec, Herstal, Belgium). This antibody was affinity purified and used at the dilution of 1:1000 for immunoblots, or 1:30 for immunocytochemistry. No cross-react was observed with NCX1. (ii) Anti-vesicular acetylcholine transporter (anti-VACht): rabbit anti-serum (Phoenix, Belmont, California, USA) was used at 1:1000. α-Bungarotoxin conjugated with Alexa Fluor 647 (Molecular Probes, Eugene, Oregon, USA) was used at 1:50. (iii) Anti-NCX1 and anti-plasma membrane calcium ATPase type 1 (anti-PMCA1): rabbit anti-sera anti-NCX1 and anti-PMCA1 (Swant, Bellinzona, Switzerland) were used at 1:1,000. (iv) Anti-desmin: mouse anti-desmin monoclonal antibody (Molecular Probes) was used at 1:100.

Secondary antibodies. (i) Peroxidase–conjugated goat anti-rabbit (Biosource International, Camarillo, California, USA) and sheep anti-mouse (Amersham, Buckinghamshire, United Kingdom) were used at the dilution of 1:25,000 for immunoblots. (ii) FITC or Texas Red–conjugated donkey anti-rabbit were used at the dilution of 1:200 (Jackson ImmunoResearch Laboratories, West Grove, Pennsylvania, USA).

Membrane preparation and immunoblot analysis. A supernatant containing membrane proteins of skeletal muscle homogenates was analyzed by SDS/PAGE. Detection by ECL Plus detection reagents (Amersham) was performed as described by the manufacturer. At least two different muscle homogenates were analyzed for each immunodetection. Protein loading was determined with a mouse anti-desmin monoclonal antibody (Molecular Probes).

Measurements of cytosolic [Ca2+]. Fibers from flexor digitorum brevis (FDB) muscles were isolated, and loaded with 1 µM Fura-PE3/AM (Calbiochem, San Diego, California, USA) according to De Backer et al. (12). Solutions used were previously described (13). To stimulate Ca2+ influx via Na/Ca exchange, all added Na+ to the solution was replaced with N-methyl-D-glucamine (NMDG) (13).

Histological and histochemical analysis. Gastrocnemius muscle cryosections (15 µm thick) were prepared according to Vanderwinden et al. (14) and processed for H&E staining.

Immunofluorescence. Double or triple immunofluorescence was performed according to Vanderwinden et al. (15). A minimum of two different preparations was used for each combination of markers. In some immunofluorescence experiments, nuclei were stained with 5 µM propidium iodide (Molecular Probes) in Tris-HCl 0.05 M (pH 7.4), containing 0.5 mg/ml ribonuclease A, for 2 hours in the dark at room temperature.

Confocal microscopy. Optical sections through regions of interest were collected sequentially for each fluorochrome as described (15).

Evans blue injection. Evans blue dye (EBD; Sigma-Aldrich, St. Louis, Missouri, USA; 10 mg/ml PBS, 50 µl/10 g body weight) was injected into the retro-orbital vein; 12 to 24 hours after injection, mice were killed, skinned, and visually inspected for blue coloration as described (16). Cryosections of gastrocnemius muscle were examined by confocal microscopy.

Electrophysiological investigations. Mice were anesthetized with chloral hydrate (loading dose: 0.4 mg/kg of body weight intraperitoneally, followed by administration of 0.4 µg/min using a microinjection pump) and fixed on a table (17). The sciatic nerve and the gastrocnemius were exposed as described (17). The stimulating cathode was a steel needle electrode inserted close to the proximal portion of the sciatic nerve at a distance of about 3 mm from the reference electrode (anode). EMG responses were amplified, filtered, and displayed using a NeuroMax 4 device (Xltek, Oakville, Ontario, Canada).

Recruitment curve. The sciatic nerve was stimulated at increasing intensities and the peak-to-peak amplitude of the compound muscle action potential (CMAP) of the gastrocnemius muscle was measured. The mean
CMAP was calculated at intensities from 0.05 to 10 mA (steps of 0.01 mA until maximal response, followed by steps of 0.5 mA) using a Digitimer DS7 stimulator (Digitimer, Hertfordshire, United Kingdom). Filters were set at 30 Hz–1.5 KHz.

