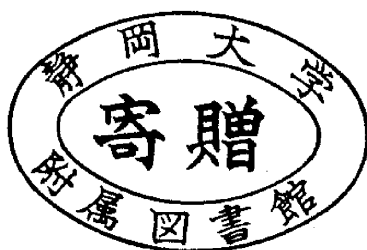


乳腺および腎臓における
グルコルチコイド代謝の比較生理学的研究

(研究課題番号 07806037)

平成7年度～平成8年度科学研究費補助金(基盤研究(C)(2))研究成果報告書



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は し が き

本報告書は平成7、8年度の2年間にわたる科学研究費補助金（基盤研究C）『乳腺および腎臓におけるグルココルチコイド代謝の比較生理学的研究』の研究成果をまとめたものである。

腎臓のようなミネラルコルチコイドの標的器官には、グルココルチコイドの11位水酸基をケトン基に変換するステロイド11 β 水酸基脱水素酵素が存在する。この酵素の存在は、逆に、ミネラルコルチコイドの標的器官であることの証明につながると考え、本研究では乳腺、腎臓、および卵管のグルココルチコイド代謝を、マウスとウズラを用いて比較検討した。

高等動物の電解質代謝および水の排泄・分泌は非常に複雑に調節されている現象であるが、本研究ではこれを分子レベルで解明することによって、少しでも理解を深めようと努力した。

本報告書が内分泌学を研究するものにとって参考になれば幸いである。

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Abstract Corticosteroid action is modulated by the NAD⁺-dependent 11 β -HSD-2, which oxidatively inactivates corticosterone, allowing the nonselective mineralocorticoid receptor to be occupied by mineralocorticoids. The present study was conducted to show that 11 β -HSD-2 is present in the reproductive tract of egg-laying birds.

The microsome fraction was prepared from homogenates of various parts of oviduct of the Japanese quail, and incubated with [³H]corticosterone for 1 hr at 39 C in the presence or absence of 1 mM NAD⁺ or NADP⁺. The reaction was stopped, and the steroid was extracted by ethyl acetate. Separation of steroids was performed using a thin-layer chromatography developed in a chloroform-methanol (95:5) system. Activity of 11 β -HSD-2 was determined by measuring the conversion of [³H]corticosterone to [³H]11-dehydrocorticosterone.

As the results, the shell gland and albumen secreting region of the oviduct showed high activity of 11 β -HSD-2. In these tissue, [³H]corticosterone was exclusively metabolized to [³H]11-dehydrocorticosterone. On the other hands, infundibulum and isthmus of the oviduct and the vagina contain low activity, indicating that these parts might not be a target of the mineralocorticoids.

Key words: 11 β -Hydroxysteroid dehydrogenase, Oviduct, Japanese quail, Corticosterone, Aldosterone

Corticosteroid action is modulated by the enzyme 11β -hydroxysteroid dehydrogenase (11β -HSD) (Edwards *et al.*, 1988; Funder *et al.*, 1988). Two isozymes of 11β -HSD are now recognized: 11β -HSD-1 is NADPH/NADP⁺-dependent enzyme with a K_m in the micromolar range (Monder and White, 1993; Funder, 1993; Seckl, 1993). 11β -HSD-2 is NAD⁺-dependent enzyme with a K_m in the nanomolar range. In the liver, 11β -HSD-1 is predominant and converts 11-dehydrocorticosterone to corticosterone (Lakshmi and Monder, 1988). On the other hand, in the kidney, the NAD⁺-dependent 11β -HSD-2 oxidatively inactivates corticosterone (Rusvai and Naray-Fejes-Tosh, 1993), allowing the nonselective mineralocorticoid receptor (Krozowski and Funder, 1983) to be occupied by mineralocorticoids (Albiston *et al.*, 1994; Krozowski *et al.*, 1994; Rusvai and Naray-Fejes-Tosh, 1993).

The present study was conducted to show that 11β -HSD-2 is present in other sodium-transporting epithelia such as those in the mammary gland of mammals and in the reproductive tract of egg-laying birds.

Materials and Methods

Animals:

Japanese quail were obtained from Tokai-Yuki at 6 wk of age. Individually caged birds were maintained under a lighting schedule of 14 h light (0500 to 1900 h) and 10 h dark

and provided *ad libitum* access to water and a commercial quail diet.

