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著者	Shiomi Kosuke, Yamazaki Atsuko, Kagawa Mitsuyoshi, Kiyomoto Masato, Yamaguchi Masaaki
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Par6 regulates skeletogenesis and gut differentiation in sea urchin larvae

Kosuke Shiomi^{1, §}, Atsuko Yamazaki^{1, †}, Mitsuyoshi Kagawa¹, Masato Kiyomoto², and Masaaki

Yamaguchi^{1, *}

¹Division of Life Science, Graduate School of Natural Science and Technology, Kanazawa University,

Kakuma, Kanazawa 920-1192, Japan

²Marine and Coastal Research Center, Ochanomizu University, Kouyatsu, Tateyama 294-0301, Japan

[§]Present address: Department of Regenerative Medicine, National Institute for Longevity Sciences,

Gengo, Oobu 474-8522, Japan

[†]Present address: Research Center for Marine Biology, Tohoku University, Asamushi, Aomori 039-3501,

Japan

K. Shiomi and A. Yamazaki contributed equally to this study.

* Author for correspondence

E-mail: myama@kenroku.kanazawa-u.ac.jp

Tel: +81-76-264-6233

Fax: +81-76-264-6215

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Abstract

Partitioning-defective (*par*) genes were originally identified as genes that are essential for the asymmetric division of the *Caenorhabditis elegans* zygote. Studies have since revealed that the gene products are part of an evolutionarily conserved PAR-aPKC system involved in cell polarity in various biological contexts. In this study, we analyzed the function of *par6* during sea urchin morphogenesis by morpholino-mediated knockdown and by manipulation swapping of the primary mesenchyme cells (PMCs). Loss of Par6 resulted in defects in skeletogenesis and gut differentiation in larvae. Phenotypic analyses of chimeras constructed by PMC swapping showed that Par6 in non-PMCs is required for differentiation of archenteron into functional gut. In contrast, Par6 in both PMCs and ectodermal cells cooperatively regulates skeletogenesis. We suggest that Par6 in PMCs plays an immediate role in the deposition of biomineral in the syncytial cable, whereas Par6 in ectoderm may stabilize skeletal rods via an unknown signal(s).

Introduction

Partitioning-defective (*par*) genes, which were originally identified in *Caenorhabditis elegans*, regulate asymmetric division of the zygote (Kemphues et al. 1988; Guo and Kemphues 1995; Watts et al. 1996; Tabuse et al. 1998). Subsequent studies have revealed that the gene products are part of an evolutionarily conserved PAR-aPKC system involved in cell polarity in various biological contexts. The PAR-aPKC system converts initial polarity cues in the establishment of complementary membrane domains along the polarity axis. In most cases, this is achieved by mutually antagonistic interactions between the aPKC-PAR3-PAR6 complex and PAR1 or PAR2 localized opposite (Suzuki and Ohno 2006; Goldstein and Macara 2007).

In sea urchins, the Par6-aPKC complex mediates establishment of the apicobasal polarity of eggs and early embryos via directed insertion of basolateral membrane in the cleavage furrow (Alford et al. 2009). On the other hand, Prulière et al. (2011) showed that aPKC localized at the base of cilia controls ciliogenesis in sea urchin embryos during the blastula and gastrula stages. In this study, we analyzed the function of *par6* in the morphogenesis of sea urchin larvae by both morpholino-mediated knockdown and manipulation through swapping the primary mesenchyme cells (PMCs). We show that Par6 is required

for skeletogenesis and gut differentiation in sea urchin larvae, and suggest a novel function of Par6 in the biomineralization.

Materials and methods

Animals and embryos

Adult *Hemicentrotus pulcherrimus* were collected near the Noto Marine Laboratory, Kanazawa University, the Marine and Coastal Research Center, Ochanomizu University, and the Research Center for Marine Biology, Tohoku University. Gametes were collected by intracoelomic injection of 0.5 M KCl. Embryos were cultured at 15°C in Jamarin U artificial seawater (ASW; Jamarin Laboratory).

Cloning of *Hp-p16*

Total RNA was purified using Sepasol-RNA I Super (Nacalai Tesque) and DNase I (Takara Bio Inc.). A partial fragment of *Hp-p16* was isolated by PCR using GoTaq DNA Polymerase (Promega) and cDNA from embryos at the mesenchyme blastula and gastrula stages. The primers used were as follows: *p16*-forward, 5'-CCTTCATCGCTCTACTTGCTTTTCATCGC-3' (coding for TFIALLAFIA in the signal sequence), and *p16*-reverse, 5'-ATACCAAACGCCATGCCGCTCTTCT-3' (coding for QKSGMAFGI in

the transmembrane domain and the N-terminal flanking region). The 3'-fragment of *Hp-p16* cDNA was obtained by RACE using the GeneRacer Kit (Invitrogen), *p16*-forward primer (described above), and KOD FX DNA Polymerase (TOYOBO). The PCR products were cloned into the pTA2 vector (TOYOBO) and sequenced using the Genetic Analyzer 3100 (Applied Biosystems) with the Big Dye Terminator v3.1 Cycle Sequencing Kit (Applied Biosystems). Predicted Hp-P16 sequence was 96% identical to residues 13-173 of Sp-P16 (Accession No. NM_214646). The sequence has been deposited in the DNA Data Bank of Japan as *Hp-p16* (Accession No. BAK74802).

Preparation of constructs used for in vitro transcription

Modified pBluescript RN3 (pBluescript RN3'; Nishimura et al. 2004) was used to make expression constructs. For rescue experiments, a truncated construct lacking the 21 bp following the initiation methionine codon was generated by inverse-PCR, using *Hp-par6* in pBluescript RN3' (Shiomi and Yamaguchi 2008) and KOD FX DNA polymerase (TOYOBO). The primers used were as follows: Truncate-*par6*-forward, 5'-GCGAAAAATGGATCCACGACAAT-3'; and Truncate-*par6*-reverse, 5'-CATTTGAAGCGGGAAATCATCAAT-3'. We confirmed the sequence of the construct before transcription.

Microinjection

The morpholino oligonucleotide (Mpar6) and a standard control oligo were obtained from Gene Tools.

The sequence and positions of Mpar6 with respect to the translation initiation site were as follows:

5'-CCATGATGCCACCGAGTCCCGCAT-3' (+1 to +25). Capped RNA was transcribed from a

linearized template plasmid using mMessage mMachine T3 Kit (Ambion) according to the

manufacturer's protocol. RNA lacking an Mpar6-target site was diluted to 3.3 pg/ μ l in 40% glycerol; ~5

μ l of each solution was injected into fertilized eggs as described by Gan et al. (1990).

Microsurgery

Before microsurgery, the embryos were treated with double-strength Jamarin U for 1-2 min to stop their

swimming and placed in micromanipulation chambers filled with ASW. Micromanipulation to remove

PMCs was carried out according to the method of Ettensohn and McClay (1988). A microneedle loaded

with ASW was inserted into the blastocoel of a mesenchyme blastula at 24 h, and PMCs were washed

away by a stream of ASW. In PMC-swapping experiments, donor embryos were stained with 5 μ g/ml

rhodamine B isothiocyanate (RITC; Research Organic Inc.) for 1.5 h, after which a microneedle was

inserted into the blastocoel and PMCs were sucked out. After PMCs were collected from two or three embryos, approximately 60 PMCs were injected into the blastocoel of one host embryo.

RT-PCR

To estimate the expression levels of marker genes in embryos at the mesenchyme blastula stage, cDNA was synthesized from total RNA using ReverTra ACE (Toyobo) and random 9mer oligonucleotides. PCR was carried out using ThermoStart Taq DNA polymerase (ABgene). For each PCR reaction, cDNAs derived from 0.2 and 0.004 embryos were used to amplify markers and *ubiquitin*, respectively. Primer sequences were as follows: *delta*-F, 5'-GTACGTGTCGCAATGAAG-3'; *delta*-R, 5'-AACAGTGGTCACGGATCT-3'; *sm50*-F, 5'-GGCTAGTATTGTAGCCTT-3'; *sm50*-R, 5'-GGCGAATCCGTTAGGATA-3'; *endo16*-F, 5'-GCAACTTCCGATCATGTTGT-3'; *endo16*-R, 5'-GCGATTCTCCTTGTACTC-3'; *ubiquitin*-F, 5'-CTCACAGGCAAGACCAT-3'; and *ubiquitin*-R, 5'-GTCTCTCAATTTGTCTATAGCT-3'.

Whole-mount *in situ* hybridization (WMISH)

WMISH was performed according to the method of Minokawa et al. (2004) for *par6* or to that of Hibino et al. (2004) for *sm30*, *p16*, and *apkc*. Digoxigenin (DIG)-labeled probes were transcribed using DIG-labeling mix (Roche) from *Hp-apkc*, *par6* (Shiomi and Yamaguchi 2008), *sm30* (Kitajima et al. 1996), and *Hp-p16* cDNAs (this study).

