Signals from the head and germinative region differentially regulate regeneration competence of the tapeworm *Hymenolepis diminuta*.

Elise McCollough Nanista[1], Landon Elizabeth Poythress[1], Isabell Reese Skipper[1], Trevor Haskins[1], Marieher Felix Cora[2] and Tania Rozario[1]*

[1]University of Georgia, Athens, GA, USA

[2]University of Puerto Rico-Cayey, PR, USA

Running title: Tapeworm regeneration competence

* corresponding author

tania.rozario@uga.edu

500 D.W. Brooks Dr., Athens, GA 30602, USA

ORCIDS:
EMN: 0009-0002-3844-876X
LEP: 0009-0007-5836-5714
IRS: 0009-0009-1869-7292
TH: 0009-0007-1528-3807
MFC: 0009-0004-9971-0826
TR: 0000-0002-9971-5211

Keywords: regeneration, tapeworm, planaria, flatworm, Wnt signaling

Abstract

Competence to regenerate lost tissues varies widely across species. The rat tapeworm,

Hymenolepis diminuta, undergoes continual cycles of shedding and regenerating thousands of

reproductive segments to propagate the species. Despite its prowess, *H. diminuta* can only

regenerate posteriorly from a singular tissue: the neck or germinative region (GR). What cells and

signaling pathways restrict regeneration competence to the GR? In this study, we show that the

head regulates regeneration competence by promoting maintenance of the GR and inhibiting

proglottid formation in a distance-dependent manner. Anterior-posterior (A-P) patterning within

the GR provide local signals that contribute to these responses. *βcat1* is necessary for stem cell

maintenance, proliferation and proglottidization. On the other hand, *sfrp* is necessary for

maintaining the GR at its proper length. Our study demonstrates that the head organizes a balance

of pro- and anti-regeneration signals that must be integrated together and therefore control

competence to regenerate.

Summary statement

This study explores contradictory roles of the tapeworm head in providing pro- and anti-

regeneration signals, which can form a tipping point zone that is refractory to regeneration.

Introduction

Regeneration occurs over a wide range of biological scales from regeneration of axons at the cellular scale to whole body regeneration in planarians and hydra (Bely and Nyberg, 2010). Tapeworms represent a fascinating case study in regeneration. They descend from a monophyletic clade (Neodermata) exclusively populated by parasites within the Platyhelminthes phylum (Laumer et al., 2015). As a sister group to planarians, it is possible that shared regenerative abilities have enabled tapeworms to achieve their monstrous growth potential and reproductive prowess. Many species of intestinal tapeworms shed large pieces of their body to disperse the developing embryos for consumption by an intermediate host, while continuously replenishing the lost tissue. Studies with the rat tapeworm, *Hymenolepis diminuta*, demonstrated that anterior fragments regenerated reproductive segments (proglottids) after serial rounds of amputation and transplantation into new host intestines for over a decade (Goodchild, 1958; Read, 1967). This greatly prolonged the normal lifespan of *H. diminuta* suggesting that this parasite may not inherently need to age and die.

By culturing *H. diminuta in vitro*, we were able to determine its range of regenerative abilities (Rozario et al., 2019) (Fig. 1A). Head regeneration was never observed. Posterior regeneration of new proglottids occurred but only when the neck/germinative region (GR) was retained. Neither the head nor proglottids could regenerate new proglottids. Hence, the GR is the only regeneration-competent tissue. We confirmed that similar to the serial transplantation experiments described above, 2 mm anterior fragments of *H. diminuta* regenerated proglottids after serial rounds of amputation and growth *in vitro*. However, proglottid regeneration was finite when the head region was removed indicating that persistent regeneration does require head-dependent signals.

Proglottid regeneration depends on GR-resident stem cells, but this population is not functionally unique (Rozario et al., 2019). When stem cells were depleted by a lethal dose of irradiation, proglottid regeneration from the GR was inhibited. Transplantation of donor cells into irradiated GRs rescued regeneration, as long as cycling stem cells were present in the donor pool. Interestingly, the rescue occurred with donor cells from the GR as well as with donor cells from

regeneration-incompetent parts of the body. Hence, while stem cells are necessary for
regeneration, signals within the GR are paramount.

A plethora of unknown signals likely operate in the GR and control stem cell proliferation,
survival, differentiation, as well as proglottidization and competence to regenerate. Here we show
that regeneration competence is tightly linked to signals from the head. The head plays seemingly
contradictory roles as it is necessary to maintain the GR but negatively regulates proliferation and
proglottid regeneration. We find two genes typically associated with Wnt signaling, *βcatenin1* and
sfrp, as critical mediators of stem cell maintenance and GR length, respectively. Our study
demonstrates that a balance of pro- and anti-regeneration signals must be overcome to enable GR
regeneration and then initiation of proglottids. *H. diminuta* provides a fascinating example of how
extrinsic signals can promote or restrict the ability to regenerate.

Results

H. diminuta can regenerate the GR.

Anterior fragments containing the head, GR and a few immature proglottids are competent to regenerate new proglottids (Rozario et al., 2019). However, it is unclear if the GR can regenerate itself and subsequently proglottidize. After acclimating worms to *in vitro* culture conditions for 3 days, we amputated within the GR (0.5 mm) vs. outside the GR (2 mm) and grew the anterior fragments for 15 days. In both cases, proglottids regenerated (Fig. 1B). At 15 dpa, GR lengths in regenerates from both groups were comparable at 1.01±0.08 mm and 1.07±0.05 mm (Fig. 1C). The 0.5 mm fragments regenerated GRs within 2-4 days before new proglottids were added (Fig. 1D). The rate of proglottid regeneration was comparable between both fragment types, at 14-15 proglottids per day (Fig. 1D). Thus, when amputated within the GR, *H. diminuta* is capable of regenerating the posterior GR and subsequently proglottids.

The GR is patterned by differentially expressed genes along the anterior-posterior (A-P) axis (Rozario et al., 2019). If *H. diminuta* can regenerate its GR, then A-P polarized patterns should be restored following amputation. We performed whole mount *in situ* hybridization (WISH) for three posterior-enriched transcripts: *βcatenin1* (*βcat1;* WMSIL1_LOCUS14475), *frizzled5/8* (*fzd5/8;* WMSIL1_LOCUS9542), and *tumor necrosis factor receptor-16* (*tnfr16;* WMSIL1_LOCUS10339) and one anterior-enriched transcript: *secreted frizzled related protein* (*sfrp;* WMSIL1_LOCUS14856) (Fig. 1E; uncut). The antisense riboprobes used showed distinct patterns while matched sense riboprobes showed minimal background (Fig. S1). Worms were amputated within the GR (0.5 mm) and at 0 dpa, all markers were reduced or lost (Fig. 1E). A-P polarized patterns were restored by at least 10 dpa demonstrating normal GR patterning (Fig. 1E).

GR regeneration fails after serial amputation.

We investigated if GR regeneration occurs after serial amputation. Newly harvested worms were acclimated *in vitro* for 3 days, 0.5 mm anterior fragments were obtained (cut 1) and grown for 18 days before re-amputation (cut 2) (Fig. 2A). Subsets of each group were fixed to capture the variation and reproducibility of the amputations performed. We find that tissues extend slightly after heat-killing and fixation resulting in slightly longer 0 dpa fragments: 0.63±0.07 mm and 0.61±0.16 mm after each amputation (Fig. 2B; means). While fragment length did vary there was

no statistically significant difference at each timepoint. The GR lengths were longer at cut 1
(1.32±0.11 mm) than at cut 2 (1.09±0.08 mm) yet the percentage of GR length retained was close
to the target range of 50% and the mean differences were not statistically significant (48±9% after
cut 1 and 56±11% after cut 2) (Fig. 2C; means). When pooled together, there was a bias toward
increased % of GR length retained after cut 2 (Fig. 2C; all), indicating that despite our best efforts,
the average amputation position in the GR may have been slightly different between the two
timepoints. After cut 1, almost all fragments regenerated GRs and proglottids (Fig. 2D). After cut
2, many fragments failed to regenerate at all producing inconsistent regeneration rates per
experiment (Fig. 2D). Worms that did regenerate after cut 2 were significantly shorter (Fig. 2E)
and produced fewer proglottids (Fig. 2F). Thus, GR regeneration can fail under certain conditions.
126

Signals from the head regulate proglottid regeneration and GR maintenance.
When amputating within the GR, the wound site is closer to the head. We suspected that the head
asserts a negative effect on regeneration. After head amputation, regenerates exhibited increased
growth and proglottid regeneration (Fig. 3A) indicating that signals from the head do inhibit
growth. Using 1 hr uptake of the thymidine analog F-*ara*-EdU (Neef and Luedtke, 2011) to
quantify the density of proliferating cells in the GR, we found that head amputation increased
proliferation, which can explain the increased growth (Fig. 3B). In uncut worms, proliferation
density in the GR increases with distance from the head (Fig. 3C) further supporting our conclusion
that the head provides anti-proliferation signals. Upon head amputation and *in vitro* culture for 3
days, the increasing proliferation density along the GR A-P axis was no longer significant (Fig.
3D). However, the curves did not fully flatten. While signals from the head do regulate
proliferation in the GR, the effects are slow to dissipate or mediated by additional factors resident
within the GR.
140

141   Many regenerating animals respond to wounding by increasing proliferation locally. Following
142   amputation in the GR, we quantified proliferation at both anterior- and posterior-facing wounds
143   and found increased proliferation (Fig. S2). However, the local increase in proliferation was
144   equivalent regardless of head presence (Fig. S2). Thus, amputation itself does not explain the
145   increased proliferation in -head worms but could contribute to increased proliferation.
146

Previous results have shown that GR maintenance depends on the head; after head amputation, the

GR is eventually lost, the regenerates become fully proglottidized and cannot continue adding

proglottids (Rozario et al., 2019). To further characterize how the head affects proglottid

regeneration and GR maintenance, four amputation schemes were compared: + head, ½ head, -

head, and -0.5 mm anterior (Fig. 4A). Making transverse cuts through the head was technically

challenging so a sampling of ½ head fragments were fixed immediately and stained with anti-

SYNAPSIN antibodies to visualize the nervous system. Though the amount of head tissue

removed did vary, the brain was always retained (Fig. S3A: yellow arrowheads). After making the

four different amputations at the anterior region, we assayed proglottid regeneration following

serial amputation. All worms were amputated posteriorly to obtain fragments of the same length

(2 mm), that were grown *in vitro* for 14 days, then re-amputated for 3 more cycles (Fig. 4A). The

posteriors were collected to determine the number of proglottids regenerated after every cycle. As

previously reported (Rozario et al., 2019), tapeworms regenerated after serial amputation when the

head was present (Fig. 4B). The ½ head regenerates consistently displayed increased proglottid

regeneration compared to +head regenerates (Fig. 4B). Both +head and ½ head regenerates

maintained their GRs throughout 2 months of *in vitro* culture. When the head was amputated

completely (-head), proglottid regeneration eventually ceased though some regenerates still had

GRs after the 4th amputation (Fig. 4B). These -head regenerates had significantly shorter GRs (Fig.

