Shanghai, China

⁴Laboratory Center, Affiliated Eye and ENT Hospital of Fudan University, Shanghai, China
 ⁵Institute of Stem Cell and Regeneration Medicine, Institutions of Biomedical Science, Fudan University, Shanghai, China

⁶Key Laboratory of Hearing Science, Ministry of Health, EENT Hospital, Fudan University, Shanghai, China

Effect of JNK inhibitor SP600125 on hair cell regeneration in

Yingzi He^{1,4,*}, Chengfu Cai^{2,*}, Shaoyang Sun³, Xu Wang³, Wenyan Li¹, Huawei Li^{1,5,6} ¹Department of Otorhinolaryngology, Key Laboratory of Hearing Science, Ministry of Health, EENT Hospital, Fudan University,

²Department of Otolaryngology Head and Neck Surgery, the First Affiliated Hospital, Xiamen University, Xiamen, China
³Kev Laboratory of Metabolism and Molecular Medicine, the Ministry of Education, Department of Biochemistry and Molecular

^{*}These authors contributed equally to this work

Correspondence to: Huawei Li, email: hwli@shmu.edu.cn

zebrafish (Danio rerio) larvae

Keywords: JNK, SP600125, hair cell regeneration, zebrafish, Wnt

Received: January 25, 2016 **Accepted:** June 29, 2016

Published: July 12, 2016

ABSTRACT

The c-Jun amino-terminal kinase (JNK) proteins are a subgroup of the mitogen-activated protein kinase family. They play a complex role in cell proliferation, survival, and apoptosis. Here, we report a novel role of JNK signalling in hair cell regeneration. We eliminated hair cells of 5-day post-fertilization zebrafish larvae using neomycin followed by JNK inhibition with SP600125. JNK inhibition strongly decreased the number of regenerated hair cells in response to neomycin damage. These changes were associated with reduced proliferation. JNK inhibition also increased cleaved caspase-3 activity and induced apoptosis in regenerating neuromasts. Finally, JNK inhibition with SP600125 decreased the expression of genes related to Wnt. Over-activation of the Wnt signalling pathway partly rescued the hair cell regeneration defects induced by JNK inhibition. Together, our findings provide novel insights into the function of JNK and show that JNK inhibition blocks hair cell regeneration by controlling the Wnt signalling pathway.

INTRODUCTION

Cochlear hair-cell loss in humans causes progressive and permanent hearing loss [1–3]. Nonmammalian vertebrates, however, are capable of regenerating lost sensory hair cells after injury [4–6]. The zebrafish (*Danio rerio*) has an important mechanosensory system – the lateral line system – that helps the fish avoid obstacles and predators [7]. The lateral line neuromasts are made up of a group of sensory hair cells and supporting cells [8], and the hair cells in neuromasts are similar to the mammalian inner ear sensory hair cells in terms of both morphology and function [9, 10]. Zebrafish are able to regenerate hair cells occurs within 48 hours [6, 11, 12]. The new hair cells in neuromasts usually arise through proliferation and differentiation of non-sensory supporting cells during the process of proliferative regeneration [6, 11–13]. Thus, numerous studies have used the lateral line system to investigate the mechanisms of hair cell differentiation and regeneration [14–17].

The c-Jun amino-terminal kinase (JNK) signalling pathway is a member of the large mitogen-activated protein kinase family. Upon activation, JNK executes its functions in cell growth, differentiation, proliferation, apoptosis, and inflammatory responses [18–21]. Subsequent studies suggested that JNK is implicated in the incidence and progression of cancer, the immune response in mammals, and apoptosis in a variety of experimental models. It has become an important and promising target for the treatment of a wide variety of diseases such as chronic inflammation, diabetes, and tumours [22–26]. The JNK family is encoded by three genes, *jnk1*, *jnk2*, and *jnk3*. The *jnk1* and *jnk2* genes are ubiquitously expressed, whereas the *jnk3* gene is restricted to the brain, heart, and testes [20, 27–29]. It has been reported that JNK signal is related to many physiological and pathological processes, such as neuron sprouting [30], tubulin dynamics in migrating neurons [31], and the progression of cancer [32]. More recently, JNK has emerged as an important regulator of the processes of regeneration. In planarians, the conserved JNK signalling cascade is required for regeneration of posterior tissues. Loss of JNK function blocks planarian posterior regeneration because the stem-cell dependent Wnt signalling expression fails to establish itself after posterior injury [33]. Two recent studies show that JNK activity is required for wound healing, for driving stem cell mitosis, and for correctly triggering cell death during planarian regeneration [34, 35]. However, the specific function of the JNK pathway in hair cell regeneration is still not well understood.

The purpose of this study was to investigate the effects of JNK on hair cell regeneration. We show that JNK inhibition with SP600125 reduced the numbers of hair cells, decreased cellular proliferation, and induced cell death in the zebrafish lateral line neuromast following neomycin-induced hair cell loss. We further provide evidence that SP600125 attenuated the expression of genes related to Wnt activation. The phenotype of regenerating hair cells induced by JNK inhibition can be partly rescued by over-activation of the Wnt signalling pathway. These results suggest that JNK supports the regenerative proliferation of hair cells by controlling the Wnt signalling pathway.

