

# Two functionally distinct myosin heavy chain isoforms in slow skeletal muscle fibres

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**Abstract** The head part of the myosin heavy chain (MHC) represents the essential component of the molecular force-generating system of muscle [1–3]. To date, three fast but only one slow MHC isoforms have been identified in adult mammalian limb muscles [4,5]. We show here two functionally different slow MHC isoforms, MHCII $\beta$  and MHCII $\alpha$ , coexisting in a considerable fraction of slow fibres of rabbit plantaris muscle. The two isoforms exhibit distinct electrophoretic mobilities and different kinetic properties. Thus, as it is known for the fast muscle, also the slow muscle seems to use different MHC isoforms in order to fulfil different functional demands.

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**Key words:** Myosin heavy chain isoforms; Muscle fibre types; Slow skeletal muscle; Muscle mechanics

## 1. Introduction

Myosin contains the essential component of the molecular force-generating system of muscle [1–3]. In skeletal muscle it exists as distinct isoforms in a variety of fibre types best classified according to their myosin heavy chain (MHC) complement (for review ref. [6]). Accordingly, three fast fibre types have been identified in mammalian limb muscles: types IIB, IID(X), and IIA containing MHCIIb, MHCII $\alpha$ (x), and MHCII $\alpha$ , respectively [4,5]. A clear correlation exists between the kinetics of force responses following stepwise length changes of maximally Ca<sup>2+</sup>-activated muscle fibres and their MHC isoforms [7–9]. This correlation suggests distinct functional properties of the various MHC isoforms.

Only one slow fibre type (type I) has to date been characterized by its MHC complement in mammalian limb muscles. Its MHCI is thought to be identical to the  $\beta$ -cardiac MHC isoform [10]. Previous observations on the variable distribution of the two slow essential (alkali) light chains, LC1sa and LC1sb, in type I fibres pointed to the possibility that the type I fibres might not represent a homogeneous population [11,12]. This possibility is supported by recent findings on the electrophoretic separation of MHCI into two bands, i.e. the original MHCI, and MHCII $\alpha$ , a slightly faster moving band [13,14]. Because these two electrophoretically distinct protein bands might represent two slow MHC isoforms, we

have tried in the present study to characterize their identity in a combined micromechanical and biochemical approach. For this purpose, the kinetics of force responses following stepwise length changes [2] were studied in skinned single type I fibres, electrophoretically identified by their MHC isoforms [5].

## 2. Materials and methods

### 2.1. Mechanical measurements

The methods of the mechanical experiments and the apparatus are more extensively described in one of our earlier publications [15]. The experiments were performed on glycerinated skinned fibres from rabbit plantaris muscle at 22°C. The temperature of the bath solutions was continuously checked using an ALMENO digital thermometer with a micro thermocouple IT-21. The attachment points for the muscle fibre ends on the mechanical apparatus were two vertically orientated epoxy carbon fibres of 100  $\mu$ m tip diameter. They were glued on silicon plates of force transducer elements (AE 801, SensoNor, Norway). One element, the force sensor (resonance frequency  $10 \pm 0.5$  kHz), was connected mechanically to a micrometer screw and electrically to a force bridge amplifier. The other element was a dummy, glued on the lever arm of a stepping motor. Rapid changes of the fibre length ( $< 0.5$  ms) were carried out by a feedback controlled stepping motor based on a Ling vibrator. The ability to make rapid solution changes ( $< 0.2$  s) was provided by a cuvette transporting system. Laser diffractometry (He-Ne laser, 632.8 nm, 4 mW) was used to measure the sarcomere length of the myofibrillar bundles before, during, and after activation of the fibres.

