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Characterization of a single prolactin (PRL) receptor in tilapia (*Oreochromis niloticus*) which binds both PRL_I and PRL_{II}

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ABSTRACT

In tilapia, there are two forms of prolactin (PRL) whose effects on sodium and chloride movements differ and depend on the living environment of the fish. To see whether different receptors or the same receptor mediates these different effects, we have characterized the specific binding of both forms of tilapia (ti)PRL in two osmoregulatory organs, the gill and kidney. Two recombinant tiPRLs were used for this analysis. The recombinant hormones had the same properties as the native hormones in a tilapia gill radioreceptor assay. Specific binding to gill and kidney membranes was increased by optimizing the quality of the tissue preparations (physiological state of fish, membrane preparation)

and the incubation conditions (pH, salt concentrations, temperature, time). Under these optimized conditions, we detected only one class of high affinity PRL receptor in gill and kidney. Its binding affinity was higher for tiPRL_I than for tiPRL_{II} in both gill and kidney (for tiPRL_I the respective affinity values were 2.9 and 2.3×10^{10} per M, for tiPRL_{II} they were 1.9 and 0.5×10^{10} per M). In competition studies, tiPRL_I was more potent, followed by tiPRL_{II} and ovine (o)PRL. tiGH and oGH did not significantly displace either tiPRL. The receptor we have characterized thus recognizes quite specifically both tiPRLs.

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INTRODUCTION

Although in fish, prolactin (PRL) is pleiotropic and exhibits a spectrum of biological activities, its primary and most studied role in these animals is water and electrolyte homeostasis (for reviews see Clarke & Bern, 1980; Hirano *et al.* 1986). In several fish species, nucleotide or polypeptide sequence analyses have revealed the existence of two distinct, albeit similar, genes. In tilapia, however, two quite different PRL molecules are synthesized and released from the pituitary: their sequence identity is only 69%, one variant (tilapia (ti)PRL_I, 188 amino acids) being more similar to other fish PRLs than to the other variant (tiPRL_{II}, 177 amino acids) (Yamaguchi *et al.* 1988; Rentier-Delrue *et al.* 1989).

Recent studies of the osmoregulatory roles of tiPRL during adaptation to a hyperosmotic environment have revealed that the two tiPRL forms perform different biological functions; only tiPRL_I exerts a dose-related sodium-retaining effect in

tilapia adapted to brackish water (Auperin *et al.* 1994a), whereas both tiPRL forms restore plasma Na⁺ levels to normal in hypophysectomized fish kept in fresh water (Specker *et al.* 1985). In addition, tiPRL_{II} is significantly less potent than tiPRL_I in restoring plasma Cl⁻ levels in such fish (B. Auperin, F. Rentier-Delrue, J. A. Martial & P. Prunet, unpublished observations). These results lead to the interesting question of whether these different functions might be associated with different tiPRL receptor forms.

Attempts have been made to characterize tiPRL receptors in various tilapia organs using ovine (o)PRL (Edery *et al.* 1984; Dauder *et al.* 1990). The specific binding measured in osmoregulatory organs (gill and kidney) was low, however, and ¹²⁵I-labelled oPRL could be significantly displaced from liver membranes by fish or mammalian growth hormone (GH) (Dauder *et al.* 1990; Prunet & Auperin, 1994). As suggested by Dauder *et al.* (1990), it is possible that 'both tiPRL and tiGH

binding sites recognize mammalian PRL'. Since the interaction between mammalian PRL and fish receptors may produce artifacts, full characterization of PRL receptors in tilapia species requires the development of homologous radioreceptor assays using tiPRL_I and tiPRL_{II} as ligands.

We have developed such assays, thanks to the recent production of two recombinant tiPRL forms exhibiting full biological activity (Swennen *et al.* 1991; Auperin *et al.* 1994a). The aim of the present study was to fully characterize the tiPRL receptors. We have focused on the different osmoregulatory functions of the two tiPRL forms and have therefore characterized the PRL receptors mainly in two major osmoregulatory organs, the gill and kidney. We have revealed only one class of tiPRL receptor in these organs, to which tiPRL_I binds with a higher affinity than tiPRL_{II}.

MATERIALS AND METHODS

Hormones

Recombinant tiPRL hormones were produced according to Rentier-Delrue *et al.* (1989). Recombinant tiGH was a gift from J. Smal (Eurogentec, Seraing, Belgium). Native tiPRLs (tiPRL_I and tiPRL_{II}) were generously provided by Prof. H. A. Bern (Department of Integrative Biology, University of California, Berkeley, CA, U.S.A.). oGH (NIDDK-oGH-15) and oPRL (NIDDK-oPRL-19) were gifts from the NIDDK and the National Hormone and Pituitary Program (Bethesda, MD, U.S.A.).

Animals

Tilapia specimens (*Oreochromis niloticus*) were obtained from IDESSA (Bouaké, Ivory Coast), weighing less than 100 mg. They were maintained at the Rennes laboratory in fresh water until reaching the desired weight (30–100 g).

Previous studies have shown that transfer to brackish water induces a significant increase in the number of free PRL receptors within 24 h (Auperin *et al.* 1994b). Thus, all tissues were collected from fish transferred to brackish water (21‰) and kept there for 36–48 h.

Membrane preparations

Gills or kidneys were frozen in liquid nitrogen and stored at -70°C before use. Frozen gills were homogenized in preparation buffer (2.5 mM Tris-HCl buffer, pH 7.5, containing 1.5 mM MgCl₂,

1.5 mM CaCl₂, 0.25 M sucrose) plus 1 mM phenylmethylsulphonyl fluoride (PMSF) (7.5 ml/g frozen tissue) with an Ultraturrax homogenizer at maximal speed for 15-s periods separated by intervals of a few minutes for cooling. The homogenate was layered over 10 ml sucrose (1 M) and centrifuged for 10 min at 400 g at 4°C . The supernatant, without the interface, was layered over 5 ml sucrose (1.8 M) and centrifuged for 1 h 45 min at 120 000 g and 4°C in a zonal rotor. The interface between the 1.8 M sucrose and the supernatant was collected, diluted with gill assay buffer (2.5 mM Tris-HCl buffer, pH 7.5, containing 1.5 mM MgCl₂ and 1.5 mM CaCl₂) and centrifuged at 120 000 g for 20 min at 4°C in a zonal rotor. The pellet was resuspended with a glass homogenizer (Dounce 250 μm) in gill assay buffer with trypsin inhibitor (0.05%, w/v) at a final concentration of 2 g initial tissue/ml. The membrane preparations were stored at -30°C until assayed.