Repetitive nerve stimulations. The CMAPs were recorded in response to direct supramaximal stimuli applied to the sciatic nerve (18). During each test, five stimuli were given, and this procedure was repeated twice (filters: 30 Hz–1 KHz). The fractional decrease (decrement) or increase (increment) in amplitude of the CMAP was calculated using the peak-to-peak amplitude of the first and fifth responses. The results of three test series were averaged and a positive decremental or incremental response was defined as a decrease or an increase of more than 10% of the fifth compared to the first response (19).

Postexercise facilitation. The testing program was the following: (i) a single repetitive stimulation at 10 Hz was recorded at rest (t0); (ii) the sciatic nerve was electrically stimulated at 10 Hz for 20 minutes (exercise); (iii) CMAPs were recorded after 10-Hz repetitive stimulations, immediately after exercise (t20), and every 30 seconds (t20+n) for 5 minutes; (iv) after a resting period of 30 minutes, the same animals were tested with the same program at 30-Hz stimulation.

Targeted disruption of the Ncx3 gene. (a) Structure of the WT and the targeted alleles. The second exon (shaded boxes), the neomycin resistance cassette (black box), the probes used in DNA and RNA hybridization analysis (black bars underneath the second exon), as well as the DNA fragments generated after digestion with KpnI + SpeI or KpnI alone are represented. K, KpnI; H, HindIII; S, SpeI; E, EcoRI; B, BamHI. (b) and (c) DNA hybridization analysis of KpnI + SpeI– (b) and KpnI– (c) digested genomic DNA isolated from ES clones using the depicted probe. (d) RNA hybridization analysis. Messenger RNA (0.5 µg/lane) was hybridized with Ncx3 or GAPDH RNA probes. (e) RT-PCR analysis. Ncx3 amplicon (~600 bp) and housekeeping gene HPRT amplicon (~250 bp) were simultaneously amplified from gastrocnemius muscle messenger RNA by one-step RT-PCR. (f, g, and h) Western blot analysis. (f) Membrane fractions (50 µg/lane) from gastrocnemius muscle were analyzed with anti-NCX3–specific antibody. (g) Immunodetection of membrane fractions from Ncx3+/+ (lanes 1–3) and Ncx3−/− (lanes 4–6) gastrocnemius muscle analyzed with anti-NCX1– and anti-PMCA1–specific antibodies. Amount of protein loaded: lanes 1 and 4, 50 µg; lanes 2 and 5, 15 µg; lanes 3 and 6, 5 µg. (h) Membrane fractions (20 µg/lane) from FDB muscle were analyzed with anti-NCX1–specific antibody. Protein loading in (f–h) was determined by using a mouse anti-desmin monoclonal antibody.

NCX activity in FDB muscle. [Ca2+]i was measured in FDB fibers loaded with Fura-PE3. (a) Response to extracellular Na+ removal (replaced by NMDG). Graphs are representative of [Ca2+]i transients from Ncx3+/+ (n = 7) and Ncx3−/− (n = 9) FDB fibers. (b) Response to caffeine and TBQ in the presence (Krebs) or in the absence of extracellular Na+. Graphs are representative of five experiments with Ncx3+/+ FDB fibers. (c) Response to caffeine and TBQ in Ncx3−/− fibers in the absence of extracellular Na+. The graph is representative of five experiments. (d) Comparison of the Na/Ca exchange activity in response to caffeine and TBQ in Ncx3+/+ (n = 7) and Ncx3−/− (n = 7) FDB fibers. Results are displayed as the mean (bold traces) ± SEM (thin traces).
**Stimulated single-fiber electromyography.** The cathode was inserted close to the proximal sciatic nerve, and a stimulated single-fiber electromyography (SFEMG) electrode (Medtronic 26G-40 mm, core diameter: 25 µm; Medtronic, Minneapolis, Minnesota, USA) was introduced into the gastrocnemius muscle using a micromanipulator. Single-fiber action potentials (SFAPs) were recorded using dedicated software at filter settings of 500 Hz–10 KHz. Amplifier gain was 200 µV–1 mV/division adjustable. Adequate current flow typically ranged from 0.2 to 2.7 mA. Single fibers were selected as described (20). For each fiber, one response corresponds to 50 consecutive SFAPs. These measures were repeated one to four times to calculate the mean consecutive discharge (MCD) of the potential latencies (17). In order to assess the relationship between MCD values and increasing discharge rates (21–23), stimuli were delivered successively at 10 and 20 Hz (WPI A310-A365 stimulator; World Precision Instruments, Hertfordshire, United Kingdom).

