# Myoskeletin, a factor related to Myocardin, is expressed in somites and required for hypaxial muscle formation in *Xenopus*

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ABSTRACT *Myoskeletin* was identified as a gene induced by activin in animal cap explants of *Xenopus laevis*. This gene encodes a protein related to the transcription factor Myocardin. Whereas Myocardin is expressed in the heart and is known to be involved in heart and smooth muscle formation, Myoskeletin is expressed in the somites and in hypaxial muscle precursors as they migrate away from the somites during tadpole stages. Myoskeletin is required for hypaxial muscle formation, as reduction of its expression through injection of an antisense morpholino oligonucleotide leads to suppression of hypaxial muscle formation. In overexpression experiments in animal caps, Myoskeletin is capable of inducing multiple genes including skeletal muscle, cardiac muscle and smooth muscle-specific genes. We conclude that Myoskeletin is a somite and hypaxial muscle-specific member of the Myocardin family that is required for hypaxial muscle formation.

#### KEY WORDS: myocardin, skeletal muscle, hypaxial muscle, somite

In the early development of the Xenopus laevis embryo, mesodermal cells are specified through inductive processes that depend on cell-cell signaling factors. Signaling by TGF- $\beta$ -related factors of the nodal/activin group is essential in this process (Heasman, 2006; Wardle and Smith, 2006; Whitman, 2001). Some aspects of the complex processes that lead to mesoderm specification and patterning can be recapitulated in a culture system in which explants derived from the animal region of the blastula, so-called animal caps, are exposed to activin. While untreated animal caps develop as atypical ectoderm, exposure to activin leads to differentiation of multiple mesodermal derivatives (Asashima et al., 1990; Smith, 1987; Smith et al., 1990). We have made use of this system to survey the global changes in gene expression that take place during mesoderm induction by activin with the aid of the DNA microarray technology. Among the genes whose expression is strongly induced by activin in the animal cap system we identified a novel member of a transcription factor family that plays a role in the differentiation of muscle, a major derivative of the mesoderm.

Muscle formation has been extensively studied in multiple organisms. Several classes of transcription factors are involved in the process of the specification and differentiation of this tissue

and in the elaboration of the various muscle subtypes such as skeletal, cardiac and smooth muscle. The joint action of varied combinations of factors from different families in the regulation of many muscle-specific genes provides a prime example for the combinatorial principle in gene regulation in general (Davidson, 2006). Several members of the basic helix-loop-helix (bHLH) class of factors are involved in muscle differentiation, including MyoD, Myf5, myogenin and MRF4 (Berkes and Tapscott, 2005; Pownall et al., 2002; Tapscott, 2005). While early studies were strongly focused on the bHLH class of factors, other transcription factor families are likewise essential in this process. Mef2, a MADS-box (MCM-1, Agamous, Deficiens, SRF) family protein, is required for muscle formation and specifically for the transcriptional regulation of multiple muscle-specific genes (Nava and Olson, 1999). In addition, SRF (Serum Response Factor) has an important role in muscle formation, as reported in the Xenopus system some time ago (Mohun et al., 1991). As the ubiquitous expression of SRF raises questions about its specific role in muscle cell differentiation, it was particularly interesting when a

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*Abbreviations used in this paper*: bHLH, basic helix-loop-helix; MyoC, myocardin; MyoS, myoskeletin; SRF, serum response factor.

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cell type specific partner protein of SRF was discovered. Myocardin (MyoC), as this factor was named, binds SRF, leads to strong gene activation and has a specific role in the differentiation of cardiac and smooth muscle cells (Chen et al., 2002; Small et al., 2005; Teg Pipes et al., 2006; Wang et al., 2001; Wang and Olson, 2004). Clearly, SRF also has a role in skeletal muscle differentiation, but MyoC is not expressed in these cells and two related proteins, MRTF-A and MRTF-B, also do not appear to have a specific role in skeletal muscle (Teg Pipes et al., 2006). Here we report the identification, among multiple genes induced by activin in animal caps, of an additional member of the MyoC family with strong and specific expression in skeletal muscle. During progress of this work we learned of related experiments by S. M. Meadows, A. S. Warkman, E. M. Small and P. A. Krieg (personal communication) and in agreement with these authors we name this protein Myoskeletin (MyoS). Our results suggest that MyoS is critical for the formation of the hypaxial muscles, a set of muscles that derive

from the somites and migrate to form the body wall and limb musculature (Buckingham et al., 2003). The discovery of MyoS fills an apparent gap in our understanding of the manner in which SRF controls the differentiation of various types of muscle cells.