S3B) indicating that complete proglottidization was imminent. Previously, we observed rapid loss

of the GR after head amputation, but anterior GR tissue was also removed in those experiments

(Rozario et al., 2019). In this study, we were careful to make minimal head amputations that

retained as much anterior GR and found that the GR persists for much longer than previously

reported but does eventually recede. On the other hand, fragments in which both the head and

anterior GR were removed (-0.5 mm) became fully proglottidized by the second amputation or

earlier (Fig. 4B). In sum, GRs were maintained if the brain region was retained but GRs were lost

when additional anterior tissue was removed (Fig. 4C).

The number of proglottids regenerated also depended on amputations at the head though our

analysis was complicated by GR loss in -head and -0.5mm fragments. After the first amputation,

a clear dose-dependent response was detected; when more head tissue was retained, the number of

proglottids decreased (Fig. S3C). This trend was maintained after serial rounds of amputation when

comparing +head and ½ head regenerates where GRs were not lost (Fig. S3D). After the 4th round

of amputation, the remaining head area plotted against number of proglottids for +head and ½ head

regenerates revealed a negative correlation between head tissue abundance and proglottidization

(Fig. 4D). We also measured the effect of head tissue abundance on proliferation within the GR

using F-*ara*-EdU. A general trend of increasing proliferation with decreasing head and anterior

tissue emerged though differences between +head and ½ head were not statistically significant

(Fig. 4E-F). Taken together, the head effects on maintaining the GR and regulating proliferation

can be separated. Signals from the brain region are necessary to maintain the GR whereas

proliferation and proglottidization are negatively regulated by head-dependent signals in a dose-

dependent manner.

***βcatenin1* is necessary for proglottid regeneration and stem cell maintenance.**

To begin to ascertain the molecular regulators of GR maintenance, stem cell proliferation and

proglottid regeneration rate, we turned to the Wnt signaling pathway. Wnt signaling determines

A-P polarity in many systems (Petersen and Reddien, 2009) and plays roles in regulating cell

proliferation, survival, and differentiation (Nusse and Clevers, 2017). Expression of multiple

members of the Wnt signaling pathway were either anterior-enriched or posterior-enriched by

RNA sequencing along the A-P axis of the GR (Rozario et al., 2019). As βcatenin protein is a key

downstream effector of Wnt signaling, we decided to examine its function. *H. diminuta* has three

βcatenin paralogs as do other parasitic flatworms (Montagne et al., 2019). BLASTp comparisons

of putative *H. diminuta βcatenin* paralogs with published sequences for other flatworms

(*Schmidtea mediterranea*, *Schistosoma mansoni* and *Echinococcus multilocularis*) demonstrated

that *Hdim-βcat1* (WMSIL1_LOCUS14475) is the rat tapeworm ortholog of *βcat1* (Table S1).

Expected conserved domains including multiple armadillo repeats and the DSGxxSxxx[S/T]xxxS

motif for CKI/GSK-phosphorylation necessary for targeting by the destruction complex

(Montagne et al., 2019; Valenta et al., 2012) are present (Fig. S4A). Capacity to bind α-catenin

cannot be ruled in or out at present (Fig. S4B; discussed below). Thus *Hdim-βcat1* likely mediates

Wnt signaling and may also function in cell adhesion.

In 6-day-old adults, *Hdim-βcat1* (hereafter *βcat1*) expression is largely absent in the head and

anterior GR but increases toward the posterior GR (Fig. 5A). *βcat1* is broadly expressed in many

tissues, in transverse stripes at proglottid boundaries and within accessory reproductive structures

(Fig. S5). We injected double-stranded RNA (dsRNA) targeting *βcat1* throughout the GR and

grew 2 mm anterior fragments *in vitro*. A dramatic growth defect was captured just 7 dpa/10 days

post injection (dpi) (Fig. 5B). Decreased growth was evident by 3 dpa and this RNAi treatment

was ultimately lethal. The proglottids formed were extremely small and impossible to quantify but

there was a marked decrease in worm lengths (Fig. 5C). RNAi of *βcat1* reduced transcript levels

to 8.7±2.0% (Fig. 5D). In canonical Wnt signaling, *βcat1* is a transcription factor with many targets

including *axins*, though *bona fide* targets in tapeworms have not been determined. Tapeworm *axins*

are highly divergent, but previous work using exogenous expression of AXIN1 and AXIN2 from

the fox tapeworm *E. multilocularis,* showed evidence that both AXINs function canonically in the

Wnt signaling pathway (Montagne et al., 2019). In *H. diminuta*, *axin1* (WMSIL1_LOCUS3986)

is broadly expressed in overlapping patterns with *βcat1* whereas *axin2* (WMSIL1_LOCUS6411)

is highly tissue-specific (Fig. S5). After *βcat1* RNAi, *axin1* expression is markedly reduced to

18.8±11.1% of normal levels (Fig. 5D). This suggests that canonical Wnt signaling is inhibited

following *βcat1* RNAi though other roles for *βcat1* cannot be ruled out.

Given the strong decrease in growth after *βcat1* RNAi, we hypothesized that cycling stem cells

require *βcat1*. Proliferation was significantly decreased in *βcat1* RNAi GRs compared to *GFP*

RNAi controls (Fig. 5E). WISH for two known stem cell markers in *H. diminuta*: *minichromosome*

maintenance complex component-2 (*mcm2;* WMSIL1_LOCUS11918) and *laminB receptor* (*lbr;*

WMSIL1_LOCUS1236) (Rozario et al., 2019) showed decreased expression after *βcat1* RNAi

(Fig. 5F). Thus, *βcat1* is required to maintain stem cells in the GR through regulation of

proliferation and/or other mechanisms.

GR maintenance requires *sfrp*.

We hypothesized that *βcat1* activity may be counterbalanced by anterior-localized Wnt inhibitors.

Previous RNA-sequencing analyses showed that *sfrp* expression is enriched in the anterior GR

(Rozario et al., 2019). Like other tapeworms, *H. diminuta* has one true *sfrp* family member and a

second highly divergent member, *sfrp-like* (WMSIL1_LOCUS8400), which was not investigated

further in this study. Interpro (Blum et al., 2020) predicts that Hdim-SFRP retains both frizzled

and netrin domains albeit with some variation (Fig. S6; discussed below). Since SFRP can inhibit

Wnt signaling (Leyns et al., 1997; Rattner et al., 1997; Wang et al., 1997), we pursued its potential

role in regeneration.

By WISH, *sfrp* expression is strikingly anterior-enriched with sparce expression in the head and

strong expression throughout the GR that tapers off posteriorly (Fig. 6A). In adult tapeworms, *sfrp*

expression is largely absent from the strobilated body except at lateral ends of proglottids (Fig.

S5). We hypothesized that *sfrp* RNAi would promote proglottid regeneration as opposed to *βcat*

RNAi, which reduced proglottid regeneration. We injected *sfrp* dsRNA into the head, amputated

2 mm anterior fragments, reinjected dsRNA into the head and collected worms at 10-12 dpa (Fig.

6B). Contrary to our expectations, a modest decrease in worm lengths and proglottid regeneration

were observed (Fig. 6C-D). Strikingly, the GRs were significantly shortened following *sfrp* RNAi

(Fig. 6E-F). This is the first time GR length has been affected by RNAi of any target gene. Injecting

dsRNA into the head does not preclude diffusion into the GR but the knockdown is likely to be

anteriorly biased. We injected dsRNA throughout the GR and observed the same phenotypes (Fig.

S7A-C). Both injection schemes resulted in similar *sfrp* knockdown efficacy (Fig. S7D).

Our results indicate that *sfrp* is necessary for proper GR maintenance but is unlikely to be involved

in head-dependent inhibition of proliferation or proglottidization. Instead, *sfrp* could mediate head-

dependent GR maintenance. Upon head amputation, we observed a modest 32.2±14.0% decrease

in *sfrp* expression 3 days after head amputation (Fig. S7E). By WISH, the domain of *sfrp*

expression after head amputation was often reduced as the GR shrinks though it was highly

variable at 3 dpa (Fig. S7F). Head amputation does not acutely abolish *sfrp* expression but may be

required for long-term stable expression of *sfrp* at the GR.

Decreased proglottid regeneration after *sfrp* RNAi does not support the hypothesis that *sfrp*

functions as a Wnt inhibitor. Proliferation within the GR was not significantly changed after *sfrp*

RNAi (Fig. S7G). However, the shortening of the GR would bring the posterior boundary of the

GR closer to the head. Thus, *sfrp* RNAi may indirectly exacerbate the anti-proglottidization effects

from the head.

Amputation distance from the head determines whether pro- or anti-regeneration signals dominate.