The solutions contained 60 mM HEPES, 8 mM Na<sub>2</sub>H<sub>2</sub>ATP, 10 mM sodium creatine phosphate, 20 units/ml creatine phosphokinase, 40 g/l dextran T-500 and 1 mM free Mg<sup>2+</sup>. In addition, the relaxation solution (pCa  $> 9$ ) contained 50 mM EGTA. The activation solution (pCa 4.7) contained 50 mM Ca-EGTA, and the preactivation solution (low Ca<sup>2+</sup> buffering capacity, pCa 7) 50 mM hexamethylenediamine-N,N,N',N'-tetraacetic acid (HDTA). The pH was adjusted to 7.10 in all solutions at 22°C. The pCa ( $-\log[\text{Ca}_{\text{free}}^{2+}]$ ) of the solutions was determined with a Ca<sup>2+</sup>-selective electrode. The solution for storing the muscle fibres at  $-20^\circ\text{C}$  was a relaxation solution containing 50% (v/v) glycerol (pH 6.9).

After attachment of the skinned muscle fibres with the tissue glue Vetseal (Braun, Melsungen, FRG), the fibre ends were fixed by superfusion (3–5 s) with a fine, rapidly downward flowing stream of pigmented glutaraldehyde solution (8% (v/v) glutaraldehyde and 5% (w/v) Toluidine blue, fixative). For this purpose, the fibre was bathed in a low ionic strength rigor solution (10 mM imidazole, 2.5 mM EGTA, 7.5 mM EDTA, 134 mM potassium propionate, pH 6.8) with lower specific mass than the fixative. This fixation procedure created a sharp boundary between the functional part of the muscle fibre and the fixed fibre ends. Control experiments ([15], Hilber and Galler, unpublished) have shown that this method considerably improves the maintenance of sarcomere order and the stability of the mechanical properties during prolonged activations.

The length of the preparations was always adjusted to exactly the slack position (final active fibre length, 2.5–3.5 mm) and the sarcomere length was recorded (2.0–2.4  $\mu$ m). Subsequently, the fibre was treated with 1% (v/v) Triton X-100 in a relaxation solution (15 min; 22°C). After transferring the fibre from the preactivation solution to the maximal activation solution, a series of quick ( $< 0.5$  ms) stretches and releases (range 0.5 to 6 nm per half sarcomere) was applied to

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**Abbreviations:** ATP, adenosine 5'-triphosphate; HEPES, 4-(2-hydroxyethyl)-1-piperazineethane-sulfonic acid; EGTA, ethylene-bis(oxyethylenetriolo)tetraacetic acid; MHC, myosin heavy chain; HDTA, hexamethylenediamine-N,N,N',N'-tetraacetic acid; pCa, negative logarithm of free Ca<sup>2+</sup> concentration

detect the time course of resulting force transients. Measurements were only taken for evaluation if the sarcomere length signal during and after maximal activation was unchanged and of comparable quality as under relaxed conditions.

### 2.2. Biochemical analysis

After completion of the mechanical measurements, the unfixed part of the muscle fibre was cut from the apparatus for biochemical analysis. The fibre fragment was dissolved in 11  $\mu$ l sodium dodecyl sulphate (SDS) lysis buffer (62 mM Tris-HCl, pH 6.8, 10% glycerol (v/v), 2.3% SDS (w/v), 5%  $\beta$ -mercaptoethanol (v/v), 19% sucrose (w/v)) and heated at 65°C for 15 min. A 2.5  $\mu$ l aliquot of this extract was applied to a 5–8% polyacrylamide gradient gel according to the method previously described [5]. Densitometric evaluation was used in order to quantify the ratio of different MHC isoforms in single fibres.

## 3. Results

Electrophoretic analyses on rabbit plantaris muscle fibres detected a considerable fraction of type I fibres (21 out of 37) displaying, in addition to MHC I, a band of slightly higher mobility (Fig. 1). According to [13] and [14], this band was designated as MHC Ia. The isoform with the lower electrophoretic mobility (the former MHC I) was designated as MHC Ib. According to densitometric evaluation of the gels, MHC Ib represented the major isoform in all type I fibres investigated in plantaris muscle. The relative concentration of MHC Ia maximally amounted to 30% of total MHC isoforms. Fibres containing only MHC Ib were termed type Ib; fibres containing MHC Ib > MHC Ia were designated as type Ib/Ia.

