

The complete nucleotide sequences of cDNA clones coding for human myosin light chains 1 and 3

Ulla Seidel, Eva Bober[§], Barbara Winter⁺, Susan Lenz, Peter Lohse and H.H.Arnold*

Department of Toxicology, Medical School, University of Hamburg, Grindelallee 117, 2000 Hamburg 13, [§]Institut für Humangenetik, Universität Hamburg, 2000 Hamburg, FRG and ⁺Department of Biochemistry, National Cancer Institute, Bethesda, MD, USA
Submitted May 15, 1987

Accession no. Y00352

Full length clones for human fast fiber type myosin light chains 1 and 3 were isolated from a fetal skeletal muscle cDNA library cloned in lambda GT 11. The nucleotide sequences were determined in M13 subclones and proven to be identical except for 201 and 79 nucleotides at the 5' end of LC1 and LC3, respectively. The coding sequences are approximately 80 % homologous to chicken (1), mouse (2) and rat (3). For these organisms it had been shown that both mRNAs are generated from two separate transcripts of one gene by an alternative splicing pathway which utilizes two different exons each for the N-terminus and five common exons for the body and C-terminus of the proteins. A similar arrangement is suggested for man by the presented cDNA sequences.

<p>1 MLC 1 TCAAAGCCAGTGCTCTGAAGATGGACCCCTTTAAAAAAGAAATGGCACAAA 55 GAAAGACGTGAAGAAACCTGGCGCTGGCGCTGCCACGCCCGCACCGCCACCC 115 TGCACCTGGCCCTGCCCCAGGCCAAACCCAAAGAAGAAAAATGACCTCTGCCTAA 1 MLC 3 GAGCCTAACTACTCTTAGCCTCCCTGCTGTGTTGCTGCCACGCCCTCCAT 175 GATGGAGTTCTCTAAGGAACAGCAGGATGAATTCAGAGGAGCAATTCCTCTTTGACAG 53 CATGTCTCTCAGTCTGACACAGATTCGCTGAATTCAGAGGAGCAATTCCTCTTTGACAG 235 AACAGGTGATTCCAAGATCACTTTAAGCCAGGTCGGTGAATGCTCTCGAGCCCTGGGCAC 113 AACAGGTGATTCCAAGATCACTTTAAGCCAGGTCGGTGAATGCTCTCGAGCCCTGGGCAC 295 AAATCCCAACCAATTCAGAGGTCAGGAAAGTTCTGGAAAACCCCAACCAATGAAGAGCTCAA 173 AAATCCCAACCAATTCAGAGGTCAGGAAAGTTCTGGAAAACCCCAACCAATGAAGAGCTCAA 353 TGCACGAAAAATGAGCTTTGAACAATTTCTGCCTATGATGCAAGCCATTCACCAACA 215 TGCACGAAAAATGAGCTTTGAACAATTTCTGCCTATGATGCAAGCCATTCACCAACA 413 GGCACAGCCACCTATGAAAGCTTTCTGAGGCTCTGCGCTGTCTTTGACAAAGGAGCCAA 293 GGCACAGCCACCTATGAAAGCTTTCTGAGGCTCTGCGCTGTCTTTGACAAAGGAGCCAA 473 TGGCAGCATCTAGGGTGCCTGAACCTCCGCCATGCTTAGCCACCCCTGGTGAAGAGTGA 553 TGGCAGCATCTAGGGTGCCTGAACCTCCGCCATGCTTAGCCACCCCTGGTGAAGAGTGA</p>	<p>535 AGAGGAAGAAGTGGAAAGCCCTGATGGCAGGTCAAGAAAGCTCCAAATGGCTGCATCAACTA 413 AGAGGAAGAAGTGGAAAGCCCTGATGGCAGGTCAAGAAAGCTCCAAATGGCTGCATCAACTA 695 CGAAGCTTTTGTCAAGCACATCACTCTAATCTGAAATGGAGCTCTCAAGAAACAGCATTT 473 CGAAGCTTTTGTCAAGCACATCACTCTAATCTGAAATGGAGCTCTCAAGAAACAGCATTT 655 TTAGGAAGACTGGCTGGAAACTTATTTAATCAACCCATGACAAACTTCCAGATCTGT 593 TTAGGAAGACTGGCTGGAAACTTATTTAATCAACCCATGACAAACTTCCAGATCTGT 715 TTACCATCATTCAGGAAAAAACAAGCAATCTGGACGGTTCAAGACTCAGCAACTCCCTGAA 593 TTACCATCATTCAGGAAAAAACAAGCAATCTGGACGGTTCAAGACTCAGCAACTCCCTGAA 775 TTTTATACATCTTCAGTTTTCTCTGAAATGAAATTCATACACAAACAATGTCCTCC 653 TTTTATACATCTTCAGTTTTCTCTGAAATGAAATTCATACACAAACAATGTCCTCC 835 TGCTCTAGATGAGAAAGATAAAATATTGACAACTCAAATCAAAGCAGCCTCTTTATTA 713 TGCTCTAGATGAGAAAGATAAAATATTGACAACTCAAATCAAAGCAGCCTCTTTATTA 895 TCTACCATGAATCAACGAAACACTCTTAAACAATAAATCAATAAACAATTTGGTCACT 773 TCTACCATGAATCAACGAAACACTCTTAAACAATAAATCAATAAACAATTTGGTCACT 955 CTGAAAAAAAAA 833 CTGAAAAAAAAA</p>
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Acknowledgement: We thank J. Bergmann for excellent technical help

*To whom correspondence should be addressed

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