Given these results, we hypothesized that the regeneration failure we observed after serial amputation within the GR (Fig. 2) was precipitated when the posterior wound was too close to the head, which allowed the growth inhibitory effects of the head to dominate. To test this, anterior fragments were allowed to regenerate after 0.5 mm amputation within the GR and then cut a second time at 4 different distances from the head. All samples were cut at ~1mm, then subsets were progressively shaved closer toward the head (Fig. 7A; cuts A-D). Subsets of fragments were fixed at 0 dpa and we verified that cut A-D progressively retained 87±5%, 64±7%, 47±9%, and 22±4% of the GR (Fig. 7B). For each of three experiments, proglottid regeneration plummeted at either cut B or C (Fig. 7C). Groups with the lowest regeneration success corresponded to 0 dpa subsets where the posterior wound position clustered at ~60% GR length (Fig. 7C-D; matched colored arrows). This suggests that there is a tipping-point region within the GR that is refractory to regeneration following amputation. Surprisingly, the smallest fragments (cut D) regenerated equivalently to the longest fragments (cut A) (Fig. 7C-D and Fig. S7). Fragments from cut A and D achieved different absolute lengths at 18 dpa but the fold change in growth was comparable (Fig. 7E). There was no significant difference is GR length between regenerates from cut A or D (Fig. 7F). Thus, amputations within the GR can result in successful regeneration but not always, with more failed incidences when the wound site was at intermediate distances from the head. These results suggest that there are yet more signals to be uncovered that explain how regeneration is promoted and restricted in tapeworms.

Discussion

A working model for GR and proglottid regeneration.

We propose the following working model (Fig. 8) to synthesize our current findings. The head provides signals that promote and restrict regenerative ability. These two roles can be separated anatomically and through molecular regulators. Proliferation and proglottidization rate are negatively regulated by the head in a dose-dependent and distance-dependent manner with increased proliferation farther from the head. Stem cells and germ cells are the only proliferative population in flatworms (Baguñà, 2012; Bolla and Roberts, 1971; Collins et al., 2013; Ishan et al., 2025; Koziol et al., 2014) and proglottid regeneration is a stem cell-dependent process (Rozario et al., 2019). We find that *βcat1* is expressed in a posterior-biased pattern at the GR and that RNAi of *βcat1* results in loss of proliferative stem cells. We speculate that the head may produce direct or indirect inhibitor(s) of unknown posterior signals that promote *βcat1* activity. Alternatively, the head-dependent inhibitors of regeneration may act independently of *βcat1*. On the other hand, the head positively regulates regeneration by promoting GR maintenance. Without the brain-region, the GR is gradually lost and when the anterior GR is also removed, GR loss happens quickly. RNAi of *sfrp* results in shortened GRs indicating that *sfrp* contributes to GR maintenance. While *sfrp* expression is clearly anterior-enriched, it is unclear if *sfrp* expression is promoted/maintained by signals from the head.

Taken together, our model provides an explanation for the highly mixed results in regeneration outcomes when re-amputations were made within the GR. When the posterior wound site was far enough from the inhibitors in the head, regeneration could occur. When the posterior wound site was close to the head, we speculate that the targets of the head-dependent inhibitors were not present. The head could promote proper growth of the GR, and at a sufficient distance from the head, proglottids regenerated. However, when the posterior wound site was at intermediate distances from the head, the inhibitors dominated leading to failure to regenerate. We speculate that the pro- and anti-regeneration contributions of the head result in a tipping point zone that is refractory to regeneration (Fig. 8). The exact players, direct and indirect relationships, as well as additional molecular mechanisms predicted by our working model still need to be tested. Alternatives to our model certainly exist. For example, GR and proglottid regeneration may occur

322 from a wide range of posterior amputation sites with sporadic failures. These failures may be
323 caused by anterior inhibitory signals that are not always sufficient to block regeneration. Future
324 experiments will help determine if regeneration in *H. diminuta* is largely stochastic or tightly
325 regulated by spatially distinct molecular interactions.
326
Why GR regeneration did not fail significantly after the first round of amputations is unclear.
Tapeworm dimensions change when grown *in vitro*. Despite acclimating worms *in vitro* for 3 days
before all amputations, the spatial distribution and concentration of factors from the head to GR
could have been different at cut 1 vs. cut 2 (Fig. 2). GR regeneration competence does not merely
dissipate over time as the smallest fragments (cut D) successfully regenerated proglottids and the
GR after the second round of amputation (Fig. 7). The head may be acting as an organizer that sets
up a signaling environment within the GR that impacts regeneration positively and negatively in
ways that we are just beginning to discover.
335
Differing roles for *βcat1* in flatworm regeneration.
In *H. diminuta*, *βcat1* is necessary to maintain stem cell abundance and RNAi of *βcat1* results in
a dramatic inhibition of growth, proglottid regeneration and ultimately death. In the liver fluke,
Fasciola hepatica, RNAi of *Fhep-βcat1* stunted juvenile growth, decreased proliferation and
caused worm death (Armstrong et al., 2025), phenocopying our observations. Thus, roles for *βcat1*
in regulating stem cell proliferation and/or survival may be a common theme in parasitic flatworms
as it is in many metazoans and in cancers (Mohammed et al., 2016; Steinhart and Angers, 2018) .
343
In addition to canonical Wnt signaling, β-catenin proteins function in cell adhesion through linking
cadherins with the actin cytoskeleton via α-catenin in many but not all species. In *S. mediterranea,*
there is complete separation of function by two catenin paralogs: Smed-βCAT1 mediates Wnt
signaling whereas Smed-βCAT2 mediates cell adhesion (Chai et al., 2010; Su et al., 2017).
Accordingly, Smed-βCAT1 does not have α-catenin binding sites (Chai et al., 2010; Su et al.,
2017). The presence/absence of α-catenin binding sites in βCAT1 from tapeworms is more
ambiguous. Previously, a likely α-catenin binding region demarcated by 10 putative critical
residues (Aberle et al., 1996; Pokutta and Weis, 2000) within a 26 amino acid span was identified

for βCAT1 and βCAT2 in flatworms (Montagne et al., 2019). When compared to human βcatenin

(CTNB1), we and others find strong conservation of this α-catenin binding region in βCAT2 from

multiple flatworms (Fig. S4B) (Montagne et al., 2019). While Smed-βCAT1 is clearly very

divergent at this site, two tapeworms (*H. diminuta* and *E. multilocularis*) show more similarity to

the human reference (Fig. S4B) (Montagne et al., 2019). Thus, we cannot rule out the possibility

that βCAT1 in tapeworms functions in both Wnt signaling and cell adhesion, unlike their free-

living counterparts.

The role of *βcat1* in posterior identity is well conserved across many taxa (Petersen and Reddien,

2009). In planarian species that are deficient in head regeneration, knockdown of *βcat1* rescues

head regeneration indicating that failure to regenerate the head in wild type worms was due to the

inability to overcome the posteriorizing effects of *βcat1* (Liu et al., 2013; Sikes and Newmark,

2013; Umesono et al., 2013). *H. diminuta* cannot regenerate its head and we initially assumed that

tapeworm *βcat1* would be acting in a homologous fashion to planarians. Because *βcat1* is required

for stem cell maintenance in *H. diminuta,* potential roles in posteriorization could not be

uncovered. RNAi manipulations of other Wnt signaling components that are beyond the scope of

this study will help reveal a more holistic understanding of Wnt signaling in this system. However,

tapeworms and planarians do differ in that *Smed-βcat1* does not appear to affect stem cells since

after RNAi and amputation, blastemas form and there is no growth defect (Gurley et al., 2008;

Iglesias et al., 2008; Petersen and Reddien, 2008).

The differences in utilization of *βcat1* between different species of platyhelminths is potentially

very interesting. Perhaps segregating functions in posteriorization vs. stem cell maintenance in *S.*

mediterranea has enabled nimble remodeling of axes that makes this species so remarkable at

regenerating from almost any fragment. Intestinal tapeworms need the suckers at the head to attach

to the intestinal wall. Thus, maintenance of the head and proglottid regeneration may not be easily

decoupled in tapeworms. Considering the wider role for βcatenin in regulating stem cell

proliferation and survival across metazoa, it would be easy to assume that planarian phenotype is

divergent. However, studies in the acoel *Hofstenia miamia*, a sister group to all bilaterians, show

that RNAi of *Hmia-βcat1* is necessary for maintaining posterior identity but functions in regulating

stem cells, proliferation, or growth are not supported (Srivastava et al., 2014; Tewari et al., 2019).

Thus, the planarian phenotype may be ancestral. A deeper understanding of how catenin paralogs

have functionally specialized may reveal how strategies to regulate axial identity and stem cell

proliferation/survival have evolved in regeneration-competent species.

GR maintenance and the role of *sfrp*.

The mechanism by which *sfrp* maintains the GR is unknown. While *sfrp* is primarily known as a

Wnt antagonist, it can bind and stimulate frizzled receptors, interact with other signaling pathways

and modulate cell-matrix interactions by binding integrins (Bovolenta et al., 2008; Mii and Taira,

2011). There is evidence that *sfrps* can stimulate Wnt and BMP signaling in the zebrafish retina

(Holly et al., 2014). In *Drosophila*, *sfrps* can modulate Wnt signaling in a biphasic manner:

inhibiting Wnt signaling when expressed at high levels while promoting Wnt signaling when

expressed at low levels (Üren et al., 2000). In *H. diminuta*, the *sfrp* RNAi phenotypes were not the

opposite of the *βcat1* RNAi phenotypes as one might predict if *sfrp* and *βcat1* were acting solely

as Wnt antagonist and Wnt effector respectively. In many flatworms, Wnt ligands are secreted

from both anterior and posterior domains (Armstrong et al., 2025; Koziol et al., 2016; Reddien,

2018; Soria et al., 2020). Expression of *sfrp* throughout the GR may be required as a permissive

signal to tune Wnt signaling to a moderately low level so that it can be acted upon differentially

by other modulators. It is also possible that *sfrp* modulates *βcat1* activity differentially in a

tissue/cell-specific manner that we have yet to uncover. Alternatively, *sfrp* may function

independently of Wnt signaling. Future studies investigating RNAi phenotypes of other

antagonists, ligands and frizzled receptors will help resolve if/how other Wnt signaling components

regulate GR maintenance.

