

RESEARCH

Open Access



La-related protein 6 controls ciliated cell differentiation

Zarko Manojlovic^{1,2}, Ryan Earwood¹, Akiko Kato¹, Diana Perez¹, Oscar A. Cabrera¹, Ruth Didier¹, Timothy L. Megraw¹, Branko Stefanovic^{1*} and Yoichi Kato^{1*}

Abstract

Background: La-related protein 6 (LARP6) is an evolutionally conserved RNA-binding protein. Vertebrate LARP6 binds the 5' stem-loop found in mRNAs encoding type I collagen to regulate their translation, but other target mRNAs and additional functions for LARP6 are unknown. The aim of this study was to elucidate an additional function of LARP6 and to evaluate the importance of its function during development.

Methods: To uncover the role of LARP6 in development, we utilized Morpholino Oligos to deplete LARP6 protein in *Xenopus* embryos. Then, embryonic phenotypes and ciliary structures of LARP6 morphants were examined. To identify the molecular mechanism underlying ciliogenesis regulated by LARP6, we tested the expression level of cilia-related genes, which play important roles in ciliogenesis, by RT-PCR or whole mount in situ hybridization (WISH).

Results: We knocked down LARP6 in *Xenopus* embryos and found neural tube closure defects. LARP6 mutant, which compromises the collagen synthesis, could rescue these defects. Neural tube closure defects are coincident with lack of cilia, antenna-like cellular organelles with motility- or sensory-related functions, in the neural tube. The absence of cilia at the epidermis was also observed in LARP6 morphants, and this defect was due to the absence of basal bodies which are formed from centrioles and required for ciliary assembly. In the process of multi-ciliated cell (MCC) differentiation, *mcidas*, which activates the transcription of genes required for centriole formation during ciliogenesis, could partially restore MCCs in LARP6 morphants. In addition, LARP6 likely controls the expression of *mcidas* in a Notch-independent manner.

Conclusions: La-related protein 6 is involved in ciliated cell differentiation during development by controlling the expression of cilia-related genes including *mcidas*. This LARP6 function involves a mechanism that is distinct from its established role in binding to collagen mRNAs and regulating their translation.

Keywords: LARP6, Cilia, *mcidas*, *Xenopus*

Background

The cilium is a motile or non-motile cellular protrusion that is essential for cell physiology, development and organ homeostasis [1–3]. The ciliary motile function generates fluid flow in a directional manner in multicellular organisms, and also serves to propel cells such as protists and spermatozoa. Non-motile primary cilia sense physical and biochemical extracellular signals in a variety of

contexts. Ciliary defects are associated with a wide spectrum of severe human diseases, called ciliopathies [4].

Recent progress in RNA biology has shown that RNA-binding proteins (RBPs) are important post-transcriptional regulators [5]. However, the only known RBP linked to ciliogenesis thus far is Bicaudal C (BicC) [6, 7]. BicC mediates the alignment of cilia at the left–right organizer, possibly by regulating the activity of the canonical Wnt pathway through Dvl2 and/or RNA silencing in P-bodies [6]. Although the post-transcriptional regulation by RBPs is likely an important level of regulatory control for ciliogenesis, the RBPs involved and their regulatory mechanisms remain largely undiscovered.

*Correspondence: branko.stefanovic@med.fsu.edu;
yoichi.kato@med.fsu.edu

¹ Department of Biomedical Sciences, Florida State University College of Medicine, 1115W. Call Street, Tallahassee, FL 32306-4300, USA
Full list of author information is available at the end of the article

LARP6 is a member of the LARP family of RBPs which carries a La motif (LAM) and a RNA recognition motif (RRM) [8–10]. LARP family member proteins have been implicated in regulating transcription and/or mRNA translation as well as maturation of tRNAs. The RNA-binding properties of LARP proteins are determined by subtle sequence variations within their LAM and RRM domains. LARP6 was originally reported to be involved in the programmed cell death of intersegmental muscle in the moth *M. sexta* [11]. Recently, vertebrate collagen mRNAs including alpha1 (I), alpha2 (I) and alpha1 (III) have been identified as binding targets for LARP6 [12]. LARP6 regulates the generation of heterotrimeric type I collagen molecules by interacting with the 5' stem-loop structure in the 5' UTR of these collagen mRNAs and affecting their translation. The binding to the 5' stem-loop requires the LAM–RRM domain (La-module) of LARP6 and is strictly sequence specific [12]. LARP6 also associates with non-muscle myosin filaments, vimentin intermediate filaments, RNA helicase A, serine–threonine kinase receptor-associated protein (STRAP) and FK506 binding protein 3 (FKBP3) [13–17]. These interactions regulate coordinated translation of collagen mRNAs to achieve 2:1 ratio of the type I collagen subunits.

A role of LARP6 in myogenesis has also been described. LARP6 stimulated the differentiation and self-renewal of mouse C₂C₁₂ cells during in vitro myogenesis and the formation of myotubes in zebrafish embryos [18]. These functions of LARP6 are probably unrelated to binding collagen mRNAs but it is unclear if a role of LARP6 in myogenesis is through the translational regulation of some other target mRNAs. Since target mRNAs other than type I collagen mRNAs have not been identified to date, the principle role of LARP6 is currently thought to be the regulation of collagen synthesis in terminally differentiated fibroblasts and myofibroblasts.

Here, we report that LARP6 is required for the formation of cilia in *Xenopus* embryos. Knockdown of LARP6 in *Xenopus* embryos resulted in neural tube closure defects coincident with loss of cilia, both primary cilia and floor plate cilia, in the neural tube. Furthermore, MCCs at the epidermis were also absent in LARP6 morphants. Our results demonstrate that the loss of the transcription regulator *mcidas*, a critical factor in centriole assembly and ciliogenesis in MCCs, is in part responsible for the lack of ciliary basal bodies formation in the epidermis of LARP6 morphants. However, additional factors whose expression is controlled by LARP6 appear to also be involved in this process. These data show that LARP6 is a regulatory factor which is required for the differentiation of ciliated cells.

Methods

Embryo manipulations

Eggs were artificially fertilized using testis homogenates and cultivated in 0.1 × Marc's Modified Ringer's solution (MMR) [19]. Embryos were staged according to Nieuwkoop and Faber [20].

DNA constructs

Xenopus laevis LARP6 (accession number DR716536) clone for whole mount in situ hybridization (WISH) probes and microinjection were purchased from Thermo Scientific. To generate WISH probes, LARP6-*EcoRI/NotI* fragments were sub-cloned into pBluescript KS vector. To generate microinjection RNA, *X. laevis* LARP6 was isolated by PCR with following primers and was sub-cloned into pCS2+-2HA vector (pCS2+xlL6-2HA). LARP6: forward 5'-GCGGATCCATTCCTGCCTGATAGAAGCT-3', reverse 5'-GCGTCGACATAAACACTGAACGGGGCTTA-3'. pExpress1 *X. laevis* *mcidas* (accession number BC124892) was used for the synthesis of *mcidas* mRNA. Other plasmids for WISH probes including *sox2* and *mcidas* or for microinjection including human LARP6 (HA-hL6), Δ1-300, Δ1-478, Δ135Δ264 and LARP6-GFP (L6-GFP) were described in previous publication [12, 15, 21, 22].

