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Bruno Della Gaspera, Alice Mateus, Yannick Andéol, Laure Weill, Frédéric Charbonnier, et al.. Lineage tracing of sclerotome cells in amphibian reveals that multipotent somitic cells originate from lateral somitic frontier. *Developmental Biology*, 2019, 453, pp.11 - 18. 10.1016/j.ydbio.2019.05.009 . hal-03488174

HAL Id: hal-03488174

<https://hal.science/hal-03488174v1>

Submitted on 20 Dec 2021

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**LINEAGE TRACING OF SCLEROTOME CELLS IN AMPHIBIAN REVEALS THAT
SOMITIC MULTIPOTENT CELLS ORIGINATE FROM LATERAL SOMITIC
FRONTIER**

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ABSTRACT

The two somite compartments, dorso-lateral dermomyotome and medio-ventral sclerotome are major vertebrate novelties, but little is known about their evolutionary origin. We determined that sclerotome cells in *Xenopus* come from lateral somitic frontier (LSF) by lineage tracing, ablation experiments and histological analysis. We identified *Twist1* as marker of migrating sclerotome progenitors in two amphibians, *Xenopus* and axolotl. From these results, three conclusions can be drawn. First, LSF is made up of multipotent somitic cells (MSCs) since LSF gives rise to sclerotome but also to dermomyotome as already shown in *Xenopus*. Second, the basic scheme of somite compartmentalization is conserved from cephalochordates to anamniotes since in both cases, lateral cells envelop dorsally and ventrally the ancestral myotome, suggesting that lateral MSCs should already exist in cephalochordates. Third, the transition from anamniote to amniote vertebrates is characterized by extension of the MSCs domain to the entire somite at the expense of ancestral myotome since amniote somite is a naive tissue that subdivides into sclerotome and dermomyotome. Like neural crest pluripotent cells, MSCs are at the origin of major vertebrate novelties, namely hypaxial region of the somite, dermomyotome and sclerotome compartments. Hence, change in MSCs properties and location is involved in somite evolution.

INTRODUCTION

Somites are embryonic transitory structures located in dorsal paraxial mesoderm on both side of neural tube and notochord (Bothe et al, 2007; Sambasivan and Tajbakhsh, 2007). Somites segment and differentiate in an antero-posterior direction during development. The somite subdivision into dermomyotome, myotome, and sclerotome forms the basis of the musculoskeletal system organization that gives rise to muscle (from dermomyotome and myotome), and also bone and tendon (from sclerotome). Somites are specific to Chordates and are major contributors of vertebrate evolutionary novelties. Indeed, vertebrae that gave the name of vertebrate subphylum originate from sclerotome. At the phylotypic stage (Irie and Kuratani, 2011), the most conserved stage of vertebrate development, somite compartmentalization is similar between the distinct classes of vertebrates (della Gaspera et al, 2018).

Interestingly, the initial phase of compartmentalization differs between amniotes and anamniotes suggesting that anamniote somite organisation is closest to the chordate ancestor. In amniotes, the newly formed somites are naïve structures composed of multipotent somitic cells (MSCs) that divide into dermomyotome dorsally and sclerotome ventrally (Yusuf and Brand-Saberi, 2006; Buckingham and Vincent, 2009). Next, myotome cells arising from the dermomyotome sit between dermomyotome and sclerotome. In contrast, in anamniotes, compartmentalization begins before somite formation and is organized in medio-lateral way (Devoto et al, 1996; Hinitz et al, 2009; della Gaspera et al, 2012a). The ancestral myotome appears first medially from the presomitic mesoderm whereas the dermomyotome appears laterally before becoming dorsal structure (Hollway et al, 2007, della Gaspera et al, 2012b). However, the origin of sclerotome in the anamniotes is still unclear. In a primitive chordate, the cephalochordate amphioxus, the somite compartmentalization is similar to anamniote vertebrates, with an ancestral myotome positioned medially. Although the lateral area is poorly characterized, derivative sclerotome-like cells have been recently identified (Mansfield et al, 2015). Therefore, understanding the sclerotome origin in anamniotes may provide an advanced picture of the evolutionary history of somite compartmentalization. Here, we investigated the origin of sclerotome cells in amphibians and showed by lineage tracing, histological analysis, tissue ablation and gene marker studies that sclerotome cells come from the lateral somitic frontier (LSF).

MATERIALS AND METHODS

Ethics statement

This work uses early *Xenopus laevis* and axolotl (*Ambystoma mexicanum*) embryos. All experimental procedures used in this study followed the recommendations of the “Comité National de Réflexion Ethique sur l’Expérimentation Animale” of the Ministry of Higher Education and Research and were approved by local Animal Care and Use Committees.

Lineage tracing experiments and immunofluorescence

Stage 13 embryos were injected using pulled-glass capillaries either with wheat germ agglutinin, WGA-rhodamine or -fluorescein or both (dissolved at 200 µg/mL, Vector laboratories) to be analysed at stages 23 and 28. Single injections were performed at LSF or at medial level of presomitic mesoderm. Double injections with WGA-rhodamine and -fluorescein were performed at medial and lateral regions respectively. Embryos were photographed at stage 13 and fixed in 4% formaldehyde at stage 28. The fate of stained cells was directly analysed on 50 µm trunk transverse vibratome sections or after immunofluorescence with the muscle-specific 12/101 or MF20 monoclonal antibodies used at 1:300 dilution for 1 hour (Developmental studies hybridoma bank-University of Iowa). The primary antibody was recognized by Alexa fluor donkey 488 anti-mouse secondary antibody (Life technologies) used at 1:300 dilution for 1 hour. In some cases, vibratome sections were incubated with Hoechst 33258 (Sigma) at 1:2000 dilution for 5 minutes before being analysed. Embryos were staged according to Nieuwkoop and Faber (1994), and Schreckenber and Jacobson (1975).