Antibodies and whole-mount immunostaining

For immunostaining, we fixed embryos and larvae according to the method of Hibino et al. (2004). Fixed embryos were washed with PBS containing 0.1% Triton X-100 (PBT), blocked with 2% bovine serum albumin (BSA), and then incubated for 2 h with goat polyclonal anti-PAR6 antibody (diluted 1:100 with PBT; Santa Cruz, sc-14405; Alford et al. 2009), the B2C2 monoclonal antibody (diluted 1:500 with PBT; Leaf et al. 1987), or the P4 monoclonal antibody (diluted 1:200 with PBT; Shimizu et al. 1988). Although the reactivity of the P4 antibody with PMCs is quite similar to that of the B2C2 antibody, Shimizu-Nishikawa et al. (1990) showed that these antibodies react with the same group of glycoproteins, including MSP130, but their epitopes are different. Alexa Fluor 488-conjugated anti-goat IgG or anti-mouse IgG (Molecular Probes) was used as the secondary antibody.

Results

Par6 is accumulated along skeletal rods and in digestive tract cells in larvae

The distribution of Par6 protein in *H. pulcherrimus* larvae was determined using an antibody, which specifically recognizes Par6 in a variety of vertebrate and sea urchin species (Alford et al. 2009).

Although *par6* is ubiquitously expressed during the development (Shiomi and Yamaguchi 2008; Fig. 1a–e), in pluteus larvae, Par6 signals were detected along the skeletal rods (Fig. 1i–k); Par6 appeared to be localized in the syncytial cable of PMCs, since few signals were observed in the PMC cell body or spicule space, where calcite had been dissolved during fixation (asterisk and inset in Fig. 1i; also see Discussion). In addition, Par6 was distributed in endodermal cells forming the boundary between the midgut and hindgut (arrows in Fig. 1f,g,i,j), and cells encircling the anus (arrowheads in Fig. 1h,k).

We injected a morpholino oligo against *par6* mRNA (Mpar6) into fertilized eggs. Larvae including Mpar6 did not exhibit the Par6 accumulations observed in controls (Fig. 2o,p), indicative of the efficacy of Mpar6 on translational inhibition.

Par6 is required for skeletogenesis and gut differentiation

Mpar6-injected zygotes cleaved normally, including unequal fourth cleavages that formed micromeres, and developed into mesenchyme blastulae via swimming blastulae (Fig. 2a,h). Similar to controls, embryos including Mpar6 (Mpar6 embryos) expressed early marker genes, including *delta* (presumptive secondary mesenchyme cell), *sm50* (PMC), and *endo16* (endomesoderm), indicating that early endomesoderm specification occurred in Mpar6 embryos (Fig. 3a).

Morphological defects in Mpar6 embryos emerged after gastrulation during two processes: skeletogenesis and gut differentiation. When controls developed into prism larvae at 48 h (Fig. 2c,d), Mpar6 embryos formed spot-like or triradial spicule rudiments within the ventrolateral PMC clusters (Fig. 2j,k). However, subsequent growth of the skeleton was severely impaired (Fig. 2m); the spicule largely remained as it was in Mpar6 embryos at 48 h. We note that Mpar6 embryos, especially after gastrulation, had a thicker ectoderm wall than uninjected controls, which reduced the size of the experimental embryo/larva (Fig. 2i,j,l).

The larval skeleton is elaborated through interactions between PMCs and ectodermal cells (Armstrong et al. 1993; Ettensohn and Marinda 1993; Guss and Ettensohn 1997; Hardin and Armstrong 1997). Duloquin et al. (2007) demonstrated that localized VEGF acts as both a guidance cue for the formation of ventrolateral PMC clusters and a signal for the differentiation of skeletogenic cells. On the

other hand, Röttinger et al. (2008) showed that FGF signaling plays an essential role in the guidance of PMC migration and the direction of skeletal morphogenesis. However, defects in skeletogenesis induced by *Mpar6* injection were unlikely due to either the loss of skeletogenic cell differentiation or a lack of ectodermal guidance/direction cues required for skeletogenesis, since in *Mpar6* embryos, mesenchyme cells expressed the PMC-specific P4 antigen (Fig. 3j; Shimizu et al. 1988) and VEGF/FGF-dependent skeletogenic genes, such as *sm50* and *sm30* (Fig. 3a,m; Duloquin et al. 2007; Röttinger et al. 2008), and because their descendants expressing the B2C2 antigen (Fig. 3l; Leaf et al. 1987) were allocated on the inner ectoderm wall in a pattern that prefigures the rod branching patterns, i.e., the anterolateral, transverse, postoral, and body rods (Fig. 3l). Furthermore, skeletogenic cells expressed *p16* that is essential for skeletal rod elongation (Fig. 3n,o; Cheers and Etensohn 2005). However, the size of the PMC ring in *Mpar6* larvae was smaller than in controls, probably due to dwarfish ectoderm patterning and/or a lack of accompanying growth of skeletal rods in the experimental larvae.

Another defect in *Mpar6* larvae was lack of gut differentiation. Although gastrulation was delayed for several hours, *Mpar6* embryos developed archenteron, from the tip of which the secondary mesenchyme cells (SMCs) had been delaminated (Fig. 2i,j). Subsequently, *Mpar6* larvae developed SMC-derived tissues, such as pigment cells (Fig. S1a), circumesophageal muscle cells, and coelomic

pouches, although the number of pigment cells was lower than in controls in some batches. In contrast, the larva did not develop a fully differentiated gut in either morphology or function; the segmentation of archenteron was repressed, and the mid/hindgut region lacked alkaline phosphatase activity (Fig. 2n).

Co-injection of *par6* mRNA without the Mpar6-target sequence (~15 pg/egg) rescued Mpar6 larvae from defects in skeletogenesis and gut differentiation, but partially in ectoderm patterning.

Co-injected zygotes developed into pluteus larvae with a slightly small skeleton, but normal in pattern (Fig. 2r,u) as well as tripartite gut with alkaline phosphatase activity (Fig. 2s,v). However, the larva was smaller and distinct from control larvae in shape (arrowhead in Fig. 2q). In contrast, zygotes injected with mRNA alone developed normally (Fig. S1c,d). These observations indicate a requirement for Par6 for skeletogenesis and gut differentiation in sea urchin larvae; they also suggest that higher and/or longer doses of Par6 may be required for correct ectoderm patterning, since co-injection of a lower dose of mRNA (~10 pg/egg) rescued morphants from defects in skeletogenesis and gut differentiation, but not from dwarfish ectoderm patterning (Fig. S1b,b').

Par6 controls skeletogenesis in PMCs and ectoderm cooperatively

Skeletogenesis in sea urchin larvae is controlled by both PMCs and ectodermal cells (Armstrong et al.

1993; Ettensohn and Marinda 1993; Guss and Ettensohn 1997; Hardin and Armstrong 1997; Duloquin et al. 2007; Röttinger et al. 2008). On the other hand, Hamada and Kiyomoto (2003) demonstrated that unidentified signals from PMCs regulate gut differentiation. When PMCs are deleted from the mesenchyme blastula, both the constrictions in archenteron and the expression of alkaline phosphatase are delayed. To clarify whether defects in skeletogenesis and gut differentiation in *par6* morphants are due to loss of Par6 in PMCs or the rest of embryos, or both, we carried out PMC-swapping experiments using mesenchyme blastulae at 24 h after fertilization.

Sham controls, which consisted of a PMC-deleted host embryo and PMCs transplanted from sibling rhodamine-labeled embryos, developed into pluteus larvae indistinguishable from the unmanipulated controls (Fig. 4a–f), although one and two of the 13 chimeras had an abnormal skeletal pattern and extra rods, respectively (Table 1). All transplanted PMCs examined were arranged along the skeletal rods in the chimeras. To examine the effects of morpholino injection on eggs we constructed chimeric embryos composed of a PMC-deprived host and PMCs including a control morpholino, and confirmed that their phenotypes were practically identical to those of the sham control (3/3 cases, data not shown). Thus, we analyzed the phenotypes of embryos with Mpar6 present only in PMCs (Fig. 4g–l). The chimera developed into gastrula-like larvae with spicule rudiments on two ventrolateral positions at 48 h

(24 h after manipulation; Fig. 4h,h'; 9/11 cases). However, skeletal rod elongation from the rudiments was delayed in the chimera, although PMC descendants were arranged in a ring pattern with two clusters plus two cables that will form the anterolateral rods, like in control chimeras (Fig. 4h,i). This defect in skeletogenesis continued during the next 24 h; the skeleton remained stunted, and the branching pattern was irregular (Fig. 4k; Fig. S2; Table 1). When compared to spicules in *par6* whole morphants (Fig. 2m), however, restoration of skeletal rod elongation was evident in the chimera (Fig. 4k; Fig. S2). This observation suggests that Par6 cooperatively regulates skeletogenesis in both PMCs and the rest of the embryo (probably in the ectoderm). Noticeably, the chimera lacked a connection of bilateral body rods at the vertex (Fig. 4k'; Fig. S2; Table 1), although PMC descendants were correctly allocated in the region (arrowhead in Fig. 4l'). In contrast, the chimera developed a tripartite gut with alkaline phosphatase activity (Fig. 4t; Table 1), indicating that Par6 is not involved in gut differential signals from PMCs.