Microinjection of synthetic mRNA and Morpholino Oligos

Capped synthetic mRNAs were generated by in vitro transcription with SP6 polymerase, using the mMessage mMachine kit (Ambion/Life Technologies). For microinjections, embryos were injected with 5–10 nl of the specified amount of mRNA in 3% Ficoll in 0.1 × MMR and cultured in 0.1 × MMR until the desired stage. The site of injection was determined based on the cell fate map of *Xenopus* embryos [23]. *nucβ-gal* mRNA for WISH and membrane RFP (memRFP) mRNA as well as membrane GFP (memGFP) mRNA for immunohistochemistry were injected as a lineage tracer. Antisense Morpholino Oligos (MO) and standard control MO were purchased from Gene Tools. Sequences of MO used in this work are followed. Control MO (conMO): CCTCTTACCTCAGTTA CAATTTATA, LARP6 MO (L6MO): TCTCCTCGGGCT CCTCCATGTCACT. Mismatches between LARP6 S, whose sequence was used in this study, and LARP6 L (accession number BJ030119) are four nucleotides including missing three nucleotides in the middle of MO.

Semi-quantitative RT-PCR analysis

Total RNA was isolated with TRIzol reagent (Life Technologies) according to the manufacturer's instructions. Semi-quantitative RT-PCR was performed as described previously [15, 24]. Sequences of primers used; LARP6: forward 5'-TGTGCGCAAAAACAAGTCTC-3', reverse 5'-TTGTTACAAGGGCTACTCC-3'; *mcidas*: forward

5'-TAGAGGGCGCACAAATTACCT-3', reverse 5'-GCCA CCAGTGGTTTTAATGG-3; mab21-l3: forward 5'-ATG CCCTGGCTGATAAGTTG-3', reverse 5'-TAGAATTGT AGGCGGGTTGG-3; gmnc: forward 5'-CTCCAAACCT TGGGACTGAA-3', reverse 5'-TCACTTCTGGCAGT GATGC-3. Other primers were described in previous publication [25, 26].

Beta-galactosidase staining and whole mount in situ hybridization

Embryos were fixed with MEMFA (0.1 M MOPS, 2 mM EGTA [pH8.0], 1 mM MgSO₄ and 3.7% formaldehyde) containing 0.02% Triton-X for 30 min at room temperature. Galactosidase activity was visualized with the RedGal substrate (Research Organics) in staining buffer (5 mM K₃[Fe(CN)₆], 5 mM K₄[Fe(CN)₆], 2 mM MgCl₂ in PBS). After staining, embryos were re-fixed with MEMFA for 30 min. WISH was performed as described previously [27–29] using BM purple (Roche Applied Science) for the chromogenic reaction. RNA probes for WISH were labeled with Digoxigenin (Roche Applied Science).

Whole mount fluorescence immunohistochemistry

Published procedures were used for staining [30, 31] and thick sectioning for the neural tube [32] with minor modifications. Embryos were fixed in MEMFA for either 2.5 h at room temperature or overnight at 4 °C for anti-acetylated- α -tubulin, anti-B9D1 and anti- γ -tubulin antibodies or in methanol for 48 h at 4 °C for anti-centrin antibody. Fixed embryos were dehydrated completely in methanol at -20 °C for at least several hours and rehydrated consecutively with PBS. After rinsing in PBT (0.1% Triton X-100 in PBS), embryos were incubated with 10% goat serum in PBT at room temperature for at least 1 h. Samples were incubated with mouse anti-acetylated- α -tubulin antibody (1:500, Sigma), rabbit anti-B9D1 antibody (1:50, abnova), rabbit anti- γ -tubulin antibody (1:50, abcam) and mouse anti-centrin antibody (1:50, Millipore) overnight at 4 °C. Primary antibodies were recognized with Cy2 donkey anti-mouse IgG antibody (1:500, Jackson ImmunoResearch) and Alexa Flour 648 goat anti-rabbit IgG antibody (1:500, Life Technologies), respectively. Antibodies were diluted in 10% goat serum in PBT. Images were taken by confocal microscopy.

Immunoblotting

Xenopus embryos were homogenized in lysis buffer [20 mM Tris-HCl: pH 8.0, 5 mM MgCl₂, 1 mM EDTA, 50 mM KCl, 10% glycerol, 0.5% Triton-X, Protease Inhibitor Cocktail (Roche Applied Science)], and embryonic protein extracts were used for immunoblotting with rabbit anti- γ -tubulin (1:1000, abcam), mouse anti-centrin

(1:1000, Millipore) and rabbit anti- β -actin (1:2000, Thermo Scientific) antibodies.

Statistical methods

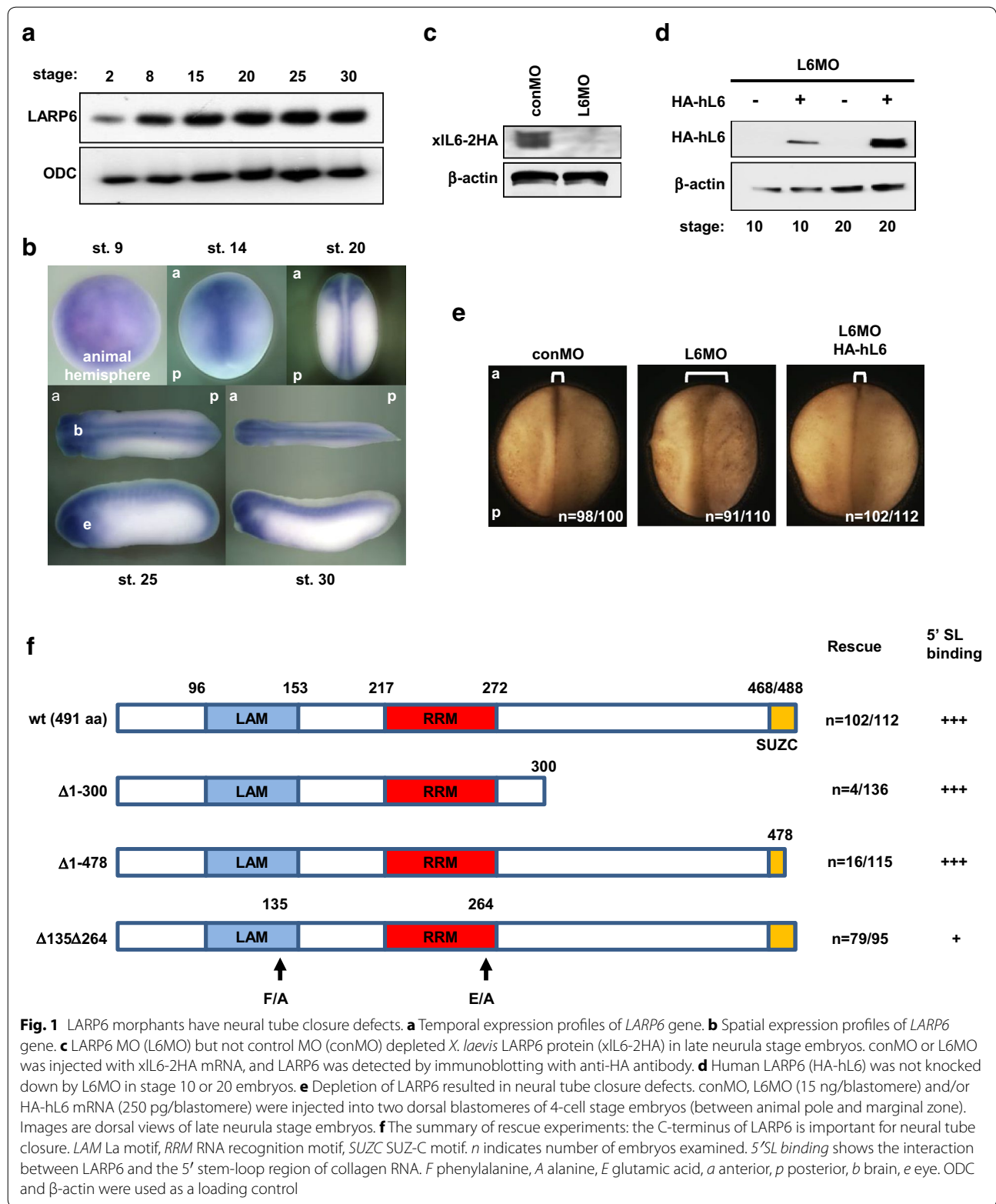
Statistical analyses were carried out using Student's *t* test to calculate *P* values. A *P* value <0.05 was considered to be significant.

