Eggshell and egg yolk proteins in fish: hepatic proteins for the next generation: oogenetic, population, and evolutionary implications of endocrine disruption
Augustine Arukwe*¹ and Anders Goksøyr²,³

Address: ¹Great Lakes Institute for Environmental Research, University of Windsor, Ontario, 401 Sunset Avenue, Windsor, N9B 3P4, Canada, ²Biosense Laboratories AS, Thormøhlenstg. 55, N-5008, Bergen, Norway and ³Department of Molecular Biology, University of Bergen, N-5020 Bergen, Norway

Email: Augustine Arukwe* - arukwe@uwindsor.ca; Anders Goksøyr - anders@biosense.no
* Corresponding author

Abstract
The oocyte is the starting point for a new generation. Most of the machinery for DNA and protein synthesis needed for the developing embryo is made autonomously by the fertilized oocyte. However, in fish and in many other oviparous vertebrates, the major constituents of the egg, i.e. yolk and eggshell proteins, are synthesized in the liver and transported to the oocyte for uptake. Vitellogenesis, the process of yolk protein (vitellogenin) synthesis, transport, and uptake into the oocyte, and zonagenesis, the synthesis of eggshell zona radiata proteins, their transport and deposition by the maturing oocyte, are important aspects of oogenesis. The many molecular events involved in these processes require tight, coordinated regulation that is under strict endocrine control, with the female sex steroid hormone estradiol-17β in a central role. The ability of many synthetic chemical compounds to mimic this estrogen can lead to unscheduled hepatic synthesis of vitellogenin and zona radiata proteins, with potentially detrimental effects to the adult, the egg, the developing embryo and, hence, to the recruitment to the fish population. This has led to the development of specific and sensitive assays for these proteins in fish, and the application of vitellogenin and zona radiata proteins as informative biomarkers for endocrine disrupting effects of chemicals and effluents using fish as test organisms. The genes encoding these important reproductive proteins are conserved in the animal kingdom and are products of several hundred million years of evolution.

Introduction
Teleost fish comprise more than 21,000 species, the largest group of vertebrates, inhabiting a wide variety of marine and freshwater environments from the abysses of the deep sea to high mountain lakes. Through more than 200 million years of evolution, this group has adapted to their habitats by adopting a diverse array of reproductive strategies [1]. A common principle for all fish, however, is the production of large yolky eggs through the development of the oocyte. The formation, development and maturation of the female gamete and ovum (oogenesis) are intricate processes that require hormonal co-ordination. Oocyte growth is normally divided into four main stages, primary growth, formation of cortical alveoli, the vitellogenic period, and final maturation [2].

Oocytes are female ovarian cells that go through meiosis to become eggs. They are derived from oogonia, mitotic cells that develop from primordial germ cells migrating into the ovary early in embryogenesis [3]. In teleost fishes,
full-grown postvitellogenic oocytes in the ovary are physiologically arrested at the G2/M border in first meiotic prophase and cannot be fertilized. In order for fertilization to occur, the oocytes must complete the first meiotic division and full-grown oocytes will resume their first meiotic division under appropriate hormonal stimulation. First meiotic division involves the breakdown of the germinal vesicle (GVBD: germinle vesicle, GV, is the oocyte nucleus), chromosome condensation, assembly of the first meiotic spindle, and extrusion of the polar body. These cells, often termed primary oocytes, become secondary oocytes after the first meiotic division, and then undergo the second meiotic division to become mature eggs. Histologically, the primary growth stage may be separated into several stages [4]. The nucleus first contains one nucleolus, thereafter multiple nucleoli and later a "circum nuclear ring" of ribonuclear material develops, which may contain a distinct yolk nucleus (Balbiani's vitelline body). Towards the end of the vitellogenic period, or by the beginning of the final maturation, the germinal vesicle (nucleus), which in the early stages is centrally located, moves to the periphery next to the micropyle [4]. Thus, the position of the germinal vesicle and the oocyte size may be used to estimate the start of final maturation.

In adult fish, the ovaries are generally paired structures attached to the body cavity on either side of the dorsal mesentery, except in lampreys [5] and some teleosts [6], where the two ovaries fuse into a single structure during development. In hagfish [5] and some elasmobranchs [7], only one ovary develops to adult. The structure of the growing ovarian follicle is remarkably similar in most fishes. The developing oocyte is located in the centre of the follicle and is surrounded by steroid producing follicle cells. The follicle cell layer generally consists of an inner sublayer, the granulosa cell layer, and one or two outer sublayers of theca cells. The theca and granulosa cell layers are separated by a basement membrane. Between the surface of the oocyte and the granulosa cell layer there is an acellular layer, the zona radiata or eggshell. During oocyte development, the zona radiata proteins (Zrp) are sequestered from circulating plasma and deposited in this position. At the same time, the oocyte is being filled with yolk proteins (lipovitellin, phosvitin), derived from vitellogenin (Vtg), another plasma protein found in sexually maturing female fish. Both of these protein groups, the Zrp and Vtg, so important constituents of the mature oocyte, are synthesized in the fish liver under endocrine regulation through the hypothalamic-pituitary-gonadal-liver axis. Herein, we will discuss the functional and developmental aspects of these hepatic-derived proteins, their regulation and role in oocyte maturation and fish reproduction. In addition, the use of these proteins as sensitive predictive and prognostic indicators for environmental endocrine disrupting chemicals will also be discussed.

**Endocrine regulation of oogenic proteins**

Pituitary gonadotropins (GtHs) and ovarian steroid hormones regulate oocyte growth and maturation in teleosts and other vertebrates [8]. Environmental changes, such as water temperature and photoperiod provide the cues to the central nervous system that triggers the maturation processes (Fig. 1). In response, the hypothalamus secretes gonadotropin-releasing hormone (GnRH). As the central regulator of hormonal cascades, GnRH stimulates the release of GtHs from the pituitary (Fig. 1). Although several GtHs have been identified from the teleost brain extract [9], two GtHs (GtH I & II) structurally similar to human follicle-stimulating hormone (FSH) and luteinising hormone (LH), respectively, are secreted from the teleost brain [10]. GtH I (FSH) is involved in vitellogenesis and zona genesis, while GtH II (LH) plays a role in final oocyte maturation and ovulation [8,10]. GtH secretion is regulated through a feedback mechanism by estradiol-17β (E2) and testosterone [9]. Several feedback mechanisms also act on the gonadal development through the hypothalamus-pituitary-gonadal-liver axis, because these organs produce substances influencing each other, leading to gonadal development and spawning [9,10]. GnRH release is inhibited by dopamine, which in turn is affected by steroid levels [9]. In addition to being a precursor for E2 and exerting feedback signals to the brain, testosterone is known to enhance stimulatory effects of gonadotropins in vitro [11]. Testosterone may also be involved in oocyte development [12], through the initiation of GVBD during final oocyte maturation [13].

E2 is the major estrogen in female teleosts, but large amounts of the androgen, testosterone, is also produced by the ovary. The ovarian two-cell model synthesizes E2 and testosterone, where the theca cells synthesize testosterone, which is subsequently aromatized by cytochrome P450 aromatase (CYP19) to E2 by the granulosa cells [8,14]. E2 stimulates the production of Vtg and eggshell Zr-protein by the liver of female fish [15–19], as described below.

**Egg yolk proteins**

In oviparous animals, accumulation of yolk materials into oocytes during oogenesis and their mobilization during embryogenesis are key processes for successful reproduction. As mentioned above, most oocyte yolk proteins and lipids are derived from the enzymatic cleavage of complex precursors, predominantly Vtg and very low-density lipoprotein [1,3,20,21]. Yolk is then stored until the late stages of oogenesis, and is mobilized in the embryo to facilitate the hydration process in buoyant eggs and provide the nutrients for embryogenesis [21,22]. Vitellogenesis is defined as E2-induced hepatic synthesis of egg yolk protein precursor, Vtg, its secretion and transport in blood to the ovary and its uptake into maturing oocytes [1,23–
Vtg is a bulky (MW; 250–600 kDa) and complex calcium-binding phospholipoglycoprotein (ibid.). The classification of Vtg as phospholipoglycoprotein indicates the crucial functional groups that are carried on the protein backbone of the molecule, namely, lipids, some carbohydrates, and phosphate groups [23,27]. In addition, the ion-binding properties of Vtg serve as a major supply of minerals to the oocytes.

Oocyte growth in fish is due to the uptake of systemic circulating Vtg, which is then modified by, and deposited as yolk in the oocyte [28] (Fig. 2). Vtg is selectively sequestered by growing ovarian follicles by receptor-mediated endocytosis before deposition in the oocyte [23,29,30]. These specific oocyte Vtg receptors are clustered in clathrin-coated pits. Coated vesicles fuse with golgian lysosomes in the outer ovoplasm of the oocytes and form multivesicular bodies [31]. The golgian lysosomes contain cathepsin D, which process Vtg into yolk proteins [32]. Vtg is an important source of nutrients for egg and larvae, making the vitellogenesis an important developmental process. In addition, teleost eggs contain maternal sex steroids [33], cortisol, and other lipophilic hormones like thyroxin that may enter the egg through Vtg [30,34]. It is not well understood which biological role(s) hormones in eggs play. However it has been hypothesized that they may act as metabolites or as synergists with other substances during early development.

**Eggshell proteins**

The envelope surrounding the animal egg plays significant roles in the reproductive and developmental processes; firstly as an interface between the egg and sperm, and secondly as an interface between the embryo and its environment [35]. The egg envelope is a major structural determinant of the eggshell in fish, and is often referred to as *zona radiata* because of its striated appearance under the light microscope [16] (Fig. 2). In mammals, these proteins function as sperm receptors and undergo a hardening process (also in fish) after fertilization. This process is important for the prevention of polyspermy, because the fish eggshell contains only one narrow canal or microple through which sperm gain access to the egg. In fish, the egg envelope is much thicker than in mammals, providing physical protection from the environment and playing a role in diffusive exchange of gases [35]. The microple is closed within minutes after the eggs are activated by exposure to fresh water, which initiates a cortical reaction necessary for development of fertilized eggs [36]. Ionic concentration of the medium lower than 0.1 M is needed for complete activation [37]. After activation, the *zona radiata* takes up water, gains resistance to breakage and can support up to 100 times more weight than oviductal eggs [38,39].