Tissue preparation:

The reproductive tract and kidney of Japanese quail were isolated immediately after decapitation, finely chopped with scissors and washed several times with an ice-cold saline. The tissue was homogenized in 5 vol of homogenizing buffer (0.25 M sucrose, 50 mM Tris-HCl and 5 mM magnesium chloride, pH 7.4) with a Polytron-type homogenizer for 10 sec. The microsome fraction was prepared from homogenates by centrifugation at 10,000 x g for 20 min at 4 C, followed by a high speed centrifugation of the supernatant at 105,000 x g for 60 min at 4 C. Microsomes were resuspended in the homogenizing buffer.

Enzymatic analysis:

11 β -HSD-2 activity in microsome fraction was determined by measuring the conversion of [³H]corticosterone to [³H]11-dehydrocorticosterone. [1, 2, 6, 7-³H]Corticosterone (spec. act., 3.22 TBq/mmol) was purchased from Amersham International plc, England. About 12.3 kBq of ³H-labeled corticosterone dissolved in ethanol was transferred into an incubation tube. After addition of 50 μ l of propylene glycol, the solvent was evaporated under reduced pressure immediately before incubation. An appropriate amount of microsome fraction, whose protein content had been previously measured, was added to each tube. NAD⁺ and NADP⁺ was dissolved in the homogenizing

buffer and added to the tube so that the final concentration of the cofactor was 1 mM. Then the final volume of the incubation mixture was adjusted to 0.5 ml per tube by the homogenizing buffer. The above mixture was incubated at 39 C in a Dubnoff-type incubator under aerobic condition. For control experiments the mixture excluding the microsome fraction was incubated simultaneously under the identical condition.

Separation of the metabolites:

At the end of incubation, 1 ml of ethyl acetate was added to the incubation mixture to terminate further enzymatic reaction. The ethyl acetate layer was separated and reserved as an extract. The residual aqueous layer was extracted five-times more to complete extraction of the steroids. The extracts were pooled and an aliquot of the extract was spotted on a thin layer of silica gel GF (E. Merck, Darmstadt, Germany), together with corticosterone and 11-dehydrocorticosterone as markers. After the development with chloroform: methanol (95:5) system, the markers on the chromatogram were detected under an ultraviolet lamp (wavelength, 253 nm). Radioactive areas were detected on another chromatogram by the fluorographic technique. The radioactive area were scraped off from the plates, and the steroids were eluted from the absorbent with chloroform: ethanol (1:1) mixture.

Quantitation of the metabolites, protein and enzyme activity:

Radioactivities were measured by a liquid scintillation spectrometer, whose counting efficiency of tritium was about 50 %. Protein contents in microsome fraction was measured by the copper-Folin method. The enzyme activities were tentatively expressed as amount in pmole of 11-dehydrocorticosterone produced for 1 hr per mg of microsome fraction.

Results and Discussion

As shown in Fig. 1, 11-dehydrocorticosterone was obtained as the major metabolite of corticosterone after the incubation with microsome fraction of the kidney. No metabolites were detected in the absence of cofactor. Corticosterone was also metabolized to 11-dehydrocorticosterone by the microsome fraction of the whole oviduct.

The regional distribution of the enzyme was shown in Fig.2. Highest activity was observed in the shell gland in the incubation with NAD as the cofactor. The albumen secreting region of the oviduct also showed high activity of 11β -HSD-2. On the other hands, infundibulum and isthmus of the oviduct and the vagina contain low activity, indicating that these parts might not be a target of the mineralocorticoids.