Results

LARP6 morphants have neural tube closure defects

Currently, the best defined function of LARP6 in vertebrate is the translational regulation of type I collagen synthesis by binding to the 5' stem-loop sequence of collagen mRNAs [12]. In *Xenopus* embryos, *LARP6* is maternally expressed and continues to be expressed throughout early *Xenopus* embryonic development (Fig. 1a). *LARP6* is strongly detected in the neural plate at the neurula stages as well as the central nervous system such as brain, spinal cord and eyes at the tailbud stages (Fig. 1b). Since type I collagen expression begins at the late neurula stage in *Xenopus* embryos [33], the expression pattern suggests that LARP6 may have additional functions during *Xenopus* development.

To assess the function of LARP6 in development, LARP6 protein was knocked down by injecting Morpholino Oligos against LARP6 (L6MO) (Fig. 1c). The effect of L6MO is highly specific to *Xenopus* LARP6, because it could not block the synthesis of the homologous human LARP6 protein (HA-hL6) expressed in *Xenopus* embryos (Fig. 1d). When L6MO was injected into the animal hemisphere of dorsal blastomeres at the 4-cell stage, the neural tube of LARP6 morphants failed to close (Fig. 1e). In addition, we could not test the role of LARP6 in myogenesis which was reported in moth, zebrafish and mice [11, 18], because the microinjection of LARP6 into the future dorsal mesoderm arrested their development by the tailbud stage. This neural tube closure defect of LARP6 morphants was rescued by co-injection of human LARP6, indicating that LARP6 is required for neural tube closure. Since the human LARP6 protein can restore this function, *Xenopus* LARP6 is its functional homolog. In addition to the La-module, LARP6 carries an SUZC motif (SUZC) [8, 9]. The SUZC in other RNA-binding protein is believed to contribute to mRNA substrate recognition [34]. To understand what domains are necessary for the function of LARP6 in neural tube closure, a rescue of the phenotype was performed with three different LARP6 mutants, Δ 1-300, Δ 1-478 and Δ 135 Δ 264 (Fig. 1f). The interactions between these mutants and the 5' stem-loop of type I collagen mRNA were previously assayed and summarized [12, 15, 21]. Following co-injection of L6MO and these mutant mRNAs, the neural tube formation of embryos was examined. Interestingly, Δ 135 Δ 264



and wt but not $\Delta 1-300$ and $\Delta 1-478$ rescued neural tube closure defects in LARP6 morphants, indicating that the SUZC is required for the function of LARP6 in neural tube closure (Fig. 1f). Because this domain does not participate in binding collagen mRNAs and neural tube closure temporally precedes type I collagen expression, the molecular mechanism underlying this function is likely different from that for LARP6 in collagen synthesis.

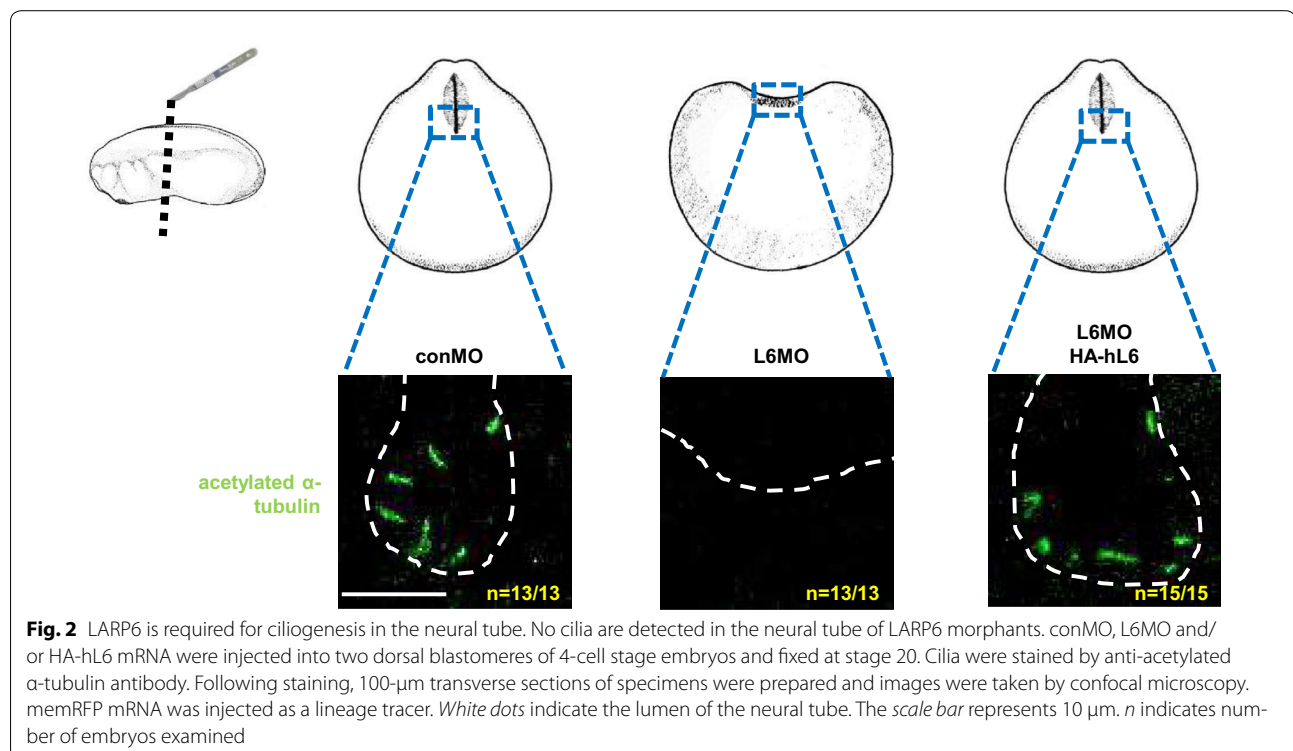
LARP6 morphants lack cilia at the neural tube

Neural differentiation, which forms the neural plate, is a fundamental embryonic process that leads to the development of the neural tube [35]. Therefore, we first examined neural differentiation in the neural plate. The expression of *sox2*, a pan-neural marker [36], was not changed in LARP6 morphants (Additional file 1: Figure S1a), indicating that LARP6 is not involved in neural differentiation at the neural plate. After the neural plate is formed, it lengthens along the anterior–posterior axis and narrows itself so the subsequent bending will form a tube. In *Xenopus* embryos, the convergent extension controls lengthening and narrowing of the neural plate [35]. Therefore, we tested if LARP6 controls the convergent extension by animal cap assay (Additional file 1: Figure S1b). The elongation of animal caps induced by activin was not blocked by knockdown of LARP6, suggesting that LARP6 does not control the convergent extension.