RESULTS

JNK inhibition disrupts the regeneration of lateral line hair cells

After 400 µM neomycin treatment for 1 h, most of the hair cells in the lateral line were eliminated, but regeneration occurred rapidly over the following 48 h. To investigate the effect of JNK inhibition on hair cell regeneration, neomycin-treated larvae were placed in 6-well plates and exposed to different doses of SP600125 during recovery periods of 24 h or 48 h. Specific labelling of newly generated hair cells was confirmed using the transgenic zebrafish line Brn3c:mGFP. Our results showed that SP600125 significantly decreased the number of regenerated hair cells after neomycin damage. In the 24 h group, 5.66 ± 0.11 GFP-positive hair cells were found in neuromasts (n = 100) of the control larvae (Figure 1A2), but the mean value of GFP-positive hair cells per neuromast was 4.8 ± 0.22 (n = 40), 3.62 ± 0.15 (n = 60), and 2.91 \pm 0.15 (*n* = 32) in the 5 μ M treated, 10 μ M treated (Figure 1B2), and 15 µM treated fish, respectively (Figure 1E; p < 0.05). At 48 h post-treatment, there were apparent differences in the number of regenerated hair cells between the untreated larvae and the larvae treated with SP600125. The mean number of GFP-positive hair cells per neuromast was 10.64 ± 0.18 in untreated fish (n = 72; Figure 1C2), 7.46 ± 0.25 (n = 28) in 5 µM treated fish, 5.81 ± 0.18 (n = 32) in 10 µM treated fish (Figure 1D2), and 4.59 ± 0.24 (n = 32) in 15 µM treated fish (Figure 1E; p < 0.05). Therefore, we conclude that the hair cell regeneration process in larval neuromasts is severely impaired in the presence of SP600125.

JNK inhibition decreases cellular proliferation in neuromasts

Because the non-sensory supporting cells within the neuromast are the major source of newly regenerated sensory hair cells after neomycin injury [6, 12], we next determined whether SP600125 has any effect on the proliferation of hair cells in neuromasts during the regeneration phase. After neomycin damage, 5 dpf zebrafish larvae were incubated in fresh egg water containing 10 mM BrdU with or without SP600125 at different doses for 24 h or for 48 h. By BrdU incorporation, we observed that the regeneration-associated cell proliferation was significantly inhibited by inhibiting JNK signalling with SP600125. Among the 24 h groups, fewer BrdU-labelled cells were found in SP600125-treated groups compared with the controls (Figure 1A4, 1B4, and 1F; p < 0.05). After 48 h of continuous BrdU incorporation, there were significant differences in the number of BrdU-positive cells per neuromast between the control larvae and the larvae treated with SP600125 (Figure 1C4, D4, and 1F; p < 0.05) indicating that SP600125 significantly decreased the proportion of neuromast cells undergoing active cell division.

To distinguish the new mitotically regenerated hair cells from cell proliferation, we double-labelled the zebrafish larvae with anti-BrdU and anti-GFP antibodies at 24 h and 48 h after neomycin damage. Our analysis showed that SP600125-treated larvae had fewer BrdU-positive hair cells in the regenerating neuromasts. In control larvae at 24 h post-treatment, the ratio of BrdU and GFP double-labelled cells to the total number of GFP-positive hair cells in the neuromast was 0.46 ± 0.016 (*n* = 100 neuromasts), while the ratio in 10 µM SP600125-treated fish was significantly lower at 0.12 ± 0.026 (n = 60 neuromasts, p < 0.001) (Figure 1H). Among the 48 h groups, a significant increase in BrdU-positive and GFP-positive hair cells was observed in control fish, while there was a significant decrease in BrdU-positive hair cells per neuromast after exposure to 10 μM SP600125 (Figure 1H; *p* < 0.001).

To further explore the effect of JNK inhibition, we quantitatively assessed the Sox2-labeled supporting cells in the neuromasts after treatment with different concentrations of SP600125 and observed a dose-dependent reduction in the number of stained supporting cells (Figure 1A3, 1B3, 1C3, 1D3 and 1G; p < 0.001). We next evaluated the rates of supporting cell division among different groups after neomycin insult by BrdU



Figure 1: SP600125 decreases regeneration of hair cells in zebrafish lateral line neuromasts. (A-D) We treated 5 dpf Tg(Brn3c:mGFP) zebrafish with 400 µM neomycin for 1 h and then treated them for 24 h or 48 h with 10 µM SP600125 and subsequently imaged GFP-positive hair cells (green), Sox2-positive supporting cells (red), and BrdU-positive replicating cells (white). SP600125 significantly decreased the numbers of GFP-positive hair cells and Sox2-positive supporting cells in neuromasts as well as reduced the proportion of cells in S-phase as indicated by BrdU staining. Scale bars = $10 \mu m$. Higher magnification of hair cells and supporting cells of the neuromast taken from z-stacks show that hair cells and supporting cells in untreated controls and SP600125-treated animals had no observable morphological differences though there were fewer GFP-positive and Sox2-positive cells in the neuromasts of larvae treated with SP600125. (E) Quantification of the number of hair cells in control and SP600125-treated larvae at 24 hours and 48 hours after neomycin incubation. (F) Quantification of replicating cells in control and SP600125-treated larvae at 24 hours and 48 hours after neomycin incubation. (G) Quantification of the number of Sox2-positive cells in control and SP600125-treated larvae at 24 hours and 48 hours after neomycin incubation. In the 24-hour group, n = 100 control neuromasts, $n = 40.5 \mu M$ SP600125-treated neuromasts, n = 6010 μ M SP600125-treated neuromasts, and $n = 32.15 \mu$ M SP600125-treated neuromasts. In the 48-hour group, n = 72 control neuromasts, $n = 28.5 \,\mu\text{M}$ SP600125-treated neuromasts, $n = 32.10 \,\mu\text{M}$ SP600125-treated neuromasts, and $n = 32.15 \,\mu\text{M}$ SP600125-treated neuromasts. *p < 0.05. (24-hour group: One-way ANOVA; GFP+ cells: F3, 228 =71.15, p < 0.05; Sox2+ cells: F3, 228 = 38.48, p < 0.05; BrdU+ cells: F3, 228 = 172.5, p < 0.05. 48-hour group: One-way ANOVA; GFP+ cells: F3, 160 = 184.9, p < 0.05; Sox2+ cells: F3, 160 = 90.65, p < 0.05; BrdU+ cells: F3, 160 = 365.5, p < 0.05). Bars are mean \pm s.e.m. (H, I) Quantification of the ratio of BrdU-positive hair cells and the ratio of BrdU-positive supporting cells in control and SP600125-treated larvae at 24 hours and 48 hours after neomycin incubation. Bars are mean \pm s.e.m. In the 24-hour group, n = 100 control neuromasts and $n = 60.10 \,\mu\text{M}$ SP600125-treated neuromasts. In the 48-hour group, n = 72 control neuromasts and $n = 32.10 \mu$ M SP600125-treated neuromasts. **p < 0.001. (24-hour group: BrdU+ HCs: unpaired t test, two-tailed, t = 11.54, df = 158, p < 0.001; BrdU+ SCs: unpaired t test, two-tailed, t = 10.5, df = 158, p < 0.001. 48-hour group: BrdU+ HCs: unpaired t test, two-tailed, t = 16.74, df = 102, p < 0.001; BrdU+ SCs: unpaired t test, two-tailed, t = 4.922, df = 102, p < 0.001). Bars are mean \pm s.e.m.