Par6 in non-PMCs regulates gut differentiation

Next, we analyzed the phenotypes of complementary chimeras composed of a PMC-less host including *Mpar6* and normal PMCs. Like *par6* whole morphants (Fig. 2i, j, l), the chimera had a thicker ectoderm wall and was therefore smaller than control or complementary chimeras (Fig. 4m,p), supporting the

notion that Par6 is required for correct ectoderm patterning. At 48 h, some chimeras generated extra small spicules in addition to two spicules in the ventrolateral PMC clusters (Fig. 4n'; 4/6 cases); the latter two constituted most of the larval skeletal rods, while the former gave rise to small spines on the body rods (see A1 and B3 in Fig. S3). Unlike complementary chimeras, the chimera formed a connection of bilateral body rods at the vertex (Fig. 4q,q'; Fig. S3; Table 1). In contrast, skeletal rod elongation was retarded in the oral side of chimeras (Fig. 4q,q'; Fig. S3; Table 1); the defect was evident in both the postoral rods and anterolateral rods toward the oral hood, when compared to those of complementary chimeras (Fig. 4k,k'; Fig. S3). These observations not only support the notion that Par6 cooperatively regulates skeletogenesis in PMCs and ectoderm, but also indicate that ectoderm produces an unidentified Par6-dependent signal(s) that is necessary for skeletal rod elongation particularly in the oral field. On the other hand, the chimera did not develop tripartite gut with alkaline phosphatase activity (Fig. 4u; Table 1), indicating that Par6 in non-PMCs is required for gut differentiation. Distribution of Par6-accumulating cells in the digestive tract (Fig. 1f-k) suggests that Par6 may control gut differentiation via the apical constriction.

Expression of *apkc* is not affected in *par6* morphants, but in cells interspersed in the ciliary band

To examine whether loss of Par6 affects the *apkc* expression, we analyzed the *apkc* expression in *par6* morphants. In controls, zygotic *apkc* expression starts in the vegetal plate, followed by expression restricted to PMCs and the archenteron tip during gastrulation (Fig. 5a,b), and finally, the expression fades by the late gastrula stage. From the prism stage, the second phase of *apkc* expression starts in three cell types: ectodermal cells interspersed in the ciliary band, skeletogenic cells at the vertex, and stomach cells (Fig. 5c). The *apkc*-expressing cells in the ciliary band do not appear to match the serotonergic cells (four to six cells in the apical ganglion; Bisgrove and Burke 1986) or synaptotagmin-expressing cells (flask-shaped cell bodies on the basal side of ectoderm; Nakajima et al. 2004), although *apkc*-expressing cells may partially overlap the neural cells. In *par6* morphants, the first and second phases of *apkc* expression occurred as they do in controls (Fig. 5d–f), except for the expression in ectodermal cells in the oral hood (arrowhead in Fig. 5f). This *apkc* downregulation induced by loss of Par6 is probably indirect, since Par6 is not expressed in the ciliary band of larvae (Fig. 2o), and because dwarfish ectoderm patterning at earlier stages is a constant phenotype observed in *par6* morphants (Fig. 2i; Fig. 3l) and chimeras with loss of Par6 in non-PMC tissues (Fig. 4m). Defects of the chimera in skeletal rod elongation in the oral field may be due to the *apkc* downregulation, since the defect is a unique phenotype of chimeras with loss of Par6 in non-PMC tissues, including ectoderm.

Discussion

Par6 may function outside the polarity complex

Par6 is a component of the apical polarity complex that contains Par3 and aPKC (Suzuki and Ohno 2006; Goldstein and Macara 2007). Defects in *par6* morphants or chimeric larvae, i.e., retardation of skeletal rod elongation and gut differentiation, largely coincide with zygotic *apkc* domains (Shiomi and Yamaguchi 2008; Fig. 5a–c). Prulière et al. (2011) have shown the aPKC expression in *Paracentrotus* larvae by immunostaining; aPKC is distributed in the ciliary band ectoderm and stomach endoderm, but not in PMCs or syncytial cables. We confirmed that the expression pattern is essentially conserved in *Hemicentrotus* larvae (Fig. S4). The lack of co-expression of Par6 and aPKC in the syncytial cable or ciliary band suggests that Par6 or aPKC may control skeletogenesis outside the polarity complex.

In *Hemicentrotus* larvae, however, skeletogenic cells at the vertex exhibit relatively stronger aPKC signals than other skeletogenic cells (Fig. S4b',e). This expression may support Par6 function as the polarity complex in skeletogenesis, since one of the definite roles of Par6 in PMCs appears to generate a connection of bilateral body rods at the vertex (Fig. 4k'; Fig. S2; lack of the skeletal element is a unique phenotype of chimeras with loss of Par6 only in PMCs), and because skeletogenic cells at the

vertex exclusively express *apkc* (arrow in Fig. 5c). This in turn implies that distinct mechanisms may work in skeletogenesis between the oral and aboral sides of larvae. Alternatively, aPKC activity is required only for the initial step of subcellular localization processes of Par6 into the syncytial cable, since *apkc* is transiently expressed in PMCs (Fig. 5). Furthermore, the aPKC expression pattern is puzzling; the protein is distributed throughout the ciliary band (Fig. S4), but the mRNA is restricted to isolated cells in this band (Fig. 5c). The pattern suggests that *apkc* expression may oscillate in individual cells in the band. To examine the aPKC function in skeletogenesis, we carried out morpholino-mediated *apkc* knockdown; however, it resulted in embryonic lethality at the blastula stage (Shiomi and Yamaguchi, unpublished).

Par6 function in PMCs

Spicule formation takes place in a membrane-delineated compartment (vacuole) enclosed in the syncytial cable formed by fusion of filopodia of PMCs (Okazaki 1960). Spicule matrix proteins, such as SM30 and SM50, are tethered to calcite deposited in the vacuole (Seto et al., 2004), whereas MSP130 and P16 are localized on the surface of PMCs and their filopodia (Anstrom et al. 1987; Cheers and Etensohn 2005). In contrast, Par6 seems to be accumulated in the cortex and/or cytoplasm of the syncytial cable (Fig.

1f-k), although further examination is needed to define its precise localization. Loss of Par6 causes defects in the skeletal rod elongation, but practically no effect on expressions of spicule matrix proteins or related genes (Fig. 3), suggesting that Par6 may play a more immediate role in the deposition of biomineral. Like Par6, P16 acts at relatively late step in skeletogenesis; P16 is not required for PMC specification, ingress, migration, or fusion, but essential for skeletal rod elongation (Cheers and Etensohn 2005). Cheers and Etensohn (2005) suggest that P16 may be a receptor for ectoderm-derived signals required for skeletal rod elongation; alternatively, P16 may function in the plasma membrane or an intracellular membrane compartment to regulate calcium transport. Since phenotypes of *p16* knockdown resemble those of chimeric embryos with *Mpar6* present only in PMCs, it seems possible that Par6 relays signals transduced by P16 or cooperates with P16 to deposit biomineral.

Par6 function in ectoderm

From phenotype comparisons between whole *par6* morphants and chimeric larvae (Fig. 2m; Fig. 3e,k,q), it is evident that ectoderm emanates a Par6-dependent signal(s) that promotes skeletogenesis. In chimeras with loss of Par6 in PMCs, skeletal rods are often irregular in branching pattern or in shape (Fig. 4k; Fig. S2; Table 1), whereas no abnormal branching pattern is observed in chimeras with loss of Par6 in

non-PMCs (Fig. 4q; Fig. S3; Table 1). Furthermore, zygotes injected with *par6* mRNA develop into larvae with normal skeleton (Fig. S1c,d). These observations suggest that, unlike VEGF or FGF, the ectoderm-derived signal does not act as positional or directional guidance cues in skeletogenesis, but appears to maintain or stabilize spicule materials that have been deposited in the vacuole. This may explain observations that loss of the signal decreases skeletal rods in size in whole morphants compared to chimeras with loss of Par6 in PMCs (Fig. 2m; Fig. 4k) and that Par6 does not appear to be localized in ectoderm overlying growing skeletal rods (Fig. 2o).

Biom mineralization in echinoderms

Biom mineralized tissues are considered to have arisen independently in vertebrates and echinoderms (Bottjer et al. 2006; Livingston et al. 2006). Their matrix proteins may be an example of evolutionary convergence. Secreted proteins associated with tooth and bone in vertebrates are believed to have arisen from an ancestral gene encoding secreted protein that is acidic and cysteine-rich (Kawasaki and Weiss 2006). In contrast, many of the secreted biom mineralization proteins in echinoderms probably arose from an ancestral C-lectin domain-containing protein (Livingston et al. 2006). On the other hand, aPKC is involved in at least two independent signaling cascades that control cell polarity and cell survival; in the

cascades, aPKC function is modulated by different partner proteins, Par6 and p62, respectively (Moscat and Diaz-Meco 2000). In biomineralization of mice, p62 is an essential mediator of aPKC function during osteoclastogenesis (Durán et al. 2004), while Par6 regulates skeletogenesis in sea urchin larvae; however, the requirement of aPKC function for skeletogenesis remains to be clarified. Further studies will reveal unique molecular mechanisms of biomineralization in sea urchins, which may reveal another role of Par6 and/or aPKC.