Frozen kidneys were homogenized in kidney assay buffer (10 mM Tris-HCl, 1.5 mM MgCl₂) plus 1 mM PMSF. The amount of buffer added for this step was 5 ml/g frozen tissue. Homogenization was performed with an Ultraturrax homogenizer at maximal speed for two 10-s periods, with an interval of a few minutes for cooling. The homogenate was sieved through cheesecloth and centrifuged for 30 min at 3200 g and 4°C . The pellet was resuspended with a glass homogenizer in assay buffer and recentrifuged under the same conditions. The pellet was then resuspended in assay buffer with 0.05% trypsin inhibitor at 2 g initial tissue/ml. The membrane preparations were stored at -30°C until assayed.

Skin was prepared like gills; liver was prepared like kidney.

For the gill membrane preparations, the membrane protein concentration was determined by the method of Lowry *et al.* (1951) and was adjusted to approximately 6 mg/ml (initial tissue concentration 2 g/ml).

Iodination of recombinant tiPRLs

Na¹²⁵I (specific activity 0.1 mCi/ μl) was purchased from Amersham (Les Ullis, France). The two tiPRLs were iodinated by the chloramine-T method (Greenwood *et al.* 1963; with modifications introduced by Martal, 1972). Radioiodinated tiPRL was separated from free ¹²⁵I⁻ using a 30 cm gel filtration column (Fractogel TSK HW-55; Merck) previously equilibrated with 20 mM phosphate buffer, 0.15 M NaCl, 1% BSA, pH 7.5. The specific activity of ¹²⁵I-labelled tiPRL was 40–50 $\mu\text{Ci}/\mu\text{g}$ PRL for both PRLs.

Receptor binding assays

A membrane preparation (100 μ l) diluted in assay buffer plus one isoform of labelled tiPRL (8000–10 000 c.p.m.) in 50 μ l assay buffer were incubated in duplicate with (non-specific binding) or without (total binding) 1 μ g tiPRL (final volume 250 μ l). Assay tubes were rocked for 20–22 h at 12 °C. The assays were stopped by adding 2 ml ice-cold assay buffer. The tubes were centrifuged for 30 min at 3200 *g* and the contents decanted. The bound radioactivity in the pellet was measured with a Packard Multiprias 2 counter. Specific binding (B) was calculated as the difference between total and non-specific binding and results are expressed as percentages of the total radioactivity (T) originally added to each tube ($B/T \times 100$).

Calculations

The quality of the labelled hormone was analysed in membrane saturation experiments. In these experiments, 60% of the labelled tiPRL_I and 30% of the labelled tiPRL_{II} were found to bind to the membrane. The calculated specific activities and amounts of hormone added to each tube were therefore corrected by these factors as described by Garnier (1980) for Scatchard analysis (Scatchard, 1949).

Saturation assay data are presented as Scatchard plots and analysed with the Ligand program in order to estimate the affinity constants (K_a values) and binding capacities (B_{max} values).

Statistical methods

Results are presented as means \pm s.e.m. Differences between the K_a and B_{max} values calculated for the tiPRL_I and tiPRL_{II} tracers were evaluated by Mann–Whitney analysis. A probability level of <0.05 was considered significant.

RESULTS

Binding to different tissue preparations

Gill and kidney exhibited considerable specific binding which was higher for ¹²⁵I-labelled tiPRL_I than for ¹²⁵I-labelled tiPRL_{II}. The skin and liver preparations exhibited low specific binding, the levels being the same for both labelled hormones (Fig. 1).

In gill and kidney membranes, tiPRL_I and tiPRL_{II} binding was fully characterized. The preparation and incubation conditions were optimized to ensure high specific tiPRL_I and tiPRL_{II} binding.

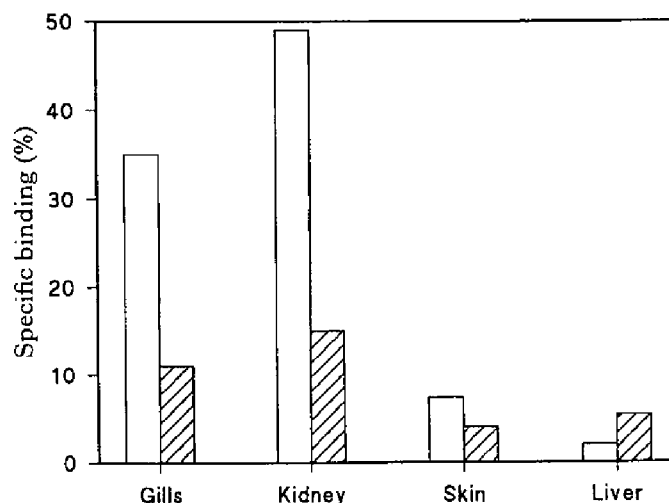


FIGURE 1. Specific binding ($B/T \times 100$, where B =specific binding and T =amount of radioactivity added to the tube) of ¹²⁵I-labelled tiPRL_I (open bars) and ¹²⁵I-labelled tiPRL_{II} (hatched bars) to tilapia organs. Tissue preparations (200 mg wet tissue per tube) were incubated as described in the Materials and Methods. Histograms are the means of duplicate determinations.

Membrane preparation and incubation conditions

For both kidney and gill membranes, specific binding was pH-dependent, binding being highest in the pH range from 6.5 to 8 (data not shown). Studies were performed at pH 7.5, a value yielding the highest ratio of specific to non-specific binding.

