The contribution of marine biology to biomedical research: past, present, future

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"Ask the fish of the sea, and they will declare unto thee" (Job 12:8)

INTRODUCTION

This contribution focusses on the role marine biology – or more precisely marine species or marine model systems - have played in the elucidation of basic physiological processes. Many of these have turned out to be of utmost significance in understanding the function of the human body in health and disease or in developing therapeutical means to cure human maladies. The particular usefulness of marine models in a variety of these seminal discoveries can be traced back to three main reasons: (a) Abundance of species, size of biological model system, and ready accessibility lead to an ease of experimentation. (b) In several marine species, that represent early steps in mammalian evolution, organ structure is often quite simple and organ function is highly specialized. Such "unifunctionality" contrasts to the "multifunctionality" usually found in mammalian and human tissues and facilitates the investigation of a particular function in a defined, homogeneous cell population. (c) Most importantly, it has become clear, mainly through the recent advances in molecular biology and cell biology, that biodiversity in cellular function does not require an incomprehensible number of functionally different units, it can be reduced to the existence of a limited number of families of closely related molecules that are employed by nature to perform basic cellular functions (Kinne, 1991a). These similarities allow us to draw conclusions from findings on marine organisms as to the function of a human organ with even more certainty.

In the following, mainly contributions of marine models to renal physiology and pathophysiology will be highlighted, but it should be emphasized that the role of marine models was equally important in other areas of cell research. In the area of oogenesis, spermatogenesis, cytokinesis, and reproduction, numerous studies on the eggs of sea urchins, sand dollars, and snails have laid a firm basis for our understanding of cell division and its temporal and spatial organization (Rappaport, 1991). Elasmobranch testes have recently been discovered to provide an ideal model system for the studies of different phases of spermatogenesis and the viviparous dogfish is a suitable model for

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E. Kinne-Saffran & R. K. H. Kinne

investigations of the hormonal regulation of reproduction (Callard, 1991; Koob & Callard, 1991). In nerve physiology, the giant axon of the squid was – and continues to be – one of the prime model systems in which the properties of numerous ion channels and ion pumps have been characterized and their role in nerve conduction identified (Boron & Knakal, 1992; Hodgkin & Huxley, 1952). Also the Na-K-ATPase, the primary pump maintaining intracellular ion homeostasis and cell volume, was first described by Skou in the leg nerves of crabs (Skou, 1957), thereby providing the first link of an ATP-consuming cellular reaction to the translocation of inorganic electrolytes across cell membranes (Skou, 1989). This enzyme was later found by Schatzmann (1967) to be inhibited by "cardiac glycosides" which are used for the treatment of heart failure. Until today, the sodium chloride-secreting rectal gland of the shark is one of the richest sources of this enzyme for biochemical and biophysical studies (Medzihradsky et al., 1967). The rectal gland also, very early on, played an important role in elucidating the mechanisms of hormonal regulation of salt transport and intracellular signalling – an area still pursued vigorously using this organ as model system (Schofield et al., 1991).

MARINE BIOLOGY AND RENAL PHYSIOLOGY

In Figure 1, a scheme of a mammalian nephron is represented with its proximal tubule, Henle's loop, the distal tubule and the collecting duct. In all these parts of the kidney, studies on marine organisms have contributed essential information on the function of these segments and the cellular and molecular basis for their function.