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Figure legends

Fig. 1 Distributions of *par6* mRNA and Par6 protein. **a–e** Expressions of *par6* analyzed by WMISH. This gene is ubiquitously expressed in the egg (**a**), 32-cell stage embryo (**b**), mesenchyme blastula (**c**), gastrula (**d**), and pluteus larva (**e**). **f–k** Distributions of Par6 in larvae (72 h) analyzed by immunostaining. **f, i** Ventral views of aboral half of a larva. **g, h, j, k** Ventrolateral views. Signals are distributed along skeletal rods (**i–k**); Par6 seems to be localized in the syncytial cable of PMCs, since few signals are observed in the PMC cell body or spicule space, where calcite has been dissolved during fixation (asterisk and inset in **i**). In addition, signals are detected in endodermal cells forming the mid-hindgut boundary (arrows in **f, g, i, j**) and cells encircling the anus/blastopore (arrowheads in **h, k**). Insets in **i, j**, and **k** show high-magnification images around the asterisk, arrow, and arrowhead regions, respectively.

Fig. 2 Par6 is required for skeletogenesis and gut differentiation in sea urchin larvae. **a–g** Control embryos and larvae. **h–n** Embryos and larvae derived from zygotes injected with Mpar6. Embryos including Mpar6 developed almost normally up to the mesenchyme blastula stage (**a, h**). Although there was a delay in gastrulation for several hours, the embryo developed into gastrulae, which had a thicker ectoderm wall, and were thus smaller than controls (**b, i**). In larvae including Mpar6, growth of the

skeletal rods was severely impaired, although spicule rudiments formed within the ventrolateral PMC clusters (**k, m**). In addition, the larva did not develop a fully differentiated gut in either morphology or function (**l, n**); growth of the midgut and segmentation of archenteron were repressed (**l, n**), and the mid/hindgut region lacked alkaline phosphatase activity (**n**). **o, p** Par6 distributions of control and Mpar6 larvae. In a control larva (**o**), Par6 is accumulated in syncytial cables (asterisk), endodermal cells (arrow), and cells around blastopore (arrowhead), whereas a larva including Mpar6 does not exhibit either of the accumulations (**p**). **q–v** Larvae including Mpar6 and *par6* mRNA without Mpar6 target sequence. Co-injected zygotes developed into pluteus larvae with a slightly smaller skeleton, but normal in pattern (**q, r, t, u**), as well as tripartite gut with alkaline phosphatase activity (**s, v**). Inset in **s** shows a high-magnification image around the mid/hindgut region, where arrowhead indicates the hindgut with alkaline phosphatase activity. However, the larva was distinct from control larvae in shape; the posterodorsal side of the aboral ectoderm wall was fallen (arrowhead in **q**), when compared to controls (**e, j, k, n, v** Dorsal views. **o, p, t, u** Ventral views

Fig. 3 Morpholino-mediated *par6* knockdown did not affect the expression of early endomesoderm specification genes or the differentiation of the skeletogenic cell. **a** The expressions of marker genes in

embryos including *Mpar6*. Total RNA was extracted from control embryos and embryos including *Mpar6* at the mesenchyme blastula stage (24 h). The expression of marker genes *delta* (presumptive SMC), *sm50* (PMC), *endo16* (endomesoderm), and *ubiquitin* (ubiquitous) was analyzed by RT-PCR. Numerals below indicate the number of PCR cycles. Expression levels of these genes in embryos including *Mpar6* were comparable to those in control embryos. **b–e, i–l** Expression of the P4 (**b–e**) and B2C2 antigens (**i–l**) analyzed by immunostaining. In either controls (**b–e**) or *par6* morphants (**i–l**), PMCs expressing the P4 antigen (**b, c, i, j**) ingressed at 24 h, and skeletogenic cells expressing B2C2 antigen (**d, e, k, l**) were allocated on the inner ectoderm wall in a pattern that prefigures the rod-branching patterns in larvae at 48 h. However, the PMC-ring in *par6* morphants was smaller than that in controls (**e, l**). **f–h, m–o** Expressions of *sm30* (**f, m**) and *p16* (**g, h, n, o**) analyzed by whole-mount *in situ* hybridization (WMISH). Like in controls (**f**), *sm30* was expressed in skeletogenic cells in the *par6* morphant (**m**). In either controls or *par6* morphants, *p16* was expressed in PMCs at the mesenchyme blastula stage (26 h; **g, n**). At the pluteus stage (72 h), the expression was restricted to PMCs at sites of skeletal rod elongation, such as the tips of the postoral, anterolateral, and body rods (**h, o**), although skeletal rod elongation was severely impaired in *par6* morphants (**o**). **d–f, k–m** Dorsal views. **g, n** Vegetal views. **h, o** Ventral views

Fig. 4 Development of chimeric embryos constructed by PMC swapping. Illustrations on the left indicate combinations of a PMC-less host (upper, from normal or Mpar6-injected embryos) and transplanted PMCs (lower in parentheses, from normal or Mpar6-injected embryos), in which cells including Mpar6 are blue. **a–u** Lateral views. **a'–r'** Ventral views. **a–f, s** Control chimeras composed of a PMC-less host and transplanted PMCs labeled with rhodamine. The chimera developed normally like unmanipulated controls. **g–l, t** Chimeras composed of a PMC-less host and PMCs including Mpar6. Although PMCs were arranged in a ring pattern with two ventrolateral clusters in the chimera, rod elongation from spicule rudiments was delayed (48 h; **h, h'**), and the branching pattern was irregular in most larvae (72 h; **k, k'**). Note that the chimera lacked a connection of bilateral body rods at the vertex (**k'**), in spite of correct allocation of skeletogenic cells in the region (arrowhead in **l'**). Arrowhead in **l** indicates rhodamine-positive cells in the gut region. The chimera developed a tripartite gut with alkaline phosphatase (AP) activity (**t**). **m–r, u** Chimeras composed of a PMC-less host including Mpar6 and normal PMCs. The chimera had a thicker ectoderm wall (**m, p**) and thus was smaller than control chimeras (**a, d**). At 48 h (24 h after manipulation), some chimeras generated extra small spicules (small arrowheads in **n'**) in addition to two spicules in ventrolateral PMC clusters (large arrowheads in **n'**). The latter two constituted most of the larval skeletal rods at 72 h (**q**). Note that rod elongation was retarded in

the oral field, whereas the connection of bilateral body rods was formed at the vertex (**q'**). Arrowhead in **r** indicates rhodamine-positive cells in the gut region. The chimera did not develop a functional gut (**u**), unlike complementary chimeras (**t**)

Fig. 5 Expression of *apkc* in *par6* morphants analyzed by WMISH. **a, b, d, e** Lateral views. **c, f** Dorsal views. **a–c** Control embryos and larvae. The first phase of zygotic *apkc* expression starts in the vegetal plate at the blastula stage (**a**). Expression is maintained in PMCs and the archenteron tip during gastrulation (**b**), but fades at the late gastrula stage. During the prism to pluteus stages, the second phase of expression starts in three cell types (**c**): ectodermal cells interspersed in the ciliary band, skeletogenic cells at the vertex (arrow in **c**), and stomach cells. **d–f** Embryos and larvae including Mpar6. The first and second phases of *apkc* expression occurred in *par6* morphants as they did in controls (**d–f**), except for expression in ectodermal cells in the oral hood (arrowhead in **f**)

References

Alford LM, Ng MM, Burgess DR (2009) Cell polarity emerges at first cleavage in sea urchin embryos.

Dev Biol 330:12-30. doi:10.1016/j.ydbio.2009.02.039

Anstrom JA, Chin JE, Leaf DS, Parks AL, Raff RA (1987) Localization and expression of msp130, a primary mesenchyme lineage-specific cell surface protein in the sea urchin embryo. *Development* 101:255-265

Armstrong N, Hardin J, McClay DR (1993) Cell-cell interactions regulate skeleton formation in the sea urchin embryo. *Development* 119:833-40

Bidgrove BW, Burke RD (1986) Development of serotonergic neurons in embryos of the sea urchin, *Strongylocentrotus purpuratus*. *Dev Growth Differ* 28:569-74.
doi:10.1111/j.1440-169X.1986.00569.x

Bottjer DJ, Davidson EH, Peterson KJ, Cameron RA (2006) Paleogenomics of echinoderms. *Science* 314:956-960. doi:10.1126/science.1132310

Cheers MS, Etensohn CA (2005) P16 is an essential regulator of skeletogenesis in the sea urchin embryo. *Dev Biol* 283:384-396. doi:10.1016/j.ydbio.2005.02.037

Duloquin L, Lhomond G, Gache C (2007) Localized VEGF signaling from ectoderm to mesenchyme cells controls morphogenesis of the sea urchin embryo skeleton. *Development*:2293-2302.
doi:10.1242/dev.005108

Durán A, Serrano M, Leitges M, Flores JM, Picard S, Brown JP, Moscat J, Diaz-Meco MT (2004) The

