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	作成者: Suzuki, Shunsuke, Kasai, Kentaro, Yamauchi,
	Kiyoshi
	メールアドレス:
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1	Characterization of little skate (Leucoraja erinacea) recombinant transthyretin:
2	zinc-dependent 3,3',5-triiodo-L-thyronine binding
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4	Shunsuke Suzuki ^{a†} , Kentaro Kasai ^{a†} , Kiyoshi Yamauchi ^{a,b*}
5	
6	^a Department of Biological Science, Graduate School of Science, Shizuoka University, Shizuoka 422-8529,
7	Japan
8	^b Green Biology Research Division, Research Institute of Green Science and Technology, Shizuoka
9	University, Shizuoka 422-8529, Japan
10	
11	
12	†Equally contributing author.
13	*Corresponding author. Green Biology Research Division, Research Institute of Green Science and
14	Technology, Shizuoka University, Shizuoka 422-8529, Japan. Tel.: +81 54 238 4777, fax: +81 54 238
15	0986.
16	E-mail address: fgyfm536@gmail.com (S. Suzuki), mitsuya59451192@gmail.com (K. Kasai),
17	sbkyama@ipc.shizuoka.ac.jp (K. Yamauchi).
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19	Abbreviations: BSA, bovine serum albumin; EDTA, ethylenediaminetetraacetic acid; HIU,
20	5-hydroxyisourate; HIUHase, 5-hydroxyisourate hydrolase; HPLC, high-performance liquid
21	chromatography; IC ₅₀ , 50% inhibitory concentration; ICP-OES, inductively coupled plasma optical
22	emission spectroscopy; IPTG, isopropyl β -D-1-thiogalactopyranoside; K_d , dissociation constant; MBC,
23	maximun binding capacities; PAGE, polyacrylamide-gel electrophoresis; reverse T3 or rT3,
24	3,3',5'-triiodo-L-thyronine; PCR, polymerase chain reaction; SDS, sodium dodecyl sulfate; T3,
25	3,3',5-triiodo-L-thyronine; T4, L-thyroxine; TH, thyroid hormone; TTR, transthyretin.
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28 Abstract

Transthyretin (TTR) diverged from an ancestral 5-hydroxyisourate hydrolase (HIUHase) by gene 29 duplication at some early stage of chordate evolution. To clarify how TTR had participated in the thyroid 30 31 system as an extracellular thyroid hormone (TH) binding protein, TH binding properties of recombinant little skate Leucoraja erinacea TTR was investigated. At the amino acid level, skate TTR showed 37-46% 32 identities with the other vertebrate TTRs. Because the skate TTR had a unique histidine-rich segment in the 33 N-terminal region, it could be purified by Ni-affinity chromatography. The skate TTR was a 46-kDa 34 homotetramer of 14.5 kDa subunits, and had one order of magnitude higher affinity for 35 3,3',5-triiodo-L-thyronine (T3) and some halogenated phenols than for L-thyroxine. However, the skate 36 TTR had no HIUHase activity. Ethylenediaminetetraacetic acid (EDTA) treatment inhibited [¹²⁵I]T3 37 binding activity whereas the addition of Zn^{2+} to the EDTA-treated TTR recovered [¹²⁵I]T3 binding activity 38 in a Zn²⁺ concentration-dependent manner. Scatchard analysis revealed the presence of two classes of 39 binding site for T3, with dissociation constants of 0.24 and 17 nM. However, the high-affinity sites were 40 completely abolished with 1 mM EDTA, whereas the remaining low-affinity sites decreased binding 41 capacity. The number of zinc per TTR was quantified to be 4.5–6.3. Our results suggest that skate TTR has 42 tight Zn^{2+} -binding sites, which are essential for T3 binding to at least the high-affinity sites. Zn^{2+} binding to 43 the N-terminal histidine-rich segment may play an important role in acquisition or reinforcement of TH 44 binding ability during early evolution of TTR. 45 46 Keywords: transthyretin, thyroid hormone, divalent cation, endocrine disruption, 5-hydroxyisourate 47

- 48 hydrolase, Leucoraja erinacea
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53 1. Introduction

Plasma thyroid hormone (TH) binding proteins ensure the adequate distribution of THs in circulation 54 to tissues. Plasma TH-binding proteins consist of thyroxine-binding globulin, transthyretin (TTR) and 55 56 albumin (Robbins, 1996), all of which appeared during vertebrate evolution. However, the phylogenetic distribution of these TH-binding proteins is species-specific. TTR acts as a major TH-binding protein in 57 58 some fish (Santos and Power, 1999; Yamauchi et al., 1999). In addition, lipoproteins and albumin are also known as major TH-binding proteins with low affinity for THs in immature salmonids (Cyr and Eales, 59 1992; Richardson et al., 2005). However, there is a lack of information about the TH binding properties of 60 TTR from lower vertebrates such as elasmobranchs. 61 62 Chondrichthyes or cartilaginous fishes are an extant group that diverged from the earliest gnathostomes about 395 million years ago (Benton, 2005). Most elasmobranchs, including skates, have 63 important ecological positions as top- or meso-predators within marine food webs. However, little is known 64 about the non-natural ligands of skate TTR, including agricultural, industrial, and pharmaceutical chemicals 65

66 that may be expected to displace THs from TTR and act as endocrine disruptors.

The TTR gene arose from an ancestral 5-hydroxyisourate hydrolase (HIUHase) gene by neo-functionalization after gene duplication at some early stage of chordate evolution (Ramazzina et al., 2006; Zanotti et al., 2006). Because this event occurred around the time that the thyroid system was established, the ancestor of TTR is likely to have acquired the ability to bind TH soon after gene duplication. Intriguingly, only two point mutations in zebrafish or amphioxus HIUHases resulted in the acquisition of TH binding activity and the loss of 5-hydroxyisourate (HIU) hydrolysis activity (Cendron et al., 2011; Li et al. 2013).

The N-terminal region of TTR forms a random-coiled structure with high conformational flexibility (Hamilton et al., 1993), and has important roles in the specificity and affinity of TH binding (Prapunpoj et al., 2006). It has been proposed that the N-terminal region has evolved from a longer and more hydrophobic region in lower vertebrates to a shorter and more hydrophilic region in eutherians, by unidirectional changes (Aldred et al., 1997; Prapunpoj et al., 2000), with functional changes in TH-binding specificity from for 3,3',5-triiodo-L-thyronine (T3) to for L-thyroxine (T4) (Yamauchi et al., 1993; Chang et al., 1999). Through the survey of several expressed sequence tag and genome databases of elasmobranch

fishes, we found TTR cDNA in the little skate *Leucoraja erinacea* database (<u>http://skatebase.org</u>), and noticed that such unidirectional changes may not be true for the skate TTR. Because the skate TTR has a highly hydrophilic stretch containing four histidine residues in the N-terminal region as if the skate TTR has a histidine tag. From this unique structure we supposed that functional analyses of the TTR gene superfamily in primitive gnathostomes may provide a good model for understanding the processes of neo-functionalization found in duplicated genes.

As a first step towards elucidating how TTR has participated in the thyroid system as a TH-binding protein during vertebrate evolution, we investigated TH binding properties and HIUHase activity of the recombinant skate TTR.

