

# Localization of Arginine Biosynthetic Enzymes in Renal Proximal Tubules and Abundance of mRNA during Development

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**ABSTRACT.** Argininosuccinate synthetase and argininosuccinate lyase catalyze the conversion of citrulline to arginine in kidney. Immunohistochemical staining of mouse kidney sections with antibodies to these two enzymes, compared with the staining patterns of known markers for proximal tubules, demonstrated that these enzymes are localized within the proximal tubules. The relative abundance of mRNA encoding argininosuccinate synthetase and argininosuccinate lyase during fetal and postnatal development of mouse kidney was also determined. Changes in relative abundance of these mRNA in kidney are coordinate during development, paralleling the developmental profile of phosphoenolpyruvate carboxykinase mRNA, which is also expressed in proximal tubules. Although relative abundances of the mRNA are comparable in liver and kidney of adult mice, the profiles of mRNA abundance during development of these two organs are distinct. The results indicate that these enzymes and their corresponding mRNA can serve as useful markers for examining the differentiation and development of renal proximal tubules *in vivo* and in cultured explants. (*Pediatr Res* 29: 151-154, 1991)

## Abbreviations

AS, argininosuccinate synthetase  
AL, argininosuccinate lyase  
 $\gamma$ -GTP,  $\gamma$ -glutamyltranspeptidase  
TH, Tamm-Horsfall glycoprotein  
PEPCK, phosphoenolpyruvate carboxykinase

The kidney is a major site of arginine synthesis *in vivo* (1). Adequate supplies of arginine are essential for maintenance of normal growth rates after birth (2-4) and recovery from protein malnutrition (5), indicating the physiologic significance of this metabolic pathway. Conversion of citrulline to arginine in the kidney is catalyzed by AS [L-citrulline: L-aspartate lyase (AMP-forming), EC 6.3.4.5] and AL (L-argininosuccinate arginine lyase, EC 4.3.2.1) (6). These enzymes were reported previously to be localized predominantly within the renal medulla (7). However, mRNA encoding these enzymes were found recently

to be localized exclusively in the renal cortex (8). Because knowledge of the localization of these enzymes within the kidney is essential for understanding their role in renal metabolism, immunohistochemical techniques were used to establish that AS and AL are expressed in the proximal tubules of the renal cortex.

Given that AS and AL are localized in the proximal tubules, it was of interest to determine the relative abundance of the corresponding mRNA as the metanephros develops into nephrons. Changes in mRNA abundance during this period might reflect the degree of tubular cell differentiation or might also indicate responses to changes in circulating hormones or nutritional factors. Because AS and AL have half-lives of several days (9, 10), changes in relative abundance of the corresponding mRNA, which apparently have half-lives of several hours (11, 12), are more timely indicators of changes in gene expression during kidney development. Developmental profiles of relative abundance for AS and AL mRNA have recently been determined for liver (13), where these enzymes form part of the urea cycle. Our study demonstrates that AS and AL mRNA undergo coordinate increases in abundance during kidney development, but the developmental profiles for these mRNA differ for liver and kidney.

## MATERIALS AND METHODS

**Animals.** Female Swiss-Webster albino mice, made pregnant by timed matings, were supplied by Hilltop Lab Animals, Inc. (Scottsdale, PA) and Tyler Laboratories (Bellevue, WA). Experimental protocols were approved by the Animal Care and Use Committees of the University of Pittsburgh and the University of Washington.

**Isolation and analysis of RNA.** Kidneys were rapidly dissected from fetuses and neonates of the indicated ages and immediately frozen on dry ice. Kidneys for each time-point were pooled before extraction of RNA. Total kidney RNA was isolated and analyzed by Northern blot as previously described (11, 12). Relative abundance of specific mRNA was determined by liquid scintillation spectrometry of hybridized RNA bands after Northern blot analysis (11, 12). Cloned rat cDNA probes used for hybridization corresponded to AS (8), AL (14), PEPCK (15), glyceraldehyde-3-phosphate dehydrogenase (16), and  $\beta$ -tubulin (17).