**In vitro microelectrophysiology.** Hemidiaphragm muscles with their associated phrenic nerves were isolated, and electrophysiological recordings were done according to Minic at al. (24). Evoked endplate potentials (EPPs) and/or spontaneous miniature endplate potentials (MEPPs) were recorded with intracellular microelectrodes filled with 3 M KCl solution (8–12 MΩ resistance) using conventional techniques (24). The motor nerve was stimulated via a suction electrode with current pulses of 0.1-ms duration and supramaximal voltage (typically 3–8 V) at different stimulating frequencies. Synaptic potentials were recorded in about four to six NMJs chosen randomly in each preparation. If necessary, µ-conotoxin GIIIB was used to prevent muscle contraction. The EPP amplitudes recorded on junctions treated with µ-conotoxin GIIIB were normalized to –75 mV membrane potential and corrected for nonlinear summation (25, 26). The quantal content of EPPs (m) was calculated at each junction by dividing the mean normalized and corrected EPP amplitude by the mean normalized MEPP amplitude.

**Twitch tension measurements.** Contractile responses of isolated mouse hemidiaphragms were elicited by stimulating the phrenic nerve with supramaximal current pulses (0.15-ms duration) at various frequencies (20, 40, 60, 80, and 100 Hz), as previously described (27).

**Endurance and coordination analyses.** Mice were tested blindly using littermate controls. The wire test was based on the taut wire test used by Gomez (18). For testing, mice were placed with front claws gripping the wire and observed for a period of 180 seconds of suspension. The chronograph was stopped any time the animal fell or reached a rest area and restarted when the animal was placed again on the wire. Number of successes (when an animal reached a rest area) and number of falls from the wire were recorded. The average scores for successes and falls were calculated as described (28). Each animal was tested only once. The Rota-Rod (7600; Ugo Basile, Comerio, Italy) rotated at a constant speed of 32 rpm. The mice were acclimatized to the treadmill by being placed on it three times the day before the experiment. Each mouse was placed on the rotating bar, and the duration the mouse remained on the bar was measured, up to a maximum of 300 seconds. Three consecutive sessions were conducted. After a rest period of 1 hour, the same mice were tested for endurance over the course of 2,000-second maximums.

**Statistics.** The mean ± SD and mean ± SEM were calculated using a statistical program (GraphPad Instat; GraphPad Software, San Diego, California, USA). A Student’s t test was used for simple comparison of means. For multiple comparisons, data were analyzed by ANOVA, followed by the Bonferroni multiple comparisons procedure.

**Results**

**Generation of Ncx3-deficient mice.** A mutant allele was constructed as described in Methods (Figure 1a). After electroporation with the targeting vector, the recombinant embryonic stem (ES) clones were identified by DNA hybridization (Figure 1, b and c) and used to produce chimeric mice. Transmission of the mutant allele produced Ncx3 heterozygous mice that were mated to generate NCX3 null mutants (Ncx3+/–). Ncx3+/– mice were viable and fertile; they appeared grossly normal and showed normal body weight (data not shown). The absence of NCX3 messenger RNA in hybridization assay and in RT-PCR (Figure 1, d and e), as well as the absence of NCX3 immunoreactivity (NCX3-ir) (Figure 1f) in Ncx3+/– gastrocnemius muscle, confirmed that the insertion of the neomycin resistance cassette in the second exon of the Ncx3 gene created a null mutation.

**Figure 3**

Histological alterations in Ncx3+/– muscles. H&E-stained sections from 3-month-old Ncx3+/– (a) and Ncx3–/– (b) gastrocnemius muscle: focus of fiber necrosis and cellular infiltrate in Ncx3–/– mice. Scale bar: 50 µm. Localization of EBD in cryosections of gastrocnemius muscles: 3-month-old EBD-injected Ncx3+/– (c) and Ncx3–/– (d) mice were examined after 12–24 hours. Red stained structures are nuclei, green stained areas are cytosol, and blue stained areas are EBD-positive fibers. Scale bar: 100 µm.
Reduction of Na/Ca exchange activity in Ncx3−/− FDB fibers.