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92 2. Materials and Methods

- 93 2.1. Reagents
- 94 3,3'[¹²⁵I],5-Triiodo-L-thyronine (81 TBq/mmol, carrier free) was purchased from PerkinElmer
- 95 (Waltham, MA, USA). 3,3',5-Triiodo-L-thyronine, T4, D-T3, 3,3',5'-triiodo-L-thyronine (reverse T3 or
- ⁹⁶ rT3), 3,3',5-triiodothyroacetic acid, 3,3',5,5'-tetraiodothyroacetic acid, diiodo-L-tyrosine,
- 97 monoiodo-L-tyrosine, benzo[*a*]pyrene and pentachlorophenol were obtained from Sigma-Aldrich (St. Louis,
- MO, USA). Bisphenol A, 2,4,6-triiodophenol, 2,4,6-tribromophenol, 2,4,6-trichlorophenol,
- 99 pentabromophenol, *o-t*-butylphenol, mirex, malathion (diethyl
- 100 2-dimethoxyphosphinothioylsulfanylbutanedioate, analytical standard), benzophenone, *n*-butylbenzyl
- 101 phthalate, bis(2-ethylhexyl) adipate, bis(2-ethylhexyl) phthalate, dicyclohexyl phthalate, di-*n*-butyl
- 102 phthalate, tributyltin (IV) chloride, α -benzoepin (α -endosulfan;
- 103 6,7,8,9,10,10-hexachloro-1,5,5a,6,9,9a-hexahydro-6,9,methane-2,4,3-benzodiathiepine), dicofol (Kelthane,
- 104 2,2,2-trichloro-1,1-bis(4-chloro-phenyl)ethanol, analytical standard), urea, Candida sp. uricase and
- 105 trimethylamine N-oxide were purchased from Wako Pure Chemical Industries (Osaka, Japan).
- 106 3,3',5,5'-Tetrabromobisphenol A, 3,3',5,5'-tetrachlorobisphenol A and 3,5-diiodo-L-thyronine were from
- 107 Tokyo Chemical Industry (Tokyo, Japan). 2,4-Dichlorophenoxyacetic acid, 4-nonylphenol, di-2-ethyl
- 108 phthalate and 2,4-dinitrophenol were purchased from Kanto Chemicals (Tokyo, Japan). Isopropyl

β-D-1-thiogalactopyranoside (IPTG) was obtained from Nacalai Tesque (Kyoto, Japan). 2-Isopropylphenol 109 was from Lancaster. Methoprene (isopropyl (E,E')-(RS)-11-methoxy-3,7,11-trimethyldodeca-2,4-dienoate) 110 was from Ehrenstorfer Quality (Augsburg, Germany). Ioxynil (3,5-diiodo-4-hydroxybezonitril, analytical 111 112 standard) and acetochlor (2-chloro-N-(ethoxymethyl)-N-(2-ethyl-6-methylphenyl)acetamide) were obtained from Riedel-de Haën (Seeize, Germany). Ni-resins were from Bio-Rad (Ni-IMAC Profinity, Hercules, CA, 113 114 USA) and Invitrogen (ProBond Nickel-Chelating Resin, Carlsbad, CA, USA). Co-resin was obtained from Clontech (Talon Metal Affinity Resin, Mountain View, CA, USA). All other chemicals used in this study 115 were either chromatography grade or the highest grade available and were purchased from Wako Pure 116 Chemical Industries or Nacalai Tesque. 117

All chemicals tested as endocrine disrupting chemicals were dissolved in dimethylsulfoxide or 0.5 N NaOH to concentrations of 2–5 mM. These chemicals were then diluted with an appropriate buffer. For TTR pull-down assay, a blank assay without proteins with solvent alone at less than 1.0% (v/v) was done. The solvent did not affect the recovery of TTR in the pull-down assay. For competitive [¹²⁵I]T3 binding assay, a control assay without test chemicals was performed in the presence of the solvent alone and at less than 0.4% (v/v). The solvent did not affect the competitive [¹²⁵I]T3 binding assay.

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125 2.2. Protein expression and purification

The skate TTR cDNA (accession no. CV221819, 607 bp) was amplified by polymerase chain reaction 126 127 (PCR) with sense primer 5'-GTTCATATGCCACATAGTCACGGCGACC-3' (64-83) with a recognition site for NdeI (underlined), and antisense primer 5'-GTAGGATCCGGTTTTCACGAGGTGTTAAG-3' 128 (468–449) with a recognition site for BamHI (underlined). PCR was carried out in 50 µL of solution 129 containing 0.2 mM dNTP, 0.2 µM of each primer, 1 ng template DNA and 1.25 U DNA polymerase 130 (PrimeSTAR HS, TAKARA, Siga, Japan), using the following protocol: 98°C (15 s), and 20 cycles of 131 98°C (10 s), 65°C (5 s) and 72°C (30 s), followed by 72°C (5 min). Amplicons (64–449) were cloned 132 between the NdeI and BamHI sites of a pET3a expression vector (Novagen, Madison, WI). Plasmids for 133 expression were transformed into Escherichia coli Rosetta 2(DE3)pLysS (Novagen). Bacteria were usually 134 grown in 50 mL LB medium with 50 μ g/mL ampicillin and 34 μ g/mL chloramphenicol at 37°C until the 135 absorbance at 600 nm reached 0.8. The temperature was lowered to 18°C, 0.1 mM IPTG was added, and 136

incubation was continued for 16 h. Bacteria were pelleted by centrifugation $(1,200 \times g)$ and stored at -35°C until used.

After suspending the bacterial pellet with 5 mL of homogenization buffer (1 mM imidazole, 0.3 M 139 140 NaCl, 50 mM sodium phosphate, pH 8.0, 1 mM benzamidine hydrochloride, and 1 mM phenylmethylsulfonyl fluoride), the cells were disrupted by sonication 20 times for 10 s on ice at the range 141 142 of 5 (UltraS homogenizer, VP-30S, TAITEC, Saitama, Japan), and the lysate was centrifuged at $18,000 \times g$ for 15 min at 4°C. Recombinant proteins were isolated from the other proteins in the supernatant by nickel 143 or cobalt affinity chromatography (0.5 mL of the resin), by stepwise elution with various concentrations of 144 imidazole (5, 20, 60, 150, 250 and 500 mM) in 0.3 M NaCl and 50 mM sodium phosphate, pH 8.0. Peak 145 fractions were immediately applied to a Cellulofine GCL-2000 sf column $(1.5 \times 93 \text{ cm}, \text{Seikagaku Co.},$ 146 Tokyo, Japan), which had been equilibrated with 20 mM Tris, pH 7.5. Purified protein was stored in 10% 147 glycerol at -85° C for later use. The yields of the recombinant proteins were 0.6–0.8 mg/50 mL bacterial 148 culture. The recombinant rainbow trout HIUHase with three histidine residues at the N-terminal end and 149 the recombinant skate TTR were also purified by ammonium sulfate precipitation, hydrophobic interaction 150 151 chromatography on Phenyl Cellulofine column (1.0×10 cm, Seikagaku Co.) and then gel filtration chromatography on Cellulofine GCL-2000 sf column, as described previously (Kasai et al., 2013). 152

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154 2.3. Protein analyses

155 Protein concentration was measured by the micro-Lowry method (Jain et al., 2002) with bovine serum albumin (BSA) as the standard. Proteins was electrophoresed on an sodium dodecyl sulfate 156 (SDS)-polyacrylamide gel (15%) (Laemmli, 1970) with molecular markers: phosphorylase b (97 kDa), 157 BSA (66 kDa), ovalbumin (45 kDa), carbonic anhydrase (30 kDa), trypsin inhibitor (20.1 kDa) and 158 159 α -lactalbumin (14.4 kDa). The gel was stained and proteins were visualized with Coomassie Brilliant Blue. The molecular mass of TTR was estimated by high-performance liquid chromatography (HPLC) on 160 YMC-Pack Diol 120 (500 × 8.0 mm, YMC, Kyoto, Japan) and YMC-Guardpack Diol-12 (30 × 8.0 mm), 161 which were pre-equilibrated and eluted with 50 mM Tris-HCl, pH 7.5, and 140 mM NaCl, at flow rate of 1 162 mL/min at 25°C. 163

165 2.4. Ligand binding assay using a metal chelate affinity resin (TTR pull-down assay)