For the proximal tubule, in 1923, the question was solved whether mammalian kidneys have the capability to excrete compounds from the blood into the primary urine (Marshall & Vickers, 1923) in addition to the – at that time – widely-accepted functions of filtration in the glomerulum and reabsorption along the tubule (Cushny, 1917). Definite proof of secretory processes in this segment could be obtained by the use of the goosefish



Fig. 1. Schematic representation of a mammalian (human) nephron indicating some of the main segments where marine biology has considerably contributed to elucidating the cellular and molecular mechanism of their function

(Lophius piscatorius), i.e. an aglomerular fish, in which no filtration occurs, and substances transferred from the blood to the urine must have passed the renal cell in a secretory direction (Marshall & Grafflin, 1928). This process was first found for a variety of organic dyes and later led to the development of contrast media to trace the urinary tract, or to measure renal blood flow. This route is still employed for the targeting of antibiotics in renal infection or of diuretics to their intratubular site of action. The cellular mechanism underlying tubular secretion was also first unveiled using fish models, such as the flounder, (Forster & Taggart, 1950; Kinter, 1966). In Figure 2, studies are shown in which



Fig. 2. ³H-chlorphenol red autoradiograph of proximal tubules dissected from the kidney of the winter flounder (*Pseudopleuronectes americanus*) and incubated for 60 min at 20°C in 10 μM tritiated chlorphenol red. Individual tubules are shown both in cross and longitudinal sections in a low-magnification dark-field photomicrograph (autographic silver grains appear as white dots). For further details see Kinter, 1975 (reprinted with kind permission)

proximal tubules dissected from the flounder kidney were incubated in a bath containing a radioactively labelled weak organic acid (Kinter, 1975). The subsequently obtained autoradiograph clearly establishes that during secretion by the cells, weak organic acids are first accumulated intracellularly and then further accumulated within the tubular lumen. The driving force for the latter accumulation could be identified in flounder kidney brush border vesicles – and later also in mammalian kidneys – to be provided by the electrical potential across the brush border membrane (Eveloff et al., 1979). The mechanism of accumulation at the basal pole of the cell is depicted in Figure 3 (Kinne, 1988a; Pritchard, 1990; Shimada et al., 1987). Indirect coupling between a sodium gradient-driven organic acid uptake system (e.g. for glutarate) and an exchange of



Fig. 3. Transport scheme for the secretion of the weak organic acid p-aminohippurate (PAH) across a proximal tubular cell (upper right panel). Uptake from blood into the cell involves indirect coupling to a sodium-A⁻ cotransport system. (A⁻ stands for glutarate or other dicarboxylic acids). Transfer across the luminal membrane involves an electrogenic transporter driven by the electrical potential difference across the cell membrane. The two other schemes show additional kinds of indirect coupling of sodium cotransport transport systems in the renal secretion of sulfate in winter flounder and organic cations (S) in mammals. For further information see Kinne, 1988a (reprinted with kind permission)

intracellularly accumulated glutarate with a weak organic acid, comprises the sequence of events that ultimately lead to the intracellular accumulation of the weak organic acid. Such a mechanism has proved to be operating also for example in the excretion of uric acid in crustaceans (A. Nies et al., unpubl. obs.) as well as in mammals (Maxild et al., 1981).

Figure 4 depicts the mechanisms involved in active chloride transport in the shark rectal gland and in the mammalian thick ascending limb of Henle's loop. This model was first proposed in 1985 (Epstein & Silva, 1985) and experimentally proven in studies on the rectal gland and the flounder intestine (Kinne, 1988b; Kinne, 1991b). The essential elements of this model are the Na-K-2Cl cotransporter which is responsible for the intracellular accumulation of chloride, the chloride channel through which chloride leaves the cell at the opposite side, the potassium channel which allows potassium to leave the cells, and the above described Na-K-ATPase. This enzyme maintains the sodium gradient across the cell membranes and provides the primary driving force for the active transcellular chloride transport. It is interesting to note that one of the first indications of the involvement of the Na-K-2Cl cotransporter in active chloride transport in the kidney, was an observation made by us during a stay at the Biologische Anstalt Helgoland. In perfusion studies using rectal glands of *Scilliorhinus canaliculus* it could be shown that compounds strongly promoting salt secretion (or inhibiting salt reabsorption) in the mammalian kidney also strongly inhibited chloride transport in the rectal gland, as