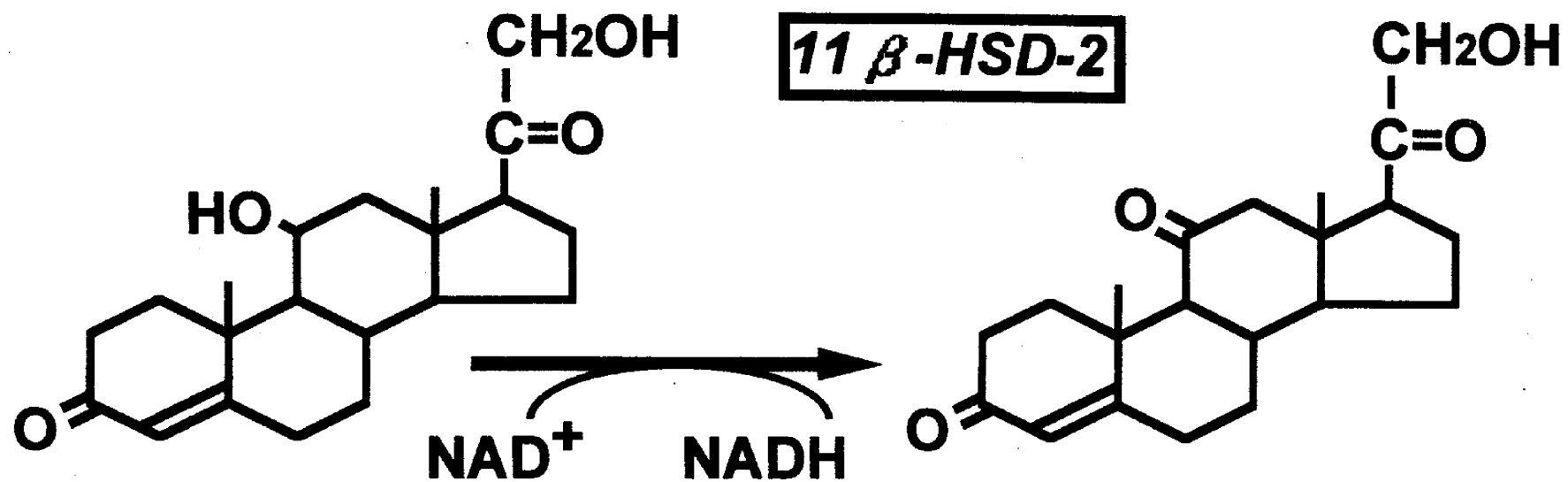
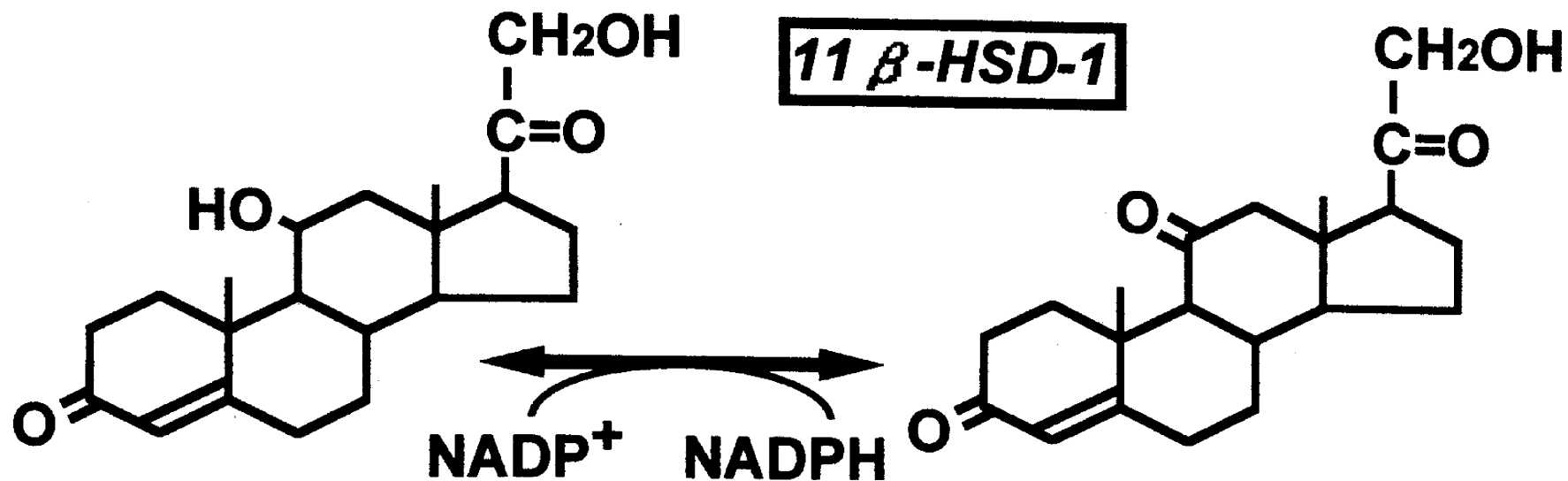
The 11β -HSD-2 enzyme converts glucocorticoids to receptor-inactive metabolites, endowing specificity on the mineralocorticoid receptor and modulating glucocorticoid

access to the glucocorticoid receptor. The 11β -HSD-2 enzyme has previously been localized in high amounts in classical sodium-transporting epithelia, the ileum and term placenta in humans (Smith *et al.*, 1996; Krozowski *et al.*, 1995; Kyosseff *et al.*, 1996; Brown *et al.*, 1996), but studies in the rat have been limited to the detection of message (Whorwood *et al.*, 1995; Roland *et al.*, 1995).