The effect of changing the MgCl₂ and CaCl₂ concentrations was also investigated for gill membranes. For this organ, low concentrations of MgCl₂ (1.5 mM) and CaCl₂ (1.5 mM) were required (data not shown). For kidney preparations, only effects of MgCl₂ were investigated, and in this study the highest binding was obtained at a low MgCl₂ concentration (1.5 mM) (data not shown). For both kidney and gill membranes, NaCl (150 mM) in the incubation medium caused a 50% decrease in specific tiPRL_I and tiPRL_{II} binding. Thus no NaCl was added to the incubation medium.

Saturation experiments

Specific binding of the tiPRLs to gill membranes increased linearly with the membrane concentration in the ranges 10–150 μ g protein/tube for tiPRL_I and 10–250 μ g protein/tube for tiPRL_{II}. We were thus able to increase specific binding to 40–45% and 25–30% respectively (Fig. 2a). Likewise, in kidney preparations specific binding of both PRLs increased linearly as the membrane concentration

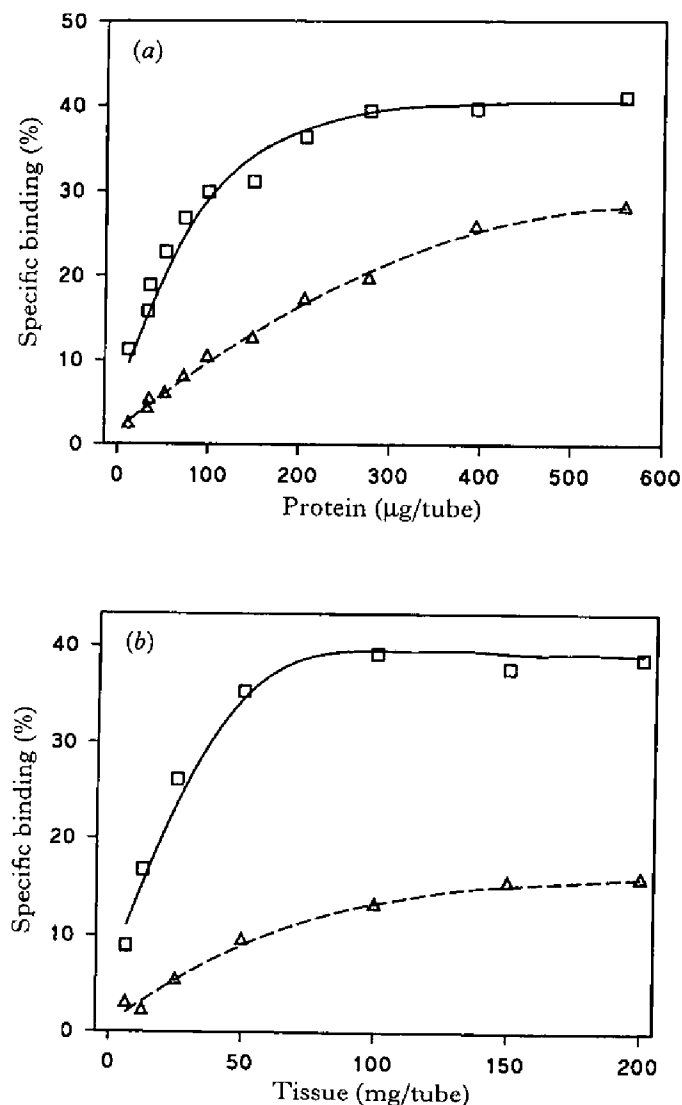


FIGURE 2. (a) Specific binding ($B/T \times 100$, where B =specific binding and T =amount of radioactivity added to the tube) of ^{125}I -labelled tiPRL_I (\square) and ^{125}I -labelled tiPRL_{II} (\triangle) to gill membranes as a function of protein concentration ($\mu\text{g}/\text{tube}$). (b) Specific binding ($B/T \times 100$) of ^{125}I -labelled tiPRL_I (\square) and ^{125}I -labelled tiPRL_{II} (\triangle) to kidney membrane preparations as a function of kidney tissue concentration (mg/tube). Tissue preparations were incubated as described in the Materials and Methods. Values are means of duplicate determinations.

increased from 10 to 50 mg tissue/tube, reaching 40–45% for tiPRL_I and 10–15% for tiPRL_{II} (Fig. 2b).

Association studies

Figure 3a shows the effects of incubation time and temperature on specific binding to gill membranes of tiPRL_I . With tiPRL_I , apparent equilibrium was

reached after 40 h at 4 °C, 20 h at 12 °C, or 8 h at 25 °C. Thereafter, the specific binding remained stable for over 40 h at 4 °C and 4 h at 12 °C, but decreased rapidly at 25 °C. A similar picture was observed with tiPRL_{II} , although at 4 °C the specific binding never quite reached the levels obtained at 12 and 25 °C (data not shown).

Figure 3b shows the effects of incubation time and temperature on the specific binding of tiPRL_I to kidney membranes. Equilibrium was reached after 20, 20 and 50 h at 18, 12 and 4 °C respectively. Thereafter, specific binding remained stable until the end of the experiment (100 h) except at 18 °C, at which temperature it decreased slowly after 50 h.

Specificity experiments

The hormonal specificity of tiPRL_I and tiPRL_{II} binding was further analysed. With gill membrane preparations, binding of ^{125}I -labelled tiPRL_I or ^{125}I -labelled tiPRL_{II} was inhibited when the homologous unlabelled tiPRL was added. This effect increased with the concentration of unlabelled hormone (Fig. 4). Interestingly, native tiPRL_I and tiPRL_{II} purified from pituitary tissues exhibited the same potency as recombinant tiPRL_I and tiPRL_{II} in inhibiting the binding of ^{125}I -labelled tiPRL_I to gill membrane preparations (Fig. 5). Other hormones were also tested. The results of these experiments, shown in Fig. 4, clearly show that: (1) tiPRL_I was the best competitor for both ^{125}I -labelled tiPRL_I (Fig. 4a) and ^{125}I -labelled tiPRL_{II} (Fig. 4b), unlabelled tiPRL_{II} being 8–10 times less potent (calculated at 50% displacement); (2) oPRL could also compete for binding, but it was 30 times less potent than tiPRL_I and 5 times less potent than tiPRL_{II} (calculated at 50% displacement); (3) tiGH and oGH did not significantly displace ^{125}I -labelled tiPRL_I or ^{125}I -labelled tiPRL_{II} from their binding sites on gill membranes.

