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The *Xenopus* tadpole: an *in vivo* model to screen drugs favoring remyelination

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Abstract

Background: In MS development of screening tools for remyelination-promoting molecules is timely.

Objective: A *Xenopus* transgenic line allowing conditional ablation of myelinating oligodendrocytes has been adapted for *in vivo* screening of remyelination-favoring molecules.

Methods: In this transgenic, the green fluorescent protein reporter is fused to *E. coli* nitroreductase and expressed specifically in myelinating oligodendrocytes. Nitroreductase converts the innocuous pro-drug metronidazole to a cytotoxin. Spontaneous remyelination occurs after metronidazole-induced demyelinating responses. As tadpoles are transparent, these events can be monitored *in vivo* and quantified. At the end of metronidazole-induced demyelination, tadpoles were screened in water containing the compounds tested. After 72h remyelination was assayed by counting numbers of oligodendrocytes per optic nerve.

Results: Among a battery of molecules tested, siponimod, a dual agonist of sphingosine-1-phosphate receptor 1 and 5, was among the most efficient favoring remyelination. Crispr/cas9 gene editing showed that the promyelinating effect of siponimod involves the sphingosine-1-phosphate receptor 5.

Conclusion: This *Xenopus* transgenic line constitutes a simple *in vivo* screening platform for myelin repair therapeutics. We validated several known pro-myelinating compounds, and demonstrated that the strong remyelinating efficacy of siponimod implicates the sphingosine-1-phosphate receptor 5.

Introduction

Oligodendrocytes are the myelin-forming cells of the CNS. During embryonic development, cells of the oligodendrocyte lineage emerge from the ventricular zone. Successive developmental stages transform the neural stem cells into proliferating oligodendrocyte progenitors, which migrate into the parenchyma as oligodendrocyte precursor cells (OPCs) before differentiating, successively into a pre-oligodendrocyte, then a mature postmitotic oligodendrocyte and finally a myelin-forming oligodendrocyte. These developmental stages, which are associated with different capacities of proliferation, migration, differentiation, survival, maturation, and finally myelin deposition, are cell autonomous, but also depend on a variety of extrinsic factors.

In multiple sclerosis (MS) remyelination often fails despite OPC presence around demyelinating areas.¹ Repair of demyelinated lesions could enhance successful re-ensheathing of naked axons by oligodendrocyte processes. Existing MS models (mostly rodent and primates) are not ideal for monitoring *in vivo* demyelination and remyelination, hence the need to develop simpler strategies for larger scale screening of compounds favoring myelin repair.² We developed a transgenic *Xenopus laevis* permitting live imaging of demyelination and remyelination by conditionally triggering apoptotic ablation of myelin forming oligodendrocytes.³ This line, *MBP-GFP-NTR*, expresses green fluorescent protein (GFP) fused to *E. coli* nitroreductase (NTR) under control of the mouse 1.9kb myelin basic protein (MBP) regulatory sequence. In the mouse, this portion of the proximal upstream region of *MBP* gene drives reporter expression specifically in myelinating oligodendrocytes.⁴ Like the mouse, the transgenic *MBP-GFP-NTR X. laevis* expresses the *GFP-NTR* transgene specifically in myelin-forming oligodendrocytes.

The NTR enzyme converts the nitro radical of prodrugs, including metronidazole, to a highly cytotoxic hydroxylamine derivative. In metronidazole-exposed transgenic *MBP-GFP-NTR* tadpoles demyelination is restricted to the CNS, without axonal damage.³ The extent of demyelination depends on the concentration and duration of metronidazole exposure.

Spontaneous remyelination occurs and is nearly complete 8 days after cessation of metronidazole. As embryos are transparent, demyelination and remyelination can be monitored and quantified *in vivo* under a fluorescent microscope.³ We used this model to screen molecules favoring remyelination. Among the battery of molecules tested, one of the most potent to accelerate remyelination was siponimod, which was recently evaluated in a Phase III trial in secondary-progressive MS.⁵

Siponimod is a dual agonist of sphingosine-1-phosphate receptors, S1PR1 and S1PR5. Sphingosine-1-phosphate (S1P) has been identified as the endogenous ligand of the five known S1P receptors (S1PR1, S1PR2, S1PR3, S1PR4 and S1PR5) a family of G-protein coupled receptors.⁶ In the rodent CNS, the S1PR5 receptor is expressed only by cells of the oligodendroglial lineage and its activation is important in the oligodendrocyte maturation. It was therefore suggested that S1PR5 may play a key role in myelin sheath repair.⁷ Here we questioned whether the remyelinating effect of siponimod implicated S1PR5. We show that, in tadpoles with Crispr/Cas 9 gene deleted S1PR5, siponimod no longer promotes remyelination. Another S1PR5 agonist (UC-42-WP04) also favored remyelination, while an S1PR1 agonist (AUY954) had no effect. The data provide strong evidence that siponimod-stimulated remyelination implicates S1PR5 activation and illustrate that *MBP-GFP-NTR Xenopus* is an efficient and simple experimental model to screen molecules favoring remyelination.

Materials and methods

Animals

Xenopus tadpoles were staged according to Nieuwkoop and Faber.⁸ *X. laevis* is an allotetraploid composed of two separate diploid genomes ($2n=36$). *X. laevis* has nine pairs of chromosomes named Xla1S and Xla1L through Xla9S and Xla9L where S (short) and L (long) refer to their relative lengths. Tadpoles of either sex were anesthetized in 0.05% MS-222 (Ethyl-3-aminobenzoate methanesulfonate; Sigma-Aldrich) before quantification of GFP+ cells, and returned to normal water to recover. Before brain and spinal cord dissection, tadpoles were euthanized in 0.5% MS-222. Animal care was in accordance with institutional and national guidelines. All animal procedures conformed to the European Community Council 1986 directive (86/609/EEC) as modified in 2010 (2010/603/UE) and have been approved by the ethical committee of the French Ministry of Higher Education and Research. Transgenic tadpoles were treated between stages NF 50 and 55, corresponding to pre-metamorphosis, a stage of ongoing myelination.

Antibodies

Mouse anti-MBP (directed against *Xenopus* MBP, 1:1000, kindly provided by Dr. Saburo Nagata), chick anti GFP (1:1000, Aves Lab, USA), rabbit anti-S1PR5 antibody (1:100 Imgenex, USA), goat anti-human Sox10 (1:50 R&D systems). Corresponding secondary antibodies conjugated Alexa 350, Alexa 488, Alexa 594 and Alexa 647 (1: 1000, Invitrogen). For CLARITY we used chicken anti-GFP, Avès Labs, 1:400 as primary antibody and goat anti-chicken Alexa Fluor 488, Invitrogen, 1:600 as secondary one.

CLARITY and Imaging using a high refractive index solution are detailed in the supplementary materials.

Immunostaining of clarified samples

CLARITY-processed brains were incubated in blocking solution overnight at 4°C. Samples were further incubated in staining solution with primary antibodies (anti-GFP) for 7d at room temperature with under gentle agitation. After 2 washes with PBT, samples were incubated in a staining solution with secondary antibody for 7d at room temperature. Samples were subsequently washed for 48h in PBT.

Immunolabeling on tissue sections

Tadpoles were fixed by immersion in 4% paraformaldehyde and treated for immunostaining as described.³

Electron microscopy and morphometry are detailed in the supplementary materials.

Metronidazole preparation and use

Metronidazole (10mM, Fluka) was dissolved in filtered tap water containing 0.1% DMSO (Sigma Aldrich) and added to aquaria water for 10 days. Tadpoles were maintained in 600 ml of metronidazole solution (maximum 10 tadpoles/600 ml) at 23.5°C in complete darkness (metronidazole is light-sensitive) and changed every 72h throughout treatment. For regeneration experiments, metronidazole-treated animals were allowed to recover for up to 8 days in either normal water or containing compounds to be tested at different concentration in ambient laboratory lighting (12h light 12h dark).