Marine biology and biomedical research



Fig. 4. Schematic representation of transcellular active chloride secretion in rectal gland cells of elasmobranchs and active chloride reabsorption in the thick ascending limb of Henle's loop in mammalians (modified after Epstein & Silva, 1985, with kind permission)

documented in Figure 5 (Kinne & Kinne-Saffran, 1979). This link led to a working model for active chloride transport in the mammalian thick ascending limb – and the operation of all the above mentioned transport systems could also be demonstrated in the kidney (Greger, 1985). It was further shown that these transport systems have similar properties – as depicted for the Na-K-2Cl cotransporter in Table 1 (Kinne, 1988a) – but that their cellular localization is different, in order to enable the mammalian cells to reabsorb chloride rather than to secrete chloride as the rectal gland cells do. To this end a Na-K-2Cl cotransporter and a potassium channel are known to be transferred into the luminal cell membrane, whereas a chloride channel is in the contraluminal membrane. Such transposition raises interesting questions on the molecular identity of the transport systems and the nature of the signals controlling their intracellular sorting and targeting (Simons & Fuller, 1985) – questions currently being investigated – again using marine model systems.

Another group of diuretics, the thioziazides, are known to inhibit salt transport in the late distal tubule (see Figure 6; [Ellison et al., 1987]), a segment of very high cell heterogeneity in the mammals. Here the discovery of a similar transport system in flounder urinary bladder (Table 2; [Stokes et al., 1984]), a rather simple epithelium, will undoubtedly lead to a detailed characterization of the transport system and to a much better understanding of the mechanism of action of these drugs.

Finally, recent studies on the collecting duct have concentrated on the role of organic

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Rectal gland	$\sim 17 \times 10^4$ $\sim 1000/s$	4.3 mMNa > Li = NH ₄	15 mM K $\ge \text{NH}_4 = \text{K} = \text{Rb} = \text{choline} > \text{Cs}$	overall 75 mM Br = $Cl >> NO_3$	$Br > Cl > NO_3$	~ 0.5 × 10 ⁻⁵ M bumetanide > piretanide > furosemid 43 000-50 000 (affinity chromatography)
Rabbit TALH	$\sim 5 \times 10^4$ ~ 6000/s	1.3 mM Na > Li >> NH4 >> K	0.3 mM K > NH4 > Cs >> Na >> choline	$\sim 1.0 \text{ mM}$ Br = Cl >> NO ₃ = SCN	> 15 mM Br > Cl > NO ₃ \approx SCN	~ 2.5 × 10 ⁻⁶ M burnetanide > piretanide > furosemide 80 000–90 000
	Number per cell Turnover rate Sodium hinding site	Affinity Specificity Potassium binding site	Affinity Specificity Chloride binding site 1	Affinity Specificity Chloride binding site 2	Affinity Specificity Interaction with loop diuretics	Affinity (bumetanide) Specificity Apparent molecular weight (radiation inactivation)

Table 1. Properties of the Na-K-2Cl cotransporter in secretory and absorptive epithelia (Reprinted with permission from Kinne, 1988a, Pergamon Dress 1 rd)

E. Kinne-Saffran & R. K. H. Kinne

Marine biology and biomedical research



Fig. 5. Effects of various "loop diuretics" on the rate of fluid secretion (i.e. sodium chloride) in "in situ perfused" rectal glands of *Scilliorrhinus caniculus* (L.). Loop diuretics are known to inhibit chloride reabsorption in human kidney at the thick ascending limb of Henle's loop. Secretion at various concentrations of the diuretics is given in percent of control. For further details see Kinne & Kinne-Saffran, 1979 (reprinted with kind permission)



Fig. 6. Effect of chlorothiazide (CTZ) on salt reabsorption in distal tubules of rabbit kidney. Note that sodium and chloride transport are inhibited to the same extent, suggesting the presence of an electroneutral NaCl cotransport in this renal segment. For further information see Ellison et al., 1987 (reprinted with kind permission)

