# Identification and Characterization of Antidepressant-Sensitive Serotonin Transporter Proteins Using Site-Specific Antibodies

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Serotonin (5HT) transporters (SERTs) are responsible for clearance of synaptic and plasma 5HT and are molecular targets for multiple therapeutic and addictive compounds. Recently brain and peripheral SERT cDNAs have been cloned and characterized functionally in transfected cells. Antipeptide (S365) and anti-fusion protein (CT-2) antibodies, directed at epitopes poorly conserved among other Na<sup>+</sup>/Cl<sup>-</sup> cotransporters, have been prepared to facilitate the identification and characterization of SERT proteins in native and transfected cells. Immunoprecipitations and immunoblots of rat/human SERT-transfected HeLa cells reveal specific SERT-immunoreactive glycoproteins absent from extracts of vector-transfected cells and absent when incubations were conducted using peptide- or fusion protein-absorbed antibody. In SDS-PAGE of membranes prepared from rat midbrain and cortex, SERTs migrate as single 76 kDa polypeptides with a relative abundance consistent with the known distribution of 5HT neurons and axonal projections. SERT-immunoreactive proteins are also detectable in platelet and pulmonary membranes, sites of peripheral 5HT uptake, but not in liver. Our studies also indicate that brain and platelet SERTs are formed from identical polypeptides differing significantly in their extent of N-linked glycosylation. Immunocytochemistry performed on rat brain sections with CT-2 antibody revealed SERT expression associated with brainstem raphe nuclei in a pattern virtually identical to that obtained by labeling adjacent sections with 5HT antisera. SERT-immunoreactive fibers were found to be widely distributed throughout the rodent brain, with highest density in forebrain regions known to receive a dense serotonergic innervation. In a similar manner, CT-2 antibody also detects endogenous expression of human SERT proteins, providing an opportunity for future studies on the modulation of transporter protein expression in neurologic and psychiatric disorders.

Key words: midbrain, platelet, immunoblot, glycosylation, immunocytochemistry, raphe nuclei]

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Na<sup>+</sup>- and Cl<sup>-</sup>-coupled serotonin (5-hydroxytryptamine, 5HT) transporters (SERTs) exist on the plasma membrane of serotonergic presynaptic terminals and play a prominent role in terminating the action of synaptically released 5HT (Ross et al., 1967; Kuhar et al., 1972; O'Reilly and Reith, 1988; Bruns et al., 1993). Blockade of 5HT transport by tricyclic and heterocylic antidepressants results in an acute elevation of extracellular 5HT and an amelioration of a variety of neuropsychiatric disorders (Fuller and Wong, 1989; Barker and Blakely, 1994; Owens and Nemeroff, 1994). SERTs are also molecular targets for cocaine and addictive amphetamines, including 3,4-methylenedioxymethamphetamine (MDMA, "ecstasy") (Blakely et al., 1991a; Hoffman et al., 1991). In addition to their synthesis by serotonergic neurons, SERTs are also displayed on plasma membranes of platelets (Sneddon, 1973; Rudnick, 1977), pulmonary endothelium (Gillis, 1985), and placental brush border epithelia (Balkovetz et al., 1989), where they participate in systemic 5HT homeostasis (see Fozard, 1989).

Recently, cDNAs encoding functional SERTs have been cloned from rat brain (rSERT; Blakely et al., 1991a) and basophilic leukemia (RBL) cells (Hoffman et al., 1991), mouse brain (Chang et al., 1994), and human placenta (hSERT; Ramamoorthy et al., 1993). Sequence identity of brain and RBL cDNAs indicates that neural and non-neural SERTs are encoded by a common gene, an hypothesis supported by the identification of single SERT loci in human (Ramamoorthy et al., 1993) and mouse (Gregor et al., 1993). Furthermore, SERTs encoded by human platelet and brain RNAs appear to be identical (Lesch et al., 1993b,c). Rodent and human SERTs are predicted to be encoded by subunits of 630 amino acids with >90% cross-species identity. Hydropathy analysis suggests that SERTs exhibit a topology of 12 transmembrane domains (TMDs) with a large extracellular loop between TMDs 3 and 4 that contains sites for N-linked glycosylation. NH<sub>2</sub>- and COOH-termini are predicted to lie in the cytoplasm. This topology is characteristic of other members of the Na<sup>+</sup>/Cl<sup>-</sup>-dependent transporter gene family whose substrates include other neurotransmitters, neuromodulators, and osmolytes (Amara and Kuhar, 1993; Rudnick and Clark, 1993). SERTs are most closely related to the norepinephrine and dopamine transporters (NETs and DATs), with greatest sequence divergence present in NH<sub>2</sub>- and COOH-termini as well as in the large TMD3-4 extracellular loop (Amara and Kuhar, 1993; Rudnick and Clark, 1993).

Whereas brain and peripheral SERTs are predicted to be encoded by identical polypeptides, little information exists to describe the extent and diversity of SERT post-translational modifications, nor have SERT proteins been visualized *in situ*. These

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issues are magnified by the frequent use of platelet SERTs as a model for the behavior of brain SERTs in disease states (Stahl, 1977; Da Prada et al., 1988; Ellis and Salmond, 1994). Alterations in 5HT transport and/or antidepressant binding have been reported in platelet and/or brain from patients with major neurologic and psychiatric disorders, including depression and suicide (Tuomisto et al., 1976; Meltzer et al., 1981; Paul et al., 1981; Stanley et al., 1982), geriatric depression (Nemeroff et al., 1988), Parkinson's disease (Raisman et al., 1986; D'Amato et al., 1987b), alcoholism (Daoust et al., 1991), schizophrenia (Kaplan and Mann, 1982; Joyce et al., 1993), and progressive supranuclear palsy (Chinaglia et al., 1993). However, the inability to evaluate structural changes in SERT protein has limited interpretation of these observations at a molecular level. Furthermore, the cellular localization of SERTs has, until now, been indirectly inferred from <sup>3</sup>H-5HT uptake studies (Björkland et al., 1973; Azmitia and Marovitz, 1980) or autoradiographic paradigms with brain sections using radiolabeled serotonin transporter antagonists (Fuxe et al., 1983; D'Amato et al., 1987a; De Souza et al., 1987; Hrdina et al., 1990; Chen et al., 1992). Although this approach can provide sensitive quantitative analyses of SERTs in post-mortem brain sections, interpretation relies upon the absolute identity of SERT antagonist-binding sites with transporter proteins (D'Amato et al., 1987a; Marcusson, 1990; Duncan et al., 1992), and the autoradiographic technique provides only limited subcellular resolution. Thus, controversy exists as to whether SERTs in brain are exclusively neuronal or are expressed as well on glial membranes surrounding serotonergic synapses (Anderson et al., 1992). Below we present our efforts to identify, characterize, and visualize SERT proteins in rat and human brain and peripheral tissues using specific, sitedirected polyclonal antibodies.