#### Isolation and characterization of MyoS

In an effort to identify novel genes involved in the induction of mesodermal tissues in the Xenopus embryo we compared the transcriptome of control animal caps with that of animal caps that were treated with activin, using DNA microarray technology. A large number of genes were upregulated in this experiment, among them a gene with sequence similarity to the previously identified Xenopus MvoCgene (Small et al., 2005): the new gene is named MyoS (see above). The increase in MyoS expression after activin induction was about 20-fold in four repeats of the microarray assay.

The predicted MyoS protein shows highest overall sequence similarity to Xenopus MyoC, with somewhat lower similarity to the related proteins MRTF-A and MRTF-B (Table 1). An alignment of the four MyoC family members in Xenopus is shown in Figure 1. Much of the sequence identity is concentrated in putative conserved domains including basic, glutamine(Q)-rich, SAP (SAF-A/B, Acinus and PIAS) and leucine zipper domains; the basic and Q-rich domains are embedded in a larger region of high sequence conservation. MyoS contains most of the conserved functional domains of the MyoC family including the basic and Q-rich domains that mediate interactions between MyoC and SRF (Teg Pipes et al., 2006). MyoS thus is likely to represent a new functional member of the Myocardin factor family.

#### Embryonic expression of MyoS

MyoS RNA is present as a maternal component in the egg, decreases during early to mid

gastrula and subsequently increases through tailbud and tadpole stages (Fig. 2A). The maternal MyoS expression does not show a distinct pattern so that in situ hybridization through gastrulation is not effective in visualizing this RNA (Fig. 2B). Staining becomes intense in the forming somites during neurula stages and intensifies in somites and presomitic mesoderm at least up to stage 41 (Fig. 2C-E). At about stage 37/38, MyoS expression is also seen clearly in the hypaxial muscles and in discrete regions in the head (Fig. 2E). The hypaxial muscles, which will form the body wall and limb musculature, arise in Xenopus from about stage 26 in the ventro-lateral areas of the somites, become more clearly defined by stage 28/29 and begin to migrate from the somites during subsequent development (Martin and Harland, 2001; Martin and Harland, 2006). Because of the strong expression of MyoS in the somites, initial stages of its expression in hypaxial precursors cannot be distinguished, but MyoSRNA is clearly visualized when these cells leave the somites (Fig. 2D). The expression pattern of