Recent studies have implicated cilia in the neural plate to be critical for the process of the neural tube closure [25, 37–39]. Therefore, cilia in the neural tube of LARP6 morphants were examined. After depletion of LARP6 by injection of L6MO, cilia were detected by immunostaining for acetylated α -tubulin [25]. Complete absence of cilia was observed in the neural tube of LARP6 morphants (middle column in Fig. 2). To show the specificity of this defect, an HA-hL6 mRNA was co-injected with L6MO to test for rescue. As we expected, cilia in the neural tube were restored upon supplementation with HA-hL6 expression (right column in Fig. 2). In addition to these ciliary morphologies, we also tested the gene expression of several cilia-related genes, such as *ARL13B*, *TTC25*, *Foxj1*, *RFX2*, *RFX4* and *RFX7* [25, 39–41]. The expression level of most cilia-related genes, with exception of *RFX2*, was reduced (Additional file 1: Figure S1c). These results indicated that neural tube closure defects in LARP6 morphants are coincident with disruption of neural tube ciliogenesis.

LARP6 is required for the differentiation of MCCs

Next, we tested whether the function of LARP6 was restricted to control ciliogenesis only in the neural tube or not. Here, we assessed the well-characterized cilia on the MCCs of the epidermis. When LARP6 was knocked down by L6MO, no cilia were observed at the epidermis



(Fig. 3a), indicating that LARP6 is also involved in MCC ciliogenesis. To examine the specificity of L6MO effect in MCCs, rescue experiments with hL6 and $\Delta 1-478$ were performed (Fig. 3b). Co-injection of hL6 but not $\Delta 1-478$ mRNA restored MCCs in LARP6 morphant epidermis (Fig. 3b bottom panels), indicating that L6MO effect is specific to ciliary morphologies and the SUZC is also important for MCC differentiation. To further characterize the MCCs in LARP6 morphants, we examined the transition zone marker B9D1/MSKR1 [42–44] as well as the centriole/basal body markers γ -tubulin [45, 46] and centrin [47, 48] in LARP6 morphant MCCs. The transition zone and basal bodies are at the base of cilia, and serve as the ciliary gate or the ciliary base, respectively. Although γ -tubulin and centrin positive centrioles were detected at the LARP6 morphant epidermis (arrowheads in Fig. 3d, e), multiple signals of all three markers were not detected (Fig. 3c–e). In addition, the protein level of γ -tubulin and centrin was not changed in LARP6 morphants when analyzed by immunoblotting (Fig. 3f). These results suggest that LARP6 is required for the formation of basal bodies in the MCCs of the epidermis.

While LARP6 is expressed at the nucleus and the cytoplasm in mammalian cells [12], the localization of LARP6 in ciliated cells has not been reported previously. To examine the localization of LARP6 in MCCs, we expressed GFP-fused human LARP6 (hL6-GFP) with mRFP, which can label cell membranes and cilia [49], in epidermis of *Xenopus* embryos. LARP6 was dominantly detected in the cytoplasm but not cilia of MCCs (Fig. 3g), suggesting that LARP6 may control translation of ciliary or cilia-related factors in MCC differentiation.

***mcidas* is a downstream factor of LARP6**

To define the role of LARP6 within the established molecular network of MCC ciliogenesis, we first examined the connection to the Notch signaling pathway. Notch signaling has been previously shown to restrict the formation of MCCs at the epidermis [50] and suppress the expression of the transcription regulator *mcidas* which initiates MCC differentiation [51, 52]. When Notch signaling is inhibited by a dominant negative form of the co-transcriptional activator mastermind 1 (dn-mam1) [53], the density of MCCs slightly increased (~2 times) [50, 54]. To test if LARP6 controls MCC differentiation in a Notch-dependent manner, we examined the effect of LARP6 depletion on ciliogenesis at the epidermis when Notch signaling was blocked by dn-mam1. Expression of dn-mam1 alone slightly increased the number of MCCs, but co-injection of dn-mam1 and L6MO completely blocked

the formation of MCCs (Fig. 4a, b). If LARP6 blocks Notch signaling to promote MCC differentiation, we expected that increased number of MCCs by injection of dn-mam1 would not be changed. However, co-injection of L6MO resulted in absence of cilia, indicating that LARP6 controls MCC differentiation downstream of Notch signaling, or in a Notch-independent manner. Furthermore, the expression of an active form of Notch receptor, the Notch receptor intracellular domain (NICD), did not suppress the expression of *LARP6*, in contrast to what was observed for *mcidas* (*multicilin*) [51] (Additional file 1: Figure S1d), suggesting that the expression of *LARP6* is not regulated by Notch signaling. Taken together, these results implicate that LARP6 is likely to control MCC differentiation in a Notch-independent manner rather than downstream of Notch signaling.

We next examined whether LARP6 controls the expression of *mcidas* in MCCs, as *mcidas* is a key factor that promotes MCC differentiation. The expression of *mcidas* was inhibited in LARP6 morphants (Fig. 4c), further suggesting that LARP6 is required for MCC differentiation at the early stage of this process. In addition, we did not observe an interaction between LARP6 and *mcidas* mRNA (data not shown). To test if *mcidas* mediates the function of LARP6 in MCC differentiation, rescue experiments with *mcidas* in LARP6 morphant MCCs were performed (Fig. 4d). Co-injection of *mcidas* mRNA restored MCCs in LARP6 morphant epidermis, but the number of MCCs was less than in *mcidas*-expressed epidermis (Fig. 4d, bottom panels). This result indicates that *mcidas* is functionally downstream of LARP6 but additional downstream factors of LARP6 are required for MCC differentiation. In two recent studies, *mab21* family gene *mab21-like3* (*mab21-l3*) and geminin coiled-coil domain containing (*gmnc*) have been shown to control the expression of *mcidas* and regulate MCC ciliogenesis [55, 56]. Therefore, we assessed if LARP6 has a connection with these molecules to control the expression of *mcidas*. The expression of *mab21-l3* and *gmnc* genes was not changed in LARP6 morphants (Fig. 4e) and the interaction between LARP6 and mRNAs of *mab21-l3* or *gmnc* was not detected by immunoprecipitation (data not shown), indicating that the regulation of *mcidas* gene expression by LARP6 is not mediated by *mab21-l3* and *gmnc* expression.

Taken together, our results show that LARP6 controls the early stage of MCC differentiation through a mechanism that appears to be independent of Notch signaling (Fig. 4f) and *mcidas* is an important downstream factor of LARP6 in this process.