In order to study Na/Ca exchange activity in muscle fibers, we examined Ca2+ signaling in FDB fibers. In Ncx3+/+ FDB fibers, a typical response to the superfusion of low-Na medium (where Na+ is replaced by NMDG), was observed: the withdrawal of extracellular Na+ was followed by a rise of about 7 ± 3 nM (mean ± SEM, n = 6) in [Ca2+], reflecting the reverse Na/Ca exchange activity (Figure 2a). When Na+ was reintroduced into the bath, basal [Ca2+], was restored, illustrating forward exchange activity. In Ncx3−/− FDB fiber, no reverse exchanger activity was observed under these conditions: [Ca2+], remained unchanged when the low-Na+ medium was superfused (Figure 2a). We then studied the forward mode of the exchanger after elevating [Ca2+], with 15 mM caffeine, which stimulates sarcoplasmic reticulum (SR) Ca2+ release, and 1 μM 2,5-di(tert-butyl)-1,4-benzoquinone (TBQ), which inhibits SR Ca2+ ATPases (SERCA) (Figure 2b). As expected (13), this treatment produced a transient peak in [Ca2+], followed by a very slow decline. This second phase was inhibited when extracellular Na+ was replaced by NMDG, suggesting that the decline was due to an efflux of Ca2+ through Na/Ca exchange (Figure 2b). In contrast to the observation made at a basal [Ca2+], where reverse Na/Ca exchange activity was absent (Figure 2a), a forward Na/Ca exchange activity was detected in Ncx3−/− FDB fibers incubated with caffeine and TBQ (Figure 2c). Indeed, like in Ncx3+/+ fibers, the secondary decline in [Ca2+], present in Ncx3−/− fibers was also suppressed in the absence of extracellular Na+. However, the Na/Ca exchange activity was significantly lower in Ncx3−/− than in Ncx3+/+ FDB fibers (Figure 2d); treatment of FDB fibers with caffeine and TBQ induced a peak of similar magnitude in Ncx3+/+ and Ncx3−/− cells, but the secondary phase was significantly slower in the latter ([Ca2+], after 275 s: 252 ± 20 nM and 326 ± 27 nM in Ncx3+/+ and Ncx3−/− mice, respectively; Student’s t test: P < 0.01, n = 7 in both groups). Basal [Ca2+], was not significantly different between control and mutant mice (53 ± 2 nM, and 59 ± 2 nM in Ncx3+/+ and Ncx3−/− fibers, respectively; Student’s t test: P > 0.5, n = 7 in both groups). The data correspond to the mean ± SEM.

Histological alterations in Ncx3−/− skeletal muscles. Cross-sections of muscle from Ncx3+/+ and Ncx3−/− mice were stained with H&E and examined by light microscopy (Figure 3, a and b). Foci of necrotic fibers, representing less than 10% of the total fibers, associated with mononuclear infiltrates were observed in Ncx3−/− muscles (Figure 3b). The presence of damaged fibers in Ncx3−/− muscles was confirmed with EBD. After EBD injection, inspection revealed in four of six Ncx3−/− mice a streaky blue coloration of some skeletal muscles, with marked individual variation in both intensity and extent (data not shown). EBD uptake in muscle fibers was confirmed by confocal microscopy (Figure 3, c and d). In any given muscle, EBD-positive fibers occurred either singly or in small clusters, and their number varied from animal to animal but was always inferior to 10% of the total fibers. Conversely, in Ncx3+/+ mice, no EBD uptake into skeletal muscles was observed by gross inspection (n = 6),

Absence of compensatory overexpression of NCX1 and PMCA1 proteins in Ncx3−/− gastrocnemius muscles. Since NCX1 and PMCA1 are also involved in the control of Ca2+ homeostasis and well expressed in skeletal muscles (10, 29), their expression in membrane protein extracts isolated from Ncx3+/+ and Ncx3−/− gastrocnemius muscle was analyzed by Western blotting. No compensatory overexpression was detected in Ncx3−/− gastrocnemius muscles: NCX1- and PMCA1-ir’s were very similar in both genotypes (Figure 1g). Similar results were obtained when analyzing the presence of NCX1 in Ncx3+/+ and Ncx3−/− FDB fibers (Figure 1h).
and less than 0.1% of total fibers was shown to be EBD positive by microscopic observation (Figure 3c).