Quick (<0.5 ms) stretches and releases of 0.5–6 nm per half sarcomere were applied to maximally  $\text{Ca}^{2+}$ -activated skinned muscle fibres ( $187 \pm 79 \text{ kN cm}^{-2}$ ,  $n = 37$ ) in order to detect differences which could be correlated to their MHC isoform complement. As in studies on rat muscle fibres [7,8], the kinetics of force responses during fibre stretches correlated closely with the kinetics during fibre releases. However, only stretch response measurements are presented here because of the greater precision with which they can be detected. A stretch of the muscle fibres induced an instantaneous force rise followed by a decay and a secondary (delayed) transient force increase (i.e. stretch activation; Fig. 2A). For evaluation, the time from the beginning of the stretch to the peak value of the delayed force increase ( $t_2$  value) was chosen. This parameter is representative for other kinetic parameters of force responses following stepwise length changes [8], and, because of its independence of the length change amplitude [7,16], it is one of the most useful parameters to distinguish

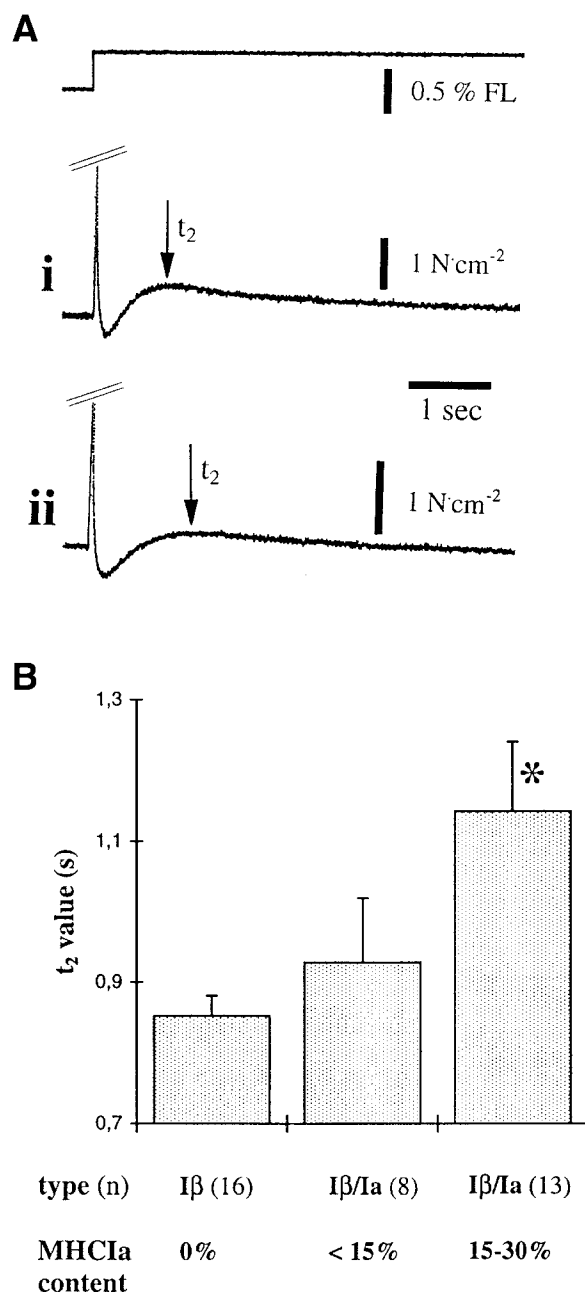


Fig. 2. (A) Original recordings of fibre length (upper panel) and force (lower panels) during quick stretch experiments (2 nm per half-sarcomere) on maximally  $\text{Ca}^{2+}$ -activated skinned rabbit muscle fibres. (i) represents a fibre containing only the MHC Ib isoform (former MHC I); (ii) represents a fibre which contains both the MHC Ib and the MHC Ia isoform. (B)  $t_2$  values (mean  $\pm$  SE) of type Ib and Ib/Ia fibres. The type Ib/Ia fibres are divided into two groups depending on their MHC Ia content. \* means significantly different from type Ib fibres ( $p < 0.01$ , unpaired  $t$ -test).