and Sox2 immunolabelling. After both 24 h and 48 h of regeneration, the ratio of BrdU and Sox2 double-labelled cells to the total number of Sox2-labeled cells in the 10 μ M SP600125-treated neuromasts was drastically reduced (Figure 1A3, 1B3, 1C3, 1D3 and 1I; p < 0.001), and this most likely explains the reduction in the number of newly regenerated hair cells in the SP600125-treated fish. Taken together, these results show that JNK inhibition has a significant negative impact on proliferation in the regenerating neuromast.

JNK inhibition induces apoptosis in neuromast cells

In addition to its role in regenerative cell proliferation, JNK signalling has also been implicated in cellular apoptosis [36]. To evaluate the effect of SP600125 on apoptosis, we labelled zebrafish larvae with anti-cleaved caspase-3 antibody. As shown in Figure 2A, we occasionally detected cleaved caspase-3-positive cells in controls. However, the emergence of cleaved

caspase-3-positive cells became frequent in larvae treated with 15 μ M SP600125 for 48 h (Figure 2B, 2C; p < 0.001). This was shown in the western blot analysis of proteins from zebrafish larvae (Figure 2D). However, we cannot rule out the possibility that these proteins might also be affected in other tissues where JNK is expressed because the proteins used for immunoblot analysis are isolated from the whole larvae, not only the neuromasts. We further performed a TUNEL analysis on control and SP600125-treated larvae to investigate the roles of JNK in apoptosis. At 48 h post-treatment, 15 μ M SP600125-treated larvae had significantly greater numbers of TUNEL-positive cells when compared to controls (Supplementary Figure S1). Thus, apoptosis appears to contribute to the regeneration defect caused by the JNK inhibitor.

JNK inhibition inhibits the Wnt signalling pathway

Wnt signalling is an important player in controlling the regeneration period, and a number of genes have



Figure 2: SP600125 induces apoptosis in neuromasts. (A–B) Cleaved caspase-3 staining in the neuromast from a control larva (A) and 15 μ M SP600125-treated larvae (B). Scale bar = 10 μ m. (C) SP600125 treatment increased the numbers of cleaved caspase-3-positive cells. Bars are mean \pm s.e.m. *n* = 48 control neuromasts and *n* = 48 15 μ M SP600125-treated neuromasts. ***p* < 0.001. (unpaired *t* test, two-tailed, *t* = 4.051, df = 94, *p* < 0.001). (D) After treatment of larvae with 15 μ M SP600125 for 48 h, protein extracts were prepared and subjected to western blot assay using antibodies against caspase-3 and cleaved caspase-3. β -Actin was included as the loading control.

been shown to be required for this process by potentially interacting with this pathway [37-39]. Thus we investigated whether JNK inhibition reduces hair cell regeneration by disrupting the regulation of the Wnt pathway in the neuromast (Figure 3). We examined the expression of the Wnt/ β -catenin targets axin2, β -catenin1, and β -catenin2 (ctnnb1 and ctnnb2 according to the Zebrafish Information Network) as well as β -catenin transcription factor *tcf7l2* in the neuromast using in situ hybridization. After neomycin treatment, axin2, ctnnb1, ctnnb2, and tcf7l2 were up-regulated in the control larvae, indicating that the Wnt pathway is active during hair cell regeneration. In contrast, treatment with SP600125 after exposure to neomycin resulted in a striking down-regulation of the expression of these genes, which was confirmed by the western blot analysis (Supplementary Figure S2). To investigate these findings in more detail, we performed in situ hybridization experiments with Wnt/β-catenin pathway genes in non-neomycin-treated control larvae and SP600125-treated larvae. Our analysis showed that larvae with undamaged hair cells showed no significant difference in the expression of Wnt/ β -catenin signaling components at the time examined with or without SP600125 treatment (Supplementary Figure S3).

These results led to the possibility that JNK supports hair cell regeneration by regulating the Wnt signaling pathway, and we hypothesized that overactive Wnt signalling should rescue the JNK inhibition phenotype. For these studies, we used *apc* mutant zebrafish in which loss of *apc* leads to constitutive Wnt signalling activation. We stained the *apc* mutants with FM1-43FX to determine if hair cell mechanotransduction channels were functional. As shown in Figure 4, at 24 h

post-treatment, Wnt overexpression partly reversed the effect of SP600125 on hair cell regeneration. We wanted to determine whether exogenous activation of the Wnt pathway could rescue the effect of SP600125 on hair cell regeneration. To test this, the canonical Wnt activator BIO $(1 \mu M)$ was applied during the regenerative process following neomycin-induced hair cell death along with either 10 μ M SP600125 or DMSO as the control treatments. At 24 h post-treatment, neuromasts in both types of neomycin-treated larvae (neomycin alone or neomycin/SP600125) contained significantly fewer hair cells than neomycin/BIO larvae (Supplementary Figure S4; p < 0.0001) indicating that BIO treatment has a significant effect on hair cell regeneration. Furthermore, neomycin/BIO/SP600125-treated larvae had significantly more hair cells in neuromasts when compared with neomycin/SP600125-treated animals (Supplementary Figure S4; p < 0.05), but had fewer hair cells when compared with neomycin-treated animals (Supplementary Figure S4; p < 0.0001), indicating that the SP600125-induced hair cell regeneration defect could be partly rescued by BIO treatment. These results are consistent with rescue studies using apc mutants. All these findings suggested that JNK might support hair cell regeneration by controlling the Wnt pathway.