In eutherian mammals and fish, the *zona* proteins are composed of three-four distinctly conserved glycoproteins, but the differences in nomenclature and terminology complicates comparison. Several of the genes that encode the *zona* proteins have been characterized. For example, the exon-intron maps and coding sequence of mouse, pig and human homologues of *zona pellucida*, Zp2 [40–42], and mouse, human and hamster Zp3 [43–46] are well conserved. Thus, it has become increasingly clear that the proteins of the *zona pellucida* are conserved among eutherian mammals and that the proteins of the egg envelope are conserved among teleostean fish.

It has recently become more apparent that the proteins from the mammalian egg envelope are distinctly related to those of the teleostean envelope [47,48]. It was found that the synthetic site of *Zr-protein* is the liver in most teleost species. For example, rainbow trout, cod, and Atlantic salmon [18,19,48], medaka, *Oryzias latipes* [49–51], winter flounder, *Pseudopleuronectes americanus* [52], and...
gilthead seabream, Sparus aurata [53], synthesize Zr-protein in the liver. Other species, such as carp, Cyprinus carpio [54,55], and pipefish, Syngnathus scovelli [56] appear to synthesize Zr-protein in the ovary. Hence, the primary sequence of Zr-proteins is known in many teleost species, including winter flounder [52], medaka [49,50], carp [54,55], Atlantic salmon [48], and rainbow trout [57–59]. Recently, the full genomic sequences of medaka Zrp genes (choriogenin L and H) were reported [60]. The genes were 2142 and 2643 bp long, and contained eight and seven exons, respectively. The H form was reported to contain a much longer exon 1 due to the presence of seven proline-rich amino acid tandem repeats. Similar repeats in the N-terminal region of Zrp genes have been reported from other fish species [48].

Zonagenesis is the E$_2$-induced hepatic synthesis of eggshell proteins, zona radiata proteins (Zrp), their secretion and transport in blood to the ovary and uptake into maturing oocytes

**Terminology**

In fish, the major portion of the egg envelope (i.e. the inner layer) has been varyingly labeled as pellucid or vitelline membrane, zona pellucida, chorion, eggshell, primary, secondary and tertiary envelope, zona radiata ( interna and externa) or vitelline envelope [61–64] and some have suggested the term choriogenin for the precursor proteins found in plasma [50]. Comparative ultrastructural analysis of zona radiata from six salmonid species showed basic similarities, but species differences in the structure of zona radiata interna [65]. Since 1989, several reports have demonstrated the hepatic synthesis of precursor proteins of the inner layer subunits under the influence of estrogen, at least in most species [16–19,51,66]. Despite the confusing terminology used to designate this very
important class of structural protein in teleost fish and its critical role in development, there is still no commonly accepted term for these proteins [59]. However, the use of the above named terms has basically been for descriptive, structural and functional purposes. In the present context, the term "zona radiata proteins" (Zr-proteins) will be used to identify the constituent proteins of the inner layer of the envelope that surrounds the oocyte of the ovulated teleost egg. We have used zona radiata proteins, a descriptive term, to designate these proteins because of the striated appearance of this structure in light microscope, in accordance with the recommendations of Oppen-Berntsen [16]. We also use the term to describe the soluble protein monomers found in synthesizing liver cells and circulating in plasma.

**Molecular mechanisms for oogenic protein gene expression**

Vitellogenesis and zonagenesis are crucial for the reproduction of oviparous animals. The cellular and molecular events that occur in tissues that produce oogenic proteins and in the ovary provide ideal systems for the study of several fundamental biological processes [67]. For example, the abundantly transcribed Vtg genes are being used to analyze stage-, sex-, tissue- and hormone-specific gene expression. One research area that has received a lot of attention in recent time is xenobiotic modulation of gene expression in organisms (see later). Thus, selective gene expression is considered to be central to our understanding of cellular differentiation and the regulation of developmental processes [68]. The term gene expression is not always well defined, but most often it is used to indicate a change in the nature of, or rate at which, different genes are transcribed [15]. Recent advances in studies of the organization of eukaryotic genomes have also focused attention on the importance of structural features of expressed and unexpressed genes and on the post-transcriptional mechanisms that would determine the processing of primary transcripts into the correct messenger sequences [69,70].

Figure 3 shows an order of the molecular mechanisms that lead to the production of Zr-protein and Vtg in the hepatocyte: (1) E2 produced by the ovarian follicular cells in response to GtH I is transported in plasma attached to sex hormone binding globulins (SHBGs: [71–76]) and enters the liver cells by either diffusion or receptor-mediated uptake. The physiological functions of the SHBGs are not fully understood. It is generally believed that these proteins play a role in the regulation of steroid amount available to target tissues and protect steroids from rapid metabolic degradation [77,78]. In addition to their role as sex steroid carriers, it has been proposed that SHBGs are involved in cellular signal transduction that involves nuclear steroid receptors through specific SHBGs membrane receptors in different sex steroid sensitive tissues [for review see, [78]]. (2) In the liver, E2 is retained in target cells by high affinity binding to a specific steroid-receptor protein, the E2-receptor (ER: [80]). In the absence of a ligand the ER is found as a monomer in association with heat shock protein 90 (hsp90). In the ligand binding process, the ER dissociates from hsp90 and usually goes through dimerization prior to translocation of the complex into the nucleus, involving a complex of coregulator proteins (more details on the molecular biology of ER forms and the events taking place in this process can be found in reviews such as [80–83]). (3) The hormone-receptor complex binds tightly in the nucleus at estrogen responsive elements (ERE) located upstream of, or within the estrogen-responsive genes in DNA. (4) This results in the activation or enhanced transcription of Vtg genes and subsequent increase and stabilization of Vtg messenger RNA (mRNA). At present, ERE for Zr-protein genes have not been identified in fish, although their response to E2 is very similar to that of the Vtg genes. Given the speculation that different ERs on the DNA may be temporarily masked by associated proteins, thus resulting in sequential or partial induction of various estrogenic responses [84], it is possible that there may be subtle differences in the responsive elements for Zr-protein and Vtg. (5) Zr-protein and Vtg precursors are synthesized and modified extensively in the rough endoplasmic reticulum (RER); (6) modified Zr-protein and Vtg are secreted into the serum for transport to the ovary. (7) In the ovary, Zr-protein and Vtg are incorporated to serve different functions (see later).

The post-translational modifications occurring to the Zr-proteins prior to secretion into the systemic tracks are not well understood. However, more is known about Vtg post-translational modifications in teleost fish. Prior to secretion into the blood stream, the biochemical information concerning Vtg clearly indicates that substantial post-translational modification must occur in the liver cell to reach the end product seen in the serum. Several changes in hepatic morphology such as proliferation of RER and Golgi apparatus also accompany estrogen stimulation. Firstly, the protein backbone of the Vtg is synthesized on membrane bound ribosomes. Vtg shares this feature with other proteins destined for secretion from the hepatocytes [85]. Thereafter, the Vtg molecule is lipidated, glycosylated and phosphorylated. Although some information exists concerning the nature and extent of modifications of the Vtg molecule, rather limited information is available for fish with respect to the mechanisms, sequential events or location of these transformations.

Several metabolic changes occur during Vtg synthesis in the maturing female fish. This is reflected in the pronounced increases in liver weight, RNA contents, lipid
deposition, glycogen depletion, increases in plasma protein, calcium and magnesium and phosphoprotein contents [86,87]. These parameters can be used as indicators of plasma Vtg levels. In addition, Vtg and gonadal maturation are energetically very expensive processes, since the fullgrown gonads account for about 25% of the total weight of a mature female fish. The uptake of Vtg by growing oocytes is rapid, specific and saturable, and occurs by receptor-mediated endocytosis [88,89]. Vtg receptors (VTGRs) have been identified in the ovary of a number of fish species [see 3, [90–92]], and was recently cloned and sequenced in rainbow trout and winter flounder [93–95]. The fish VTGRs are 70–80% similar to the chicken very low-density lipoprotein receptor VLDLR (ibid.). The enzymatic cleavage and processing of Vtg into oocyte yolk proteins and lipids is mediated by serine proteases and cathepsins found in ovary extracts [21,94]. After uptake, the Zr monomers are cross-linked by a transglutaminase reaction to form the rigid structure of the fish eggshell inner layer [16].

Figure 3
Simplified diagram of estradiol-17β (E2) or E2-mimic stimulated oogenic protein synthesis. Eggshell zona radiata proteins and the egg yolk protein precursor, vitellogenin are synthesized and secreted by the hepatocyte. They are transported in blood to the ovary and incorporated into maturing oocytes in female teleosts.
**Effects of xenobiots on oogenic protein synthesis**

The terms environmental estrogens, endocrine disruptors, endocrine modulators, eco-estrogens, environmental hormones, xenoestrogens, hormone-related toxicants, and phytoestrogens all have one thing in common, namely, they describe synthetic chemicals and natural plant or animal compounds that may affect the endocrine system (the biochemical messengers or communication systems of glands, hormones and cellular receptors that control the body's internal functions) of various organisms. Many of the effects caused by these substances have been associated with developmental, reproductive and other health problems in wildlife and laboratory animals [for reviews, see [97–100]]. There is also growing concern that these compounds may be affecting humans in similar ways [101,102].