Biochemical evidence for parasitic flatworm SFRP proteins as Wnt inhibitors has not been

confirmed though both frizzled and netrin domains are readily identifiable. There are conflicting

assertions as to whether all 6 conserved cysteine residues typically found in the netrin domain are

present (Armstrong et al., 2025; Koziol et al., 2016; Riddiford and Olson, 2011). We detect all 6

cysteine residues, but the 6th residue is displaced by three animo acids in flatworms, including free-

living *S. mediterranea,* but not humans (Fig. S6). *Smed-sfrp1* is generally accepted as a Wnt

inhibitor (Doddihal et al., 2024; Hill and Petersen, 2018; Reuter et al., 2015; Scimone et al., 2016;

Stückemann et al., 2017; Sureda-Gómez et al., 2015). Thus, it remains possible that a muted or

more ambiguous role in Wnt signaling by *sfrp1* is a shared feature in free-living and parasitic

flatworms.

Rationales for A-P polarized signals regulating regeneration.

Our work has revealed that a balance of opposing signals at the GR can determine regeneration

competence. It makes intuitive sense that initiation of proglottids must be inhibited close to the

head if the GR is to be maintained as an unsegmented tissue. A recent study from the mouse bile

duct tapeworm, *Hymenolepis microstoma,* has described discrete expression of Wnt and Hedgehog

signaling components in a signaling quartet (SQ) within the transition zone from GR to fully

elaborated proglottids (Jarero et al., 2024). It is possible the SQ initiates proglottid formation and

represents the unknown targets of head-dependent inhibitors predicted by our study. How the SQ

responds to head-dependent signals and vice versa is currently unknown and warrants future

investigation.

Our study also shows that regulation of proliferation dynamics is dependent on signals from the

head. While amputation-induced proliferation does occur in *H. diminuta*, the enduring increase in

proglottid regeneration after ½ head removal and dose-dependent effects of head tissue abundance

on proliferation all suggest that the head provides signals that downregulate proliferation. What

purpose do anti-proliferation signals serve in the GR? One intriguing possibility is that signals

from the head could differentially regulate stem cell potency, self-renewal, asymmetric division or

other cell cycle dynamics. There is already evidence for different cycling rates in stem cells of

parasitic flatworms (Herz et al., 2024; Wang et al., 2018). Future investigation will help us

understand how head-dependent signals may influence stem cell behaviors.

In short, the head and GR constitute a signaling environment that influences regeneration

competence. Furthermore, the anterior GR contains a minimal cohort of stem cells and extrinsic

signals that are required to regenerate the GR and seed new proglottids. Understanding these

signals could enrich our knowledge of how stem cells and regeneration are regulated in parasitic

flatworms and beyond.

Study limitations

Our study relies on RNAi, which cannot eliminate gene expression. We also use *in vitro* culture conditions that necessarily deviate from conditions *in vivo*. How environmental signals influence our results is unknown. We have also only explored the contributions of two genes and tapeworm regeneration is likely to be a complex process involving interconnecting networks of interactions.

Materials and Methods

<u>Animal care and use</u>

The life cycle of *H. diminuta* was maintained in house using mealworm beetles (*Tenebrio molitor*) and Sprague-Dawley rats for the larval and adult stages respectively. Rats were fed 200-400 infective cysticercoids by oral gavage in ~0.5 mL 0.85% NaCl. Typically, adult tapeworms were collected 6 days post infection by euthanizing infected rats in a CO_2 chamber and flushing the small intestine with 1X Hanks Balanced Salt Solution (HBSS: 21-023-CV; Corning). Rodent care was performed in accordance with protocols approved by the Institutional Animal Care and Use Committee (IACUC) of the University of Georgia (A2023 10-019-Y1-A0).

<u>*In vitro* parasite culture</u>

Tapeworms were harvested 6 days post-infection and acclimated to *in vitro* culture conditions for 3 days before any amputations were made. We used biphasic cultures as described before (Rozario et al., 2019). 2.5X MEM essential amino acids (M5550; Sigma-Aldrich) was added to the liquid culture: Working Hanks 4 [WH4: 1X HBSS (MT21023CV; Corning), 4 g/L glucose (158968; Sigma-Aldrich), 1X antibiotic-antimycotic (15240062; Fisher)]. This ameliorated inconsistencies between batches of blood used for culturing but was not strictly necessary.

<u>Cloning genes for riboprobe and dsRNA synthesis</u>

Cloning was performed as previously described (Collins et al., 2010). All primers and accessions are described in Table S2. Plasmids are available upon request. Briefly, adult *H. diminuta* cDNA was used to PCR amplify products that were ligated into pJC53.2 plasmid via TA cloning. Riboprobes for *in situ* hybridization were generated by *in vitro* transcription reactions with either T3 or SP6 RNA polymerase (Ishan et al., 2025) . For RNAi, dsRNA was generated as previously described (Rouhana et al., 2013) using T7 RNA polymerase.

<u>*In situ* hybridization and other staining</u>

Heat-kills were performed by swirling tapeworms in 75°C water for a few seconds before fixing (4% formaldehyde in Phosphate Buffered Saline with 0.3% TritonX-100 (PBSTx)) for 30 min-2 hr at room temperature or overnight at 4°C. Worms were either dehydrated into methanol and

stored (-30°C) or used directly for staining. For simple phenotyping of growth and regeneration,

worms were stained in 1 μg/mL DAPI (D9542; Sigma-Aldrich)/PBSTx overnight at 4°C and

cleared in mounting solution (80% glycerol/10 mM Tris pH7.5/ 1 mM EDTA) overnight before

mounting and imaging. Previously published methods were employed for *in situ* hybridization

(Rozario et al., 2019) and immunostaining (Rozario and Newmark, 2015). Anti-SYN antibodies

(3C11; Developmental Studies Hybridoma Bank) were used at 1:200 with Tyramide Signal

Amplification (TSA).

<u>F-*ara*-EdU uptake and staining</u>

F-*ara*-EdU pulses were performed in 0.1 μM F-*ara*-EdU (T511293; Sigma-Aldrich)/1%

DMSO/WH4 at 37°C for 1 hr. Staining was performed as previously described (Ishan et al., 2025)

with the following specifics: 1) 15 min proteinase-K digestion (10 μg/mL in 0.1% SDS/PBSTx)

and 2) 10 min TSA reaction. F-*ara*-EdU+ cells were quantified using Imaris (Oxford Instruments)

spot finder from 3D confocal z-stacks. Automatic spot discovery (with background subtraction)

was enabled then manually adjusted to label all visible spots without false positives. All F-*ara*-

EdU data is represented as densities by normalizing to the area of the quantified fields.

<u>Imaging and image processing</u>

Confocal imaging was performed on a Zeiss LSM 900 with the following objectives: 20X/Plan-

Apochromat/0.8 M27/FWD=0.55mm and 63X/Plan-Apochromat/1.40 Oil DIC M27. WISH and

DAPI-stained worms were imaged on a Zeiss AxioZoom V16 Microscope. Fiji (Schindelin et al.,

2012) was used for brightness/contrast adjustments, maximum-intensity projections and

measurements of lengths, areas, and numbers of proglottids.

<u>RNAi</u>

Knockdowns were performed as previously described (Rozario et al., 2019). Tapeworms were

microinjected with dsRNA (1-1.5 μg/μl in HBSS), 2 mm anterior fragments were amputated 3

days later and cultured *in vitro*. Injections were performed using a Femtojet 4i (Eppendorf) at 500

hPa for 0.3-1s. Injection of *GFP* dsRNA (Roberts-Galbraith et al., 2016) was used as a negative

control. Experimenters were blinded to the dsRNA used throughout microinjections, staining and

image analysis.

<u>qRT-PCR</u>

To exclude genomic DNA, primers were designed to flank introns or span an intron-exon junction.

Amplicon range was 133-277 bp. All primers were tested on cDNA +/- reverse transcriptase and

gel electrophoresis to confirm the presence of a single product. Primer efficiencies were calculated

from 2-fold serial dilutions of cDNA. All primers are reported in Table S2.

For RNA extractions, 3 mm anteriors from 4-5 worms were cut, submerged in 200 µL TRIzol

Reagent (15596026; Invitrogen) and processed according to manufacturer's instructions with these

exceptions: 1) samples were homogenized with a motorized pestle twice in a semi-solid state then

centrifuged at >12K RCF for 5 min to obtain 180 µL clean supernatant, 2) RNA pellets were

washed twice with 75% ethanol-DEPC and 3); pellets were resuspended in 40 µL nuclease-free

water. DNAse (M6101; Promega) treatment was performed for 1 hr at 37°C. RNA was cleaned

using RNA Clean & Concentrator-5 kit (R1016; Zymo Research) and the concentration measured

on a Nanodrop spectrophotometer. cDNA synthesis was performed, according to manufacturer's

protocol, using Oligo(dT)$_{20}$ primers and iScript Select cDNA Synthesis Kit (1708897; Bio-Rad)

with undiluted RNA. cDNA was further diluted with 5-25 µL nuclease-free water depending on

the number of reactions needed.

qRT-PCR was performed using GoTaq MasterMix (A6001; Promega) according to manufacturer's

instructions and scaled to 25 µL reactions with 2 µL cDNA template for each of 3 technical

replicates. Each reaction had 0.4 µM primer pairs, 0.3 µM CXCR in 1X master mix. 7500 Real

Time PCR System (Applied Biosystems) was used for 40 amplification cycles: 15 s at 95°C and 1

min at 60°C. Relative gene expression change was calculated using the PFAFFL equation

(Hellemans et al., 2007; Vandesompele et al., 2002) to account for different primer efficiencies.

Internal normalization was done on the geometric mean of two previously published endogenous

controls: *60S ribosomal protein L13* (*60Srpl13;* WMSIL1_LOCUS3984) and *myosin heavy chain*

(*mhc;* WMSIL1_LOCUS8394) (Rozario et al., 2019) .

<u>Statistical Analysis</u>

Graph Pad Prism 10 was used for all statistical analyses. All experiments were repeated at least
twice. Error bars, statistical tests, number of replicates (N) and sample sizes (n) are indicated in
corresponding figure legends. P-values: ns= not significant, * = $p<0.05$, **=$p<0.01$, ***=$p<0.001$,
****=$p<0.0001$.