- atypical PKC-interacting protein p62 is an important mediator of RANK-activated osteoclastogenesis. *Dev Cell* 6:30-309. doi:10.1016/S1534-5807(03)00403-9
- Ettensohn CA, McClay DR (1988) Cell lineage conversion in the sea urchin embryo. *Dev Biol* 125:396-409. doi:10.1016/0012-1606(88)90220-5
- Ettensohn CA, Malinda KM (1993) Size regulation and morphogenesis: a cellular analysis of skeletogenesis in the sea urchin embryo. *Development* 119:155-167
- Gan L, Zhang W, Klein WH (1990) Repetitive DNA sequences linked to the sea urchin spec genes contain transcriptional enhancer-like elements. *Dev Biol* 139:186-196. doi:10.1016/0012-1606(90)90287-S
- Goldstein B, Macara IG (2007) The PAR proteins: fundamental players in animal cell polarization. *Dev Cell* 13:609-622. doi:10.1016/j.devcel.2007.10.007
- Guo S, Kemphues KJ (1995) *par-1*, a gene required for establishing polarity in *C. elegans* embryos, encodes a putative Ser/ Thr kinase that is asymmetrically distribution. *Cell* 81:611-620. doi:10.1016/0092-8674(95)90082-9
- Guss KA, Ettensohn CA (1997) Skeletal morphogenesis in the sea urchin embryo: regulation of primary mesenchyme gene expression and skeletal rod growth by ectoderm-derived cues. *Development*

124:1899-1908

Hamada M, Kiyomoto M (2003) Signals from primary mesenchyme cells regulate endoderm

differentiation in the sea urchin embryo. *Dev Growth Differ* 45:339-350.

doi:10.1046/j.1440-169X.2003.00702.x

Hardin J, Armstrong N (1997) Short-range cell-cell signals control ectodermal patterning in the oral

region of the sea urchin embryo. *Dev Biol* 182(1):134-49. doi:10.1006/dbio.1996.8436

Hibino T, Harada Y, Minokawa T, Nonaka M, Amemiya S (2004) Molecular heterotopy in the expression

of Brachyury orthologs in order Clypeasteroidea (irregular sea urchins) and order Echinoidea (regular

sea urchins). *Dev Genes Evol* 214:546-558. doi:10.1007/s00427-004-0437-x

Kawasaki K, Weiss KM (2006) Evolutionary genetics of vertebrate tissue mineralization: the origin and

evolution of the secretory calcium-binding phosphoprotein family. *J Exp Zool B Mol Dev Evol*

15:295-316. doi:10.1002/jez.b.21088

Kemphues KJ, Priess JR, Morton DG, Cheng NS (1988) Identification of genes required for cytoplasmic

localization in early *C. elegans* embryos. *Cell* 52:311-320. doi:10.1016/S0092-8674(88)80024-2

Kitajima T, Tomita M, Killian CE, Akasaka K, Wilt FH (1996) Expression of spicule matrix protein gene

SM30 in embryonic and adult mineralized tissues of sea urchin *Hemicentrotus pulcherrimus*. *Dev*

Growth Differ 38:687-695. doi:10.1046/j.1440-169X.1996.t01-5-00012.x

Leaf DS, Anstrom, JA, Chin JE, Harkey MA, Showman RM, Raff RA (1987) Antibodies to a fusion protein identify a cDNA clone encoding msp130, a primary mesenchyme-specific cell surface protein of the sea urchin embryo. Dev Biol 121:29-40. doi:10.1016/0012-1606(87)90135-7

Livingston BT, Killian CE, Wilt F, Cameron A, Landrum MJ, Ermolaeva O, Sapojnikov V, Maglott DR, Buchanan AM, Etensohn CA (2006) A genome-wide analysis of biomineralization-related proteins in the sea urchin *Strongylocentrotus purpuratus*. Dev Biol 300:335-348. doi:10.1016/j.ydbio.2006.07.047

Minokawa T, Rast JP, Arenas-Mena C, Franco CB, Davidson EH (2004) Expression patterns of four different regulatory genes that function during sea urchin development. Gene Expr Patterns 4:449-456. doi:10.1016/j.modgep.2004.01.009

Moscat J, Diaz-Meco MT (2000) The atypical protein kinase Cs. Functional specificity mediated by specific protein adapters. EMBO Rep 1:399-403. doi:10.1093/embo-reports/kvd098

Nakajima Y, Kaneko H, Murray G, Burke RD (2004) Divergent patterns of neural development in larval echinoids and asteroids. Evol Dev 6:95-104. doi:10.1111/j.1525-142X.2004.04011.x

Nishimura Y, Sato T, Morita Y, Yamazaki A, Akasaka K, Yamaguchi M (2004) Structure, regulation, and function of *micro1* in the sea urchin *Hemicentrotus pulcherrimus*. *Dev Genes Evol* 214:525-536.

doi:10.1007/s00427-004-0442-0

Okazaki K (1960) Skeleton formation of sea urchin larvae II. Organic matrix of the spicule. *Embryologia* 5:283-320. doi:10.1111/j.1440-169X.1960.tb00096.x

Prulière G, Cosson J, Chevalier S, Sardet C, Chenevert J (2011) Atypical protein kinase C controls sea urchin ciliogenesis. *Mol Biol Cell* 22:2042-53. doi:10.1091/mbc.E10-10-0844

Röttinger E, Saudemont A, Duboc V, Besnardeau L, McClay D, Lepage T (2008) FGF signals guide migration of mesenchymal cells, control skeletal morphogenesis and regulate gastrulation during sea urchin development. *Development* 135:353-65. doi:10.1242/dev.014282

Seto J, Zhang Y, Hamilton P, Wilt F (2004) The localization of occluded matrix proteins in calcareous spicules of sea urchin larvae. *J Struct Biol* 148:123-30. doi:10.1016/j.jsb.2004.04.001

Shimizu K, Noro N, Matsuda R (1988) Micromere differentiation in the sea urchin embryo: expression of primary mesenchyme cell specific antigen during development. *Dev Growth Differ* 30:35-47.

doi:10.1111/j.1440-169X.1988.00035.x

Shimizu-Nishikawa K, Katow H, Matsuda R (1990) Micromere differentiation in the sea urchin embryo:

immunochemical characterization of primary mesenchyme cell-specific antigen and its biological role. *Dev Growth Differ* 32:629-636. doi: 10.1111/j.1440-169X.1990.00629.x

Shiomi K, Yamaguchi M (2008) Expression patterns of three Par-related genes in sea urchin embryos. *Gene Expr Patterns* 8:323-330. doi:10.1016/j.gep.2008.01.002

Suzuki A, Ohno S (2006) The PAR-aPKC system: lessons in polarity. *J Cell Sci* 119:979-987

Tabuse Y, Izumi Y, Piano F, Kemphues KJ, Miwa J, Ohno S (1998) Atypical protein kinase C cooperates with PAR-3 to establish embryonic polarity in *Caenorhabditis elegans*. *Development* 125:3607-3614

Watts JL, Etemad-Moghadam B, Guo S, Boyd L, Draper BW, Mello CC, Priess JR, Kemphues KJ (1996) par-6, a gene involved in the establishment of asymmetry in early *C. elegans* embryos, mediates the asymmetric localization of PAR-3. *Development* 122:3133-3140