xMyoS_ xMyoC_ xMRTF-A xMRTF-B	60 
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	MSERNLIPSPVVKKTPAPFPEQNGNLGQNKVEDFMKIKGQNKLHKVAALKMHFPED LVSQGLMP-PLKSPAAFAQEQRKNVDRAKAEDYLKHKIRSRPEILNMQILQDPANE LENQGIMP-PLKSPAAFHEQRRSLERARTEDYLKRKIRSRPERAELVRMHILEETSAE LVDQGIMP-PLKSPAAFHEQIKSLERARTENFLKHKIRSRPNRSELVRMHILEETLAE *****
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	180 QLQISD SSAQAAQIKLKRARLADDLNERIALRPGPLELVEKNIIPVESTVKEVFKGNQVNFSKSVD PSLQAKQIKLKRARLADDLNEKISQRPGPMELVVKNILPVETSLKEVIIDVDYPEVVD PSLQATQLKLKRARLADDLNEKIAQRPGPLELVEKNILPVDLSVKEAITVSQTNLPENLD *
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	240 AFSFEDDISSSS-SSSTSSSPRFAPSPGLSLNLSPTSTNTVFQLDLPQIIEVNQPN AFAFEEDSSNDGLSPEKEPSENSPVLNKASFQETKDLETLN-SLHSGLTQNHSQEHDSDA NSSFDEDSS-DALSPE-QPASQESQGSIPSPIENRPSETTQIPALSPSHAFSCVQFGTDA TLSFDEDSS-DALSPE-QPASQESQGSAASPGEMKTSDSSS-PVSNTTIQCQTVSSPLPD ****
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	basic domain <u>3</u> 00 VRTV-TEAETLVTSRPATHNPTQASSTVPKVTVKPSDVG <mark>KIQRPKKPKDTKPKVKKLK</mark> YH QDGSSIQSHSCSLHSESQLSPSMSASAAVKSKSPIDVKN-RHKTKDIKPKVKKLKYH FNQDSLQSTAITISNGLTASICKSLPALVKQSQPKPSFEKSQRIKKPKEPKPKVKKLKYH FFKP-VPTADLTTRSPLSCIVSKPGPALIKQTQPK-HTEKP-RSKKSKDPKPRVKKLKYH
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	Q-rich domain 360 QYIPPDQKAEKAPVAMDAAYSRLLQQQQIFLQLQILNQQQN-PTFCVQTVHPLTTS-IPA QYIPPDQKAEKSPPPMDSAYARLLQQQQLFLQLQILSQQQQQQ QYIPPDQKGEKIEEEMDSNYARLLQQQQLFLQLQIINQQH-QHYNYQTILPAPPKPLPD QYIPPDQKGEKIEEEMDSNYARLLQQQQLVLQLQILSQQHSTLTTSRKSYPAPLKS ******** ** ** ** ** ****************
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	420 DQVISFTGAPPSSAPAINLSPAPGTAAVTTAPTSTVPSPMKTEMLPANIDDLTVSELRQH DLIRNS-NTSSVNSPSLSPVKTTFSGQAN-VSMKAGLLPSNLDDLKVSELRQQ QQNTNSSSTTTVRSMST-VAPSTLATPTITRQNSNVAVGGRTGPLPHNLDEMKVAELKLE QKKQNTINTTICNGNAG-APPAQCSVNRQNS-VPCK-KTGPLPSSLDDMKVAELKME * * * * * * * * * * * * * * * * * * *
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	SAP domain 480 LRKRGLPVSGTKPSLLERLRPYQIPRAKTIPAPIQSAGLMTPIIELSAFPKQSVCD LRIGLPVSGTKTSLMERLRPFQECSGN-TVPTYGEITTVTFPVTPNGTLSGYQSHASAG LKHRGLPVSGTKIDLIERLKASQDPSTATAASAKPT-PVQQAKPPEVVPIVSSSC LKLRGLPVSGTKMDLIERLKPFQDFSSNGVSPSSANTVNITNPACNTTDDATTAFSTSAL * ******** * * *** * * * *
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	540 STVPTLCTFQTVPSP-PSGEVPQETSET-ACSMPE MLSNGFYQFGSTSSTP-PISPASSDFS-VSGS-LPDTFSDG-PMSSPQ LTTREPIKLCSTSSTP-PGSPCPSEVSVVSMDEVSMISDALGETVACPVTQ INSSSPTPSVSIGNNQTMLDGINSPLPMSPTPSEQSNFSSEDTN-ITDTFAEILTMMSPS * * * * * * * * * * * * * * * * * * *
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	zinc finger <u>600</u> STEMPATVQDKDTPMQDIEDDHV <mark>LMEKQKVIENLTWKLKQEQKQ</mark> FGLHPSPIHLSAEESLMNSMNSGTYQVELEGIDAERDKMLVEKQKVINELTWKLKHEQRQ QVQ-QNPAAEKSPPDARDKDLMLREKDRQIEELTQRLKQKQEL QFMNTSPL-KVNEDSMGATPGN-TPNVELDAVEKDRKLQEKEKQIEELKRKLEQEQKL * *** *******************************

*MyoS* is thus distinct from that of other members of the family, none of which is expressed in somites and hypaxial muscles (Teg Pipes *et al.,* 2006). Further, the heart, which is the site of strong and specific *MyoC* expression (Small *et al.,* 2005), is entirely negative for *MyoS* at all stages examined.

#### Interference with MyoS expression inhibits hypaxial muscle formation

To obtain insight into the function of MyoS in embryonic development we used an antisense morpholino oligonucleotide (MO) designed to bind to the region surrounding the initiation codon of the protein (see Experimental Procedures). MyoS-MO or a control MO (C-MO) was injected into the equatorial region of both 2-cell stage blastomeres. Injection of 20 ng MyoS-MO into each blastomere led to absence or strong reduction in the formation of hypaxial muscles as visualized by the antibody 12/101 (Kintner and Brockes, 1984), whereas the C-MO-injected em-