Iodothyronine or chemical binding was carried out by the TTR pull-down assay using a metal chelate 166 affinity resin (Kasai et al., 2013). Briefly, TTR (0.5 µM) was incubated with iodothyronines or chemicals 167 168 $(0.5 \ \mu\text{M})$ in 500 μL of Tris-buffered saline, pH 7.5, for 1 h at 4°C, and the mixture was further incubated with Co-resin (5 µL bed volume) for 1 h at 4°C in a 600-µL plastic tube using a rotator. After a brief 169 170 centrifugation, the TTR-ligand complex bound to the resin in the pellet was immediately recovered by the addition of 20 µL of elution buffer (500 mM imidazole, 0.3 M NaCl, and 50 mM sodium phosphate, pH 171 8.0). The TTR-ligand complex released from the resin was mixed with the same volume of 1% acetic acid 172 in methanol to extract the ligand from the complex. The mixture was agitated then centrifuged at 173 $12,000 \times g$ for 10 min at 4°C. The ligand (15 µL) in the supernatant was analyzed by reverse phase-HPLC 174 on a C₁₈ analytical column (Mightysil RP-18 GP, 250 mm × 4.6 mm, 5 mm particle diameter, Kanto 175 Chemical, Tokyo, Japan) with a UV detector at 254 nm, as described previously (Kasai et al., 2013). The 176 amount of the ligand was quantified by comparison with the standards (4-100 pmol), with the lower limits 177 of quantitation for iodothyronines and environmental chemicals of 2.0-7.2 pmol and 6-14.2 pmol. 178 respectively, which were three times higher than the lower limits of detection. 179

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181 2.5. [¹²⁵I]T3 binding assay

Unlike bony fishes, cartilaginous fishes retain a high concentration of organic osmolytes, such as urea 182 183 (300–400 mM), in their body fluids to maintain a high plasma osmolality that is similar to that in the external environment. To counteract destabilization of proteins by urea, cartilaginous fishes also have 184 185 methylamines, e.g., trimethylamine N-oxide (Yancey, 2005). Recombinant TTR (7.5 ng/tube, 0.53 nM) was incubated with 0.1 nM [¹²⁵I]T3 in 250 µL of buffer (20 mM Tris, pH 7.5, 280 mM NaCl, 360 mM urea, 186 and 90 mM trimethylamine N-oxide) in the presence or absence of test compounds (cations, unlabeled 187 iodothyronines or environmental chemicals) for 1.0 h at 4°C, unless otherwise noted. In some cases, 188 189 various concentrations of TTR were pre-incubated with 1 mM ethylenediaminetetraacetic acid (EDTA). After dilution with the buffer, TTR (0.53 nM) was used for $[^{125}I]T3$ binding assay in the presence or 190 absence of cations. For the Scatchard plot of T3 binding to both high-affinity and low-affinity sites, skate 191 TTR was incubated with 0.1 nM [¹²⁵I]T3 in the presence or absence of increasing concentrations of 192

193	unlabeled T3. The dissociation constant (K_d) value for high affinity sites was expected to be 10 ⁻¹⁰ M from
194	this experiment. To estimate more accurately the K_d value, TTR was incubated with [¹²⁵ I]T3 (0.052 to 0.4
195	nM) in the presence or absence of 5 μ M of unlabeled T3. [¹²⁵ I]T3 bound to proteins was separated from
196	free [¹²⁵ I]T3 by the polyethyleneglycol method (Yamauchi et al., 1993). The radioactivity of the samples
197	was measured in a gamma counter (2480 WIZARD ² ; PerkinElmer). The amount of [¹²⁵ I]T3 bound
198	non-specifically was derived from the radioactivity of samples incubated with 5 μ M unlabeled T3. The
199	non-specific binding value was subtracted from the amount of total bound [¹²⁵ I]T3 that was derived from
200	the radioactivity of samples incubated without unlabeled T3, to give the value of specifically bound
201	$[^{125}I]T3$. The K_{ds} and maximum binding capacities (MBCs) for T3 binding were determined from
202	Scatchard plots for a single class (Scatchard, 1949) or for two classes (Rosenthal, 1967).

203

204 2.6. Determination of the average number of divalent cations per TTR

TTR samples were prepared by Co-affinity chromatography and gel filtration chromatography on 205 Cellulofine GCL 2000-sf column. Alternatively, to avoid the effects of Co-resin on the metal contents of 206 TTR, TTR samples were also prepared by ammonium sulfate precipitation, hydrophobic interaction column 207 chromatography, and gel filtration column chromatography, as shown previously (Kasai et al., 2013). The 208 209 content of divalent cations in TTR samples (1.5 mL in 4 mM Tris, pH 7.5, 0.16-0.65 mg/mL) was quantified three times by inductively coupled plasma optical emission spectroscopy (ICP-OES) (Optima 210 211 3300 DV, PerkinElmer) using calibration curves of zinc, copper, nickel, cobalt and iron standards (0, 0.01, 0.05, 0.1, 0.5, 1.0, 5.0 mg/L in 0.1 mol/L HNO₃) (Wako). The lower limits of quantitation were 0.011 212 213 mg/mL, and the lower limits of detection were 0.002 mg/mL.

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215 2.7. HIU hydrolysis assay

HIU hydrolysis assay was conducted according to the previous method (Kasai et al., 2013). Enzyme activity was measured at 25–27°C by following the decrease in absorbance at 312 nm as HIU was hydrolyzed (Lee et al., 2005). 5-Hydroxyisourate was generated *in situ* by the addition of 50 μ L of 0.32 μ M uricase in 50 mM potassium phosphate buffer, pH 7.5, to 400 μ L of 100 μ M uric acid in the same buffer. When the amount of HIU reached a maximum (2.4 min after starting the reaction), 10 μ L of HIUHase (2

μg) or TTR (2 μg) was added to the mixture in a microcuvette. Absorbance at 312 nm was immediately 221 monitored at 0.2 min intervals using a spectrophotometer (U-3210, Hitachi, Japan). The control assay was 222 carried out similarly except that HIUHase or TTR solution was replaced by phosphate buffer. As HIU is 223 224 not stable, changes in absorbance at 312 nm does not reflect correctly the enzymatic hydrolysis of HIU. Therefore, the enzyme activity was expressed as the difference in absorbance at 312 nm between the 225 226 presence (spontaneous + enzymatic hydrolysis of HIU) and the absence (spontaneous hydrolysis of HIU) of proteins. 227 228 2.8. Statistics 229 The data are the mean \pm standard error of the mean ($n \ge 3$), unless otherwise noted. Differences 230 between groups were analyzed by a one-way analysis of variance, with the Scheffe's test for multiple 231 comparisons. P < 0.05 was considered statistically significant. 232 233 234 235 3. Results 3.1. Comparison of TTR amino acid sequences 236 The skate TTR amino acid sequence shared 37-46% identities with the other vertebrate TTR amino 237 acid sequences shown in Fig. 1, although the skate TTR amino acid sequences shared only 24-27% 238 239 identities with the amphioxus and trout HIUHase amino acid sequences (data not shown). Significant variations of amino acid residues were detected at the N-terminal regions, where the skate and lamprey 240 241 TTR had a histidine-rich segment of high hydrophilicity, HSHGDHH and HKSHESH, respectively. Thirty 242 residues were invariant in the amino acid sequences among the 9 TTR sequences. 243 3.2. Molecular features of recombinant skate TTR proteins 244Skate TTR could be purified by nickel-affinity chromatography (Fig. 2A) and gel filtration 245 chromatography. Skate TTR molecule may be a slightly compact tetramer. The molecular mass of the skate 246 TTR under native conditions was estimated to be ~46 kDa by gel filtration column chromatography (Fig. 247 2B). This value was somewhat smaller than the molecular mass of the tetramer (56.8 kDa) estimated from 248

the amino acid sequence. The molecular mass for the TTR polypeptide was 14.5 kDa on

250 SDS-polyacrylamide-gel electrophoresis (PAGE) (Fig. 2C).

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252 3.3. Binding of iodothyronines and environmental chemicals to recombinant skate TTR using TTR

- 253 pull-down assay
- 254 3,3',5-Triiodo-L-thyronine bound to skate TTR then extracted with methanol was eluted at the same
- retention time (11.46 min) on reverse phase-HPLC as the T3 standard, as shown previously (Kasai et al.,
- 256 2013). The peak of T3 was negligible when the same pull-down assay was done without TTR (blank).

257 Skate TTR exhibited high binding activity for natural ligands T3 and T4, and related compounds, with the

- following rank order affinity: $D-T3 \ge 3,3',5,5'$ -tetraiodothyroacetic acid, T3, T4 and
- 3,3',5-triiodothyroacetic acid \geq reverse T3 > 3,5-diiodo-L-thyronine (Fig. 3A). With regard to

environmental chemicals (Fig. 3B), skate TTR showed binding activity for triiodophenol and ioxynil, but

261 not for benzophenone and 2,4-dinitrophenol.

