

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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DjMSH1	1	$\texttt{MKT}\underline{\texttt{RNFDIDCLL}QPRPVIHARSEPVTTIHHRSNIKCPISSDSQWSNEVVNSLEYRWINN}$	60
DjMSH2	1	MEGKSISDFSISKLITDVTHETQSNFLIGKKKFST	35
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		N-bern Arm Helix I	
DjMSH1	61	IITNENYEPSSQKFS.QIQNHCQISTPIHFRTSPNKCTLRKHKENRKPRTPFTTQQLIQL	119
DjMSH2	36	DDSAENFIHRDLFYPWYFMELSKASNNFQKNVSNQKCNLRKHKENRKPRTPFTTQQLMEL	95
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		Heliy T Heliy TT THYD Heliy TTT	
DjMSH1	120	EKKFHQKQYLSIAERAEFSSSLGIJEIQVKIWFQNRRAKSKRLQEAEIDKLRIDHTAALF	179
DjMSH2	96	ENKFLTKQYLSISERAEFSTNISLTETQVKIWFONRRAKAKRLEETELEKYRFIKRPAL.	155
2		* ** **********************************	
DjMSH1	180	HLSKDCDISNQITAFRAPSLIPIIPALFSQKSYETSRLLNTL	221
DjMSH2	156	EDAELYLSNNQNCLPNSIMNDENWNNHKIPKNFRSISLFDGHIDKLSSTTSIPGFVS	211
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DjMSH1	221	221	
D-1MSH2	212	KFTSSCSSAYO 222	

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	← → ←		→
Hs MSX1	CTLRKHKTNRKPRTPFTTA	QLLALERKFRQKQYLSIAERAEFSSSLSLTETQVKIWFQ	NRRAKAKRLQEAELEKLK
Hs MSX2			
Mm MSX1			
Mm MSX2	s		
Mm MSX3			
Gq MSX1			
xl xHOX-7	.P		
Xl XHOX-7'	s		I
Dr MSXA	. P		RF.
Dr MSXB	s	s	F.
Dr MSXC			
Dr MSXD	.P		
Dr MSXE	.P		
Bf MSX	Q	AN	
Ci MSXB	. S P S. E	SR.QD	HFV.
Sk MSX	s	AN	
Зр МЗН		AN	
Ame H17	Q	sKETHH.	I
Dm MSH	.NQ	SKE	I.
Ce VAB-15	.MN	ISQSAQ	vvvv.
Pd MSX	VQQ	AN	
Hr Le-MSX	Q	NER	s
Dj MSH1	Q	QKH	SIDR
Dj MSH2	.NQ	MENLTSTN	YR
Smed MSH1	Q	IQKH	SIDR
Smed MSH2	.NQ	MENMTSTN.N	YR
Hvi MSH	.F	TQKRSL.EL.R	QSKI.E
Hvu MSH	.IA	QKRSS.L.VM.R	QRTI.ETA
Ami MSX3	.Qs	S	H

C_flenking

Fig. S1. *msh/msx*-related genes in planarians. (A) Alignment of amino acid sequences of Dj/MSH1 and Dj/MSH2. Analysis of sequence similarity was performed with CLUSTAL W software. The regions flanking the homeodomain in 5' and 3' are boxed. The homeodomain is boxed and in grey. The region related to the en1 core motif, starting with a conserved Phenylalanine (position 6 and 9 in Dj/MSH1 and Dj/MSH2, respectively) is underlined. (B) Amino acid alignment of the homeodomain and flanking regions of representative MSH/MSX proteins from vertebrate and invertebrate species. Identical amino acid residues are indicated by dots with respect to the Hs MSX1 sequence; the incomplete sequence of HrLe-MSX is indicated by hyphens. Sequences were obtained from the EMBL/ GenBank. AmeH17: Apis mellifera H17 (M29491); AmiMSX3: Acropora millepora MSX3 (EF044217); Bf/MSX: Branchiostoma floridae MSX (AJ130766); CeVAB-15: Caenorhabditis elegans VAB-15 (AF286218); CiMSXB: Ciona intestinalis MSXB (AJ515411); Dj/MSH1: Dugesia japonica MSH1 (AM293348); Dj/MSH2: Dugesia japonica MSH2 (AM293349); DmMSH: Drosophila melanogaster MSH; DrMSXA: Danio rerio MSXA (NM131274); DrMSXB: Danio rerio MSXB (U16311); DrMSXC: Danio rerio MSXC (NM131272); DrMSXD: Danio rerio MSXD (NM131276); GrMSXE: Homo sapiens MSX2 (NM002449); HviMSH: Hydra viridis MSH (X64629); HvuMSH: Hydra vulgaris MSH (AJ271008); MmMSX1: Mus musculus MSX1 (AF308572); MmMSX2: Mus musculus MSX2 (NM013601); MmMSX3: Mus musculus MSX3 (AF060229); PdMSX: Platynereis dumerilii MSX (AM114785); SkMSX: Saccoglossus kowalevskii (DQ431008); SmedMSH1: Schmidtea mediterranea MSH2 (AM293347); SpMSH: Strongylocentrotus purpuratus MSH (NM214613); XIXHOX-7: Xenopus laevis HOX-7 (P35993); XIXHOX-7': Xenopus laevis HOX-7 (Q04281).



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 con*trols.* 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 µm in (A-J, M,N), and 100 µm 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 Disyt highlighted abnormal dorsal outgrowth of the nervous structures in the head region (E-G). Eye alterations could be better detected afterin situ hybridization with the photoreceptor marker Diops that, following prolonged detection, may reveal the presence of Diops 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 regeneratis 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'-CGGRTNMRIRWITGGTT-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'-TCGCGGATCCTGGTTTMARAAYMGNMGIGCAA-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
DjEF2	1) 5'-CCAACAAGTCCACATATGTT-3' 2) 5'-GCAATCGAAGACGTTCCATGTG-3' 3) 5'-CCAGGAAAAGTTGTTATAGTCCCAGTTT-3'
Djmsh1	1) 5'-CCAAAGATTGTGATATCTCC-3' 2) 5'-CCATACAGCTGCTCTGTTCCATTTA-3' 3) 5'-GGGCTCTGAAAGCAGTAATTTGATT-3'
Djmsh2	1) 5'-CCTGGTATTTTATGGAATTAT-3' 2) 5'-CTGATGATTCCGCTGAAAATTTCATTCA-3' 3) 5'-ATTGCACTTTTGATTTGACACATTTTTTTGAA-3'
DjBMP	1) 5'-CTTTGGCAATGAATCATTTC-3' 2) 5'-GAAAACCGAAACATTTAGGACCAGTT-3' 3) 5'-GAATACCAACCGAAGACCATGTTTG-3'
DjMCM2	1) 5'-CTCTACTGCTTCATAAGTCTG-3' 2) 5'-GGCAGGTGAAACATTGGGATCA-3' 3) 5'-GGCTACCGACATTCCTTTGGT -3'