**Histology and immunohistochemistry.** Fresh murine tissue was obtained from young adults (2 mo) and fetuses or neonates at gestational d 13, 15, 17, 19, 21, and 23. Tissue was fixed in 3.5% paraformaldehyde (pH 7.4) for 1 h at 4°C. Tissue was then washed, dehydrated through graded acetone, and infiltrated and embedded with Immunobed plastic embedding medium (Polysciences, Warrington, PA). For light microscopy, 3- $\mu$ m sections

Received March 14, 1990; accepted October 15, 1990.

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Supported in part by Grants DK 33144 (S.M.M.) and DK 34891 (E.D.A.) from the National Institutes of Health.

were cut on an ultramicrotome, mounted on glass slides, and stained with hematoxylin.

Immunohistologic techniques were used on serial sections of murine renal tissue to clearly localize the antigenic expression of AS and AL to specific nephron segments. Discrete tubular segments were identified by specific staining with segment-specific antibodies and lectins as previously described (18–20). Proximal tubules were identified by staining with affinity-purified antibody to the brush border enzyme  $\gamma$ -GTP and the lectin Lotus tetragonolobus. Thick ascending limb-early distal tubular segments were identified by staining with affinity-purified antibody to TH and collecting tubules were identified by staining with the lectin Dolichos biflorus. Rabbit antibodies to AS and AL were prepared as described previously (21, 22). Antibodies to  $\gamma$ -GTP and TH were kindly provided by Drs. N. P. Curthoys (Colorado State University) and John R. Hoyer (University of Pennsylvania School of Medicine), respectively, and prepared by their previously described methods (23, 24). Biotinylated lectins were obtained from Sigma Chemical Company (St. Louis, MO) and all other antisera were DAKO immunochemicals (Accurate Chemical and Scientific Corporation, Westbury, NY). A total of six to eight kidneys for each gestational stage were used in the current studies, with five to six sets of serial sections assessed per kidney.

The specific immunostaining procedure used was our previously described postembedding technique specifically developed for immunolocalization of antigens and lectins in plastic sections of fetal murine tissue (19, 20, 25, 26). Trypsinized Immunobed sections were incubated with primary antibody (anti- $\gamma$ -GTP, anti-TH at 1:100; anti-AS and -AL at 1:50) or biotinylated lectins (1:400) for 48 h at 4°C. For antibodies, this was followed by sequential incubations with bridging swine anti-rabbit IgG (1:20) and rabbit peroxidase-antiperoxidase-complex (1:100). For lectins, avidin peroxidase-antiperoxidase complex (1:600) replaced the bridging antibody step. In addition to substitution of primary antibody with preimmune rabbit serum, controls included staining after absorption of primary antibodies with purified antigens and primary biotinylated lectins with avidin as well as deletion of specific reagents at each step of the procedure. In such controls, no significant background staining was observed. Slides were photographed using Nomarski differential interference contrast.

## RESULTS AND DISCUSSION

Immunohistochemical staining of mouse kidney sections with  $\gamma$ -GTP antibody (17; Fig. 1c) and Lotus tetragonolobus lectin (Fig. 1a) confirmed that these reference markers are localized to the proximal tubules. By comparison, AS and AL are also localized in the proximal tubules (Fig. 1b and d). These results agree with a recent analysis of arginine biosynthesis rates in microdissected nephron segments from rat kidney (27) as well as the finding that AS and AL mRNA were present in renal cortex but not in medulla or papilla (8). Although AS and AL are soluble cytosolic enzymes, the immunohistochemical staining appeared more intense near the luminal face of the tubules, suggesting that uptake of citrulline from the glomerular filtrate and its subsequent conversion to arginine may be closely associated spatially. AS or AL immunohistochemical staining did not correspond to Dolichos biflorus lectin-stained collecting tubules (Fig. 1e), Tamm-Horsfall protein stained tubules (not shown), or glomeruli. The presence of AS and AL in the proximal tubules—rather than in the renal medulla, as originally reported (7)—is reasonable in that resorption of citrulline from the glomerular filtrate and its conversion to arginine for release into the blood are colocalized.