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263 3.4. [¹²⁵I]T3 binding activity of recombinant skate TTRs

264 3.4.1. Effects of divalent cations

To test the possibility that the skate TTR has significant affinity for metal ions, [¹²⁵I]T3 binding was 265 performed in the presence of 1 mM divalent cations. However, any cations tested did not show statistically 266 significant effects on [¹²⁵I]T3 binding (Fig. 4A). We next pre-incubated skate TTR (0.53–2640 nM) with 267 EDTA (1 mM) at various molar ratios followed by [¹²⁵I]T3 binding assay at 0.53 nM of TTR, to determine 268the most effective molar ratio of TTR/EDTA for eliciting the chelating action of EDTA (Fig. 4B). [¹²⁵I]T3 269 binding was decreased by more than a half at the TTR pre-incubation concentration of 264 nM 270 (TTR/EDTA molar ratio of 1/3,800), and by about 90% at the TTR pre-incubation concentration of 26.4 271 nM (TTR/EDTA molar ratio of 1/38,000). Using the TTR pre-treated with 1 mM EDTA at the TTR/EDTA 272 molar ratio of 1/38,000, the effects of divalent cations (1 mM) on [¹²⁵I]T3 binding to TTR (0.53 nM) was 273 investigated (Fig. 4C). Zn^{2+} , Ni²⁺ and Mn²⁺ (each 1 mM) recovered [¹²⁵I]T3 binding activity up to 274 60–90% of that of the TTR not pre-treated with EDTA. Ca^{2+} and Mg^{2+} had little effect or were not effective. 275 The effects of Fe^{2+} , Co^{2+} , and Cu^{2+} on $[^{125}I]T3$ binding could not be determined, because of the formation 276

of insoluble materials under our assay conditions (data not shown). Addition of 20 μ M Zn²⁺ activated partially [¹²⁵I]T3 binding in spite of the presence of the residual EDTA (final 20 μ M; Fig 4D). [¹²⁵I]T3 binding activity reached a plateau at more than 50 μ M Zn²⁺.

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281 3.4.2. Competitive inhibition of $[^{125}I]T_3$ binding to recombinant skate TTR by iodothyronines and other 282 compounds

[¹²⁵I]T₃ binding was investigated in the presence of various concentrations of iodothyronines and their related compounds (Fig. 5). The most powerful competitor was 3,3',5-triiodothyroacetic acid with 50% inhibitory concentration (IC₅₀) of 1.0 ± 0.2 nM, which was two times less than the IC₅₀ for T3 (Table 1).

286 The rank order binding affinity was 3,3',5-triiodothyroacetic acid \geq T3 \geq

 $D-T3 \ge 3,3',5,5'-tetraiodothyroacetic \ acid > T4 > reverse \ T3 > 3,5-diiodo-L-thyronine > diiodo-L-tyrosine > diiodo-L-tyr$

and monoiodo-L-tyrosine. T3 was 13 times more potent than T4. Acetic acid analogs

(3,3',5-triiodothyroacetic acid and 3,3',5,5'-tetraiodothyroacetic acid) were 2-3 times more potent than the
 corresponding iodothyronines (T3 and T4). Iodotyrosines had no affinity for skate TTR.

As a primary screening of environmental chemicals that interfere with $[^{125}I]T_3$ binding to skate TTR in vitro, inhibitory effects of chemicals on $[^{125}I]T_3$ binding was investigated at 1 µM concentration. Out of 29 chemicals tested, 8 chemicals, all of which were phenolic compounds consisting of one or two phenolic rings with several halogen atoms, strongly inhibited $[^{125}I]T_3$ binding (Fig. 6A). Phenolic compounds without halogen atoms, phthalates and the other chemicals tested were not significant competitors, even if they had benzene or phenolic ring(s).

[¹²⁵I]T₃ binding in the presence of varying concentrations of the eight chemicals that strongly inhibited [¹²⁵I]T₃ binding was tested (Fig. 6B). The most effective inhibitor of [¹²⁵I]T₃ binding to skate TTR was pentabromophenol, with an IC₅₀ of 1.4 ± 0.3 nM, which was between those for 3,3',5-triiodothyroacetic acid and T3 (Table 1). The rank order binding affinity was pentabromophenol \geq triiodophenol \geq pentachlorophenol \geq 3,3',5,5'-tetrabromobisphenol A \geq 3,3',5,5'-tetrachlorobisphenol A and ioxynil \geq

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304 3.4.3. Kinetics of
$$[^{125}I]T_3$$
 binding to skate TTR

trichlorophenol and tribromophenol.

Scatchard plots revealed the presence of two classes of binding sites for T3 (Fig. 7), with K_{d1} and K_{d2} 305 of 0.52 ± 0.03 nM and 17 ± 3 nM, and MBC₁ and MBC₂ of 2.2 ± 1.0 pmol/µg protein and 19 ± 5 pmol/µg 306 protein, respectively. To determine more precisely the K_{d1} value, we conducted Scatchard analysis at lower 307 concentrations and within a more narrow range of [¹²⁵I]T3, which showed the K_{d1} value of 0.24 ± 0.04 nM 308 (Fig. 7 inset). In the presence of 1 mM EDTA, the Scatchard plot showed a single class of binding sites, 309 where high-affinity sites were completely abolished. The K_d value was estimated to be 21 ± 1 nM, which 310 nearly corresponded to that for the low-affinity sites of the TTR untreated with EDTA, with decreased 311 312 MBC value, 1.4 ± 0.1 pmol/µg protein (Table 2). 313

314 3.5. HIU hydrolysis activity of recombinant skate TTR and recombinant rainbow trout HIUHase

As the production of HIU, a substrate for HIUHase, reached a peak at 2.4 min after the addition of uricase into the reaction mixture (Fig. 8), we investigated HIUHase activity from this time point by adding trout HIUHase or skate TTR. Hydrolysis of HIU by the trout HIUHase was clearly detected at the concentration of 4.4 μ g/mL. Activity (difference in absorbance at 312 nm between the samples in the absence and the presence of HIUHase) peaked at 2.4–4.0 min and decreased thereafter. However, there was no significant effect of the skate TTR on the hydrolysis of HIU at the same concentration (Fig. 8).

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322 *3.6.* Contents of divalent cations in skate TTR molecule

The zinc content measured by using ICP-OES was 4.47–6.25 mol/mol TTR (Table 3). Additionally, nickel and copper were also detected at 0.40 and 0.36–2.29 mol/mol TTR, respectively. The contents of cobalt and iron were below detectable levels. The TTRs purified using Co-resin and hydrophobic interaction chromatography had similar total metal contents, 6.76–7.01 mol/mol TTR. The same fractions on gel chromatography of the cell extract, which were obtained from *E. coli* harboring intact pET3a, did not contain detectable levels of the metals analyzed.

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331 4. Discussion

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The present study demonstrates that skate TTR is a zinc-dependent TH-binding protein. This feature is

supported by the presence of a histidine-rich segment in the N-terminal region, the ability to be purified by metal chelate affinity chromatography, inactivation of T3 binding by EDTA and recovery of T3 binding activity by divalent cations. Skate TTR had high-affinity and low-affinity sites for T3 with K_{ds} of 0.24 nM and 17 nM, but not HIU hydrolysis activity. All of the high affinity and most of the low affinity sites for T3 were abolished by the treatment of EDTA. Analysis by IPC-OES revealed that the skate TTR molecule contains 4.47–6.25 zinc ions. Therefore, zinc ion may be essential for the skate TTR to bind T3.