The detailed mechanisms by which xenoestrogenic compounds mediate their induction of oogenic proteins is not fully understood, but it is known that they can bind with high affinity to the ER (as agonists) and initiate cell synthetic processes typical of natural estrogens. Some compounds also have the ability to bind to the receptor, but not eliciting estrogenic activities (as antiestrogens or antagonists), thereby blocking the binding site of natural estrogens [103–105]. During ovarian recrudescence, incorporation of oogenic proteins accounts for the major growth of the developing oocytes. A probable indirect measure of altered hepatic oogenic protein synthesis in fish exposed to xenobiotics is reduced or increased gadosomatic index (GSI). A more direct quantification of these alterations can be obtained from plasma, hepatic and ovarian oogenic protein concentrations [106]. Modern and advanced molecular biology techniques are revolutionizing the process of oogenic protein quantitation in oviparous species [99].

Laboratory studies have been conducted to evaluate the impact of fish exposure to toxicants on ovarian development. Several effects have been observed and these include inhibition of oocyte development and maturation, increased follicular atresia of both yolked and previtellogenic oocytes, abnormal yolk deposition and formation within oocytes, and abnormal egg maturation and production [for reviews, see [98,99,102,106–108]].

Wester and Canton [109] observed the development of testis-ova in males and induced vitellogenesis in either sex of medaka (Oryzias latipes) exposed to β-HCH, demonstrating estrogenic effects of this compound. Similar responses have been observed when medaka was exposed to 4-nonylphenol (NP) and to bisphenol in more recent studies [110–112].

In designing a bioassay for xenoestrogens, toxicologists and biologists have used the induction of Vtg and Zr-protein in male and juvenile oviparous vertebrates as an effective and sensitive biomarker for xenoestrogens [113–118]. Using juvenile Atlantic salmon (Salmo salar) and different doses of NP, we saw that NP treatment significantly elevated plasma levels of Zr-protein and Vtg in a two week in vivo study, with the former showing more sensitivity to the xenoestrogen compound [115]. Higher sensitivity of Zr-protein when compared with Vtg evaluated with indirect ELISA has also been observed in with juvenile Atlantic salmon treated with different doses of an oil refinery treatment plant effluent [[115], Fig. 4] and with E2 [119]. In both these studies, induced Zr-protein levels were apparent at lower E2 doses, while Vtg was only induced at higher E2 doses, thus indicating differential induction of both proteins as was observed using NP [115]. However, it could be argued that the differences in sensitivity could arise from different affinities of the antibodies used in the assays. Attempts to resolve this issue have focused on the development of quantitative assays for the two protein groups and their mRNAs (see below). In a recent study with medaka, Lee et al. [51] reported a differential sensitivity of the two zona radiata precursor genes choriogenin H and L, respectively, with choriogenin L mRNA responding at lower doses of estrogen than mRNA of the H form. Unfortunately, however, they did not compare the response directly with Vtg mRNA. In the study of Yadetie et al. [120], no clear differences were observed in the response of Vtg and Zrp mRNA levels of salmon exposed to NP. However, Celius et al. [57], employing a quantitative real time polymerase chain reaction assay (qPCR) for rainbow trout Vtg and Zrp, reported that Zrp mRNA was more responsive than Vtg mRNA to low doses of E2 and the mycoestrogen α-zearenol.

Furthermore, a large number of in vivo studies have also reported Vtg induction by xenobiotic estrogens in fish and amphibians, e.g. jobling et al. [121] using rainbow trout (Oncorhynchus mykiss) and alkylphenolic chemicals; Donohoe and Curtis [122] using juvenile rainbow trout, o, p'-DDT and o, p'-DDE; Schwager et al. [123] using rainbow trout, common carp (Carpio carpio) and NP; and Janssen et al. [124] using flounder (Platichthys flesus) and polluted harbour sediment [reviewed in [99,102]]. All these studies showed significant elevations of Vtg at the tested dose of the chemicals. In other studies, Sumpter and Jobling [125], Pelissero et al. [126], Jobling and Sumpter [127], Celius et al. [128], have reported the in vitro induction of yolk protein synthesis (in a dose-dependent manner) of several environmental chemicals, including alkylphenol ethoxylate (APE) metabolites [129]. Both in vitro and in vivo studies have been used to study oogenic protein synthesis in fish. In a few studies where the two approaches have been directly compared, it has been shown that in...
vitro assessments for estrogenicity underestimate the in vivo response [114]. This is particularly evident with chemicals that require metabolic activation (proestrogens) or are capable of substantial bioaccumulation. In addition, they do not provide information on possible physiological alterations. Given that in vitro systems lack the complex metabolic processes that are typical of in vivo systems, the former system should only be used as a supplement to the latter system, and short-term in vivo assays using plasma Vtg measurements in small test fishes have been suggested to screen individual existing or new chemicals for estrogenic potency (ibid.).

Endocrine disruptors can also target other sites of the hypothalamus-pituitary-gonad-liver axis (Fig. 1), e.g. pituitary GtH release or ovarian aromatase activity [130,131]. However, this aspect is outside the scope of this review.

Use of Vtg/Zrp as biomarkers in chemical product testing

The increased awareness that chemicals in the environment can cause endocrine disruption in wildlife and, possibly, humans, has lead international organizations such as OECD to consider developing new test methodologies for detecting EDCs. These methods will eventually be used as standard test procedures in the toxicity testing of new and existing chemicals. Recent work in OECD and the US Environmental Protection Agency has focused on reviewing available methods for detecting endocrine disrupting effects of chemicals in wildlife, including fish. An implementation of Vtg as a core endpoint in a piscine short-term endocrine disrupter screen for chemicals, in combination with e.g. gross morphology and histology, is suggested. The tests should be applicable to different species, in particular zebrafish (*Danio rerio*), fathead minnow (*Pimephales promelas*), and medaka (*Oryzias latipes*) [132]. These fish share several attributes that make them ideal test species for reproductive toxicity testing, including small size at maturity, relatively short generation times, asynchronous spawning, and overall ease of culture. Sensitive and quantitative immunoassays for Vtg in these species have recently been developed in our laboratory [133].

Oogenic protein assays

Depending on the target organ or tissue, a wide variety of assays have been developed to measure oogenic protein expression in fish. These include radioimmunoassays; enzyme-linked immunosorbent assays (ELISAs) and immunohistochemistry using monoclonal and polyclonal antibodies (Abs), RNA protection assay and transcript analysis by Northern blotting or various variants of
polymerase chain reaction (PCR). Recently, the use of real-time (quantitative) PCR is increasingly becoming a valuable tool in oogenic protein analysis. In plasma samples, these assays vary in their sensitivity, but some have the ability to detect very low levels of protein expression, i.e. 1 ng/ml or less [134–137]. Vtg assays based on polyclonal antibodies are generally restricted for use with the homologous species, but some antibodies do cross-react with Vtg in other species (e.g. [135,138,139]) (Fig. 5).

The basic principle of a radioimmunoassay (RIA) is the use of radio labeled Abs or antigens (Ags) to detect Ag:Ab reactions. The Abs or Ags are labeled with the $^{125}$I (iodine-125) isotope, and the presence of Ag:Ab reactions is de-

Figure 5
Cross-reactivity of a monoclonal zebrafish (Danio rerio) vitellogenin antibody to different cyprinid fish species. Monoclonal mouse anti-zebrafish vitellogenin IgG JE-10D4 (Biosense Laboratories AS, Bergen, Norway) was used to probe a Western blot with samples of: (1) Pre-stained molecular weight standard (Bio-Rad), (2) purified zebrafish Vtg, (3) whole-body homogenate sample of estradiol-17\(\beta\) (E\(_2\)) treated zebrafish, (4) whole-body homogenate sample of control zebrafish, (5) plasma sample of E\(_2\) treated carp (Cyprinus carpio), (6) plasma sample of control carp, (7) plasma sample of E\(_2\) treated fathead minnow (Pimephales promelas), (8) plasma sample of control fathead minnow, (9) plasma sample of E\(_2\) treated roach (Rutilus rutilus), (10) plasma sample of control roach. Reproduced with permission from Biosense Laboratories AS.
tected using a gamma counter. RIA techniques are well developed for egg yolk (Vtg) analysis (e.g. [140,141]), but have not been developed for the zona radiata proteins. Because this technique requires the use of radioactive substances, RIAs are more and more being replaced by other immunologic assays such as ELISAs, that over the last decade have reached similar levels of sensitivity.

The ELISA technique is a sensitive laboratory technique widely used to detect and quantitate Ags or Abs in a variety of biological samples. It can be quantitative (with a standard curve) or semi-quantitative (without a standard curve). The two most widely used principles for quantitative detection of proteins are the competitive ELISA and the sandwich ELISA techniques [142].

In addition to the general issues of antibody specificity and sensitivity, there are some specific challenges related to the development of quantitative immunoassays for the oogenic proteins Vtg and Zrp. For Vtg, although it is relatively easily purified from plasma of estrogenized fish (where it can reach levels of 50–150 mg/ml), it is an inherently unstable protein. The instability of Vtg is due to its role as a precursor for shorter peptide fragments, and it is very sensitive to proteolytic breakdown into these fragments. Care must therefore be taken during sampling to avoid proteolytic breakdown by adding suitably protease inhibitors [96]. This instability leads to some problems with immunization, since breakdown products may be more immunogenic than Vtg itself. In addition, it creates an important problem for the use of Vtg as a standard in quantitative assays, since users must ensure that each batch of standard is stored under conditions that prevent breakdown, and is quantitated in a consistent manner (see below). In our own laboratory, we have had success in finding conditions for stabilizing Vtg by lyophilization, although this has not been a straightforward task, and different species behave differently in this process (Goksøyr, Nilsen, Berg et al., unpublished results).

The dynamic range of Vtg concentrations found in fish plasma creates another problem. Plasma Vtg can vary maybe 100 million-fold, from a few ng/ml in unexposed male fish, to the 50–150 or above mg/ml found in estrogenized salmonids (e.g. [136]). To be able to quantitate this enormous range in blind samples, the working range of the assay should preferably be as wide as possible. Nevertheless, even with an assay covering several hundredfold variation, all samples need to be serially diluted at least 3–4 times to ensure that at least one dilution falls within the working range of the assay. Many of the recent assays published obtain this range (e.g. [133]).