A corresponding study conducted with kidney membranes yielded results similar to the above.

Scatchard analysis

The characteristics of tiPRL_I and tiPRL_{II} binding to gill and kidney membrane preparations were determined by Scatchard analysis of the saturation curves. As shown in Figs 6 and 7, binding of either tiPRL was a saturable process. The Scatchard plots reveal a single class of high affinity binding site in both gill and kidney tissues. This single class of receptors, moreover, exhibited a higher affinity for tiPRL_I than for tiPRL_{II} in gill. This difference was clearer when kidney membranes were used

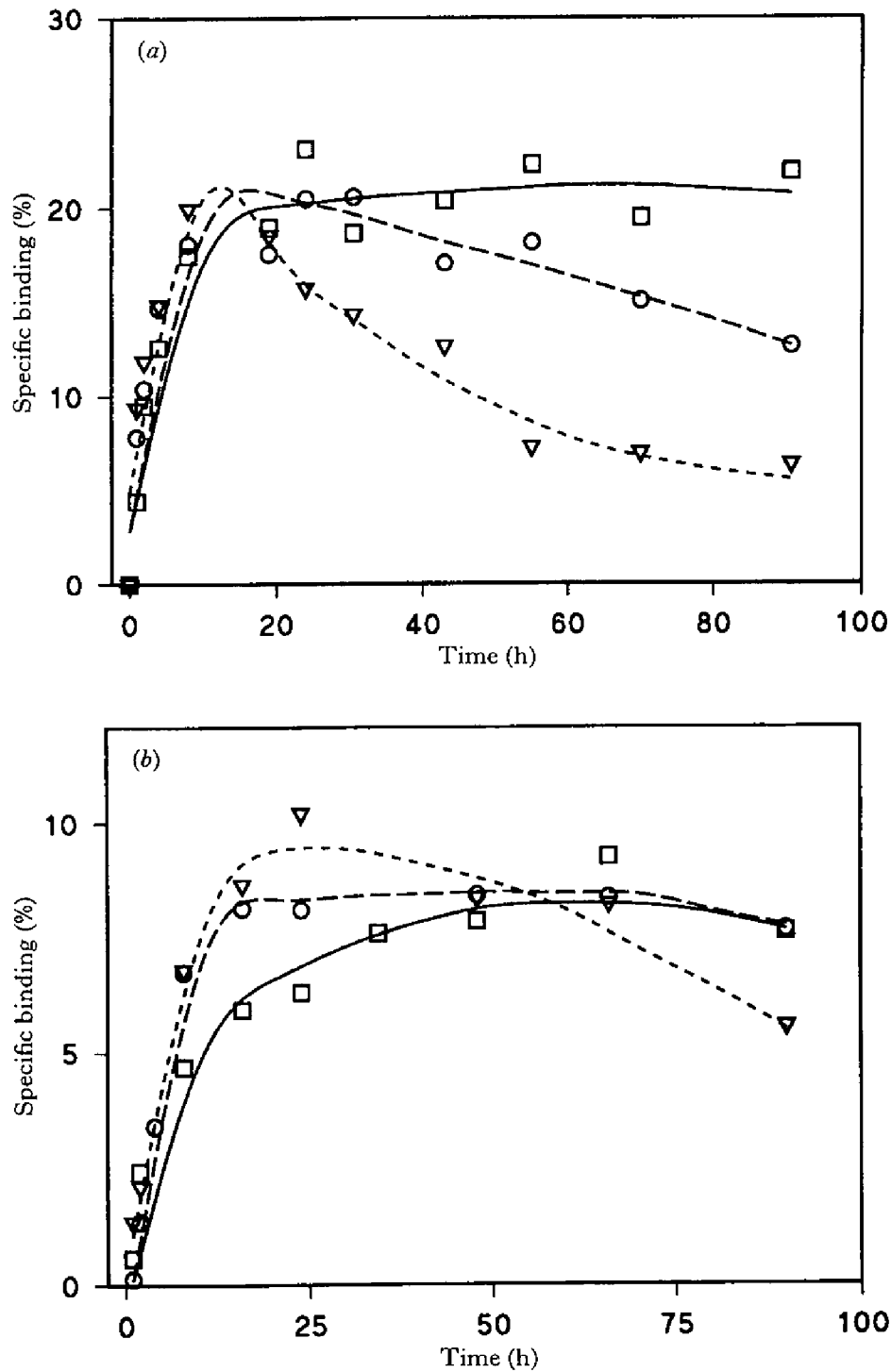


FIGURE 3. (a) Effects of time and temperature (\square , 4 °C; \circ , 12 °C; ∇ , 25 °C) on specific binding ($B/T \times 100$, where B =specific binding and T =amount of radioactivity added to the tube) of ^{125}I -labelled tiPRL_I to gill membrane preparations (40 µg protein/tube). (b) Effects of time and temperature (\square , 4 °C; \circ , 12 °C; ∇ , 18 °C) on specific binding ($B/T \times 100$) of ^{125}I -labelled tiPRL_I to kidney membrane preparations (15 mg wet tissue/tube). Values are means of duplicate determinations.