E. Kinne-Saffran & R. K. H. Kinne

Table 2. Effect of hydrochlorothiazide on simultaneously determined sodium and chloride tracer fluxes in the urinary bladder of the winter flounder (*Pseudopleuronectes americanus*). Mucosa-to-serosa denotes the fluxes measured when tracer ions were present at the surface representing the lumen of the bladder; serosa-to-mucosa depicts fluxes when tracer ions were present at the outside of the bladder. Net fluxes represent the difference between the fluxes found in the mucosa-to-serosa direction and those observed in the serosa-to-mucosa direction. Note the similarity of sodium and chloride net fluxes and the parallel inhibition by hydrochlorothiazide. Modified after Stokes et al., 1984, with kind permission

		I	Flux			
	Mucosa-to-serosa $(n = 8)$		Serosa-to-m	Net fluxes		
	J _{Na}	J _{Cl}	J _{Na}	J _{Cl}	J _{Na}	$J_{\rm Cl}$
	μM/c	$m^2 \cdot h$	μM/c	$m^2 \cdot h$		
Control	1.70	2.45	0.52	1.34	1.18	1.11
	± 0.25	± 0.30	± 0.07	± 0.12		
HCTZ	0.40	0.69	0.23	0.29	0.17	0.40
(0.1 mM)	± 0.05	± 0.24	± 0.02	± 0.04		
P	< 0.005	< 0.005	< 0.002	< 0.001	-	_
J _{Na} = sodium	flux; $J_{Cl} = chl$	oride flux				

osmolytes in the volume regulation of these cells (Garcia-Perez & Burg, 1991). During the formation of concentrated urine, these cells are exposed to salinities which are similar to or exceed those encountered by marine organisms. These organic osmolytes have for a long time already been identified in marine organisms (see Table 3; [Yancey et al., 1982]) and are also found in the mammalian papilla (Table 4). The regulation of their intracellular concentrations according to the external osmolarity has been extensively studied, for example in skate erythrocytes (Goldstein & Brill, 1991). Again, numerous similarities have become apparent, and it is to be expected that the cross-talk between disciplines – as in the other examples mentioned above – will yield fruitful and important insights into general physiological mechanisms.

CONCLUDING REMARKS

In the future, marine biological models will continue to foster progress in the understanding of molecular, cellular and systemic processes in biomedicine. Undoubtedly, a major impact will be made by studies on the molecular biology of basic physiological and pathophysiological events. At the same time, however, an equally strong emphasis should be placed on integrative aspects in marine biology and biomedicine alike; only the integration of knowledge obtained at various levels of complexity and in various areas of research will lead to a thorough understanding of life on this planet and to the development of strategies to protect it.

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52

Marine biology and biomedical research

Table 3. Organic osmolytes in marine species. Taken from Yancey et al., 1982; copyright 1982 by the AAAS, with kind permission. For references see Yancey et al. 1982