**Immunolocalization of NCX3 in muscle and nerve fibers as well as at the neuromuscular junction.** Immunofluorescence studies were performed on 3-month-old Ncx3+/+ and Ncx3–/– gastrocnemius cryosections. NCX3-ir was detected in all muscle fibers in Ncx3+/+ but not in Ncx3–/– animals (Figure 4, a and b). In addition, NCX3-ir was also observed in nerve ramifications present between muscle fibers (Figure 4c). Confocal studies of cross-sectioned myofibers stained with NCX3 antisera, VAChT antisera (a marker for nerve terminals) and Alexa fluor 647-α-bungarotoxin (a specific probe for nicotinic acetylcholine receptors) indicated the presence of NCX3-ir at the NMJ (Figure 4, d–f).

**EMG alterations in Ncx3–/– mice.** An extensive EMG analysis was performed on Ncx3+/+ and Ncx3–/– mice. First, recruitment curve determination was performed. At increasing stimulus intensities, alterations in the recruitment curve recordings were detected: significantly reduced CMAPs were observed in Ncx3–/– mice, as compared with a decrement of 13.57 ± 0.43% in Ncx3+/+ mice at HRRS, where the graph is representative of one animal (Figure 5a). The slope of the curve was also significantly reduced in the mutant mice (P < 0.05 from 0 to 5 mA). The latency of the muscle fiber responses was identical in both genotypes (data not shown). Second, responses evoked in the gastrocnemius muscle with repetitive nerve stimulation (RNS) at 10 and 30 Hz were analyzed (Figure 5, b and c). At a low rate of repetitive stimulation (LRRS), a decremental CMAP was recorded in Ncx3–/– mice (mean decrement ± SEM: 11.78 ± 1.19%) and HRRS, where the graph is representative of one animal (Figure 5e). The recovery was quick.

The graphs showed the presence of NCX3-ir at the NMJ (Figure 4, d–f). In contrast, a marked increment was observed at HRRS in Ncx3–/– mice (Figure 5e). The recovery was quicker in the control group whatever the frequency. The ratio between the percent of change recorded at HRRS divided by the percent of change recorded at LRRS showed the prominent postexercise facilitation (Figure 5f). Finally, stimulated SFEMG was performed in Ncx3+/+ and Ncx3–/– mice. SFAPs were recorded following stimulation, and the MCDs of the potential latencies were calculated. Sixty-seven responses were recorded from 23 Ncx3+/+ single fibers, and 88 responses from 22 Ncx3–/– single fibers. MCDs were significantly different between Ncx3+/+ (range: 30–84 µs; mean ± SEM = 58.8 ± 0.6 µs) and Ncx3–/– mice (range: 47–130 µs; mean ± SEM = 82.3 ± 0.8 µs; intergroup comparison of MCDs expressed as at the neuromuscular junction (LRRS), a decremental CMAP was recorded in Ncx3–/– mice (mean decrement ± SEM: 11.78 ± 1.19%) and HRRS, where the graph is representative of one animal (Figure 5a). The slope of the curve was also significantly reduced in the mutant mice (P < 0.05 from 0 to 5 mA). The latency of the muscle fiber responses was identical in both genotypes (data not shown). Second, responses evoked in the gastrocnemius muscle with repetitive nerve stimulation (RNS) at 10 and 30 Hz were analyzed (Figure 5, b and c). At a low rate of repetitive stimulation (LRRS), a decremental CMAP was recorded in Ncx3–/– mice (mean decrement ± SEM: 11.78 ± 1.19%) and HRRS, where the graph is representative of one animal (Figure 5e). The recovery was quick.

