

SUPPLEMENTARY MATERIAL

corresponding to:

**Two *msh/msx*-related genes, *Djmsh1* and *Djmsh2*,
contribute to the early blastema growth
during planarian head regeneration**

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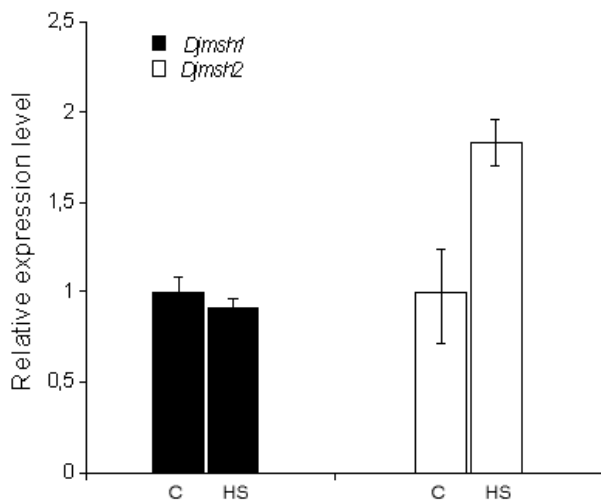


Fig. S2. Relative expression level of *Djmsh1* and *Djmsh2* in intact planarians after heat shock treatment, analyzed by real time RT-PCR. Heat shock was performed in 30 intact planarians by overnight incubation at 28°C. The specimens were then sacrificed for RNA extraction. *Djmsh1* expression level did not change after heat shock treatment, while *Djmsh2* expression was strongly activated, indicating that *Djmsh2* is a stress-responsive gene. Expression levels are indicated in relative units, assuming the value of untreated specimens (control) as unitary. Each value is the mean ± s.d. of three independent samples, analyzed in duplicate. C, control; HS, heat shock.

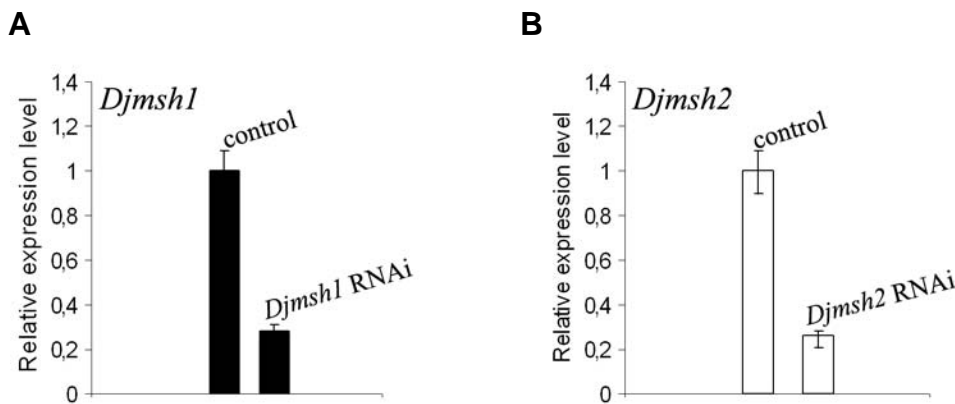


Fig. S3. RNAi-mediated downregulation of *Djmsh1* and *Djmsh2* in regenerating planarians, as visualized by real time RT-PCR. (A) Expression level of *Djmsh1* in water-injected controls and *Djmsh1* RNAi animals. (B) Expression level of *Djmsh2* in water-injected controls and *Djmsh2* RNAi animals. In all experiments, the expression levels are indicated in relative units, assuming as unitary the value of water-injected controls. Each value is the mean ± standard deviation of two independent RNAi experiments, performed in duplicate.