(occurrences)				
A. Polyhydric a	lcohols-polyols			
Cyanobacteria				
Synechococcus sp.	głucosylglycerol			
Fungi				
Saccharomyces rouxii	arabitol			
Asteromyces cruciatus	arabitol, glycerol, mannitol			
Lichens				
Lichina pygmeae	mannosidomannitol			
Unicellular algae				
<i>Dunaliella</i> spp.	glycerol			
Chlorella pyrenoidosa	sucrose			
Ochromonas malhamensis	isofloridoside			
Multicellular algae				
Fucus spp.	mannitol			
Vascular plants				
Gossypium hirsutum L.	glucose, fructose, sucrose			
Insects (freeze-tolerant or -resistant)				
<i>Eurosta solidaginis</i> (Diptera)	glycerol, sorbitol			
Bracon cephi (Hymenoptera)	glycerol			
Crustaceans				
Artemia salina (emerging larvae)	glycerol, trehalose			
Vertebrates				
Hyla versicola	glycerol			
P. Amino peide and p	mine acid derivatives			
B. Amino acius anu a	mino acid derivatives			
Klobsiolla aerogenes	glutamic acid, proline			
Salmonella oranienburg	glutamic acid, proline			
Streptococcus faecalis	y-aminobutyric acid proline			
Protozoa	r anniosatyrie acia, promite			
Miemiensis avidus	glycine alanine proline			
Vacular plants	gryeme, diamie, promie			
Spartina townsendii	betaine			
Atriplex spongiosa	betaine			
Aster tripolium	proline			
Mesembryanthemum nodiflorum	proline			
Invertebrates	promite			
All phyla of marine vertebrates				
Balanus nubilus (barnacla)	various amino acids			
nonanas maninas (narmacie)	various amino acids			
Friocheir sinensis (crah)	Tarious ammo actus			
Eriocheir sinensis (crab)	various amino acide			
Eriocheir sinensis (crab) Parastichopus sp. (echinoderm) Sonia officinalie (mollusk)	various amino acids various amino acids			
Eriocheir sinensis (crab) Parastichopus sp. (echinoderm) Sepia officinalis (mollusk) Cuclostomos	various amino acids various amino acids			
Eriocheir sinensis (crab) Parastichopus sp. (echinoderm) Sepia officinalis (mollusk) Cyclostomes Murino glutinosa (hagfich)	various amino acids various amino acids			
Eriocheir sinensis (crab) Parastichopus sp. (echinoderm) Sepia officinalis (mollusk) Cyclostomes Myxine glutinosa (hagfish) Amphihia	various amino acids various amino acids various amino acids			
Eriocheir sinensis (crab) Parastichopus sp. (echinoderm) Sepia officinalis (mollusk) Cyclostomes Myxine glutinosa (hagfish) Amphibia Buío marinus	various amino acids various amino acids various amino acids various amino acids			
Eriocheir sinensis (crab) Parastichopus sp. (echinoderm) Sepia officinalis (mollusk) Cyclostomes Myxine glutinosa (hagfish) Amphibia Bufo marinus	various amino acids various amino acids various amino acids various amino acids			

Osmolyte system (occurrences)	Principal osmolytes
C. Urea and	methylamines
Cartilaginous fishes (elasmobranchs)	
Vertebrates	
Squalus acanthias (dogfish)	urea, trimethylamine-N-oxide
Dasyatis americana (ray)	urea, trimethylamine-N-oxide
Raja erinacea (ray)	urea, amino acids
D. Urea: est	ivating forms
Mollusks	
Bulimulus dealbatus	
Lungfishes: African and South American	
Amphibians	
Scaphiopus couchi (spadefoot toad)	
E. Inorg	anic ions
Archaebacteria	
Halobacterium spp.	K ⁺

Table 3 (continued)

Table 4. Osmolytes in renal inner medulla of antidiuretic animals. Units are mmol/kg protein or mmol/kg wet weight (in brackets). GPC = glycerophosphorylcholine. Taken from Garcia-Perez & Burg, 1991, with kind permission. For references see Garcia-Perez & Burg, 1991

Species	Urea	Sodium	Sorbitol	Inositol	GPC	Betain
Rabbit			[7]	[11]	[13]	[56]
	[269]		[21]		[21]	[35]
	[346]	[279]	[80]	[16]	[41]	[42]
	1.017	1.890	221	97	195	235
			[21]	[20]		
Rat				• •	[49]	
	[380]	[248]	854	331	2.498	132
	4.405			178	517	214
			145	214	443	
Vole	[349]	[171]	[26]	[12]	[43]	[19]
Deer mouse	[695]	[278]	[19]	[14]	[65]	[21]
Pocket mouse	[1.129]	[350]	[53]	[14]	[71]	[50]
Sheep			[54]			
Dog					[67]	
	[376]	[304]		[14]		
			[2]	5	[18]	

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