We discovered that a histidine-rich segment, HSHGDHH, located in the N-terminal region of skate 339 TTR may act as a natural histidine tag, which facilitated the purification of this protein by Ni-affinity 340 chromatography and may also provide a novel ligand binding assay (TTR pull-down assay) for 341 investigating the ligand interaction. The histidine-rich segments of the skate and lamprey TTRs resemble 342 343 the zinc-binding motif of metalloproteases (HEXXH or HEXXHXXGXXH/D) (Cerdà-Costa and Gomis-Rüth 2014), but are quite different from the metal-binding motif of metallothionein with 344 cysteine-rich motifs (Vašák aand Meloni, 2011). TTR is a homotetramer consisting of a dimer of dimers 345 (subunits A and B, and subunits C and D). As the N-terminal regions of two subunits (A and D, or B and C) 346 are located in the vicinity of the entrance of either hormone binding pocket (Hamilton et al., 1993), the two 347 histidine-rich segments in the N-terminal regions are highly likely to act as a histidine tag comparable with 348 the ordinary 6× histidine-tag. In this study, we do not exclude the possibility of the presence of several 349 binding sites for divalent cations in skate TTR molecule besides the N-terminal region. 350

351 Divalent cation binding to skate TTR is necessary to elicit strong T3 binding activity that may occur at physiological conditions, in which Zn^{2+} is the best possible candidate cation. [¹²⁵I]T3 binding to skate TTR 352 353 was inactivated by EDTA and re-activated by divalent cations in vitro. When skate TTR (264 nM) was pre-treated with EDTA (1 mM) at the molar ratio of 1/3,800, [¹²⁵I]T3 binding to the skate TTR was 354 partially inhibited. This suggests that most of the zinc ions involved in T3 binding do not dissociate from 355 the TTR of the micromolar range. The concentration of Zn^{2+} that was needed to reactivate the [¹²⁵I]T3 356 binding activity, ~0.05 mM, was less than the physiological range (0.1–0.2 mM) of Zn^{2+} in fish plasma 357 (Sturrock et al. 2013). Scatchard analysis revealed the presence of two classes of binding sites. [¹²⁵]T3 358 binding to all of the high affinity sites and most of the low affinity sites were lost with the addition of 1 359 mM EDTA. This finding leads us to the possibilities that Zn^{2+} binding is essential for skate TTR to express 360

at least high affinity sites for T3 binding and that Zn^{2+} at the concentration in plasma may modulate T3 binding to skate TTR in vivo. In this study we determined the average number of metal ions per TTR. If skate TTR samples consist of a mixture of the TTR components with different content of Zn^{2+} or the Zn^{2+} content of skate TTR in circulation differs locally, a variety of the number of Zn^{2+} per TTR may increase the functional diversity in T3 binding activity. Future study will need to address the physiological meanings of Zn^{2+} binding to TTR on the skate thyroid system.

This is the first report showing the divalent-dependent TH binding activity of TTR, although there are 367 several reports regarding the effects of metal ions on the other functions of human TTR. Zn²⁺ binding 368 decreases monomer stability to enhance TTR aggregation at an acidic pH (Palmieri et al., 2010) or amyloid 369 370 formation of a mutant TTR (Wilkinson-White and Easterbrook-Smith, 2007), and decreases the affinity of TTR for RBP4 (Palmieri et al., 2010). Zn^{2+} is also essential for expressing the metalloprotease activity of 371 human TTR, which was inhibited by EDTA or 1,10-phenanthroline and reactivated by divalent cations (Liz 372 et al., 2012). Zinc content in human TTR was ~0.2 per TTR molecule (Liz et al., 2012), which was one 373 order less than that in the skate TTR (4.47-6.25 per TTR). In addition, the skate TTR contained copper at 374 0.36–2.29 per TTR, and nickel at 0.4 per TTR. This means that four fifths of human TTR is independent of 375 zinc-dependent functions. However, in the skate TTR, because of the high content of zinc, T3 binding 376 activity which is highly zinc-dependent may be fully affected by the zinc content. Therefore, the deficiency 377 of metal ions such as zinc may influence the functions of TTR to greater extent in skates compared with 378 379 humans.

Zn²⁺ binding may be an ancient property of the TTR/HIUHase superfamily with high diversity. In the 380 TTR family, Zn^{2+} binding is experimentally shown so far in the skate and human protein alone. Among the 381 four Zn²⁺ binding sites (ZBS1 to ZBS4) proposed in human TTR (Palmieri et al., 2010), only ZBS1 (C10 382 and H56 in Fig. 1) is conserved in the vertebrate TTRs investigated. Among ten amino acid residues that 383 constitute the four Zn²⁺ binding sites in the human TTR, four are identical and one is similar to the 384 corresponding amino acid residues in the skate TTR. The histidine-rich segments found in the skate and 385 lamprey TTRs are not detected in the bony fish and tetrapod TTRs (Fig.1) or HIUHases of various 386 vertebrates. However, in the HIUHase family, our previous study (Kasai et al., 2013) showed that Zn^{2+} (50 387 μM) inhibits significantly the HIU hydrolysis activity of rainbow trout HIUHase. The protein structure of E. 388

coli HIUHase revealed the presence of zinc ions bound to the functional sites (Lundberg et al., 2006). All 389 of these circumstances suggest that Zn^{2+} binding may be a common property of the HIUHase/TTR 390 superfamily. Interestingly, the number of histidine residues in TTR polypeptide has changed during 391 392 vertebrate evolution: 11 for lamprey and skate, 7–8 for bony fishes, 6 for amphibians, 5 for reptiles and a bird and 4 for humans (Fig. 1). It is likely that TTR has evolved from a strong to weak divalent cation 393 394 binder (Wang et al., 2013). Although there is no experimental data to explain why the histidine content in TTR sequences is high in lamprey and skate TTRs and continuously decreased during vertebrate evolution, 395 some of these histidine residues may participate in divalent cation binding and contribute to the acquisition 396 or reinforcement of TH binding activity in an ancestor of the agnathan and cartilaginous fish TTRs soon 397 after TTR diverged from ancestral HIUHase. 398

Skate TTR showed the highest affinity for T3 among the vertebrate TTRs studied so far, with K_{d1} of 399 10^{-10} M, which is comparable to those of nuclear TH receptors ($K_d = 10^{-11} - 10^{-10}$ nM) (Oppenheimer et al., 400 1996). Giving that plasma TH levels are 10^{-9} – 10^{-8} M in several elasmobranchs: the whitetip reef shark 401 Triaenodon obesus (Crow et al., 1999), dogfish Squalus acanthias (Leary et al., 1999) and Atlantic stingray 402 Dasyatis Sabina (Volkoff et al., 1999), skate TTR may be functional as a T3-binder rather than T4-binder. 403 The preference for T3 over T4 was also detected in bony fish (Kasai et al., 2013; Kawakami et al., 2006; 404 Santos and Power, 1999; Yamauchi et al., 1999), amphibian (Prapunpoj et al., 2000; Yamauchi et al., 1993), 405 reptilian (Prapunpoj et al., 2002) and bird (Chang et al., 1999), but not in mammalian TTRs (Chang et al., 406 407 1999), suggesting that TTR has originally evolved as a T3-binder during the early evolution of fish and changed as T4-binder during the early evolution of mammals (Richardson, 2007; 2015). 408

409 We applied the TTR pull-down assay for ligand binding to skate TTR. Ligand preference of TTR tested by this assay agreed well with that tested by the $\int^{125} \Pi T3$ competitive binding assay for endogenous 410 and exogenous ligands. The TTR pull-down assay is a convenient and a non-radioisotopic method that can 411 estimate directly and semi-quantitatively the ligand-TTR interaction. Furthermore, the TTR-pull-down 412 assay can detect several potent chemicals with different retention times on HPLC at the same time. The 413 414 drawback is that relatively high concentrations of ligands and TTR are needed to quantify their amounts by HPLC, and that the accuracy of this assay is not high. Therefore, this assay may be suitable for the primary 415 screening that allows to estimate semi-quantitatively environmental contaminants with significant affinity 416

417 for TTR and with significantly high absorption spectrum in the ultraviolet or visible region.