Whole-mount *in situ* hybridization and immunohistochemistry

cDNA for whole-mount *in situ* hybridization (ISH) probes were prepared by RT-PCR with the following forward and reverse primers: 5’-GAGCTGGACTCCAAAATGGC-3’ and 5’-TGGCAACGCATGCATGCAGT-3’ for *Twist1*, 5’-ATCCCAGCCAACGTTACCCT-3’ and 5’-GATGAAGATGGGAGTGGAAG-3’ for *Pax1*, 5’-ACCACATACAACGCCATGGG-3’ and 5’-GTTACTTTGTGTGTCCTCCAG-3’ for *Pax9*, 5’-AAAGATGCCACCAAGGAGGA-

3' and 5'-GTCCACACAATCCAGGTACA-3' for *FoxC1*, 5'-GGAAGATCGGCTCCTTAAAG-3' and 5'-ACGTTGGGAAAAGTCTGCTG-3' for *FoxC2* or 5'-GACGAGCTCGACTCCAAGAT-3' and 5'-AATGGAGTGTGGGCAATGTC-3' for axolotl *Twist1*. PCR products were cloned into pGEM-T (Promega). cDNAs of *Meox2*, *Tcf15*, *MyhE3*, *Mef2c* and *Myod1* were previously described (della Gaspera et al, 2009 and 2012b). RNA probes synthesis and whole mount *in situ* hybridization were carried out as previously described (Harland, 1991). Following *in situ* hybridization and depigmentation, embryos were fixed at room temperature for 1 hour in 4% formaldehyde, 0.25% glutaraldehyde PBS (Phosphate Buffer Saline) solution before embryos were sectioned transversally. When both *in situ* hybridization and 12/101 immunostaining were carried out, embryos were first stained using *in situ* hybridization and sectioned before immunohistochemistry. The 12/101 antibody was used at 1:1000 dilution and detected with an alkaline phosphatase-conjugated anti-mouse antibody (Vector Laboratories) diluted at 1:1000 followed by reaction with BCIP (Sigma-Aldrich).

Embryos section

Following *in situ* hybridization, fixed embryos were embedded in a PBS solution with 3% agarose, 1% porcine gelatin B (Sigma) and 0.01% glutaraldehyde. In the other cases, embryos were embedded without adding glutaraldehyde. 40 µm vibratome sections were performed transversally and serially, in an orderly way from anterior to posterior side. For histological analysis, embryos sections were incubated successively with 12/101 antibody at 1:300 dilution for 1 hour, Alexa fluor donkey 488 anti-mouse secondary antibody at 1:300 dilution for 1 hour, diI (Molecular Probes) at 10 µg/mL or WGA-rhodamine at 20 µg/mL for 15 minutes and Hoechst 33258 (Sigma) at 1:2000 dilution for 5 minutes.

Ablation experiments

Dechorionated embryos were oriented and immobilized in 0.4X MMR 1% agarose gel slits. Using sharpened needles, antero-posterior incision in the ectoderm was performed at stages 17-18 on the dorsal side of the embryo at the medial (Abl. M) or lateral level (Abl. L). Ectoderm and mesoderm were separated from each other on the lateral incision side and the superficial part of the mesoderm was removed. The sham-operated (Abl. C) embryos were treated without removing the mesoderm. The embryos were fixed at stages 28, 32 or 45.

RESULTS

In order to highlight the fate of LSF cells specified by *Meox2* expression (della Gaspera et al, 2012b and Fig. 1) during *Xenopus* development, we first double injected stage 13 embryos with fluorescent dyes, WGA-fluorescein (green) in the LSF and WGA-rhodamine (red) in the medial presomitic mesoderm (Fig. 1A a, b and c). At stage 28, as shown in three distinct embryos in Figure 1, cells located at the somite periphery are stained by the WGA-fluorescein, suggesting that LSF migrate to these regions. Cells located to the most dorso-lateral region correspond to the dermomyotome as previously showed (della Gaspera et al, 2012b) while the ventral and medial cells near the neural tube and notochord could be sclerotome cells (Fig. 1A g, h and i) (Sánchez and Sánchez, 2013). As expected (della Gaspera et al, 2012a and b), the medial-injected cells (WGA-rhodamine/red) gave rise to internal somitic cells (Fig. 1A d, e and f). Co-staining with a nuclear marker showed clearly that cells between the notochord and the myotome (Fig. 1B and C) were labeled after injection in the LSF (Fig. 1B), but never after injection in the medial presomitic mesoderm (Fig. 1C), indicating that non-muscle cells of lateral origin were present in ventral and medial region of the somites. Indirect immunofluorescence with two muscle-specific antibodies (green) either 12/101 (Fig. 1D a and a') or MF20 (Fig. 1D d and d') on embryos previously injected with WGA-rhodamine in the LSF (Fig. 1D b, e, b' and e') confirmed no co-localization of the rhodamine labelled cells with the myotome. In addition, *in situ* hybridization experiments revealed that *Twist1* (Fig. 1E a) and *Pax1/Pax9* mRNAs (Fig. 1F a), two sclerotome markers (Fleming et al, 2015), were expressed in the same region. Finally, a co-staining of fluorescent LSF-derived cells with *Twist1* (Fig. 1E b and c) or *Pax1/Pax9* cRNA probes (Fig. 1F c and d) on serial transverse sections (Fig. 1F) identified these cells as sclerotome ones. Hence, our results suggest that cells from LSF migrate around primitive myotome dorsally and ventrally to give rise to dermomyotome and sclerotome cells.