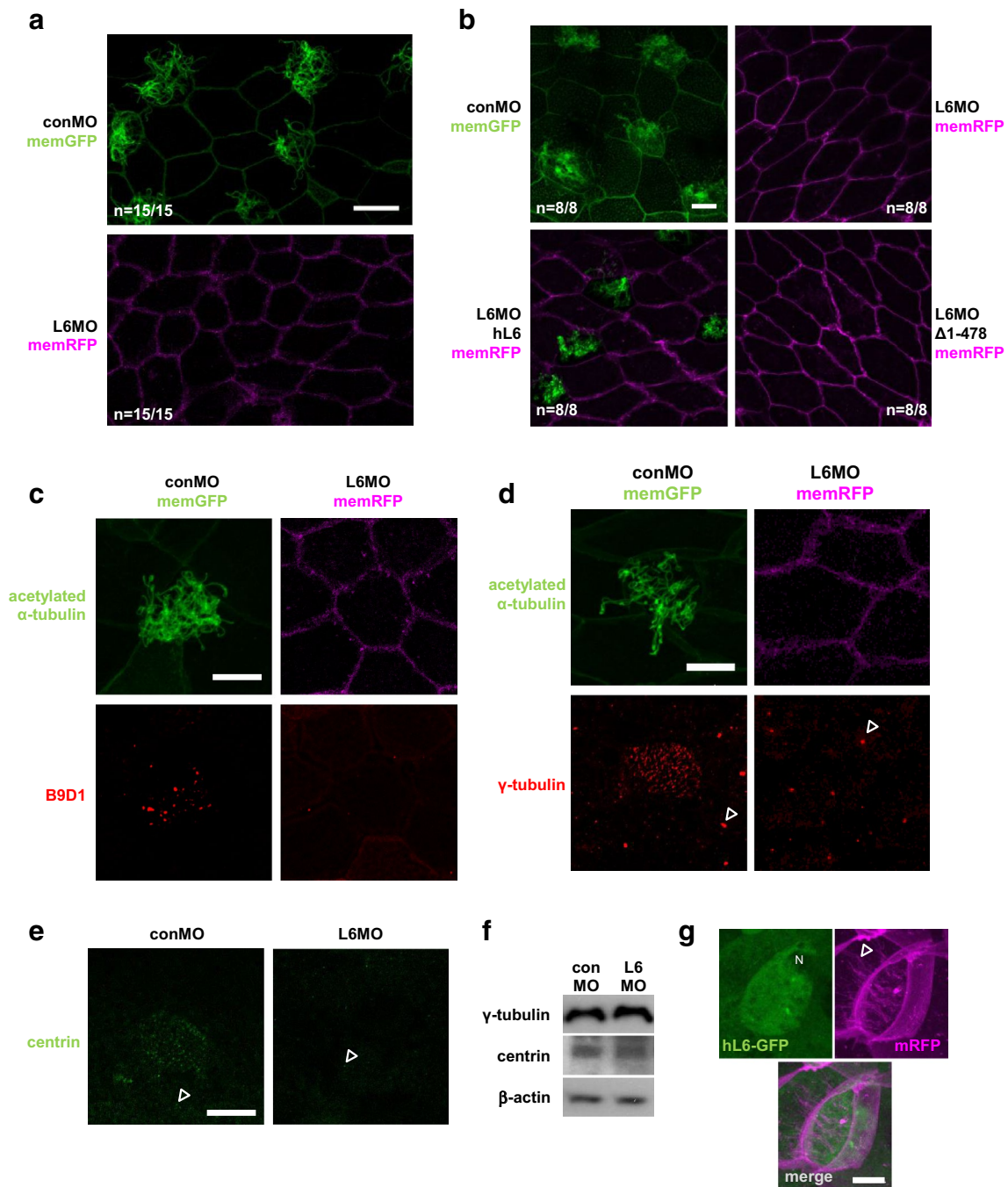


Fig. 3 LARP6 is necessary for MCC differentiation. **a** Absence of cilia on epidermal MCC of LARP6 morphants. L6MO (15 ng/blastomere) with memRFP mRNA or conMO with memGFP mRNA were injected into the *right side* or the *left side*, respectively. Embryos were fixed at stage 25. Cilia were stained with an anti-acetylated tubulin antibody. Following staining, lateral explants were prepared from embryos and images from the *right* (L6MO) or *left* (conMO) lateral side were taken by confocal microscopy. **b** Rescue experiments with hL6 or Δ 1-478 mRNA in LARP6 morphant MCCs. **c–e** B9D1, γ -tubulin and centrin proteins were not detected at the LARP6 morphant epidermis. Axonemes and ciliary transition zones or basal bodies were stained by an anti-acetylated tubulin and anti-B9D1, anti- γ -tubulin or anti-centrin antibody, respectively. Following staining, lateral explants were prepared from embryos and images were taken by confocal microscopy. *Arrowheads* indicate examples of centrioles. **f** The depletion of LARP6 did not change total amounts of γ -tubulin and centrin proteins in embryos. **g** The localization of LARP6 in MCCs. *N* nucleus. An *Arrowhead* indicates an example of a cilium. *Scale bars* 20 μ m in **(a)**, 10 μ m in **(b–e, g)**. *n* indicates number of embryos examined