DISCUSSION

Previous work in zebrafish showed that JNK activity is required for regeneration-associated cell proliferation and cell cycle activation of the blastema cells [40]. However, little was known regarding the effect of JNK inhibition on hair cell regeneration. In this



Figure 3: *In situ* hybridization of Wnt pathway-related genes in SP600125-treated and untreated larvae at 24 h after **neomycin treatment.** Expression of *axin2*, *ctnnb1*, *ctnnb2*, and *tcf7l2* is significantly decreased in SP600125-treated neuromasts during the regeneration period. (*axin2*: n = 28 neuromasts from control animals, n = 20 neuromasts from SP600125-treated larvae; *ctnnb1*: n = 20 neuromasts from control animals, n = 16 neuromasts from SP600125-treated larvae; *ctnnb2*: n = 18 neuromasts from control animals, n = 16 neuromasts from SP600125-treated larvae; *tcf7l2*: n = 22 neuromasts from control animals, n = 20 neuromasts from SP600125-treated larvae; *tcf7l2*: n = 22 neuromasts from control animals, n = 20 neuromasts from SP600125-treated larvae; *tcf7l2*: n = 22 neuromasts from control animals, n = 20 neuromasts from SP600125-treated larvae; *tcf7l2*: n = 22 neuromasts from control animals, n = 20 neuromasts from SP600125-treated larvae; *tcf7l2*: n = 21 neuromasts from control animals, n = 16 neuromasts from SP600125-treated larvae; *tcf7l2*: n = 21 neuromasts from control animals, n = 20 neuromasts from SP600125-treated larvae; *tcf7l2*: n = 21 neuromasts from control animals, n = 20 neuromasts from SP600125-treated larvae; *tcf7l2*: n = 21 neuromasts from control animals, n = 16 neuromasts from one representative neuromast are shown. Scale bar = $10 \mu m$.

report, we provide evidence that JNK supports hair cell regeneration in larval neuromasts and that JNK inhibition by SP600125 results in fewer hair cells and supporting cells in regenerative neuromasts.

Several studies have suggested a role for the JNK pathway in cell-cycle regulation [19, 21], and the JNK inhibitor SP600125 is well known to induce G_2/M phase cell-cycle arrest and apoptosis in various cancer cells [24, 25, 41]. Our results agree well with previous reports showing that inhibition of JNK by SP600125 leads to a drastic decrease of proliferation as assessed by BrdU staining in zebrafish. Given that the cell proliferation defect in the regenerating neuromasts of SP600125-treated fish could also be due to induction of apoptosis, we

performed a cell death analysis by TUNEL analysis and cleaved caspase-3 staining. Our present data clearly show that SP600125 treatment significantly induced apoptosis in the neuromasts.

Recent studies have suggested that the Wnt signalling pathway plays pivotal roles in a wide variety of developmental processes by controlling cell proliferation, cell fate determination, and several cellular polarization events [42–45], and it is also heavily implicated in regeneration processes in several tissues [46, 47]. In this report, it has been shown that Wnt pathway genes *axin2*, *ctnnb1*, *ctnnb2*, and *tcf7l2* are stimulated in neuromast cells at 24 h of regeneration, and this supports previous studies that Wnt signalling activation increases the degree of hair



Figure 4: Decreased hair cell regeneration in JNK inhibition zebrafish can be partly rescued by over-activating Wnt signalling. (A–B) The FM1-43FX labelled hair cells are significantly increased in *apc* mutant zebrafish compared to control larvae in SP600125-treated and untreated larvae at 24 h after neomycin treatment. Scale bar = 10 µm. (C) Quantification of FM1-43FX-positive hair cells. *p < 0.05; **p < 0.001. (n = 53 neuromasts from the non-neomycin-treated 5 dpf control animals, n = 44 neuromasts from the non-neomycin-treated *apc* mutant zebrafish at 5 dpf, n = 72 neuromasts from the control larvae at 24 h following neomycin damage, n = 11 neuromasts from *apc* mutant larvae at 24 h following neomycin damage, n = 21 neuromasts from 15 µM SP600125-treated larvae at 24 h following neomycin damage. Con 5 dpf vs. *apc* mutant 5 dpf: unpaired *t* test, two-tailed, t = 11.39, df = 95, p < 0.001; Con neo-24 h vs. *apc neo*-24 h: unpaired *t* test, two-tailed, t = 2.767, df = 81, p = 0.007; Con neo SP600125-24 h vs. *apc* neo SP600125-24 h: unpaired *t* test, two-tailed, t = 5.186, df = 82, p < 0.0001).

cell regeneration after damage [46, 47, 50]. However, JNK inhibition with SP600125 decreases the expression of these Wnt signalling genes, raising the possibility that activated Wnt signalling might cooperate with the JNK pathway to support hair cell regeneration. Our results suggest that inducing constitutively active Wnt either in *apc* mutant zebrafish or with BIO treatment could partly rescue the hair cell-less phenotype caused by abrogation of JNK signalling during regeneration. However, we did not observe complete regeneration in neomycin/SP600125-treated/ BIO-recovery larvae, which suggests that other molecular mechanisms or signaling pathways might be involved in the JNK-induced hair cell regeneration.

