Figure S1. The genome of zebrafish encodes three Mi-2 orthologues.

Schematic representation of zebrafish Mi-2 proteins, Chd4a (NP_001038323.1), Chd4b (XP_685699.4) and Chd3 (XP_696641.3). Conserved domains determined with the Conserved Domain Database (NCBI) are shown.
Figure S2. One of the three Mi-2 homologues, chd4a, is induced during regeneration in the adult caudal fin.

*In situ* hybridization with chd4a, chd4b, and chd3 mRNA antisense probes on whole regenerating adult caudal fins at 3 dpa. No signals were detected with the corresponding sense RNA probes. Arrowheads indicate the amputation plane. Scale bar: 100 μm.
Figure S3. *chd4a* is expressed during regenerative outgrowth in adult caudal fin.

*In situ* hybridization with *chd4a* mRNA antisense probe on cryosections at 8 hpa and 24 hpa and on whole fins at 2 dpa, 3 dpa, 7 dpa and in uninjured fin. *chd4a* expression was first observed at 2 dpa with a robust signal at 3 dpa. Arrowheads indicate the amputation plane. Scale bars: 100 μm.
Figure S4. *chd4a, chd4b* and *chd3* are expressed in developing zebrafish embryos.

Whole-mount *in situ* hybridization with *chd4a, chd4b* and *chd3* RNA antisense probes in zebrafish embryos at 1-2 cell stage, 6 hours post-fertilization (hpf), 12 hpf and 24 hpf. The expression pattern of *chd4a, chd4b* and *chd3* in embryos suggest that they might also play a role during embryonic development. Scale bars: 50 μm.
Figure S5. The splice blocking antisense chd4a MOSP efficiently impairs the splicing of chd4a transcript.

The splice-blocking chd4a MOSP was predicted to target the exon 8 - intron 8 boundary leading to an aberrant splicing of chd4a transcript. (A) The level of correctly spliced chd4a mRNA was determined by quantitative RT-PCR in morphant embryos. The control MO or the splice blocking antisense chd4a MOSP was injected into 1-2 cell stage embryos and RNA from 24 hpi embryos was extracted. The level of correctly spliced chd4a mRNA was significantly reduced in chd4a morphant embryos compared to control embryos, indicating that the splicing of chd4a transcript was impaired. Error bars represent the s.e.m. * P<0.001 (B) Unprocessed chd4a transcript is schematically represented. The primers prCP46, prCP47 and prCP82 used to amplified segments of the incorrectly spliced chd4a RNA are indicated. (C) Segments of the intron 8 of chd4a transcript were amplified by PCR with the indicated primers from cDNA synthetized from RNA of control embryos or chd4a MOSP morphant embryos.
Figure S6. Zebrafish Hdac1 and human HDAC1/HDAC2 are highly conserved

Amino acid sequence alignment of zebrafish Hdac1 (NP_775343), human HDAC1 (NP_004955) and human HDAC2 (NP_001518). The catalytic domain (highlighted in red) of the three proteins is highly conserved (93% identity). The active center of HDAC1/2 contains a zinc ion, that binds to three conserved residues (indicated in blue).
Figure S7. MGCD0103 treatment does not affect the general health of zebrafish.

(A) Schematic representation of fin regeneration progression and MGCD0103 treatment. Fish were treated with DMSO or MGCD0103 starting from the time of amputation. (B) Images of fish before and 10 days after DMSO or MGCD0103 treatment. The general health of fish treated with MGCD0103 for 10 days was not obviously altered compared with fish incubated with DMSO-containing water.
**Figure S8. Hdac1 inhibition after blastema formation is sufficient to impair regenerative outgrowth.**

(A) Schematic representation of fin regeneration progression and MGCD0103 treatment. Fish were treated with DMSO or MGCD0103 starting from 3 dpa. (B) Whole caudal fins at 3 dpa and at 7 dpa treated with DMSO or 5 μM MGCD0103 starting from 3 dpa. Arrowheads indicate the amputation plane. Scale bar: 100 μm.
Figure S9. The effects of MGCD0103 treatment are not reversible.

(A) Schematic representation of fin regeneration progression and MGCD0103 treatment. Fish were treated with DMSO or MGCD0103 for 10 days starting from the time of amputation and then fish were transferred into normal fish water for 10 additional days. (B) Corresponding images of whole caudal fins just after amputation (0 hpa), at 10 dpa or at 20 dpa. Effects of MGCD0103 treatment were not reversible. In some cases, a few rays were able to resume the regeneration process. Arrowheads indicate the amputation plane.
Figure S10. Hdac1 inhibition and chd4a knockdown do not result in the activation of the apoptosis marker caspase-3.

(A-F) Longitudinal sections of fin regenerates stained with active caspase-3 antibody (green) and with DAPI (blue). Only a few cells positive for active caspase-3 are present in control fins (A,C). Fins treated with MGCD0103 (B) or injected with chd4a MO (D) do not display any increase of apoptosis at 4 dpa. (E-F) As a positive control, the activation of caspase-3 was assessed in fin regenerates at 1 dpa treated with the chemical inhibitor of Igf1 receptor NVP_AEW541. Inhibition of IGF signaling blocks fin regeneration and results in the increase of cell apoptosis in the wound epidermis (Chablais et al., 2010). Dashed lines indicate the amputation plane. Scale bars: 100 μm.
**Figure S11.** *wnt5b, lef1* and *msxb* are expressed in *chd4a* MO-injected and MGCD0103-treated fins.

(A-B) *In situ* hybridization with *wnt5b, lef1* and *msxb* mRNA antisense probes in *chd4a* MO-injected (A) and in MGCD0103-treated (B) fin regenerates at 4 dpa. Scale bar: 100 μm.
Figure S12. *chd4a* knockdown does not affect Tenascin C expression.

(A-B) Longitudinal sections of fin regenerates at 4 dpa injected with control (A) or *chd4a* (B) MOs stained with Tenascin C antibody (red) and with DAPI (blue). Mesenchymal remodeling was not altered in *chd4a* MO-injected fins. Dashed lines indicate the amputation plane. Scale bars: 100 μm.
Figure S13. *chd4a* knockdown reduces osteoblast proliferation.

(A-B) Longitudinal sections of fin regenerates at 4 dpa injected with control (A) or *chd4a* (B) MOs triply stained with BrdU (green), ZNS5 antibody (red) and DAPI (blue). Dashed lines indicate the amputation plane. Scale bar: 100 μm. (C) Percentage of ZNS5-positive cells that were also positive for BrdU relative to the total number of ZNS5-positive cells in fin regenerates at 4 dpa injected with control or *chd4a* MOs. Error bars represent the s.e.m. n = 17. * P<0.01.
Figure S14. Hdac1 inhibition after blastema formation is sufficient to block reactivation of osteocalcin expression.

(A) Caudal fins of osteocalcin:GFP transgenic fish at 3 dpa and 7 dpa treated with DMSO or MGCD0103 starting at 3 dpa. osteocalcin:GFP expression was not reactivated in fin regenerates treated with MGCD0103 after blastema has been formed. (B) Uninjured caudal fins of osteocalcin:GFP transgenic fish before and 10 days after MGCD0103 treatment. MGCD0103 treatment does not alter osteocalcin:GFP expression in uninjured fins. Constant exposure times were used. Scale bars: 100 μm.