## **Materials and Methods**

Production of SERT antisera. For production of antipeptide antibody S365, a synthetic peptide corresponding to amino acids 388-401 (EMRNEDVSEVAKDA, see Fig. 1) and containing an additional COOH-terminal cysteine for carrier conjugation was synthesized and purified by reverse-phase HPLC at the Emory University Microchemical Facility. S365 peptide was coupled to keyhole limpet hemocyanin (KLH) or bovine serum albumin (BSA) carrier proteins (Sigma, St. Louis, MO) by maleimide cross-linking to free sulfhydryl groups as previously described (Melikian et al., 1994). For production of antifusion protein antibody CT-2, a 102 bp segment of rSERT encoding the terminal 34 amino acids of the transporter's COOH-terminus was amplified by polymerase chain reaction (Saiki, 1987) using oligonucleotides containing restriction sites compatible with the pGEX2T fusion protein expression plasmid (Pharmacia, Piscataway, NJ; Smith et al., 1988). Oligonucleotides utilized were 5'-GCGCGCGGATCCCTGAT-CAGCACTCCGGGGACA-3' (sense) and 5'-GCGCGCGAATTCT-TACACAGCATTCATGCTGAT-3' (antisense). After digestion with BamHI and EcoRI, the amplified fragment was cloned into pGEX2T and sequence verified via the dideoxy chain termination sequencing method of Sanger (1977). The resulting plasmid encodes a modified glutathione S-transferase (GST) fusion protein fused in frame with the SERT COOH-terminal fragment. SERT fusion protein was harvested from transformed E. coli strain BL21 after IPTG induction and purified by affinity chromatography following the methods described by Levey et al. (1991). The purified fusion protein migrated as a single band by SDS-PAGE as revealed by Coomassie blue staining.

Rabbit immunization and antibody purification. Female New Zealand white rabbits were maintained and immunized with peptide conjugates or fusion protein by Spring Valley Labs (Sykesville, MD). For peptide antibody production, primary immunization with S365 peptide–KLH conjugate (1000 µg) emulsified in complete Freund's adjuvant was followed by 14 and 28 d booster injections (500 µg) in incomplete Freund's adjuvant. Thereafter, monthly boosts were given as 500 µg in incomplete Freund's adjuvant. Blood was collected, and serum isolated

1–2 weeks following boosts and stored at  $-20^{\circ}$ C. Studies reported reflect sera harvested 2–3 months after the initial immunization. For fusion protein antibody production, primary immunizations with SERT fusion protein (500  $\mu$ g), emulsified as described above, were repeated 2 weeks later, followed by monthly 100  $\mu$ g boosts. Sera were obtained and stored as above. All studies reported with CT-2 antibody were conducted with sera derived 6 weeks after the initial immunization.

Immunoreactivity of S365 antisera was determined by ELISA on BSA-peptide conjugates as described by Harlow and Lane (1988). S365 antisera was utilized for immunoprecipitation without further purification. CT-2 antibody immunoreactivity was assessed by immunoprecipitations of rSERT- or hSERT-transfected cells using unpurified serum and by immunoblots (see below) after purification by affinity chromatography. For initial purification of fusion protein antisera, a column was made by coupling bacterial lysates (124 mg protein) from BL21 cells previously induced to express only GST with a 1:1 mixture (2.5 ml each) of Affi-10 and Affi-15 beads (Bio-Rad, Hercules, CA) according to manufacturer's methods. A SERT fusion column was made by coupling 4 mg fusion protein onto 0.5 ml of Affi-15 beads. Serum was first preadsorbed over the GST/BL21 column and the flow through was applied to the SERT fusion protein column. After washing the column with 10 mm Tris, pH 8.0, the anti-SERT component was eluted with 100 mm glycine, pH 3.0, and neutralized with 1/10 volume of 1 м Tris, pH 8.0. O.D. was used to monitor and quantitate the protein concentration of eluted fractions (1 O.D. = 1  $\mu$ g/ $\mu$ l protein).

Transient expression of transporter cDNAs in HeLa cells. rSERT, hSERT, human NET (hNET), and human DAT (hDAT) cDNAs were transfected into HeLa cells using the vaccinia-T7 expression system (Fuerst et al., 1986; Blakely et al., 1991b). pBluescript SKII (Stratagene, La Jolla, CA) transfections were performed in parallel as a negative control for transporter expression. Briefly, 6-well tissue culture plates were seeded with HeLa cells (American Type Tissue Collection, Rockville, MD) 1 d before transfection. Media was removed and virus (VTF<sub>7-3</sub>, 10 pfu/cell) was added to cells in OPTI-MEM (Life Technologies, Gaithersburg, MD) supplemented with 55 μM 2-mercaptoethanol for 30 min, followed by addition of DNA as a liposome suspension with a 3:1 lipid to DNA ratio (Lipofectin; Life Technologies). Immunoprecipitations were performed using  $3 \times 10^5$  cells transfected with 5 μg DNA, whereas immunoblots utilized 10° cells transfected with 10 μg DNA. Transfected cells were harvested 7 or 14 hr following transfection for immunoprecipitation and immunoblots, respectively. Transport of 5-hydroxy-G-3H-tryptamine creatinine sulfate (8.6 Ci/mmol; Amersham, Buckinghamshire, England), 1-ring 2,5,6-3H-norepinephrine (NE, 48.6 Ci/mmol; DuPont-New England Nuclear, Wilmington, DE), and 7,8-3H-dopamine (DA, 45 Ci/mmol; Amersham) was assayed as previously described (Blakely et al., 1991a; Pacholczyk et al., 1991; Ramamoorthy et al., 1993).

Immunoprecipitation. Four hours post-transfection, cells were washed with prewarmed Met/Cys-free media (ICN, Irvine, CA) and labeled with 500 μl prewarmed Tran-35S label (1131 Ci/mmol, 50 μCi/ml; ICN) in Met/Cys-free media (3 hr, 37°C). For experiments examining the effects of tunicamycin on SERT biosynthesis in HeLa cells, tunicamycin (10 µg/ml; Sigma) or vehicle were added to cells both at the time of transfection as well as during the media change to initiate metabolic labeling. Labeled media was removed and whole cell extracts prepared by washing cells three times in Na phosphate-buffered saline, pH 7.4, and solubilizing in RIPA buffer (10 mm Tris, pH 7.4, 150 mm NaCl, 1 mm EDTA, 0.1% SDS, 1% Triton X-100, 1% Na deoxycholate) supplemented with protease inhibitors (1 mg/ml soybean trypsin inhibitor, 1 mm o-phenanthroline, 1 μg/ml leupeptin, 1 mm iodoacetamide, 1 μm pepstatin A, and 250 µM PMSF; all from Sigma) for 30 min at 4°C with shaking. S365 or CT-2 antisera (5 µl) were added to the labeled cell extracts (300 µl) and samples incubated with continuous mixing (1 hr, 22°C). Protein A-Sepharose beads (Pharmacia) were preblocked by incubation with unlabeled HeLa cell extract (1 hr, 22°C), followed by three washes and resuspension to 30 mg/ml in RIPA buffer. Blocked beads (100 µl) were added to labeled cell extracts and incubated for either 1 hr at 22°C or overnight at 4°C, followed by washing as above, and bound protein was eluted into 100 µl Laemmli sample buffer (15 min, 22°C). Following bead centrifugation, supernatants were subjected to SDS-PAGE (Laemmli, 1970). Gels were soaked in Entensify enhancing solution (Dupont-New England Nuclear), dried, and exposed to x-ray film at -80°C. Control immunoprecipitations were conducted with preimmune serum and S365 serum incubated with S365 peptide  $(10 \mu M)$ .