Quantification of GFP⁺ cells

GFP was detected directly by fluorescence in live embryos using an AZ100 Nikon Zoom Macroscope. In the optic nerve, the total number of GFP⁺ cells was counted, from the

emergence of the nerve (i.e., after the chiasm) to the retinal end before and after metronidazole treatment and after treatment with the compound to be tested on the same embryos. Double-blind counts were performed independently by two researchers. Data were compared to control untreated animals of the same developmental stage. Data presented are the mean \pm SEM of number of GFP+ cells counted on at least 6 tadpoles per condition.

Preparation of sgRNAs, generation of SIP5 null mutant and PCR primers are described in the supplementary materials.

Statistical Analysis

Prism v5 software was used for statistical analyses. Results are expressed as means \pm SEM. To assess differences between more than two groups when comparing efficiency of remyelinating drugs, parametric two way ANOVA was used (after normality test passed using Shapiro Wilk normality test) followed by an uncorrected Fisher's LSD test. Other *P*-values for comparison of means are from Student's two-tailed *t*-tests. Analysis of covariance was used to analyze regression lines. Statistical significance was set at $p < 0.05$. Significant differences are indicated by asterisks: * $p < 0.05$, ** $p < 0.01$, and *** $p < 0.001$.

Results

Oligodendrocyte visualization in MBP-GFP-NTR transgenic Xenopus

In the brain stem, GFP positive cells are detected from stage 43 onward. This corresponds to the developmental stage when myelination markers such as proteolipid protein and MBP are first immunodetected in the hindbrain (stage 42/43) before spreading throughout the brain and spinal cord by stages 46/47.⁹ Electron microscope studies on optic nerve¹⁰ showed myelination to begin in the middle portion at stage 48/49, with numbers of myelinated axons increasing sevenfold between stage 50 and 57. Distribution of myelin forming oligodendrocytes at stage 48 and 52 is

shown (Figure 1A-C). Oligodendrocytes populate the ventral half of the hindbrain and spinal cord on each side of the mid-line. At this early developmental stage oligodendrocytes have not yet migrated long distances from their origin and are myelinating the major descending fibre tracts. This is well illustrated by the 3D-reconstruction of whole-mount CNS preparations performed after tissue clarification, i.e. replacement of light-scattering lipids of the brain and spinal cord by a transparent hydrogel matrix (Supplementary file Movie 1).

*Conditional ablation and spontaneous repair in the MBP-GFP-NTR *Xenopus tadpole**

Stage NF52-53 MBP-GFP-NTR transgenic *X. laevis* were treated with metronidazole (10mM) introduced in the aquarium water to induce conditional demyelination. Screening experiments focused on the embryonic optic nerve, easily monitored by live imaging. The number of GFP⁺ oligodendrocytes is moreover small and very reproducible at each developmental stage, facilitating quantification. After 10 days metronidazole, the number of GFP⁺ cells per optic nerve significantly decreased from 13.9 ± 1.0 to 2.3 ± 2 ($p < 0.01$) (Figure 1D upper panel). The extent of demyelination was further investigated using electron microscopy (EM) showing the depletion of myelin around axons, and the presence of myelin debris (Supplementary Figure S2 A, B). After 10 days metronidazole exposure, tadpoles were returned to normal water and recovery occurred rapidly: oligodendrocytes per optic nerve reached 8.3 ± 0.9 and 13.3 ± 1.5 at R3 and R8, respectively (Fig 1D lower panel). Of note, during the 10 days of metronidazole treatment at 23°C, tadpoles progress from stage 52 to 55 (see <http://www.xenbase.org/anatomy/alldev.do>); therefore during the recovery period not only demyelinated axons undergo remyelination, but it is likely that in addition some previously naked axons are also myelinated.

*Transgenic MBP-GFP-NTR *Xenopus tadpoles* as a screening platform for molecules enhancing remyelination*

We reasoned that our model could serve for screening potential remyelination-favoring molecules by quantifying remyelination rates as a function of increases in GFP+ cell numbers following demyelination. Since spontaneous repair was often complete at R8, we chose to evaluate repair efficiency only at R3, i.e., 3 days after metronidazole removal. We compared the promyelinating effect of 20 compounds for which an effect on myelination (or remyelination) has been described (Table 1). Among the molecules tested, the three most efficient were, by increasing order: siponimod (3nM), UC-42-WP04 (10nM) and clemastine (100nM)(Figure 2). It is noteworthy that UC-42-WP04, a selective S1PR5 agonist,¹¹ was amongst the most efficient remyelinating drugs, while AUY954, a selective S1PR1 agonist,¹² had no effect. Since siponimod is a dual S1P1 and S1P5 receptor agonist, we investigated further siponimod's remyelinating potential.

Xenopus oligodendrocytes express S1P5 receptor

As oligodendrocytes in the mouse brain selectively express S1PR5,⁷ we questioned whether S1PR5 expression was also restricted to oligodendrocytes in *X. laevis*. We investigated oligodendroglial expression of S1PR5 in *X. laevis* by triple immunolabeling of brainstem tissue sections of *MBP-GFP-NTR* tadpoles (stage 52-53) using a rabbit anti-S1PR5 antibody, a chicken anti-GFP to label mature oligodendrocyte and a goat anti-Sox-10 to label all cells from the oligodendroglial lineage. We observed a perfect cellular co-localisation of Sox10 and S1PR5 (21.5 ± 6.5 cells/mm²) and all the GFP+ cells were S1P5+ (14.5 ± 5.3 GFP+/S1PR5+ cells/mm²). Since, in the CNS, Sox10 is an established marker of the oligodendroglial lineage, we concluded that the S1P5+/Sox10+/GFP- cells are most likely OPCs. Altogether our data suggest that in the *Xenopus* CNS, as in the mouse, S1P5 expression is restricted to the oligodendrocyte lineage and that all myelin-forming oligodendrocytes (GFP+ cells) express S1PR5 (Figure 3).

Detection of myelinated fibers following siponimod treatment

To verify that increased numbers of myelinating oligodendrocytes translated into increased myelination, brain stem tissue sections were immunolabeled with anti-MBP, an established myelin marker. Transgenic *MBP-GFP-NTR X. laevis* (n=18)(stage 52-53) were treated for 10 days with metronidazole (n=15), then returned to normal water (n=3) or water containing siponimod (3nM) for either 3 days (n=3) or 8 days (n=3) prior to MBP immunostaining (Figure 4). After 10 days of metronidazole treatment a nearly complete demyelination was achieved (compare D0 and D10 in Figure 4). After metronidazole removal, exposure to siponimod for 3 days (Figure 4 Siponimod R3) resulted in a dramatic increase in myelinated internodes by comparison with spontaneous remyelination (Figure 4 Control R3). At R8 the extent of myelinated fibers was back to the level prior to demyelination (Figure 4 D0, Siponimod R8, and control R8. EM examination of tadpoles following 6 days of recovery confirmed a partial remyelination of MTZ-induced demyelinated axons (Supplementary Figure S2 C), in agreement with the return to nearly normal level of the number of GFP⁺ cells. Evaluation of the g-ratios representing myelin thickness with respect to axon diameter revealed a reduction in myelin thickness, suggestive of remyelination, in siponimod remyelinated animals ($g = 0.85 \pm 0.04$) versus controls not subjected to demyelination ($g = 0.82 \pm 0.04$, $p < 0.001$) (Supplementary Figure S2 D).