At 13 d of gestation, no immunoreactivity for AS or AL could be demonstrated in developing tubular elements distributed throughout the metanephric mesenchyme. This was in contrast to immunostaining for  $\gamma$ -GTP, which, as previously reported, was localized to the apical surface of developing renal vesicles (23). Immunoreactivity for AS and AL in proximal tubular

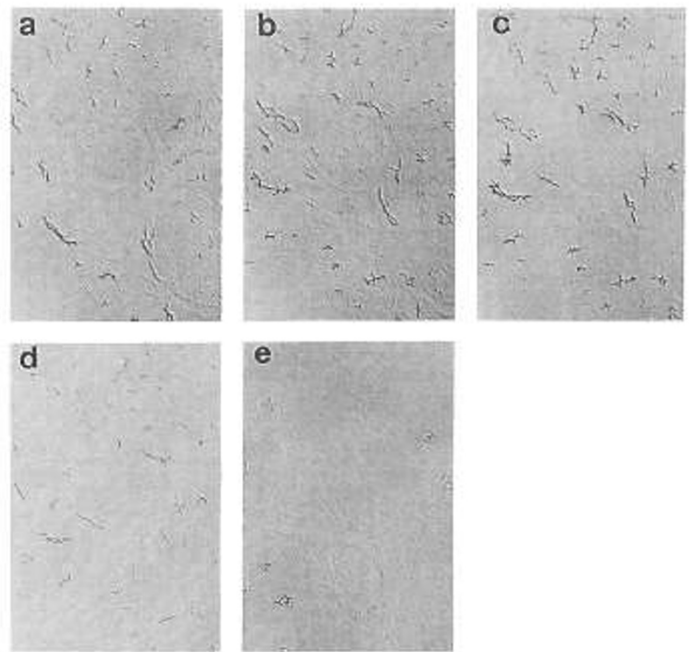


Fig. 1. Localization of AS and AL in young adult mouse kidney. Serial sections were stained, in order, for Lotus tetragonolobus lectin (a), AL (b),  $\gamma$ -GTP (c), AS (d), and Dolichos biflorus lectin (e). Original magnification  $\times 100$ .

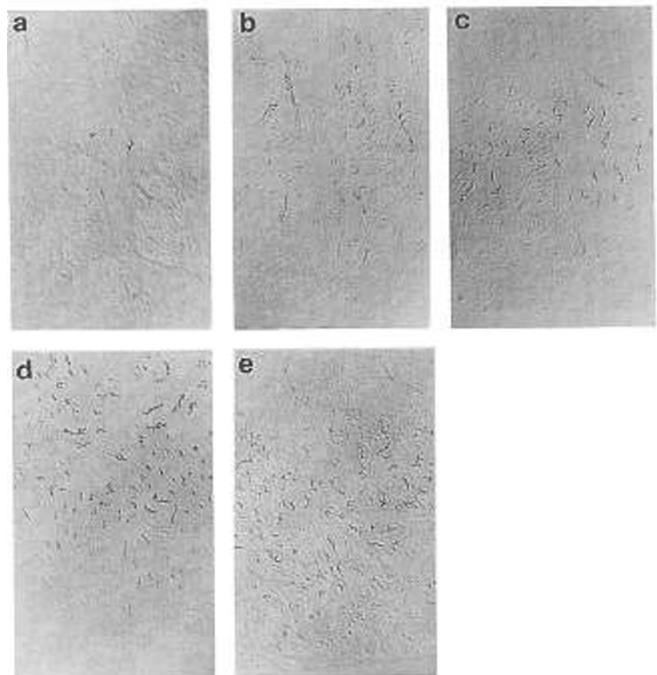


Fig. 2. Immunocytochemical localization of AL during fetal development. Kidney sections were fixed and stained as noted in Materials and Methods. At d 15 (a), sparse immunoreactivity was present in a population of subcortical proximal tubules. At progressive stages of nephrogenesis—d 17 (b), 19 (c), 21 (d), and 23 (e)—immunoreactivity increased and staining became localized to cortical regions of mature proximal tubules. This pattern of immunocytochemical staining was identical to that for AS. Original magnification  $\times 100$  (a) or  $\times 66$  (b–e).

