Cloning and characterization of *Vasa* gene expression pattern in adults of the Lusitanian toadfish *Halobatrachus didactylus*

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Supplement. Lusitanian toadfish cDNA *Vasa* products and Genbank accession numbers of the sequences used to perform the molecular phylogenetic analysis

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D  S  R  K  G  R  A  F  P  D  N  S  V  V  T  Q  P  N  D  P  A  V  A  D  D  D  E  W  E  *
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Fig. S1. Nucleotide and predicted amino acid sequences of Lusitanian toadfish *Vasa* cDNA (GenBank accession number JX849133), shown in lower and upper case letters, respectively. Nucleotides are numbered to the right. Arginine/glycine-rich regions in the N-terminal region are boxed with a discontinuous line. Eight consensus sequences for the DEAD-protein family are framed by a solid black line. The region amplified by qPCR and used as probe in *in situ* hybridization (ISH) is emphasized in gray, primers OLIGOVAF and OLIGOVAR are represented by a solid black line above them

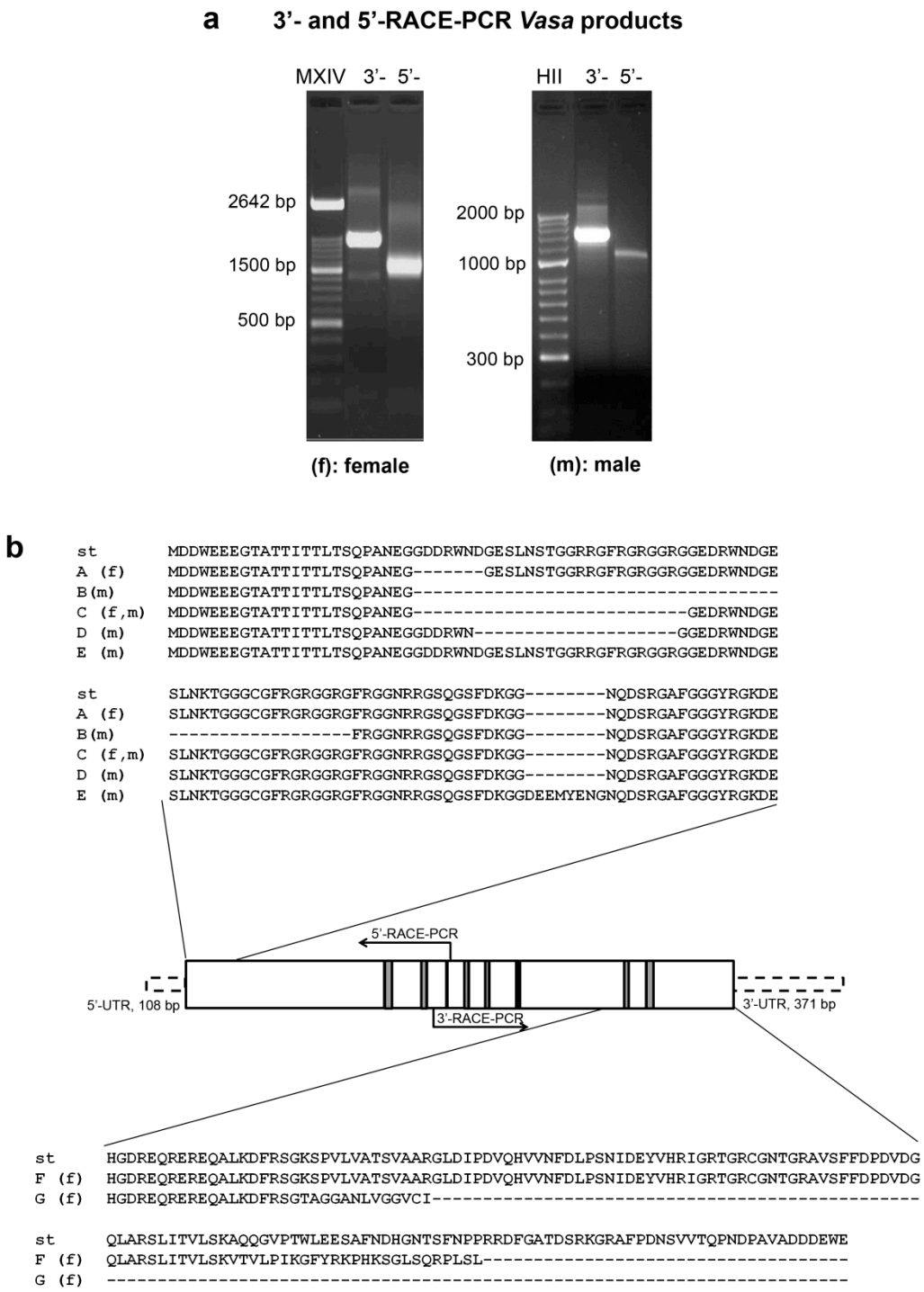

Fig. S2. (a) Lusitanian toadfish 3'- and 5'-RACE *Vasa* products from ovary and testis. MXVI: molecular weight Marker XIV (Roche Applied Science). HII: molecular weight Marker Hyperladder II (Bioline). (b) Diagram of the standard *Vasa* transcript with the 8 conserved domains for DEAD-box proteins in gray. The alignments of protein sequences are shown in the regions in which the minority products A (from JX849140), B (JX849137), C (JX849136), D (JX849139), E (JX849138), F (JX849141), and G (JX849142) differ from the standard sequence (st). Hyphens represent deleted amino acids

Table S1. Genebank accession numbers of the amino acid sequences from the DEAD-box protein family used to perform the molecular phylogenetic analysis