The assay also needs to be robust and reproducible, and current experience in our laboratory demonstrates that the sandwich type ELISA is more robust and reproducible over the working range of the assay compared to the competitive format.

The method used to quantify the standard must be consistent and reliable. For Vtg, many different methods are presented. In some cases, Vtg is weighed after a lengthy purification procedure. Others have used different protein quantification methods such as Lowry [143], Bradford [144], or the simple A280 absorbance measure. In all these cases, the sample needs to be quantitated towards a known sample. When bovine serum albumin (BSA), ovalbumin, or Immunoglobulin G is used, an assumption is made that Vtg behaves more or less similar to the chosen standard. Generally, this is not the case, and some laboratories develop their own "gold standard" of Vtg, which is used as the standard in quantitation. Again, this gold standard needs to be verified, and this can be done by quantitative amino acid analysis. In this case, one may want to take into account the non-proteinaceous parts of the Vtg, i.e. the lipid, phosphate, and carbohydrate parts. The lipid and phosphate parts have been reported for some species to represent 15–20% and 0.6–0.8%, respectively (e.g. [27]), whereas the carbohydrate portion is not well studied. In general, however, the protein part of the molecule is calculated to represent around 65–75% of the weight of the whole molecule, depending on species. The most important aspect of a protein to be used as a standard in an immunoassay is of course that the epitope(s) involved in the immunoassay maintain their stability. This can only be checked by a quality control using the immunoassay itself, so the question becomes a "hen or egg" issue. One way to manufacture a Vtg standard that maintains both proteolytic and epitope stability is to produce a synthetic peptide fragment that contains the epitope(s) of interest.

For Zrp, the challenges are somewhat different. Zrp are found in lower concentrations in plasma compared to Zrp, but recent analyses show that they may reach levels of 1–10 mg/ml in estrogenized rainbow trout [145]. The protein is much more stable than Vtg, probably due to the different natures of their fate in the oocyte. Whereas Vtg needs to be broken down to fulfill its role as nutrient for the embryo, the Zrp needs to be incorporated into the eggshell intact. In the eggshell, the Zrps will cross-link by a transglutaminase reaction to form the robust zona radiata structure upon fertilization and hardening [146]. The solubilization of Zrp from eggshells requires harsh conditions (ibid.), whereas it is more easily obtained from plasma. Although polyclonal antibodies for Zrp have been developed and used for some time [115,119], monoclonal antibodies (MAbs) to Zrp have only recently become available [147]. Screening a large panel of MAbs, it has become clear that the α- and β-form of Zrp are closely
related to each other, whereas the γ-form is structurally more different (Fig. 6; Berg, Bringsvor, Nilsen, Goksøyr, unpublished results). We have also shown that combining a γ-specific MAb with a polyclonal Zrp-antibody can be used to develop a quantitative sandwich ELISA for γ-Zrp, where the standard γ-form can be purified from plasma using the same MAb in immunoaffinity chromatography [145]. Because of the close similarity between the two other isomers, this has proven more difficult for the α- and β-form. However, comparing their relative responses both in ELISA and Western blots, it becomes clear that the α- and β-form are more responsive to estrogens than the γ-form of Zrp (Berg, Bringsvor, Nilsen, Goksøyr, unpublished results).

Oogenic mRNAs can be assayed by reverse transcriptase polymerase chain reaction (RT-PCR, e.g. [25]), or quantitative PCR techniques (qPCR, [57]). qPCR is a rather new method for the quantification of target mRNA sequences. Unlike conventional PCR, qPCR systems are probe-based PCR product detection. During amplification, annealing of the probe to its target sequence generates a substrate that is cleaved by the 5’ nuclease activity of Taq DNA polymerase when the enzyme extends from an upstream primer into the region of the probe. This dependence on polymerization ensures that cleavage of the probe occurs only if the target sequence is being amplified. The development of fluorogenic probes made it possible to eliminate post-PCR processing for the analysis of probe degradation. The probe is an oligonucleotide with both a reporter fluorescent dye and a quencher dye attached. While the probe is intact, the proximity of the quencher greatly reduces the fluorescence emitted by the reporter dye by fluorescence resonance energy transfer (FRET) through space.

Probe design and synthesis has been simplified by the finding that adequate quenching is observed for probes with the reporter at the 5’ end and the quencher at the 3’ end. The qPCR has several advantages compared to other hybridization techniques. This includes; fluorogenic probes over DNA binding dyes require specific hybridization between probe and target to generate fluorescent signal. Thus, with fluorogenic probes, non-specific amplification due to mis-priming or primer-dimer artifact does not generate signal. Another advantage of fluorogenic probes is that they can be labeled with different, distinguishable reporter dyes. By using probes labeled with different reporters, amplification of two distinct sequences can be detected in a single PCR reaction. The disadvantage of fluorogenic probes is that different probes must be synthesized to detect different sequences.

Other mRNA targeting assays for oogenic proteins, such as the RNA protection assay [148], have also been developed.

Cellular localization of hepatic oogenic protein synthesis has also been demonstrated using immunohistochemical analysis of exposed fish with specific antibodies [149,150] (Fig. 7). Immunohistochemistry is a valuable tool in the studies of estrogen and estrogen mimicking compound induced hepatic synthesis of Vtg and Zrp in oviparous vertebrates, especially in situations where blood samples are difficult to collect, e.g. when studying small-sized species. Although this technique is time-consuming, localization of Vtg in liver sections may provide insight into responses of different cell types that are important for understanding the role and mechanisms of the estrogens and estrogen mimicking compounds.

Effects and interactions of complex chemical mixtures

There are many potential xenobiotics and xenoestrogens in aquatic systems (e.g. pharmaceuticals, pesticides and personal care products). Thus, in the environment, chemical interactions have profound consequences since organisms, including fish, are exposed to complex mixtures of environmental pollutants [117]. These complex interactions have only recently become the focus of systematic investigations. There is no doubt that biomarkers (of exposure to environmental hazards, of effects to environmentally-induced cellular/molecular changes and of genetic susceptibility) are revolutionizing the science of risk assessment. Biomarker measurements have the ability to improve our accuracy, reliability and scientific basis for the quantitative assessment of environmental health risks. The relative importance of the influence of contaminants on biological systems is not well-understood or quantified mechanistically in complex chemical mixtures.

For example, exposure of juvenile rainbow trout to different doses of E2 and CYP1A-inducers showed both elevation and reduction of plasma Vtg levels, depending on relative ratios of the test compounds [151–153]. In a recent study, exposure of juvenile salmon to an estrogen mimic (NP) and a CYP1A-inducer with documented anti-estrogenic activity (3,3’4,4’-tetrachlorobiphenyl; PCB-77) resulted in the potentiation of NP-induced synthesis of Vtg and Zr-proteins [117]. In addition, this study also showed that the reported effect depends on NP and PCB-77 ratios, seasonal factors and in which order the two compounds were given. Using the natural estrogen (E2) in fish and mammals, the antiestrogenic effects of aryl hydrocarbon receptor (AhR) agonists are paralleled by the induction of CYP1A-dependent monoxygenase activities such as EROD [151,152], several E2 hydroxylase activities and aryl hydrocarbon hydroxylase (AHH) [104,154,155]. AhR agonists do not competitively bind to the steroid hor-
Specificity of Atlantic salmon (Salmo salar) zona radiata protein antibodies. A plasma sample from estradiol-17β treated salmon was probed with different monoclonal Zrp antibody supernatants and the polyclonal mouse antiserum. (1) Clone 2C4, showing equal specificity for the α- and β-isomer, (2) clone 3D7, showing highest specificity for the α-isomer, (3) clone 7F2, a γ-specific clone, (4) clone 8C4, a predominantly α-specific clone, and (5) polyclonal mouse antiserum, showing reactivity with all three isomers. (Berg, Nilsen, Goksøyr, unpublished results).
mone receptors nor do steroid hormones bind to the AhR [156]. Therefore, the molecular mechanisms of interaction between ER and AhR agonists need to be explored in more details.

**Possible consequences of precocious Vtg and Zrp induction**

Reproductive development is a continuous process throughout ontogeny. Consequently, it is susceptible to the effects of xenoestrogens and/or xenobiotics at all stages of the life-cycle, including fertilization, embryonic development, sex differentiation, oogenesis or spermatogenesis, final maturation, ovulation or spermiation, and spawning. Thus, the sensitivity to a particular compound will vary depending on the stage of reproductive development [157].

Understanding the general principles by which chemical substances or foreign compounds (xenobiotics) interfere with fish reproduction is particularly important for meeting the larger objectives in aquatic reproductive toxicology, as it is impossible to empirically determine the biological specificity or how every compound affect the reproductive life-history strategy of every species. Here we will briefly discuss the specific effects that can be extrapolated from a precocious hepatic synthesis of Vtg and Zrp.

Given the energetic cost of reproduction and the long decision time, it seems most likely that xenobiotically-induced hepatic Zrp and Vtg synthesis may cause an imbalance in the reproductive strategy of a given fish population. The reason is that an organism can only acquire a limited amount of energy for which several processes compete directly; an increase in the energetic allocation to one process must result in a decrease in energy allocation to others [158–160]. Thorpe [161] suggested that during maturation, the internal responses that are synchronized by external signals depend upon some genetically determined performance threshold, and that maturation processes will continue if this performance exceeds a set point at this critical time. Furthermore, maturation has developmental priority over somatic growth, and in salmonids survival after spawning implies a chance dependent balance between stored energy and that spent on reproduction [162]. Therefore, xenoestrogen-induced Vtg

---

**Figure 7**

Immunohistochemical localization of vitellogenin (Vtg) in liver sections of control (a), nonylphenol- (b) and estradiol-17β-treated (c) juvenile Atlantic salmon (*Salmo salar*). Cellular Vtg levels were detected with mouse monoclonal antibody (BN-5) against salmon Vtg. Yellow colors show strong Vtg-specific staining and as demonstrated primarily in endothelial cells, hepatic sinusoids, and cytoplasm of hepatocytes (labeled C). Blue stains show nuclei of hepatocytes. Goat anti-mouse horseradish peroxidase (GAM-HRP) was used as secondary antibody. Reproduced from Arukwe et al. [143] with permission from Taylor and Francis [http://www.tandf.co.uk](http://www.tandf.co.uk).
and Zrp synthesis outside the normal maturation period may result in wasteful use of stored energy resources. The ecological implication of this might be failure in the reproduction of affected individual fish, and in the long-term affecting recruitment of the entire population. Another possible deleterious effect is that high Vtg and/or Zrp concentrations might cause kidney failure and increased mortality rates as a result of metabolic stress [163]. Furthermore, although not yet demonstrated, there is a possibility that the reduced testicular growth could reduce fertility [121].