epithelial cells appeared at 15 d of gestation and increased in intensity throughout fetal and postnatal nephrogenesis as exemplified in Figure 2. At no stage did other developing tubular segments, glomeruli, or Bowman's capsule react with anti-AL or anti-AS. This developmental pattern of immunolocalization to

proximal tubules was identical to that of  $\gamma$ -GTP (19). Appearance of AS and AL immunoreactivity at 15 d gestation correlates morphologically with the development of the microvillar apparatus and apical-vacuolar network in proximal tubular epithelial cells (28).

To determine whether the developmental increase in immunoreactivity for AS and AL primarily reflected regulation at translational or posttranslational *versus* pretranslational steps in gene expression, relative abundances of the corresponding mRNA during renal development were determined. Messenger RNA for AS and AL were readily measurable at all stages of kidney development, with relative abundances of 2 and 7%, respectively, as early as d 15 of gestation (Fig. 3). Expression of these two mRNA was strikingly coordinate throughout the fetal and neonatal periods. The developmental increases in AS and AL mRNA are specific, because relative mRNA abundance of the constitutively expressed glyceraldehyde-3-phosphate dehydrogenase is unchanged from d 15 of gestation to adulthood (Fig. 3). This specificity in developmental expression is further dem-

onstrated by the relative abundance of the  $\beta$ -tubulin mRNA, which also undergoes little change throughout fetal and neonatal periods but declines sharply in adults, presumably reflecting the much lower level of cell proliferation. The increases in AS and AL mRNA abundance during gestation and neonatal life are consistent with the increased immunoreactivity of these enzymes during this period, although the qualitative nature of the immunostaining precludes precise comparisons between relative mRNA and protein abundances. The increases in mRNA abundance likely reflect increased transcription of these genes, although regulation at posttranscriptional steps cannot be ruled out. The developmental increase in mRNA abundance probably reflects both an increased concentration of these mRNA per cell and an increase in the number of expressing cells. Notably, parturition had no impact on the steady linear increase in abundance of these mRNA during this period. In contrast, mRNA for PEPCK, which is also expressed in proximal tubules (29), sharply increased in relative abundance at birth in both kidney and liver (Fig. 3). Thus, developmental regulation of gene expression for AS and AL in kidney is similar but not identical to that of PEPCK, probably reflecting differences in responses to hormonal changes occurring at birth. This conclusion is in agreement with results of studies on hormonal regulation of AS and AL mRNA abundance in rat kidney (8) and effects of chromosomal deletions on abundance of AS, AL, and PEPCK mRNA in mouse kidney (30).

Because AS and AL are also expressed in liver as components of the urea cycle, it was of interest to compare the tissue-specific patterns of expression in the same animals. The development profiles for AL mRNA in liver and kidney are similar but not identical (Fig. 3). In contrast, the profiles for AS mRNA in liver and kidney are strikingly different. These comparisons suggest that there are cell-specific differences in intrinsic developmental programs or in responses to hormones or other factors. Organ-specific differences in AS and AL gene expression have been observed previously with regard to effects of certain chromosomal deletions on AS and AL mRNA levels in mouse liver and kidney (30). These differences raise the possibility that kidney-specific defects in AS or AL expression may exist. Unlike deficiencies in hepatic AS or AL, which result in hyperammonemia and death if untreated, renal AS or AL deficiency might result in modest growth retardation in infants, possibly accompanied by mild citrullinemia. Therefore, these defects would probably escape detection unless the diet were seriously deficient in arginine.

