

sulfonation not only of the alicyclic amines but also dehydroepiandrosterone. However, it had little sulfonating activity toward 2-naphthol, desipramine and aniline. The apparent kinetic parameters obtained for the enzymatic sulfonation of PTHP were as follows:  $K_m$  382  $\mu$ M and  $V_{max}$  278 nmol/mg of protein/min.

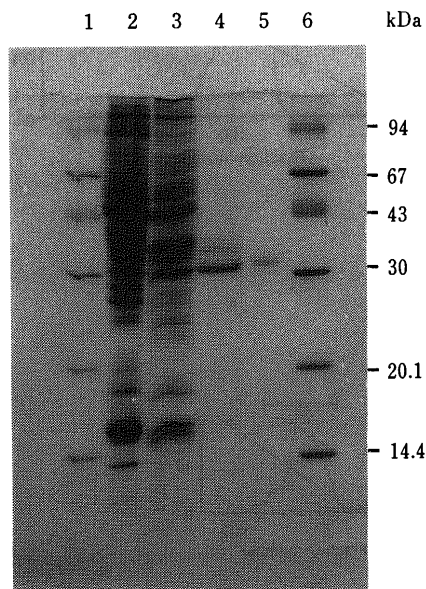


Fig. 2. SDS-PAGE of Purified Alicyclic Amine *N*-Sulfotransferase

Lanes 1 and 6 contain the molecular mass standard proteins, including phosphorylase b (94000), bovine serum albumin (67000), ovalbumin (43000), carbonic anhydrase (30000), soybean trypsin inhibitor (20100) and  $\alpha$ -lactalbumin (14400). Lane 2 contains the protein of female rat liver cytosol. Lanes 3 and 4 contain the active alicyclic amine *N*-sulfotransferase fraction recovered DE-52 and PAP-agarose chromatography, respectively. Lane 5 contains the purified alicyclic amine *N*-sulfotransferase. Staining was performed with Coomassie blue.

TABLE II. Substrate Specificity of Purified NST-1

Substrate	Concentration (mM)	Activity of NST-1 (nmol/mg protein/min)	
		pH 7.4	pH 10
PTHP	2.0	44	212
Tiaramide	5.0	120	165
DETR	5.0	18	77
Desipramine	0.5	N.D.	3.7
Aniline	5.0	N.D.	N.D.
2-Naphthol	0.5	5.1	4.8
Dehydroepiandrosterone	0.05	391	368

The activity was assayed with 100 mM phosphate buffer (pH 7.4) or 100 mM glycine-NaOH buffer (pH 10) containing 0.2 mM PAPS and the indicated concentration of substrates. N.D.; not detected ( $<0.4$  nmol/mg protein/min).

**NH<sub>2</sub>-Terminal Amino Acid Sequence** The results of the NH<sub>2</sub>-terminal amino acid sequence analysis of NST-1 is shown in Fig. 3. The NH<sub>2</sub>-terminal amino acid sequence of NST-1 exhibited a high homology with those of STa<sup>14)</sup> and BAST I.<sup>15)</sup> The sequence differed by two (2nd and 12th) and by four (2nd, 12th, 17th and 23rd) amino acids from those of STa and BAST I, respectively.

**Sex Difference in Hepatic Level of NST-1** Immunoblot analysis of male and female rat liver cytosol was performed using rabbit polyclonal antibodies raised against purified NST-1, in order to obtain direct evidence as to whether the enzyme played an important role in the previously demonstrated sex difference (female  $\gg$  male). The male and female rat liver cytosols contained a M.W. 30500 protein, corresponding to NST-1, and there was a marked difference in the cytosolic levels of NST-1 between female and male rat livers (female  $\gg$  male) (Fig. 4). Other immunoreactive proteins were also observed in both cytosols.

## DISCUSSION

We report here the purification of a form of female rat liver sulfotransferase conjugating PTHP. Purification of an alicyclic amine *N*-sulfotransferase was performed by a three-step procedure. NST-1 had a higher affinity towards PAP-agarose and hydroxylapatite-HPLC than the other alicyclic amine *N*-ST isozymes (Fig. 1).

NST-1 catalyzed sulfonation not only of the alicyclic amines but also the endogenous steroid dehydroepiandrosterone, which is a typical substrate for rat liver hydroxysteroid ST.<sup>1,2)</sup> NST-1 had little sulfonating activity towards 2-naphthol, which is a typical substrate for rat liver aryl ST.<sup>1,2)</sup> The aryl STs have been demonstrated to be unable to catalyze sulfonation of endogeneous steroids.<sup>1,2)</sup> Therefore, NST-1 could be an isozyme of hydroxysteroid ST. Thus, this study provides the first direct evidence that the hydroxysteroid ST participates in enzymatic alicyclic amine *N*-sulfonation. The purified enzyme exhibited no activity toward an alkylamine (desipramine) and an arylamine (aniline). These amines might be catalyzed by other isozymes. Recently, an amine *N*-sulfotransferase sulfonating 2-naphthylamine was purified from guinea pig liver.<sup>4)</sup> However, this purified enzyme catalyzed not only of arylamines but also alicyclic amines. These results suggest that this purified enzyme has different characteristics from NST-1.

Recently, Ogura *et al.* reported the N-terminal amino

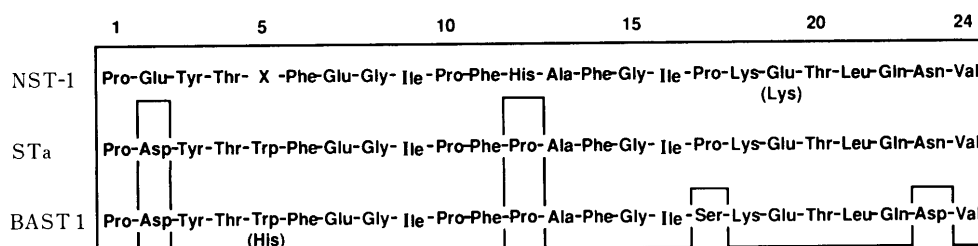


Fig. 3. Comparison of NH<sub>2</sub>-Terminal Amino Acid Sequences of NST-1, STa and BAST I

In the sequence of NST-1, "X" indicated an unidentified amino acid. The sequences of STa and BAST I have been reported by Ogura *et al.* and Barnes *et al.*, respectively.<sup>14,15)</sup>