We showed previously that LSF expresses *Meox2* and *Tcf15* at stage 13 (della Gaspera et al., 2012b) and two early mesoderm and somitic markers, *Foxc1* and *Foxc2*, which are broadly expressed in the trunk mesoderm but not in the notochord. The LSF cells probably migrate medially by extension-convergence, then at stage 18, extend into lateral somitic domain (della Gaspera et al, 2012b). The lateral somitic domain still expresses the same markers than LSF, while the ancestral myotome differentiates first medially (stage 16) and next dorso-laterally (stages 17/18) (della Gaspera et al, 2012a). To characterize molecularly

the migrating sclerotome progenitors, we performed here *in situ* hybridization experiments, with the LSF markers *Meox2* and *Tcf15* and sclerotome-expressed genes, *Foxc1*, *Foxc2* and *Twist1* (Fleming et al, 2015) in comparison with myotome marker *Myod1*.

Twist1 was expressed at stage 20 in the lateral somitic domain and this expression was stronger in cells bordering the myotome (Fig. 2A). In the following stages, during myotome growth, dermomyotome progenitors migrate dorsally and continued to express LSF markers and *Twist1* (Fig. 2A). While dermomyotome maturation, the expression of *Meox2*, *Tcf15* and *Twist1* fell, and *Foxc1* and *Foxc2* expressions were restricted to hypaxial and epaxial regions (Fig. 2A). Concerning the sclerotome cells, only *Twist1* was expressed in the cell layers that were located ventrally and medially at stage 26. *Foxc1* and *Foxc2* mRNAs started to accumulate in the sclerotome cells only at stage 34 (Fig. 2A). Considering that somites differentiate in an antero-posterior way, we analysed *Twist1* expression in transverse sections from posterior to anterior direction at stage 28 (Fig. 2B). *Twist1* expression domain shifted over somite differentiation, from lateral region caudally, to medial location anteriorly (Fig. 2B a to f). In the same way, histological analysis in transverse sections, with myotome marker and *diI* to mark all cells, from posterior to anterior direction, allowed to easily visualize a non-muscle cells population which seems to migrate ventrally from lateral to medial region (Fig. 2C). In the less differentiated somite, at posterior region, these cells were located laterally (Fig. 2C). These cells seem to migrate between the myotome and endoderm at slightly more anterior region and, between the myotome and the notochord, at the most anterior region (Fig. 2C). We equally observed these seemingly moving cells (Fig. 3A), and shift of *Twist1* expression from lateral to medial domain (Fig. 3B), over time from stages 18 to 25, at the anterior trunk level. Altogether, these results suggest that lateral *Twist1* expressing cells migrate ventrally to give rise to sclerotome.

We next followed experimentally the sclerotome progenitors expressing *Twist1* from lateral to medial somite side. LSF-injected cells at stage 13 give rise to lateral somitic domain at stage 18 (della Gaspera et al, 2012b) which express *Twist1* (Fig. 3B). We fixed LSF-injected embryos at stage 22/23, which were also stained with myotome marker (Fig. 3C) or with *Twist1* ISH-probe (Fig. 3D) on the same slides. We observed a co-localization of the tracer with *Twist1* expressing cells in the lateral to medial direction (Fig. 3D) indicating that these cells move ventrally from lateral to medial side of the somite (Fig. 3C and D). To confirm experimentally the origin of sclerotome cells, we proceeded to the ablation of lateral somitic domain at stage 17, when *Twist1* begin to be expressed, and showed that the absence of these cells dramatically affected sclerotome formation since no *Twist1* expression at stage 28 was

detected (Fig. 3E) while immunohistochemistry with 12/101 antibody indicated that the myotome was faintly affected. Neither the sham-operated embryos, nor the ablation of medial somite led to a decrease in *Twist1* expression. We already showed that ablation of LSF leads to absence of dermomyotome and particularly to *Pax3* expression at hypaxial region (della Gaspera et al, 2012b). Here, we confirmed that ablation of LSF prevented the development of hypaxial muscles derived from somite (Fig. 3F). Since the syndetome, which give rise to tendons and ligaments is derived from sclerotome in amniotes and anamniotes (Brent et al, 2003; Bricard et al, 2014), we also analyzed the expression of *Mef2c*, a syndetome marker in *Xenopus* (della Gaspera et al, 2009) and probably in mice (Edmondson et al, 1994). *Mef2c* expression was strongly decreased after the ablation of lateral somitic domain (Fig. 3G). Altogether, the co-staining of the tracer with *Twist1*-expressing cells and the LSF ablation experiments confirms that the sclerotome/synetome originates from lateral somitic domain. Moreover, hypaxial muscles stems also from lateral somitic domain.