The promyelinating effect of siponimod is mediated through S1PR5

Siponimod is a potent and selective agonist of S1P receptors with an EC₅₀ of 0.39nM for S1P1 and 0.98nM for S1P5 receptors, respectively.¹³ We questioned which receptor implicated the pro-myelinating effect of siponimod. As remyelination was observed with the selective S1PR5 agonist (UC-42-WP04), but not with the selective S1PR1 agonist (AUY954) we favored the possibility that siponimod induced remyelination through S1PR5. If this were the case, siponimod should lose efficiency after S1P5 receptor deletion. We took advantage of the clustered regularly interspaced short palindromic repeats (CRISPR)-Cas system (Crispr/Cas9) to test this hypothesis.¹⁴ To determine the exact mutations of *s1pr5-S* and *s1pr5-L* the genomic DNA from 6

F0 embryos, identified after the T7 endonuclease-based mutation detection, was extracted and sequenced. Each embryo presented different degrees of deletion (between 2 to 169 bp) leading to shift of the reading frame and premature STOP codon, for both *s1pr5-L* and *s1pr5-S* genes (Figure 5A). Each of these embryos at stage 55 was submitted to metronidazole demyelination and treated with siponimod. The pro-remyelinating effect of siponimod was no longer observed in the S1PR5^{-/-} embryos (Figure 5B).

To confirm oligodendroglial S1P5 receptor loss, mutated tadpoles were euthanized after the demyelination-remyelination experiments, and immunolabeled with anti-S1PR5 antibodies on brain stem sections. As illustrated, (Figure 5C), S1PR5 protein was not immuno-detectable in GFP+ cells in any of the six *MBP-GFP-NTR; s1pr5^{-/-}* tadpoles examined.

Discussion

For the past 15-20 years significant therapeutic progress has resulted from developing anti-inflammatory strategies in MS. These treatments are efficient in diminishing numbers of relapses, but their influence on long-term disability is unclear. Loss of the myelin sheath perturbs normal axon functioning and persistent demyelination renders them vulnerable to irreversible damage leading to axonal transection, with consequent disability accumulation. Spontaneous remyelination can occur, but most often endogenous repair mechanisms are insufficient, notably during aging. The experimental demonstration that environmental clues can restore remyelination potential of aging OPCs indicates that pharmacological approaches may be developed to restore, increase or accelerate remyelination.¹⁵ Several groups have recently developed *in vitro* screening approaches for small molecules that drive OPC differentiation.^{16, 17, 18} These *in vitro* high throughput screens can identify hit molecule candidates to be further validated in *in vivo* experimental models of demyelination.

Rodents are the preferred animal models, with primary demyelination induced either by

detergents to disorganize the myelin sheath (*e.g.* lyssolecithin) or by killing oligodendrocytes with toxins (such as cuprizone or ethidium bromide). In other models demyelination is secondary to an auto-immune attack as in MOG-induced EAE.¹⁹ Toxin-induced models of demyelination do not mimick MS, but they allow the fundamental biology of remyelination to be explored without the confounding and complicating involvement of an autoimmune process.²⁰ Although rodents may be considered as the “standard” experimental model to investigate demyelination and remyelination, they are costly, time consuming and not ideal for longitudinal investigation in the same animal. Although the last decade has seen considerable progress for *in vivo* imaging of live cells in the normal and demyelinated CNS,²¹ technical issues strongly limit imaging in deep layers of mouse brain, such as myelinated axonal tracts. Tadpole transparency allowed us to overcome these limitations by using transgenic *Xenopus* expressing fluorescent reporter and a suicide protein in oligodendrocytes.

We present results on 20 molecules previously described, usually *in vitro*, as candidates to promote remyelination. The *Xenopus* results are in good agreement with reports on pro-myelinating efficacy of clemastine,¹⁷ benztropine¹⁶ and retinoic acid²² in mammalian cells. A recent small size phase 2 study of clemastine in chronic optic neuritis showed some positive results. Concerning siponimod no data on remyelination is reported, but a recently completed phase 3 study showed some positive results in secondary progressive MS.⁵ Fingolimod, (acting on S1PR1, S1PR2, S1PR3 and S1PR5, and which is used in the clinic) also promoted remyelination, a finding in agreement with previous data in culture.²³

In contrast, clobetasol and miconazole, known to be efficient remyelinating candidates *in vitro* as well as *in vivo* in the mouse¹⁸ were without effect in the *Xenopus* model. One possible explanation is that these molecules were screened for their ability to generate new oligodendrocytes, while in the *MBP-GFP-NTR* *Xenopus* model there is no need to generate new OPCs since remyelination results from maturing OPCs (Sox10+ cells) already present in the optic nerve prior to demyelination, without requiring additional production of OPCs.²⁴ Generation of new

oligodendrocytes is only one possible mechanism to favor remyelination. It has been described that OPCs are present in some chronic lesions of MS, so remyelination is not only limited by an absence of precursors or their failure to generate oligodendrocytes, but by the defective maturation of OPCs.²⁵

However, an advantage of *Xenopus* as opposed to other aquatic models, lies in myelin biology and the fundamental difference in the molecular composition of myelin between teleosts and tetrapods. Teleosts, and zebrafish in particular, use P0 as major myelin protein both in the CNS and the peripheral nervous system (PNS). In contrast, starting with amphibians (including *Xenopus*), a dramatic evolutionary change in myelin composition between PNS and CNS occurred. In tetrapods (including amphibians and mammals) the major myelin protein in the PNS remains P0, while in the CNS it is PLP and MBP. This event occurred in amphibians and has been maintained throughout evolution up to mammals, including humans.

An advantage of the *Xenopus* model is read-out rapidity: after 10 days of demyelination, remyelination effects were seen by 3 days recovery (hence a total of 13 days), while the cuprizone model in rodent requires at least 6 weeks for demyelination and an additional 3 weeks for recovery. A final advantage of the *Xenopus* model is that assessing demyelination and remyelination is performed on the same animals and does not require immunohistochemistry. Although the *Xenopus* model, as described, is more indicated as a medium throughput screen, it has the potential to be automatized for high-throughput screens. Live imaging of remyelination (automated counting of GFP+ cells) is theoretically possible through the modification of existing automated devices such as the VAST (Vertebrate Automated Screening Technology, Union Biometrica, MA, USA). This device allows the automated pumping from multiwell plates and automated imaging at cellular level of whole zebrafish embryos. This system is limited by the valve and motor pin diameter, which has a maximum of 700 micron diameter. To be

transposable to *Xenopus* tadpoles would require adapting the technology by using larger size capillaries. Overall, the procedure is far more efficient when compared with the efforts, time and cost necessary to perform the same type of experiment in mice. However, whilst ideal for screening small molecules, a limit of the *MBP-GFP-NTR* *Xenopus* model is that it is not designed to test antibodies, such as anti-Lingo, which would require to be injected into the tadpole and not simply added to the swimming water. Another limitation is that the *Xenopus* tadpole model may not be adapted to screen drugs aimed at overcoming the decline in remyelination that occurs with advancing age particularly for long-term demyelinating diseases such as MS.

In addition to the rapidity with which the test can be performed, another appreciable advantage of the *Xenopus* model is using Crispr/Cas9 technology so the molecular mechanism of a given compound can be dissected, with ease and precision.¹⁴ In the case of siponimod, not only S1P5 receptor was directly deleted in the *MBP-GFP-NTR*- lineage, but the effect of the deletion was tested on 6 different KO animals. Three arguments strongly support that the remyelinating potency of siponimod is mediated through the S1P5 receptor, first, another S1PR5 agonist improved remyelination; second, in the absence of S1PR5 siponimod was no longer active and, third, a S1PR1 agonist was inefficient. We did not attempt to delete S1P1 because it is established that within endothelial cells this receptor is required for the coverage of vessels by vascular smooth muscle cells²⁸ and lethality occurs in embryos of S1P1 receptor-deficient mice, due to rupture of the unreinforced vessels.²⁷ However, although unlikely, we cannot definitively exclude that some of the benefit from siponimod could be from its potential to act through a different mechanism than a S1PR5 agonist.