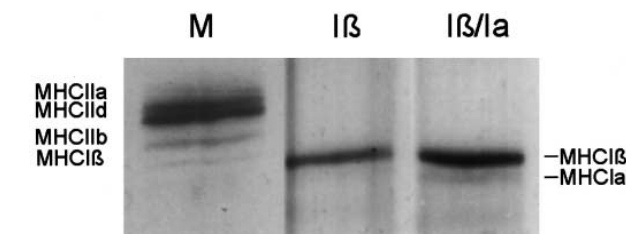


Fig. 1. Silver-stained polyacrylamide gradient gel electrophoreses of MHC isoforms from rabbit plantaris slow muscle fibres for which mechanical properties were previously measured. The label at the top of each lane represents the fibre type. The left lane is an electrophoresis of rabbit gastrocnemius muscle and serves as a marker (M).

between fibres with different MHC isoforms. Type Ib fibres ( $852 \pm 29 \text{ ms}$ , SE,  $n = 16$ ) exhibited significantly ( $p < 0.01$ , unpaired  $t$ -test) smaller  $t_2$  values than type Ib/Ia fibres ( $1067 \pm 75 \text{ ms}$ ,  $n = 21$ ). This difference is remarkable since the mean of the relative MHC Ia content in type Ib/Ia fibres was rather small (15% of total MHC content). The differences of the  $t_2$

values are further displayed in Fig. 2B where the type I $\beta$ /Ia fibres were divided into two groups depending on their MHC Ia content.

#### 4. Discussion

Subtypes of fast muscle fibres display a strong correlation between the MHC isoform complement and the kinetics of force responses following stepwise length changes. This is most conspicuous with regard to the  $t_2$  value of stretch activation [7,8,16]. In hybrid fibres containing two fast MHC isoforms, the  $t_2$  values depend on the relative proportion of each isoform [7,16]. The present results indicate that a similar relationship exists in slow fibres exhibiting both the MHC I $\beta$  and MHC Ia bands. It appears, therefore, that the MHC Ia band truly represents an additional slow myosin heavy chain isoform associated with specific functional properties. This novel finding indicates that a considerable fraction of slow muscle fibres contains two functionally distinct MHC isoforms.

The MHC Ia isoform characterised in the present study appears to be distinct from the  $\alpha$ -cardiac MHC isoform because the latter has a lower electrophoretic mobility than the  $\beta$ -cardiac isoform. As judged from its electrophoretic mobility, the MHC Ia also seems to differ from the  $\alpha$ -cardiac-like MHC isoform detected in the diaphragm and chronically stimulated fast-twitch muscles of rabbit [17]. Furthermore, the  $\beta$ -cardiac MHC isoform (predominantly present in cardiac ventricle) displays slower functional characteristics than the  $\alpha$ -cardiac MHC isoform [18]. Taken together, our data suggest that MHC Ia is a new MHC isoform with slower properties than MHC I $\beta$  (or the  $\beta$ -cardiac MHC).

Force responses following stepwise length changes may be related to the elementary process of force generation [2,19–21]. According to this interpretation, our results suggest that the two slow MHC isoforms exhibit different kinetic properties during the force-generating process. Thus, mammalian slow muscle fibres represent a heterogeneous population composed of fibres characterised by the expression of one or two MHC isoforms and the kinetic properties associated with each

isoform. This presumably enables slow muscles to fulfil differing functional demands.

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