To ensure that the lateral origin of sclerotome cells is a general anamniote feature, we then performed whole-mounted hybridization for *Twist1* on the urodelan amphibian, axolotl, at stage 31 (Fig. 4A) and compared to myotome staining by 12/101 antibody (Fig. 4B). *Twist1* expression was also analyzed in transverse sections from posterior to anterior direction as somite differentiates (Fig. 4A and B, a to g). As in *Xenopus* embryos, *Twist1* was detected in non-muscle cells which seem to migrate ventrally from lateral region to medial region (Fig. 4A and 4B). As in *Xenopus* embryos, in the less differentiated somite at posterior level, *Twist1* expressing cells were located laterally. At slightly more anterior level, *Twist1* expressing cells seem to migrate between the myotome and endoderm (Fig. 4A and B) before migrating between myotome and notochord. At the same time, *Twist1* mRNA level also decreased sharply, in contrast to *Xenopus*. Despite this difference, these results strongly suggest that the sclerotome cells have also a lateral origin in axolotl.

DISCUSSION

The combination of lineage tracing, histological analysis, ablation experiments, and expression markers studies shows that sclerotome cells originate from LSF at early neurulation in *Xenopus*. *Twist1* marks ventrally the migrating sclerotome cells and *Twist1* expression in axolotl confirms that the lateral origin of sclerotome cells is a general anamniote trait. Moreover, LSF is made up of MSCs since (1) early somite markers as *Meox2* and *Tcf15* are strongly expressed at LSF (della Gaspera et al, 2012b) and (2) both sclerotome and dermomyotome (della Gaspera et al, 2012b) derive from LSF.

Twist1 marks ventrally the migrating sclerotome cells but is not restricted to these cells in somites. In amniotes, *Twist1* has been shown expressed in whole undifferentiated somites before to be restricted to the sclerotome (Füchtbauer, 1995; Gitelman, 1997). In *Xenopus*, we show here that *Twist1* is first expressed in the lateral somitic domain and next both in the migrating dorsal and ventral cells until stage 23-24, before to be restricted to the sclerotome at later stages (Fig. 2A). The lateral somitic domain and next, the dorsal part of the somite that gives rise to the functional dermomyotome at stage 28, also express other early patterning somite genes like *Foxc1*, *Foxc2*, *Meox2* and *Tcf15* until stage 23-24. It has been shown that these somite patterning genes could participate to the dermomyotome but also to the sclerotome formation (Kume et al, 2001; Mankoo et al, 2003; Wilson-Rawls et al, 2004; Linker et al, 2005). Hence, although the sclerotome precursors expressing *Twist1* seem to take the ventral route to form the sclerotome medially, we cannot exclude that some dorsal sclerotome cells come from the dorsal route. Moreover, in zebrafish, a dorsal sclerotome domain with specific markers has been recently characterized (Ma et al, 2018).

The discovery of MSCs at LSF in addition to the knowledge about ancestral myotome and dermomyotome provides a more precise description of somite compartmentalization in *Xenopus* (della Gaspera et al, 2012a and b), making it easier comparisons between chordate species, and allows reconstructing the evolutionary history model of somite compartmentalization (Fig. 4D). MSCs should be already present in amphioxus since lateral somite cells, like in *Xenopus*, migrate dorsally and ventrally in order to give rise to both external cells and sclerotome-like cells respectively (Mansfield et al, 2015). The transition

from cephalochordates to vertebrates is characterized by appearance of new somitic compartments, namely the hypaxial region of dermomyotome, the dermomyotome itself and the sclerotome. Our results show that all these new somite novelties in vertebrates derive from MSCs. MSCs or MSC-daughter cells evolution could be related to the modification of their cell potency state, allowing MSCs to acquire new cell identities and/or to redeploy pre-existing genetic program. Indeed, the new set of genes inherited from the double whole genome duplication (2R hypothesis) that took place during this evolutionary period (Panopoulou and Poustka, 2005), could participate to the evolution of MSCs and somite novelties (Shimeld and Holland, 2000; Furlong and Holland, 2002). It is the MSCs location that changes at the transition from anamniote to amniote vertebrates. We previously proposed that the transition from anamniotes to amniotes is associated to the loss of ancestral myotome (della Gaspera et al, 2012a). Hence, the transition from anamniotes to amniotes is characterized by global extension of MSCs to the entire somite at the expense of ancestral myotome (Fig. 4D). In amniotes, the whole newly formed somite is a naïve structure that expresses the same markers as *Xenopus* MSCs at LSF (Chal and Pourquié, 2017, Nakajima et al, 2018). The evolutionary advantages of the MSCs domain extension remain to be determined but it could be related to the appearance of a larger sclerotome and facilitating the formation of its derivatives. It is well known that the muscle development begins earlier in anamniotes than in amniotes probably because the water tadpole must quickly find its own feeding. In *Xenopus* somite, we identified three successive myogenic waves during embryogenesis, on the basis of myogenic regulatory factors expression (della Gaspera et al, 2012a). The two first ones give rise to the primary myotome, formed by mononucleated fibers until stage 45 (Kiełbówna and Daczewska, 2005). The third one participates to the myotome growth. More precisely, the first -medial and lateral- myogenic wave is at the origin of the ancestral myotome, which disappeared per se in amniotes. The second wave, at the hypaxial and epaxial lips of the dermomyotome, is similar to those observed at the beginning of amniote myogenesis, with strong expression of *Myf5* and *Myog*. Hence, we hypothesized that the first -medial and lateral- myogenic wave was lost, at least in mice (della Gaspera et al, 2012a). However, it seems that the remnants of first wave could still be present in amniotes, as in the chick (Fig. 4D). Indeed, muscle pioneer cells observed precociously in medial somite during chick myotome formation seem to be the remnant of the first medial myogenic wave (Kahane et al, 1998). Moreover, amniote dermomyotome can generate the primary myotome not only from its epaxial and hypaxial borders but also from its anterior and posterior borders (Gros et al, 2004). Although epaxial and hypaxial myogenesis presents similarities between

Xenopus and amniotes, as explained above, myogenic cells arising from anterior and posterior borders have until now no equivalents in *Xenopus*. One hypothesis is that the first lateral wave in *Xenopus* could evolve to give rise to these anterior and posterior populations. Therefore, further studies are needed to find whether specific features are shared between these two myogenic populations.