Skate TTR has significant high affinity for several halogenated phenolic compounds that structurally 418 resemble THs (Howdeshell, 2002). Tetrabromobisphenol A and pentabromophenol were the most potent 419 420 chemicals in halogenated phenolic compounds with two phenolic rings and with a single ring, respectively. The structural preference of ligand binding was well conserved in vertebrate evolution (Kudo et al., 2006; 421 422 Meerts et al., 2000; Morgado et al., 2007; Yamauchi et al., 2003). Furthermore, in the structure-activity relationships, halogen species-dependency (Br > Cl in pentahalogenated phenols; $I > Br \approx Cl$ in 423 trihalogented phenols; Br > Cl > no halogen in bisphenol A) and halogen number-dependency (penta-424 >tri- in brominated and chlorinated phenols) are similar with previous studies using human (Meerts et al., 425 2000), chicken and amphibian TTRs (Kudo et al., 2006; Yamauchi et al., 2003). 426 Production of brominated phenols, used as a flame retardant or its intermediate, or as a wood 427 preservative, may result in the release of brominated phenols to the aquatic environment for several 428 decades (WHO, 2005). The annual consumption by humans of tetrabromobisphenol A is 89,400 tons in 429 Asia, which is 56% of the total market demand in 2001 (Birnbaum and Staskal, 2004). As skates spend 430 431 time in the sediments of the sea, where they feed on various invertebrates including decapod crustaceans, amphipods and polychaete worms, they are ecologically situated in the top or middle of the benthic food 432 web. This ecological position may expose skates to hydrophobic and persistent organic pollutants retained 433 in the benthic environments with a relatively high bioaccumulation. Therefore, investigating the inhibitory 434 435 effects of persistent halogenated compounds on TH binding to skate TTR may be useful to assess the threat on the thyroid system from environmental chemicals in benthic environments. 436

In conclusion, we have demonstrated that skate TTR is a strong T3-binder in vitro, and that T3 binding to skate TTR is divalent cation-dependent and competitively inhibited by environmental phenolic compounds with halogens. The unique structure of a histidine-rich segment in the N-terminal region may act as a natural histidine tag, raising the possibility that the TH binding activity of the skate TTR may be modulated by Zn^{2+} in plasma. We proposed that a histidine-rich segment in the N-terminal region of TTR may play an important role in acquisition or reinforcement of TH binding activity via metal binding during early vertebrate evolution.

445 Conflict of interest stat

446 The authors declare that there are no conflicts of interest.

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585 Figure legends

Fig. 1. Comparison of transthyretin (TTR) sequences with possible zinc-binding sites. Mature TTR amino 586 acid sequences from lamprey (Manzon et al., 2007), skate (accession no. CV221819), trout (Yamauchi et 587 588 al., 1999; accession.no.CX256523), seabream (Santos and Power, 1999), toad (Prapunpoj et al., 2000), bullfrog (Yamauchi et al., 1993, 1998), crocodile (Prapunpoj et al., 2002), chicken (Duan et al., 1991) and 589 590 human (Kanda et al., 1974) were aligned. The N-terminal ends of the lamprey, skate and seabream TTRs are predicted, whereas those of the other TTRs are based on Edman degradation. The numbering of amino 591 acid is based on that of human TTR. Features of secondary structure of human TTR (Hamilton et al., 1993) 592 are indicated above the lamprey sequence. Histidine-rich segments in the disordered N-terminal region of 593 the skate and lamprey TTRs are boxed. A segment that resembles a zinc-binidng motif of metalloprotease 594 (HEXXHXXGCCH/D) (Cerdà-Costa and Gomis-Rüth, 2014) is underlined. Four zinc-binding sites (ZBSs 595 1-4) proposed in human TTR (Palmieri et al., 2010) are also boxed: C10 and H56 for ZBS1, H88, H90 and 596 E92 for ZBS2, H31, D74 and E72 (or E62 at acidic pH), H31, E61 and D74 for ZBS4. Marks (*) 597 underneath the sequence alignment indicate positions where amino acid residues are invariant, among the 9 598 599 sequences. The numbers of histidine residues in the TTR sequences are shown on the right.

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Fig. 2. Purification of recombinant skate transthyretin (TTR) by metal chelate affinity chromatography and 601 gel chromatography. (A) SDS-PAGE of the fractions of metal chelate affinity chromatography. The 602 603 recombinant skate TTR was separated from the other bacterial proteins in cell extract by metal chelate affinity chromatography on Ni-resin, and then eluted with the buffer containing 150 mM imidazole (an 604 605 arrow in fraction E4). Lanes from left to right: ppt, pellet of cell lysate; sup, supernatant of cell lysate; FT, flow-through fraction; W1 to W4, washing fractions; E1 to E6, elution fractions with buffer containing 5, 20, 606 60, 150, 250, 500 mM imidazole, respectively. (B) High performance gel chromatography of purified 607 recombinant skate TTR. Protein was applied to YMC Diol 120 High performance gel chromatography 608 column (8 × 500 mm) in 50 mM Tris-HCl, pH 7.5, and 140 mM NaCl, at 1 mL/min. Elution was monitored 609 at absorbance at 280 nm. Markers: Blue dextran (void volume, V_0); apoferritin (Fer, 440 kDa); alcohol 610 dehydrogenase (ADH, 158 kDa); bovine serum albumin (BSA, 68 kDa); ovalbumin (OVA, 45 kDa); and 611 myoglobin (Mb, 17 kDa). (C) SDS-PAGE of the peak fraction of Cellulofine GCL-2000 sf gel filtration 612

613 chromatography. Arrow, band of skate TTR.

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Fig. 3. Ligand-binding to recombinant skate transthyretin (TTR) determined by the TTR pull-down assay 615 616 using metal chelate affinity resin. After incubating TTR (14.2 μ g, 0.5 μ M) with each ligand (0.5 μ M) (A, iodothyronines and their acetic acid analogs; B, environmental chemicals) in 500 μ L of the incubation 617 buffer, the ligand bound to TTR was analyzed by reverse phase-HPLC as described in the Materials and 618 methods. The vertical axis of graphs indicates the amount of the bound ligand that was recovered in 40 µL 619 of the extraction buffer. Blank data obtained from samples in the absence of TTR were subtracted from 620 those obtained from samples in the presence of TTR. Each value is the mean \pm standard error of the mean 621 (n = 5-6 for panel A; n = 3 for panel B). Different letters indicate significantly different means (p < 0.01). T3, 622 3,3',5-triiodo-L-thyronine; T4, L-thyroxine; Triac, 3,3',5-triiodothyroacetic acid; Tetrac, 623 3,3',5,5'-tetraiodothyroacetic acid; rT3, 3,3',5'-triiodo-L-thyronine, T2, 3,5-diiodo-L-thyronine, D-T3, 624 3,3',5-triiodo-D-thyronine, TIP, 2,4,6-triiodophenol, 2,4-DNP, 2,4-dinitrophenol. 625 626 Fig. 4. Effects of ethylenediaminetetraacetic acid (EDTA) and divalent cations on [¹²⁵I]T3 binding to 627

recombinant skate transthyretin (TTR). (A) Effect of divalent cations of [¹²⁵I]T3 binding to skate TTR. TTR 628 (0.53 nM) was incubated with 0.1 nM [¹²⁵I]T3 in the presence or absence (control) of 1 mM each divalent 629 cation for 1 h at 4°C. (B) Effect of pre-treatment of TTR at various concentrations with 1 mM EDTA on 630 ¹²⁵]T3 binding. TTR (0.53–2640 nM) was pre-incubated with or without (control) 1 mM EDTA for 30 min 631 for 4°C. After dilution 1, 5, 50, 500, or 5,000 times with the buffer, TTR (0.53 nM) was incubated with 0.1 632 nM [¹²⁵I]T3 for 1 h at 4°C. (C) Effects of divalent cations on [¹²⁵]T3 binding to the TTR pre-treated with 633 EDTA. TTR (26.4 nM) was pre-incubated with or without (control) 1 mM EDTA for 30 min for 4°C. After 634 dilution 50 times with the buffer (final concentration of EDTA = 0.02 mM), TTR (0.53 nM) was incubated 635 with 0.1 nM [¹²⁵I]T3 in the presence or absence of 1 mM of each cation for 1 h at 4°C. Binding activity of 636 the TTR that was treated with neither EDTA nor cation (EDTA-, cation -) is indicated as the control (100%). 637 (D). Zn²⁺-dependent activation of [¹²⁵]T3 binding to TTR. TTR (26.4 nM) was pre-incubated with 1 mM 638 EDTA for 30 min for 4°C. After dilution 50 times with the buffer, TTR (0.53 nM) was incubated with 0.1 639 nM $[^{125}I]T3$ in various concentrations of Zn²⁺ for 1 h at 4°C. $[^{125}I]T3$ binding activity of the TTR that was 640

untreated with 1 mM EDTA (*open circle*) is indicated as control 100%. Each value is the mean \pm standard error of the mean of triplicate determinations. Experiments were repeated two or three times. Asterisks indicate significant differences in T3 binding activity between the reference sample [cation (-) in *panel A*; EDTA (-) in *panel B*; EDTA(+) and cation (-) in *panels C* and *D*] and each test sample. *, *p* < 0.05; **, *p* < 0.01.