Many extensive cross-regulatory interactions between JNK and both canonical and noncanonical Wnt signalling pathways have been reported previously in a number of processes in many organisms, including both the promotion and repression of Wnt activity [51–55] For example, previous work has shown that Wnt5a, a noncanonical Wnt ligand, is capable of activating JNK. Appropriate activation of JNK is able to function downstream of Wnt5a to correct convergent extension movements in Xenopus [56]. Wnt7a stimulates JNK activation and c-Jun phosphorylation in non-small cell lung cancer cells [57]. More evidence for the well-characterized crosstalk between JNK and the canonical Wnt signalling pathway has now appeared in mammalian cells and zebrafish [51, 58]. The connection between these two pathways might result in multiple correlations of expression among phosphorylated c-Jun (p-c-Jun), TCF4, and β-catenin target genes. Several groups have reported that the p-c-Jun protein cooperates with the HMG family transcription factor TCF4 to form a functional transcription complex that cooperatively enhances their transcriptional activity in the presence of β -catenin, and this complex plays a critical role in mouse intestinal tumor development. Abrogation of JNK signalling by pharmacological inhibition leads to a failure in c-Jun phosphorylation and inhibition of the phosphorylation-dependent interactions between c-Jun and TCF4 [58]. In addition, Gan and colleagues demonstrated a novel role of JNK signalling in regulating the Wnt signalling via direct interaction of c-Jun with Dishevelled, a pivotal regulator of the canonical Wnt signalling pathway. c-Jun can function as a scaffold protein within the β -catenin–TCF transcription complex to bridge Dvl to TCF on the Wnt target gene promoters and stimulate gene transcription [51]. The crosstalk between JNK and the canonical Wnt pathways in regeneration processes will provide important directions for future studies regarding how JNK regulates hair cell regeneration.

In conclusion, our present study supports a novel role for JNK in hair cell regeneration. JNK inhibition with SP600125 decreases hair cell regeneration in larval zebrafish, an this defect can be partly rescued by over-activation of the Wnt signalling pathway. Taken together, our study provides new insights into the mechanisms of hair cell regeneration in the zebrafish lateral line and proposes that JNK activation might represent a useful therapeutic strategy in the treatment of hearing loss.

MATERIALS AND METHODS

Zebrafish embryos and drug administration

Zebrafish embryos were maintained in our facility according to standard procedures. The *apc* mutant transgenic line was obtained from Professor Xu Wang. The ages of the zebrafish larvae are described as days post fertilization (dpf). SP600125 (Sigma-Aldrich, St Louis, MO, USA) was dissolved in dimethyl sulfoxide (DMSO, Sigma-Aldrich) at a stock concentration of 50 mM and further diluted to the desired concentrations in fresh egg water. Dose-response data were obtained by treating larvae with SP600125 (5 μ M, 10 μ M, and 15 μ M) after hair cell damage. For hair cell damage, neomycin sulphate (Sigma-Aldrich) was added to 5 dpf larvae at a final concentration of 400 μ M and incubated for 1 h. This was followed by three rinses in fresh egg water, and the zebrafish larvae were allowed to recover for 24 h or 48 h at 28.5°C.

Cell proliferation and analysis

Proliferating cells in the lateral line neuromasts were labelled by adding 10 mM 5-bromo-2-deoxyuridine (BrdU; Sigma-Aldrich) to the fresh egg water for 24 h or 48 h at 28.5°C. Larvae were then fixed with 4% PFA overnight at 4°C, and BrdU incorporation was detected by fluorescent immunostaining. The fixed larvae were washed three times in PBS containing 0.5% Triton X-100 (PBT-2) and placed in 2 N HCl for 0.5 h at 37°C. Larvae were blocked in 10% normal goat serum for 1 h at room temperature and incubated with the monoclonal primary anti-BrdU antibody (1:200 dilution; Santa Cruz, Dallas, TX, USA. Cat. no. sc-32323) overnight at 4°C. The next day, larvae were washed three times for 10 min each with PBT-2 and then incubated with the secondary antibody for 1 h at 37°C. Fluorescently labelled larvae were imaged with a Leica confocal fluorescence microscope (TCS SP5; Leica, Wetzlar, Germany). Images were processed using Photoshop software (Adobe).

Immunohistochemistry

For immunohistochemistry analysis, larvae were fixed with 4% PFA and were permeabilized with PBT-2 for 30 min followed by incubation in blocking solution for 1 h. The following antibodies were used as primary antibodies: anti-GFP (1:1000 dilution; Abcam, Cambridge, UK); anti-Sox2 (1:200 dilution; Abcam); and anti-cleaved caspase-3 (1:500 dilution; Cell Signaling Technology Inc., Danvers, MA, USA). The larvae were washed three times with PBT-2 and incubated with secondary antibodies to detect primary antibodies. Nuclei were labelled with 4,6-diamidino-2- phenylindole (DAPI; 1:800 dilution; Invitrogen, Carlsbad, CA, USA) for 20 min at room temperature.

FM1-43FX labelling

To visualize and image the hair cells in lateral line neuromasts, the vital dye FM1-43FX (Molecular Probes, Eugene, OR, USA) was applied at a concentration of 3 μ M to live larvae for 45 s in the dark. After quickly rinsing three times with fresh water, the larvae were anesthetized in 0.02% MS-222 and fixed with 4% PFA in PBS for 2 h at room temperature or overnight at 4°C.

Western blot analysis

Total protein was isolated with the AllPrep DNA/RNA/Protein Mini Kit (QIAGEN, Hilden, Germany) according to the manufacturer's instructions. Protein concentrations were measured using a BCA protein kit (Thermo Fisher Scientific, Rockford, IL), and proteins were separated on SDS-polyacrylamide gels and transferred onto PVDF membranes (Immobilon-P; Millipore, Bedford, MA, USA). The membranes were blocked with 5% nonfat dried milk in TBST (50 mM Tris-HCl (pH 7.4), 150 mM NaCl, and 0.1% Tween-20) for 1 h at room temperature and then blotted overnight with primary antibodies at 4°C. The following antibodies were used as primary antibodies: anti-cleaved caspase-3 (1:1000 dilution; Cell Signaling Technology), anti-caspase-3 (1:1000 dilution; Abcam), anti-axin2 (1:1000 dilution; Abcam), and anti-tcf7l2 (1:1000 dilution; Abcam).