**Figure 5**
Electromyographic findings in Ncx3–/– mice. (a) Recruitment curves of muscle responses evoked in the gastrocnemius muscle by stimulation of the sciatic nerve at increasing intensities. Mean CMAPs obtained in Ncx3+/+ (n = 5; filled circles) and Ncx3–/– (n = 5; open circles) mice are illustrated. (b and e) Effects of repetitive stimulation of the sciatic nerve at 10 (b) and 30 Hz (e) in Ncx3+/+ and Ncx3–/– mice. The graphs show representative of all the mice tested in the same group, except for Ncx3–/– mice at HRRS, where the graph is representative of one out of the 4/8 mice with an incremental CMAP. (d and e) Time course of the fifth response amplitude/first response amplitude ratios during LRRS (d) and HRRS (e) before and after exercise (20 minutes of repetitive stimuli) in Ncx3+/+ (n = 5; black bars) and Ncx3–/– (n = 5; gray bars) mice. (f) Ratios of the values illustrated in (d) and (e) (filled circles, Ncx3+/+ mice; open circles, Ncx3–/– mice). (g and h) MCDs expressed in microseconds and blocking rates expressed in percent in Ncx3+/+ mice (n = 22 fibers; MCDs, filled circles; blocking rate, open circles) and Ncx3–/– mice (n = 23 fibers; filled triangles, MCDs; open triangles, blocking rate). Statistics (in a, d–h): mean ± SEM; *P ≤ 0.05; **P ≤ 0.01.
difference, $P < 0.01$) (Figure 5, g and h). Moreover, although 5 of 23 Ncx3−/− fibers presented an abnormal neuromuscular blocking (blocking rate ranged from 6 to 25%), none of the Ncx3+/− fibers showed a blocking rate higher than 5% (Figure 5, g and h). In order to assess the relationship between MCD values and increasing firing rates (21–23), 21 fibers from Ncx3+/− mice and 9 fibers from Ncx3+/+ mice were stimulated successively at 10 and 20 Hz, respectively. A significant decrease of the mean MCDs from 77.2 to 64.4 µs was observed in Ncx3−/− fibers when stimulated at 10 and 20 Hz, respectively. By contrast, at the same frequencies, the mean MCDs increased from 43.6 to 75.9 µs in Ncx3+/− fibers (ANOVA, $P < 0.01$, genotype X frequency).

Electrophysiological alterations in isolated Ncx3−/− neuromuscular preparations. The NMJ of Ncx3−/− mice was explored with electrophysiological techniques in order to detect eventual changes in synaptic parameters. Spontaneous quantal transmitter release, measured as the frequency of MEPPs in a standard physiological solution, did not differ significantly in Ncx3−/− and Ncx3+/− NMJs ($n = 8$, Student’s $t$ test: $P > 0.3$) (Figure 6a). In Ncx3−/− junctions, the mean amplitude of MEPPs was $0.86 \pm 0.11$ mV ($n = 8$), whereas in Ncx3+/− muscles it was $0.95 \pm 0.16$ mV ($n = 8$, four different muscles; Student’s $t$ test: $P > 0.2$).

To test the eventual presynaptic effects of the absence of NCX3, the quantal content of EPPs was determined in Ncx3+/− and Ncx3−/− NMJs. In isolated hemidiaphragms equilibrated in standard physiological solution supplemented with 2.2 µM µ-conotoxin GIIIB, nerve stimulation at a low rate (0.1 Hz) evoked full-sized EPPs (Figure 6b, left panel). In Ncx3−/− junctions, the mean amplitude of these EPPs was $25.7 \pm 1.8$ mV ($n = 4$), and their mean quantal content was $32.6 \pm 2.5$ ($n = 4$). These values did not differ significantly from Ncx3+/− junctions (Student’s $t$ test: $P > 0.1$). Thus, under the conditions tested, the basic quantal release parameters were not significantly different in Ncx3+/− and Ncx3−/− NMJs. However, further experiments in Ncx3−/− NMJs revealed that EPP amplitudes during continuous stimulation depended on nerve stimulation frequency and pulse number in a given train.

Indeed, at 10 Hz nerve stimulation, there was an initial decrease of EPP amplitudes that attained a plateau over an expanded time frame (Figure 6b, right panel). When analyzed as a function of time or pulse number, the decline of EPP amplitudes was found to be greater at 20 Hz than at 50 Hz stimulation frequency (data not shown).

These results are in marked contrast to Ncx3+/− NMJs, in which the decline of EPP amplitudes was greater at high than at low nerve stimulation frequencies. A comparison of EPP amplitudes evoked at 50 Hz in Ncx3+/− and Ncx3−/− NMJs also revealed that EPPs were initially slightly facilitated and thereafter depressed in the Ncx3−/− NMJ, whereas in the Ncx3+/− NMJ, facilitation was more marked and lasted longer; several peaks of facilitation were observed superimposed to the EPP depression. These results indicate that transmitter mobilization and synaptic vesicle turnover were differently affected in Ncx3+/− and Ncx3−/− nerve terminals following high-frequency stimulation. This was further substantiated by the finding that delayed transmitter release, measured as the number of quantal releases in the 100 ms after a train of nerve stimulation at 50 Hz, was significantly and markedly increased in Ncx3+/− NMJs ($11.4 \pm 1.8$ Hz; $n = 3$) when compared with Ncx3−/− NMJs ($2.2 \pm 0.4$ Hz; $n = 3$; Student’s $t$ test: $P < 0.001$). The data correspond to the mean ± SD.