Our results point out that change in MSCs properties and location could partly explain both the transition from cephalochordates to vertebrates and from anamniote to amniote vertebrates, underlining the particular role of some kind of multipotent embryonic cells during vertebrate evolution. In the same way, neural crest cells are vertebrate-specific and pluripotent cells that give rise to numerous cell types (Buitrago-Delgado et al, 2015). Neural crests also contribute to the major vertebrate novelties like MSCs do (Shimeld and Holland, 2000). This raises the question, how and to what extent, acquisition or modification of the cell potency state in a specific cell type could be essential to vertebrate evolution.

Acknowledgments:

We thank Dr Anne Simon for providing some reagents.

Competing interests:

The authors declare no competing or financial interests.

Funding:

This work was supported by INSERM (Institut National de la Santé et de la Recherche Médicale) and Université Paris Descartes. This research did not receive any specific grant from funding agencies in the public, commercial, or not-for-profit sectors.

REFERENCES

- Bothe, I., Ahmed, M.U., Winterbottom, F.L., von Scheven, G., Dietrich, S., 2007. Extrinsic versus intrinsic cues in avian paraxial mesoderm patterning and differentiation. *Dev. Dyn.* 236, 2397–2409. <https://doi.org/10.1002/dvdy.21241>
- Brent, A.E., Schweitzer, R., Tabin, C.J., 2003. A somitic compartment of tendon progenitors. *Cell* 113, 235–48. [https://doi.org/10.1016/S0092-8674\(03\)00268-X](https://doi.org/10.1016/S0092-8674(03)00268-X)
- Bricard, Y., Ralli re, C., Leuret, V., Lefevre, F., Rescan, P.-Y., 2014. Early fish myoseptal cells: insights from the trout and relationships with amniote axial tenocytes. *PLoS One* 9, e91876. <https://doi.org/10.1371/journal.pone.0091876>
- Brunet, T., Lauri, A., Arendt, D., 2015. Did the notochord evolve from an ancient axial muscle? The axochord hypothesis. *Bioessays* 37, 836–50. <https://doi.org/10.1002/bies.201500027>
- Buckingham, M., Vincent, S.D., 2009. Distinct and dynamic myogenic populations in the vertebrate embryo. *Curr. Opin. Genet. Dev.* 19, 444–53. <https://doi.org/10.1016/j.gde.2009.08.001>
- Buitrago-Delgado, E., Nordin, K., Rao, A., Geary, L., LaBonne, C., 2015. NEURODEVELOPMENT. Shared regulatory programs suggest retention of blastula-stage potential in neural crest cells. *Science* 348, 1332–5. <https://doi.org/10.1126/science.aaa3655>
- Chal, J., Pourqui , O., 2017. Making muscle: skeletal myogenesis in vivo and in vitro. *Development* 144, 2104–2122. <https://doi.org/10.1242/dev.151035>
- Della Gaspera, B., Armand, A.-S., Lecolle, S., Charbonnier, F., Chanoine, C., 2012. Mef2d acts upstream of muscle identity genes and couples lateral myogenesis to dermomyotome formation in *Xenopus laevis*. *PLoS One* 7, e52359. <https://doi.org/10.1371/journal.pone.0052359>
- Della Gaspera, B., Armand, A.-S., Sequeira, I., Chesneau, A., Mazabraud, A., L colle, S., Charbonnier, F., Chanoine, C., 2012. Myogenic waves and myogenic programs during *Xenopus* embryonic myogenesis. *Dev. Dyn.* 241, 995–1007. <https://doi.org/10.1002/dvdy.23780>
- Della Gaspera, B., Armand, A.-S., Sequeira, I., Lecolle, S., Gallien, C.L., Charbonnier, F., Chanoine, C., 2009. The *Xenopus* MEF2 gene family: evidence of a role for XMEF2C in larval tendon development. *Dev. Biol.* 328, 392–402. <https://doi.org/10.1016/j.ydbio.2009.01.039>
- Della Gaspera, B., Chesneau, A., Weill, L., Charbonnier, F., Chanoine, C., 2018. *Xenopus* SOX5 enhances myogenic transcription indirectly through transrepression. *Dev. Biol.* 442, 262–275. <https://doi.org/10.1016/j.ydbio.2018.07.025>