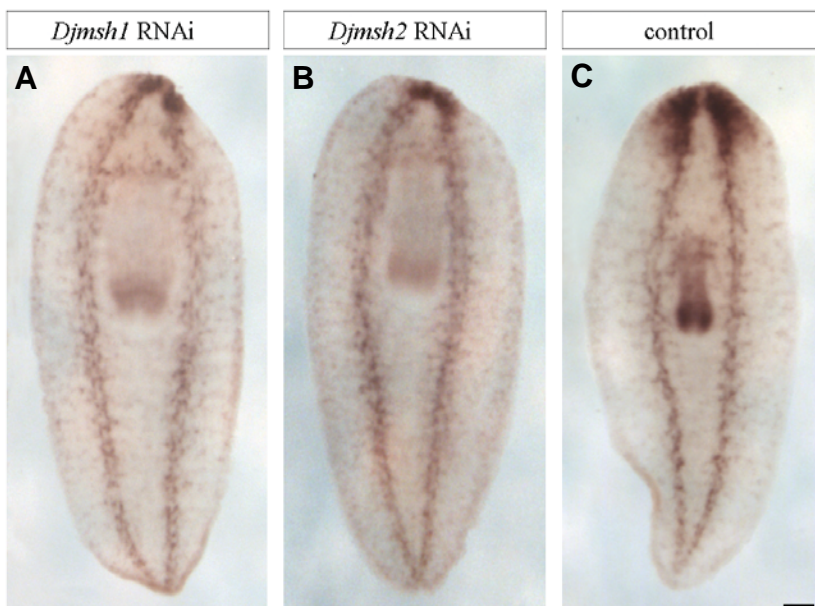


Fig. S4. Whole mount *in situ* hybridization with *Djsyt* of representative phenotypes of planarian fragments regenerating a head, 6 days after transection. (A) *Djmsh1* RNAi. (B) *Djmsh2* RNAi. (C) Water-injected control. Scale bar, 100 µm.

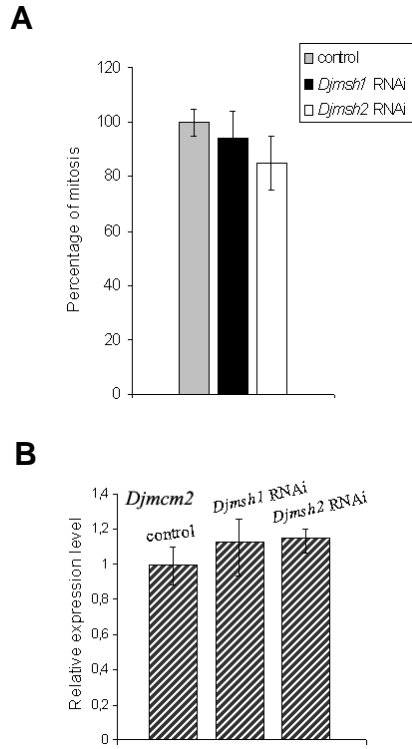


Fig. S5 (Left). Effects of *DjmsH1* and *DjmsH2* RNAi on intact planarians. (A) Analysis of the percentage of mitoses in intact planarians injected with *DjmsH1* or *DjmsH2* dsRNA and in samples injected with water (control). The number of mitotic metaphases was normalized to the number of total cells and the values indicated in the graph are average \pm s.d. of three independent samples, assuming as 100% the value of the control. (B) Expression level of *DjmcM2*, analyzed by real time RT-PCR in water-injected controls and in *DjmsH1* and *DjmsH2* RNAi planarians. In all experiments, the expression levels are indicated in relative units, assuming as unitary the value of the controls. Each value is the mean \pm s.d. of two independent RNAi experiments, carried out in duplicate.