A recently completed Phase III randomized, controlled study of siponimod in secondary progressive MS has met its primary endpoint, compared with placebo.⁵ Obinata and co-workers have reported the genetic variations of the *s1pr1* coding region from exon sequencing of about 12,000 individuals and their functional consequences, suggesting that individual genetic variations

of *s1pr1* can influence receptor function and, therefore, infer differential interaction with S1PR1-targeted therapeutics.²⁸ Concerning *s1pr5*, deletion of *s1pr5* has no apparent evident clinical phenotype in either mouse or *Xenopus*. Whether patients with *s1pr5* mutation or deletions may differ in their remyelination capacity, disease severity and/or response to siponimod is an as yet unexplored hypothesis.

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Author Contribution

A.M. and B.Z. conceived the studies and designed the experiments. A.M. performed most of the experiments. S.S. carried out the immunohistochemistry experiments using S1PR5 antibodies. L.A. and Q.V. carried out screening of drugs. J.B.F. carried out the CLARITY experiments. C.G. provided Cas9 protein and advices on Crispr/Cas9 genome editing. D.D., J.L.T., C.L. and B.D. provided technical support and intellectual inputs. B.Z. wrote the paper with inputs from all co-authors.

Declaration of Conflicting Interests

B.Z. received research support and travel grant from Novartis Pharma, which own patent rights to siponimod (BAF312). B.D. is a founder of the company WatchFrog and D.D. is an employee of WatchFrog. C.L. has participated to advisory boards for Vertex, Roche, Biogen, Novartis, Genzyme.