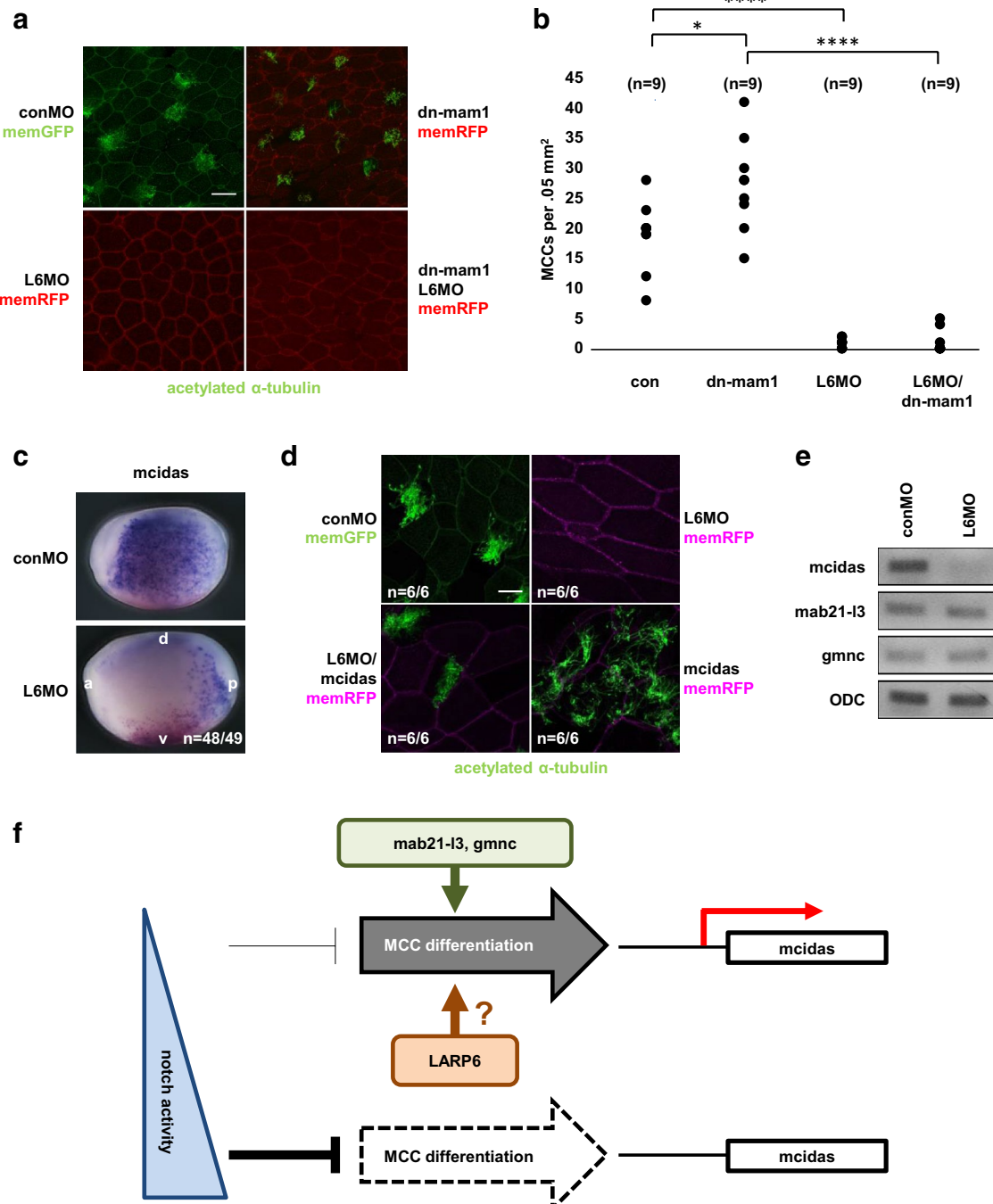


Fig. 4 LARP6 regulates the expression of *mcidas* in ciliogenesis. **a** The depletion of LARP6 blocked ciliogenesis in a Notch-independent manner. dn-mam1 RNA and/or L6MO with memRFP mRNA or conMO with memGFP mRNA were injected into the *right side* or the *left side*, respectively. Embryos were fixed at stage 25. Cilia were stained with an anti-acetylated tubulin antibody. Following staining, lateral explants were prepared from embryos and images were taken by confocal microscopy. The scale bar represents 20 μ m. **b** Quantification of number of MCCs from **a**. Three embryos were randomly chosen from each of three independent experiments. * ≤ 0.05 , **** ≤ 0.0001 . **c** *mcidas* expression in LARP6 morphants at stage 14 by WISH. Images from the left (conMO) or right (L6MO) lateral side were taken from the same embryo. Red-gal signals indicate L6MO injected side. a anterior, p posterior, d dorsal, v ventral. **d** Rescue experiments with *mcidas* mRNA (500 pg/blastomere) in LARP6 morphant MCCs. **e** LARP6 does not control the expression of *mab21-l3* and *gmnc* genes. L6MO was injected into two blastomeres of 2-cell stage embryos and RNAs were isolated at stage 13 for semi-quantitative RT-PCR. **f** The model of LARP6 function in ciliogenesis. LARP6 controls the expression of *mcidas* in a Notch-independent manner. n indicates number of embryos examined

Discussion

The results presented here reveal a role of La-related protein 6 (LARP6) for ciliated cell differentiation in *Xenopus* embryos. Knockdown of LARP6 resulted in neural tube closure defect, which is coincident with the absence of cilia, and blocking MCC differentiation at the epidermis. Although LARP6 appears to control MCC differentiation through a yet-discovered molecular network, *mcidas* is an important downstream factor of LARP6 in this process. Since the molecular mechanisms underlying the structure and function of cilia are well-conserved among the vertebrates [3], this newly discovered role of LARP6 in ciliated cell differentiation is likely conserved in humans and may be involved in pathogenesis of human ciliopathies.

Knockdown of LARP6 in *Xenopus* embryos resulted in neural tube closure defects. The first step of neural tube closure is neural differentiation in dorsal ectoderm of embryos which becomes the neural plate. Neural differentiation, which was represented by the expression of *sox2*, was not changed in LARP6 morphants. After neural differentiation, the neural plate buckles, rolls up and then fuses to form a hollow tube called the neural tube [35]. The convergent extension is involved in shaping of the neural plate of *Xenopus* embryos which is required for the formation of a tube. Our study indicated that LARP6 was not required for the convergent extension. These results suggested that neural tube closure defects in LARP6 morphants are caused by neither the inhibition of neural differentiation in dorsal ectoderm nor shaping of the neural plate. Therefore, neural tube closure defects of LARP6 morphants may be caused by the loss of cilia.

LARP6 has the LAM and the RRM, collectively termed the La-module, which is required for binding to the 5' stem-loop of type I collagen mRNAs [12], as well as the SUZC in the extreme C-terminus. Although the SUZC is not necessary for binding the 5' stem-loop of collagen mRNAs, this motif was required for neural tube closure and MCC differentiation. Moreover, LARP6 mutant $\Delta 135\Delta 264$, which weakly interacts with the 5' stem-loop of collagen I mRNAs [15], could rescue neural tube closure defect of LARP6 morphants similar to wild-type LARP6. These data indicate that LARP6 may regulate mRNAs for ciliated cell differentiation through a different mechanism than the mechanism which regulates collagen synthesis. However, at present we have not yet identified the target mRNAs for LARP6 required for ciliated cell differentiation. The identification of LARP6 target mRNAs in development will be critical to uncover the molecular mechanism underlying ciliated cell differentiation controlled by LARP6.

The SUZC of LARP6 is required for the differentiation of ciliated cells at the neural tube and MCCs. Since this motif in other protein is generally believed to contribute

to mRNA substrate recognition [34], factor(s) interacting with the SUZC of LARP6 must be important for the differentiation of ciliated cells and may determine the target mRNA(s) in this process. Serine–threonine kinase receptor-associated protein (STRAP) has been previously shown to interact with the SUZC of LARP6 [16]. STRAP was initially identified as a novel WD40 domain-containing protein which interacts with both T β R-I and T β R-II serine–threonine kinase receptors and negatively regulates gene expression from TGF- β -responsive promoters [57]. Although homozygous STRAP mutant embryos are embryonically lethal with many defects between embryonic day 10.5 and 12.5 [58], these embryos present neural tube closure defects. Therefore, STRAP seems to be a possible partner of LARP6 in the differentiation of ciliated cells. However, further studies are required to reveal the importance of this SUZC and identify interacting factor(s) during the differentiation of ciliated cells.

mcidas has been reported to control centriole assembly and promote MCC differentiation [51, 52]. MCCs with multiple basal bodies were lost in the epidermis of LARP6 morphants, and *mcidas* appears to mediate this function of LARP6 partially. Although *mcidas* could restore MCCs in LARP6 morphants, the number of MCCs induced by *mcidas* was reduced in LARP6 morphants, indicating that additional factor(s) may be needed for complete restoration of MCCs in LARP6 morphants. Therefore, LARP6 appears to control the expression of *mcidas* and additional factor(s) which cooperate with *mcidas* in MCC differentiation. In addition, *mcidas* is not expressed in the neural tube [51], but cilia were absent in LARP6 morphant neural tube. This also indicates that LARP6 controls the expression of additional factor(s) which are necessary for the differentiation of ciliated cells. For instance, our study showed that the expression of *RFX4* and 7, which are required for neural tube ciliogenesis [25], is controlled by LARP6. Furthermore, while the expression of *mcidas* and MCC differentiation are negatively regulated by Notch signaling [51], the increase of MCC formation induced by inhibition of Notch signaling was completely blocked by knockdown of LARP6. These results indicate that LARP6 likely controls the expression of *mcidas* and MCC differentiation in a Notch-independent manner. *mab21-13* and *gmnc* proteins are known to regulate the expression of *mcidas* and MCC differentiation [55, 56], however, we did not observe any changes in the expression level of these genes in LARP6 morphants and the interaction between LARP6 and their mRNAs. Therefore, LARP6 is likely involved in *mcidas* expression and MCC differentiation in a yet-uncovered mechanism (Fig. 4f). The identification of LARP6 target mRNAs will help us to understand the molecular mechanism of ciliated cell differentiation.

Conclusions

In summary, the work presented here strongly suggests that LARP6 is involved in the differentiation of ciliated cells and that this function is independent of the regulation of type I collagen synthesis. Moreover, recent single-nucleotide polymorphisms' analysis from four genome-wide association studies indicate that a polymorphism at or near LARP6 is linked to the risk of type 2 diabetes [59]. Since Bardet–Biedl syndrome, a ciliopathic human genetic disorder, has been reported to be linked to obesity and diabetes [60], dysfunction of LARP6 may result in ciliopathies associated with type II diabetes. In this aspect, our study opens new avenues to investigate the pathogenesis of ciliopathies.

Additional file

Additional file 1: Figure S1. LARP6 controls the expression of cilia-related genes in a Notch-independent manner.

Abbreviations

LARP6: La-related protein 6; RBP: RNA-binding protein; BisC: Bicaudal C; LAM: La motif; RRM: RNA recognition motif; STRAP: serine–threonine kinase receptor-associated protein; FKBP3: FK506 binding protein 3; MCC: multi-ciliated cell; SUZC: SUZ-C motif; NICD: Notch receptor intracellular domain; mab21-l3: mab21 family gene mab21-like3; gmnc: geminin coiled-coil domain containing.

