

Expressions of Raldh3 and Raldh4 during zebrafish early development

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Abstract

Retinoic acid (RA) plays crucial roles in vertebrate embryogenesis. Four retinal dehydrogenases (Raldhs) that are responsible for RA synthesis have been characterized in mammals. However, only Raldh2 ortholog is identified in zebrafish. Here, we report the identification of *raldh3* and *raldh4* genes in zebrafish. The predicted proteins encoded by zebrafish *raldh3* and *raldh4* exhibit 70.0% and 73.5% amino acid identities with mouse Raldh3 and Raldh4, respectively. RT-PCR analyses reveal that both *raldh3* and *raldh4* mRNAs are present in early development. However, whole mount *in situ* hybridization shows that *raldh3* mRNA is first expressed in the developing eye region of zebrafish embryos at 10-somite stage. At 24 hpf (hours post fertilization), *raldh3* mRNA is expressed in the ventral retina of eye. At 36 hpf, the mRNA is also expressed in otic vesicle in addition to ventral retina, and it maintains its expression pattern till 2 dpf (days post fertilization). At 3 dpf, *raldh3* mRNA becomes very weak in ventral retina but is present in otic vesicle at a high level. At 5 dpf and 7 dpf, *raldh3* is no longer expressed in eyes but is expressed in otic vesicles at a very low level. *raldh4* mRNA is initially detected in developing liver and intestine regions at 2 dpf embryos. Its expression level becomes very high in these two regions of embryos from 3 dpf to 5 dpf. Analysis on the sections of 5 dpf embryos reveals that *raldh4* is expressed in the epithelium of intestine. At 7 dpf, *raldh4* mRNA is only weakly expressed in the epithelium of intestinal bulb.

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1. Results and discussion

Retinoic acid (RA) is a morphogen that is essential to vertebrate embryogenesis (Lander, 2007). *In vivo*, animals synthesize RA by first converting the precursor vitamin A into retinal using retinol dehydrogenase, then oxidizing retinal into RA irreversibly using retinal dehydrogenases (Raldh) (Ross et al., 2000). Four Raldhs including Raldh1, Raldh2, Raldh3 and Raldh4 have been characterized in mammals (Glover et al., 2006). Raldh2, a major retinoic acid generating enzyme in the early mouse embryo (Zhao et al., 1996), is mainly expressed in mesoderm during early gastrulation, and is later expressed in undifferentiated som-

ites, the optic vesicles and differentiating limbs (Niederreither et al., 1997). Mice with *Raldh2* deletion die at midgestation with a shorter anterior–posterior axis, open neural tube, absence of limb buds (Niederreither et al., 1999), and asymmetric somitogenesis (Vermot et al., 2005; Vermot and Pourquie, 2005). Raldh3 is primarily localized in ventral retinas of the developing eyes, otic vesicles and olfactory placodes in early mouse embryos (Li et al., 2000; Mic et al., 2000). *Raldh3*-null mice die at birth from respiratory distress due to choanal atresia, displaying malformations in ocular and nasal regions (Dupé et al., 2003), and mild defects in retinas (Dupé et al., 2003; Molotkov et al., 2006). Raldh1 is expressed in dorsal retinas and mesencephalic flexures (McCaffery et al., 1991; Mic et al., 2000). Genetic ablation of *Raldh1* had no apparent defects in mouse embryo development (Fan et al., 2003;

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Matt et al., 2005). However, double knock out of both *Raldh1* and *Raldh3* causes apoptosis and alters gene expressions in periocular mesenchymal cells, resulting in severe eye abnormalities in mice (Matt et al., 2005; Molotkov et al., 2006). *Raldh4* is first expressed in liver at E14.5 stage during mouse embryogenesis and present in adult mouse liver and kidney (Lin et al., 2003). The role of *Raldh4* in mice embryogenesis is unknown because targeted deletion of *Raldh4* in mouse has not been reported yet. Till now, only *Raldh2* ortholog has been characterized in zebrafish. Zebrafish *raldh2*^{-/-} mutants (*nls*, *nof*) display a truncation of the anterior–posterior axis anterior to the somites, absence of pectoral fins (Begemann et al., 2001; Grandel et al., 2002) and asymmetric somitogenesis (Kawakami et al., 2005), which are similar to defects of *Raldh2*^{-/-} mouse. In this paper, we report the identification of *raldh3* and *raldh4* genes in zebrafish and describe their expression patterns during early development.

1.1. Zebrafish has *raldh3* and *raldh4* genes

BLASTing the zebrafish EST database with *Xenopus laevis* *Raldh3* sequence (GenBank Accession No.: AY692028) using TBLASTN program, we found a candidate EST (DR729671) of zebrafish *Raldh3*. Employing RACE-PCR strategy, we cloned a cDNA of zebrafish *raldh3* with the length of 1590 base pairs (bp). The sequence was deposited in GenBank under Accession No. EF375713. The *raldh3* cDNA consists of a complete