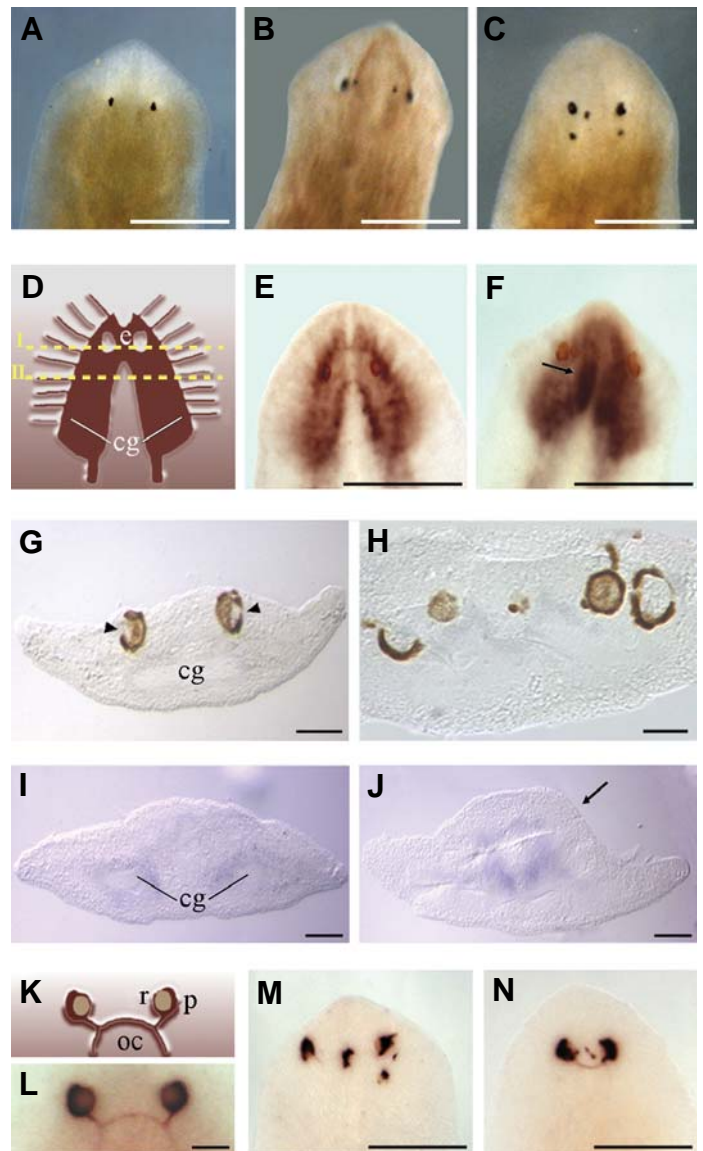


Fig. S6 (Right). Functional analysis of *Djbmp* during anterior regeneration. (A-C) Brightfield images of a water-injected control (A) and representative *Djbmp* RNAi phenotypes (B,C), 6 days after transection. (D) Schematic drawing of the planarian brain, composed by a mass of nerve cells, the cephalic ganglia (cg), e: eyes. I and II indicate paraffin section levels. (E,F) Representative images of whole-mount in situ hybridization with *Djsyt* of a water-injected control (E) and a *Djbmp* dsRNA-injected planarian with abnormal cephalic ganglia (arrow) (F), 15 days after transection. (G-J) Transverse paraffin sections. (G) Control depicted in (E) (arrowheads indicate the eyes), and (H) the *Djbmp* dsRNA-injected planarian depicted in (F), at the level I. (I) Control depicted in (E) and (J) the *Djbmp* dsRNA-injected planarian depicted in (F), at the level II. Arrow shows the dorsal outgrowth. (K) Schematic drawing of the planarian visual system. p, pigmented eye cup; r, rhabdomeric photoreceptors; oc, optic chiasma. (L-N) Whole-mount in situ hybridization with *Djops* of a water-injected control (L) and representative *Djbmp* RNAi phenotypes with multiple eyes (M) and abnormal chiasma (N), 15 days after transection. Scale bars, 500 μ m in (A-J, M, N), and 100 μ m in (L). According to the *Djbmp* RNAi results recently obtained and demonstrating that *Djbmp* knockdown results in additive ventral structures in the dorsal side of regenerants (Orii and Watanabe, 2007), our functional experiments confirmed that *Djbmp* acts as a key signaling molecule in the dorso-ventral patterning of neural structures during head regeneration. In fact we observed that, while 20-25% of planarians died as a consequence of *Djbmp* RNAi, about 70% of the injected specimens regenerated heads with morphological defects in the dorsoventral (DV) axis and in the organization of the cephalic nervous structures. Multiple eyes were also frequently observed (B,C). A further analysis of these phenotypes by using the panneural marker *Djsyt* highlighted abnormal dorsal outgrowth of the nervous structures in the head region (E-G). Eye alterations could be better detected after in situ hybridization with the photoreceptor marker *Djops* that, following prolonged detection, may reveal the presence of *Djops* transcripts also in the photoreceptor axons. Using this approach, multiple eyes with altered organization of the optic cup and/or abnormal connections between eyes and brain were observed (L-N).