Authors' contributions

YK and BS conceived the project. ZM, RE and DP carried out the morphological analysis of embryos and cilia. AK and DP gave support in multiple experiments including whole mount in situ hybridization and molecular biological experiments. RD gave support in taking confocal images. ZM and RE prepared figures. YK, CO, TM and BJ wrote the manuscript. All authors read and approved the final manuscript.

Author details

¹ Department of Biomedical Sciences, Florida State University College of Medicine, 1115W. Call Street, Tallahassee, FL 32306-4300, USA. ² Present Address: Department of Translational Genomics, Keck School of Medicine of University of Southern California, Los Angeles, CA 90089-9601, USA.

Acknowledgements

We would like to thank Dr. M. Brum for critical suggestions in manuscript and Dr. B. J. Mitchell for critical suggestions in immunohistochemistry. This work was supported by NIH Grants (GM087641 and GM036556 to YK).

Competing interests

The authors declare that they have no competing interests.

Ethics approval and consent to participate

This work was approved by Animal Care and use committee (ACUC) at the Florida State University on May 28, 2014. The Protocol Number is 1422.

Received: 22 October 2016 Accepted: 16 February 2017

Published online: 23 March 2017

References

- Ishikawa H, Marshall WF. Ciliogenesis: building the cell's antenna. *Nat Rev Mol Cell Biol.* 2011;12(4):222–34.

- García-Gonzalo FR, Reiter JF. Scoring a backstage pass: mechanisms of ciliogenesis and ciliary access. *J Cell Biol.* 2012;197(6):697–709.
- Choksi SP, Lauter G, Swoboda P, Roy S. Switching on cilia: transcriptional networks regulating ciliogenesis. *Development (Cambridge, England).* 2014;141(7):1427–41.
- Hildebrandt F, Benzing T, Katsanis N. Ciliopathies. *N Engl J Med.* 2011;364(16):1533–43.
- Marchese D, de Groot NS, Lorenzo Gotor N, Livi CM, Tartaglia GG. Advances in the characterization of RNA-binding proteins. *Wiley Interdiscip Rev RNA.* 2016;7:793–810.
- Maisonneuve C, Guilleret I, Vick P, Weber T, Andre P, Beyer T, Blum M, Constam DB. Bicaudal C, a novel regulator of Dvl signaling abutting RNA-processing bodies, controls cilia orientation and leftward flow. *Development (Cambridge, England).* 2009;136(17):3019–30.
- Tran U, Pickney LM, Ozpolat BD, Wessely O. Xenopus Bicaudal-C is required for the differentiation of the amphibian pronephros. *Dev Biol.* 2007;307(1):152–64.
- Stavraka C, Blagden S. The La-related proteins, a family with connections to cancer. *Biomolecules.* 2015;5(4):2701–22.
- Bousquet-Antonelli C, Deragon JM. A comprehensive analysis of the Lamotif protein superfamily. *RNA.* 2009;15(5):750–64.
- Bayfield MA, Yang R, Maraia RJ. Conserved and divergent features of the structure and function of La and La-related proteins (LARPs). *Biochim Biophys Acta.* 2010;1799(5–6):365–78.
- Valavanis C, Wang Z, Sun D, Vaine M, Schwartz LM. Acheron, a novel member of the Lupus Antigen family, is induced during the programmed cell death of skeletal muscles in the moth *Manduca sexta*. *Gene.* 2007;393(1–2):101–9.
- Cai L, Fritz D, Stefanovic L, Stefanovic B. Binding of LARP6 to the conserved 5' stem-loop regulates translation of mRNAs encoding type I collagen. *J Mol Biol.* 2010;395(2):309–26.
- Cai L, Fritz D, Stefanovic L, Stefanovic B. Nonmuscle myosin-dependent synthesis of type I collagen. *J Mol Biol.* 2010;401(4):564–78.
- Challa AA, Stefanovic B. A novel role of vimentin filaments: binding and stabilization of collagen mRNAs. *Mol Cell Biol.* 2011;31(18):3773–89.
- Manojlovic Z, Stefanovic B. A novel role of RNA helicase A in regulation of translation of type I collagen mRNAs. *RNA.* 2012;18(2):321–34.
- Vukmirovic M, Manojlovic Z, Stefanovic B. Serine–threonine kinase receptor-associated protein (STRAP) regulates translation of type I collagen mRNAs. *Mol Cell Biol.* 2013;33(19):3893–906.
- Manojlovic Z, Blackmon J, Stefanovic B. Tacrolimus (FK506) prevents early stages of ethanol induced hepatic fibrosis by targeting LARP6 dependent mechanism of collagen synthesis. *PLoS ONE.* 2013;8(6):e65897.
- Wang Z, Glenn H, Brown C, Valavanis C, Liu JX, Seth A, Thomas JE, Karlstrom RO, Schwartz LM. Regulation of muscle differentiation and survival by Acheron. *Mech Dev.* 2009;126(8–9):700–9.
- Peng HB. *Xenopus laevis*: practical uses in cell and molecular biology. Solutions and protocols. *Methods Cell Biol.* 1991;36:657–62.
- Nieuwkoop PD, Faber J. Normal table of *Xenopus laevis* (Daudin). Amsterdam: North-Holland Publishing Co; 1967.
- Stefanovic L, Longo L, Zhang Y, Stefanovic B. Characterization of binding of LARP6 to the 5' stem-loop of collagen mRNAs: implications for synthesis of type I collagen. *RNA Biol.* 2014;11(11):1386–401.
- Tozser J, Earwood R, Kato A, Brown J, Tanaka K, Didier R, Megraw TL, Blum M, Kato Y. TGF-beta signaling regulates the differentiation of motile cilia. *Cell Rep.* 2015;11(7):1000–7.
- Moody SA. Fates of the blastomeres of the 16-cell stage *Xenopus* embryo. *Dev Biol.* 1987;119(2):560–78.
- Kato Y, Shi Y, He X. Neuralization of the *Xenopus* embryo by inhibition of p300/CREB-binding protein function. *J Neurosci.* 1999;19(21):9364–73.
- Manojlovic Z, Earwood R, Kato A, Stefanovic B, Kato Y. RFX7 is required for the formation of cilia in the neural tube. *Mech Dev.* 2014;132:28–37.
- Sakano D, Kato A, Parikh N, McKnight K, Terry D, Stefanovic B, Kato Y. BCL6 canalizes Notch-dependent transcription, excluding Mastermind-like1 from selected target genes during left–right patterning. *Dev Cell.* 2010;18(3):450–62.
- Harland RM. In situ hybridization: an improved whole-mount method for *Xenopus* embryos. *Methods Cell Biol.* 1991;36:685–95.
- Takada H, Hattori D, Kitayama A, Ueno N, Taira M. Identification of target genes for the *Xenopus* Hes-related protein XHR1, a