(Table 1). In both tissues, the binding capacity for tiPRL_I was about twice that obtained for tiPRL_{II}, although these differences are not significant due to

the variability of the measurements (Table 1). Moreover, gill and kidney binding capacities were of similar magnitudes when the B_{max} values were

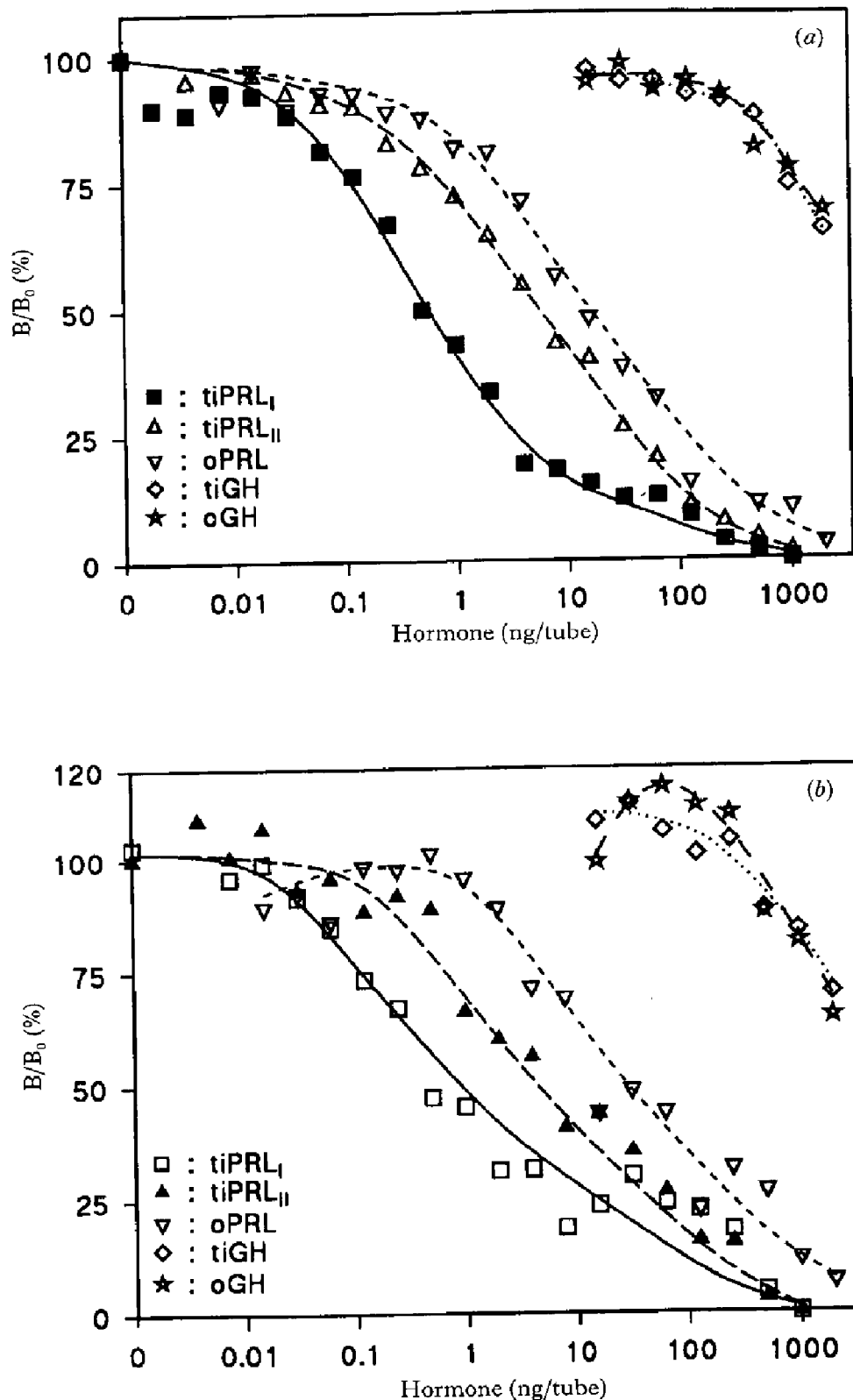


FIGURE 4. Competition curves for (a) ^{125}I -labelled tiPRL_I and (b) ^{125}I -labelled tiPRL_{II} binding to gill membrane preparations (80 μg protein/tube), obtained by increasing the concentration (expressed in ng/tube) of various hormone preparations. Binding is expressed as a ratio of the ^{125}I -labelled tiPRL specific binding in the presence of competition (B) to the ^{125}I -labelled tiPRL specific binding in absence of competition (B_0), multiplied by 100 to obtain a percentage. Tissue preparations were incubated as described in the Materials and Methods. Values are means of duplicate determinations.