- Devoto, S.H., Melançon, E., Eisen, J.S., Westerfield, M., 1996. Identification of separate slow and fast muscle precursor cells in vivo, prior to somite formation. *Development* 122, 3371–80. <https://doi.org/10.1103/PhysRevB.61.16956>
- Edmondson, D.G., Lyons, G.E., Martin, J.F., Olson, E.N., 1994. Mef2 gene expression marks the cardiac and skeletal muscle lineages during mouse embryogenesis. *Development* 120, 1251–63.
- Fleming, A., Kishida, M.G., Kimmel, C.B., Keynes, R.J., 2015. Building the backbone: the development and evolution of vertebral patterning. *Development* 142, 1733–44. <https://doi.org/10.1242/dev.118950>
- Füchtbauer, E. -m, 1995. Expression of M-twist during postimplantation development of the mouse. *Dev. Dyn.* <https://doi.org/10.1002/aja.1002040309>
- Furlong, R.F., Holland, P.W.H., 2002. Were vertebrates octoploid? *Philos. Trans. R. Soc. Lond. B. Biol. Sci.* 357, 531–44. <https://doi.org/10.1098/rstb.2001.1035>
- Gitelman, I., 1997. Twist Protein in Mouse Embryogenesis. *Dev. Biol.* 189, 205–214. <https://doi.org/10.1006/dbio.1997.8614>
- Gros, J., Scaal, M., Marcelle, C., 2004. A two-Step mechanism for myotome formation in chick. *Dev. Cell.* 6, 875–882. <https://doi.org/10.1016/j.devcel.2004.05.006>
- Harland, R.M., 1991. In situ hybridization: an improved whole-mount method for *Xenopus* embryos. *Methods Cell Biol.* 36, 685–695.
- Hinits, Y., Osborn, D.P.S., Hughes, S.M., 2009. Differential requirements for myogenic regulatory factors distinguish medial and lateral somitic, cranial and fin muscle fibre populations. *Development* 136, 403–14. <https://doi.org/10.1242/dev.028019>
- Hollway, G.E., Bryson-Richardson, R.J., Berger, S., Cole, N.J., Hall, T.E., Currie, P.D., 2007. Whole-somite rotation generates muscle progenitor cell compartments in the developing zebrafish embryo. *Dev. Cell* 12, 207–19. <https://doi.org/10.1016/j.devcel.2007.01.001>
- Irie, N., Kuratani, S., 2011. Comparative transcriptome analysis reveals vertebrate phylotypic period during organogenesis. *Nat. Commun.* 2, 248. <https://doi.org/10.1038/ncomms1248>
- Kahane, N., Cinnamon, Y., Kalcheim, C., 1998. The origin and fate of pioneer myotomal cells in the avian embryo. *Mech. Dev.* 74, 59–73. [https://doi.org/10.1016/S0925-4773\(98\)00066-5](https://doi.org/10.1016/S0925-4773(98)00066-5)
- Kielbówna, L., Daczewska, M., 2005. The origin of syncytial muscle fibres in the myotomes of *Xenopus laevis* - A revision. *Folia Biol. (Krakow).* 53, 39–44. <https://doi.org/10.3409/1734916054663401>
- Kume, T., 2001. The murine winged helix transcription factors, Foxc1 and Foxc2, are both required for cardiovascular development and somitogenesis. *Genes Dev.* 15, 2470–2482. <https://doi.org/10.1101/gad.907301>

- Linker, C., 2005. Beta-catenin-dependent Wnt signalling controls the epithelial organisation of somites through the activation of paraxis. *Development* 132, 3895–3905. <https://doi.org/10.1242/dev.01961>
- Ma, R.C., Jacobs, C.T., Sharma, P., Kocha, K.M., Huang, P., 2018. Stereotypic generation of axial tenocytes from bipartite sclerotome domains in zebrafish. *PLOS Genet.* 14, e1007775. <https://doi.org/10.1371/journal.pgen.1007775>
- Mankoo, B.S., Skuntz, S., Harrigan, I., Grigorieva, E., Candia, A., Wright, C.V.E., Arnheiter, H., Pachnis, V., 2003. The concerted action of Meox homeobox genes is required upstream of genetic pathways essential for the formation, patterning and differentiation of somites. *Development* 130, 4655–4664. <https://doi.org/10.1242/dev.00687>
- Mansfield, J.H., Haller, E., Holland, N.D., Brent, A.E., 2015. Development of somites and their derivatives in amphioxus, and implications for the evolution of vertebrate somites. *Evodevo* 6, 21. <https://doi.org/10.1186/s13227-015-0007-5>
- Nakajima, T., Shibata, M., Nishio, M., Nagata, S., Alev, C., Sakurai, H., Toguchida, J., Ikeya, M., 2018. Modeling human somite development and fibrodysplasia ossificans progressiva with induced pluripotent stem cells. *Development* 145, dev165431. <https://doi.org/10.1242/dev.165431>
- Nieuwkoop, P.D. and Faber, J., 1994. Normal table of *Xenopus laevis* (Daudin).
- Panopoulou, G., Poustka, A.J., 2005. Timing and mechanism of ancient vertebrate genome duplications -- the adventure of a hypothesis. *Trends Genet.* 21, 559–67. <https://doi.org/10.1016/j.tig.2005.08.004>
- Sambasivan, R., Tajbakhsh, S., 2007. Skeletal muscle stem cell birth and properties. *Semin. Cell Dev. Biol.* 18, 870–82. <https://doi.org/10.1016/j.semcdb.2007.09.013>
- Sánchez, R.S., Sánchez, S.S., 2013. Characterization of pax1, pax9, and uncx sclerotomal genes during *Xenopus laevis* embryogenesis. *Dev. Dyn.* 242, 572–9. <https://doi.org/10.1002/dvdy.23945>
- Schreckenberg, G.M., Jacobson, A.G., 1975. Normal stages of development of the axolotl. *Ambystoma mexicanum*. *Dev. Biol.* 42, 391–400. [https://doi.org/10.1016/0012-1606\(75\)90343-7](https://doi.org/10.1016/0012-1606(75)90343-7)
- Shimeld, S.M., Holland, P.W., 2000. Vertebrate innovations. *Proc. Natl. Acad. Sci. U. S. A.* 97, 4449–52. <https://doi.org/10.1073/pnas.97.9.4449>
- Wilson-Rawls, J., Rhee, J.M., Rawls, A., 2004. Paraxis Is a Basic Helix-Loop-Helix Protein That Positively Regulates Transcription through Binding to Specific E-box Elements. *J. Biol. Chem.* 279, 37685–37692. <https://doi.org/10.1074/jbc.M401319200>
- Yusuf, F., Brand-Saberi, B., 2006. The eventful somite: patterning, fate determination and cell division in the somite. *Anat. Embryol. (Berl)*. 211 Suppl, 21–30. <https://doi.org/10.1007/s00429-006-0119-8>