Membrane preparations and immunoblots. Twelve hours after transfection, HeLa cells were washed, scraped into PBS, and harvested by centrifugation. Cell pellets were resuspended in 10 mm Tris, 1 mm EDTA (TE), pH 7.2, containing protease inhibitors, homogenized, assayed for protein content, and stored at  $-20^{\circ}$ C at  $\sim 1$  mg/ml protein. Prior to SDS-PAGE, membranes were collected at  $21,000 \times g$ . For characterization of SERT expression in primary tissues, male Sprague-Dawley rats were killed by rapid decapitation, and tissues dissected, and homogenized in TE buffer plus protease inhibitors. Homogenates were centrifuged at low speed  $(20 \times g)$  to remove particulate debris, and membranes were pelleted from the supernatants at  $21,000 \times g$  for 20 min. Membrane pellets were resuspended in TE buffer plus protease inhibitors and stored at -80°C. Purified membranes from rat platelets were prepared as described by Nemeroff et al. (1988). Sample protein concentration was determined by the Bradford method (Bio-Rad) using BSA as standard. Membrane pellets were solubilized in 10% SDS for 1 hr at room temperature followed by treatment with Laemmli sample buffer for 20 min. Solubilized proteins were separated by SDS-PAGE (10% separating gel) and then electroblotted onto PVDF membrane (0.45 µm, Millipore, Bedford, MA) at 150 mA for 16 hr in transfer buffer (25 mm Tris, 192 mm glycine). Subsequently, blots were preblocked with 5% nonfat dried milk in PBS and incubated with affinitypurified CT-2 antibody (1 µg/ml) in new blocking solution overnight at 4°C. Blots were incubated with horseradish peroxidase (HRP)-conjugated goat anti-rabbit IgG (1:10,000; Bio-Rad) for 1 hr at room temperature and developed using enhanced chemiluminescence (Amersham). Prestained molecular mass markers (Bio-Rad) were used to determine  $M_r$ , of immunoreactive species. For preabsorption experiments, CT-2 antibody was preincubated with or without SERT fusion protein-conjugated Affi-15 beads (1 µg/ml antibody to 100 µg/ml fusion protein) at room temperature for 1 hr prior to use.

To examine the extent of N-linked glycosylation of rSERT, membranes from rat brain (50 µg membrane protein), platelet (50 µg membrane protein), and rSERT cDNA-transfected HeLa cells (25 µg whole cell protein) were pelleted, resuspended in 20 µl TE containing protease inhibitors, and incubated with PNGase F (800 U/µl; New England Biolabs, Beverly, MA) at 37°C for 4 hr. Deglycosylated membranes were repelleted, dissolved in SDS as described above, and subjected to SDS-PAGE/immunoblot.

Immunocytochemistry. Male Sprague–Dawley rats (n = 6) were deeply anesthetized with Nembutal (200 mg/kg, Abbot Laboratories, Chicago, IL) and perfused through the aorta with Ca2+-free Tyrode's solution (137 mm NaCl, 2.7 mm KCl, 0.83 mm MgCl<sub>2</sub>, 3.5 mm Na, HPO<sub>4</sub>, 8.3 mm NaH<sub>2</sub>PO<sub>4</sub>, 5.6 mm D-glucose, pH 7.4) containing 50 units/ml heparin sulfate (Sigma), followed by 0.1 M Na phosphate-buffered 4% paraformaldehyde (Fisher, Pittsburgh, PA), pH 7.6, with or without 0.15-0.3% glutaraldehyde (Electron Microscopy Sciences, Fort Washington, PA). Animals were then perfused sequentially with 10% and 30% sucrose dissolved in 0.1 M Na phosphate buffer. Brains were removed and cryoprotected in 30% sucrose for 48 hr at 4°C and then sectioned at 40 µm on a freezing stage sliding microtome. Tissue sections were collected into 0.05 M Tris-buffered saline (TBS), pH 7.4. Sections were preblocked in TBS containing 4% normal goat serum (NGS, Organon Teknika, West Chester, PA) and 0.1% Triton X-100 (Sigma) for at least 1 hr at 4°C, and then incubated with affinity-purified CT-2 antibody (0.25-0.5 µg/ml in TBS containing 4% NGS) for 48 hr at 4°C. SERT immunoreactivity was detected with the Vectastain ABC Elite kit (Vector Laboratories, Burlingame, CA) using biotinylated goat anti-rabbit IgG (1:200; 16 hr at 4°C) and the avidin-biotin-peroxidase complex (1:50; 4 hr at 4°C). HRP reaction product was processed with 0.5 mg/ml diaminobenzidine (Sigma) and 0.01% H<sub>2</sub>O<sub>2</sub>. The developed sections were mounted on Superfrost Plus slides (Fisher), dried and dehydrated through graded ethanols, soaked in Histo-Clear (National Diagnostics, Atlanta, GA), and coverslipped for microscopic examination. Control incubations consisted of omission of primary antibody or incubations conducted with fusion protein (100 µg/ml)-absorbed CT-2 antibody. Adjacent sections were also incubated with polyclonal rabbit anti-5HT antibody (1:20,000; Incstar, Stillwater, MN), and immunoreactivity visualized as described for CT-2 antibody.

Human brains were obtained postmortem from four individuals (range of post-mortem interval from 2.5 to 9 hr), and 1 cm blocks of mesopontine tegmentum or temporal lobe were fixed by immersion in cold 4% paraformaldehyde in 0.1 M sodium phosphate buffer, pH 7.2, for 24–36 hr. Blocks were then transferred to 10% sucrose in the same buffer for at least 3 d and then sectioned and processed for SERT

immunoreactivity with CT-2 antibody as described above, with the following modifications. Sections were pretreated with 3% hydrogen peroxide and 1% Triton in 0.05 M TBS for 30 min to inhibit endogenous peroxidase, rinsed five times in TBS, and then processed in avidin D and biotin solutions (Blocking Kit, Vector Laboratories, Inc.) to reduce background. A mixture of 4% each human and goat sera were used in preblock and other antibody-containing steps.

#### Results

Characterization of SERT antibodies

Comparative analysis of rodent and human SERT cDNA inferred amino acid sequences revealed that the loops between several putative TMDs as well as the NH2 and COOH tails are regions highly conserved across SERTs but divergent with other gene family members. These regions are also most likely to be exposed to aqueous compartments for antibody recognition. Accordingly, a synthetic peptide (S365), derived from a putative external loop between TMDs 7 and 8 (Fig. 1) that is absolutely conserved across cloned rat (Blakely et al., 1991a; Hoffman et al., 1991), mouse (Chang et al., 1994), and human (Ramamoorthy et al., 1994) SERTs, was coupled to KLH for rabbit immunization. S365 antisera, but not preimmune sera (Fig. 2), specifically immunoprecipitated a 61 kDa protein species from 35S-Met/Cys metabolically labeled rSERT-transfected HeLa cells. Immunoprecipitation products were absent from extracts prepared from control vector (pBluescipt SKII)-transfected HeLa cells. In addition, recovery of the 61 kDa species from rSERT-transfected HeLa cells was abolished by co-incubation of S365 antibody with the immunizing peptide. Co-incubation of antisera with a different peptide derived from a distinct region of the rSERT sequence failed to block immunoprecipitation (data not shown). S365 antisera also immunoprecipitated an equivalently sized protein from hSERT-transfected HeLa cells. Further attempts, however, to utilize the S365 antibody for immunoblot or immunocytochemical detection of endogenous SERT proteins proved unsuccessful.

To improve our chances of identifying more powerful SERT antibodies, a GST fusion protein comprising 34 amino acids of the rSERT COOH-terminus was synthesized in *E. coli* as described in Materials and Methods. The region selected for fusion protein production (see Fig. 1) is absolutely conserved between rat and mouse, exhibits only three amino acid differences from rat to human, and possesses only 29% identity with the COOH-terminus of the human norepinephrine transporter (Pacholzyk et al., 1991), the most closely related SERT homolog. Following a repetitive absorption procedure to remove *E. coli*—and GST-targeted antibodies, SERT fusion protein—specific antibodies (CT-2) were purified by affinity chromatography. On immunoblots, affinity-purified CT-2 antibody detected the SERT-GST fusion protein but not GST itself nor any proteins derived from *E. coli* lysates (Fig. 3A).