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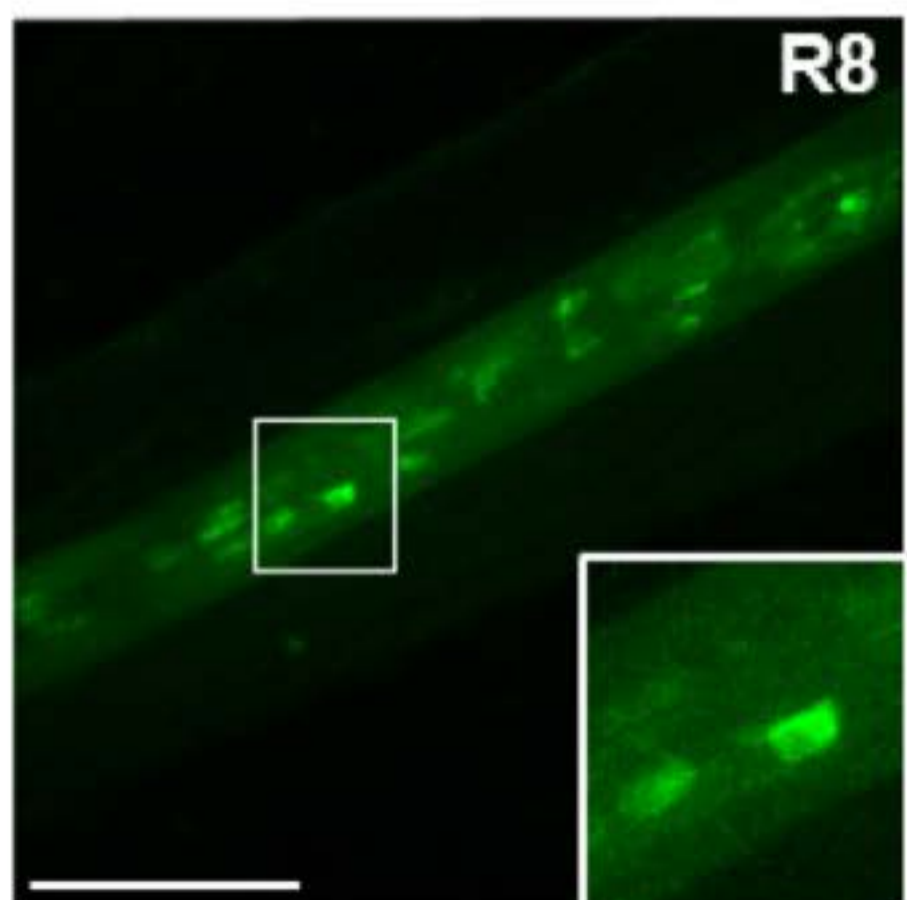
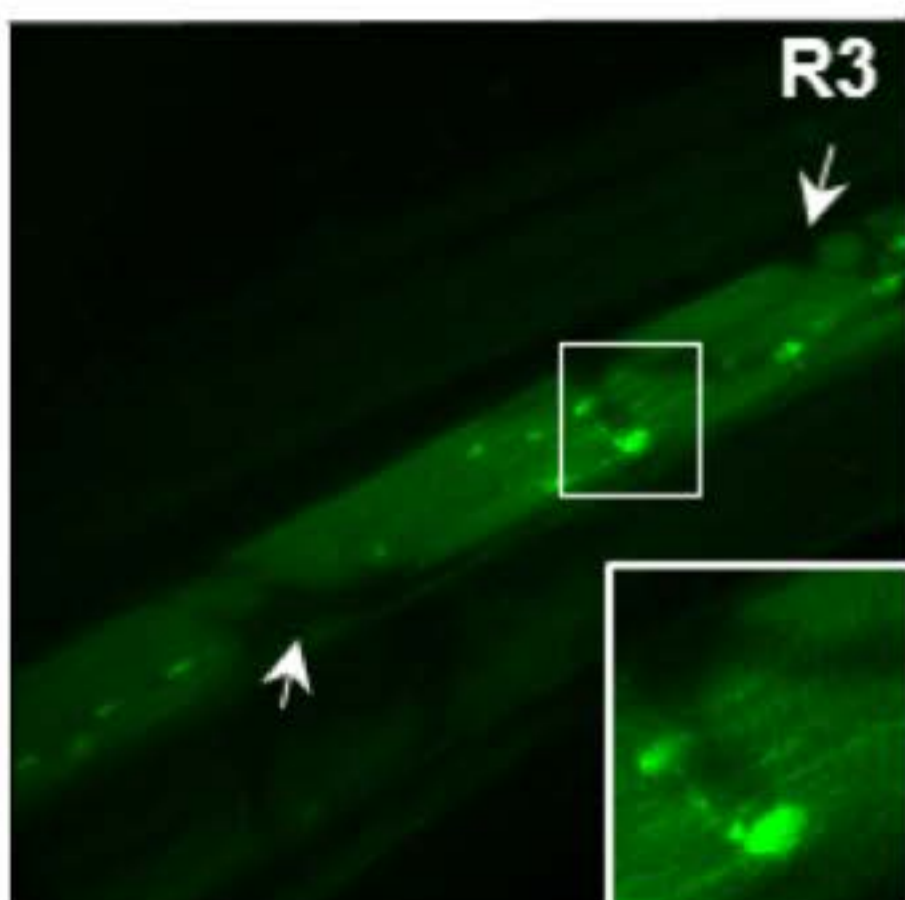
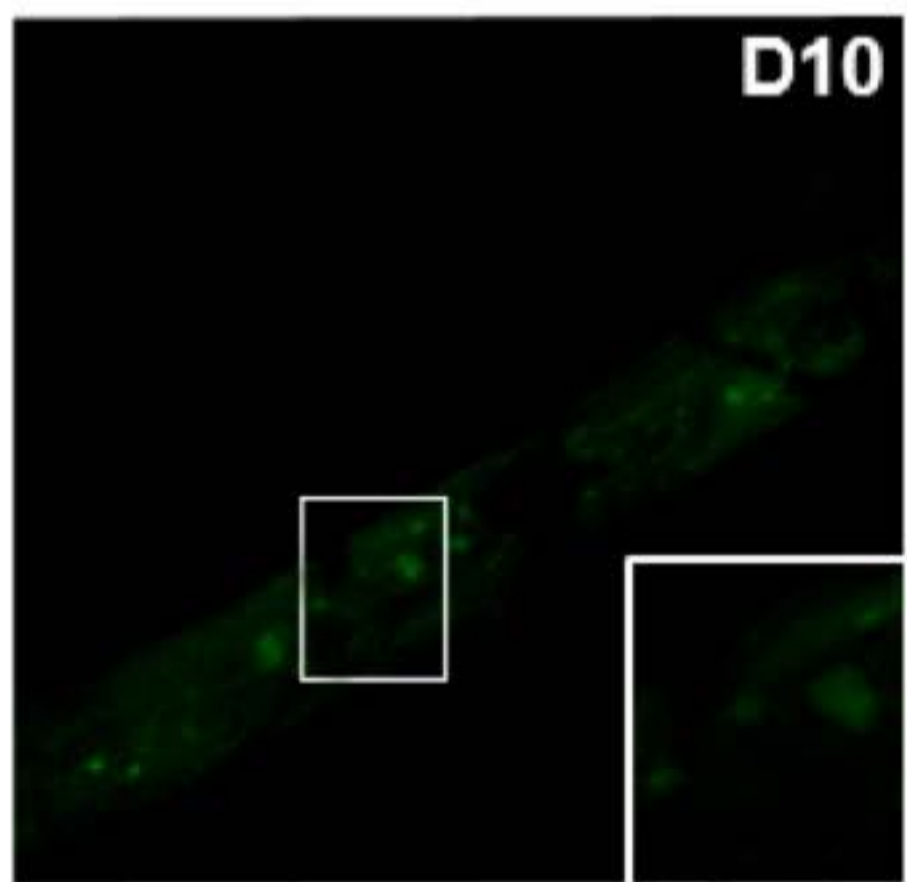
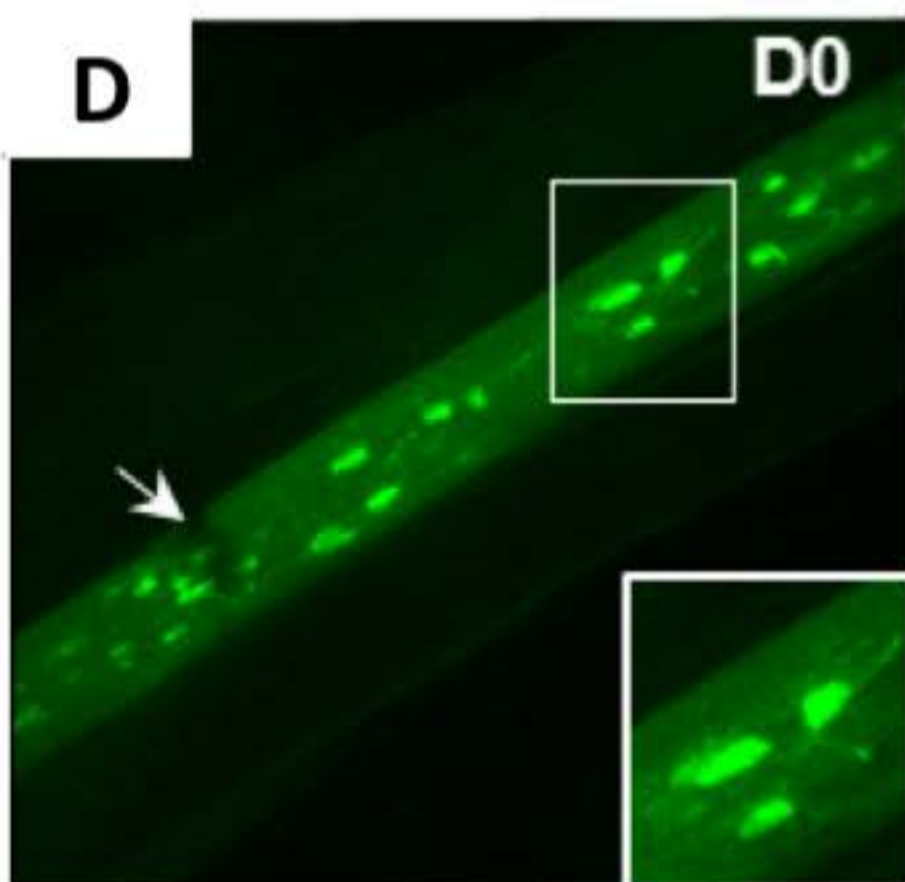
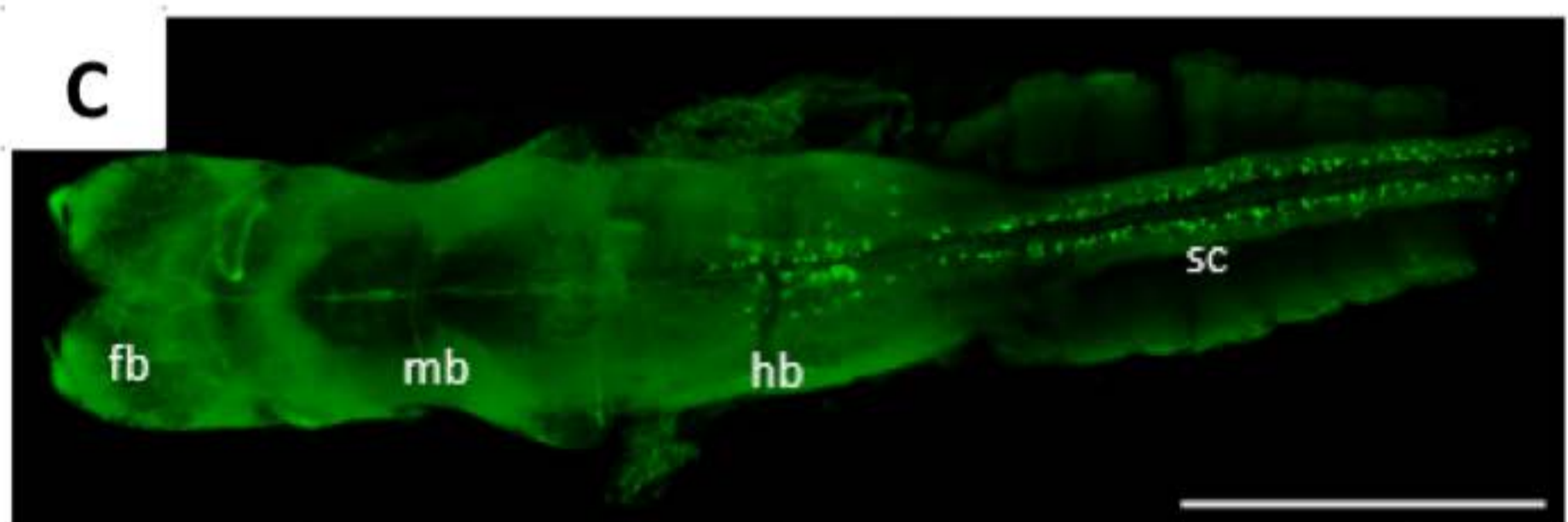
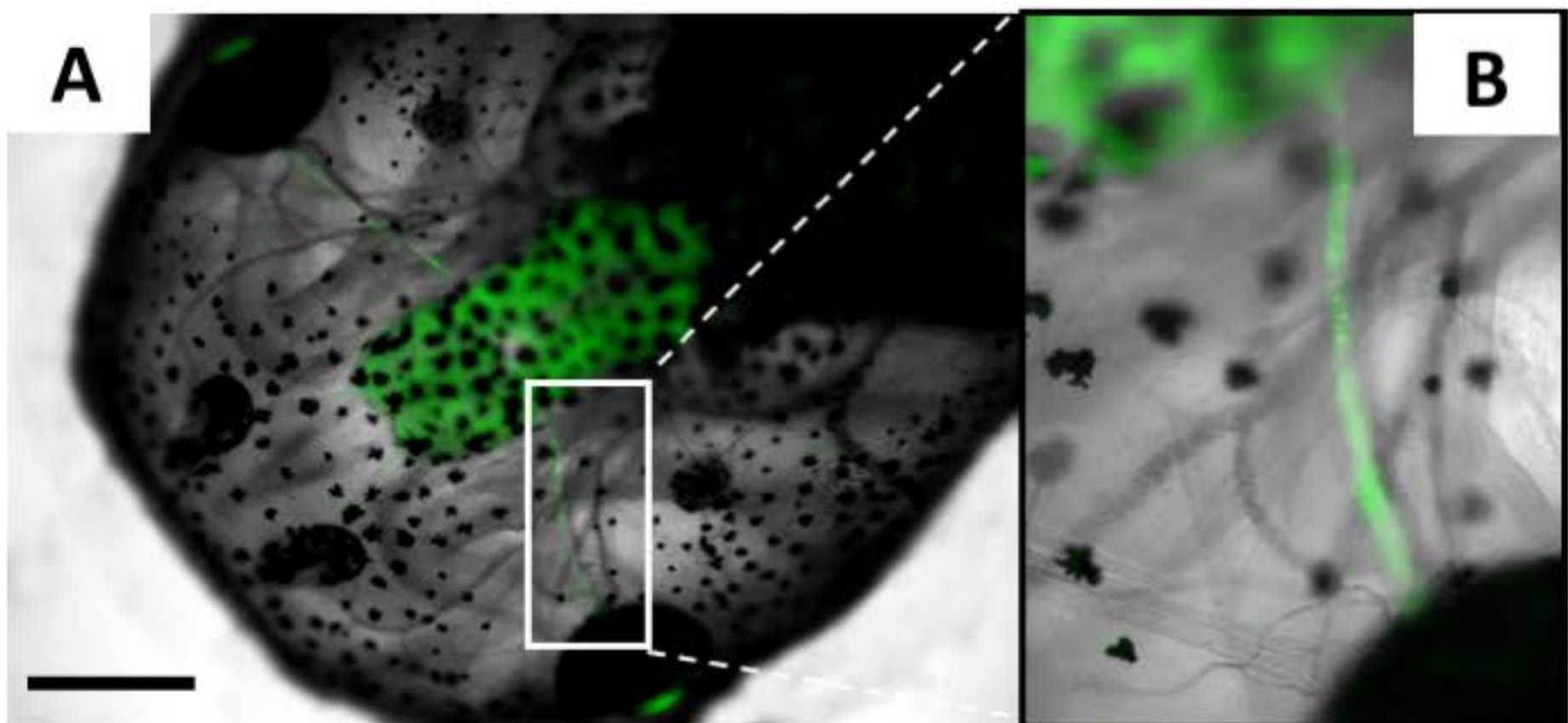
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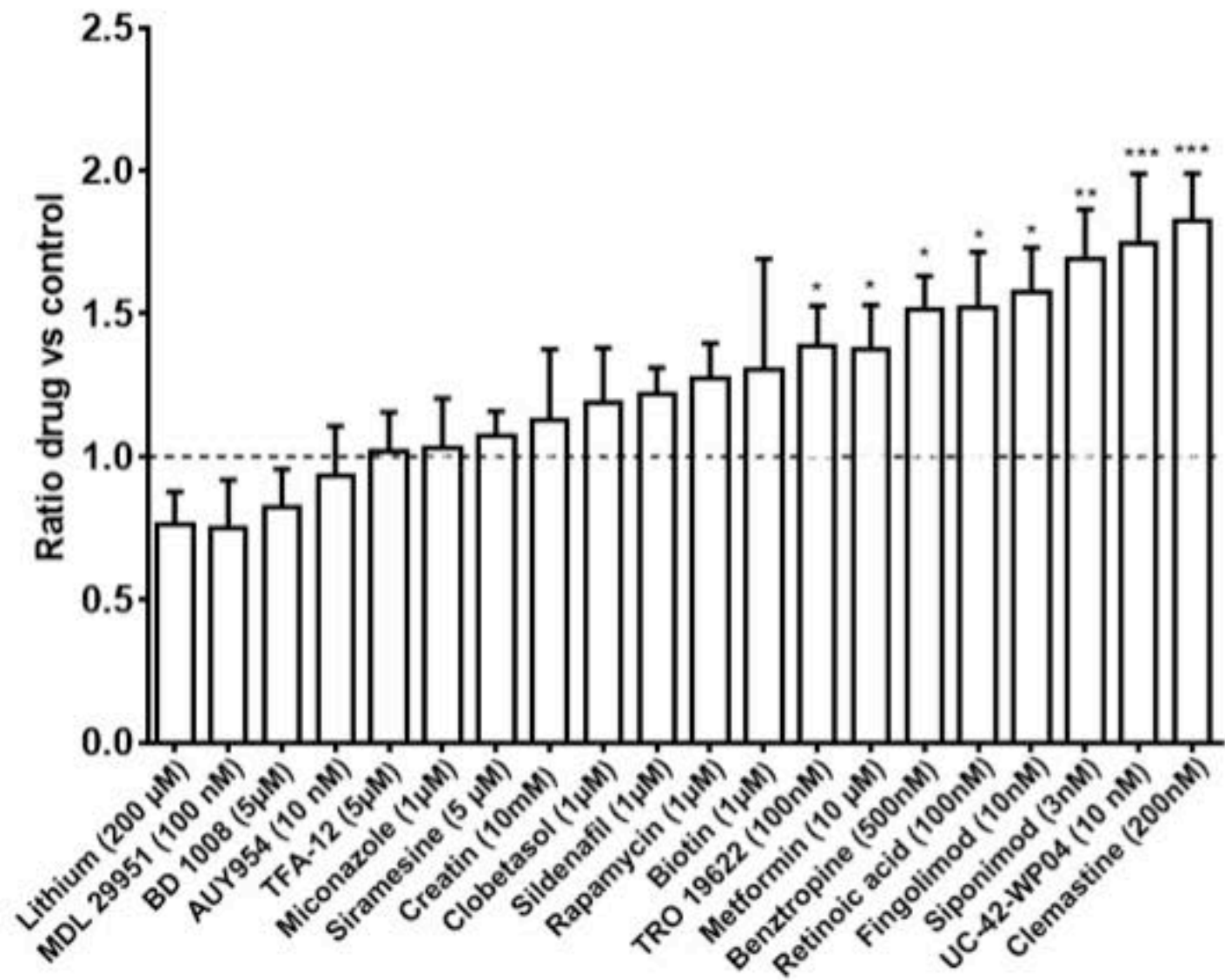