	660
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	AEDLRVELEMHKRLKNRHKKEKLISRVHIKQEVD VEELRLQLQKRKVDPQDKQPASQHFFGVPIKQEDV VERLRQQLEQEKRTPQHSTDDQQALILAVKQEPLPL VEVLKKQLELEKRGQQQQPCSNVLLKMEPKHFNLQIKEETEAPDCQNSKQPVGSGGQILG ***
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	720 RTSPTSCCAKAQTEETPFLYTISNDRTELKPPI SSCPFAAKQMALKVQVNNVTEKLGVGRAQPITCLV TVDSINKKASSIVKQELNTAIICQQ
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	780 QEPQINANLNETYMVFCPPSCDLIGQDFELPMQITASPES-P NNQCMDASTQNSIMSSTFLSPQCS-PQHSPMGTTNSPQHISLPPSP SSGIEGKVDNSAGTKLVFTLTNPSSQLPEENRQIVLQKVPTP-PSSLHPNNSLP-KQE TAQLLLPLSIKAANGVQLSMVQAQPHTVNPAPAQLSTAAATTTT-LLSAPPKQSAPPTQD * * *
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	840 RSFEEELQEAIQKAQMALC NNHY-LASVSPNSQGDRHSGSPNVNNR-LHPAQAAGS VLLSCCALQNQKPALQLVPGTVLSLSSSNLQPMLN-MNGFQKWHGEALDSLQKQLV KFTPHLLNQNQQ-IRKLCPSATSGNVFSYPQNPVTAVPQSFSASISTSAQPQRSTQLTAV *
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	900 ESHFPPNQNNLPISYNEQIEKKGNKETAPPKSPKQITRS HNESPATPPQQPEPEPPHSIFLTHSSPQWSKNPPGYDEAMKQQPNS-CED QNGPTSLHEKSSTPPQLQQFIVQQHPLFSSPTTKSKDPPRYEEAIKQARNNPASQPEV
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	960 EHLMCTDELMHIDPVSNMIEHHSSAALVPQHGHFSKPLYCCSDA QQMDELLDVLIESGEIPANARDDRSCAQKHHQLNVPSANSNPS-TSNVH GRPGCLQAVDFFDVLIKNLDIPSEFKDYLVPCLKQTSPSHQAAQMVP-QVEMA SNAHSQQMDDLFDLIIKSGEMSPLIKEPSPISKMRPVTANVTTMPVNTVVSRPPPQIHMA
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	1020 PPVASTIVSSS-ILEFPVSGHYDLLS-GQDNLSVIFAPDQI LPYDNCSSHGDGHLEVLLNSHSPLGRTSELRLLKIGNDDPQFENVLSGFSGKSTE PPPSPIHSALG-RLEDFLESS-TGTPLLRGHQDGPSSMPLIDDLHS PPLSLESTNSLSVSLESQLEAFLDGTLPSGNN-IPHLESNSDDRETFSLIDDLNN *
xMyoS_ xMyoV_ xMRTF-A xMRTF-B	1080 EIPSSPEHHEHDLSP-SPSSTS-SSS-APFDPADWLEALTSGSASNFGPG EILPSRENLDASLSSMETQMSPSSAEGTALQHVSFTESPWETMEWLDLTPPSSATCLSSL QMLSSLAILDHPPSPMDTSDLHFSPIGNSLG-LDISEPPLDGMDWLELSEP-PAMNLTPL DLLQNTAMLEHESTPMETTDTPFTAN-SCLS-LDLADANLDNMEWLDLTMPNSSSSLTPL
xMyoS_ xMyoC_ xMRTF-A xMRTF-B	SPVGSSIFSTDIFDSPDLSINRMIDLMVEQW TTTGPSIFNTDFLDVTDLNLNTAMDLHLEHW STFTPSVFSTDFLDSHDLHL-HWDSCL SSTLPSMFSTDFLDSNDLHL-HWE *********************************

*MyoC* (AAT90392) (Small et al., 2005), *MRTF-A* (Q8AYC2) and *MRTF-B* (Q8AYC1). Residues conserved in all sequences are indicated by an asterisk, those conserved in three sequences by a small circle. Conserved motifs are indicated, including the basic, Qrich, SAP and Zinc Finger domains. [The first part of this fig. begins on the previous page.]

#### TABLE 1

#### SEQUENCE SIMILARITY IN MYOCARDIN FAMILY PROTEINS

	MyoS	МуоС	MRTF-A	MRTF-B
MyoS		30	26	24
MyoC	30		30	31
MRTF-A	26	30		38
MRTF-B	24	31	38	

bryos were normal (Fig. 3A,B). No substantial inhibition on the formation of somites was observed. As *MyoS* is strongly expressed in the hypaxial muscles we used this gene as an additional marker for the differentiation of these groups of cells. As seen in Figure 3C,D, injection of MyoS-MO, but not C-MO, strongly inhibited the expression of *MyoS* in the hypaxial muscles. An attempt was made to rescue hypaxial muscle differentiation

after MyoS-MO injection by coinjection of mRNA in which the recognition site was modified (see Experimental Procedures). In this experiment, in which 20ng MO was injected into one blastomere at the 2-cell stage, the fraction of embryos that developed hypaxial muscles was substantially increased when mutated mRNA was coinjected with MyoS-MO (Fig. 4A,B).