Acknowledgements

We are sincerely thankful to Phil Newmark and Melanie Issigonis (Morgridge Institute for

Research, WI). Phil's unwavering support in the initialization of these projects and continued

advice was instrumental in our success. We are also grateful for Melanie's critical evaluation of

our manuscript and valuable feedback. We thank the Biomedical Microscopy Core at the

University of Georgia, Athens, GA for necessary microscopy services.

Competing interests

No conflicts to report.

Funding

This work was funded by a National Institute of Allergy and Infectious Disease (NIAID) grant

(DP2 AI 154416-01) to Tania Rozario.

Data and resource availability

All relevant data and resources can be found within the article and its supplementary information.

References

Aberle, H., Schwartz, H., Hoschuetzky, H. and Kemler, R. (1996). Single Amino Acid Substitutions in Proteins of the armadillo Gene Family Abolish Their Binding to α-Catenin*. *J. Biol. Chem.* **271**, 1520–1526.

Armstrong, R., Marks, N. J., Geary, T. G., Harrington, J., Selzer, P. M. and Maule, A. G. (2025). Wnt/β-catenin signalling underpins juvenile *Fasciola hepatica* growth and development. *PLOS Pathog.* **21**, e1012562.

Baguñà, J. (2012). The planarian neoblast: the rambling history of its origin and some current black boxes. *Int J Dev Biol.* **56**, 19–37.

Bely, A. E. and Nyberg, K. G. (2010). Evolution of animal regeneration: re-emergence of a field. *Trends Ecol Evol.* **25**, 161–170.

Blum, M., Chang, H.-Y., Chuguransky, S., Grego, T., Kandasaamy, S., Mitchell, A., Nuka, G., Paysan-Lafosse, T., Qureshi, M., Raj, S., et al. (2020). The InterPro protein families and domains database: 20 years on. *Nucleic Acids Res.* **49**, D344–D354.

Bolla, R. I. and Roberts, L. S. (1971). Developmental physiology of cestodes. IX. Cytological characteristics of the germinative region of *Hymenolepis diminuta*. *The Journal of parasitology* **57**, 267–277.

Bovolenta, P., Esteve, P., Ruiz, J. M., Cisneros, E. and Lopez-Rios, J. (2008). Beyond Wnt inhibition: new functions of secreted Frizzled-related proteins in development and disease. *J. Cell Sci.* **121**, 737–746.

Chai, G., Ma, C., Bao, K., Zheng, L., Wang, X., Sun, Z., Salò, E., Adell, T. and Wu, W. (2010). Complete functional segregation of planarian β-catenin-1 and -2 in mediating Wnt signaling and cell adhesion. *The Journal of biological chemistry* **285**, 24120–24130.

Collins, J. J., Hou, X., Romanova, E. V., Lambrus, B. G., Miller, C. M., Saberi, A., Sweedler, J. V. and Newmark, P. A. (2010). Genome-wide analyses reveal a role for peptide hormones in planarian germline development. *PLoS biology* **8**, e1000509.

Collins, J. J., Wang, B., Lambrus, B. G., Tharp, M. E., Iyer, H. and Newmark, P. A. (2013). Adult somatic stem cells in the human parasite *Schistosoma mansoni*. *Nature* **494**, 476–479.

Doddihal, V., Mann, F. G., Ross, E. J., McKinney, M. C., Guerrero-Hernández, C., Brewster, C. E., McKinney, S. A. and Alvarado, A. S. (2024). A PAK family kinase and the Hippo/Yorkie pathway modulate WNT signaling to functionally integrate body axes during regeneration. *Proc. Natl. Acad. Sci.* **121**, e2321919121.

Goodchild, C. G. (1958). Transfaunation and Repair of Damage in the Rat Tapeworm, *Hymenolepis diminuta*. *J. Parasitol* **44**, 345–351.

Gurley, K. A., Rink, J. C. and Alvarado, A. S. (2008). β-catenin defines head versus tail identity during planarian regeneration and homeostasis. *Science* **319**, 323–327.

Hellemans, J., Mortier, G., Paepe, A. D., Speleman, F. and Vandesompele, J. (2007). qBase relative quantification framework and software for management and automated analysis of real-time quantitative PCR data. *Genome Biol.* **8**, R19.

Herz, M., Zarowiecki, M., Wessels, L., Pätzel, K., Herrmann, R., Braun, C., Holroyd, N., Huckvale, T., Bergmann, M., Spiliotis, M., et al. (2024). Genome-wide transcriptome analysis of *Echinococcus multilocularis* larvae and germinative cell cultures reveals genes involved in parasite stem cell function. *Front. Cell. Infect. Microbiol.* **14**, 1335946.

Hill, E. M. and Petersen, C. P. (2018). Positional information specifies the site of organ regeneration and not tissue maintenance in planarians. *eLife* **7**, e02238.

Holly, V. L., Widen, S. A., Famulski, J. K. and Waskiewicz, A. J. (2014). Sfrp1a and Sfrp5 function as positive regulators of Wnt and BMP signaling during early retinal development. *Dev. Biol.* **388**, 192–204.

Iglesias, M., Gomez-Skarmeta, J. L., Saló, E. and Adell, T. (2008). Silencing of *Smed-βcatenin1* generates radial-like hypercephalized planarians. *Development* **135**, 1215–1221.

Ishan, M., Skipper, I. R. and Rozario, T. (2025). Combined fluorescent *in situ* hybridization and F-ara-EdU staining on whole mount *Hymenolepis diminuta*. *Biol. Methods Protoc.* bpaf011.

Jarero, F., Baillie, A., Riddiford, N., Montagne, J., Koziol, U. and Olson, P. D. (2024). Muscular remodeling and anteroposterior patterning during tapeworm segmentation. *Dev. Dyn.* **253**, 998–1023.

Koziol, U., Rauschendorfer, T., Rodríguez, L. Z., Krohne, G. and Brehm, K. (2014). The unique stem cell system of the immortal larva of the human parasite *Echinococcus multilocularis. EvoDevo* **5**, 10.

Koziol, U., Jarero, F., Olson, P. D. and Brehm, K. (2016). Comparative analysis of Wnt expression identifies a highly conserved developmental transition in flatworms. *BMC biology* **14**, 10.

Laumer, C. E., Hejnol, A. and Giribet, G. (2015). Nuclear genomic signals of the "microturbellarian" roots of platyhelminth evolutionary innovation. *eLife* **4**, e05503.

Leyns, L., Bouwmeester, T., Kim, S.-H., Piccolo, S. and Robertis, E. M. D. (1997). Frzb-1 Is a Secreted Antagonist of Wnt Signaling Expressed in the Spemann Organizer. *Cell* **88**, 747–756.

Liu, S. Y., Selck, C., Friedrich, B., Lutz, R., Vila-Farré, M., Dahl, A., Brandl, H.,
Lakshmanaperumal, N., Henry, I. and Rink, J. C. (2013). Reactivating head regrowth in a
regeneration-deficient planarian species. *Nature* **500**, 81–84.

Mii, Y. and Taira, M. (2011). Secreted Wnt "inhibitors" are not just inhibitors: Regulation of
extracellular Wnt by secreted Frizzled-related proteins. *Dev., Growth Differ.* **53**, 911–923.

Mohammed, M. K., Shao, C., Wang, J., Wei, Q., Wang, X., Collier, Z., Tang, S., Liu, H.,
Zhang, F., Huang, J., et al. (2016). Wnt/β-catenin signaling plays an ever-expanding role in
stem cell self-renewal, tumorigenesis and cancer chemoresistance. *Genes Dis.* **3**, 11–40.

Montagne, J., Preza, M., Castillo, E., Brehm, K. and Koziol, U. (2019). Divergent Axin and
GSK-3 paralogs in the beta-catenin destruction complexes of tapeworms. *Dev. Genes. Evol.*
11, 1–14.

Neef, A. B. and Luedtke, N. W. (2011). Dynamic metabolic labeling of DNA in vivo with
arabinosyl nucleosides. *P.N.A.S.* **108**, 20404–20409.

Nusse, R. and Clevers, H. (2017). Wnt/β-Catenin Signaling, Disease, and Emerging
Therapeutic Modalities. *Cell* **169**, 985–999.

Petersen, C. P. and Reddien, P. W. (2008). *Smed-βcatenin-1* is required for anteroposterior
blastema polarity in planarian regeneration. *Science* **319**, 327–330.

Petersen, C. P. and Reddien, P. W. (2009). Wnt signaling and the polarity of the primary body
axis. *Cell* **139**, 1056–1068.

Pokutta, S. and Weis, W. I. (2000). Structure of the Dimerization and β-Catenin- Binding
Region of α-Catenin. *Mol. Cell* **5**, 533–543.

Rattner, A., Hsieh, J.-C., Smallwood, P. M., Gilbert, D. J., Copeland, N. G., Jenkins, N. A.
and Nathans, J. (1997). A family of secreted proteins contains homology to the cysteine-rich
ligand-binding domain of frizzled receptors. *Proc. Natl. Acad. Sci.* **94**, 2859–2863.

Read, C. P. (1967). Longevity of the tapeworm, *Hymenolepis diminuta*. *J Parasitol* **53**, 1055–6.

Reddien, P. W. (2018). The Cellular and Molecular Basis for Planarian Regeneration. *Cell* **175**,
327–345.

Reuter, H., März, M., Vogg, M. C., Eccles, D., Grífol-Boldú, L., Wehner, D., Owlarn, S.,
Adell, T., Weidinger, G. and Bartscherer, K. (2015). β-Catenin-Dependent Control of
Positional Information along the AP Body Axis in Planarians Involves a Teashirt Family
Member. *Cell Rep.* **10**, 253–265.

Riddiford, N. and Olson, P. D. (2011). Wnt gene loss in flatworms. *Dev. Genes. Evol.* **221**,
187–197.

Roberts-Galbraith, R. H., Brubacher, J. L. and Newmark, P. A. (2016). A functional
genomics screen in planarians reveals regulators of whole-brain regeneration. *eLife* **5**,
a000505.