	Species	Accession	Species	Accession
Vasa (DDX4)	*Danio rerio*	AAI29276	*Ctenopharyngodon idella*	ACR61400
	Gobiocypris rarus	AFA45124	*Kryptolebias marmoratus*	AGA16734
	Cyprinus carpio	AAL87139	*Leucopsarion petersii*	BAD04052
	Carassius gibelio	AAV70960	*Osphronemus goramy*	ACV69940
	Carassius auratus	AAX22126	*Dicentrarchus labrax*	ADK79106
	Seriola quinqueradiata	ADD91316	*Trachurus japonicus*	BAG72093
	Nibea mitsukurii	ACV32355	*Oreochromis aureus*	AEO36953
	Oreochromis niloticus	BAB19807	*Salvelinus leucomaenis*	ACA33927
	Thunnus orientalis	ABY77970	*Auxis thazard*	ADD81194
	Katsuwonus pelamis	ADD81192	*Scomber australasicus*	ADD81190
	Auxis rochei	ADD81193	*Euthynnus affinis*	ADD81191
	Scomber japonicus	ACV32356	*Oryzias latipes*	NP_001098146
	Sebastes schlegelii	AEP68013	*Silurus meridionalis*	ACD62525
	Clarias gariepinus	ADK94762	*Monopterus albus*	ABA54551
	Cynoglossus semilaevis	ADX41681	*Paralichthys olivaceus*	AEY68604
	Solea senegalensis	AFN89212	*Scophthalmus maximus*	AFQ38974
	Pagrus major	BAJ25759	*Oncorhynchus mykiss*	NP_001117665
	Gadus morhua	ADV36250	*Culter ilishaeformis*	AGG53839
	Lateolabrax japonicus	AFI61840	*Xenopus laevis*	NP_001081728
	Salmo salar	AFH41530	*Homo sapiens*	NP_077726
	Larimichthys crocea	AFW17056	*Gallus gallus*	NP_990039
	Misgurnus anguillicaudatus	BAJ19133		
PL10 (DDX3)	*Danio rerio*	NP_571016	*Xenopus laevis*	BC044972
	Homo sapiens	AAC34298		
DDX17	*Gallus gallus*	XP_416260	*Danio rerio*	XP_001923830
	Homo sapiens	CAG30318	*Xenopus laevis*	NP_001082679
DDX5	*Homo sapiens*	NP_004387	*Gallus gallus*	NP_990158
	Xenopus laevis	AAH82849	*Danio rerio*	NP_997777
	Salmo salar	ACN11269		
DDX41	*Homo sapiens*	NP_057306	*Meleagris gallopavo*	XP_003210383
	Salmo salar	NP_001133799		