The origins of the head musculature are complex. As described in the chick, most of these muscles arise from non-segmented head mesoderm (Noden and Francis-West, 2006). In *Xenopus*, Martin and Harland (2005) showed that certain head muscles arise from cells that migrate from anterior somites in a similar way as hypaxial cell precursors. The partial sensitivity of head muscles to MyoS-MO may be due to their multiple developmental origins.

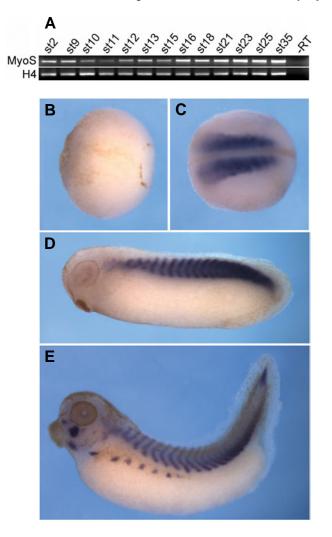
The experiments described above indicate that MyoS function is required for the formation of hypaxial muscles in the *Xenopus* embryo. The lack of effect of the MyoS-MO on somitic muscle differentiation might indicate that hypaxial muscles are more sensitive to diminished MyoS concentrations, while somitic muscle may require only low levels of this factor which might persist in the MO-injected embryos.

## MyoS can activate many muscle-specific genes in animal explants

Animal explants are suitable for testing the ability of a transcription factor to activate the expression of various genes. We used this system to examine the ability of MyoS to induce the expression of muscle-specific genes that are not expressed in untreated animal caps. As seen in Fig. 5, injection of 1 or 2ng *MyoS* RNA into the animal region of the embryo led to the induction of SRF, the putative partner of MyoS. This activation suggests that a positive feedback loop is activated by *MyoS* expression. Further, some regulatory genes involved in

muscle formation were induced including Nkx2.5 and, rather weakly, Mef2A. In contrast, MyoD and Myf5 were not induced at a detectable level. Likewise, other members of the MyoS family, MyoC, MRTF-A and MRTF-B were not induced. Genes characteristic for differentiating muscle cells of different types were activated, including skeletal and cardiac actin, cardiac troponin and smooth muscle genes including actin and SM22. In contrast several myosin chain genes were induced weakly or not at all. It is notable that cardiac and smooth muscle genes were induced effectively even though MyoS is not expressed in the heart or in smooth muscle cells, at least not during embryogenesis up to stage 41. In distinction to these results, MyoC activated smooth muscle and cardiac muscle, but not skeletal muscle genes after ectopic expression in animal explants (Small et al., 2004). Our observations may imply that under conditions of overexpression, MyoS cross activates target genes specific for the related factor MyoC. Similar observations have been reported for other members of the Myocardin family (Du et al., 2004; Selvaraj and Prywes, 2004; Wang et al., 2002).

Activation of several muscle-specific genes in animal explants in response to *MyoS* RNA injection did not lead to elongation of the explants (Fig. 6). Elongation is usually observed when mesoderm is induced by signaling factors such as activin and involves the activation of the *Xbra* gene, which is not activated by *MyoS* 



injection (Fig. 5A). Thus *MyoS* overexpression appears to activate only a portion of the genetic program that leads to full tissue differentiation.

## MyoS is a novel transcription factor involved in muscle differentiation

The results reported in this paper indicate that Myoskeletin is a novel member of the transcription factor family of which the founding member is Myocardin. As the expression of MyoS and *MyoC* is non-overlapping in the *Xenopus* embryo it is likely that these two related factors have similar functions in the regulation of muscle gene expression in different types of muscle. As the role of MyoC in cardiac and smooth muscle differentiation has been well documented (Chen et al., 2002; Small et al., 2005; Teg Pipes et al., 2006; Wang et al., 2001; Wang and Olson, 2004), the question arose whether skeletal muscle can be formed without participation of a factor of this class. It is now clear that somites and hypaxial precursors express a member of the MyoC family, MyoS, during their differentiation, suggesting that MyoS is involved in the regulation of muscle-specific gene expression in these cells. This suggestion could be supported by the finding that a morpholino antisense oligonucleotide targeting MyoS can suppress the formation of hypaxial muscles in the embryo. The reason why we did not observe a reduction in somite formation in the MO-injected embryos is not presently clear. It is possible that low levels of MyoS that persisted in the Mo-injected embryos were sufficient for somite differentiation. Alternatively an additional, as yet undiscovered factor in the MyoC family, might compensate for the reduction of MyoS levels in the somites after MyoS-MO

