

RETRACTED: Effect of histone deacetylase inhibitors trichostatin A and valproic acid on hair cell regeneration in zebrafish lateral line neuromasts

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In humans, auditory hair cells are not replaced when injured. Thus, cochlear hair cell loss causes progressive and permanent hearing loss. Conversely, non-mammalian vertebrates are capable of regenerating lost sensory hair cells. The zebrafish lateral line has numerous qualities that make it well-suited for studying hair cell development and regeneration. Histone deacetylase (HDAC) activity has been shown to have an important role in regenerative processes in vertebrates, but its function in hair cell regeneration in vivo is not fully understood. Here, we have examined the role of HDAC activity in hair cell regeneration in the zebrafish lateral line. We eliminated lateral line hair cells of 5-day post-fertilization larvae using neomycin and ther treated the larvae with HDAC inhibitors. To assess hair cell regeneration, we used 5-bromo-2-deoxyuridine (BrdU) incorporation in zebrafish larvae to label mitotic cells after hair cell loss. We found that pharmacological inhibition of HDACs using trichostatin A (TSA) or valproic acid (VPA) increased histone acetylation in the regenerated neuromasts following reomycin-induced damage. We also showed that treatment with TSA or VPA decreased the number of supporting cells and regenerated hair cells in response to hair cell damage. Additionally, BrdU immunostaining and western blot analysis showed that TSA or VPA treatment caused a significant decrease in the percentage of S-phase cells and induced p21^{Cip1} and p27^{Kip1} expression, both of which are likely to explain the decrease in the amount of newly regenerated hair cells in treated embryos. Finally, we showed that HDAC inhibitors induced no observable cell death in neuromasts as measured by cleaved caspase-3 immunohistochemistry and western blot analysis. Jaken together, our results demonstrate that HDAC activity has an important role in the regeneration of hair cells in the lateral line.

Keywords, histone deacetylase inhibitors, trichostatin A, valproic acid, zebrafish, hair cell, regeneration

INTRODUCTION

Loss of sensory hair cells (HCs) in the inner ear as a result of aging, ototoxic drugs, or noise is the primary cause of hearing disorders in humans and most other mammals. The majority of HCs do not regenerate in the mammalian cochlea, but a limited amount of HC regeneration can occur in the neonatal mouse cochlea and vestibular sensory epithelium (Forge et al., 1993; Kawamoto et al., 2009; Burns et al., 2012; Chai et al., 2012; Golub et al., 2012; Kelly et al., 2012; Shi et al., 2013; Cox et al., 2014). In contrast, non-mammalian vertebrates such as birds, amphibians, and fish produce HCs throughout their lives and are capable of regenerating HCs after injury (Corwin and Cotanche, 1988; Balak et al., 1990; Harris et al., 2003). Significant progress has been made in elucidating the cellular and molecular mechanisms of sensory HC death and regeneration. Previous studies have reported that newly regenerated HCs in the auditory and vestibular sensory epithelia usually arise through proliferation

and differentiation of progenitor cells during the process of proliferative regeneration (Corwin and Cotanche, 1988; Warchol and Corwin, 1996; Stone and Rubel, 2000). These progenitor cells are believed to be non-sensory supporting cells (SCs) that surround HCs in the sensory epithelia. Upon HC damage, SCs adjacent to the dying HCs reenter the cell cycle and divide asymmetrically during mitosis and give rise to new HCs and SCs. Alternatively, SCs have also been shown to have the ability to spontaneously convert into new HCs through an unusual process called direct transdifferentiation, which is the phenotypic conversion of SCs into HCs without cell cycle reentry (Adler and Raphael, 1996; Adler et al., 1997; Baird et al., 2000; Roberson et al., 2004).

In zebrafish, HCs are found in both the inner ear and the lateral line system. The zebrafish inner ear is responsible for sound detection and balance. Much of the work on HC regeneration in zebrafish has focused on the inner ear. One recent study investigated the effect of aminoglycosides on the regeneration of the