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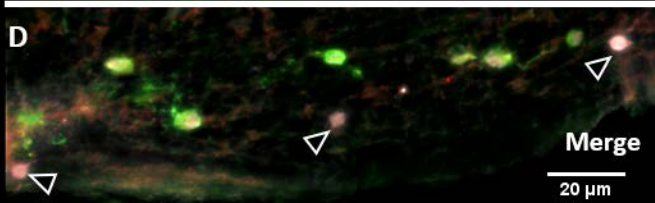
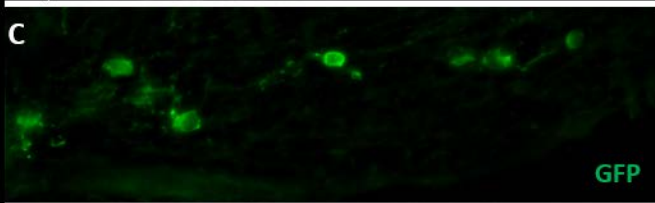
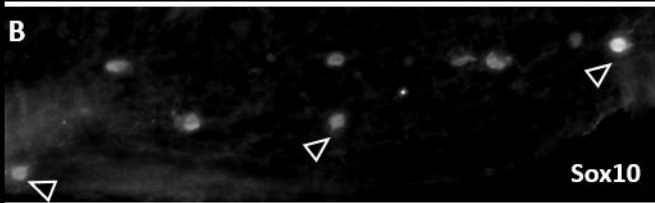
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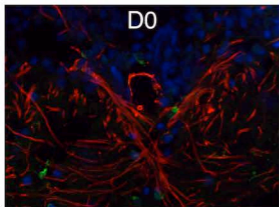
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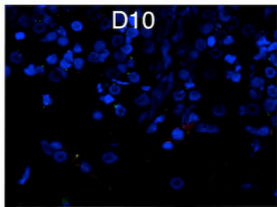