Whole-mount in situ hybridization

The probes used in *in situ* hybridization (axin2, ctnnb1, ctnnb2, and tcf7l2) were amplified by PCR from zebrafish embryo cDNA using the following primers and cloned into the pGEM-T Easy Vector (Promega, cat. no. A1360): axin2 forward: 5'-accgacaaaccaagcaca ag-3'; axin2 reverse: 5'-tccgttttgagttatgaagctct-3'; ctnnb1 forward: 5'-cccaggactacaagaagcga-3'; ctnnb1 reverse: 5'-acaggcaaggctaaggttga-3'; ctnnb2 forward: 5'-catcgag aacatccagcgtg-3';ctnnb2 reverse: 5'-tggactacactacagcc gtc-3'; tcf7l2 forward: 5'-ccctccacatctacagggag-3'; tcf7l2 reverse: 5'-tgtgttcattgccctctcct-3'. Digoxigeninlabeled antisense RNA probes were generated by in vitro transcription using T7 or SP6 RNA polymerase. Regular whole-mount in situ hybridization of zebrafish embryos was performed as previously described [17, 59]. Details of the probes used are available on request. Briefly, the embryos were depigmented with 1-phenyl-2-thiourea (PTU, Sigma-Aldrich, cat. no. P7629), euthanized in MS-222, and fixed overnight with 4% PFA at 4°C. The fixed embryos were washed in PBS with 0.1% Tween-20 (PBST) and placed in 100% methanol at -20°C for dehydration. Prior to use, they were rehydrated in a graded methanol series and washed three times for 5 min with PBST. To permeabilize the embryos, proteinase K (10 µg/mL in PBST) was added for 50 min and the embryos were refixed in 4% PFA for 20 min. After washing in PBST, the embryos were prehybridized at 65°C for ≥ 2 h in hybridization buffer. For hybridization, the labeled probes were added to the hybridization buffer at 65°C overnight. After washing for 15 min with 75%, 50%, and 25% hybridization buffer and $2 \times$ SSCT ($20 \times$ SSC, Life technologies, AM9770; 0.1% Tween-20) and for 30 min twice in $0.2 \times$ SSC at 65°C, embryos were blocked for at least 2 h at 4°C in blocking buffer (Roche cat. no. 11096176001) and were incubated with preabsorbed sheep anti-digoxigenin-AP Fab fragments (Roche cat. no. 11093274910) at a 1:4000 dilution in blocking buffer overnight at 4°C. The next day, the embryos were washed 4×30 min with 2 mg/mL BSA in PBST and 3×5 min in staining buffer (100 mM Tris (pH 9.5), 100 mM NaCl, and 0.1% Tween-20). Afterwards, the embryos were stained with BM purple AP substrate (Roche cat. no. 11 442074001) in the dark. Finally, the color reaction was stopped by adding PBST, and the embryos were observed under a bright field microscope (Nikon Instruments).

TUNEL staining

For TUNEL (Terminal deoxynucleotidyl transferasemediated dUTP nick end labeling) assays, larvae were incubated in 0.1 M glycine/PBS solution for 10 min and then rinsed with PBT-2 three times for 10 minutes each. The larvae were then processed using the *In Situ* Cell Death Detection Kit (Roche, Nutlet, NJ, USA; cat. no.11684795910) following the directions supplied by the manufacturer.

Statistical analysis

Prior to analysis, all data were first examined for normality and homogeneity of variances by the Shapiro–Wilk test and Levene's test, respectively. For statistical comparisons, differences among groups were compared using one-way ANOVA, and differences between groups were compared using an unpaired *t*-test (2-tail) (see figure legends for details). Data were analyzed using SigmaPlot (version 12.0 for Windows; Systat Software Inc., CA, USA). All data are presented as the mean \pm s.e.m. A *p*-value < 0.05 was considered statistically significant, and *p* < 0.001 was considered highly significant.

ACKNOWLEDGMENTS AND FUNDING

The authors would like to thank Dong Liu and Min Yu for their technical assistance and Yalin Huang for help with the confocal microscope. This work was supported by grants from the National Natural Science Foundation of China (Nos. 81230019, 81470687, 81500784), the Specialized Research Fund for the Doctor Program of Higher Education (20120071110077), the Construction Program of Shanghai Committee of Science and Technology (12DZ2251700), and the Fundamental Research Funds for the Central Universities (2242014R30022, 021414380037).

CONFLICTS OF INTEREST

The authors declare that they have no competing interests.

REFERENCES

- 1. Forge A, Li L, Corwin JT, Nevill G. Ultrastructural Evidence for Hair Cell Regeneration in the Mammalian Inner-Ear. Science. 1993; 259:1616–1619.
- Chai RJ, Kuo B, Wang T, Liaw EJ, Xia AP, Jan TA, Liu ZY, Taketo MM, Oghalai JS, Nusse R, Zuo J, Cheng AG: Wnt signaling induces proliferation of sensory precursors in the postnatal mouse cochlea. Proc Natl Acad Sci USA. 2012; 109:8167–8172.
- Shi FX, Hua LX, Edge ASB. Generation of hair cells in neonatal mice by beta-catenin overexpression in Lgr5positive cochlear progenitors. Proc Natl Acad Sci USA. 2013; 110:13851–13856.
- Corwin JT, Cotanche DA. Regeneration of Sensory Hair-Cells after Acoustic Trauma. Science. 1988; 240: 1772–1774.
- Balak KJ, Corwin JT, Jones JE. Regenerated Hair-Cells Can Originate from Supporting Cell Progeny - Evidence from Phototoxicity and Laser Ablation Experiments in the Lateral Line System. J Neurosci. 1990; 10:2502–2512.
- Harris JA, Cheng AG, Cunningham LL, MacDonald G, Raible DW, Rubel EW. Neomycin-induced hair cell death and rapid regeneration in the lateral line of zebrafish (Danio rerio). J Assoc Res Otolaryngol. 2003; 4:219–234.
- Montgomery J, Carton G, Voigt R, Baker C, Diebel C. Sensory processing of water currents by fishes. Philos Trans R Soc Lond B Biol Sci. 2000; 355:1325–1327.
- Metcalfe WK, Kimmel CB, Schabtach E. Anatomy of the Posterior Lateral Line System in Young Larvae of the Zebrafish. J Comp Neurol. 1985; 233:377–389.
- Raible DW, Kruse GJ. Organization of the lateral line system in embryonic zebrafish. J Comp Neurol. 2000; 421:189–198.
- 10. Nicolson T. The genetics of hearing and balance in zebrafish. Annu Rev Genet. 2005; 39:9–22.