Xenoestrogen-induced changes in Zrp synthesis appear to have a higher potential for ecologically adverse effects than Vtg induction, because critical population parameters such as offspring survival and recruitment may be more directly affected. The argument for this, is that whereas subtle changes in Vtg content would not be of great significance to the survival of the offspring, small changes in Zrp synthesis might cause the thickness and mechanical strength of the eggshell to be altered, thus causing a loss in its ability to prevent polyspermy during fertilization and to protect the embryo during development [115].

Intersex is another and a much more common condition caused by early exposure of fish larvae to estrogenic substances. The intersex condition in males usually takes the form of ovotestis. Bortone and Davis [164] have reviewed the subject, particularly with respect to the masculinisation of females caused by pulpmill effluents. Ovotestis is a partial feminization in which oocytes may appear in otherwise normal testes (Fig. 8). Little is known about the implications of this condition for reproductive functionality. Ovotestis can be induced in the laboratory by exposing fish larvae to weak estrogens like NP [110–
112], and has also been observed at prevalences ranging from 20 to 100% in wild fish populations exposed to estrogenic effluents [165,166].

**Evolutionary aspects**

From an evolutionary point of view, an appreciation of the oogenetic components will not be complete without considering how they evolved and what evolutionary factors have been driving their evolution. In the preceding discussions about oogenic proteins, it is clear that maternal influence in developmental modes that have direct roles in early embryonic patterning and gene regulation is not restricted to informational components such as maternal mRNAs and proteins as developmental biologists will generally consider it [167]. Maternal factors also include proteins and lipids that have structural or nutritive roles and that can play a large role in evolution of life histories and embryogenesis. The ability to transport fat, in the form of lipoprotein through the circulatory system by eukaryotes is one of their most significant functions right from the beginning of existence [168]. The reason and functional basis for why Vtg transport systems initially evolved provides clues into how energy in the form of water insoluble fat can be distributed from sites of synthesis and absorption to specific tissues and cells. Thus, the evolutionary advancement of storing energy in the form of fat has provided organisms with enormous advantage in adapting to environmental and developmental changes.

During vitellogenesis, oviparous species (i.e. nematodes, decapods, echinoderms, insects, fish, amphibians, birds, reptiles) display over three orders of magnitude increases in the transport of fat, for instance from the liver to oocytes, in order to facilitate egg development [3,15,67,168]. Thus, Vtgs can be regarded as ancient proteins that are normally encoded by a small variable number of genes [169]. One Vtg gene has been demonstrated in sea urchin and silkworm, three in the chicken, four in *Xenopus laevis* and six in *Caenorhabditis elegans* (ibid.). However, more recent analysis uncovered 20 Vtg genes and ten pseudogenes in rainbow trout [170]. Previously, comparative studies based on amino acid sequences and on gene organization support the hypothesis of a common evolutionary origin of the vertebrate and invertebrate Vtg genes [171–176]. More recently, this comparative analysis was extended to include other species groups in order to elucidate the events responsible for the relatively rapid diversification of the Vtg gene family [169].

For example, Vtg genes of the fruit fly, *Drosophila melanogaster*, are not obviously related to the Vtg genes of chicken, *X. laevis* and *C. elegans* [67]. The chicken and *Xenopus* genes both have 34 introns and have the same exon-intron organization. These genes are distantly related to the *C. elegans* gene with only four introns, whose positions are apparently conserved in vertebrates, suggesting their presence at the same positions in the ancestral Vtg gene [177,178]. It has been suggested that prokaryotes and lower eukaryotes have streamlined their genome by intron sequence elimination. Given that this is true, the structural organization of the contemporary vertebrate Vtg gene may be more representative of the earliest gene than those of the invertebrates [67,177,178]. This assumption is supported by the study of Mouchel et al [169], which suggests that almost half of the splicing junctions identified in invertebrates are related neither to each other nor to vertebrate genes. The observed differences between vertebrate and invertebrate Vtg gene structure may also be explained by the “intron-late” theory [179,180], which hypothesizes that introns became inserted more recently, therefore assuming that insertions may be specific to each lineage. Irrespective of what the explanation may be, it has been suggested that Vtg genes have been re-organized through multiple insertions and deletions of intervening sequences during the evolution of the various lineages. Characterization of the Vtg region in the genome of the rainbow trout, *e.g.*, revealed that this locus contains twenty complete genes and ten pseudogenes per haploid genome [170]. The Vtg genes differed from each other by insertion, deletion and rearrangement events, although, at the sequence level, they showed a high degree of similarity. Fluorescent in situ hybridization (FISH), pulsed-field gel electrophoresis (PFGE) and Southern analysis indicated that all gene copies are contained in a single 1,500-kb region, and that most of the genes form tandem arrays separated by a conserved 4.5-kb intergenic region. The presence of large reiterated fragments indicates that this region has been subjected to several amplification events. The presence of a retroposon element in Vtg intron 9 appeared to be responsible for the silencing of at least nine of the ten pseudogenes (ibid.).

It has become increasingly clear that the proteins of the *zona pellucida* are conserved among eutherian mammals and that the proteins of the *zona radiata* are conserved among teleostean fish. In most fish, sperm lack an acrosome and penetrate the *zona radiata* surrounding fish eggs via a discrete micropyle [40]. Most commonly, the micropylar channel is sufficiently narrow to permit the passage of a single sperm, and subsequent fusion with the plasma membrane induces the cortical granule reaction, resulting in a block to polyspermy [52]. In contrast, a prerequisite to successful fertilization in all vertebrates is penetration of sperm through an acellular envelope surrounding ovulated eggs. In mammals, sperm bind to the *zona pellucida*, the mammalian equivalent of fish *zona radiata*. Following the induction of the acrosome reaction and release of lytic enzymes, sperm penetrate the *zona* and fuse with the egg’s plasma membrane, triggering the post fertiliza-
tion block to polyspermy [181]. More recently, it has become apparent that, although critical for speciation, the proteins from the mammalian egg envelope are distinctly related to those of the teleostean envelope [48]. Recently, the mouse zona proteins was successfully incorporated into the extracellular envelope surrounding Xenopus eggs, showing that they have been sufficiently conserved through 350 million years of evolution [182]. In general, the exon conservation at the same region in mammalian zona pellucida and fish zona radiata protein suggests that not only has this protein domain been duplicated in mammals, but that it has been conserved and used as an egg envelope protein in species that diverged 650 million years ago.

How ancient are these important components of eukaryotic reproduction? Recently, Walther [183] summarized his hypothesis that the oocyte and the sperm represent cellular lineages dating back to the two prokaryotic cell domains (eubacteria and archa, respectively), which gradually evolved ever more complete but reversible coalescence or syngamy instead of permanent fusion to form an equilibrium between the two moneric prokary and the prototypic zygote under photosensitive polar conditions two billion years ago. The now commonly accepted theory of endosymbiosis as the origin of eukaryotic cells was presented independently by Jostein Goksøyr [184] and Lynn Margulis (Sagan) [185] to account for a eukaryotic cell with organelle. Walther's theory instead inserts a primordial syngamy to a dimeric prokaryotic cell (termed A-KARYON), and proposes that such sexual syngamy was the origin of symbiosis leading to organelles. According to this theory further endosymbiosis [184,185] created in one event the eukaryotic nucleolmma and the outer membrane of the mitochondrion, as the second step in cell evolution from moneric prokary to dimeric eukarya. This theory depicts cell evolution by a dynamic interaction between only two moneric species in a unique event in cell evolution, which established sexuality as the dynamic fusion of these two cells or species. This dynamic model contrasts markedly from the view of an evolutionary past where cell evolution occurred by fusing a multitude of prokaryotic cell types to yield the static eukaryote, among which some later acquired sexuality (see also Margulis and Bermudes, [186]).

The evolution of the eggshell and egg yolk protein genes would appear to have been driven by different factors (protection vs. nutrition), but still in modern ovoiparous vertebrates they are being synthesized in close concert by the hepatic machinery under a common endocrine regulation. There is still a lot to be learnt about when these genes appeared and how they evolved in the interplay between hormones, environmental cues, speciation, reproductive strategies, and the hepatic organ as their major site of synthesis today.

Conclusions

Different reproductive strategies have evolved among vertebrates, based on energy requirement, mating behavior, gamete structures, and the specificity of recognition molecules on the surface of sperm and eggs. In teleosts, environmental changes, such as photoperiod and water temperature provide signals that are received by the central nervous system. These signals lead to oocyte growth and maturation that are regulated by pituitary gonadotropins and ovarian sex steroids. An integral part of this process is the synthesis of the oogenic proteins, Vtg and Zr-proteins. E2 is the major estrogen in female fish. E2 stimulates the production of Vtg and Zr-proteins in the liver. The genes encoding these fish reproductive proteins are conserved in the animal kingdom and are products of several hundred million years of evolution.

An increasing number of widely used chemicals and their degradation products are found now to induce precocious synthesis of oogenic proteins in fish. Convincing evidence of this effect has been obtained from studies at the molecular and cellular levels of biological organization, in addition to reports on the individual level from laboratory studies. In addition, there are numerous reports demonstrating that fish populations are adversely affected by living in, and accumulating xenoestrogens. Although xenoestrogen-induced synthesis of oogenic proteins appears to possess a potential for ecologically adverse effects, as does inhibition and elevation of biotransformation enzymes, studies are still needed of critical population parameters such as offspring survival and recruitment to validate these findings at higher levels of biological organization.