inner ear HCs of adult zebrafish (Uribe et al., 2013) and other studies showed that proliferation is involved in the inner ear and investigated the effect of noise on cell proliferation in the adult zebrafish saccule (Schuck and Smith, 2009; Sun et al., 2011; Liang et al., 2012). Previous study showed that growth hormone (GH) promotes auditory HC regeneration in the zebrafish inner ear following acoustic overexposure, and this effect is mediated through stimulating cell proliferation and suppressing apoptosis (Sun et al., 2011). It is known that the stat3/socs3a pathway is important for HC regeneration in the inner ear of adult zebrafish following acoustic trauma (Liang et al., 2012). On the other hand, zebrafish lateral line is another important mechanosensory system that enables them to detect directional water flow. This helps the fish to avoid obstacles and predators and also facilitates prey detection (Elepfandt, 1988). The lateral line comprises a set of sense organs called neuromasts that are located on the surface of the head (anterior lateral line) and the body (posterior lateral line, PLL) in species-specific patterns (Metcalfe et al., 1985). Mature lateral line neuromasts consist of central HCs and peripheral non-sensory SCs. Because the lateral line HCs in zebrafish are very similar to those in the mammalian inner ear in terms of both morphology and function (Raible and Kruse, 2000; Nicolson, 2005), numerous studies have used the lateral line system to better understand the mechanisms of HC differentiation and regeneration. Moreover, lateral line HCs have been shown to be hypersensitive to ototoxic insults such as aminoglycoside drugs and cisplatin as well as to water-borne copper (Song et al., 1995; Harris et al., 2003; Ton and Parng, 2005; Hernandez et al., 2006; Linbo et al., 2006; Ou et al., 2007; Owens et al., 2009; Giari et al. 2012). In addition, zebrafish lateral lines have retained the ability to rapidly regenerate HCs following ototoxic drugs and provide an attractive model system for investigating HC death, protection, and regeneration and for identifying genes that are crucial for HC development and regeneration. Many studies have used 5-brome 2-deoxyuridine (BrdU) to label mitotic cells and have shown that SCs in the zebrafish lateral line can proliferate and differentiate to renew both HCs and SCs within 72 h after injury. These results suggest that mitosis is the dominant regenerative mechanism in the zebrafish lateral line (Williams and Holder, 2000; Harris et al., 06; Hernandez et al., 2007; 2003; Lopez-Schier and Hudspeth, Ma et al., 2008).

The precise mechanisms responsible for initiating and maintaining-or preventing-HC regeneration in nonmammalian vertebrates are still not completely understood. It has been shown that many candidate molecules and pathways are required for HC regeneration in zebrafish, including Atoh1 (Sweet et al., 2011), Rb (Lin et al., 2013), and the Notch signaling pathway (Ma et al., 2008; Jiang et al., 2014). Recently, chromatin remodeling by post-translational histone modification has been shown to be important for modulating many biological processes including cellular development and regeneration. Acetylation and deacetylation of histones in chromatin are the most common histone modifications and are essential for many developmental processes and are governed by two classes of enzymes, histone acetyltransferases (HATs) and histone deacetylases (HDACs), respectively. Generally, HDAC activity is linked to transcriptional repression by removing the acetyl groups from conserved lysine

residues within the N-terminal histone tails. Conversely, HATs induce transcriptional activation by adding acetyl groups to these residues. The mammalian HDAC family is subdivided into the following four major classes based on their similarity to their yeast counterparts (de Ruijter et al., 2003): Class I (HDAC1, -2, -3, and -8), Class II (HDAC4, -5, -6, -7, -9, and -10), Class III (Sir2 family of NAD⁺-dependent enzymes), and Class IV (HDAC11). Class I HDACs, which are related to yeast RPD3, are widely expressed nuclear enzymes and play an important role in regulating cell survival and proliferation. Class II HDACs, which are related to yeast HDA1, can shuttle between the nucleus and the cytoplasm and have tissue-specific roles. HDAC activity has been shown to be critically involved in the regulation of multiple aspects of developmental processes, including neuronal differentiation (Yamaguchi et al., 2005) and heart (Kim et al., 2012) and liver (Farooq et al., 2008) morphogenesis. HDAC activity has also been shown to be important for modulating tissue regeneration. A recent study on Xenopus tail regeneration showed that HDAC1 is expressed during the early stage of regeneration and that pharmacological blockage of HDACs could inhibit regeneration and induce aberrant expression of genes that are known to be critical for tail regeneration (Tseng et al., 2011). It has been shown that HDAC activity is required for the regeneration of sensory epithelia in the avian utricle (Slattery et al., 2009). Histone deacetylation is a positive regulator of regenerative proliferation, and inhibition of HDACs is sufficient to prevent SCs from entering into the cell cycle.

Previous studies have found that histone acetylation states and HATs are essential regulators of development in zebrafish neuromast HCs (He et al., 2014a). However, the specific functions of HDACs in epigenetic regulation of the regeneration of HCs in the zebrafish lateral line are unknown. In order to determine whether HDACs are directly involved in HC regeneration in the zebrafish lateral line after neomycin-induced cell death, we took advantage of the Tg(Brn3c:mGFP) transgenic zebrafish embryo that expresses GFP in the HCs of the inner ear and lateral line neuromasts (Xiao et al., 2005). Our data indicated that inhibition of HDAC function by trichostatin A (TSA), a Class I and II HDAC inhibitor, affected HC regeneration in zebrafish neuromasts by altering the histone acetylation state. Our BrdU experiments also demonstrated that HDAC inhibitors suppressed proliferation of the progenitor cell population in regenerated neuromasts. In addition, we did not find any significant differences in cell death between control and treated groups over the course of HC regeneration in the lateral line neuromasts. These results suggest that inhibition of HDAC activity is required for HC regeneration in the zebrafish lateral line neuromasts and that HDACs might be potential therapeutic targets for the induction of HC regeneration and SC proliferation.