- 11. Williams JA, Holder N. Cell turnover in neuromasts of zebrafish larvae. Hear Res. 2000; 143:171–181.
- 12. Ma EY, Rubel EW, Raible DW. Notch signaling regulates the extent of hair cell regeneration in the zebrafish lateral line. J Neurosci. 2008; 28:2261–2273.
- Hernandez PP, Olivari FA, Sarrazin AF, Sandoval PC, Allende ML. Regeneration in zebrafish lateral line neuromasts: Expression of the neural progenitor cell marker Sox2 and proliferation-dependent and-independent mechanisms of hair cell renewal. Dev Neurobiol. 2007; 67:637–654.
- He YZ, Mei HL, Yu HQ, Sun S, Ni WL, Li HW. Role of histone deacetylase activity in the developing lateral line neuromast of zebrafish larvae. Exp Mol Med. 2014; 46.
- He YZ, Cai CF, Tang DM, Sun S, Li HW. Effect of histone deacetylase inhibitors trichostatin A and valproic acid on hair cell regeneration in zebrafish lateral line neuromasts. Front Cell Neurosci. 2014; 8.
- He YZ, Yu HQ, Sun S, Wang YF, Liu LM, Chen ZY, Li HW. Trans-2-phenylcyclopropylamine regulates zebrafish lateral line neuromast development mediated by depression of LSD1 activity. Int J Dev Biol. 2013; 57:365–373.
- He Y, Wu J, Mei H, Yu H, Sun S, Shou J, Li H. Histone deacetylase activity is required for embryonic posterior lateral line development. Cell Prolif. 2014; 47:91–104.
- Han J, Ulevitch RJ. Emerging targets for anti-inflammatory therapy. Nat Cell Biol. 1999; 1:E39-E40.
- 19. Davis RJ. Signal transduction by the JNK group of MAP kinases. Cell. 2000; 103:239–252.
- 20. Weston CR, Davis RJ. The JNK signal transduction pathway. Curr Opin Cell Biol. 2007; 19:142–149.
- 21. Lin A. Activation of the JNK signaling pathway: breaking the brake on apoptosis. Bioessays. 2003; 25:17–24.
- Wilhelm M, Xu ZH, Kukekov NV, Gire S, Greene LA. Proapoptotic Nix activates the JNK pathway by interacting with POSH and mediates death in a Parkinson disease model. J Biol Chem. 2007; 282:1288–1295.
- Storling J, Binzer J, Andersson AK, Zullig RA, Tonnesen M, Lehmann R, Spinas GA, Sandler S, Billestrup N, Mandrup-Poulsen T. Nitric oxide contributes to cytokineinduced apoptosis in pancreatic beta cells via potentiation of JNK activity and inhibition of Akt. Diabetologia. 2005; 48:2039–2050.
- Mingo-Sion AM, Marietta PM, Koller E, Wolf DM, Van Den Berg CL. Inhibition of JNK reduces G2/M transit independent of p53, leading to endoreduplication, decreased proliferation, and apoptosis in breast cancer cells. Oncogene. 2004; 23:596–604.
- Hideshima T, Hayashi T, Chauhan D, Akiyama M, Richardson P, Anderson K. Biologic sequelae of c-Jun NH2-terminal kinase (JNK) activation in multiple myeloma cell lines. Oncogene. 2003; 22:8797–8801.
- Johnson GL, Nakamura K. The c-jun kinase/stress-activated pathway: regulation, function and role in human disease. Biochim Biophys Acta. 2007; 1773:1341–1348.

- Yoshida S, Fukino K, Harada H, Nagai H, Imoto I, Inazawa J, Takahashi H, Teramoto A, Emi M. The c-Jun NH2-terminal kinase3 (JNK3) gene: genomic structure, chromosomal assignment, and loss of expression in brain tumors. J Hum Genet. 2001; 46:182–187.
- Gupta S, Barrett T, Whitmarsh AJ, Cavanagh J, Sluss HK, Derijard B, Davis RJ. Selective interaction of JNK protein kinase isoforms with transcription factors. Embo J. 1996; 15:2760–2770.
- Kallunki T, Su B, Tsigelny I, Sluss HK, Derijard B, Moore G, Davis R, Karin M. Jnk2 Contains a Specificity-Determining Region Responsible for Efficient C-Jun Binding and Phosphorylation. Gene Dev. 1994; 8: 2996–3007.
- Eminel S, Roemer L, Waetzig V, Herdegen T. c-Jun N-terminal kinases trigger both degeneration and neurite outgrowth in primary hippocampal and cortical neurons. J Neurochem. 2008; 104:957–969.
- Kawauchi T, Chihama K, Nabeshima Y, Hoshino M. The in vivo roles of STEF/Tiam1, Rac1 and JNK in cortical neuronal migration. Embo J. 2003; 22:4190–4201.
- Moon DO, Kim MO, Choi YH, Kim ND, Chang JH, Kim GY. Bcl-2 overexpression attenuates SP600125-induced apoptosis in human leukemia U937 cells. Cancer Lett. 2008; 264:316–325.
- Tejada-Romero B, Carter JM, Mihaylova Y, Neumann B, Aboobaker AA. JNK signalling is necessary for a Wnt- and stem cell-dependent regeneration programme. Development. 2015; 142:2413–2424.
- Tasaki J, Shibata N, Sakurai T, Agata K, Umesono Y. Role of c-Jun N-terminal kinase activation in blastema formation during planarian regeneration. Dev Growth Differ. 2011; 53:389–400.
- 35. Almuedo-Castillo M, Crespo X, Seebeck F, Bartscherer K, Salo E, Adell T. JNK controls the onset of mitosis in planarian stem cells and triggers apoptotic cell death required for regeneration and remodeling. PLoS Genet. 2014; 10:e1004400.
- 36. Liu J, Lin A. Role of JNK activation in apoptosis: a doubleedged sword. Cell Res. 2005; 15:36–42.
- Romero-Carvajal A, Navajas Acedo J, Jiang L, Kozlovskaja-Gumbriene A, Alexander R, Li H, Piotrowski T. Regeneration of Sensory Hair Cells Requires Localized Interactions between the Notch and Wnt Pathways. Dev Cell. 2015; 34:267–282
- He Y, Tang D, Cai C, Chai R, Li H. LSD1 is Required for Hair Cell Regeneration in Zebrafish. Mol Neurobiol. 2016; 53:2421–2434.
- 39. Lush ME, Piotrowski T. Sensory hair cell regeneration in the zebrafish lateral line. Dev Dynam. 2014; 243:1187–1202.
- Ishida T, Nakajima T, Kudo A, Kawakami A. Phosphorylation of Junb family proteins by the Jun N-terminal kinase supports tissue regeneration in zebrafish. Dev Biol. 2010; 340:468–479.