Figure 1. Sclerotome cells originate from lateral somitic frontier (A) Scheme of the experimental procedure: lineage tracing experiments with WGA-rhodamine injected in the medial segmental plate and with WGA-fluorescein injected in the *Meox2* expression domain at LSF. a to c: Results for three embryos double injected at stage 13. d to i: Transverse section at stage 28. WGA-rhodamine (d, e, f), WGA-fluorescein (g, h, i), merge (j, k, l). Dash lines indicate the position of bilateral symmetry plane. Horizontal lines define the limit between anterior and trunk regions. Anterior side on the top. Nc, notochord. Nt, neural tube. (B) Scheme of the experimental procedure: lineage tracing experiments with WGA-fluorescein injected in the *Meox2* expression domain at LSF. Fixed embryos were stained with Hoechst 33258 nucleus marker. Transverse section was performed at stage 28. Blue Hoechst 33258 nucleus marker (a), green WGA-fluorescein (b), merge (c). Arrows design cells lining the notochord and muscle cells. (C) Scheme of the experimental procedure: lineage tracing experiments with WGA-rhodamine injected in the medial segmental plate. Fixed embryos were stained with Hoechst 33258 nucleus marker and 12/101 specific antibody for muscle cells. Transverse section were performed at stage 28. Blue nuclear staining alone (a) with red WGA-rhodamine (b), with green muscle cells (c), merge (d). (D) Scheme of the experimental procedure: embryos were injected in LSF with WGA-rhodamine at stage 13 and then transverse trunk sections at stage 28 were submitted to indirect immunofluorescence with 12/101 (a) or MF20 antibodies (d). b and e, WGA-rhodamine staining; c and f, merge. Rectangle dotted line designs the area magnification at the right side (a', b', c' and d', e', f'). (E) Scheme of the experimental procedure: embryos were injected in LSF with WGA-fluorescein (green) and then submitted to ISH with *Twist1* probe or directly observed. a, *Twist1* expression analysis in an uninjected embryo; b-c, serial sections submitted to ISH with *Twist1* probe (b) or directly observed (c). (F) Scheme of the experimental procedure: *Pax1* plus *Pax9* expression analysis in an uninjected embryo (a). Serial sections of an embryo injected in LSF with WGA-rhodamine (red) submitted to ISH with *Pax1* and *Pax9* probes (c) compared to ISH negative control (b) or directly observed (d). Arrows design co-staining areas. Scale bars: 100 μ m.

Figure 2. *Twist1* is expressed in seemingly migrating sclerotome cells. (A) Scheme of the experimental procedure: expression of the somite patterning genes *Meox2*, *Tcf15*, *Foxc1*, *Foxc2* and *Twist1* in comparison to *Myod1* on trunk transverse sections between stages 20 and 34. The analyzed gene is indicated at the bottom right of each picture. Arrows design the

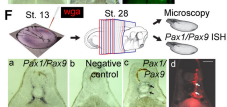
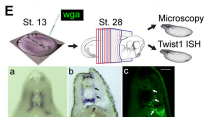
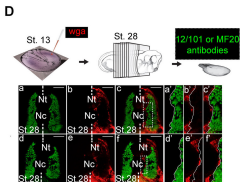
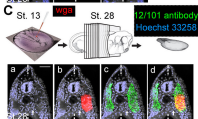
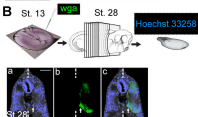
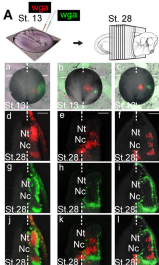
stained sclerotome. (B) Postero-anterior expression of *Twist1* at stage 28 on transverse section (a to f). (C) Somitic ventral migrating cells (arrows) on postero-anterior transverse sections (a to e). Orderly serial transverse sections stained by myotome marker (12/101 antibody, green), membrane lipophilic diI (red), and Hoechst 33258 nucleus marker (blue). DiI is used here to stain all cells. Rectangle dotted line designs the area magnification at the right side of each section (a' to e'). Scale bars: 100 μ m.