MATERIALS AND METHODS

ZEBRAFISH MAINTENANCE

Zebrafish embryos were obtained from the natural spawning of wild-type adults and were maintained in our facility according to standard procedures. The Tg(brn3c:mGFP)^{s356t} transgenic line was obtained from the laboratory of Professor Zhengyi Chen, our collaborator at Harvard University. Zebrafish larvae were staged

according to Kimmel et al. (1995) and raised at 28.5°C in Petri dishes. Ages of embryos are described as days post-fertilization (dpf).

NEOMYCIN TREATMENT AND PHARMACOLOGICAL ADMINISTRATION

Neomycin sulfate (Sigma-Aldrich, Inc., St. Louis, MO, USA) was added to a final concentration of 400 µM, and the 5 dpf larvae were incubated for 1 h. This was followed by three rinses in fresh egg water, and the larvae were allowed to recover at 28.5°C. The HDAC inhibitors trichostatin A (TSA, Sigma-Aldrich) and valproic acid (VPA, Sigma-Aldrich) were dissolved either in DMSO (TSA) or ddH2O (VPA) at stock concentrations of 500 µM and 200 mM, respectively, and then diluted to their final concentrations in fresh egg water. Dose-response data were obtained by treating larvae with TSA (0.05 μ M, 0.1 μ M, and 0.2 μ M) or VPA $(50 \,\mu\text{M}, 100 \,\mu\text{M}, \text{and} 150 \,\mu\text{M})$ after neomycin damage. Larvae were anesthetized with 0.02% MS-222 (ethyl 3-aminobenzoate methanesulfonate; Sigma-Aldrich, Inc.) for 5 min before observation and scoring. In all experiments, control groups were maintained in parallel under the same conditions but without inhibitor treatment.

HAIR CELL LABELING AND IMMUNOHISTOCHEMISTRY

For immunohistochemistry analysis, larvae were fixed in 4% paraformaldehyde (PFA) and were permeabilized with PBS containing 0.5% Triton X-100 (PBT-2) for 30 min followed by incubation in blocking solution for 1 h. The following antibodies were used as primary antibodies: anti-myosin VI (1:200 dilution; Proteus BioSciences, Ramona, CA); anti-HC soma-1 antigen (HCS1, 1:200; Developmental Studies Hybridoma Bank); anti-Sox2 (1:200; Abcam, Cambridge, UK); anti-acetylated histone H3 (1:500; Upstate Biotechnologies Inc., Lake Placid, NY, USA); anti-acetylated histone H4 (1:500; Upstate Biotechnologies Inc.); and anti-cleaved caspase-3 (1:500; Cell Signaling Technology Inc., Danvers, MA, USA). The embryos were washed three times with PBS and incubated with secondary antibodies to detect primary antibodies. Nuclei were labeled with 4,6-diamidino-2phenylindole (DAPI; Invitrogen, Carlsbad, CA, USA) for 20 min at room temperature

FM1-43FX LABELING

For staining of functional HCs within the neuromasts, the vital dye FM1-43FX (Invitrogen, r-35355)—which enters mature HCs through mechanotransduction-dependent activity—was applied at a concentration of 3 μ M to live 5 dpf larvae for 45 s in the dark. After quickly rinsing three times with fresh egg 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.

CELL PROLIFERATION ASSAY

To label proliferating cells, larvae were incubated in egg water containing 10 mM BrdU (Sigma) solution for 24 h or 48 h at 28.5°C starting at 1 h after neomycin treatment. Larvae were then fixed with 4% PFA overnight at 4°C. BrdU incorporation was detected by immunocytochemistry. The fixed larvae were washed three times in 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 Biotechnology, Inc., CA, USA) overnight at 4°C. The next day, the 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. Specimens were examined by Leica confocal fluorescence microscope (TCS SP5; Leica, Wetzlar, Germany).