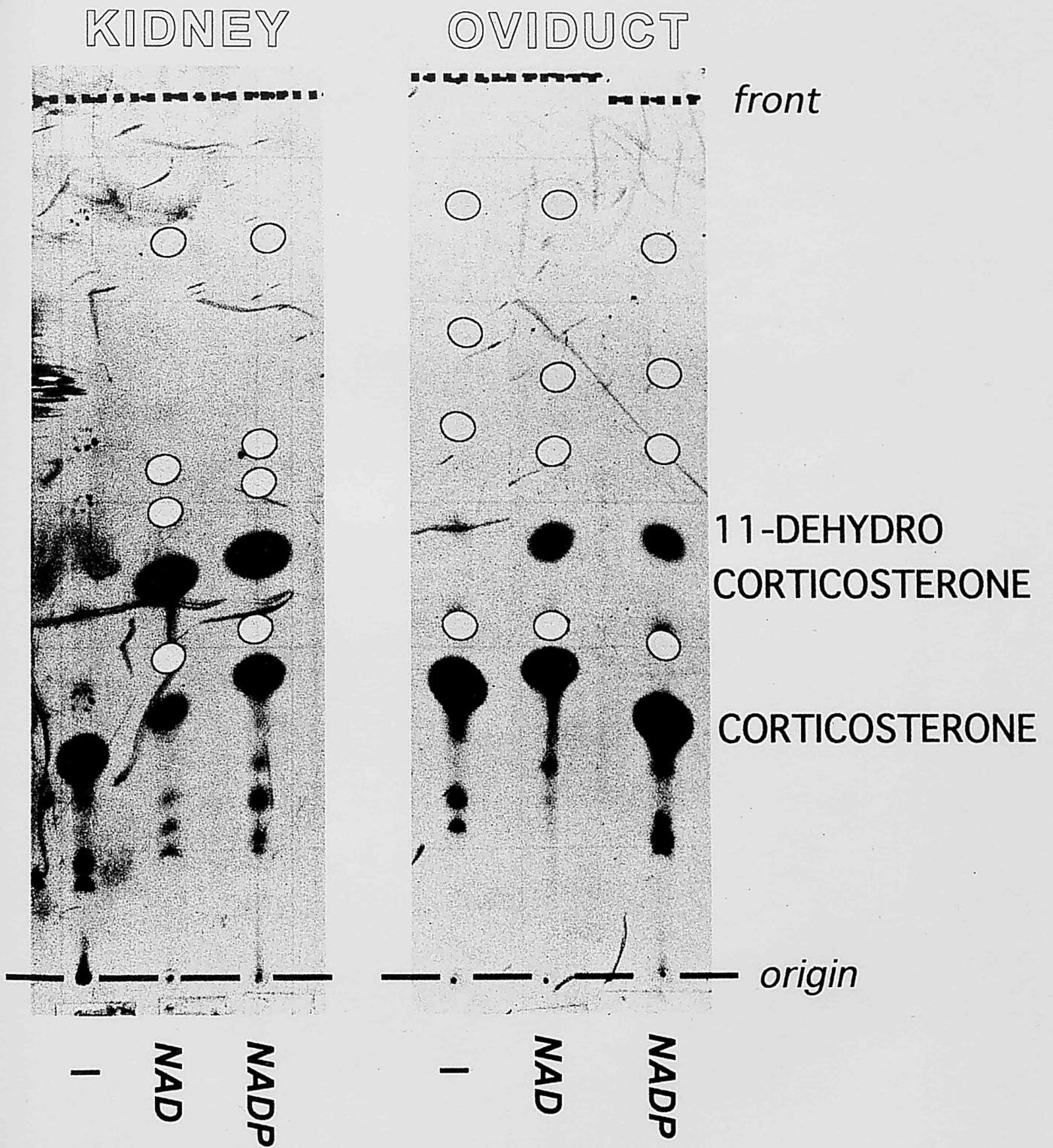
Developmental studies showed the presence of low numbers of 11β -HSD-2-positive tubules in the fetal rat kidney before birth. Studies in other species have provided evidence for fetal 11β -HSD-2 activity, with highest amounts in the kidney (Stewart *et al.*, 1994), whereas in the mouse, renal 11β -HSD-2 mRNA is present by 13 days gestation, preceding the appearance of message of mineralocorticoid receptor (Brown *et al.*, 1996).