Authors' contributions

AA and AG contributed equally to this review, with AA taking the lead.

Acknowledgements

We want to thank our collaborators and co-authors for their contribution to the work from our own laboratories that has been presented here, and Professor Bernt Walther and an anonymous referee for helpful comments to the manuscript. We are grateful to University of the Basque Country Press, Taylor and Francis, for their kind permission to use copyright-held material. We also thank Professor Charles Tyler for providing the ovotestis picture. The work of AA and AG has been sponsored by grants from the Norwegian Research Council and VISTA.

References

1. Tyler CR and Sumpter JP. Oocyte growth and development in teleost. Rev Fish Biol Fisheries 1996, 6:287-318
2. Selman K and Wallace RA. Cellular aspects of oocyte growth in teleosts. Zoos Sci 1989, 6:211-231
endocrine disruption in Japanese medaka (Oryzias latipes Blomachers 2002, 7:80-93).

24. Noguchi T, Ishizaki M, Volld A and Braunbeck T Temperature-dependent vitellogenin-mRNA expression in primary cultures of rainbow trout (Oncorhynchus mykiss) hepatocytes at 14 and 18°C. Toxical in vitro 2000, 14:531-540.

25. Silversand C and Haux C Fatty acid composition of vitellogenin from four teleost species. J Comp Physiol 1995, 164:593-599.

26. Wallace RA Vitellogenesis and oocyte growth in non-mammalian vertebrates. In: Developmental Biology (Edited by: Brower LW) Plenum Press, New York 1985, 1:127-177.

27. Wallace RA and Selman K Ultrastructural aspects of oogenesis and oocyte growth in fish and amphibians. J Electron Microsc Tech 1990, 16:175-201.

28. Specker JL and Sullivan CV Vitellogenesis in fishes: status and perspectives. In: Perspectives in Comparative Endocrinology (Edited by: Davey KG, Peter RE, Toye SS) Ottawa, National Research Council of Canada 1992, 303-318.

29. Le Menn F, Daveli B, Perellero C, Ndiaye P, Bon E, Perazzolo L and Rodriguez JN A new approach to fish vitellogenesis. In Proceedings of the 6th International Symposium on the Reproductive Physiology of Fish: 1999 July 4-9; Bergen (Edited by: Norberg B, Kjesbu OS, Taranger GL, Andersson E, Stefansson SO) John Grieg A/S, Bergen, Norway 2000, 211-222.

30. Peter RE and Yu KL Neuroendocrine regulation of ovulation in fishes: basic and applied aspects. Rev Fish Biol Fisher 1997, 7:173-197.

31. Swanson P Salmon gonadotropins: reconciling old and new data. In: Proceedings of the 4th International Symposium on Reproductive Physiology of Fish: 1991; Sheffield (Edited by: Scott AP, Sumpter JP, Kime DE, Raffi MS) Sheffield, UK 1991, 2-7.

32. Young G, Kagawa H and Nagahama Y Oocyte maturation in the amago salmon (Oncorhynchus rhodurus): In vitro effects of salmon gonadotropin, steroids and cyanoketone (an inhibitor of 3-hydroxy-5α-stereoid dehydrogenase). J Exp Zool 1982, 224:265-275.

33. Marte CL and Lam TJ Hormonal changes accompanying sexual maturation in captive milkfish (Chanos chanos Forsskål). Fish Physiol Biochem 1992, 10:267-275.

34. So YP, Idler DR, Truscott B and Walsh JM Progesterone, androgens and their glucuronides in the terminal stages of oocyte maturation in landlocked Atlantic salmon. J Steroid Biochem 1985, 23:583-591.

35. Kagawa H, Young G, Adachi S and Nagahama Y Estradiol-17β production in amago salmon (Oncorhynchus rhodurus) ovarian follicles: role of the thecal and granulosa cells. Gen Comp Endocrinol 1982, 47:440-448.

36. Tata JR and Smith DF Vitellogenesis: a versatile model for hormonal regulation of gene expression. Recent Prog Horm Res 1979, 35:479-90.

37. Oppen-Berntsen DO Oogenesis and hatching in teleostean fishes with special reference to breamshell proteins. Dr. Scient. Thesis, University of Bergen, Norway 1990.

38. Hyllner SJ, Oppen-Berntsen DO, Helvik JV, Walther BT and Haux C Estradiol-17β induces the major vitellogenic envelope proteins in both sexes in teleosts. J Endocrinol 1991, 131:229-236.

39. Oppen-Berntsen DO, Gram-Jensen E and Walther BT Zona radiata proteins are synthesized by rainbow trout (Oncorhynchus mykiss) hepatocytes in response to estradiol-17β. J Endocrinol 1992, 135:293-302.

40. Oppen-Berntsen DO, Hyllner SJ, Haux C, Helvik JV and Walther BT Eggshell zona radiata proteins from cod (Gadus morhua) extra-ovarian origin and induction by estradiol-17β. Int J Dev Biol 1992, 36:247-254.

41. Wiegand MD Composition, accumulation and utilization of yolk lipids in teleost fish. Rev Fish Biol Fish 1996, 6:259-286.

42. Kwon YJ, Prat R, Caudal C and Tyler CR Molecular characterization of putative yolk processing enzymes and their expression during oogenesis and embryogenesis in rainbow trout (Oncorhynchus mykiss). Biol Reprod 2001, 65:1701-1709.

43. Sire MF, Babin P and Vernier JM Involvement of the lysosomal system in yolk protein deposit and degradation during vitellogenesis and embryonic development in trout. J Exp Zool 1994, 269:69-83.

44. Mommsen PT and Walsh PJ Vitellogenesis and oocyte assembly. In: Fish Physiology (Edited by: Hoar WS, Randall DJ, Donaldson EM) Academic Press, New York 1988, 11A:347-406.

45. Norberg B Vitellogenesis in salmonid fish. Fin. Dr. Thesis, Department of Zoophysiology, University of Goteborg, Sweden 1989.

46. Islergen M, Yuan H, Volli A and Braunbeck T Measurement of vitellogenin gene expression by RT-PCR as a tool to identify

47. "Comparative Hepatology 2003, 2 http://www.comparative-hepatology.com/content/2/1/4" (page number not for citation purposes)
48. Oppen-Berntsen DO, Arakwe A, Yadetie F, Lorenz JB and Male R. Salmon eggshell protein expression: A marker for environmental effects. Mol Biotechnol 1999; 1253-260

49. Murata K, Sasaki T, Yasumasa S, Iuchi I, Enami J, Yasumasa I and Yamagami K. Cloning of cDNAs for the precursor protein of a low-molecular-weight subunit of the inner layer of the egg envelope (chorion) of the fish Orizias latipes. Dev Biol 1995; 167:19-77

50. Murata K, Sugiyama H, Yasumasa S, Iuchi I, Yasumasa I and Yamagami K. Cloning of cDNA and estrogen-induced hepatic gene expression for choriogenin H, a precursor protein of the fish egg envelope. Proc Natl Acad Sci USA 1997; 94:2050-2055

51. Lee C, Na J, Lee K and Park K. Choriogenin mRNA induction in male medaka, Orizyas latipes as a biomarker of endocrine disruption. Aquat Toxicol 2002, 61:233-241

52. Lyons CE, Payette KL, Price JL and Huang RCC. Expression and localization of maternal zona-pellucida gene. J Biol Chem 1993; 268:1351-23138

53. Del-Giacco L, Vanoni C, Bonsigno A, Citelli F. Identification and spatial distribution of the mRNA encoding the gp49 component of the vitelline envelope of Morone americana. Mol Reprod Dev 1996; 45:256-267

54. Chang YS, Wang SC, Tsao CC and Huang FL. Molecular cloning, structural analysis, and expression of carp ZP3 gene. Mol Reprod Dev 1996, 44:295-304

55. Chang YS, Hsu CC, Wang SC, Tsao CC and Huang FL. Molecular cloning, structural analysis, and expression of carp ZP2 gene. Mol Reprod Dev 1997, 46:258-267

56. Begovac PC and Wallace RA. Major vitelline envelope proteins in pipefish oocytes originate within the follicle and are associated with the zona pellucida of Matronus auratus. J Exp Zool 1999, 285:156-73

57. Celius T, Matthews JB, Giesy JP and Zacharewski TR. Structural and functional aspects of fish oviducal gland tissue binding of androgens related to plasma steroid levels in the brown trout Salmo trutta L. Gen Comp Endocrinol 1988, 70:334-344

58. Lidley CW and Thomas P. Partial characterization of a sex-steroid binding protein in the spotted sea trout (Cynoscion nebulosus). Bull Reprod 1994, 51:982-992

59. Övrevik J, Steensen J, Nielsen K and Tolflettes K-E. Partial characterization of a sex-steroid-binding protein in plasma from Svalbard Charr (Sal v elinus alpinus L.). Gen Comp Endocrinol 2001, 121:31-39

60. Trenbley KB. Interaction of estrogen mimics singly and in combination with plasma sex-steroid-binding proteins in rainbow trout (Oncorhynchus mykiss). Aquat Toxicol 2002, 56:215-225

61. Mayer PH, Stanczyk FZ, Namkung P, Fritz MA and Noyv MJ. Direct effect of sex-steroid-binding protein (SBP) of plasma on the metabolic clearance rate of testosterone in the rhesus macaque. J Steroid Biochem 1985, 22:739-46

62. Rosner W. The functions of corticosteroid-binding globulin and sex hormone-binding globulin: recent advances. Endocrine 1990, 5:81-90

63. Fortunati N. Sex hormone-binding globulin: Not only a transport protein. What news is around the corner? J Endocrinol Invest 1999, 22:223-233

64. Katzenellenbogen BS. Estrogen receptors: biochemicals and interactions with cell signaling pathways. Biol Reprod 1996, 54:287-293

65. Kuntz MA and Shapiro DJ. Dimerization of the estrogen receptor DNA-binding domain enhances binding to estrogen response elements. J Biol Chem 1997, 272:2799-56

66. Tamrasi A, Carlson KE, Daniels JR, Hurlt KM and Katzenellenbogen JA. Estrogen receptor dimerization: Ligand binding regulates dimer affinity and dimer dissociation rate. Mol Endocrinol 2002, 16:2706-2719

67. Ruh TS, Ruuf MF and Singh RK. Nuclear receptor sites: interaction with estrogen versus antiestrogen receptor complexes. In: Steroid receptors in health and diseases (Edited by: Moudgil VK) New York, Plenum Press 1992, 540-553.