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 bicinchoninic acid (BCA) protein assay kit, and samples were separated by 12% SDS-PAGE. After electrophoresis, the proteins were transferred onto PVDF membranes (Immobilon-P; Millipore, Bedford, MA, USA) that were blocked with 5% non-fat dried milk in TBST (50 mM Tris-HCl (pH 7.4), 150 mM NaCl, and 0.1% Tween-20) for 1h at room temperature. After washing, primary antibodies were added to blocking buffer overnight at 4°C. The following antibodies were used as primary antibodies: antiacetylated histone H3 (1:1000 dilution); anti-acetylated histone H4 (1:1000); anti-cleaved caspase-3 (1:1000); anti-p21^{Cip1} C-19 (1:1000; Santa Cruz Biotechnology, Inc., CA, USA); antip27^{Kip1} (1:500; Santa Cruz Biotechnology); anti-p53 (1:1000; Santa Cruz Biotechnology); anti-HDAC1 (1:1000; Cell Signaling Technology Inc., Danvers, MA, USA); anti-HDAC2 (1:1000; CST); anti-HDAC3 (1:1000; CST); anti-HDAC4 (1:1000; CST); nti-HDAC5 (1:1000; CST); anti-HDAC6 (1:1000; CST); anti-HDAC8 (1:1000; Abcam, Cambridge, UK); and anti-β-Actin (1:2000).

QUANTITATIVE ANALYSIS

For quantitative real time PCR (qPCR), total RNA was obtained with the AllPrep DNA/RNA/Protein Mini Kit (QIAGEN, Hilden, Germany) according to the manufacturer's instructions. cDNA was synthesized from total RNA using first-strand cDNA synthesis kit (Promega, USA). qPCR was performed on an ABI StepOneTM Real-Time PCR System (Applied Biosystems) with GoTaq qPCR Master Mix kit (Promega, USA). Glyceraldehyde-3phosphate dehydrogenase (GAPDH) was used as reference genes. Gene expression levels were normalized to GAPDH and calculated as $2^{-\Delta\Delta Ct}$ according to the manufacturer's instructions (Applied Biosystems). All primers were listed in Supplemental Table.

CELL COUNTS AND STATISTICAL ANALYSIS

Cells in the first five posterior lateral line (PLL) neuromasts (L1–L5) were counted. All statistical analysis was performed with SPSS (version 13.0 for Windows) and SigmaPlot (version 12.0 for Windows). Prior to analysis, all data were first examined for normality and homogeneity of variances by the Shapiro-Wilk test and Levene's test, respectively. Data were analyzed using either *t*-tests or analysis of variance (ANOVA) with multiple comparisons. All data are presented as the mean ± s.e.m. *p* < 0.05 was considered statistically significant and *p* < 0.001 was considered highly significant.

RESULTS

THE EFFECT OF HDAC INHIBITORS ON THE NUMBER OF REGENERATED HCs AND SCs IN THE ZEBRAFISH LATERAL LINE

To induce regeneration, we treated 5 dpf zebrafish larvae with 400 µM neomycin for 1 h to kill mature lateral line HCs. To facilitate visualization of regeneration, we used Tg(brn3c:mGFP) zebrafish that expressed GFP in differentiated HCs under the control of the pou4f3 promoter. After neomycin treatment, most of the HCs were damaged and lost but regeneration occurred rapidly in the fish over the following 2 days (Figures 1A1,B1; Supplemental Figure 1). To investigate the role of HDAC inhibition in HC regeneration, neomycin-treated larvae were placed in 6-well plates and exposed to TSA for 24 h and 48 h recovery periods. We found that larvae treated with 0.1 µM TSA for both 24 h and 48 h (Figures 1A2, B2) had fewer regenerated HCs relative to DMSO vehicle controls (Figures 1A1,B1). To quantify changes in the numbers of regenerated HCs after neomycin-induced damage, specific labeling of regeneration of HCs was observed and quantified using Myosin VI immunostaining. The numbers of HCs in neuromasts L1-L5 were counted in 6-13 fish at each time period. In the 24 h group, we found an average of five Myosin VI⁺ HCs in neuromasts of the vehicle control (Figure 1C1), but at the most only three Myosin VI⁺ HCs were seen in the TSA-treated neuromasts (Figure 1C2). At 48 h post-treatment, we found an average of 10 Myosin VI⁺ HCs in the DMSO controls (Figure 1D1) compared to at most five in the TSA-treated neuromasts (Figure 1D2).

To confirm that the reduction of regenerated HCs in the presence of TSA was related to an inhibitory effect on HDACs, we performed the same experiment with another HDAC inhibitor, VPA. Compared with controls, the group treated with 100 µM VPA for 24 h showed a decrease in the number of HCs (**Figures 1A3,C3**), and there was an even greater decrease of HCs per neuromast after exposure to 100 µM VPA for 48 h (**Figures 1B3,D3**). To further assess the concentration effect, we treated neomycindamaged larvae with TSA or VPA at varying concentrations ranging from 0.05 µM to 0.2µM or 50 µM to 150 µM, respectively. Both TSA and VPA treatment for 24 h and 48 h reduced the number of regenerated HCs in a dose-dependent manner (**Figures 1E,F**).