The bile duct and gall bladder absorb water to concentrate bile fluid (Tavoloni, 1985), but this process is not known to be mediated by mineralocorticoids.

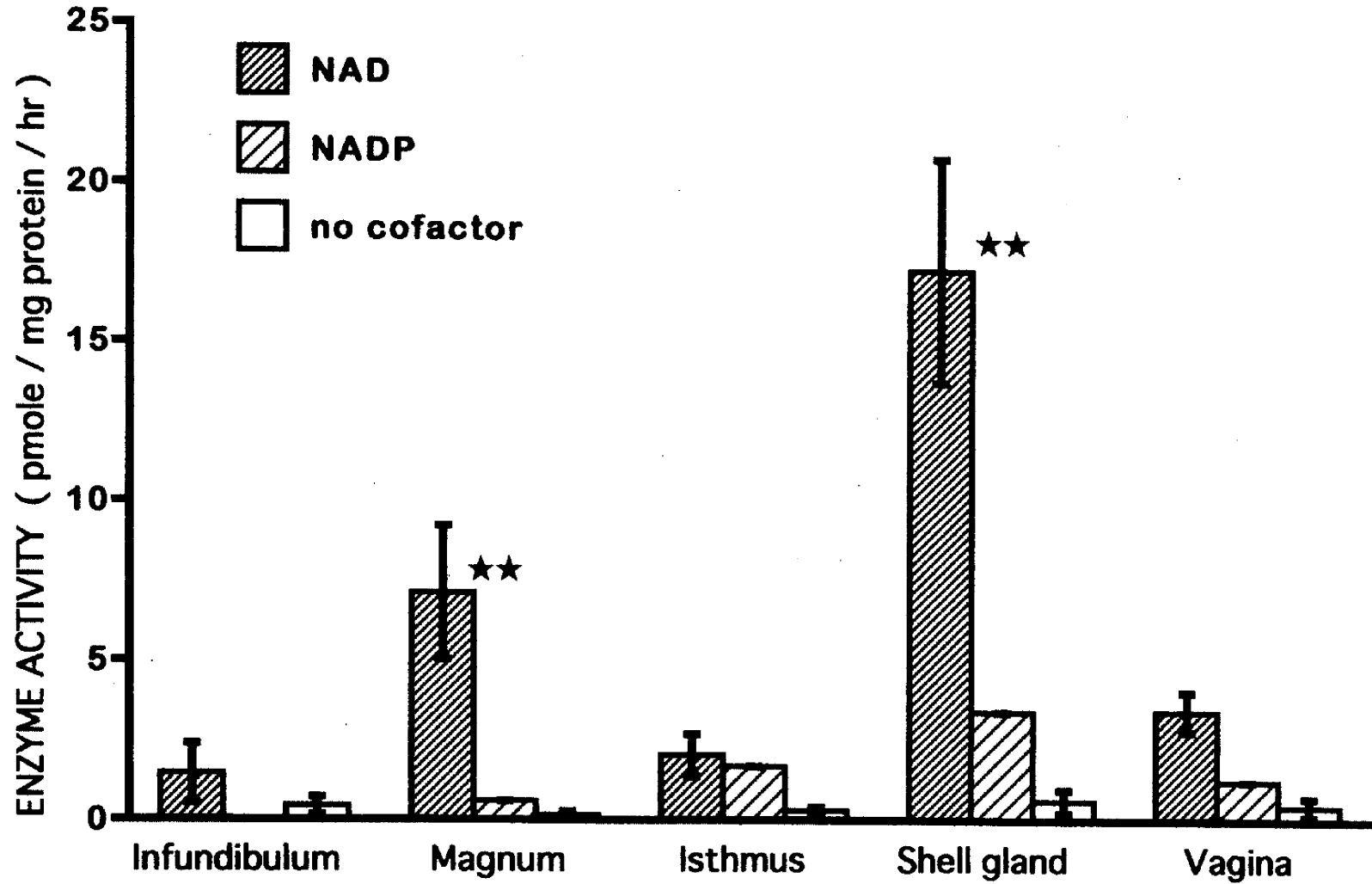
Currently, little is known about the modulation of 11β -HSD-2 activity, although recent studies suggest that it could be regulated by the protein kinase A pathway (Pasqualette *et al.*, 1996) and via corticosteroids (Li *et al.*, 1996).



Fluorography of TLC



11 β -Hydroxysteroid Dehydrogenase in Quail Oviduct



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