CDS (coding sequence) with 1542 bp long, predicted to encode a protein comprising 513 amino acids (Fig. 1). BLASTing GenBank protein database, we found that the predicted protein is identical to that encoded by **DQ300198** (Canestro et al., 2006) and the two cDNAs are mapped in the same contig of zebrafish chromosome 7, and it is encoded by 13 exons. However, the CDS of **EF375713** has nine nucleotides (9 of 1542) that are different from that of **DQ300198**. The difference indicates that single nucleotide polymorphisms (SNPs) are present in the gene. Protein alignment analysis reveals that zebrafish *Raldh3* shares high identity with other members of *Raldh* family (Table 1). It exhibits 64.5% amino acid identity with zebrafish *Raldh2*, about 70% identity with mammalian *Raldh3*, 63% identity with mammalian *Raldh1* and 34% identity with mammalian *Raldh4*. DNA sequence alignment shows that the CDS of zebrafish *raldh3* (EF375713) shares 65.5% nucleotide identity with that of zebrafish *raldh2*, 69% identity with that of mammalian *Raldh3*, about 65% identity with that of mammalian *Raldh1* and about 50% identity with that of mammalian *Raldh4*. Using Mega3.1, phylogenetic analysis on the vertebrate *Raldh* family shows that zebrafish *Raldh3* is clustered into *Raldh3* subfamily (Fig. 2). Additionally, synteny analysis reveals that zebrafish *raldh3* gene is surrounded by *asb7* gene in chromosome 7. Similarly, mouse and rat *Raldh3* genes are also linked with *Asb7* and are located in mouse chromosome 7 and rat chromosome 1, respectively (<http://www.ncbi>.

	1	11	21	31	41	51	61	71	81	91	101	111
Raldh2	HTSSEVLP	GEVKKDPAALMA	SLHMFSPV	NFKYTKIFINNE	WDSVSGVF	PTYPATGE	KICDVQ	EADKAD	VKAQAARS	ATSLSSV	WVKMDAS	EGKLI
Raldh3	MAQNGTI	ENGDSR	-----	TSPPPL	QPKITE	KIKIFINNE	WTSKGG	QPTTIN	PATGVC	KICDEE	ADKADV	EA
Raldh4	-----	-----	-----	MSKDM	KYLVLEN	-----	YIGGK	VPCS	-----	KLDS	FDPST	GHVYCK
Consensus	m	g	n	s	mp	pvk	eik	tkifinn	ewh	ss	gk	fpt
	121	131	141	151	161	171	181	191	201	211	221	231
Raldh2	ATLESLS	GKPFIL	PCFFVD	LQGI	LKTRFY	AGADK	THGSL	IRIDGE	FTLLR	HEPIG	VCGLIP	WNFFL
Raldh3	ATLESKIT	GKPFIL	MAFFVD	LSIKL	RYAGT	DKING	KTMVDEN	FVCF	KHEPI	GVCGLI	PNFFL	MLNM
Raldh4	VQAESK	IQGKIT	ITFARN	VDIP	RSAYN	FRFAS	SVLH	OTNDC	SQMDM	CLNYT	IRCP	VGVAGL
Consensus	atleskd	gkplf	affvdl	gsiktr	fyagw	dking	t p d	f	t	hepi	gvcg	iipwn
	241	251	261	271	281	291	301	311	321	331	341	351
Raldh2	PGYGP	TAGAA	SSHHG	IDKVA	FTGSTE	VGKI	QEAAG	KNLKV	TLGGK	SENTI	FADAD	FELAL
Raldh3	PGFGP	TAGAA	LAGH	NNIDK	LAFTG	STEV	QGLV	KAAAS	NNLKV	TLGGK	NPCIV	FADSD
Raldh4	FTGPR	AGD	ALVSH	PDV	FLISFT	GSTAT	ARLIT	ERSAP	HC	KLS	ELGGK	NALIF
Consensus	pg	gptaga	ai	shm	idk	aftgstev	g li	aaaa	snlkr	vtlel	ggknp	iifadad
	361	371	381	391	401	411	421	431	441	451	461	471
Raldh2	GPQVSE	EQRRV	LELTQ	SGIT	EGAKLE	CGG	-----	KAPATK	-----	GFVE	PTVFS	NVDH
Raldh3	GPQIDQ	HQF	KILAL	VD	SGKKE	GAKLE	FGG	-----	CAVEDR	-----	GLK	HTP
Raldh4	GALISK	HLQ	KYK	YITL	ALAE	GAQV	HC	GEGV	DKL	LPQQ	NI	GYSML
Consensus	gpqis	eq	kvl	li	sg	egakle	cgg	a	g	f	ptifs	vkdh
	481	491	501	511	521	531	541	551	561	571	581	591
Raldh2	CYNAM	SCC	PF	GGF	KMSG	NGREL	GEIGL	KYTEL	ITITM	MSG	KTS	-----
Raldh3	CYNAL	HA	CT	PF	GGY	KMSG	NGREL	GEYAL	AAYTE	VRAIT	IKL	SEQL
Raldh4	CNLV	RDL	NLP	FG	GM	HSG	IG	REG	GKDS	YHFT	EVKS	VTV
Consensus	cyna	q	pfgg	kmsg	ngrel	ge	l	eyte	vk	it	k	s

Fig. 1. Amino acid sequence alignment of zebrafish *Raldh2*, *Raldh3* and *Raldh4*. The amino acid identities and similarities of zebrafish *Raldh2* with zebrafish *Raldh3* and *Raldh4* are determined using Jellyfish software (www.biowire.com; version 1.4). Identical amino acid residues are crosshatched, and dashes represent gaps for alignment purposes.

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