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Emergence of ovarian 11-deoxycorticosteroid biosynthesis at ovulation time in the sea bass, Dicentrarchus labrax L.

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Summary. Steroidogenesis in the ovary of the teleost Dicentrarchus labrax has been studied in vitro at five different stages: 1) previtellogenic; 2) postvitellogenic; 3) postovulatory, one day after HCG-induced spawning; 4) postovulatory, 9 days after induced spawning; 5) atretic. Ovarian tissues were incubated with [4-14C] pregnenolone and [4-14C] progesterone for 6 hr at 15-16°C. The following metabolites were formed after ovulation: progesterone (from pregnenolone), 11-deoxycorticosterone, 11-deoxycortisol, 17α-hydroxyprogesterone, androstenedione, testosterone and estradiol-17β. Percentages of conversion were high for 11-deoxycorticosteroids and androgens and low for estradiol-17β. Before ovulation, yields of androgens were high at stage 2 but negligible at stage 1. Estradiol-17β and 11-deoxycorticosterone were obtained in small amounts at maturity but could not be detected during quiescence. Atretic ovarian tissue converted pregnenolone to progesterone but it went no further in steroid hormone biosynthesis.

Introduction.

The biosynthesis of 11-deoxycorticosterone (DOC) and/or 11-deoxycortisol (S), through 21-hydroxylation of progesterone and 17α-hydroxyprogesterone, respectively (Ortci-Flores et al., 1976), has been shown to occur in the ovaries of 6 teleost species (Colombo et al., 1973; Colombo and Colombo Belvedere, 1977), two urodeles (Colombo et al., 1977a, b) and 2 reptiles (Colombo et al., 1974; Colombo and Yaron, 1976). The information available, however, was insufficient to establish whether 11-deoxycorticosteroid production was in alternative to that of estrogens by competing, for instance, for C16 intermediates, or whether 11-deoxycorticosteroids alternated with estrogens through a shift in ovarian steroidogenesis at a certain stage of the oogenic cycle.

The observation that the teleost aromatizing system was more apparent in the vitellogenic ovary of Gobius jozo than in pre- and post-ovulatory ovaries of Diplodus annularis and Solea impar, while the reverse was true for steroid 21-hydroxylase activity (Colombo and Colombo Belvedere, 1977), could reflect either a sequence of distinct steroidogenic phases or simply a species specificity in ovarian steroid patterns. To avoid
the latter complication, we decided to use the sea bass, a commercially reared species, to study in vitro the course of ovarian steroid hormone biosynthesis during the reproductive cycle.

Materials and methods.

Sixteen females of *Dicentrarchus labrax* (mean standard length: 38.0 cm ± 2.9 SD; mean body weight: 994 g ± 225 SD) were used. Mature or spawned specimens were obtained from the SIRAP Marine Fish Hatchery at Pellestrina whereas nonbreeding animals were captured in coastal lagoons near Venice. Both ovaries were dissected out following pithing of the fish. Five ovarian stages were investigated: 1) quiescent ovary with only previtellogenic oocytes obtained from females caught outside the breeding period in April; 2) mature or postvitellogenic ovary packed with yolky oocytes and some growing follicles; 3) ovary one day after spawning, induced in mature females by injecting 1000 IU/kg body weight of human chorionic gonadotropin (HCG); 4) ovary nine days after induced spawning; 5) ovary with extensive follicular atresia as found in females confined to low salinity (5-10 p.p.m.) brackish water during ovarian maturation.

Portions of ovarian tissue from animals at the same stage were pooled in ice-cold physiologic medium of the following composition: 187.3 mM NaCl, 3.34 mM KCl, 2.30 mM CaCl₂, 3.98 mM MgCl₂, 0.37 mM MgSO₄, 1.61 mM Na₂HPO₄, 0.36 mM KH₂PO₄, 20.0 mM NaHCO₃, 5.55 mM glucose, phenol red 2 mg/liter. 2-g samples of tissue were then incubated in 50-ml Erlenmeyer flasks containing 1 μCi of either [4-14C] pregnenolone (specific activity (SA): 52.8 mCi/mmmole) or [4-14C] progesterone (SA: 52.8 mCi/mmmole) previously dissolved in 200 μl of propylene glycol, plus 20 ml of medium. Incubations were carried out in a Dubnoff metabolic shaker for 6 hr at 15-16 °C. Aliquots of 2 ml were withdrawn from the incubation medium after 7.5, 15, 30, 60, 120, 240 and 360 min. and replaced by an equal volume of fresh medium without precursor. At the end of incubation, the medium was separated from the tissue. Metabolism was terminated by mixing thoroughly with 3 volumes of ethanol.