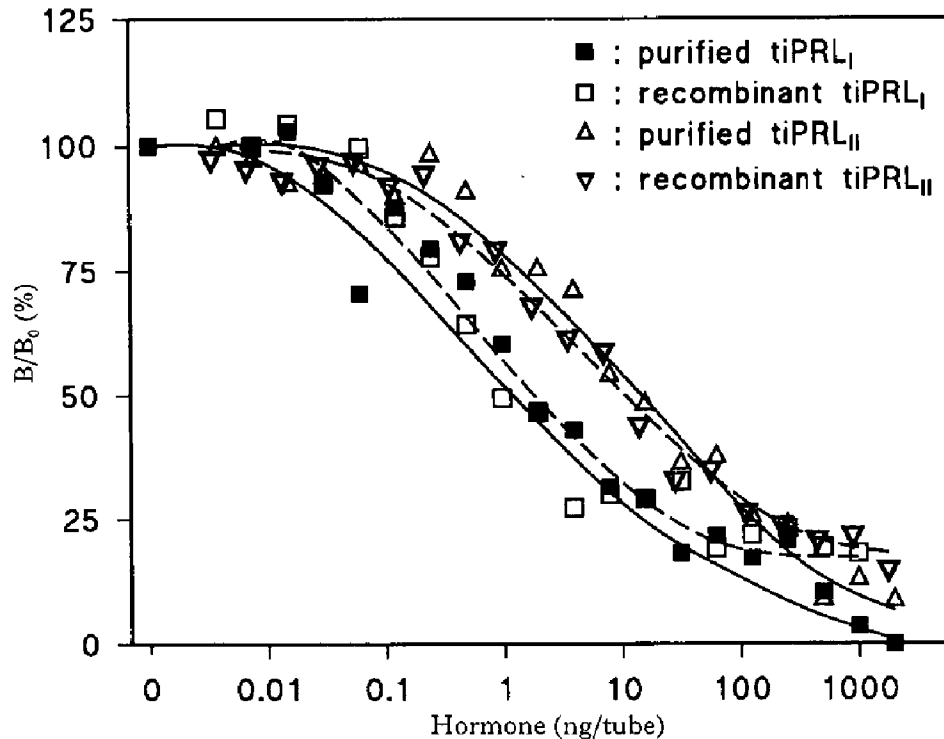


FIGURE 5. Competition curves for ^{125}I -labelled tiPRL_I binding to gill membrane preparations (130 μg protein/tube), obtained by adding increasing concentrations (expressed in ng/tube) of various hormone preparations. Binding is expressed as the ratio of ^{125}I -labelled tiPRL_I specific binding in the presence of competition (B) to ^{125}I -labelled tiPRL_I specific binding in the absence of competition (B_0), multiplied by 100 to obtain a percentage. Tissue preparations were incubated as described in the Materials and Methods. Values are means of duplicate determinations.

expressed per unit weight of initial fresh tissue (data not shown).

DISCUSSION

The present study is the first to present a complete characterization of tiPRL receptors in lower vertebrates, based on the use of homologous recombinant tiPRLs. Our results clearly establish the presence of one class of high affinity tiPRL receptors in the gill and kidney of *Oreochromis niloticus*. tiPRL_I binds to these receptors with a higher affinity than tiPRL_{II}.

The two recombinant tiPRLs previously prepared and characterized by Swennen *et al.* (1991) compete with ^{125}I -labelled tiPRL_I as effectively as do native tiPRLs for binding to gill receptors. This confirms the finding of Swennen *et al.* (1991) that the biological activities of the recombinant tiPRLs are intact, thus validating the use of these recombinant hormones to characterize tiPRL receptors in *Oreochromis niloticus* gill and kidney.

A preliminary analysis of specific binding in various organs indicated that gill and kidney seem to be the tissues showing the highest tiPRL receptor contents. This suggests that both tissues are physiological targets for the direct action of PRL, which is in agreement with the much studied osmoregulatory role of PRL in fish (Clarke & Bern, 1980). Surprisingly, the liver showed low specific binding for either tiPRL, although we tried fresh and frozen tissues and different protocols for membrane preparation (B. Auperin & P. Prunet, unpublished results).

Use of recombinant tiPRL preparations allowed us to obtain high levels for specific binding of tiPRL_I and tiPRL_{II} to both kidney and gill membrane preparations. It also appears that specific binding is increased by the use of low Tris-HCl and divalent cation concentrations, an observation already reported by Yao *et al.* (1991) for trout GH receptors. Attempts to desaturate the gill and kidney membrane preparations by MgCl_2 treatment (Kelly *et al.* 1979) failed to improve the specific binding of either tiPRL (data not shown). However, collecting tissues from fish transferred for 36–48 h