Immunoprecipitation of metabolically labeled rSERT-transfected HeLa cells with affinity-purified CT-2 antibody yielded a 61 kDa species identical in migration to that described above for S365 immunoprecipitations (data not shown). Of greater significance, CT-2 antibody proved capable of recognizing denatured rSERT protein as demonstrated by immunoblots of transfected HeLa cell membrane extracts subjected to SDS-PAGE (Fig. 3B). In short exposures, a single immunoreactive species was evident in membranes from both rat and human SERT-transfected cells, but was absent from membranes of vector- or hNET-transfected cells. CT-2 antibody also exhibited no crossreactivity with proteins from extracts of hDAT-transfected HeLa

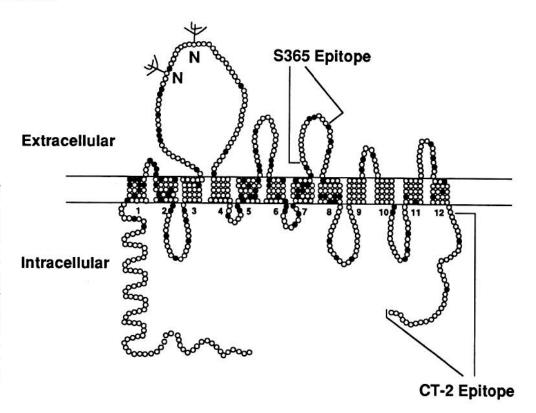


Figure 1. Schematic diagram of SERT antibody epitopes. The 630 amino acid polypeptide predicted from rat and human SERT cDNAs (Blakely et al., 1991, 1992; Hoffman et al., 1991) is shown with 12 putative transmembrane domains as indicated by hydrophobicity analysis. • represent amino acids absolutely conserved across the Na+/Cl- cotransporter gene family. N designates the position of asparagine residues in canonical N-glycosylation sites. The S365 antipeptide antibody is directed at rSERT amino acids 388-401, whereas CT-2 fusion protein antibody is directed at the final 34 amino acids (597-630) of the COOH-terminus.

cells (Pristupa et al., 1994) (data not shown). 5HT, NE, and DA transport assays conducted in parallel with immunoblots confirmed the expression of uptake in the transfected cells from which membrane extracts were prepared (data not shown).

Whereas the major immunoreactive band detected after short exposures of immunoblots from SERT-transfected HeLa cells migrated at  $\sim$ 61 kDa (Fig. 3B,C), longer film exposures re-

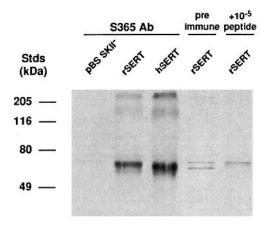


Figure 2. S365 immunoprecipitation of rSERT protein. HeLa cells were transfected with rSERT, hSERT, or pBluescript SKII (vector) DNAs, metabolically labeled, and immunoprecipitated with either S365 antisera, preimmune sera, or S365 antisera preincubated with S365 peptide (10  $\mu$ M) as described in Materials and Methods. Each lane derives from immunoprecipitation of extracts from 5  $\times$  10<sup>5</sup> HeLa cells. S365 antibody immunoprecipitates a major 61 kDa species from both rat and human SERT-transfected cells absent from vector-transfected cells or after immunoprecipitation with preimmune or peptide-blocked antisera. S365 antibody also fails to recognize the homologous epitope in hNET (data not shown). The *numbers to the left of the blot* indicate the positions of molecular mass markers in kilodaltons electrophoresed in parallel.

vealed two additional immunoreactive bands (90 and 200 kDa) absent from vector-transfected cells (Fig. 3D). The 90 and 200 kDa species were also faintly evident in immunoprecipitations of metabolically labeled transfected cells (Fig. 2). CT-2 immunoreactivity was absent from immunoblots conducted with antibody preabsorbed with SERT fusion protein (Fig. 3C,D). In addition, immunoblot immunoreactivity was not diminished by incubations conducted with an equivalent amount of the GST protein lacking the rSERT COOH-terminal fragment, nor did preimmune sera yield visible immunoreactivity (data not shown).

### Detection of endogenous SERT protein expression

The ability to detect directly unlabeled, denatured SERT proteins by immunoblot offered an opportunity to visualize endogenous SERT protein expression. Thus, membrane extracts from rat brain and selected peripheral tissues were immunoblotted with affinity-purified CT-2 antibody as described for SERT-transfected HeLa cells. Single CT-2-immunoreactive species of 76 kDa were detected in cortex and midbrain, but not in cerebellum (Fig. 4A). Among the peripheral tissues examined, platelet membranes displayed prominent SERT immunoreactivity. On short exposures, platelet immunoreactivity appeared as a single 94 kDa band. An 80 kDa CT-2-immunoreactive band was also present in pulmonary membranes, although relatively long exposures were required to visualize this species. SERT-immunoreactive bands were absent from liver membrane extracts. As with blots from transfected cells, antibody recognition of endogenous SERTs was abolished by preincubation with SERT fusion protein (Fig. 4B). Immunoblots of membranes prepared from human platelet and brain displayed immunoreactive SERT bands comparable in size to that observed with rat tissues (data not shown).

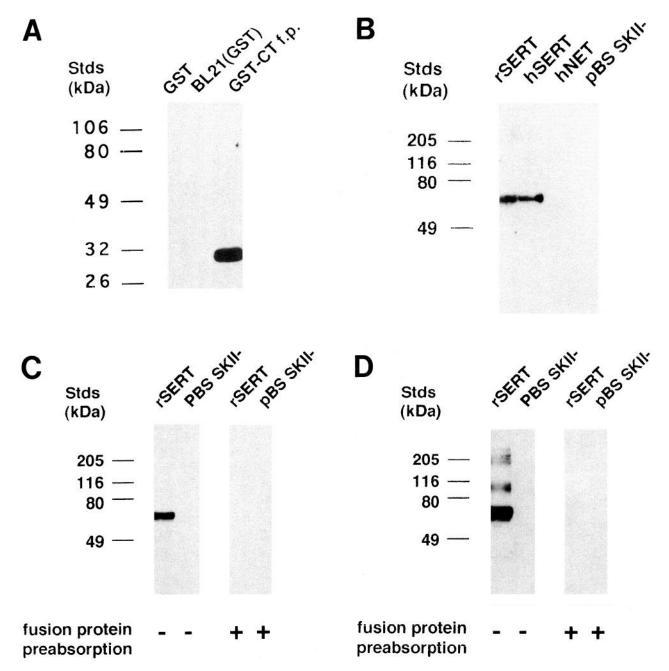


Figure 3. Characterization of SERT fusion protein antibody CT-2 by immunoblot. A, Specificity of CT-2 immunoreactivity for SERT sequences. SDS-PAGE was performed with purified glutathione-S-transferase (GST, 100 ng), total *E. coli* lysate from GST transformants [BL21(GST), 16 μg], or purified SERT fusion protein (GST-CT f.p., 100 ng), blotted and probed with CT-2 antibody (1 μg/ml) as described in Materials and Methods. The affinity-purified anti-SERT antibody recognized the fusion protein, but not the GST or the bacterial components. B, Detection of SERT expression in transfected HeLa cells. SDS-PAGE and immunoblot were performed as described in A, using solubilized membranes from 5 μg of rSERT-transfected cells, 15 μg of hSERT-transfected cells, 15 μg of hNET-transfected cells, or 15 μg of vector (PBSKII)-transfected cells. CT-2 antibody recognizes a major 61 kDa protein in cells transfected with both rat and human SERT cDNAs but not in cells transfected with hNET cDNA. C, Extracts of rSERT cDNA- or vector-transfected HeLa cells (5 μg protein for each) were subjected to SDS-PAGE and immunoblotted with affinity-purified CT-2 antibody (1 μg/ml) with or without preincubation with purified SERT fusion protein (100 μg/ml). D, Longer exposure of experiment presented in A to emphasize additional SERT species detected by CT-2 antibody. Antibody recognition of all SERT species (61, 90, and 200 kDa) was abolished by fusion protein preincubation. The numbers to the left of each blot indicate the positions of molecular mass markers in kilodaltons electrophoresed in parallel.