Rouhana, L., Weiss, J. A., Forsthoefel, D. J., Lee, H., King, R. S., Inoue, T., Shibata, N.,
Agata, K. and Newmark, P. A. (2013). RNA interference by feeding in vitro-synthesized
double-stranded RNA to planarians: methodology and dynamics. *Dev. Dyn.* **242**, 718–730.

Rozario, T. and Newmark, P. A. (2015). A confocal microscopy-based atlas of tissue
architecture in the tapeworm *Hymenolepis diminuta*. *Exp. Parasitol.* **158**, 31–41.

Rozario, T., Quinn, E. B., Wang, J., Davis, R. E. and Newmark, P. A. (2019). Region-
specific regulation of stem cell-driven regeneration in tapeworms. *eLife* **8**, e48958.

Schindelin, J., Arganda-Carreras, I., Frise, E., Kaynig, V., Longair, M., Pietzsch, T.,
Preibisch, S., Rueden, C., Saalfeld, S., Schmid, B., et al. (2012). Fiji: an open-source
platform for biological-image analysis. *Nat. Methods* **9**, 676–682.

Scimone, M. L., Cote, L. E., Rogers, T. and Reddien, P. W. (2016). Two FGFRL-Wnt circuits
organize the planarian anteroposterior axis. *eLife* **5**, 905.

Sikes, J. M. and Newmark, P. A. (2013). Restoration of anterior regeneration in a planarian
with limited regenerative ability. *Nature* **500**, 77–80.

Soria, C. L. D., Lee, J., Chong, T., Coghlan, A., Tracey, A., Young, M. D., Andrews, T.,
Hall, C., Ng, B. L., Rawlinson, K., et al. (2020). Single-cell atlas of the first intra-
mammalian developmental stage of the human parasite *Schistosoma mansoni*. *Nat. Commun.*
11, 6411.

Srivastava, M., Mazza-Curll, K. L., Wolfswinkel, J. C. van and Reddien, P. W. (2014).
Whole-Body Acoel Regeneration Is Controlled by Wnt and Bmp-Admp Signaling. *Curr.*
Biol. **24**, 1107–1113.

Steinhart, Z. and Angers, S. (2018). Wnt signaling in development and tissue homeostasis.
Development **145**, dev146589.

Stückemann, T., Cleland, J. P., Werner, S., Vu, H. T.-K., Bayersdorf, R., Liu, S.-Y.,
Friedrich, B., Jülicher, F. and Rink, J. C. (2017). Antagonistic Self-Organizing Patterning
Systems Control Maintenance and Regeneration of the Anteroposterior Axis in Planarians.
Dev. Cell **40**, 248-263.e4.

Su, H., Sureda-Gomez, M., Rabaneda-Lombarte, N., Gelabert, M., Xie, J., Wu, W. and
Adell, T. (2017). A C-terminally truncated form of β-catenin acts as a novel regulator of
Wnt/β-catenin signaling in planarians. *PLoS Genet.* **13**, e1007030.

Sureda-Gómez, M., Pascual-Carreras, E. and Adell, T. (2015). Posterior Wnts Have Distinct
Roles in Specification and Patterning of the Planarian Posterior Region. *Int J Mol Sci.* **16**,
26543–26554.

Tewari, A. G., Owen, J. H., Petersen, C. P., Wagner, D. E. and Reddien, P. W. (2019). A
small set of conserved genes, including *sp5* and *Hox*, are activated by Wnt signaling in the
posterior of planarians and acoels. *PLoS Genet.* **15**, e1008401.

Umesono, Y., Tasaki, J., Nishimura, Y., Hrouda, M., Kawaguchi, E., Yazawa, S.,
Nishimura, O., Hosoda, K., Inoue, T. and Agata, K. (2013). The molecular logic for
planarian regeneration along the anterior-posterior axis. *Nature* **500**, 73–76.

Üren, A., Reichsman, F., Anest, V., Taylor, W. G., Muraiso, K., Bottaro, D. P.,
Cumberledge, S. and Rubin, J. S. (2000). Secreted Frizzled-related Protein-1 Binds Directly
to Wingless and Is a Biphasic Modulator of Wnt Signaling*. *J. Biol. Chem.* **275**, 4374–4382.

Valenta, T., Hausmann, G. and Basler, K. (2012). The many faces and functions of β-catenin.
EMBO J. **31**, 2714–2736.

Vandesompele, J., Preter, K. D., Pattyn, F., Poppe, B., Roy, N. V., Paepe, A. D. and
Speleman, F. (2002). Accurate normalization of real-time quantitative RT-PCR data by
geometric averaging of multiple internal control genes. *Genome Biol.* **3**, research0034.1.

Wang, S., Krinks, M., Lin, K., Luyten, F. P. and Moos, M. (1997). Frzb, a Secreted Protein
Expressed in the Spemann Organizer, Binds and Inhibits Wnt-8. *Cell* **88**, 757–766.

Wang, B., Lee, J., Li, P., Saberi, A., Yang, H., Liu, C., Zhao, M. and Newmark, P. A.
(2018). Stem cell heterogeneity drives the parasitic life cycle of *Schistosoma mansoni. eLife*
7, e35449.

 Figure legends

Fig. 1. The germinative region (GR) regenerates after amputation. (A) Schematic of basic *H.*
diminuta anatomy and known regenerative ability. Head regeneration does not occur but
proglottids regenerate from fragments that retain the GR. Without the head region, proglottid
regeneration initially occurs but ceases. (B) DAPI-stained anterior fragments cut within the GR
(0.5 mm) or outside of it (2 mm) at 15 dpa. Quantification of worm length and proglottids
regenerated at 15 dpa. N= 3, n= 23, 27; t-test. (C) DAPI-stained 0.5 mm anterior regenerate at 15
dpa. Yellow arrowhead marks the first proglottid. Quantification of GR lengths from B; t-test. (D)
Quantification of proglottids regenerated every 2 days. Linear regression fitted; n= 50, 55. (E)
WISH for transcripts with A-P polarized expression patterns 3 days after acclimation *in vitro*
(uncut) then amputated at 0.5 mm and stained at 0 dpa and 10 dpa. Anterior toward the left.
Asterisks mark pigmented debris. Error bars= SD.

Fig. 2. Serial amputation within the GR negatively affects regenerative ability. (A) Scheme
for serial amputation within the GR. (B-C) Quantification of GR lengths and percentage of GR
length retained compared to GR lengths of uncut worms at 0 dpa per timepoint. Black squares=
means, gray circles= individual worms; N= 3, n= 29, 23; t-test. (D) Quantification of regeneration
success rate with F-test to compare variances (##: p=0.0015). (E) DAPI-stained regenerates 18
dpa. (F) Quantification of proglottids from worms that regenerated after cut 2. N= 3; n= 64, 25; t-
test. Error bars= SD.

Fig. 3. The head inhibits proglottid regeneration and cell proliferation. (A) DAPI-stained
regenerates from 0.5 mm amputations +/- head at 15 dpa. Quantification of proglottids regenerated
from a representative experiment. n= 7; t-test. (B) Confocal maximum intensity projections of +/-
head worms at 3 dpa immunostained for neuronal marker SYN after a 1 hr pulse of F-*ara*-EdU to
label proliferating cells. Quantification of proliferation density within the GR from a representative
experiment. Each sample was normalized to the mean proliferation density in the corresponding
+head group. n= 10, 12; t-test. (C-D) Confocal maximum intensity projections after 1 hr pulse of
F-*ara*-EdU +/- head at 3 dpa. Yellow dotted lines represent 200 μm-wide regions used for

quantification of proliferation density. H= head. N= 3, n= 11, 13; one-way ANOVA with Dunnett's
multiple comparison test. Error bars= SD.

Fig. 4. Effect of head or head-adjacent amputations on proliferation and regeneration. (A)
Scheme for results shown in B-C and Fig. S3. Step 1: Four amputation schemes tested: intact head
(+head), transverse cut through half the head while retaining the brain (½ head), head removal (-
head) and cut 0.5 mm from the anterior into the GR (-0.5 mm). Step 2: all worms were amputated
posteriorly to obtain 2 mm fragments. Step 3: fragments were grown *in vitro* for 14 days. Step 4-
5: regenerates were recut to obtain 2 mm anterior fragments that were cultured *in vitro* while the
posteriors were collected to measure proglottid regeneration. (B) Quantification of proglottids
regenerated. All samples were normalized to the mean of +head worms set to 100%. Bars indicate
means for all samples after cut 1 and only for samples with visible GRs for cut 2-4. Proglottidized
worms without GRs= red dots, n= 27, 37, 26, 19 by scheme at start. (C) Percentage of regenerates
that maintained a visible GR from B. (D) After cut 4, proglottids regenerated from +head and ½
head samples from one experiment plotted against head area remaining measured from DAPI-
stained worms. Linear regression fitted; n= 8, 12. (E-F) Quantification of F-*ara*-EdU+ cells
normalized to GR area at 3 dpa. All samples were normalized to the mean of +head worms set to
100%. N= 4, 3, 4, 3 by scheme; two-way ANOVA (mixed model) and Tukey's multiple
comparison test (E). All pooled data from E analyzed with one-way ANOVA compared to +head
with Dunnett's multiple comparison test; n= 26, 28, 26, 21 by scheme. Error bars= SD.

Fig. 5. *βcat1* is a posterior-biased factor that is necessary to maintain the cycling stem cell
population in the GR. (A) Schematic and WISH of *βcat1* expression at the GR from 6-day-old
adult. (B) RNAi scheme and DAPI-stained worms at 7 dpa. (C) Quantification of worm lengths
after RNAi at 7 dpa. Bars= means, gray circles= individual worms; N= 3, n= 24, 24, 23; one-way
ANOVA with Dunnett's multiple comparison test. (D) qRT-PCR for target gene expression (top)
after *βcat1* RNAi compared to *GFP* RNAi (set at 1). (E) Representative maximum intensity
confocal projections at the GR after 1 hr F-*ara*-EdU uptake. Dotted yellow lines= worm outlines.
Quantification of F-*ara*-EdU+ cells normalized to area from 800 μm-wide cropped region
posterior to the head at 6-7 dpi. All samples were normalized to the mean of *GFP* RNAi worms
set to 100%. Bars= means, gray circles= individual worms; N= 3, n= 20, 19, 16; one-way ANOVA

with Dunnett's multiple comparison test. (F) WISH for cycling stem cell markers *mcm2* and *lbr* at

6 dpi; N= 3. Error bars= SD.