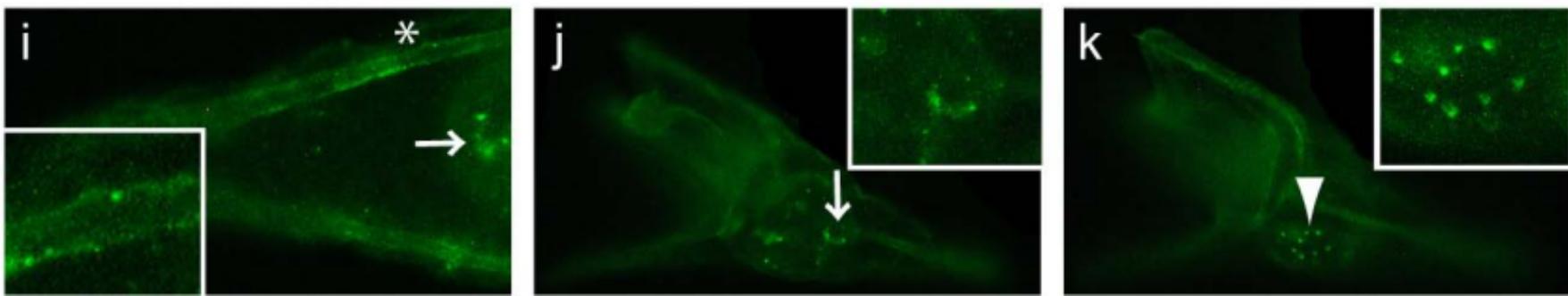
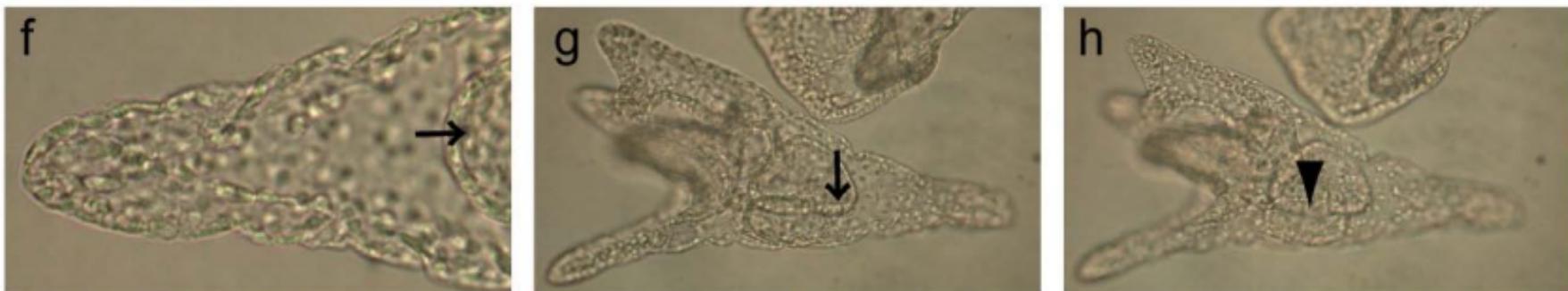
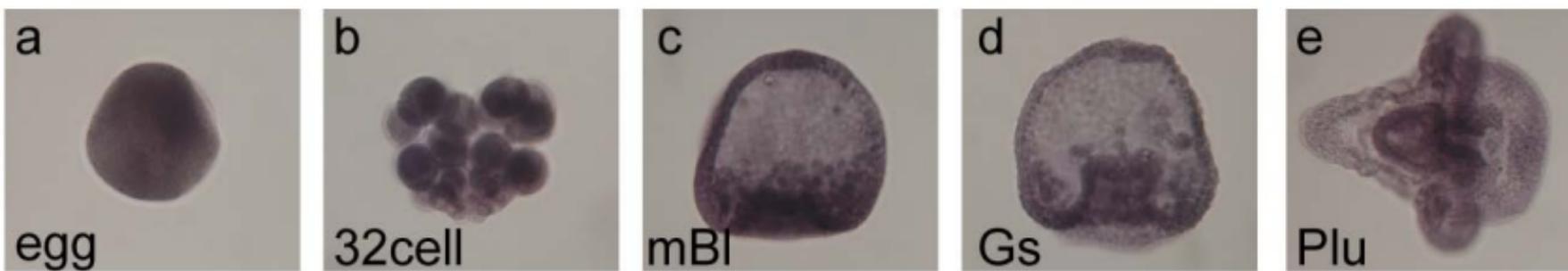
Table 1

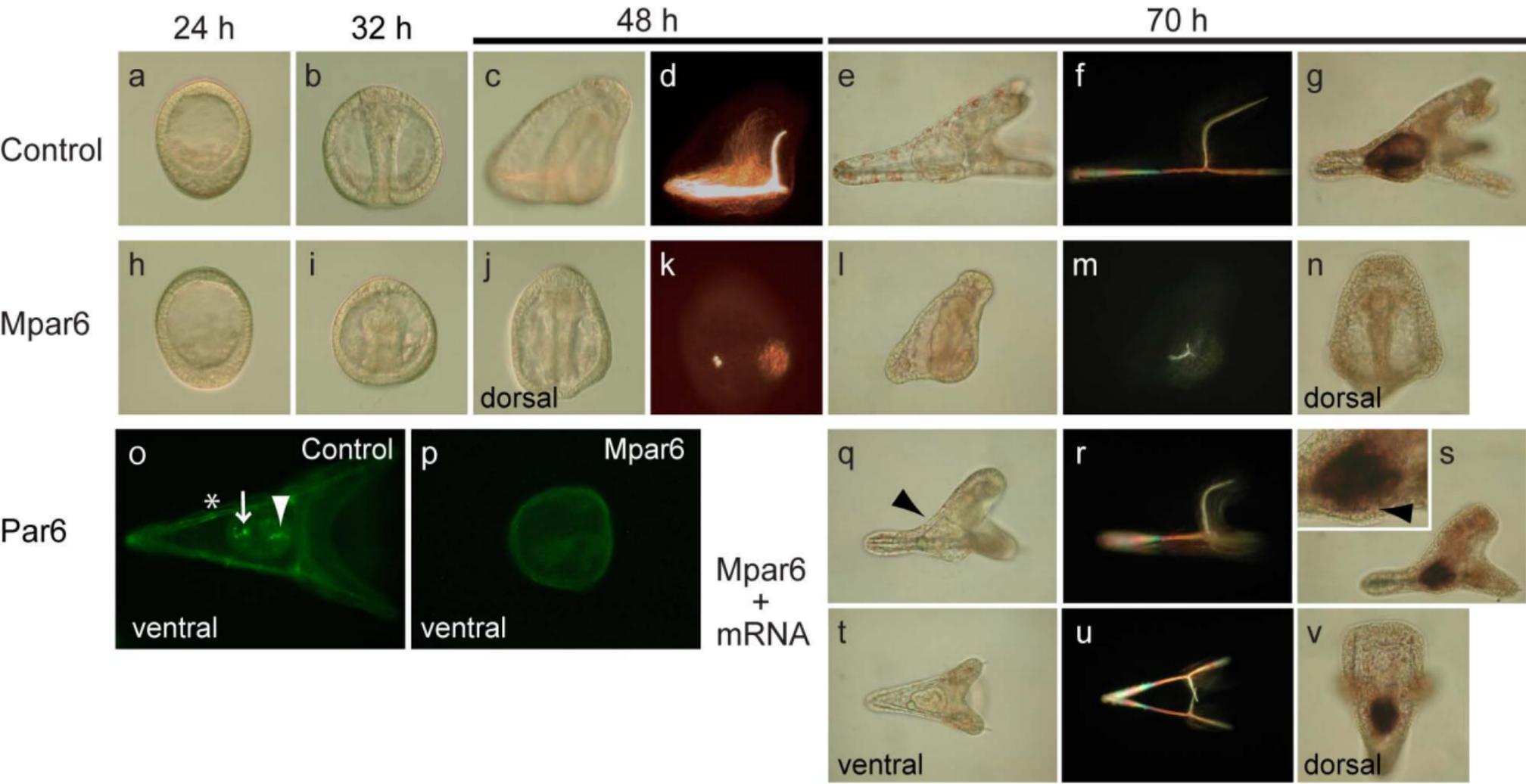
Skeletal rod formation and gut differentiation following the elimination of Par6 function in chimeric larvae composed of PMC-less hosts and transplanted PMCs

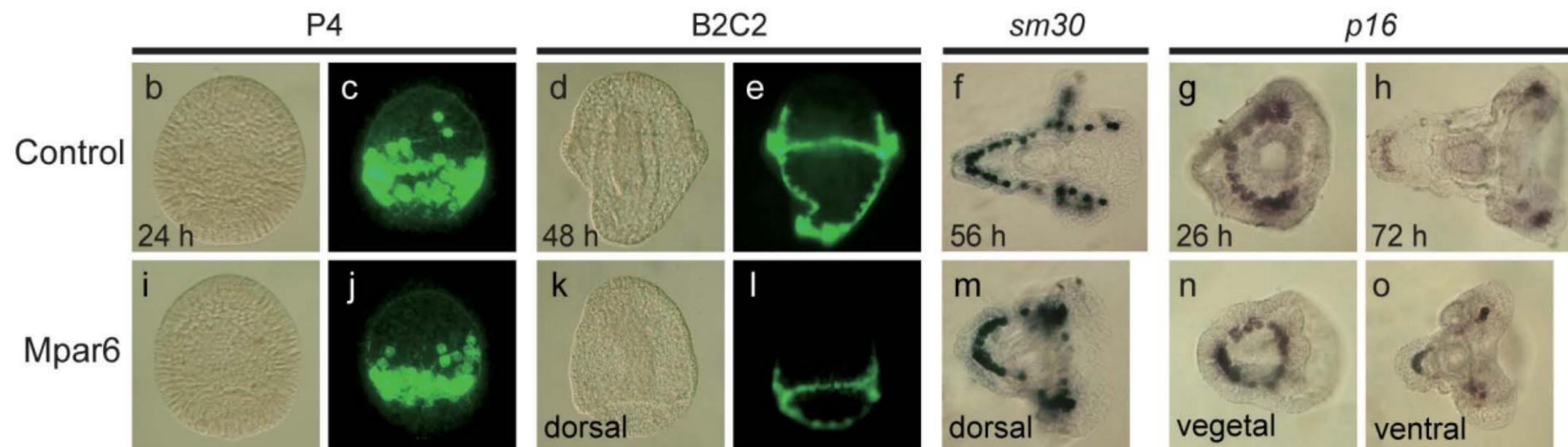
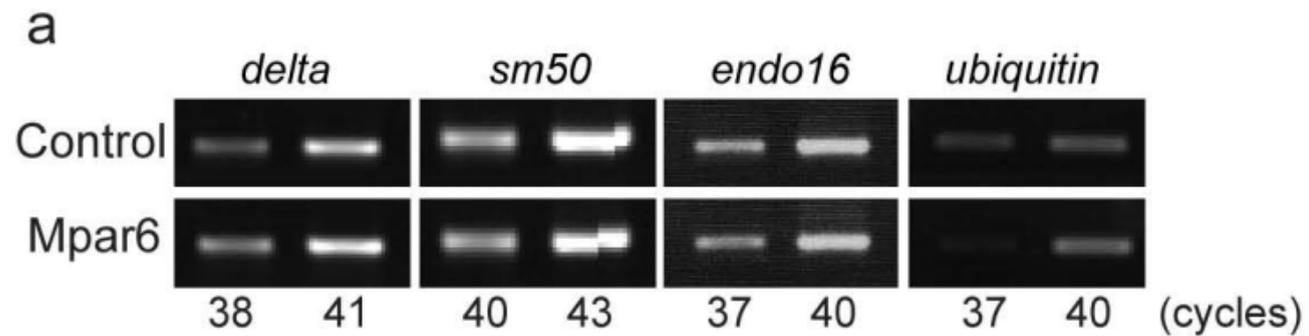
Hosts (PMCs)	Rod elongation				Abnormal rod patterns	Posterior connections	Tripartite guts	AP activity
	AL	PO	T	B				
N (N)	13/13	13/13	13/13	13/13	3/13 ^{*1}	11/13	13/13	6/6
N (Mpar6)	10/11	11/11	9/11	11/11	9/11	0/11	11/11	3/3
Mpar6 (N)	5/6	2/6	4/6	6/6	0/6	5/6	4/6	0/4 ^{*2}