To test the functionality of the regenerated HCs, larvae were stained with the vital dye FM1-43FX, which is a marker of functional mechanotransduction channels in HCs (Seiler and Nicolson, 1999). We imaged and quantified the FM1-43FX⁺ HCs in neuromasts in control and inhibitor-treated zebrafish larvae after neomycin exposure at two time points during the recovery period. Again, we found that 0.1 μ M TSA or 100 μ M VPA treatment induced a significant decrease in FM1-43FX⁺ cell numbers when compared to controls at 24 h and 48 h post-treatment (**Figure 2**). Therefore, we conclude that the regeneration process in larvae is severely impaired in the presence of HDAC inhibitors.

We next evaluated the effect of HDAC inhibitor treatment on the number of SCs in the neuromasts after neomycin exposure. In order to visualize the SCs, we identified them with an antibody against Sox2, which is highly expressed in the SCs of the neuromasts (**Figures 3A,B**). We counted the Sox2-labeled cells in the neuromasts after treatment with HDAC inhibitors at 24 h and 48 h post-treatment and observed a significant reduction in the number of stained SCs (**Figures 3C,D**).

IMPACT OF HDAC INHIBITORS ON NEUROMAST CELL PROLIFERATION

Previous studies have reported that most regenerated HCs arise from SC proliferation within 72h after neomycin treatment (Harris et al., 2003; Ma et al., 2008), so we next determined whether inhibition of HDAC affects cell proliferation during this regeneration process. After neomycin injury, zebrafish larvae were continuously incubated in fresh egg water containing 10 mM BrdU and either TSA or VPA for 24 h and 48 h. Cellular proliferation was determined by counting the number of BrdU⁺ cells in the L1-L5 neuromasts. As Figure 4 illustrated, compared with control, 24 h groups treated with 0.1 µM TSA or 100 µM VPA showed significant decrease in the number of BrdU⁺ cells per neuromast (control larvae harbor 15.9 \pm 0.5 BrdU⁺ cells, TSA treated-larvae harbor 6.9 \pm 0.2, *p* < 0.001; control larvae harbor 15.5 ± 0.5 BrdU⁺ cells, VPA treated-larvae harbor 6.3 ± 0.3 , p <0.001). In 48 h groups, there was an intensive decrease of $BrdU^+$ cells per neuromast after the exposure to TSA or VPA (control larvae harbor 24.8 \pm 0.7, TSA treated-larvae harbor 9.6 \pm 0.5, p < 0.001; control larvae harbor 25.3 \pm 0.8, VPA treated-larvae harbor 10.1 \pm 0.3, p < 0.001). These data suggest that TSA and VPA significantly inhibit the proliferation of neuromast cells.

To distinguish the newly regenerated HCs from cell proliferation, we double-stained the larvae with an anti-BrdU antibody (green in Figures 4A,B) and with an anti-Myosin VI antibody (red in Figures 4A,B) at 24 h and 48 h after neomycin damge. Our analysis showed that fish treated with TSA or VPA for 24h had fewer HCs in the neuromasts and that virtually none of them were co-labeled with the anti-BrdU antibody. However, in vehicle-only control larvae we found on average wo double-stained cells per neuromast indicating that they arose from proliferating cells. Among the 48 h groups, a significant increase in BrdU⁺ and Myosin VI⁺ cells was observed in control fish and there was a significant decrease in double-stained cells per neuromast after exposure to TSA or VPA (Figures 4A-D). Furthermore, on comparing the ratio of BrdU⁺ and Myosin VI⁺ double-labeled cells to the total number of Myosin VI⁺ cells in the different groups, fewer BrdU⁺ HCs were found in the TSA- or VPA- treated groups compared with the vehicle control groups at 24 h and 48 h post-treatment (Figures 4E,F). These findings suggest that HDAC inhibitors decreased the proportion of cells in S-phase.