The small intestine of mice also contains significant amounts of AS and AL in the perinatal period. In contrast to liver and kidney, levels of AS and AL in small intestine of mice are high at birth and decline to barely measurable values in intestine of adult mice (31). This has been interpreted to suggest that the small intestine has significant capacity for endogenous arginine biosynthesis in the perinatal period, whereas the kidney has the predominant role for arginine biosynthesis in the adult (31, 32). Overall, the results indicate a complex pattern of developmental expression for the arginine biosynthetic enzymes in different organs.

Determination of the site and development of AS and AL expression in kidney are important for a better understanding of the role of arginine biosynthesis in renal physiology and in interorgan relationships during development and in disease. These enzymes and their corresponding mRNA should also serve as useful markers for further analysis of the differentiation and development of renal proximal tubules *in vivo* and in cultured explants.

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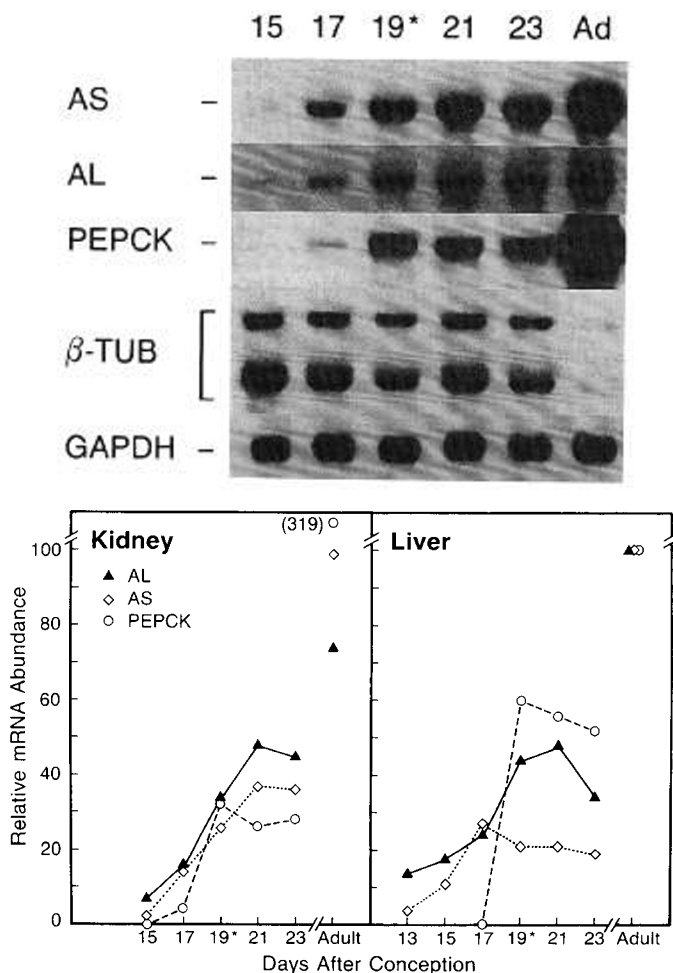


Fig. 3. Relative abundance of mRNA in fetal and neonatal mouse liver and kidney. Ten- $\mu$ g aliquots of total RNA were subjected to Northern blot analysis as described in Materials and Methods. For d 15-19, RNA was extracted from livers and kidneys pooled from 15 mice; d 21 and 23, from nine mice; and adult, from three female mice. Upper panel: autoradiogram of Northern blots. Messenger RNA for  $\beta$ -tubulin ( $\beta$ -TUB) and glyceraldehyde-3-phosphate dehydrogenase (GAPDH) served as internal controls. Days after conception are indicated at the top of the panel; Ad = adult. Lower panel: Abundances of mRNA encoding AS, AL, and PEPCK are expressed relative to the value determined for adult mouse liver. The asterisk denotes the day of birth. Values for liver mRNA are reproduced from reference 13 with permission of Academic Press.

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