68. Lewis JA, Clemens MJ and Tata JR. Morphological and biochemical changes in the hepatic endoplasmic reticulum and golgi apparatus of male Xenopus laevis after induction of egg-yolk protein synthesis by oestriadiol-17β. Mol Cell Endocrinol 1976, 11:311-329

69. Wiegand MD. Vitellogenesis in fishes. In: Reproductive Physiology of Fish (Edited by: Richter CJ). Goos HJT. Pudoc, Wageningen, The Netherlands 1982, 136-146

70. Bjornsson BTh, Hauk, C, Forlin L and Deftos LJ. The involvement of calcitonin in the reproductive physiology of the rainbow trout. J Endocrinol Invest 1968, 10:87-123

71. Opresko LK and Wiley HS. Receptor-mediated endocytosis in Xenopus oocytes. I. Characterization of the vitellogenin receptor system. J Biol Chem 1987, 262:409-415

72. Opresko LK and Wiley HS. Receptor-mediated endocytosis in Xenopus oocytes. II. Evidence for two novel mechanisms of hormonal regulation. J Biol Chem 1987, 262:4116-4123

73. Chan SL, Tan CH, Pang MK and Lam TJ. Vitellogenin purification and development of assay for vitellogenin receptor in oocyte membranes of the tilapia (Oreochromis niloticus Linnaeus 1766). Exp Zool 1991, 205:96-109

74. Tyler CR and Lancaster PA. Isolation and characterization of the receptor for vitellogenin from follicles of the rainbow trout (Oncorhynchus mykiss). J Cell Physiol 1993, 153:225-233

75. Tao Y, Bertinsky DL and Sullivan CV. Characterization of a vitellogenin receptor in white perch (Morone americana). Biol Reprod 1996, 55:646-656

76. Prat F, Coward K, Sumpter JP and Tyler CR. Molecular characterization and expression of two ovarian lipoprotein receptors
in the rainbow trout Oncorhynchus mykiss. Biol Reprod 1998, 58:1146-1153

94. Bonzaitis LM, Coward K, Davail B, Normand E, Tyler CR, Paskel F, Schneider WJ and Le Menn F Expression and localization of messenger ribonucleic acid for the vitellogenin receptor in ovarian follicles throughout oogenesis in the rainbow trout Oncorhynchus mykiss. Biol Reprod 1999, 60:1057-1066

95. Hiramatsu N, Hara A, Hiramatsu K, Fukuda H, Weber GM, Denslow ND and Sullivan CV Vitellogenin-derived yolk proteins of white perch Morone americana: Purification characterization and vitellogenin-receptor binding1. Biol Reprod 2002, 67:655-667

96. Hiramatsu N, Ichikawa N, Fukuda H, Fujita T, Sullivan CV and Hara A Identification and characterization of proteases involved in specific proteolysis of vitellogenin and yolk proteins in salmonids. J Exp Zool 2002, 292:11-25

97. Colborn T and Clement C Chemically-induced alterations in sexual and functional development: The wildlife/human connection. Advances in modern environmental toxicology Vol XXI. Princeton NJ, Princeton Scientific Publishing 1992.

98. Arukwe A and Gokseyr A Xenobiotics, xenoestrogens and reproduction disturbances in fish. Sarsia 1998, 83:225-241

99. Gokseyr A, Arukwe A, Larsson J, Cajardin MP, Hauser L, Nilsen BM, Lowe D and Matthiessen P Links between the cellular and molecular response to pollution and the impact on reproduction and fecundity including the influence of endocrine disrupters. In: Impacts of Marine Xenobiotics on European Commercial Fish-Molecular Effects and Population Responses (Edited by: Lawrence A) London, Caldwell Publishing 2003.

100. Danstra T, Page SW, Herrman JL and Meredith T Persistent organic pollutants: potential health effects? J Epidemiol Comm Health 2002, 56:824-825

101. Toppari J, Larsen JC, Christiansen P, Giwerman A, Grandjean P, Guilliette LJ Jr, Jegou B, Jansen TK, Jouannot P, Keiding N, Leffers H, McLachlan JA, Meyer O, Möller J, Raipert-De Meys E, Scheike T, Sharpe R, Sumpter J and Skakkebæk NE Male reproductive health and environmental xenoestrogens. Environ Health Perspect 1998, 106:1741-803

102. Danstra T, Barlow S, Bergman A, Kavlock R and Van Der Kraak G Global Assessment of the State-of-the-Science of Endocrine Disruptors. WHO/PCS/EDEC/02-2 2002.

103. Safe SH Environmental and dietary estrogens and human health: Is there a problem? Environ Health Perspect 1995, 103:346-315

104. Safe S and Krishnan V Cellular and molecular biology of aryl hydrocarbon (Ah) receptor-mediated gene expression. Arch Toxicol 1995, 17(Suppl)99-115

105. Alborg UG, Liversorge C, Stuer-Lauridsen R, Titus-Ernstoff L, Hsieh C-C, Hanberg A, Baron J, Trichopoulos D and Adami HO Organochlorine compounds in relation to breast cancer endometrial cancer and endometriosis: An assessment of the biological and epidemiological evidence. Crit Rev Toxicol 1995, 25:463-531

106. Kune DE The effects of pollution on reproduction in fish. Rev Fish Biol Fisher 1995, 5:52-96

107. Lam TJ Environmental influences on gonadal activity in fish. In: Fish Physiology Vol IX – Reproduction Part B (Edited by: Hoar WS, Randall DJ, Donaldson EM) New York, Academic Press 1983, 65-116

108. Susan L Effects of contaminants on teleost reproduction: past and ongoing studies. Washington, NOAA Technical Memorandum NOS OMA 29 1986.

109. Wester PW and Canton JH Histopathological study of Oryzias latipes (Medaka) after long-term [β-hexachlorocyclohexane (Lindane)] exposure. J Toxicol Environ Health 1981, 6:21-45

110. Gimeno S, Gerritsen A, Bowmer T and Komen H Molecular response to pollution and histopathological effects in juvenile Common carp (Cyprinus carpio). Aquat Toxicol 2000, 51:69-78

111. Janssen PAH, Dalesi DLWM, Lambert [JG, Vethaak AD, van Wezel AP and Goos H]Th Oestrogenic effects in the flounder Platichthys flesus after exposure to polluted harbour sediment [abstract]. 7th SETAC – Europe Annual Meeting; 1997 April 6–10, Amsterdam, The Netherlands.

112. Sumpter JP and Jobling S Vitellogenesis as a biomarker for estrogenic contaminants of the aquatic environment. Environ Health Perspect 1995, 103:173-178

113. Pelissero C, Fournier G, Génetier JL, Benneau B, Dunogues J, Gac FL and Sumpter JP Vitellogenin synthesis in cultured hepatocytes; an in vitro test for the estrogenic potency of chemicals. J Steroid Biochem Mol Biol 1993, 44:263-272

114. Jobling S and Sumpter JP Detergent components in sewage effluents are weakly estrogenic to fish: An in vitro study using rainbow trout (Oncorhynchus mykiss) hepatocytes. Aquat Toxicol 1993, 27:361-372

115. Ceulis T, Haugen TB, Grotmol T and Walther BT A sensitive zonagenic assay for rapid in vitro assessment of estrogenic potency of xenobiotics and mycotoxins. Environ Health Perspect 1999, 107:63-68

116. Tollefsen KE, Mathiesen R and Scenersen J Estrogen mimics bind with similar affinity and specificity to the hepatic estrogen receptor in Atlantic salmon (Salmo salar) and rainbow trout (Oncorhynchus mykiss). Gen Comp Endocrinol 2002, 126:4-21

117. Arukwe A and Male R Effects of 4-nonylphenol on gene expression of pituitary hormones in juvenile Atlantic salmon (Salmo salar). Aquat Toxicol 2002, 58:11-29

118. Arukwe A and Male R Effects of 4-nonylphenol on gene expression of pituitary hormones in juvenile Atlantic salmon. Dr. Scient. Thesis, University of Bergen, Norway 2001.