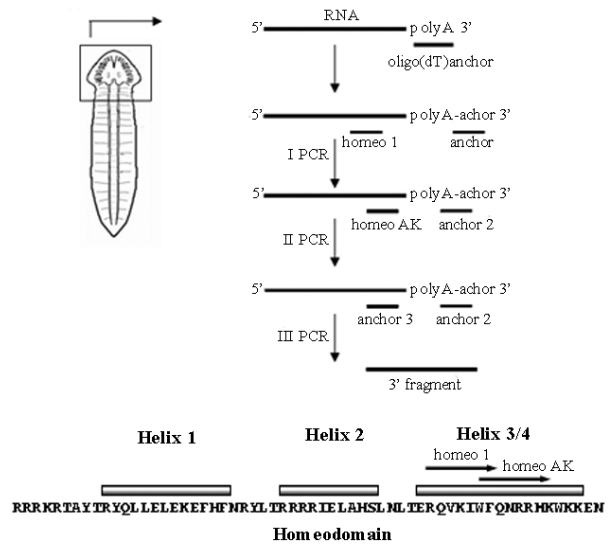
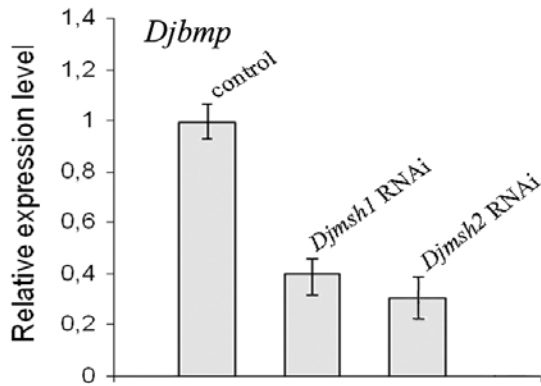


Fig. S7 (Left). *Djbmp* expression level in water-injected controls and in *Djmsh1* or *Djmsh2* RNAi planarians, 3 days after transection. In all experiments, the expression level is indicated in relative units, assuming as unitary the value of the water-injected controls. Each value is the mean \pm s.d. of two independent RNAi experiments, performed in duplicate.

Fig. S8 (Right). RT-PCR strategy utilized to isolate *Djmsh1* and *Djmsh2* from regenerating heads. cDNA obtained from RNA isolated from the head portion of anterior regenerants was amplified by using the anchor primer (Roche Applied Science) and a sense-strand primer, selected from the homeodomain region R(I/V)(K/R/Q)(I/V)WF (primer homeo 1: 5'-CGGRTNMRIWITGGTT-3'), for 35 cycles at 94°C for 30 sec, 40°C for 1 min, and 72°C for 1 min. A second round of PCR was then performed with a sense-strand primer representing the homeodomain region WF(Q/K)NRRAK (primer homeo AK: 5'-TCGCGGATCCTGGTTMARAAYMGNMGIGCAA-3'), and the primer 2 (5'-CACGCGTATCGATGTCGAC-3'). Primer homeo AK includes, at the 5' end, an adapter corresponding to primer 3 (5'-TCGCGGATCCTGGTTT-3'), the amplification reaction was conducted as described above except for the annealing temperature that was 50°C. Subsequently, a third PCR was carried out on the diluted second PCR products using the primers 2 and 3. The products of the third PCR round were *Clal/BamHI* digested and directionally cloned into the *Clal/BamHI* digested pBluescript SK vector (Stratagene). The clones were sequenced by automated fluorescent cycle sequencing (ABI).

SUPPLEMENTARY TABLE S1

PROBES AND PRIMERS USED IN REAL TIME RT-PCR

Clone	1) Probe 2) Forward primer 3) Reverse primer
<i>DjEF2</i>	1) 5'-CCAACAAGTCCACATATGTT-3' 2) 5'-GCAATCGAAGACGTTCCATGTG-3' 3) 5'-CCAGGAAAAGTTGTTATAGTCCAGTTT-3'
<i>Djmsh1</i>	1) 5'-CCAAGATTGTGATATCTCC-3' 2) 5'-CCATACAGCTGCTCTGTTCCATTTA-3' 3) 5'-GGGCTCTGAAAGCAGTAATTTGATT-3'
<i>Djmsh2</i>	1) 5'-CCTGGTATTTTATGGAATTAT-3' 2) 5'-CTGATGATTCGCTGAAAATTCATTCA-3' 3) 5'-ATTGCACTTTTGATTTGACACATTTTTTGA-3'
<i>DjBMP</i>	1) 5'-CTTTGGCAATGAATCATTTTC-3' 2) 5'-GAAAACCGAAAACATTTAGGACCAGTT-3' 3) 5'-GAATACCAACCGAAGACCATGTTTG-3'
<i>DjMCM2</i>	1) 5'-CTCTACTGCTTCATAAGTCTG-3' 2) 5'-GGCAGGTGAAACATTGGGATCA-3' 3) 5'-GGCTACCGACATTCCTTTGGT-3'