Because the number of SCs in HDAC inhibitor-treated neuromasts is different from that of control neuromasts, we next examined the rates of SC division among different groups after neomycin damage at 24 h and 48 h by BrdU and Sox2 immuno-labeling. Similar to the results described above, at 24 h after neomycin treatment we found fewer double-stained cells in the presence of TSA or VPA compared with the vehicle controls (**Figure 5A**). Furthermore, there was a significant difference between the ratio of BrdU⁺ SCs (**Figures 5C,D**). By 48 h after neomycin insult, the control larvae and the HDAC inhibitor-treated larvae showed a striking difference in the number of double-stained cells (**Figure 5B**). The ratio of BrdU⁺ SCs in the treated neuromasts were drastically reduced, and this most likely



Tg(pou4f3:gap43-GFP) zebrafish with $400\,\mu M$ neomycin for 1 h and then treated them for 24 h or 48 h with TSA or VPA and subsequently imaged GFP+ HCs (green). Nuclei are stained with DAPI and scale

bars = $10\,\mu$ m. (C and D) Lateral line HCs are stained with Myosin VI. (E and F) The average number of Myosin VI⁺ cells per neuromast (NM) in larvae treated with or without HDAC inhibitor for 24 h or 48 h after neomycin damage. Bars are mean \pm s.e.m. and n = total number of embryos. *p < 0.05.



explains the reduction in the number of newly regenerated HCs in the inhibitor-treated embryos (Figures 5C,D). Taken together, these results show that HDAC inhibition has a significant negative impact on proliferation in the regenerating neuromast.

INDUCTION OF P21CIP1 AND P27KIP1 BY HDAC INHIBITORS

HDAC plays a central role in the regulation of cell cycles (Xiao et al., 2014). We investigated the effect of HDACs on the expression of cell cycle-regulated proteins during the recovery period. The results revealed that HDAC inhibition significantly decreased the expressions of cyclin B1, cyclin D1, cyclin D3, cyclin E1, cyclin-dependent kinase (CDK) 2, and CDC2, whereas increased the expression of p21 and p27 (Supplemental Figure 2). These results suggest that HDAC inhibitors might inhibit the proliferation of neuromast cells by inducing p21 and p27, leading to G1 phase cell cycle arrest. To further confirm the effect of HDAC inhibition on p21^{Cip1} and p27^{Kip1} expression during the recovery period, embryos were treated with TSA or VPA after neomycin-induced injury and protein expression was detected using antibodies specific to p21^{Cip1} and p27^{Kip1}. The

levels of $p21^{Cip1}$ and $p27^{Kip1}$ in the inhibitor-treated groups increased compared to controls (**Figure 6**). Because it is wellknown that the $p21^{Cip1}$ gene is physiologically induced by p53 (Sherr, 1994), we next examined the expression of p53 by western blotting. The expression of p53 was not significantly altered in fish treated with TSA or VPA compared to controls, and this suggests that the induction of $p21^{Cip1}$ expression in response to HDAC inhibitors might be mediated by a p53-independent pathway.

IMPACT OF HDAC INHIBITORS ON APOPTOSIS

Previous studies have suggested that HDAC inhibitors can induce cell cycle arrest and apoptosis in many different cell types (Riester et al., 2007). Because the reduced number of regenerated HCs might also be caused by increased cell death, we further monitored cell death using an antibody against cleaved caspase-3 over the time course of HC regeneration. We did not find any significant differences in cell death in the lateral line neuromasts between control and inhibitor-treated groups (**Figure 7**), and these results suggest that the inhibition of proliferation in HDAC



inhibitor-treated neuromasts was mainly due to cell cycle arrest and not apoptosis.

HDAC INHIBITION INCREASES HISTONE ACETYLATION IN NEUROMASTS

To confirm that the deficiency in HC regeneration was, indeed, caused by the inhibition of HDAC activity, we examined the level of histone acetylation in zebrafish after treatment with $0.1 \,\mu M$ TSA or 100 µM VPA for 48 h. Western blot analysis showed that the levels of acetylated histone H3 and H4 in controls were low, but incubation with TSA and VPA resulted in the accumulation of acetylated histones H3 and H4 (Figure 8A). To more accurately localize acetylated H3 and H4 expression to the specific cell types in the neuromast, we performed immunostaining for the HC marker HCS-1 (green). In controls, both acetylated H3⁺ (red) and acetylated H4⁺ (red) cells are expressed outside the central HCs and there was little or no overlap of signals (Figure 8B). However, the acetylated H3 and H4 signals were elevated in inhibitor-treated neuromast cells, particularly in the SCs, and more overlap was seen between the signals (Figure 8B). These results provide evidence that HDAC inhibitor-mediated histone acetylation in neuromast cells might indeed be responsible for the decreased HC production. We next investigated which members of HDAC family are affected by HDAC inhibitors treatment during the recovery period. We found a significant downregulation of HDAC1, HDAC3, and HDAC4 protein levels in zebrafish treated with the VPA or TSA. Reduction of HDAC2 protein levels is also found after 48 h HDAC inhibitor treatment. Furthermore, VPA and TSA treatment do not cause a reduction in protein levels of HDACs 5, 6, and 8 (Supplemental Figure 3).