Glycosylation state of endogenous and transfected rSERT proteins

The multiple SERT proteins evident in transiently transfected HeLa cells and the disparate mobilities of endogenous brain and platelet SERT proteins prompted us to examine the presence and

extent of SERT N-linked glycosylation. Metabolic labeling of transfected HeLa cells in the presence or absence of the glycosylation inhibitor tunicamycin (Fig. 5A) followed by SERT immunoprecipitation with S365 antibody revealed an increased mobility of SERT protein by approximately 5 kDa. To demonstrate the second second

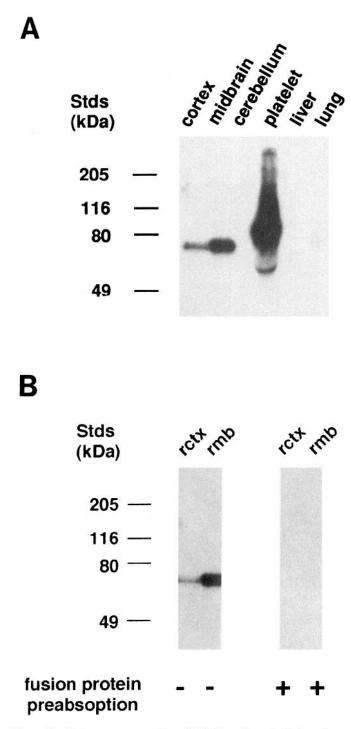


Figure 4. Endogenous expression of SERT proteins. A, Brain and peripheral SERT expression. Extracts from tissue membranes (50  $\mu g$ ) were subjected to SDS-PAGE and immunoblotted with CT-2 antibody (1  $\mu g/ml$ ) as described in Materials and Methods. The blot presented was exposed so as to reveal the species present in lung as well as in brain and platelet. B, Specificity of endogenous SERT detection by CT-2 antibody. Brain membrane extracts (50  $\mu g$ ) were electrophoresed and blotted as in A using either CT-2 antibody or CT-2 antibody preincubated with SERT fusion (100  $\mu g/ml$ ). CT-2-immunoreactive species in midbrain and cortex were absent from incubations conducted with fusion protein—absorbed antibody. The numbers to the left of each blot indicate the positions of molecular mass markers in kilodaltons electrophoresed in parallel.

strate the presence of SERT glycosylation directly, membranes from SERT-transfected HeLa cells were digested with PNGase F prior to performing immunoblots with CT-2 fusion protein antibody. As with immunoprecipitations, the major species at 61 kDa shifted to an apparent molecular mass of 56 kDa (Fig. 5B). In the data presented, blots were exposed to allow for the detection of the two more slowly migrating SERT species before and after PNGase F treatment. Whereas the most slowly migrating species at 200 kDa failed to exhibit an appreciable mobility shift, the 90 kDa form disappeared, presumably shifting to the 56 kDa form. Thus, the lower molecular mass SERTs differentially accumulated in transfected HeLa cells appears to be due to different levels of N-linked glycosylation. Like HeLaexpressed SERTs, endogenous rat brain and platelet SERTs exhibited altered mobilities after PNGase F digestion, consistent with the presence of N-linked glycosylation (Fig. 5B). Although brain and platelet SERTs exhibit distinct mobilities by SDS-PAGE prior to PNGAse F digestion or in mock digestions conducted in the absence of glycosidase, both proteins migrate equivalently at 60 kDa following PNGase F treatment. Thus differential N-linked glycosylation also appears to generate distinct CNS and peripheral SERT subunits.

The 200 kDa band present in transfected cells did not reveal an appreciable shift in relative molecular mass in our gel system. The 200 kDa SERT form may represent a transporter aggregate. Prolonged incubations at elevated temperatures often resulted in the appearance of higher  $M_r$ , species (see Fig. 5B, platelet samples treated with PNGase F) and/or a loss of immunoreactivity. The loss of immunoreactivity of the 94 kDa platelet species in mock digestions (37°C, 4 hr, no enzyme) was also observed in a repeat of this experiment and paralleled by an increase in aggregated SERT. Thus deglycosylated SERT monomers may have less of a tendency to nonspecifically aggregate, or glycosylation may be involved in multimerization. Interestingly, brain and platelet SERTs lacking N-linked glycosylation migrate consistently slower (by ~2 kDa) than do deglycosylated SERTs derived from transfected HeLa cells. The source of this difference in mobility remains to be elucidated.

Immunocytochemical detection of rat and human brain SERT

Since the CT-2 antibody specifically detected endogenous SERT proteins by immunoblot analysis, we explored its utility for visualizing the cellular localization of the transporter in situ. Immunocytochemistry performed on serial coronal sections of the entire rat brain using CT-2 antibody revealed intense staining of neurons in the brainstem raphe nuclei. As shown in Figure 6A-C, neurons of the dorsal raphe in both the lateral and medial subdivisions exhibit intense SERT immunoreactivity. At this level, neurons of the raphe pontine subdivision (B9) were also highly stained (Fig. 6D). The somatic staining filled the cytoplasm around unstained nuclei and also extended into dendritic elements. The neuropil SERT immunoreactivity was also dense and appeared comprised of puncta and varicose fibers. As is evident from Figure 6A, somatic staining was highly restricted with little or no immunoreactivity in perikarya outside of the raphe complex. Importantly, neurons of the locus ceruleus and substantia nigra that express homologous NETs and DATs, respectively, were devoid of staining although the neuropil of both areas were densely packed with SERT-immunoreactive fibers. We could detect no SERT-immunoreactive cells with the morphology of glia in any of our sections.

To validate the specificity of SERT immunoreactivity, adja-

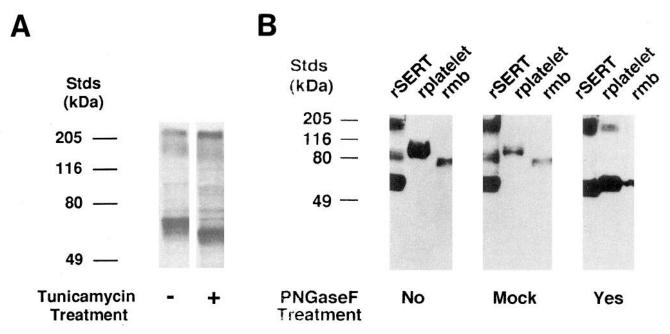


Figure 5. Analysis of rSERT N-linked glycosylation. A, Synthesis of rSERT in transfected HeLa cells in the presence or absence of tunicamycin. HeLa cells were transfected with rSERT cDNA, metabolically labeled with <sup>35</sup>S-methionine and cysteine, and immunoprecipitated with S365 antibody as described in Materials and Methods. The major rSERT species detected at 61 kDa from untreated cells shifts in mobility to 56 kDa in cells treated with tunicamycin. B, Enzymatic deglycosylation of endogenous and transfected rSERT. Membranes from rSERT-transfected HeLa cells (from 25 μg of cell homogenates, 5 μg loaded), brain (50 μg), and platelets (50 μg) were treated with PNGase F to remove N-linked oligosaccharides as described in Materials and Methods. Samples were subjected to SDS-PAGE and immunoblotted with CT-2 antibody. Exposures were developed to visualize multiple SERT species in HeLa cells. Treatment with PNGase F increased the mobility of both transfected and endogenous SERTs. Both platelet and brain SERTs exhibit a 58 kDa relative molecular mass after PNGase F digestion, whereas the transfected rSERT protein migrates as a 56 kDa species. Treatment of platelet membranes with PNGase F resulted in the appearance of SERTs at ~200 kDa, presumed to be a nonspecific aggregate. An aggregate was also present after mock treatments; however, in the experiment shown, this form resided at the interface of the stacking and resolving gel and is not shown.