- 646
- Fig. 5. Competition of iodothyronines and their related compounds for $[^{125}I]T3$ binding to recombinant skate transthyretin (TTR). TTR (0.53 nM) was incubated with 0.1 nM $[^{125}I]T3$ for 1 h at 4°C, in the
- 649 presence or absence (control) of increasing concentrations of the following compounds:
- 650 3,3',5-triiodothyroacetic acid (Triac), T3, D-T3, 3,3',5,5'-tetraiodothyroacetic acid (Tetrac), T4, reverse T3
- (rT3), 3,5-diiodo-L-thyronine (T2), 3,5-diiodo-L-tyrosine (Y2) and 3-iodo-L-tyrosine (Y1). Non-specific

binding, which was 3–4 % of total binding, was subtracted from total binding to give values for specific

binding. Each value is the mean of triplicate determinations. These experiments were repeated at least threetimes.

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Fig. 6. Inhibition of [¹²⁵I]T₃ binding to recombinant skate transthyretin (TTR) by various environmental 656 chemicals. (A) TTR (0.53 nM) was incubated with 0.1 nM [¹²⁵I]T3 in the presence of 1 µM of each 657 chemicals or solvent (dimethylsulfoxide, control) for 1 h at 4°C. Asterisks indicate significantly differences 658 in $[^{125}I]T_3$ binding activity between control and test samples containing each environmental chemical (**, 659 p < 0.01). (B) TTR (0.53 nM) was incubated with 0.1 nM [¹²⁵I]T3 for 1 h at 4°C, in the presence or absence 660 (control) of increasing concentrations of the following compounds: pentabromophenol (PBP), triiodophenol 661 (TIP), pentachlorophenol (PCP), tetrabromobisphenol A (Br4-BPA), tetrachlorobisphenol A (Cl4-BPA), 662 ioxynil, tribromophenol (TBP) and trichlorophenol (TCP). To estimate the relative potency of the test 663 chemicals against T3, the competition curve for unlabeled T3 (broken line) was inserted. Non-specific 664 binding, which was less than 6% of total binding, was subtracted from total binding to give values for 665 specific binding. Each value is the mean of triplicate determinations. These experiments were repeated at 666 least three times. 667

Fig. 7. Scatchard plots of [¹²⁵I]T3 binding to recombinant skate transthyretin (TTR). For high-affinity and 669 low affinity sites, TTR was incubated with 0.1 nM [¹²⁵I]T3 M in the presence or absence of various 670 concentrations of unlabeled excess T3 (5 µM) in the incubation buffer with (•; 5.3 nM TTR) or without (o; 671 0.53 nM TTR) 1 mM EDTA for 1 h at 4°C. For high-affinity sites, TTR (0.53 nM) was incubated with 672 125 IT3 ranging from 0.05 to 0.4 nM in the presence or absence of unlabeled excess T3 (5 μ M) in the 673 incubation buffer for 1 h at 4°C (inset). Non-specific binding, which was 12-16% of total binding, was 674 subtracted from total binding to give values for specific binding. Each value is the mean of triplicate 675 determinations. These experiments were repeated three or four times. v, molar ratio of bound [¹²⁵I]T3 to 676 TTR concentration. 677

678

Fig. 8. Time course of enzyme-dependent 5-hydroxyisourate (HIU) hydrolysis by recombinant rainbow trout 5-hydroxyisourate hydrolase (HIUHase) and recombinant skate transthyretin (TTR). HIUHase, TTR and blank (buffer) were added to the reaction solution containing HIU (*arrow*). The hydrolysis of HIU was monitored spectrophotometrically at 312 nm as described in the Materials and Methods. Enzyme-dependent HIU hydrolysis indicates the difference in absorbance at 312 nm between the samples in the absence (spontaneous degradation of HIU) and the presence of HIUHase or TTR (spontaneous + enzymatic degradation of HIU). These experiments were repeated at least three times, with similar results.

Compounds	$IC_{50} \pm $	SEM (n)		Relative potency*	
	(nM)				
3,3',5-Triiodo-L-thyronine	2.0 \pm	0.1 (3)	a,b**	1	
L-Thyroxine	26 ±	5 (3)	d	0.078	
3,3´,5´-Triiodo-L-thyronine	81 ±	6 (3)	e	0.025	
3,3´,5-Triiodo-D-thyronine	4.4 ±	0.3 (3)	b,c	0.46	
3,3',5-Triiodothyroacetic acid	1.0 ±	0.2 (3)	а	2.0	
3,3',5,5'-Tetraiodothyroacetic acid	9.7 ±	0.8 (3)	c	0.21	
3,5-Diiodo-L-thyronine	250 ±	60 (3)	f	0.008	
3,5-Diiodo-L-tyrosine	1000>>	(3)			
3-Doiodo-L-tyrosine	1000>>	(3)			
Tetrachlorobisphenol A	27 ±	5 (4)	d,e,f,g	0.074	
Tetrabromobisphenol A	9.8 ±	2.3 (4)	c,d,e	0.20	
Pentachlorophenol	5.1 ±	0.5 (3)	b,c	0.39	
Pebtabromophenol	1.4 ±	0.3 (4)	a	1.4	
Triiodophenol	2.2 ±	0.4 (3)	a,b	0.91	
Tribromphenol	49 ±	11 (3)	g	0.04	
Trichlorophenol	44 ±	8 (3)	f,g	0.05	
Ioxynil	33 ±	6 (3)	e.f.g	0.06	

Table 1. Relative potency of ligand binding to skate transthyretin.

*Relative potency is given as the ratio of IC_{50} for T3 to IC_{50} for a chemical. **Different letters indicate significantly different means and were determined by one-way analysis of variance with Scheffe's test for multiple comparisons (p < 0.01).

Chelator	Class	$K_{\rm d}$ (nM)	MBC (pmol/µg protein)
(-) EDTA	two classes (high affinity site) (low affinity site)	0.52 ± 0.03 (4) 17 ± 3 (4)	$2.2 \pm 1.0 (4) 19 \pm 5 (4)$
	one class (high affinity site)	0.24 ± 0.04 (4)	3.7 ± 1.1 (4)
(+) EDTA	one class	21 ± 1 (3)	1.4 ± 0.1 (3)

Table 2. Kinetic parameters of T3 binding to the recombinant little skate transthyretin.

Each value is the mean \pm standard error of the mean (*n*).

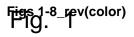
	Methods of preparations		
	HIC column	Co-resin	
Zn	4.47	6.25	
Cu	2.29	0.36	
Ni	<0.06	0.40	
Со	<0.06	(0.48)	
Fe	nd	< 0.02	
Total metal	6.76	7.01	

Table 3. Number of divalent cations in recombinant little skate transthyretin.

TTR was prepared by hydrophobic interaction column (HIC) or Co-affinity resin chromatography and then gel filtration chromatography. Data are shown as metal/TTR (mol/mol). Each value is the mean (n = 3), with less than 1% variation.

*When Co-affinity chromatography was used, it is likely that the TTR samples were contaminated with Co^{2+} released from the Co-affinity resin. Therefore, these values are not included in the total metal contents.