Before extraction, suitable amounts of carrier and tritiated steroids were added to samples. Metabolites were then extracted, chromatographed bidimensionally on thin-layer plates, autoradiographed, eluted and identified by their isopolarity and isomorphism with authentic steroids, as previously described (Colombo and Colombo Belvedere, 1977).

Results.

The following compounds were formed from both precursors by ovulated ovarian tissues (stages 3 and 4): progesterone (from pregnenolone), 11-deoxycorticosterone, 11-deoxy cortisol, 17α-hydroxyprogesterone, androstenedione, testosterone and estradiol-17β. The steroidogenic pattern was not greatly affected by the time interval elapsed from spawning (figs. 3 and 4), despite the fact that the involution of non-ovulated oocytes was more advanced after 9 days. The release of the above metabolites in the medium during incubation with either precursor was measured by the integrated
values of their yield vs time curves: it was high for 11-deoxycorticosteroids (S > DOC) and androgens (testosterone > androstenedione) but low for estradiol-17β.

Before ovulation, accumulation of androgens in the incubation medium was high at stage 2 but negligible at stage 1. Estradiol-17β and DOC (but not S) were obtained in small amounts at maturity but could not be detected during quiescence (figs. 1 and 2). On the other hand, the production of 17α-hydroxyprogesterone was always conspicuous from stages 1 to 4, especially with progesterone as a precursor. Metabolite yields in the incubated tissues were correlated according to the same patterns outlined for the

FIG. 1-4. — Autoradiographs of thin-layer chromatographic profiles of metabolites released in the medium after 6-hr incubation of sea bass ovarian tissue at quiescence (fig. 1), at maturity (fig. 2), one day after deposition (fig. 3) and 9 days after deposition (fig. 4) with [4-14C] progesterone. (0) chromatographic origin; (1) progesterone; (2) androstenedione; (3) 17α-hydroxyprogesterone; (4) testosterone; (5) 11-deoxycorticosterone; (6) 11-deoxycortisol; (7) estrone; (8) estradiol-17β. Dotted circles indicate the position of carrier steroids associated with very little or no [14C] label.
media. Tissues, however, showed greater yields of unidentified compounds of low and intermediate polarity.

At stage 5, atretic ovarian tissue converted pregnenolone to progesterone but was unable to perform further steps for steroid hormone biosynthesis. Radioactivity was instead channelled along other transformation routes but their products could not be characterized.

Discussion.

The present work demonstrates that the following steroid-converting enzymes may contribute to the ovarian steroid profiles of D. labrax: 3β- and 17β-hydroxy-steroid dehydrogenases, 5-ene-ketosteroid isomerase, 17α, 20-C21-desmolase, steroid 17α- and 21-hydroxylases and aromatizing system. Interestingly, the testicular tissue in the same species was devoid of the last two enzymatic activities (Colombo et al., 1978a).

The ovarian location of steroid 21-hydroxylase in the sea bass, a member of the family Serranidae, confirms similar findings in 3 other families of the order Perciformes (Colombo et al., 1973; Colombo and Colombo-Belvedere, 1977).

The changing patterns of steroid metabolism from ovarian quiescence to the post-ovulatory stage suggests that steroidogenic shifts occur in the ovary of D. labrax during the oogenic cycle making androgen intermediates more available for estrogen biosynthesis during follicular growth and causing a rise in 11-deoxycorticosterone formation around the time of ovulation.

The former change confirms the role of estrogens in the promotion of hepatic synthesis and release of vitellogenin, as documented in fish by recent reports (Plack et al., 1971; Campbell and Idler, 1976). On the other hand, the induction of steroid 21-hydroxylase activity concomitantly with spawning fits very well with the fact that in a wide spectrum of steroid compounds, DOC and S were the most potent inducers in vitro of meiotic maturation in goldfish (Jalabert et al., 1973) and in catfish (Goswami and Sundararaj, 1974). These observations cannot be reconciled, however, with the model proposed by Indian workers (Goswami et al., 1974) in which 11-deoxycorticosteroids are produced in the interrenal under gonadotropin stimulation and act on the ovary.

A last comment should be made on the extensive atresia likely to result from prolonged gonadotropin deprivation. In this case, suppression of ovarian endocrine activity seems to parallel the involution of gametic elements and the phagocytic conversion of follicular cells.

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Résumé. La stéroïdogenèse de l'ovaire du téléostéen Dicentrarchus labrax a été étudiée in vitro à 5 stades différents : 1) prévitellogenèse, 2) postvitellogenèse, 3) post-ovulation, 1 jour après la fraie induite par HCG, 4) post-ovulation, 9 jours après la fraie induite, 5)

References