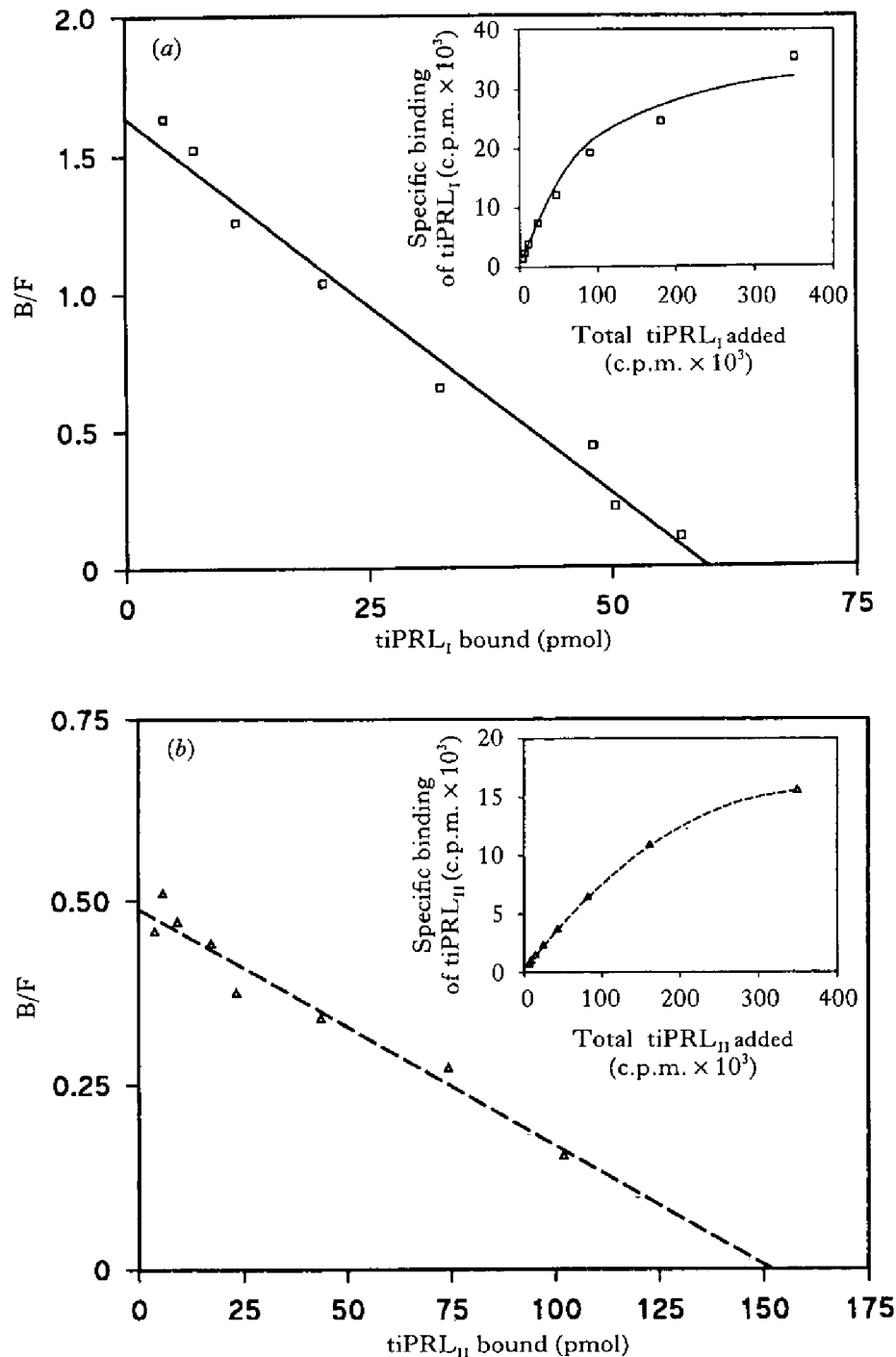


FIGURE 6. Scatchard plots of specific (a) ¹²⁵I-labelled tiPRL_I and (b) ¹²⁵I-labelled tiPRL_{II} binding to gill membrane preparations. The insets show the effect of increasing concentrations of ¹²⁵I-labelled tiPRL_I or ¹²⁵I-labelled tiPRL_{II} on specific binding, and were used for the Scatchard analysis. B/F, ratio of bound to free hormone. Tissue preparations were incubated as described in the Materials and Methods. Values are means of duplicate determinations.

to brackish water increased the specific binding of both PRLs to gill and kidney preparations. This was also shown by Auperin *et al.* (1994b) who

demonstrated an increase in the number and affinity of gill PRL receptors after transfer from fresh to brackish water.

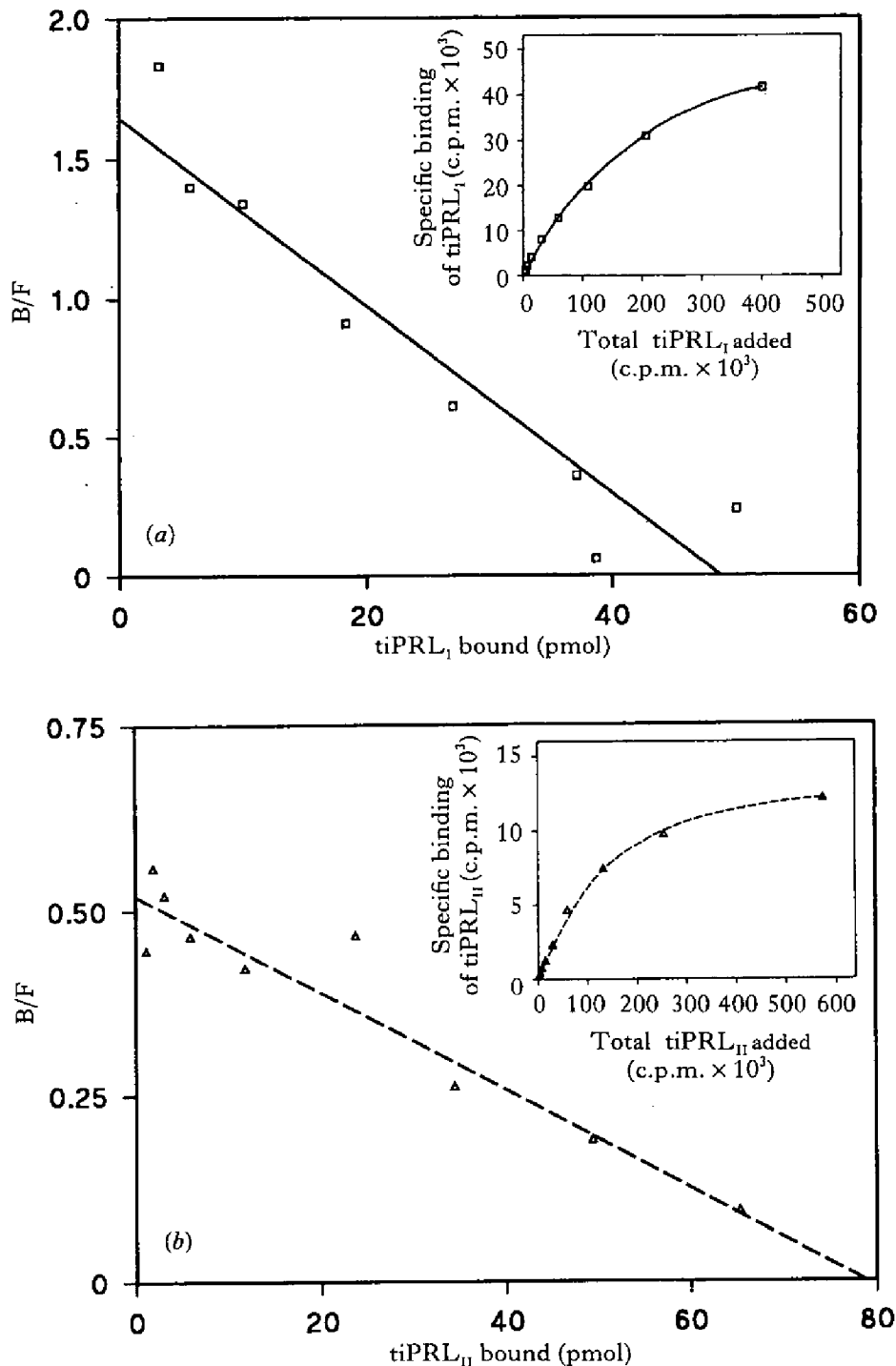


FIGURE 7. Scatchard plots of specific (a) ^{125}I -labelled tiPRL_I and (b) ^{125}I -labelled tiPRL_{II} binding to kidney membrane preparations. The insets show the effect of increasing concentrations of ^{125}I -labelled tiPRL_I or ^{125}I -labelled tiPRL_{II} on specific binding, and were used for the Scatchard analysis. B/F, ratio of bound to free hormone. Tissue preparations were incubated as described in the Materials and Methods. Values are means of duplicate determinations.