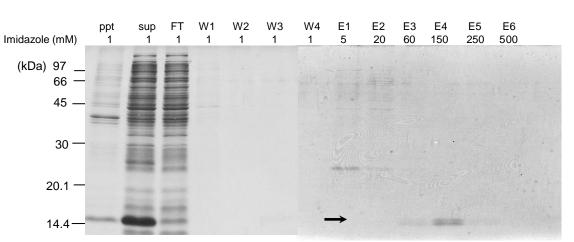
nd, not determined.



	disordered	β-strand A	β-strand B	β-strand C	β-strand D
Lamprey	ADDD <u>HKSHESH</u> EGGVK	DSCPLMVKATDSVOGK	PAAGVKLSVMKKODD	ASWKEVATGVTGKT	GESHHLISDKDFT
Skate		_ FRCPVLIKVLDALKGT			
Trout	PVDRHGESD'	THCPLMVKILDAVKGV	PAGAVALSVSRRVNG	MTWAOVASGVTDLT	GEVHNLISDODFO
Seabream	TPTDKHGGSD	FRCPLMVKILD-VKGT	PAGSVALKVSQKTAL	GGWTQIATGVTDAT	GEIHNLITEQQFP
Toad	APPGHASHGEAD	SKCPLMVKVLDAVRGI	PAANLLVNVFRQTES	GKWEQITSGKTTEL	GEIHNLTTDEQFT
Bullfrog	THGEAD	SKCPLMVKVLDAVRGI	PAAKLPVKVFKQNEI	KSWDLISSGTTSSD	GEIHNLATEEQFV
Crocodile	APLVSHGSID	SKCPLMVKVLDAVRGS	PAANVAIKVFKKTSI	GDWQEFAAGKTTEF	GEVHELTSDEKFV
Chicken	APLVSHGSVD	SKCPLMVKVLDAVRGS	PAANVAVKVFKKAAD	GTWQDFATGKTTEF	GEIHELTTEEQFV
Human	GPTGTGE	sk@plmvkvldavrgs	PAINVAVHVFRKAAD	DTWEPFASGKTSES	GELHGLTTEEFV
	*	** * * *	* *	* * *	** * *
			.	.	
	1	10 20	30	40 50	60
	β -strand E α -hei	lix β-strand	F β-strand	G β -strand H	No. of
	$ \longleftrightarrow $	→ ←→	► ←	\rightarrow	histidine
Lamprey	EGTYKVRFETQQYWT				
Skate	VGLYKFHFETGAYWS	~			
Trout		SQGTTPFHETAEVVFE			
Seabream	AGVYRVEFDTKAYWT	~			
Toad	EGVYKIEFATKAFWG	KLGLSPFHEYVDVVFT	ANDAGHRQYTIAVLI	TPYSFSSTAIVSEP	PHDDL 6
Bullfrog	EGIYKLEFATKRFWS	KLGLTPFHEYVDVVFT	ANDAGHRHYTTAVLI	TPYSFSTTAVVSDV	YKEAHV 6
Crocodile	EGIYRVEFDTSSYWK	ALGLSPFHEYADVVFT	ANDSGHRHYTIAALI	SPFSYSTTAVVSDP	PQE 5
Chicken			ANDSGHRHYTIAALI	ODEOVORDAVACOD	OE 5
CHITCKEH	EGVYRVEFDTSSYWK	JLGLSPIHLIADVVII	ANDOGIIIMITTTAALI	SFESISIIAVVSDE	QE J
Human	EGVYRVEFDTSSYWK EGIYKVEIDTKSYWK				~
				SPYSYSTTAVVTNP	~
	EGIYKVEIDTKSYWK	ALGISPEHEHAEVVFT	ANDSGPRRYTIAALI	SPYSYSTTAVVTNP	PKE 4
	EGIYKVEIDTKSYWK	ALGISPEHEHAEVVFT	ANDSGPRRYTIAALI	SPYSYSTTAVVTNP	PKE 4

Fig. 2

А



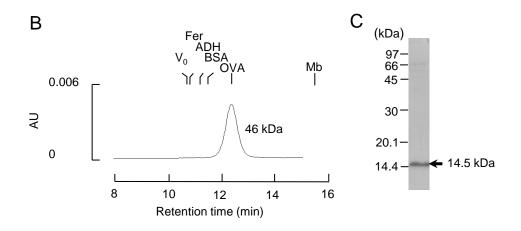
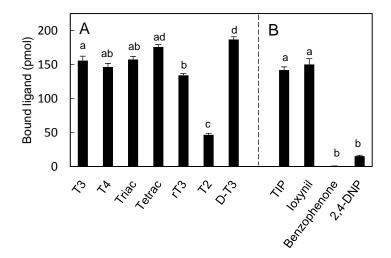
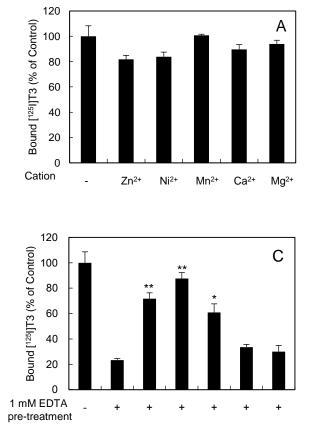


Fig. 3



Cation



Zn²⁺

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Ni²⁺

Mn²⁺

Ca²⁺

Mg²⁺

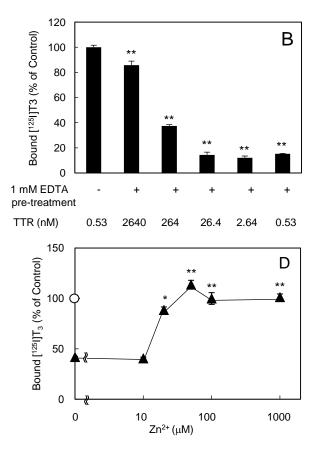
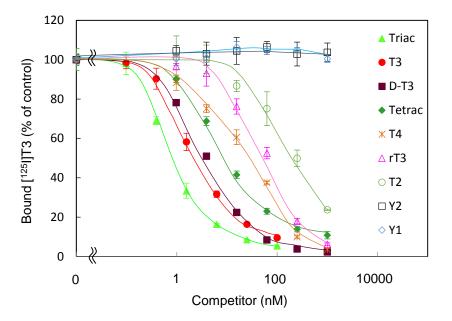
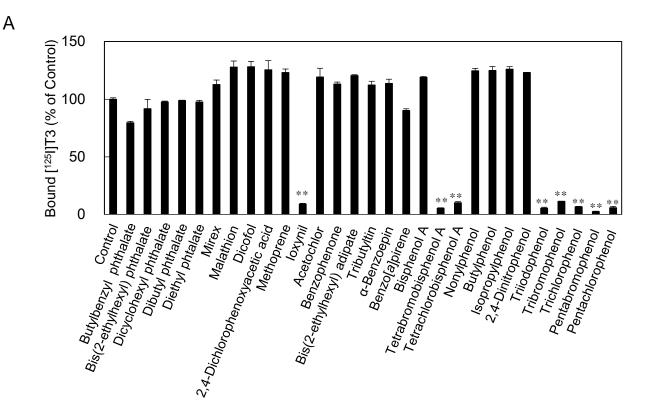
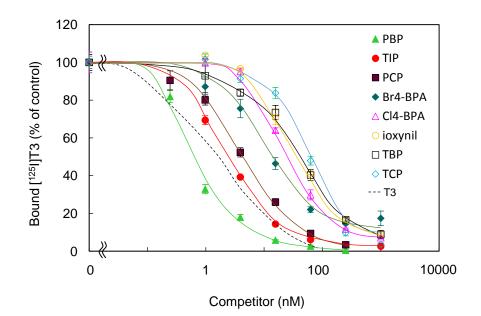


Fig. 5









В

Fig. 7

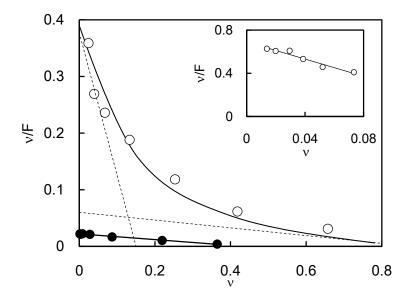


Fig. 8