Fig. 6. *sfrp* is an anterior-biased factor that regulates GR length. (A) Schematic and WISH of

sfrp expression at the GR from 6-day-old adult. (B) RNAi scheme for C-F. (C) Representative

DAPI-stained regenerates after RNAi. (D) Quantification of proglottids regenerated after RNAi

normalized to mean from *GFP* RNAi set to 100%. (E) Representative DAPI-stained images

showing GR lengths (between yellow arrowheads). (F) Quantification of GR lengths from D. For

D, F: Bars= means, gray circles= individual worms; N= 3, n = 26, 25, 25; one-way ANOVA with

Dunnett's multiple comparison test; error bars= SD.

Fig. 7. How amputations at increasing distance from the head influence competence to

regenerate. (A) Regenerates from 0.5 mm anterior fragments grown for 18 days were re-

amputated at 1 mm (cut A). A subset was fixed to quantify GR length and a subset was grown *in*

vitro for 18 days. The remaining fragments were recut to shave off ~20-25% GR (cut B) and the

process repeated successively for cut C and cut D. Darkfield images showing live fragments before

in vitro culture. (B) GR lengths measured from DAPI-stained 0 dpa fragments represented as

percentage of GR length relative to uncut worms from the same cohort. N=3, n= 20, 21, 21, 21;

two-way ANOVA (mixed model) with Tukey's multiple comparison test. (C) Percentage of

fragments that regenerated proglottids after 18 days. N= 3, n= 27, 23, 21, 22 at start. Each

experiment is represented by the same color in B and C. Colored arrows point to matched sets

representing groups with lowest regeneration success. (D-F) Quantification of worm length (D),

growth fold change (E) and GR length (F). N= 3, n= 20, 8, 10, 17; t-test. Error bars= SD.

Fig. 8. Working model for regeneration in *H. diminuta*. Overlapping regions at the head (blue

bars) play two contrasting roles: proliferation and proglottidization rate are negatively regulated

whereas GR maintenance is positively regulated. *βcat1* expression is posteriorly biased and *βcat1*

is necessary to maintain stem cells in the GR, which are required for proglottid regeneration.

Expression of *sfrp* is anteriorly biased and *sfrp* is necessary to maintain GR length. Signals that

may link the head/brain/anterior GR to *βcat1* and *sfrp* expression or activity remain to be

identified. Serial amputation within the GR revealed increased failure to regenerate the GR and

proglottids (yellow X) at intermediate distances forming a tipping point zone that is refractory to

regeneration (black shading). Both smaller and larger fragments amputated outside of the black

region regenerated equivalently (green check).

Fig. S1. Specificity of antisense riboprobes. WISH with antisense and sense riboprobes for

transcripts in Fig. 1E.

Fig. S2. Proliferation response at wound site. Worms were amputated at 0.5 mm within the GR.

Proliferation density was quantified from a 100 μm-wide box at the posterior-facing wound (Pfw)

and anterior-facing wound (Afw) at 1 dpa after 1 hr pulse of F-*ara*-EdU. Quantification from one

representative experiment, n= 6, 5, 6, 7; error bars= SD, one-way ANOVA with Tukey's multiple

comparison test.

Fig. S3. How head tissues influence regeneration. (A) Widefield fluorescent images of ½ head

fragments at 0 dpa stained with anti-SYN antibodies. Yellow arrowheads point to the

brain/cephalic ganglia. (B) GR lengths at the end of a representative serial amputation experiment.

n= 5, 8, 8; one-way ANOVA with Dunnett's multiple comparison test. (C) Quantification of

proglottids regenerated after cut 1, normalized to +head means. N= 3; n= 27, 37, 26; statistical

significance by one-way ANOVA and was the same for all 3 experiments. (D) Quantification of

proglottids regenerated normalized to +head means. Cut 2: N= 3, n= 27, 34; Cut 3: N= 2, n= 11,

16; Cut 4: N= 2, n= 13, 15; t-tests. Error bars= SD.

Fig. S4. Domain analysis of *H. diminuta* βCAT1. (A) Sequence alignment of βCAT1 from four

flatworms (*Hd: H. diminuta, Em: E. multilocularis, Sm: S. mansoni, Smed: S. mediterranea*). The

conserved DSGxxSxxx[S/T]xxxS motif for CKI/GSK-phosphorylation (red) is present as are

armadillo repeats (lilac). The boundaries of both domains are shown according to *S. mediterranea*

(Su et al., 2017). Shading indicates identical residues between 4 species (gray) or 3 species (blue).

The N- and C-termini are more divergent between all species. (B) Sequence alignment of 26 amino

acids covering a putative α-catenin binding domain (Montagne et al., 2019). 10 critical residues

for α-catenin binding (Aberle et al., 1996; Pokutta and Weis, 2000) are marked by α (green). In

reference to human CTNB1, dark gray shading highlights identical residues. Chemically similar

residues marked in light gray for aromatic (F, Y, W, H), aliphatic (V, I, L), positively charged (R,

K, H), negatively charged (D, E), polar but not charged (N, Q, T, S, Y, C) and hydrophobic (V, I,

L, M, A, F, P, W, G) residues.

Fig. S5. WISH for *βcat1, axin1, axin2* and *sfrp*. Representative micrographs spanning different

stages of development in 6-day-old worms with anterior facing left.

Fig. S6. Domain analysis of *H. diminuta* SFRP. BLASTp alignment of SFRPs from flatworms

and human SFRP-1,-2 and -5 as references (*Hd: H. diminuta, Em: E. multilocularis, Smed: S.*

mediterranea). The frizzled domain (orange) and netrin domain (blue) are marked at the most

inclusive positions reported for human SFRP1 using Interpro (Blum et al., 2020). Conserved

cysteines highlighted in yellow.

Fig. S7. Phenotypes observed after *sfrp* RNAi and head amputation. (A) Scheme for RNAi

with dsRNA injections throughout the GR. (B-C) Quantification of proglottid regeneration (B) and

GR length (C) from two independent experiments; Exp 1: n= 8, 8, 9, Exp 2: n= 8, 8, 7; one-way

ANOVA with Dunnett's multiple comparison test. (D) qRT-PCR following dsRNA injection at

the head/scolex (Sc-biased) or throughout the GR. Expression of *sfrp* (target) after *sfrp* RNAi

compared to *GFP* RNAi (set at 1); t-test. (E) qRT-PCR for *sfrp* expression compared to +head (set

at 1); t-test. (F) WISH for *sfrp* at representative worm anteriors from +/- head worms 3 dpa.

Anterior facing left. (D) Quantification of F-*ara*-EdU+ cells normalized to area from 800 μm-wide

cropped region posterior to the head or to the first proglottid. Combined from single and double

dsRNA injections throughout the GR with significant *sfrp* knockdown. All samples were

normalized to the mean of *GFP* RNAi worms set to 100%. Bars= means, gray circles= individual

worms; N= 3, n= 22, 21, 23; one-way ANOVA with Dunnett's multiple comparison test. Error

bars= SD.

Fig. S8. Comparison of regenerates from longest vs. shortest fragment groups after re-

amputation. (A) DAPI-stained regenerates at 18 dpa. (B) Proglottid regeneration at 18 dpa. N= 3,

n= 20, 17; t-test; error bars= SD.

Table S1. *βcatenin* **paralogs in** *H. diminuta.*

Table S2. Primers and transcripts used in this study.

Table S3. Source data used in this study.

Figure 2

A 6-day-old worms +3 days *in vitro*

cut 1

↓ 18 days

cut 2

↓ 18 days

?

B GR length (0 dpa)

means   all

ns   ns

length (mm)

cut 1 cut 2   cut 1 cut 2

C GR % (0 dpa)

means   all

ns   *

percentage (%)

cut 1 cut 2   cut 1 cut 2

D Regeneration success

##

percentage (%)

cut 1 cut 2

E

cut 1

5 mm

cut 2

5 mm

F Proglottid regeneration

**

proglottids

cut 1 cut 2

Figure 3

A +head   -head

Proglottids regenerated

B +head   -head   SYN   F-*ara*-EdU

Proliferation Index

C   + head

D   - head

Figure 4

Figure 5

A βcat1

B βcat1 dsRNA → 3 days → 2 mm → 7 dpa

GFP RNAi   βcat1 RNAi

5 mm   5 mm

C Worm length

D qPCR

βcat1   axin1

RNAi

E GFP RNAi   βcat1 RNAi

F-ara-EdU   F-ara-EdU

Proliferation 6-7 dpi

F RNAi: mcm2

GFP   14/14

βcat1   16/16   200 μm

lbr

GFP   15/15

βcat1   13/13   200 μm

A

sfrp

B

sfrp dsRNA — 3 days — 2 mm — *sfrp* dsRNA — 10-12 dpa

C

GFP RNAi | *sfrp* RNAi

5 mm | 5 mm

D

Proglottid regeneration

% proglottids normalized to *GFP* RNAi

ns | *

UN | *GFP* RNAi | *sfrp* RNAi

E

GFP RNAi | *sfrp* RNAi

0.5 mm | 0.5 mm

F

GR length

GR length (mm)

ns | ****

UN | *GFP* RNAi | *sfrp* RNAi

Figure 7

Figure 8

Figure S1

anterior   proglottids

βcat1 — antisense / sense

fzd5/8 — antisense / sense

tnfr16 — antisense / sense

sfrp — antisense / sense

200 μm

Amputation induced proliferation

Figure S3

A

Rostellum amputated but suckers partially intact

SYN

Amputated through suckers

SYN

B

GR after cut 4

****

ns

GR length (mm)