^{*1}Two of the three larvae had a normal branching pattern, but formed extra rods.

^{*2}Larvae with tripartite gut were examined.

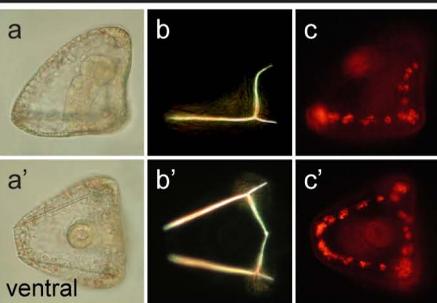




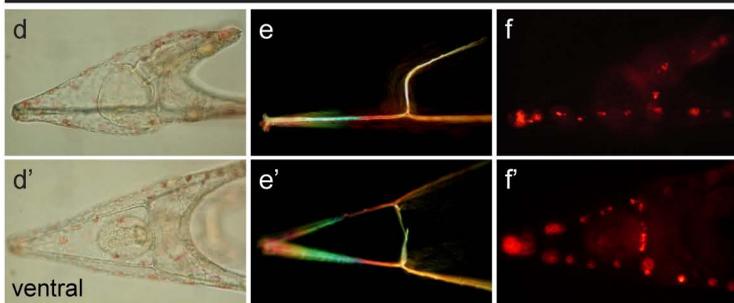


48 h

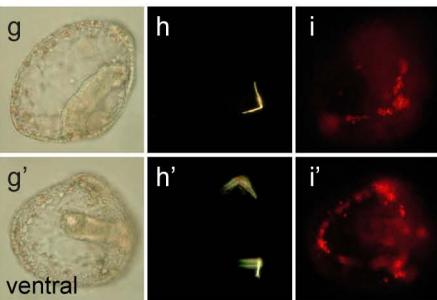
72 h

Normal
(Normal)

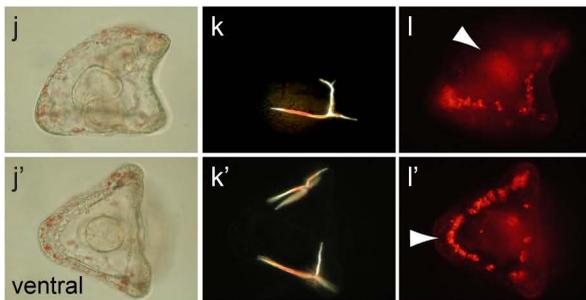
ventral



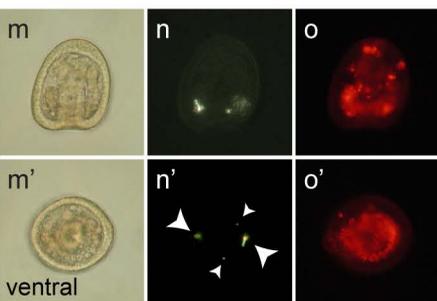
ventral

Normal
(Mpar6)

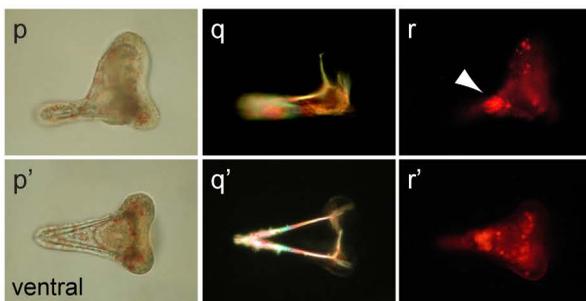
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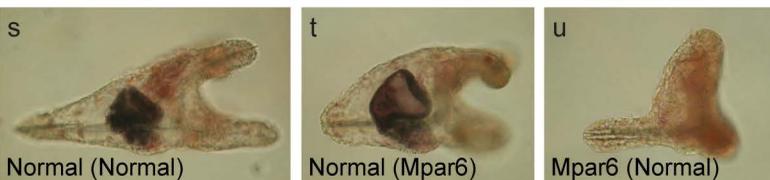
ventral

Mpar6
(Normal)

ventral



ventral

AP
activity

Normal (Normal)

Normal (Mpar6)

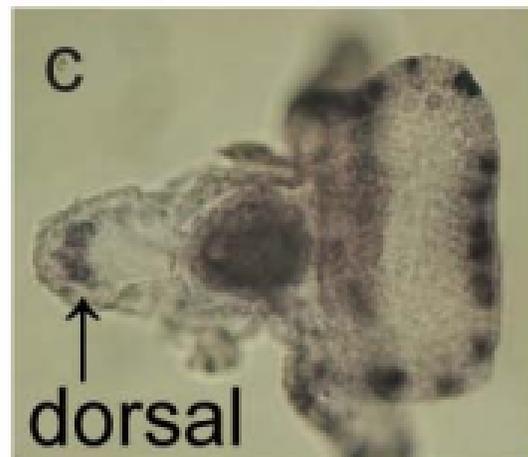
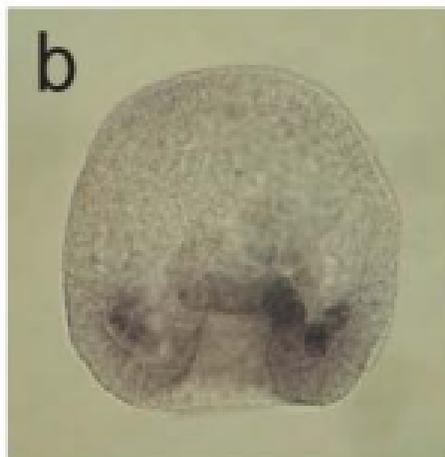
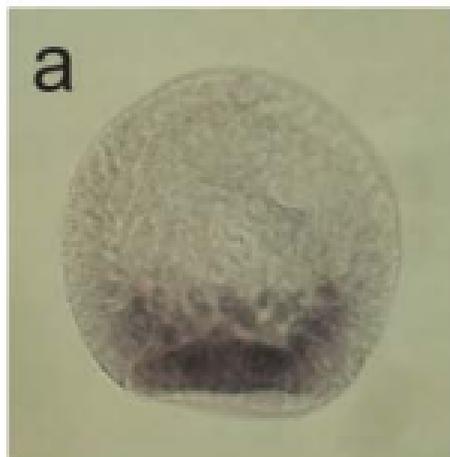
Mpar6 (Normal)