The specific binding characteristics (saturability and high affinity) of the two tiPRLs fit the criteria for receptors. Scatchard analysis revealed a single class

of tiPRL receptor, whichever tiPRL was used as a ligand. The affinity constants were always significantly higher for tiPRL_I than for tiPRL_{II} in both gill

TABLE 1. K_a and B_{max} values for ^{125}I -labelled tiPRL_I and ^{125}I -labelled tiPRL_{II} in gill and kidney membrane preparations. Tissue preparations were incubated as described in the Materials and Methods. Values are the means \pm S.E.M. of three separate determinations

Tissue	Ligand	K_a ($\times 10^{10}$) (per M)	B_{max} (fmol/mg)
Gills ($n=3$)	tiPRL _I	2.9 ± 0.2	61.7 ± 4.4
	tiPRL _{II}	$1.9 \pm 0.6^*$	34.8 ± 12.9
Kidney ($n=3$)	tiPRL _I	2.3 ± 1.1	24.5 ± 11.7
	tiPRL _{II}	$0.5 \pm 0.16^*$	13.5 ± 1.0

* $P < 0.05$ compared with K_a for tiPRL_I in the same tissue (Mann-Whitney test).

and kidney tissues. This result is confirmed by the specificity curves, in which tiPRL_I always appeared 8–10 times more potent than tiPRL_{II} in competing for both gill and kidney binding sites, whichever tiPRL was used as a ligand. Moreover, both tiPRLs totally displaced both tiPRL ligands from their binding sites, a result which further supports our hypothesis of a unique PRL receptor in these organs. However, we cannot be certain that our experimental conditions are adequate for characterizing another tiPRL receptor in gill and kidney, and that Scatchard analysis would necessarily reveal a second binding site. This would be the case particularly if the second receptor was in low concentration with a binding affinity close to the values reported for the previous PRL receptor. The initial cloning of the cDNA for the rat PRL receptor in the liver led to the identification of multiple forms of PRL receptor in mammals and human (Kelly *et al.* 1991). A similar approach should be developed to determine whether one or several PRL receptor forms are present in tilapia gill and kidney.

Interestingly, recombinant tiGH and oGH did not significantly displace either the ^{125}I -labelled tiPRL_I or the ^{125}I -labelled tiPRL_{II} ligand from gill or kidney preparations. This indicates that, although PRLs and GHs have structural and evolutionary similarities, PRL receptors in these fish tissues do not bind homologous or mammalian GH. The results of previous studies in which oPRL was used as a ligand (Edery *et al.* 1984; Dauder *et al.* 1990) have already suggested the presence of PRL-binding sites in the gill and kidney of *Oreochromis mossambicus*. Yet with this heterologous system (^{125}I -labelled oPRL displaced by tiPRL_I or tiPRL_{II}) it was not possible to clearly characterize the tiPRL receptors, as specific ligand binding was always low (2–7%). Furthermore, tiGH displaces ^{125}I -labelled oPRL bound to liver membranes with potencies of 0.8 and 5.5 compared with tiPRL_I and tiPRL_{II} respectively (Dauder *et al.* 1990). It thus appears that oPRL can also recognize tiGH receptors, as sug-

gested by Nicoll (1982) and Dauder *et al.* (1990). This makes it hard to compare these studies (Edery *et al.* 1984; Dauder *et al.* 1990) with our results.

The characteristics of the tiPRL receptor we have uncovered are very similar to those described for PRL receptors in mammalian liver and mammary gland, especially as regards their specificity (Shiu & Friesen, 1974; Djiane *et al.* 1977; Sakai *et al.* 1985). Measurements of the apparent K_a values for tiPRL_I in both tissues ($2\text{--}3 \times 10^{10}$ per M) indicate a receptor of very high affinity. These values are about 10-fold higher than those currently reported for liver and mammary gland PRL receptors in mammals. However, similar high values for K_a have recently been reported for the PRL receptor in the adrenal cortex of the pig (Klemcke *et al.* 1989). It should be also noted that the higher K_a values obtained using tiPRL_I are quite compatible with the low concentrations of plasma PRL_I reported in fresh water-adapted tilapia (Auperin *et al.* 1994a).

It was also interesting to compare the characteristics of tiPRL receptors in gill and kidney. As regards their specificity, the receptors present in the two tissues exhibit similar ranges of activity, tiPRL_I being the most potent ligand, followed by tiPRL_{II}, oPRL, and, as weak competitors, tiGH and oGH. The K_a values obtained for tiPRL_I binding were similar in the two tissues; so were the binding capacities expressed per unit weight of initial tissue. The K_a values calculated for tiPRL_{II} binding differed significantly, however, between the two tissues. This warrants a more complete characterization of the tiPRL receptor using molecular biological techniques in order to determine whether the receptors are identical in gill and kidney.

The presence of a single tiPRL receptor class in gill and kidney, with a higher affinity for tiPRL_I than for tiPRL_{II}, does not explain, or at least not simply, all the results obtained *in vivo* in tiPRL-treated fish. It can explain why tiPRL_I is more potent than tiPRL_{II} in its ability to maintain plasma Na^+ and Cl^- levels in fish adapted to brackish

water, but it does not explain why this effect is dose-related for tiPRL_I, but not for tiPRL_{II} (Auperin *et al.* 1994a). Indeed, the binding affinity of oPRL for the tiPRL receptor characterized here is lower than that of either tiPRL, yet when oPRL is injected into fish adapted to brackish water, its Na⁺- and Cl⁻-retaining effects are dose-related (Auperin *et al.* 1994a); thus, the absence of dose-related effects after tiPRL_{II} treatment cannot be explained by a lower affinity of tiPRL_{II} for kidney or gill PRL receptors. Moreover, tiPRL_I and tiPRL_{II} show similar abilities to maintain plasma Na⁺ levels in hypophysectomized fish reared in fresh water, but they affect Cl⁻ levels differently (B. Auperin, F. Rentier-Delrue, J. A. Martial & P. Prunet, unpublished observations). These results clearly show that the two tiPRL forms have different osmoregulatory effects. At the present time, it is difficult to speculate upon the mechanisms that mediate these different actions, a question which clearly deserves further study.

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