+head   1/2 head   -head

C   **After first amputation**

****

*

Normalized proglottids
to + head (%)

+ head   1/2 head   -head

D   **After subsequent amputations**

cut 2   cut 3   cut 4

**
***
*

ns
ns

*
*

Normalized proglottids
to + head (%)

+ head 1/2 head   + head 1/2 head   + head 1/2 head

Figure S4

A

B

Figure S5

Figure S6

```
                                          20                    40                    60
Hd_sfrp_HD_tophat_13573    M- - - - - - - - -  - - - -LYLLLL  IFIAITSRGN  TVGRWDWNR-  - - -VNKTSVT  TAPFGGGAFG  43
Em_sfrp_EmuJ_000838700.1   M- - - - - - - - -  - - - -LPLVIL  FAIS-AAQAN  TVGRWDWHRG  GGAKNKTAVT  SSPFGGTSLS  46
Smed_sfrp1_ABY85212.1      MEMTKI- - - -  - - - -FPLSLL  LFISNV- - - -  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  18
Human_SFRP1_sp|Q8N474.1|   MGIGRSEGGR  RGAALGVLLA  LGAALLAVGS  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  30
Human_SFRP5_sp|Q5T4F7.3|   MRAAAAGGGV  RTAALALLL-  -GALHWAPAR  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  28
Human_SFRP2_sp|Q96HF1.2|   MLQGPG- - - -  - - - - -SLLLL  FLASHCCLGS  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  21

                                          80                    100                   120
Hd_sfrp_HD_tophat_13573    RLTTADADMI  SNEPPQSPYF  SDWNRLVSG-  -YGSQ-RCYK  IPKELKLCHK  IGYDLMVLPN  100
Em_sfrp_EmuJ_000838700.1   RLTTADVHMI  SNEPPQSPYF  SDWNRLVSG-  -YGSQ-RCYK  IPRGMKLCHK  IGYDFMVLPN  103
Smed_sfrp1_ABY85212.1      - - - - - - - - - -  - - - - - -NSFI  GDWQAFQSG-  -YTID-QCYQ  IPDNFTLCSN  VGYRLMVLPN  59
Human_SFRP1_sp|Q8N474.1|   - - - - - - - - - -  ASEYDYVSFQ  SDIGPYQSGR  FYTKPPQCVD  IPADLRLCHN  VGYKKMVLPN  80
Human_SFRP5_sp|Q5T4F7.3|   CEEYDYYGWQ  AE- -PLH-GR  SYSKPPQCLD  IPADLPLCHT  VGYKRMRLPN  75
Human_SFRP2_sp|Q96HF1.2|   - - - - - - - - - -  ARGLFLFG-Q  PDF- - - - - - -  SY-KRSNCKP  IPANLQLCHG  IEYQNMRLPN  62
```

Frizzled domain

```
                                          140                   160                   180
Hd_sfrp_HD_tophat_13573    SLEHEGMDEV  ITQSEVWLTL  VNLGCHDELE  RFLCSLYAPV  CIRGYHEKLI  QPCRELCESV  160
Em_sfrp_EmuJ_000838700.1   SLEHETLDEA  ITQSEVWLTL  VNLGCHDELK  RFLCSLYAPV  CINGYHEKLI  QPCRELCESV  163
Smed_sfrp1_ABY85212.1      YIQHESIEEA  AQHSKVWMGL  VNTQCHVDIK  KFICSLYAPV  CISNQRNQKV  PPCRELCVTV  119
Human_SFRP1_sp|Q8N474.1|   LLEHETMAEV  KQQASSWVPL  LNKNCHAGTQ  VFLCSLFAPV  CLD- - - -RPI  YPCRWLCEAV  136
Human_SFRP5_sp|Q5T4F7.3|   LLEHESLAEV  KQQASSWLPL  LAKRCHSDTQ  VFLCSLFAPV  CLD- - - -RPI  YPCRSLCEAV  131
Human_SFRP2_sp|Q96HF1.2|   LLGHETMKEV  LEQAGAWIPL  VMKQCHPDTK  KFLCSLFAPV  CLDDL-DETI  QPCHSLCVQV  121

                                          200                   220                   240
Hd_sfrp_HD_tophat_13573    RAACLPTMTT  FGLGWPDIVK  CSKFPQAPQE  LCIPPNKQRN  KTVVLKPETR  -CSGCVDKPT  219
Em_sfrp_EmuJ_000838700.1   RAACLPTMTT  FGLGWPDIVK  CSKFPQAPQE  LCIPLNKHKN  KTIILNPETR  -CSGCIDRPT  222
Smed_sfrp1_ABY85212.1      KESCLPSMKM  YGFDWPTIMK  CSKFPQFKNS  LCIP- - - - -  K.VV- -PGKK  -CDLCMKIAS  170
Human_SFRP1_sp|Q8N474.1|   RDSCEPVMQF  FGFYWPEMLK  CDKFP- -EGD  VCIAMTPPNA  TEASKPQGTT  VCPPCDNELK  194
Human_SFRP5_sp|Q5T4F7.3|   RAGCAPLMEA  YGFPWPEMLH  CHKFPL-DND  LCIAVQFGHL  P-ATAPPVTK  ICAQCEMEHS  189
Human_SFRP2_sp|Q96HF1.2|   KDRCAPVMSA  FGFPWPDMLE  CDRFPQ-DND  LCIPLASSDH  LLPATEEAPK  VCEACKNKND  180
```

Netrin domain

```
                                          260                   280                   300
Hd_sfrp_HD_tophat_13573    YES-AIGSFC  TADVAVRSKV  LDLIPVPSTK  GQNTTYRIRT  DGRGGIFKM-  - - -PKNVNKE  274
Em_sfrp_EmuJ_000838700.1   YES-AIGSFC  TADVVIRAKV  LDLVPTLPSS  GHNTTHRIRT  TGRVGAFKL-  - - -PKTLNSI  277
Smed_sfrp1_ABY85212.1      YEV-IANRFC  LSPIVIRAKI  KRIIP- - -TS  GN- -AIQIIL  QKKSKFLKF-  - - -QANFSQN  220
Human_SFRP1_sp|Q8N474.1|   SEA-IIEHLC  ASEFALRMKI  KEV- - - - -KK  ENGDKKIV- -  PKKKKPLKLG  PIKKKDLKKL  246
Human_SFRP5_sp|Q5T4F7.3|   ADG-LMEQMC  SSDFVVKMRI  KEI- - - - -KI  ENGDRKLIGA  QKKKKLLKPG  PLKRKDTKRL  243
Human_SFRP2_sp|Q96HF1.2|   DDNDIMETLC  KNDFALKIKV  KEI- - - - -TY  INRDTKIILE  TKSKTIYKLN  GVSERDLKKS  235

                                          320                   340                   360
Hd_sfrp_HD_tophat_13573    A-NLDFRMTC  DCPIIKAAMT  RSKGPGRWLL  MGKLNEDGRS  VTVLHLSQPS  RQNKGIKRAI  333
Em_sfrp_EmuJ_000838700.1   V-NLEFEMEC  DCPIIRAALT  RQKGPGRWLM  MGKLNDDRRT  VTVQHISQPS  RQNAGIKRAI  336
Smed_sfrp1_ABY85212.1      A- -LELNLKC  NCTNLKFKQR  ALKG- -RWII  MAQIDSDNKA  V-VNFISKWK  RKKAEFKHSM  275
Human_SFRP1_sp|Q8N474.1|   VLYLKNGADC  PCHQLDNLS-  - - - - -HHFLI  MGR- -KVKSQ  YLLTAIHKWD  KKNKEFKNFM  298
Human_SFRP5_sp|Q5T4F7.3|   VLHMKNGAGC  PCPQLDSLA-  - - - - -GSFLV  MGR- -KVDGQ  LLLMAVYRWD  KKNKEMKFAV  295
Human_SFRP2_sp|Q96HF1.2|   VLWLKDSLQC  TCEEMNDIN-  - - - - -APYLV  MGQ- -KQGGE  LVITSVKRWQ  KGQREFKRIS  287

                                          380                   400                   420
Hd_sfrp_HD_tophat_13573    KEMRHQPDTL  CKVDLTRSSP  VVHPRTVAGR  PAANGAQISQ  EQKQQHQQLQ  PRRNGSNLSK  393
Em_sfrp_EmuJ_000838700.1   RDIRRRPDSL  CKVDLMHPNP  VAHSRAISSL  SSPSSMQSLR  PRSHQSRRYR  P- - - - -HLSP  391
Smed_sfrp1_ABY85212.1      KIIQNFGHIV  CKSKLPDGIS  L-RERYAAYL  KMNNQLKYSN  ERKLQTRKFR  - - - - - - - - -K  325
Human_SFRP1_sp|Q8N474.1|   KKMKNHE- - -  CPTFQSVF- -  - - - -K- - - - -  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  314
Human_SFRP5_sp|Q5T4F7.3|   KFMFSYP- - -  CSLYYPFFYG  AAEPH- - - - -  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  317
Human_SFRP2_sp|Q96HF1.2|   RSIRKLQ- - -  C- - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  295

                                          440                   460                   480
Hd_sfrp_HD_tophat_13573    EERRRRRQLR  RRRNE- - -AG  AQQQQHPSNV  TLSSTNSQES  QQPVNDMPIG  VNTQHPYNQQ  450
Em_sfrp_EmuJ_000838700.1   EERRRR-QLR  RRRNHPRGSS  PEAQQRAANV  TPSRKPPQLQ  QQPINDIPIL  TPTPRPFSHH  450
Smed_sfrp1_ABY85212.1      QKSRK- - - - -  - - - - - - - - -S  VKPNNRSAGI  S- - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  342
Human_SFRP1_sp|Q8N474.1|   - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  314
Human_SFRP5_sp|Q5T4F7.3|   - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  317
Human_SFRP2_sp|Q96HF1.2|   - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  - - - - - - - - - -  295
```

A

cut A → 18 dpa

cut D → 18 dpa

5 mm

5 mm

B

Proglottid regeneration

**

\# proglottids

A   D