Increased tetanus/contraction ratio at high-frequency nerve stimulation in Ncx3−/− nerve-muscle preparations. Analysis of contractile responses of isolated hemidiaphragms elicited by phrenic nerve stimulation indicated that both Ncx3+/− and Ncx3−/− nerve-muscle preparations can maintain nerve-evoked tetanic contraction at frequencies between 40 and 100 Hz, respectively, for 1 second without decline. However, the tetanus-twitch ratio was significantly higher in Ncx3−/− than in Ncx3+/− muscles (Figure 6d, $P < 0.001$; $n = 3$).

Decreased endurance and increased fatigue in Ncx3+/− mice. Ncx3−/− mice performed poorly in the wire and the Rota-Rod tests compared to Ncx3+/− mice. Indeed, after 100 seconds spent on the wire, reduction in activity (“average success score”: $1.4 \pm 0.4$ and $4.3 \pm 0.8$ for Ncx3+/− and
Ncx3+/− mice, respectively; Student’s t test: P < 0.01) and weakness of forelimbs muscles (“average falling score”: 7.1 ± 0.7 and 9.5 ± 0.3 for Ncx3+/− and Ncx3+/− mice, respectively; Student’s t test: P < 0.01) were significantly different between control and mutant groups (Figure 7, a and b). Whatever the Rota-Rod trial, Ncx3+/− mice stayed for a shorter mean duration on the rotating bar than Ncx3+/− mice (ANOVA, P < 0.001, genotype X session) (Figure 7c). Moreover, individual comparison of the Rota-Rod endurance assay confirmed that Ncx3+/− mice readily fatigue in comparison to their littermate controls. Ncx3+/− mice spent 20.3 ± 4.5 minutes on the Rota-Rod vs. 2.4 ± 1.4 minutes for Ncx3+/− mice (Student’s t test: P < 0.01). The data correspond to the mean ± SEM.

Discussion

We have generated a mouse strain with null alleles in the gene coding for NCX3, the third isoform of the Na/Ca exchanger, which is mainly expressed in brain and skeletal muscles (5). The absence of NCX3 mRNA in RNA hybridization and RT-PCR studies as well as of NCX3 protein in Western blotting and immunofluorescence analysis in skeletal muscles confirmed the null mutation in Ncx3+/− mice.

The absence of NCX3 in skeletal muscle fibers was associated with necrosis and cellular infiltrates. Our results indicated that necrosis was not the consequence of an increased basal [Ca2+]. This was expected since the exchanger had a low affinity for Ca2+ (Km = 3 µM) (30) and therefore had a minor influence on resting (low) [Ca2+]. On the contrary, the plasma membrane Ca2+ pumps had a higher affinity for Ca2+ but a lower maximal rate of transport and were therefore more suitable for regulating [Ca2+]i in the low concentration range (e.g., at rest) (31). Balnave and Allen have shown that the forward-mode Na/Ca exchange contributes to the removal of Ca2+ from fibers stimulated with TBQ and caffeine (13). This was confirmed here, and we showed that the absence of NCX3 results in a significantly lower Na/Ca exchange activity and a prolonged secondary decline in [Ca2+]i, emphasizing the high-transport capacity of this exchanger. The contribution of Na/Ca exchange to Ca2+ efflux may become crucial in conditions of high and sustained [Ca2+]i, as during fatiguing stimulation (32) or after repeated eccentric exercise (33). In these conditions, the absence of NCX3 could lead to altered Ca2+ homeostasis and fiber necrosis. The low extent of necrotic lesions in Ncx3+/− skeletal muscles (always <10% of total fibers) may result from the quantitatively low intrinsic contribution of NCX3 to the control of Ca2+ efflux in skeletal fibers and/or from the presence of other proteins, which may partially compensate for the absence of NCX3. Among these are NCX1 and PMCA1, two proteins involved in Ca2+ extrusion, which are well expressed in skeletal muscles and, in the case of NCX1, responsible for major cardiac muscle cell necrosis when lacking in mice (8). However, no increased NCX1 or PMCA1 expression was observed in Ncx3+/− muscles.