- Du LH, Lyle CS, Obey TB, Gaarde WA, Muir JA, Bennett BL, Chambers TC. Inhibition of cell proliferation and cell cycle progression by specific inhibition of basal JNK activity - Evidence that mitotic Bcl-2 phosphorylation is JNK-independent. J Biol Chem. 2004; 279:11957–11966.
- Iglesias DM, Hueber PA, Chu L, Campbell R, Patenaude AM, Dziarmaga AJ, Quinlan J, Mohamed O, Dufort D, Goodyer PR. Canonical WNT signaling during kidney development. Am J Physiol-Renal. 2007; 293:F494–500.
- 43. Maretto S, Cordenonsi M, Dupont S, Braghetta P, Broccoli V, Hassan AB, Volpin D, Bressan GM, Piccolo S. Mapping Wnt/beta-catenin signaling during mouse development and in colorectal tumors. Proc Natl Acad Sci USA. 2003; 100:3299–3304.
- 44. Chai R, Xia A, Wang T, Jan TA, Hayashi T, Bermingham-McDonogh O, Cheng AG. Dynamic expression of Lgr5, a Wnt target gene, in the developing and mature mouse cochlea. J Assoc Res Otolaryngol. 2011; 12:455–469.
- 45. Rosso SB, Inestrosa NC. WNT signaling in neuronal maturation and synaptogenesis. Front Cell Neurosci. 2013; 7.
- Head JR, Gacioch L, Pennisi M, Meyers JR. Activation of canonical Wnt/beta-catenin signaling stimulates proliferation in neuromasts in the zebrafish posterior lateral line. Dev Dynam. 2013; 242:832–846.
- 47. Jacques BE, Montgomery WHt, Uribe PM, Yatteau A, Asuncion JD, Resendiz G, Matsui JI, Dabdoub A. The role of Wnt/beta-catenin signaling in proliferation and regeneration of the developing basilar papilla and lateral line. Dev Neurobiol. 2014; 74:438–456.
- Nusse R. Wnt signaling in disease and in development. Cell Res. 2005; 15:28–32.
- 49. Clevers H. Wnt/beta-catenin signaling in development and disease. Cell. 2006; 127:469–480.
- 50. Wang T, Chai R, Kim GS, Pham N, Jansson L, Nguyen DH, Kuo B, May LA, Zuo J, Cunningham LL, Cheng AG. Lgr5+ cells regenerate hair cells via proliferation and direct transdifferentiation in damaged neonatal mouse utricle. Nat Commun. 2015; 6:6613.
- Gan XQ, Wang JY, Xi Y, Wu ZL, Li YP, Li L. Nuclear Dvl, c-Jun, beta-catenin, and TCF form a complex leading to stabilization of beta-catenin-TCF interaction. J Cell Biol. 2008; 180:1087–1100.
- 52. Liao G, Tao Q, Kofron M, Chen JS, Schloemer A, Davis RJ, Hsieh JC, Wylie C, Heasman J, Kuan CY. Jun NH2-terminal kinase (JNK) prevents nuclear beta-catenin accumulation and regulates axis formation in Xenopus embryos. Proc Natl Acad Sci USA. 2006; 103:16313–16318.
- 53. Cizelsky W, Tata A, Kuhl M, Kuhl SJ. The Wnt/JNK signaling target gene alcam is required for embryonic kidney development. Development. 2014; 141:2064–2074.
- Evason KJ, Francisco MT, Juric V, Balakrishnan S, Lopez Pazmino Mdel P, Gordan JD, Kakar S, Spitsbergen J, Goga A, Stainier DY. Identification of Chemical Inhibitors

of beta-Catenin-Driven Liver Tumorigenesis in Zebrafish. PLoS Genet. 2015; 11:e1005305.

- Lee MH, Koria P, Qu J, Andreadis ST. JNK phosphorylates beta-catenin and regulates adherens junctions. Faseb J. 2009; 23:3874–3883.
- 56. Yamanaka H, Moriguchi T, Masuyama N, Kusakabe M, Hanafusa H, Takada R, Takada S, Nishida E. JNK functions in the non-canonical Wnt pathway to regulate convergent extension movements in vertebrates. EMBO Rep. 2002; 3:69–75.
- 57. Heasley LE, Winn RA. Analysis of Wnt7a-stimulated JNK activity and cJun phosphorylation in non-small cell lung cancer cells. Methods Mol Biol. 2008; 468:187–196.
- Nateri AS, Spencer-Dene B, Behrens A. Interaction of phosphorylated c-Jun with TCF4 regulates intestinal cancer development. Nature. 2005; 437:281–285.
- Thisse C, Thisse B. High-resolution *in situ* hybridization to whole-mount zebrafish embryos. Nat Protoc. 2008; 3:59–69.