cent sections were incubated with either affinity-purified CT-2 antibody, CT-2 antibody preabsorbed with SERT fusion protein (100  $\mu$ g/ml), or without primary antibody. As shown in Figure 7, A and B, CT-2 staining is abolished by fusion protein preincubation. A similar absence of immunoreactivity was evident in CT-2 antibody–free incubations (data not shown). Adjacent sections incubated with anti-5HT antibody (Fig. 7C) labeled the same medial, dorsal, and lateral subdivisions of the dorsal raphe complex detected by CT-2 antibody, consistent with the conclusion that SERT-immunoreactive neurons are serotonergic.

In addition to cell body staining, SERT-immunoreactive fibers were observed throughout the rodent brain, particularly evident in forebrain regions known to receive a prominent serotonergic innervation, such as the hypothalamus, substantia nigra, hippocampus, amygdala, and cortex. In Figure 8 B and C, SERTimmunoreactive fibers can be seen to course throughout the rat hippocampus with highest density observed in molecular and polymorphic layers. The highest concentration of SERT-immunoreactive fibers in the hippocampal formation was observed in the molecular and polymorphic layers of the CA3 region, with reduced density in CA1 dendritic fields. The cerebellum contained few CT-2-positive fibers; particularly noticeable were a small population of highly varicose fibers terminating in rosettes in the granular layer of the flocculonodular lobe (data not shown). Like somatic staining, SERT fiber staining was eliminated by preincubation of CT-2 antibody with SERT fusion protein (data not shown). The visualization of SERT-immunoreactive fibers was also diminished in sections prepared with glutaraldehyde-containing fixative. However, 0.15-0.3% glutaraldehyde provided significantly enhanced detection of immunoreactivity in cell bodies, apparently through a reduction of fiber staining in the surrounding neuropil.

Since CT-2 antibody recognized hSERT as well as rSERT in transfected cells and native tissues, we examined the ability of the antibody to stain serotonergic neurons and fibers in selected post-mortem human brain sections. As in the rat, the raphe nuclei showed dense immunoreactivity associated with cell bodies and neuropil (Fig. 9A,B). Neurons in the locus ceruleus and substantia nigra did not exhibit SERT immunoreactivity, although varicose axons were widely distributed throughout the mesopontine tegmentum as well as in entorhinal cortex (Fig. 9C,D). Glia appeared unstained. Control sections in which primary antibody was omitted showed no significant staining, although neuromelanin content of catecholamine neurons rendered them distinctly visible.

### Discussion

Despite the importance of SERTs in the clearance of synaptic 5HT, SERT proteins have been only incompletely characterized. The availability of cloned rat (Blakely et al., 1991a; Hoffman et al., 1991), human (Lesch et al., 1993b,c; Ramamoorthy et al., 1993), and mouse (Chang et al., 1994) SERT cDNAs permitted the generation of sequence-specific antibodies to identify and localize SERT proteins in transfected cells and native tissues. SERTs are members of the Na<sup>+</sup>/Cl<sup>-</sup> cotransporter gene family and are most closely related to cocaine-sensitive NETs and DATs (Amara and Kuhra, 1993; Rudnick and Clark, 1993). However, these biogenic amine transporters exhibit substantial sequence

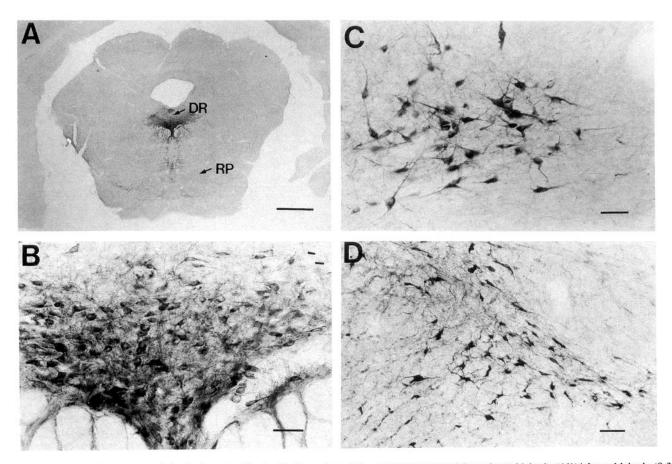


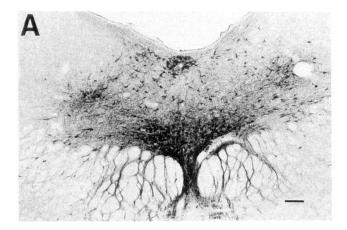
Figure 6. SERT-like immunoreactivity in the rat midbrain. Brain sections (40 μm) were prepared from formaldehyde (4%)/glutaraldehyde (0.3%)-perfused rats and stained with affinity-purified CT-2 antibody (0.5 μg/ml) as described in Materials and Methods. A, A coronal section through the rat midbrain displays intense SERT-like immunoreactivity restricted to midline neurons ventral to the third ventricle. The section shown is taken at the level of the dorsal raphe complex. B, Higher magnification of SERT-like-immunoreactive neurons in the median subnucleus of the dorsal raphe complex. D, SERT-like-immunoreactive neurons in the nucleus raphe pontis. These neurons are ventral and lateral to the dorsal raphe groups depicted in B and C. Scale bars: A, 1000 μm; B-D, 50 μm.

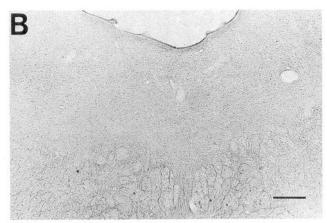
divergence in regions between putative TMDs as well as in the predicted cytoplasmic NH<sub>2</sub>- and COOH-termini. We utilized two of these regions that are highly divergent among Na<sup>+</sup>/Cl<sup>-</sup> cotransporters, but highly conserved among SERT species variants, to prepare SERT-specific antibodies. The equivalent positions to the S365 and CT-2 epitopes in hNET exhibit only 14% and 29% identity with the comparable SERT sequences, respectively, and are further divergent with other members of the Na<sup>+</sup>/Cl<sup>-</sup> cotransporter gene family. Thus, this strategy has the potential for specific recognition of SERT proteins across multiple species.