22 h

26 h

52 h

Control



Mpar6

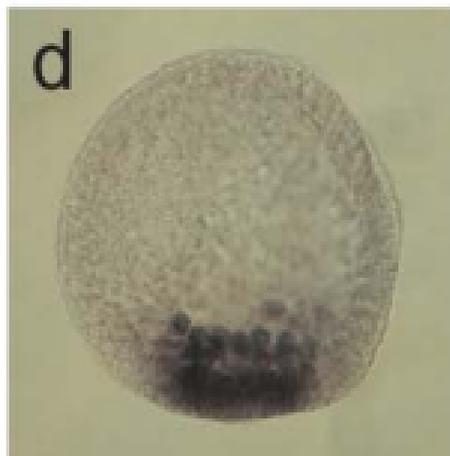
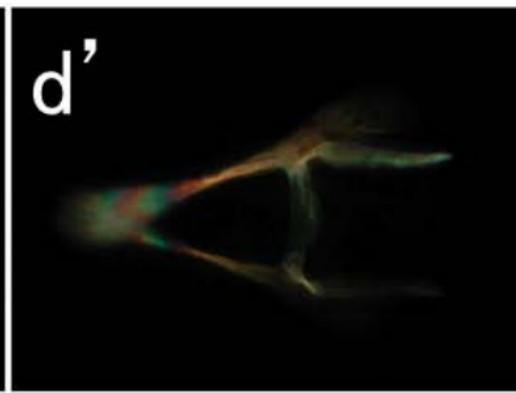
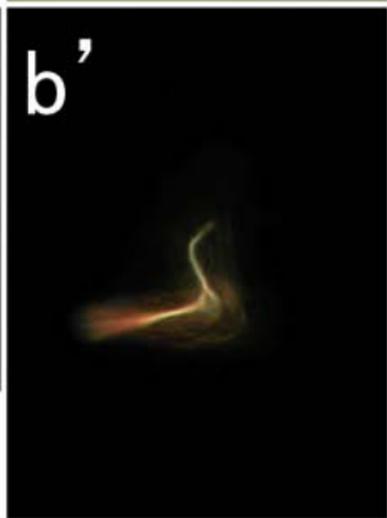
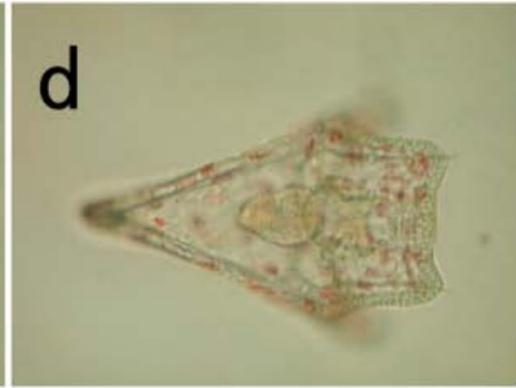
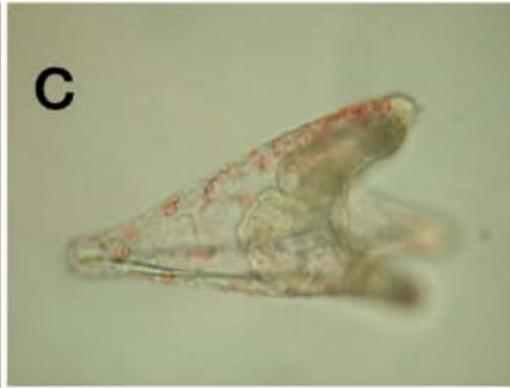
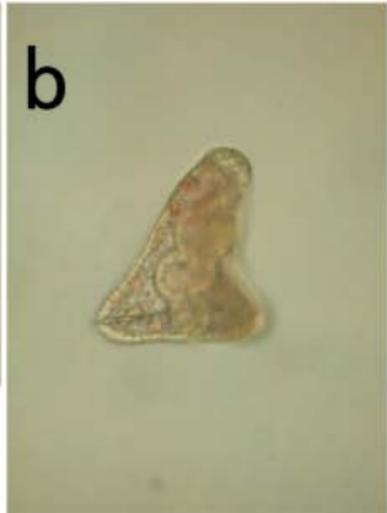


Fig. S1 Co-injection of a lower dose of mRNA injection (~10 pg/egg) rescues *Mpar6* larvae (a, a') from defects in skeletogenesis and gut differentiation, but not from dwarfish ectoderm patterning (b, b'). In contrast, injection of *Par6* mRNA (~15pg/egg) does not affect the skeletal patterning (c, c', d, d')

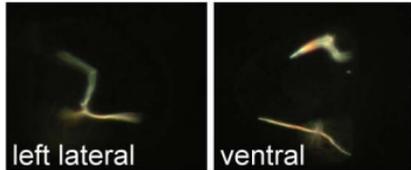
Fig. S2 Skeletal rod formation in eleven chimeric larvae composed of PMC-less hosts and PMCs including *Mpar6*. The elongation and pattern of skeletal rods are summarized in Table 1

Fig. S3 Skeletal rod formation in six chimeric larvae composed of PMC-less hosts including *Mpar6* and normal PMCs. The elongation and pattern of skeletal rods are summarized in Table 1

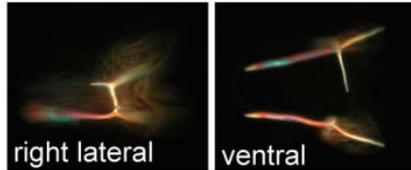
Fig. S4 Distribution of aPKC protein in larvae (72 h) analyzed by immunostaining according to the method of Prulière et al. (2011). The kinase is localized in the ciliary band ectoderm (a'–d'), coelomic sacs (b'), stomach (c'), and intestine (d'). Few signals are detected in the skeletogenic cells; however, those at the vertex show relatively stronger signals (e)



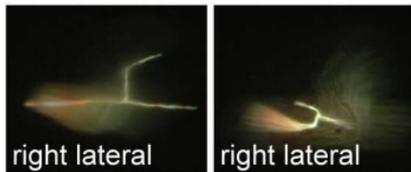
A1-1



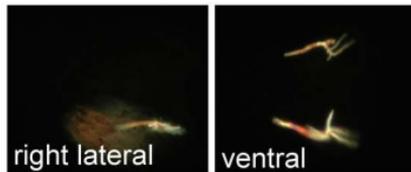
B1



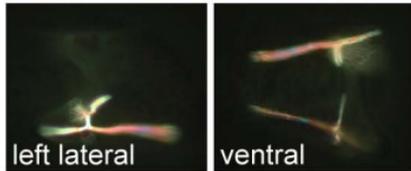
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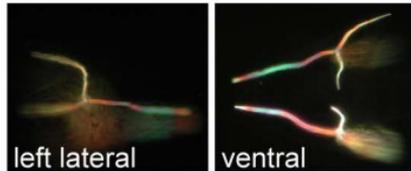
B2



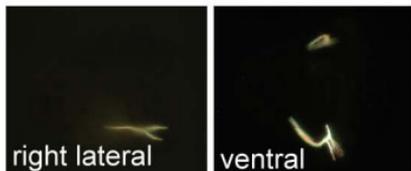
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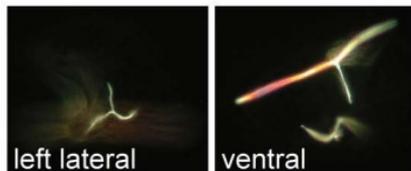
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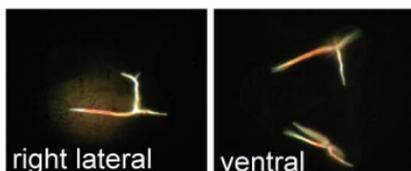
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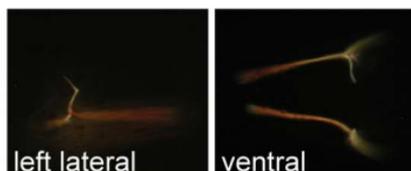
C4-1



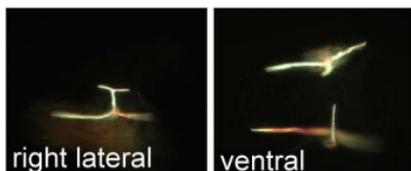
A3-2



C4-2

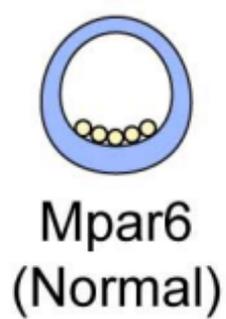


A4

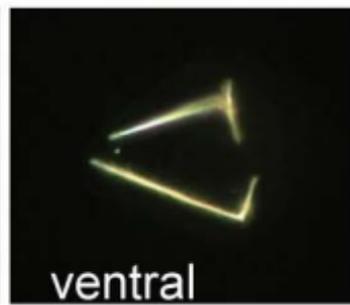


Normal
(Mpar6)

A1



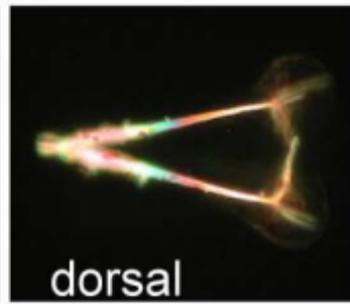
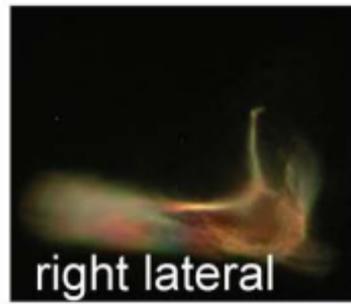
B2



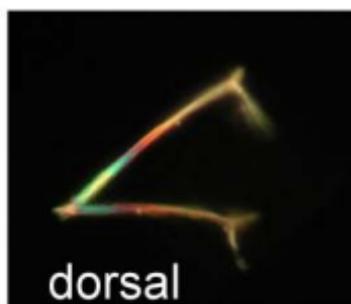
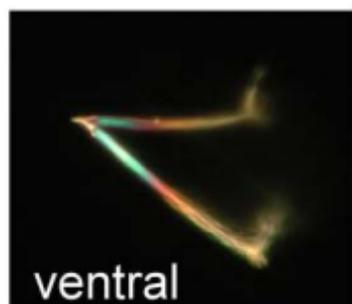
A2



B3



B1



B4