In response to low Na+-containing medium, a rather small reverse exchange was detected in Ncx3+/− skeletal fibers. In contrast, no Ca2+ influx was observed in Ncx3+/− fibers in the same condition, suggesting that in these cells, the small reverse exchange is exclusively dependent on NCX3. These unexpected results also suggest that in skeletal muscle fibers, NCX3 is the predominant functional isoform and that NCX1 activity has a strong dependence on [Ca2+]i, being inactive at low [Ca2+]i (50 nM; reverse mode) and operative at high [Ca2+]i (> 200 nM; forward mode).

For the first time, positive NCX3-ir was detected at the NMJ and in nerve ramifications between muscle fibers. Although a clear demonstration of the presence of NCX3 protein at the presynaptic level is currently lacking, our results indicate that NCX3 plays an essential role in the control of neuromuscular transmission. Indeed, major EMG abnormalities, including an incremental response and a prominent postexercise facilitation, were recorded at high-frequency nerve stimulation in Ncx3+/− muscle. The increment of the muscle fiber response was associated with a facilitation of EPP amplitudes and little depression of synaptic responses. During the 100-ms period after the high-frequency nerve stimulation, there was an elevated quantal delayed release in Ncx3+/− NMJs that was five to six times higher than in Ncx3+/− NMJs. SFEMG also revealed abnormal jitter, frequent neuromuscular blocks, and MCD improvement at higher discharge rate in the mutant mice.

Combined with NCX3 localization at the NMJ and the facilitation of the EMG and contractile responses at HRRS, our electrophysiological results suggest that Ca2+ entering motor nerve terminals may be cleared relatively slower in the absence of the NCX3, thereby enhancing residual Ca2+ and evoked and delayed quan-
tal transmitter release during RNS. Recent results from the analysis of mice lacking NCX2 support this hypothesis. Indeed, Jean et al. have recently observed a delayed Ca\(^2+\) clearance from Ncx2\(^{-/-}\) hippocampal pyramidal neurons following glutamate stimulation (9). The slower recovery of [Ca\(^{2+}\)] to basal level in presynaptic terminals of their mutant mice is associated with an enhanced short- and long-term potentiation (9).

According to Blaustein et al., Na/Ca exchange could reverse during membrane depolarization and contribute to Ca\(^{2+}\) entry in nerve cells (34, 35). We propose that in mice lacking NCX3, a decreased Ca\(^{2+}\) influx through the latter pathway may explain the low amplitude CMAPs and the decremental electromyographic responses observed at LRRS.

The main clinical features associated with impaired neuromuscular transmission are weakness and fatigability. Using wire test and Rota-Rod assays, we demonstrated that Ncx3\(^{-/-}\) mice have reduced activity, weakness of forelimb muscles, and readily fatigued in comparison to Ncx3\(^{+/+}\) mice. However, since NCX3 is also expressed in the central and peripheral nervous system, these symptoms cannot exclusively be attributed to alterations at the NMJ or the skeletal fiber and may also partially or totally result from defects in the nervous system.

In this study, we attempted to define the role of NCX3 in neuromuscular transmission and in skeletal muscles in vitro and in vivo using mice lacking NCX3. Our data indicate that Ncx3\(^{-/-}\) mice present an impaired neuromuscular transmission and a slight skeletal muscle fiber necrosis, highlighting the important role of this exchanger in motor nerve terminals and muscle fibers.

Acknowledgments

We thank K. Gillard, Huy Nguyen Tran, O. Giot, E. Marion, and S. Mellaerts for technical contributions and J.D. Fransson, J.M. Raymackers, A. de Kerchove, A. Kamagate, and Y. Sokolow for discussion. S. Sokolow was supported by the Belgian National Fund for Scientific Research (FNRS), the Belgian Telethon, and the Fondation Rose et Jean Hoguet and is a research fellow at the FNRS. S. Schurmans and J.-M. Vanderwinden are Senior Research Associates of the FNRS, and M. Manto is supported by the FNRS. This work was supported by the Belgium Telethon, a grant from the Université Libre de Bruxelles, by the Fonds de la Recherche Scientifique Médicale de Belgique (to S. Schurmans), by the Association française contre les myopathies (AFM) (to P. Gaillly and J. Molgó), by an Action de Recherche Concertée grant (00/05-260) from the General Direction of Scientific Research of the French Community of Belgium (to P. Gaillly) and by the Fondation Médicale Reine Elisabeth and the Fondation Universitaire David et Alice Van Buuren (to J.-M. Vanderwinden).