# SERT antibody specificity analyzed in transfected cells

To determine the specificity of SERT antipeptide and fusion protein antibodies, we transiently expressed rat or human SERT cDNAs in vaccinia-T7-infected HeLa cells. SERTs expressed in this manner transported 5HT with high affinity in an Na<sup>+</sup>- and Cl<sup>-</sup>- dependent manner, and are blocked by SERT-selective antagonists including paroxetine and citalopram (Blakely et al., 1991a; Hoffman et al., 1991; Ramamoorthy et al., 1993). S365 antisera specifically immunoprecipitated 61 kDa proteins from both rSERT- and hSERT-transfected HeLa cells, as expected given the equivalent length of their predicted coding sequences; however, this antibody proved an unsuitable reagent for immunoblots of membranes prepared from the same cells. The S365

epitope could be sensitive to the denaturing conditions of SDS-PAGE. This is a common characteristic of antipeptide antibodies, as evidenced in immunologic characterization of DAT (Vaughn et al., 1993) and NET (Melikian et al., 1994) proteins. However, when overexpressed in Sf9 cells, SERTs can be immunoblotted with the S365 antibody (Tate and Blakely, 1994). The level of functional expression in the baculovirus expression system is approximately fourfold greater than that obtained with SERT expression in HeLa cells (Blakely et al., 1993), and, in side by side comparisons, detection of the insect cell-expressed SERT with CT-2 serum is achieved with significantly greater dilutions of antisera compared with the S365 preparation (Tate and Blakely, 1994). These data indicate that the problem in use of the S365 antibody for immunoblotting lies with limited sensitivity (titer, affinity) rather than epitope availability. Given the predicted external location and proven antigenicity of this epitope, higher-affinity antibodies directed at this region may prove useful for SERT topology and membrane insertion analysis. The CT-2 antibody, directed at the rSERT COOH-terminus, represents a more useful reagent under a wide range of conditions for both heterologously expressed and endogenous SERTs. Thus, in both immunoprecipitation and immunoblot experiments with CT-2 antibody, a major 61 kDa species is evident and absent from cells transfected with homologous transporter plasmids





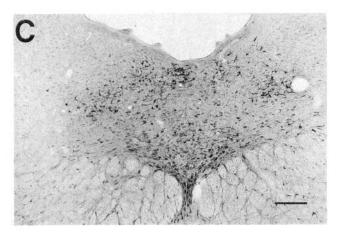


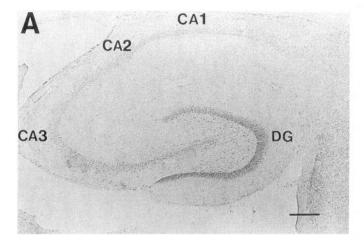
Figure 7. Specificity of SERT-like immunoreactivity. Adjacent rat brain sections (40  $\mu$ m) were prepared from formaldehyde (4%)/glutar-aldehyde (0.3%)-perfused rats and stained with 0.5  $\mu$ g/ml affinity-purified CT-2 antibody (A), CT-2 antibody preabsorbed with 100  $\mu$ g/ml SERT fusion protein (B), or 1:20,000 anti-5HT antibody (C) as described in Materials and Methods. Note that all three subdivisions (dorsal, medial, lateral) of the dorsal raphe complex are stained with either CT-2 or anti-5HT antibody. With the conditions utilized for this experiment, fiber staining is more evident with the CT-2 antibody (particularly evident in the medial subnucleus) than with the 5HT antibody. Staining with CT-2 antibody is completely abolished with fusion protein preincubation. Scale bars: 100  $\mu$ m.

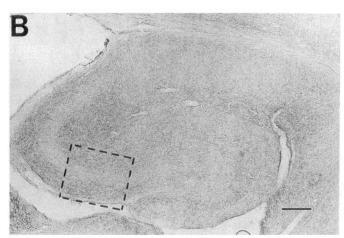
(NET, e.g., Fig. 3; DAT, data not shown) or the plasmid vector alone. Interestingly, epitope-tagged SERTs bearing additional sequences on the rSERT COOH-terminus cannot be recognized by CT-2 antibodies on immunoblots (Tate and Blakely, 1994), suggesting that this antibody's main recognition site is positioned very near SERT's most COOH-terminal amino acids.

### SERT protein from brain and peripheral tissues

In CNS, raphe nuclei express high levels of SERT mRNA (Blakely et al., 1991a; Fujita et al., 1993; Austin et al., 1994). In addition, SERT activity has been described in certain peripheral tissues, including platelets (Sneddon, 1973; Rudnick, 1977) and lung (Gillis, 1985). The ability of SERT CT-2 antibody to detect SERT proteins on immunoblots allowed us to characterize the pattern of SERT protein expression in native membranes. Immunoblots with CT-2 antibody demonstrate the presence of single immunoreactive bands in both rat brain and peripheral tissues in a pattern consistent with predicted regional abundance. Thus, SERT proteins were evident in midbrain and cerebral cortex, but not cerebellum, which receives only a sparse serotonergic innervation (Steinsbusch, 1984; Takeuchi, 1988). SERT immunoreactivity in midbrain is greater than that found in cortex, paralleling the greater relative abundance observed for midbrain SERT antagonist-binding sites (De Souza and Kuyatt, 1987; Hrdina et al., 1989). The high level of SERT protein observed in platelets is consistent with the enrichment of 5HT transporter sites revealed by radioligand binding (Plenge and Mellerup, 1991). The low abundance of SERT-immunoreactive protein in the lung seems at odds with endogenous expression by pulmonary endothelial cells (Lee et al., 1986); alternatively, it may represent expression by the small number of 5HT-immunoreactive neuroepithelial bodies lining the airway mucosa (Youngson et al., 1993).

Interestingly, SDS-PAGE mobility of SERTs from rat platelet (94 kDa) and pulmonary (80 kDa) membranes differs significantly from that exhibited by rat brain SERTs (76 kDa). SERT protein coding sequences inferred from rat brain and basophilic leukemia cell (RBL) cDNAs are identical (Blakely et al., 1991a; Hoffman et al., 1991). Similarly, identical primary amino sequences have been predicted for human placental, brain, and platelet SERTs (Lesch et al., 1993b,c; Ramamoorthy et al., 1993). These data, along with the presence of a single SERT genomic locus in human (Ramamoorthy et al., 1993) and mouse (Gregor et al., 1993), suggested to us that differential post-translational modifications underlie the differences in size of brain and peripheral SERTs. Deglycosylation experiments verified that the distinct electrophoretic mobility of endogenous SERTs is due to differential N-glycosylation. Multiple mobilities were also observed for SERTs in transfected HeLa cells and, like NETs expressed in transfected LLC-PK, cells, appear to reflect sequential stages of transporter glycosylation (Melikian et al., 1994). The functional role of differential N-linked glycosylation of brain and platelet SERTs is presently unclear. CNS DATs have also been shown to exhibit endogenous region-specific glycosylation differences (Lew et al., 1992), although no functional consequences of these differences are known. Glycosylation plays a role in receptor and transporter assembly and trafficking (Rands et al., 1990; Asano et al., 1993; Collier et al., 1993), and in some cases contributes to functional properties (Boege et al., 1988; Leconte et al., 1992). Recent studies of rSERT point mutants lacking one or both N-glycosylation sites indicate a reduction in SERT antagonist  $B_{\text{max}}$ , but not  $K_D$  (Tate and Blakely,





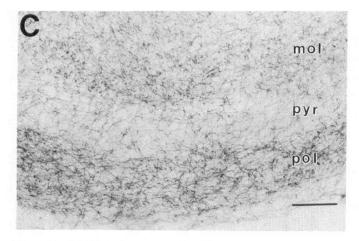


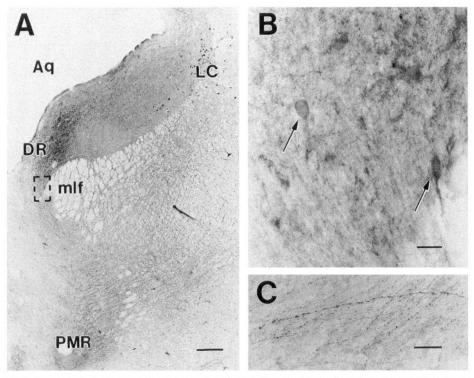
Figure 8. SERT-like-immunoreactive fiber staining in the rat hippocampal formation. Brain sections (40  $\mu$ m) were prepared from formaldehyde (4%)-perfused rats and stained with affinity-purified CT-2 antibody (0.5  $\mu$ g/ml) as described in Materials and Methods. A, Nissl stain of a horizontal section revealing the localization of neurons in the granule layer of the dentate gyrus (DG) and pyramidal neurons (CA1–CA3) of the hippocampus. B, SERT-like-immunoreactive fibers concentrated in the polymorphic and molecular layers of the hippocampus. Note the high density of fibers in the CA3 region. C, High-power bright-field micrograph of CA3 region shown in the box of B. Note the appearance of varicose enlargements on individual fibers. Scale bars: A and B, 250  $\mu$ m; C, 100  $\mu$ m.