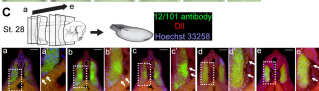
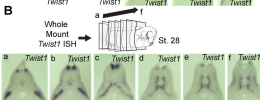
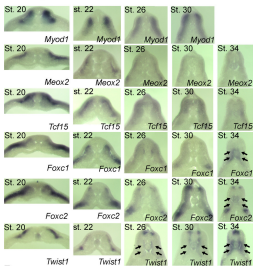
Figure 3. The WGA tracer co-localizes with *Twist1*-expressing cells in the ventral route

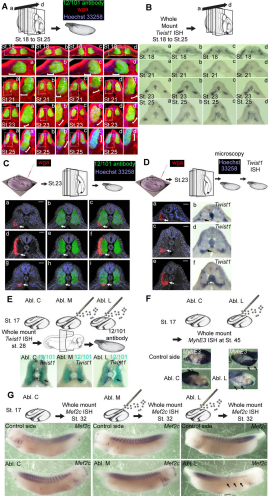
(A) Scheme of the experimental procedure: postero-anterior sections of embryos (a to d) at stages 18, 21, 23 and 25 were stained by a myotome marker (12/101 antibody, green), WGA-rodhamine (red), and Hoechst 33258 nucleus marker (blue). WGA is used here to mark all cells and to better visualize the notochord contour. Brackets design somitic ventral migrating cells. Rectangle dotted line designs the area magnification at the bottom (stage 18) or at the right side (stages 21, 23 and 25) of each section (a' to d'). Scale bars: 100 μ m. (B) Postero-anterior expression of *Twist1* at stage 18, 21, 23 and 25 on transverse sections (a to d). (C) Scheme of the experimental procedure: lineage tracing experiments with WGA-rhodamine (red) injected in the *Meox2* expression domain at stage 13, fixed and sectioned at stage 22/23, then sections were stained with a myotome marker (12/101 antibody, green) and Hoechst 33258 nucleus marker. (D) Scheme of the experimental procedure: lineage tracing experiments with WGA-rhodamine (red) injected in the *Meox2* expression domain at stage 13, fixed and sectioned at stage 22/23, then slides were stained with Hoechst 33258 nucleus marker. The same slides were next submitted to *in situ* hybridization with *Twist1* probe to evaluate co-staining of WGA tracer with *Twist1* expression. Arrows design cells labelled by WGA-rhodamine. (E), (F) and (G). Ablation of lateral somitic domain prevents sclerotome and hypaxial muscles formation. Scheme of the experimental procedure: ablation experiments of the lateral somitic domain at stage 17. Embryos were fixed and analyzed for *Twist1* (E), *MyhE3* (F) and *Mef2c* (G) expressions (arrows) and in comparison with 12/101 staining in E. Abl. C, sham-operated embryos. Abl. M, medial domain was removed. Abl. L, lateral domain was removed. For statistical analysis, number of embryos is as follows: same expression, faint decreased, strong decreased. Kruskal-Wallis test and Dunn's multiple comparisons test revealed significant difference between Abl. L with Abl. C or Abl. M, for both *Twist1* $p < 0.0001$, (Abl. C 13, 2, 0; Abl. M 13, 3, 0; Abl. L 0, 2, 13), and *Mef2c* $p < 0.0001$, (Abl. C 24, 0, 0; Abl. M 15, 6, 0; Abl. L 3, 0, 16) expressions. Mann-Whitney test revealed significant

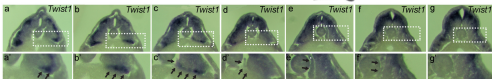
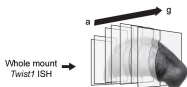
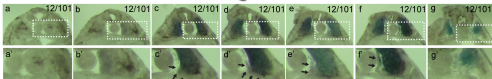
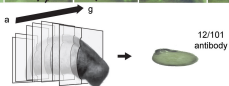
difference between Abl. L with Abl. C for *MyhE3* expression $p < 0.0001$ (Abl. C 15, 0, 0; Abl. L 0, 5, 10)

Figure 4. *Twist1* marks lateral migrating sclerotome cells in axolotl. Scheme of experimental procedure: Postero-anterior transverse sections (a to g) of axolotl embryos at stage 31 after whole mount *in situ* hybridization with *Twist1* (A) or immunohistochemistry with 12/101 antibody (B). Arrows design migrating sclerotome progenitors. Rectangle dotted line designs the area magnification at the bottom of each panel (A and B, a' to g'). (C) **Schematic representation of somite compartmentalization in *Xenopus*.** MSC domain is located at LSF at early neurulation (stage 13). Lateral *Myod1* expression domain could overlap the MSC domain. At mid-neurulation (stage 17/18), differentiation of lateral myotome is initiated and early markers of dermomyotome and sclerotome like *Pax3* (della Gaspera et al, 2012b) and *Twist1* begin to be expressed. At stage 22 lateral cells migrate dorsally to give rise the dermomyotome. Next, sclerotome progenitors expressing *Twist1* migrate ventrally to give rise to the sclerotome located medially at stage 28. (D) **Evolution of somite compartmentalization based on axochord hypothesis.** The axochord hypothesis (Brunet et al, 2015) proposes that the notochord evolves from a ventromedian muscle (the axochord in annelids) present in *Urbilateria*, the last common ancestor of bilaterian animals, and suggests that transverse muscles attached to it, could give rise to the ancestral myotome in cephalochordates. The origin of MSCs in *Urbilateria* is unknown. MSCs probably already exist in cephalochordates. The transition from anamniotes to amniotes is characterized by extension of MSCs domain at the expense of ancestral myotome. The chordate dorso-ventral axis is inverted compared with *Urbilateria*. VM, ventromedian muscle; TM, transverse muscle; M, medial somite; L, lateral somite.

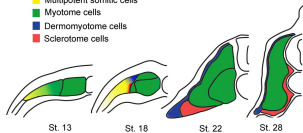






A**B****C**

- Multipotent somitic cells
- Myotome cells
- Dermomyotome cells
- Sclerotome cells

**D**