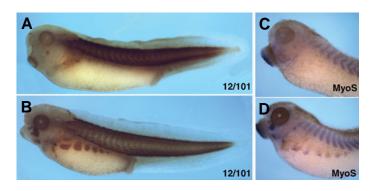
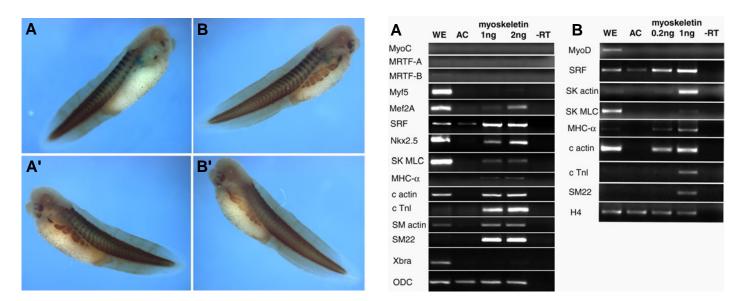


Fig. 2 (Left). Expression pattern of *MyoS*. (A) MyoS expression assayed by RT-PCR. (B-E) In situ hybridization. No pattern of expression is seen at stage 10.5 (B), but strong expression in the somites is apparent as stages 18 (C), 24 (D) and 37/38 (E). At stage 37/38, strong expression is also seen in hypaxial muscles and in head musculature.

**Fig. 3 (Right). MyoS is required for hypaxial muscle differentiation.** *MyoS-MO* **(A,C)** *or Control (C)-MO* **(B,D)** *were injected at a level of 20 ng into each blastomere at the 2-cell stage.* (A,B) *Staining with 12/101 antibody at stage 39.* (*C,D)* In situ hybridization with MyoS probe at stage 33/34. Strong reduction or complete absence of antibody staining was seen in 31/31 MyoS-MO-injected embryos **(A)**, while 34/37 *C-MOinjected embryos showed intense staining equivalent to that in uninjected embryos* **(B)**; three separate experiments were carried out. Using MyoS staining, 27/27 embryos showed strong suppression **(C)**, while 35/36 *C-MO-injected embryos showed normal expression* **(D)**.



**Fig. 4 (Left). Rescue of hypaxial muscle differentiation.** Embryos were injected into one side with MyoS-MO (20 ng) or MyoS-MO plus MyoS mRNA (0.5 ng); the injected side was marked by lacZ RNA. Embryos were stained with 12/101 antibody and for β-galactosidase. **(A)** MyoS-MO, injected side (3/16 or 20% of embryos showed medium to strong staining, 13/16 weak to negative staining); **(A')** uninjected side (16/16 strong staining). **(B)** MyoS-MO plus mut-MyoS mRNA, injected side (8/13 or 60% medium to strong staining, 5/13 weak to negative); **(B')** uninjected side (13/13 strong staining).

**Fig. 5 (Right). MyoS induces muscle genes in animal explants.** MyoS RNA was injected into the animal region of 2-cell embryos, animal caps were dissected at blastula, cultured to equivalent stage 12.5 and the expression of various genes was assayed by RT-PCR. In (A), 1 and 2ng mRNA was injected per embryo, in (B) the doses were 0.2 and 1ng. AC, uninjected animal caps; -RT, without reverse transcriptase.

injection, but might be absent or insufficient in hypaxial muscle precursors. Irrespective of these considerations it is likely that MyoS does have a role in the differentiation of muscle cells that arise in the somites during *Xenopus* embryogenesis.

#### **Experimental Procedures**

For microarray analysis, animal caps from stage 8/9 embryos were cultured with or without addition of 500pM activin until sibling embryos reached stage 11.5, for about 4hr. The explants were homogenized with Stat-60 (TEL-TEST), RNA was precipitated by isopropanol, purified using the RNeasy (Qiagen) system and treated with DNase I. Biotinylated probe was prepared with the Enzo RNA Transcript Labeling Kit following the protocol from Affymetrix. The probes were hybridized to Affymetrix *Xenopus* chips and the results analyzed using the GCOS1.1 and JMP5.1 software. Labeling and hybridization were repeated with four independent preparations of activin-treated and control RNA. Details of these experiments will be presented elsewhere.

The sequence of *myoskeletin* is based on the Image cDNA clone number 4959565, Est BM179219. The *myosleletin* sequence has been submitted to GeneBank under Accession number EF175167.