DISCUSSION

HDAC activity has been demonstrated to play an important role in the regulation of diverse physiological processes in the cell—including cell differentiation, apoptosis, migration, proliferation, and survival—via the formation of complexes with various transcription factors (for example, Sp1, Sp3, p53, and nuclear factor-kappa B) (Haberland et al., 2009). We have previously shown that HDAC activity is necessary for the control of proliferation and migration of the PLL primordium and the concomitant deposition of the neuromasts during the early stages of PLL development in zebrafish (He et al., 2014b). We have



also demonstrated that HDAC activity is involved in the development of HCs and SCs in zebrafish lateral line neuromasts (He et al., 2014a). Our previous study revealed that HDAC activity is involved in lateral line development and might have a role in neuromast formation by altering cell proliferation through the expression of cell cycle regulatory proteins. However, there is no existing report concerning the role of the HDACs in HC regeneration. In this study, we have demonstrated the role of HDAC in HC regeneration in the zebrafish lateral line. We observed that TSA and VPA administration decreased the numbers of newly regenerated HCs and SCs in the neomycin-damaged neuromast. We also showed that TSA and VPA treatment dramatically reduced cell proliferation in neuromasts as detected by BrdU immunostaining, and this activity is believed to be the primary mechanism behind the decreased regeneration of HCs in the zebrafish lateral line. It has been shown that the anti-proliferative effect of HDAC



 $100\,\mu\text{M}$ VPA for 48 h, protein extracts were prepared and subjected to

*p < 0.05.



inhibitors is responsible, at least in part, for the induction of G1 cell cycle arrest of various types of cells.

Cell cycle regulation is of pivotal importance for the control of growth and development, and deregulation of the cell cycle has been shown to be associated with proliferation defects and carcinogenesis (Massague, 2004). The coordinated interactions between Cyclins and cyclin-dependent kinases (CDKs) play key roles in promoting the progression through the different phases of the cell cycle. CDK activity is negatively regulated by two types of CDK inhibitors (CKIs), the CIP/KIP subfamily (p21^{Cip1}, p27^{Kip1}, and p57^{Kip2}) and the INK4 subfamily (p15^{lnk4b}, p16^{lnk4a}, p18^{Ink4c}, and p19^{Ink4d}) (Sherr and Roberts 1995 CKIs can inhibit a broad range of Cyclin/CDK complexes and act as key regulators of the timing of cell cycle arrest. Previous wor **k** has shown that several CKI family members are expressed in inner ear HCs and SCs during embryogenesis and that the expression of these regulators becomes more restricted with maturation. For instance, p27Kip1 remains strongly expressed in SCs during adulthood, but becomes down-regulated in mature HCs suggesting that it plays a particularly important role in maintaining differentiated SCs in a non-proliferative status (Chen and Segil, 1999). Another CIP/KIP family member, p21^{Cip1}, and the INK4 family member p19^{Ink4d} help to maintain the postmitotic state of HCs. Mutations of both p21^{Cip1} and p19^{Ink4d} in mice lead to HC death and reentry into the cell cycle, but these differentiated HCs quickly die by apoptosis (Laine et al., 2007). Together these studies reinforce the fact that direct manipulation of cell cycle regulatory genes has not yet been successfully used for the production of functional auditory sensory epithelium in mammals, and they suggest that it is important to understand the detailed mechanisms that activate cell cycle reentry in many regenerating systems. It is possible that manipulating signaling events through epigenetic regulation will be a more promising therapeutic avenue for stimulating mammalian HC regeneration.

A great number of studies have demonstrated that expression of HDACs may contribute to the development and regeneration

 In the present study, we idenof many tissues (Tsenget al., tified that protein levels of HDAC1-4 were downregulated in TSA or VPA treated zebrafish, whereas the expression of other HDAC family members (HDAC5, HDAC6, and HDAC8) were not significantly changed after 48 h TSA or VPA treatment. These results suggest that several family members of class I and II HDACs might play key roles in HC regeneration in zebrafish. Emerging vidence is increasingly showing that the intrinsic relationship between cell cycle progression and HDAC activity is essential for embryogenesis. The CKI p21^{Cip1}, which mediates cell cycle rrest, apoptosis, and differentiation, was previously shown to e a crucial target of the transcriptional corepressor HDAC1 in mouse ES cells (Lagger et al., 2002) and human tumor cells (Lagger et al., 2003; Gui et al., 2004). Lagger et al. (2002) demonstrated that HDAC1 acts as a positive regulator of proliferation in mouse ES cells and mouse embryonic fibroblasts by repressing the expression of p21^{Cip1}. Targeted deletion of HDAC1 in mice results in embryonic lethality before embryonic day 10.5 due to severe proliferation defects and suggests a key role for HDAC1 in regulating cell proliferation. Reduced proliferation rates in HDAC1-deficient embryos and embryonic stem cells (ES cells) are also correlated with elevated levels of p21^{Cip1} and p27^{Kip1} (Lagger et al., 2002). In contrast, ablation of p21^{Cip1} rescues the proliferation phenotype of HDAC1-null ES cells but not the embryonic lethality of HDAC1-deficient mice (Zupkovitz et al., 2010). Recently, the promoter regions of the p21 gene were shown to be common targets of HDAC1 and HDAC2 indicating that HDAC1 and HDAC2 directly regulate p21 gene expression to control G1 to S-phase progression (Lin et al., 2008). Combined deletion of HDAC1 and HDAC2 results in G1 cell cycle arrest and is accompanied by up-regulation of p21 in primary and oncogenic-transformed fibroblasts (Wilting et al., 2010). HDAC inhibitors such as suberoylanilide hydroxamic acid (SAHA) and TSA induce differentiation and apoptosis and alter the expression of p21 in transformed cells. Richon et al. (2000) have shown that SAHA induces the activation of the p21 gene via acetylation