- prepattern factor specifying the midbrain-hindbrain boundary. *Dev Biol*. 2005;283(1):253–67.
29. Kiyota T, Kato A, Altmann CR, Kato Y. The POU homeobox protein Oct-1 regulates radial glia formation downstream of Notch signaling. *Dev Biol*. 2008;315(2):579–92.
 30. Suzuki M, Satoh A, Ide H, Tamura K. Transgenic *Xenopus* with prx1 limb enhancer reveals crucial contribution of MEK/ERK and PI3K/AKT pathways in blastema formation during limb regeneration. *Dev Biol*. 2007;304(2):675–86.
 31. Lee C, Kieserman E, Gray RS, Park TJ, Wallingford J. Whole-mount fluorescence immunocytochemistry on *Xenopus* embryos. *Cold Spring Harb Protocols*. 2008; 2008.pdb prot4957.
 32. Becker BE, Gard DL. Visualization of the cytoskeleton in *Xenopus* oocytes and eggs by confocal immunofluorescence microscopy. *Methods Mol Biol*. 2006;322:69–86.
 33. Goto T, Katada T, Kinoshita T, Kubota HY. Expression and characterization of *Xenopus* type I collagen alpha 1 (COL1A1) during embryonic development. *Dev Growth Diff*. 2000;42(3):249–56.
 34. Song MH, Aravind L, Muller-Reichert T, O'Connell KF. The conserved protein SZY-20 opposes the Plk4-related kinase ZYG-1 to limit centrosome size. *Dev Cell*. 2008;15(6):901–12.
 35. Wallingford JB. Planar cell polarity, ciliogenesis and neural tube defects. *Hum Mol Genet*. 2006;15(2):R227–34.
 36. Mizuseki K, Kishi M, Matsui M, Nakanishi S, Sasai Y. *Xenopus* Zic-related-1 and Sox-2, two factors induced by chordin, have distinct activities in the initiation of neural induction. *Development* (Cambridge, England). 1998;125(4):579–87.
 37. Murdoch JN, Copp AJ. The relationship between sonic Hedgehog signaling, cilia, and neural tube defects. *Birth Defects Res A Clin Mol Teratol*. 2010;88(8):633–52.
 38. Vogel TW, Carter CS, Abode-lyamah K, Zhang Q, Robinson S. The role of primary cilia in the pathophysiology of neural tube defects. *Neurosurg Focus*. 2012;33(4):E2.
 39. Chung MI, Peyrot SM, LeBoeuf S, Park TJ, McGary KL, Marcotte EM, Wallingford JB. RFX2 is broadly required for ciliogenesis during vertebrate development. *Dev Biol*. 2012;363(1):155–65.
 40. Hayes JM, Kim SK, Abitua PB, Park TJ, Herrington ER, Kitayama A, Grow MW, Ueno N, Wallingford JB. Identification of novel ciliogenesis factors using a new in vivo model for mucociliary epithelial development. *Dev Biol*. 2007;312(1):115–30.
 41. Stubbs JL, Oishi I, Izipisua Belmonte JC, Kintner C. The forkhead protein Foxj1 specifies node-like cilia in *Xenopus* and zebrafish embryos. *Nat Genet*. 2008;40(12):1454–60.
 42. Williams CL, Li C, Kida K, Inglis PN, Mohan S, Semencik L, Bialas NJ, Stupay RM, Chen N, Blacque OE, et al. MKS and NPHP modules cooperate to establish basal body/transition zone membrane associations and ciliary gate function during ciliogenesis. *J Cell Biol*. 2011;192(6):1023–41.
 43. Garcia-Gonzalo FR, Corbit KC, Sirerol-Piquer MS, Ramaswami G, Otto EA, Noriega TR, Seol AD, Robinson JF, Bennett CL, Josifova DJ, et al. A transition zone complex regulates mammalian ciliogenesis and ciliary membrane composition. *Nat Genet*. 2011;43(8):776–84.
 44. Huang L, Szymanska K, Jensen VL, Janecke AR, Innes AM, Davis EE, Frosk P, Li C, Willer JR, Chodirker BN, et al. TMEM237 is mutated in individuals with a Joubert syndrome related disorder and expands the role of the TMEM family at the ciliary transition zone. *Am J Hum Genet*. 2011;89(6):713–30.
 45. Chung MI, Nascone-Yoder NM, Grover SA, Drysdale TA, Wallingford JB. Direct activation of Shroom3 transcription by Pitx proteins drives epithelial morphogenesis in the developing gut. *Development* (Cambridge, England). 2010;137(8):1339–49.
 46. Muresan V, Joshi HC, Besharse JC. Gamma-tubulin in differentiated cell types: localization in the vicinity of basal bodies in retinal photoreceptors and ciliated epithelia. *J Cell Sci*. 1993;104(Pt 4):1229–37.
 47. Mitchell B, Jacobs R, Li J, Chien S, Kintner C. A positive feedback mechanism governs the polarity and motion of motile cilia. *Nature*. 2007;447(7140):97–101.
 48. LeDizet M, Beck JC, Finkbeiner WE. Differential regulation of centriole genes during ciliogenesis in human tracheal epithelial cells. *Am J Physiol*. 1998;275(6 Pt 1):L1145–56.
 49. Brooks ER, Wallingford JB. Control of vertebrate intraflagellar transport by the planar cell polarity effector Fuz. *J Cell Biol*. 2012;198(1):37–45.
 50. Deblandre GA, Wettstein DA, Koyano-Nakagawa N, Kintner C. A two-step mechanism generates the spacing pattern of the ciliated cells in the skin of *Xenopus* embryos. *Development* (Cambridge, England). 1999;126(21):4715–28.
 51. Stubbs JL, Vladar EK, Axelrod JD, Kintner C. Multicilin promotes centriole assembly and ciliogenesis during multiciliate cell differentiation. *Nat Cell Biol*. 2012;14(2):140–7.
 52. Ma L, Quigley I, Omran H, Kintner C. Multicilin drives centriole biogenesis via E2f proteins. *Genes Dev*. 2014;28(13):1461–71.
 53. Fryer CJ, Lamar E, Turbachova I, Kintner C, Jones KA. Mastermind mediates chromatin-specific transcription and turnover of the Notch enhancer complex. *Genes Dev*. 2002;16(11):1397–411.
 54. Stubbs JL, Davidson L, Keller R, Kintner C. Radial intercalation of ciliated cells during *Xenopus* skin development. *Development*. 2006;133(13):2507–15.
 55. Takahashi C, Kusakabe M, Suzuki T, Miyatake K, Nishida E. mab21-13 regulates cell fate specification of multiciliate cells and ionocytes. *Nat Commun*. 2015;6:6017.
 56. Zhou F, Narasimhan V, Shboul M, Chong YL, Reversade B, Roy S. Gmnc is a master regulator of the multiciliated cell differentiation program. *Curr Biol*. 2015;25(24):3267–73.
 57. Datta PK, Chytil A, Gorska AE, Moses HL. Identification of STRAP, a novel WD domain protein in transforming growth factor-beta signaling. *J Biol Chem*. 1998;273(52):34671–4.
 58. Chen WV, Delrow J, Corrin PD, Frazier JP, Soriano P. Identification and validation of PDGF transcriptional targets by microarray-coupled gene-trap mutagenesis. *Nat Genet*. 2004;36(3):304–12.
 59. Strawbridge RJ, Dupuis J, Prokopenko I, Barker A, Ahlqvist E, Rybin D, Petrie JR, Travers ME, Bouatia-Naji N, Dimas AS, et al. Genome-wide association identifies nine common variants associated with fasting proinsulin levels and provides new insights into the pathophysiology of type 2 diabetes. *Diabetes*. 2011;60(10):2624–34.
 60. Forsythe E, Beales PL. Bardet-Biedl syndrome. *Eur J Hum Genet*. 2013;21(1):8–13.

Submit your next manuscript to BioMed Central and we will help you at every step:

- We accept pre-submission inquiries
- Our selector tool helps you to find the most relevant journal
- We provide round the clock customer support
- Convenient online submission
- Thorough peer review
- Inclusion in PubMed and all major indexing services
- Maximum visibility for your research

Submit your manuscript at
www.biomedcentral.com/submit