Embryo handling, whole mount in situ hybridization and antibody staining followed standard procedures. RNA for microinjection was prepared by the mMessage mMachine kit (Ambion) and MOs were obtained from Gene Tools. The sequence of the MyoS-MO is CGCTCTGAGGCCAGCAGGGTCATCT, the Control-MO is CCTCTTACCTCAGTTACAATTTATA. To generate mut-MyoSmRNA for use in rescue experiments, the sequence of the first 23 nts of the ORF was ATGACCCTGCTGGCCTCAGAGCG changed from to ATGACACTTCTGGCGTCCGAGCG. These changes generate four mismatches with the MyoS-MO but do not change the encoded amino acid sequence. MO against MyoS or an unrelated control (C) MO was injected into the equatorial region of both blastomeres at the 2-cell stage at a level of 20ng per cell. For RNA co-injection, MyoS RNA was mutated to abolish the target region while leaving the coding capacity unchanged; 20ng MyoS-MO or C-MO plus 0.5ng RNA were injected into one blastomere at the 2-cell stage together with 100ng  $\beta$ -galactosidase RNA to mark the injected side. Embryos were analyzed by *in situ* hybridization with MyoS at stages 33/34 and 39 and by staining with monoclonal antibody 12/101 (Developmental Studies Hybridoma Bank, University of Iowa) at stages 39-40.

For explant studies, RNA was injected into the animal region at the 2cell stage, animal caps dissected at stage 9 and the explants harvested at equivalent stage 12.5; incubation time was approximately 6 hr. RNA was extracted as described above and assayed by RT-PCR analysis. PCR primers used in this assay were those described by (Small *et al.*, 2005), except for *cardiac*  $\alpha$ -*actin* (Niehrs *et al.*, 1994), *MyoD* (Schohl and Fagotto, 2003) and *H4*, *ODC*, *Xbra* and *Myf5* (http://www.hhmi.ucla.edu/ derobertis/index.html).

#### Acknowledgment

We thank Stryder Meadows and Paul Krieg for making information available before publication. This work has been supported by the

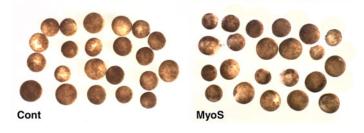


Fig. 6. Appearance of control explants and explants injected with 1 ng *MyoS* RNA. The animal explants were cultured in the same way as the explants in Fig. 5. No elongation was seen in these explants.

intramural research program of the National Institute of Child Health and Human Development, National Institutes of Health.

#### References

- ASASHIMA, M., NAKANO, H., SHIMADA, K., KINOSHITA, K., ISHII, K., SHIBAI, H. and UENO, N. (1990). Mesoderm induction in early amphibian embryos by activin A. *Roux's Arch. Dev. Biol.* 198: 330-335.
- BERKES, C. A. and TAPSCOTT, S. J. (2005). MyoD and the transcriptional control of myogenesis. Semin Cell Dev Biol 16: 585-595.
- BUCKINGHAM, M., BAJARD, L., CHANG, T., DAUBAS, P., HADCHOUEL, J., MEILHAC, S., MONTARRAS, D., ROCANCOURT, D. and RELAIX, F. (2003). The formation of skeletal muscle: from somite to limb. *J Anat* 202: 59-68.
- CHEN, J., KITCHEN, C. M., STREB, J. W. and MIANO, J. M. (2002). Myocardin: a component of a molecular switch for smooth muscle differentiation. J Mol Cell Cardiol 34: 1345-1356.
- CHRIST, B. and BRAND-SABERI, B. (2002) Limb muscle development. *Int. J. Dev. Biol.* (2002) 46: 905-914
- DAVIDSON, E. H. (2006). The regulatory genome. Amsterdam: Academic Press.
- DU, K. L., CHEN, M., LI, J., LEPORE, J. J., MERICKO, P. and PARMACEK, M. S. (2004). Megakaryoblastic leukemia factor-1 transduces cytoskeletal signals and induces smooth muscle cell differentiation from undifferentiated embryonic stem cells. *J Biol Chem* 279: 17578-17586.
- HEASMAN, J. (2006). Patterning the early Xenopus embryo. *Development* 133: 1205-1217.
- KINTNER, C. R. and BROCKES, J. P. (1984). Monoclonal antibodies identify blastemal cells derived from dedifferentiating limb regeneration. *Nature* 308: 67-69.
- MARTIN, B. L. and HARLAND, R. M. (2001). Hypaxial muscle migration during primary myogenesis in Xenopus laevis. *Dev Biol* 239: 270-280.
- MARTIN, B. L. and HARLAND, R. M. (2006). A novel role for lbx1 in Xenopus hypaxial myogenesis. *Development* 133: 195-208.
- MOHUN, T. J., CHAMBERS, A. E., TOWERS, N. and TAYLOR, M. V. (1991). Expression of genes encoding the transcription factor SRF during early development of Xenopus laevis: identification of a CArG box-binding activity as SRF. *Embo J* 10: 933-940.
- NAYA, F. J. and OLSON, E. (1999). MEF2: a transcriptional target for signaling pathways controlling skeletal muscle growth and differentiation. *Curr Opin Cell Biol* 11: 683-688.
- NODEN, D. M. and FRANCIS-WEST, P. (2006). The differentiation and morphogenesis of craniofacial muscles. *Dev Dyn* 235: 1194-1218.
- NIEHRS, C., STEINBEISSER, H. and DE ROBERTIS, E. M. (1994). Mesodermal patterning by a gradient of the vertebrate homeobox gene goosecoid. *Science* 263: 817-820.