regeneration. (A) Western blot analysis of acetylated H3 (Ace H3) and acetylated H4 (Ace H4) protein extracts from control and 0.1 μ M TSA- or 100 μ M VPA-treated larvae at 48 h after neomycin damage. β-Actin was

included as the control. Mean \pm s.e.m. for three experimental replicates. *p<0.05. **p<0.001. **(B)** Acetylation of histone H3 and H4 were stained (red) by Ace H3 and Ace H4 antibody. HCs were labeled with HCS-1 (green) antibody and nuclei were stained with DAPI (blue). Scale bar = 10 μ m. of histones H3 and H4 that are associated with the promoter and coding regions of the p21 gene. Yamaguchi et al. (2010) also demonstrated that HDAC1 and HDAC2 directly regulate the G1 to S-phase transition of the cell cycle by repressing the expression of p21 and p57. These findings indicate that multiple HDACs regulate cell cycle progression through the G1/S-phase checkpoint by repressing CKI expression, in particular through transcriptional repression of p21 in different cell types. In agreement with this, our data show that HDAC inhibitors significantly inhibited proliferation and upregulated the expression of the CKIs p21^{Cip1} and p27Kip1. This was associated with increased H3 and H4 acetylation and decreased HDAC (1, 2, 3, and 4) expression suggesting that p21^{Cip1} and p27^{Kip1} might be involved in mediating the anti-proliferative effects of HDAC inhibitors on neuromast cells. However, the underlying mechanism of HDAC in the regulation of regeneration in HCs remains unidentified. So it will be interesting and valuable to determine the possible mechanisms of HDACs, especially HDACs1-4, in HC regeneration in future studies.

Expression of the p21^{Cip1} gene is tightly controlled by a variety of factors, including the MAPK signaling pathway (Simboeck et al., 2010) and the tumor suppressor p53-which is important in the DNA damage checkpoint (Sherr, 1994). p21^{Cip1} expression has been shown to be regulated in a p53-dependent and p53-independent manner (Michieli et al., 1994; Gartel and Tyner, 1999). We next examined the expression of p53 protein. In our experiments, TSA or VPA treatment does not activate p53 but can increase p21 protein, suggesting that the effect of TSA and VPA on p21 expression might be independent of p53 status. This is consistent with the previously reported observations that HDAC inhibitors can activate p21 transcription through its promoter Sp1 sites in a p53-independent manner (Nakano et al., 1997; Sowa et al., 1997; Nuang 2000). However, it remains a possibility that both p53-dependen and p53-independent signaling pathways are involved in our study. To confirm that TSA- and VPA-mediated induction of p21 is independent of p53, we plan to investigate the stability of p53 protein in HDAC inhibitor-treated fish because it is well-accepted that p53 function is tightly controlled through stability of the protein. The stability of p53 protein is regulated predominantly by the oncoprotein Mdm2 that acts as a repressive regulator of p53 (Oliner et al., 1993), and the analysis of Mdm2 activity will also be investigated in future work.

In conclusion, our results indicate that inhibition of HDAC activity with TSA and VPA downregulates several family members of class I and II HDACs and significantly decreases the numbers of SCs and regenerated HCs in zebrafish lateral line neuromasts during the regeneration process. Furthermore, administration of HDAC inhibitors strongly inhibit cell proliferation in response to HC death. HDAC function is important for regulating the expression of cell cycle regulators as an critical component during HC regeneration, and this strengthens and extends the importance of HDAC activity that was previously shown to be important during the development of the neuromasts in zebrafish (He et al., 2014a). Our findings suggest that HDAC is a potential target for therapies aimed at promoting HC regeneration.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: http://www.frontiersin.org/journal/10.3389/fncel.2014. 00382/abstract

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