119. OECD 2nd Expert Consultation on Endocrine Disrupter Testing in Fish Tokyo, Japan 2000.

120. Nilsson BM, Eineden JE, Kristiansen SI, Nilsen MV, Berg K and Gokseyr A Development of quantitative vitellogenin-ella assays for fish test species used in endocrine disrupter screening [abstract NORDTOX 2001]. Pharmacol Toxicol 2001, 88(suppl I):P27

121. Okumura H, Aara A, Saeki F, Todo T, Shini A and Yamauchi K Development of a sensitive sandwich enzyme-linked
immunosorbent assay (ELISA) for vitellogenin in the Japanese eel Anguilla japonica Fish Sci 1995, 61:283-289

135. Tyler CR, Van der Eerden B, Sumpter JP, Jobling S and Panter G Measurement of vitellogenin as a biomarker for exposure to oestrogens in a wide variety of cyprinids. J Comp Physiol 1996, 166:418-426

136. Tyler CR, van Aerele R, Nilsen MV, Blackwell R, Maddix S, Nilsen BM, Berg K, Hutchinson TM and Goksøyr A Monoclonal antibody enzyme-linked immunosorbent assay to quantify vitellogenin for studies on environmental estrogens in the rainbow trout (Oncorhynchus mykiss). Environ Toxicol Chem 2002, 21:47-54

137. Brion F, Nilsen BM, Eidem JK, Goksøyr A and Porcher JM Development and validation of an enzyme-linked immunosorbent assay to measure vitellogenin in the zebrafish (Danio rerio). Environ Toxicol Chem 2002, 21:1699-1708

138. Benfey TJ, Donaldson EM and Owen TG An homologous radioimmunoassay for coho salmon (Oncorhynchus kisutch) vitellogenin with general applicability to other pacific salmonids. Gen Comp Endocrinol 1989, 75:78-82

139. Nilsen BM, Berg K, Arukwe A and Goksøyr A Monoclonal and polyclonal antibodies against fish vitellogenin for use in pollution monitoring, Mar Environ Res 1998, 46:153-157

140. Nilsen BM, Van Haux C Induction, isolation, and characterization of the lipid content of plasma vitellogenin from two salmon species: rainbow trout (Salmo gairdneri) and sea trout (Salmo trutta). Comp Biochem Physiol 1985, 81A:869-876

141. Sumpter JP The purification, radioimmunoassay and plasma levels of vitellogens in rainbow trout Salmo gairdneri. In Proceedings of the IXth International Symposium on Comparative Endocrinology: 1981 December 7–11; Hong Kong (Edited by: Lofts B) University Press 1985, 355-357

142. Crowther JR The ELISA Guidebook. Methods in Molecular Biology Vol 149, New Jersey, Humana Press 2001

143. Lowry OH, Rosebrough NJ, Farr AL and Randall RJ Protein measurements with Folin phenol reagent. J Biol Chem 1951, 193:265-275

144. Bradford MM A rapid and sensitive method for the quantitation of microgram quantities of protein utilizing the principle of protein-dye binding. Anal Biochem 1976, 72:248-254

145. Berg K, Bringsvør K, Nilsen MV, Walther BT, Goksøyr A and Nilsen BM Monoclonal antibodies against zona radiata proteins for purification of individual Zp-monomers and development of a quantitative Zp-ELISA [abstract PRIMO 11]. Mar Environ Res 2002, 54:745

146. Oppen-Berntsen DO, Helvik JV and Walther BT Major structural proteins of cod (Gadus morhua) eggshells and protein crosslinking during teleost egg hardening. Dev Biol 1990, 137:258-65

147. Berg K, Nilsen MV, Walther BT, Goksøyr A and Nilsen BM Monoclonal antibodies against zona radiata proteins for detection of estrogenic effects in fish [abstract NORTODX 2001]. Phar Sci Total Environ 2000, 263:828-837

148. Islinger M, Pawlowski S, Hollert H, Vollk A and Braunbeck T Measurement of vitellogenin-mRNA expression in primary cultures of rainbow trout hepatocytes in a non-radioactive dot blot/RNAase protection assay. Sci Total Environ 1999, 233:109-122

149. Arukwe A, Nilsen BM, Berg K and Goksøyr A Immunohistochemical analysis of the vitellogenin response in the liver of Atlantic salmon exposed to environmental estrogens. Biomarkers 1999, 4:373-380

150. Bieberstein U and Braunbeck T Immunohistochemical localization of vitellogenin in rainbow trout (Oncorhynchus mykiss) hepatocytes using immunofluorescence. Sci Total Environ 1999, 233:67-75

151. Anderson MJ, Miller MR and Hinton DE In vitro modulation of 17β-estradiol-induced vitellogenin synthesis: Effects of cytochrome P4501A1 inducing compounds on rainbow trout (Oncorhynchus mykiss) liver cells. Aquat Toxicol 1996, 34:327-350

152. Anderson MJ, Olsen H, Matsumura F and Hinton DE In vivo modulation of 17β-estradiol-induced vitellogenin synthesis and estrogen receptor in rainbow trout (Oncorhynchus mykiss) liver cells by [(−)-naphthoflavone. Toxicol Appl Pharmacol 1996, 137:210-218

153. Villalobos SA, Anderson MJ, Denison MS, Hinton DE, Tullis K, Kennedy IM, Jones AD, Chang DPF, Yang G and Kelley P Diosynthlike properties of a trichloroethylene combustion-generated aerosol. Environ Health Perspect 1996, 104:734-743

154. Giertych JF, Lincoln DW, Massip SJ, Dickerman HW, Bradlow HL, Niwa T and Swaneck GE Enhancement of 2- and 16α-estradiol hydroxylation in MCF-7 human breast cancer cells by 2,3,7,8-tetrachlorodibenzo-p-dioxin. Biochem Biophys Res Commun 1988, 157:515-520

155. Slupsky DC, Lincoln DW, Dickerman HW and Giertych JF 2,3,7,8-Tetrachlorodibenzo-p-dioxin causes an extensive alteration of 17β-estradiol metabolism in MCF-7 breast tumor cells. Proc Natl Acad Sci USA 1990, 87:6917-6921

156. Safe S, Astroff B, Harris M, Zacharewski T, Dickerson R, Romkes M and Siegel I 2,3,7,8-Tetrachlorodibenzo-p-dioxin (TCDD) and related compounds as antiestrogens: Characterization and mechanism of action. Pharmacol Toxicol 1991, 69:400-409

157. Donaldson EM Reproductive indices as measures of the effects of environmental stressors in fish. Amer Fishers Soc 1990, 8109-20

158. Ware DM Bioenergetics of stock and recruitment. Can J Fish Aquat Sci 1980, 37:1012-1024

159. Ware DM Power and evolutionary fitness of teleosts. Can J Fish Aquat Sci 1982, 39:3-15

160. Sibly R and Calow P An integrated approach to life-cycle evolution using selective landscapes. J Theoret Biol 1983, 102:527-547

161. Thorpe JE Reproductive strategies in Atlantic salmon Salmo salar. Aquat Fish Manag 1994, 25:77-87

162. Polliccansky D Size, age and demography of metamorphosis and sexual maturation in fishes. Amer Zool 1983, 23:57-63

163. Herman RL and Kincaid HL Pathological effects of orally administered estradiol to rainbow trout. Aquaculture 1988, 72:165-172

164. Bortone SA and Davis WP Fish intersexuality as indicator of environmental stress. Biosci 1994, 44:65-172

165. Allen MS, Miranda LE and Brock RE Implications of compensatory and additive mortality to the management of selected sportfish populations Lakes. Reserv Res Manage 1996, 3:67-79

166. Jobling S, Nolan M, Tyler CR, Brighty G and Sumpter JP Widespread sexual disruption in wild fish. Environ Sci Technol 1998, 32:2498-2506

167. Byrne M, Villinski JT, Cisternas P, Siegel RK, Popodi E and Raff RA Maternal factors and the evolution of developmental mode: evolution of oogenesis in Helicodrias erythrogramma. Dev Genes Evol 1999, 209:275-281

168. Davis RA Evolution of processes and regulators of lipoprotein synthesis: from birds to mammals. J Nutr 1997, 127:795S-800S

169. Mouchel N, Tricet V, Naimy BY, Le Penne JP and Wolff J Structure of a fish (Oncorhynchus mykiss) vitellogenin gene and its evolution and regulatory implication. Gene 1999, 197:147-52

170. Tricet V, Buisine N, Mouchel N, Moran P, Pendas AM, Le Penne JP and Wolff J Genomic analysis of the vitellogenin locus in rainbow trout (Oncorhynchus mykiss) reveals a complex history of gene amplification and retroposon activity. Mol Gen Genet 2000, 263:828-837

171. Gerber-Huber S, Nardelli D, Haefliger J, Cooper DN, Givel F, Ger mond JE, Engel J, Green NM and Wahl W Precursor-product relationship between vitellogenin and the yolk proteins as derived from the complete sequence of a Xenopus vitellogenin gene. Nucleic Acids Res 1987, 15:4737-4760

172. van het Schip FD, Samalo J, Brooks J, Ophuis J, Mojet M, Gruber M and Ab G Nucleotide sequence of a chicken vitellogenin gene and derived amino acid sequence of the encoded yolk precursor protein. J Mol Biol 1987, 196:245-260

173. Kiesth J, Nettleton M, Zöckler G, Rosebrough NJ, Lehmann T and Le Menn F Evolution of oogenesis: the receptor for vitellogenin from the rainbow trout. J Lipid Res 1998, 39:1929-1937

174. Nardelli D, Gerber-Huber S, van het Schip FD, Gruber M, Ab G and Wahl W Vertebrate and nematode genes coding for yolk
proteins are derived from a common ancestor. Biochem 1987, 26:6397-6402
178. Nardelli D, van het Schip FD, Gerber-Huber S, Haefliger J-A, Gruber M, Ab G and Wahl W Comparison of the organization and fine structure of a chicken and a Xenopus laevis vitellogenin gene. J Biol Chem 1987, 262:15377-15385
179. Rogers J Exon shuffling and intron insertion in serine proteases genes. Nature 1985, 315:458-459
180. Cavalier-Smith T Intron phylogeny: a new hypothesis. Trends Genet 1991, 7:145-148
181. Yanagimachi R Fertility of mammalian spermatozoa: its development and relativity. Zygote 1994, 2:371-372
182. Doreen S, Landsberger N, Dwyer N, Gold L, Blanchette-Mackie J and Dean J Incorporation of mouse zona pellucida proteins into the envelope of Xenopus laevis oocytes. Dev Genes Evol 1999, 209:330-339
183. Walther BT Do life’s three domains mirror the origin of sex? J Biosci 2000, 25:217-220
184. Goksør J Evolution of eukaryotic cells. Nature 1967, 214:461
185. Sagan L On the origin of mitosing cells. J Theor Biol 1967, 14:225-274
186. Margulis L and Bermudes D Symbiosis as a mechanism of evolution: Status of the endosymbiosis theory. Symbiosis 1985, 1:101-124