- POURQUIÉ, O. (2003). Vertebrate somitogenesis: a novel paradigm for animal segmentation? *Int. J. Dev. Biol.* 47: 597-603.
- POWNALL, M. E., GUSTAFSSON, M. K. and EMERSON, C. P., JR. (2002). Myogenic regulatory factors and the specification of muscle progenitors in vertebrate embryos. *Annu Rev Cell Dev Biol* 18: 747-783.
- SCHOHL, A. and FAGOTTO, F. (2003). A role for maternal beta-catenin in early mesoderm induction in Xenopus. *Embo J* 22: 3303-3313.
- SELVARAJ, A. and PRYWES, R. (2004). Expression profiling of serum inducible genes identifies a subset of SRF target genes that are MKL dependent. *BMC Mol Biol* 5: 13.
- SMALL, E. M., WARKMAN, A. S., WANG, D. Z., SUTHERLAND, L. B., OLSON, E. N. and KRIEG, P. A. (2005). Myocardin is sufficient and necessary for cardiac gene expression in Xenopus. *Development* 132: 987-997.
- SMITH, J. C. (1987). A mesoderm-inducing factor is produced by Xenopus cell line. *Development* 99: 3-14.
- SMITH, J. C., PRICE, B. M., VAN NIMMEN, K. and HUYLEBROECK, D. (1990). Identification of a potent Xenopus mesoderm-inducing factor as a homologue of activin A. *Nature* 345: 729-731.
- TAPSCOTT, S. J. (2005). The circuitry of a master switch: Myod and the regulation of skeletal muscle gene transcription. *Development* 132: 2685-2695.
- TEG PIPES, G. C., CREEMERS, E. E. and OLSON, E. N. (2006). The myocardin family of transcriptional coactivators: versatile regulators of cell growth, migration and myogenesis. *Genes Dev* 20: 1545-1556.
- WANG, D., CHANG, P. S., WANG, Z., SUTHERLAND, L., RICHARDSON, J. A., SMALL, E., KRIEG, P. A. and OLSON, E. N. (2001). Activation of cardiac gene expression by myocardin, a transcriptional cofactor for serum response factor. *Cell* 105: 851-862.
- WANG, D. Z., LI, S., HOCKEMEYER, D., SUTHERLAND, L., WANG, Z., SCHRATT, G., RICHARDSON, J. A., NORDHEIM, A. and OLSON, E. N. (2002). Potentiation of serum response factor activity by a family of myocardin-related transcription factors. *Proc Natl Acad Sci U S A* 99: 14855-14860.
- WANG, D. Z. and OLSON, E. N. (2004). Control of smooth muscle development by the myocardin family of transcriptional coactivators. *Curr Opin Genet Dev* 14: 558-566.
- WARDLE, F. C. and SMITH, J. C. (2006). Transcriptional regulation of mesendoderm formation in Xenopus. *Semin Cell Dev Biol* 17: 99-109.
- WHITMAN, M. (2001). Nodal signaling in early vertebrate embryos: themes and variations. *Dev Cell* 1: 605-617.

Received: 12th December 2006 Reviewed by Referees: 12th January 2007 Modified by Authors and Accepted for Publication: 21st February 2007 Published Online: 4th May 2007

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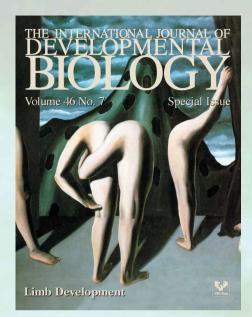
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