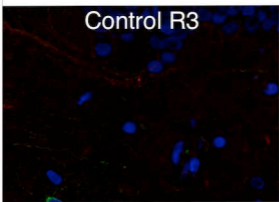
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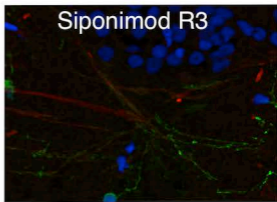
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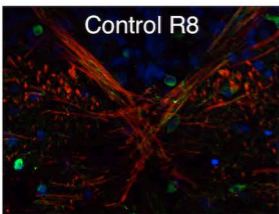
Control R3



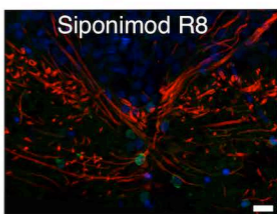
Siponimod R3



Control R8



Siponimod R8



A

S1P5.L sgRNA PAM
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 #1 ATTGCGATAGAACGCCACATTACTATGGTTCGGATGAAATGA---AGCGGGGACAAGAAGGGGGCGCATG Δ3-stop(444)
 #2 ATTGCGATAGAACGCCACATTACTATGGTTCGGATGAAAC-----AGCGGGGACAAGAAGGGGGCGCATG Δ5-stop(677)
 #3 ATTGCGATAGAACGCCACATTACTATGGTTC-----AGCGGGGACAAGAAGGGGGCGCATG Δ14-stop(677)
 #4 ATTGCGATAGAACGCCACATTAC-----AGCGGGGACAAGAAGGGGGCGCATG Δ22-stop(514)
 #5 ATTGCGATAGAACGCCACATTACTATGGTTC-----GGGGACAAGAAGGGGGCGCATG Δ17-stop(677)
 #6 GTTTCGGATAGAACGCCACATTACTATGGTTCGGAT-----GACAAGAAGGGGGCGCATG Δ16-stop(514)

S1P5.S sgRNA PAM
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 #2 ATTGCGATAGAACGTCACATTACTATGGTTCGGATGAAA-----CGGGGACAAGAAGGGTTCGCATG Δ8-stop(710)
 #3 ATTGCGATAGAACGTCACATTACTATGGTTCGGATGAA-----ACAAGAAGGGTTCGCATG Δ14-stop(710)
 #4 ATTGCGATAGAACGTCACATTACTA-----CAGCGGGGACAAGAAGGGTTCGCATG Δ19-stop(512)
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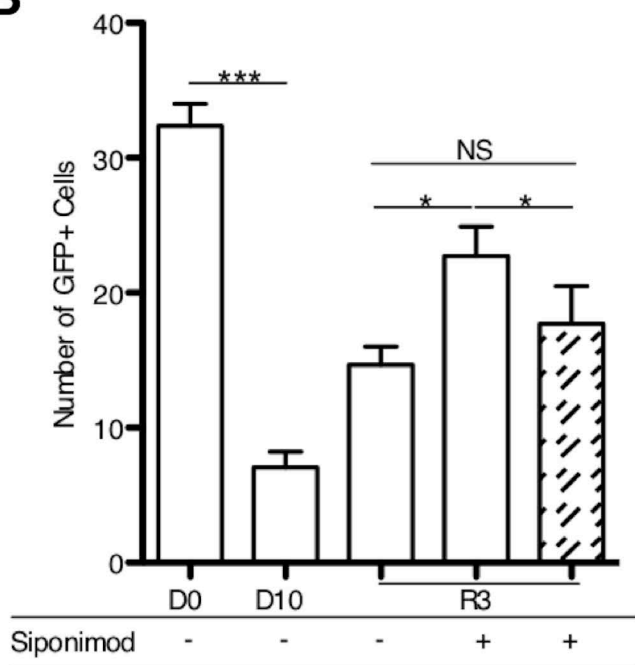
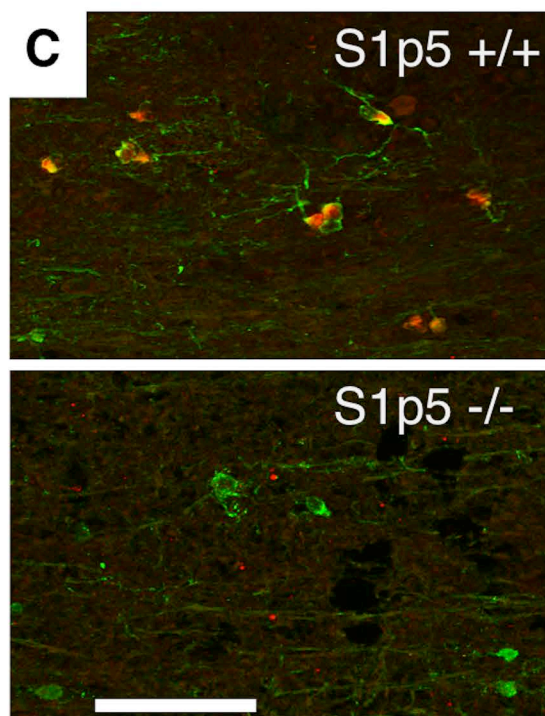
B**C**

Table 1: List of compounds tested. (a) Concentrations used *in vivo* (when available) are shown in italics

Compound tested	Known mode of action	Concentration used (a)	Reference
AUY954	Sphingosine 1 phosphate receptor 1	1.2nM// <i>1-10mg/kg/d</i>	(12) Pan et al., 2006
Benztropine	Anticholinergic	2μM// <i>2.5mg/kg</i>	(16) Deshmukh et al., 2013
BD 1008	Sigma receptor	5μM	(29) Demerens et al., 1999 & unpublished data
Biotin	Vitamin B8	<i>300mg/d</i>	(30) Sedel et al., 2015
Clemastine	Anti-histaminic anti-muscarinic	1μM / <i>10mg/kg</i>	(17) Mei et al., 2014
Clobetasol	Glucocorticoid anti-fungal	5μM// <i>10mg/kg</i>	(18) Najm et al., 2015
Creatine	Peptide derived from arginine, glycine & methionine	Ratio NAA/Cr P<0.16	(31) Fayed et al., 2006
Fingolimod	Sphingosine 1 phosphate receptor 1, 2, 3 and 5	10nM/ <i>0.5mg/d</i>	(23) Miron et al., 2010
Lithium	Alcalin bipolar disorder	<i>50mg/kg/d</i>	(32) Makoukji et al., 2012
MDL29951	NMDA antagonist	100nM	(33) Hennen et al., 2013
Metformin	Biguanide Type2 diabetes	200μM	(34) Paintlia et al., 2013
Miconazole	Imidazole anti-fungal	1μM// <i>1mg/kg</i>	(18) Najm et al., 2015
Rapamycin	Inhibitor of mTOR	15nM	(35) Tyler et al., 2014
9-cis-Retinoic acid	Retinoic receptor	50nM	(22) Huang et al., 2011
Sildenafil	Phosphodiesterase inhibitor	<i>25mg/kg/d</i>	(36) Raposo et al., 2014
Siponimod	Sphingosine 1 phosphate receptor 1 and 5	1-1000nM// <i>0.25-10mg/d</i>	(37) O'Sullivan et al., 2016
Siramesine	Sigma receptor	1μM	(29) Demerens et al., 1999 & unpublished data
TFA-12	Tocopherol derivative	<i>0.39mg/kg</i>	(38) Blanchard et al., 2013
TRO 19622 (Olesoxime)	Cholesterol derivative	<i>5ml/kg or 60-600μg/g</i>	(39) Magalon et al., 2012
UC-42-WP04	Sphingosine 1 phosphate receptor 5	3nM// <i>3mg/kg</i>	(11) Mattes et al., 2010

Supplementary files:

Supplementary materials:

CLARITY

Brains of stage 48 *MBP-GFP-NTR* tadpoles were submitted to clarification following the procedure developed by TEFOR core facility < <http://tefor.net/pages/main/>>. Briefly, fixed tadpoles and dissected brains were subjected to clarification following the CLARITY protocol (Chung & Deisseroth, 2013; Epp et al., 2015) with some tissue-specific adaptations: samples were infused in a cooled solution of extemporaneously prepared hydrogel monomers, 4% acrylamide and 0.0025% bis-acrylamide for 2d at 4°C. Hydrogel polymerization was triggered, after degassing, by replacing atmospheric oxygen with nitrogen in a desiccation chamber for 3h at 37°C. Samples were then transferred into embedding cassettes for lipid clearing. The samples were subsequently thoroughly washed in 0.01 M PBS, Tween-20 0.1%, 2d with gentle agitation.

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Imaging using a high refractive index solution

A fructose-based high refractive index solution (fHRI) was prepared and the refractive index of the solution was adjusted to 1.4571 using a refractometer (Kruss). Before imaging the clarified samples were mounted in 1% low melting point agarose covered with fHRI for imaging. Whole-mount brain fluorescence was captured using a Leica TCS SP8 laser scanning confocal microscope equipped with a Leica HC FLUOTAR L 25x/1.00 IMM motCorr objective.

Preparation of sgRNAs, and generation of S1P5 null mutant: CRISPR sgRNA were designed using the « ChopChop » web tool (<http://chopchop.cbu.uib.no/>) and in vitro transcribed as described by Nakayama et al., 2013. *X. laevis* is an allotetraploid composed of two separate diploid genomes (2n=36). *X. laevis* has nine pairs of chromosomes named Xla1S and Xla1L through Xla9S and Xla9L where S (short) and L (long) refer to their relative lengths. The recent sequencing of *X. laevis* genome has identified two homologous genes *s1pr5S* and *s1pr5L* encoding two forms (S and L) of S1P5 receptors (Fig 1A-B). To delete both S1P5 receptors we designed sgRNAs targeting both the L and S homologs, by choosing a sequence common to both receptors. Briefly, the template was prepared using a PCR-based method using as forward primer (60 nt): 5'-TAATACGACTCACTATA (T7 Promoter)-GGTTCGGATGAAACACTACAG (target sequence)-GTTT'TAGAGCTAGAAATAGCAA-3'(complimentary overhang with reverse primer). For PCR Phusion polymerase (NEB) was used using stock CRISPR-R primer (80 nt) (5'-AAAAGCACCGACTCGGTGCCACTT'TTTCAAGTTGATAACGGACTAGCCTTATT TTA ACTTGCTATTTCTAGCTCTAAAAC-3') to generate the complete 100 nt sgRNA DNA template. PCR sgRNA were transcribed using T7 MEGAShortscript kit (Ambion) using 8 µl of purified PCR product in 20 µl at 37 °C for 4 h. RNA was purified by precipitation with isopropanol. The sgRNA have been chosen on the ChopChop site <<https://chopchop.rc.fas.harvard.edu>> as having no off-target. Since it was described that off-target predictors can fail to detect some off-target sites, due to an inappropriate search algorithm (Doench et al, Nat Biotechnol. 2016;34:184–91) we also used the more recently described CRISPOR website <<http://crispor.tefor.net/>> that is able to find all validated off-targets. Using this site for GGTTCGGATGAAACACTACAG sgRNA we can find 5 off-targets sites with 3 and 4 mismatches and no mismatches in the 12 bp adjacent to the PAM. However, all these potential off targets are present in introns. We did not confirm these potential sites by sequencing.

In addition, to facilitate detection of embryos KO for *s1pr5*, we co-injected sgRNA for *slc45a2*, a gene required for pigmentation, assuming that albinos embryos would also be KO for *s1pr5* (not shown). sgRNA for *slc45a2* were designed by M. Salanga, (National Xenopus Resource, Woods Hole, MA). Oocytes from *MBP-GFP-NTR* were fertilized with spermatozooids from wild type male *Xenopus laevis* and immediately microinjected with a mixture of *slc45a2* and *s1pr5* sgRNA and Cas9 protein. Of note, in preliminary experiments we found that micro-injection in fertilized oocytes of recombinant Cas9 protein was far more efficient compared to *Cas9* mRNA to produce total albinism, as evaluated by the degree of mosaic pigmentation of the eyes. To detect *s1pr5* mutation among albinos (GFP+) tadpoles, the *s1pr5* sequence was PCR amplified and amplicons submitted to T7 endonuclease 1 digestion (Fig 1C).

References:

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PCR primers

The regions targeted by the complex Cas9 protein-sgRNA were PCR amplified with genomic primers. We used specific forward primers to amplify the *s1p5L* (**Fn1**) CTICTGGCTGGTGTGGCATA and *s1p5S* (**Fn2**) CTTACCCGGGAAGCTCAACA, respectively. Reverse primer (**Rd1**) GGTCAACAGCCAGAGACTTGGC was common to *s1p5L* and *s1p5S*. PCR parameters were: 5 min at 95 °C followed by 35 cycles of 95 °C for 30 s, 72 °C for 40 s, and 60 °C for 1 min, with a final 10 min extension.

Electron microscopy and morphometry

Larvae were fixed in a mixture of 2% paraformaldehyde, 2% glutaraldehyde, in 0.1M cacodylate buffer pH 7.4 and 0.002% calcium chloride overnight at 4°C, washed in 0.1M cacodylate buffer, and postfixed in 1% osmium tetroxide, 1% potassium ferricyanide in 0.1M cacodylate buffer. After washing in cacodylate buffer and water, larvae were incubated in 2% uranyl acetate aqueous solution at 4°C overnight. After rinsing twice in water, larvae were dehydrated in increasing concentrations of ethanol, final dehydration was in 100% acetone (twice 10min each). Samples were infiltrated with 50% acetone 50% Epon for 2h and then embedded in Epon (EMBed 812, Electron Microscopy Sciences Cat 14120) Blocks were heated at 56°C for 48h. Ultrathin sections (70 nm) were examined on an HT7700 electron microscope (Hitachi) operated at 70kV.

Images were analyzed using Image J (NIH) for *g*-ratio measurements by manually drawing 2 lines across the long and short axis of each axon and axon plus myelin (myelinated fiber). Lengths of the lines (in pixels) as generated by ImageJ were averaged across the 2 perpendicular measurements and converted to micrometer (μm) using the image scale bars. The *g*-ratio was determined by dividing the mean diameter (long and short axis) of an axon by the mean diameter of the same myelinated fiber.