1994). These findings suggest that the differences in platelet and brain SERT mobility may be inconsequential for 5HT or SERT antagonist recognition, but rather may influence transporter biosynthetic efficiency in these distinct cellular contexts. A small (2 kDa) but consistent difference in mobility exists between N-deglycosylated SERTs derived from endogenous sources or transfected cells. These data suggest the existence of additional post-translational modifications of the endogenous SERT protein absent from SERT-transfected HeLa cells. Candidate SERT modifications, including O-linked glycosylation, sulfation, acylation, and/or phosphorylation, are presently under evaluation.

Antidepressant-binding glycoproteins from rat brain and human platelets have been partially purified (Biessen et al., 1990; Graham et al., 1992; Launay et al., 1992). The size of endogenous rat brain SERTs detected with CT-2 antibody agrees well with the size of a citalopram-binding protein (73 kDa) partially purified from rat brain (Graham et al., 1992), supporting the hypothesis that a single protein comprises both the transport mechanism as well as SERT antidepressant recognition sites (Blakely et al., 1991a; Hoffman et al., 1991). However, the apparent M, of antidepressant-binding proteins partially purified from human platelets (78 kDa, Biessen et al., 1990; 68 kDa, Launay et al., 1992) is lower than what we obtained in direct platelet immunoblots. Possibly, these differences reflect different mass standards used for SDS-PAGE. In addition, we have noted that human platelet SERTs are susceptible to significant proteolysis during membrane preparation. In this regard, partially truncated SERTs expressed in HeLa cells still transport 5HT and recognize antidepressants with high affinity (Blakely et al., 1993), and thus could persevere through affinity-based purification procedures even if partially degraded. Finally, the mass of the core SERT polypeptide (68 kDa) calculated from cloned rat cDNAs is higher than that derived from deglycosylated SERTs (56 kDa), but can easily be accounted for by the anomalous mobility of highly hydrophobic transporter proteins on SDS gels (Keynan et al., 1992; Melikian et al., 1994).

SERT-like immunoreactivity in serotoninergic neurons and processes

SERT-like-immunoreactive soma in the rat brain were found to be highly restricted to serotonergic raphe nuclei in a manner virtually identical to that observed for rSERT gene expression (Blakely et al., 1991a; Fujita et al, 1993). The regional localization of SERTs in brain has previously been inferred by radioligand autoradiography using SERT antagonists such as imipramine (Fuxe et al., 1983), citalopram (D'Amato et al., 1987a; Duncan et al., 1992), and paroxetine (De Souza et al., 1987; Hrdina et al., 1990; Chen et al., 1992). Although autoradiography has limited cellular resolution, SERT distribution inferred from these studies concurs with the localization of 5HT immunoreactivity (Steinbusch, 1984; Takeuchi, 1988). In our studies, SERT-immunoreactive fibers are prominent in the same brain structures that exhibit high-level SERT antagonist binding. Thus, the cerebellum, which receives only a sparse 5HT innervation and exhibits low-density 3H-antagonist binding, possesses only a few immunoreactive SERT fibers (and undetectable levels of SERT protein on immunoblots). Within the hippocampus proper, the CA2-CA3 region has been shown to display the highest level of 3H-paroxetine and 3H-citalopram binding (Hrdina et al., 1990; Duncan et al., 1992), and similarly this region possesses a greater density of SERT-immunoreactive terminals than does CA1 or the dentate gyrus. The rat brain region with the greatest



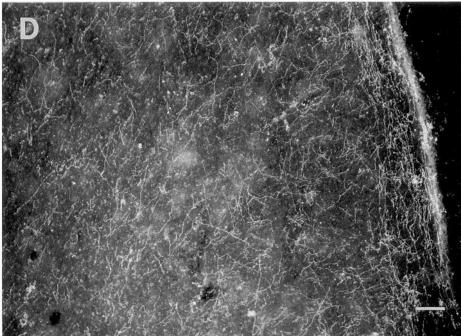


Figure 9. Immunocytochemical localization of SERT in human brain. Brains were fixed by immersion in cold 4% paraformaldehyde in 0.1 м sodium phosphate buffer, pH 7.2, for 24-36 hr prior to sectioning and staining with CT-2 antibody (0.5 μg/ml), as described in Materials and Methods. A, Bright-field photomicrograph of a coronal section through the rostral pons. Note the dense immunoreactivity in the dorsal raphe (DR) and paramedian raphe (PMR). Neurons in the locus ceruleus (LC) are unstained but visible because of neuromelanin content. B, Higher magnification of the boxed area in A, showing examples of immunoreactive neuronal perikarya (arrows) embedded in the densely stained neuropil in the DR. C, Example of immunoreactive varicose fibers in the central gray. D, Dark-field photomicrograph of entorhinal cortex, demonstrating immunoreactive fibers scattered throughout the cortical mantle. The pial surface is on the right and the deep white matter is on the extreme left. Scale bars: A, 500  $\mu$ m; B and D, 50 μm; C 15 μm.

density of SERT antagonist-binding sites has been found to be the dorsal raphe itself (D'Amato et al., 1987; De Souza et al., 1987; Hrdina et al., 1990; Chen et al., 1992; Duncan et al., 1992). Consistent with these data, SERT-immunoreactive cell bodies of the dorsal raphe are surrounded by a dense network of SERT-immunoreactive processes. Indeed, we found that cell body staining was clearly apparent but difficult to demonstrate under standard (4% paraformaldehyde) fixation conditions due to the densely stained neuropil. The addition of low concentrations of glutaraldehyde (0.15–0.3%) diminished fiber staining

and significantly improved cell soma visualization of SERT immunoreactivity.

SERT-like immunoreactivity was abolished by fusion protein preabsorption at concentrations that eliminated visualization of SERT bands detected by immunoblots. Specificity was further evident in an absence of cell body staining in brain regions densely populated with NE and DA neurons that synthesize homologous NET and DAT proteins. SERT-immunoreactive neurons and fibers were evident in both human as well as rat brain, consistent with the high homology of the CT-2 epitope across

species and with the ability of the antibody to immunoblot h-SERT from transfected cells (Fig. 3B) and human brain and platelets (data not shown). Finally, parallel staining in adjacent sections with anti-5HT antisera demonstrated a high correspondence in the cellular and axonal distribution of SERT and 5HT immunoreactivity. The SERT antibody may prove a superior reagent to 5HT-directed antibodies for the identification of sero-tonergic axons and terminals in circumstances, endogenous or induced, where reduced 5HT synthesis and/or storage may occur independently of structural alterations in neuronal processes (Takeuchi et al., 1988; Dewar et al., 1992).

SERT expression has been attributed to cultured and endogenous glial as well as neuronal elements (Kimelberg, 1986; Anderson et al., 1992). As with SERT localization by in situ hybridization, SERT distribution by immunocytochemistry appears to argue against a high level of endogenous glial SERT expression, if any, in the adult rat. Only a single SERT gene appears to be present in humans (Ramamoorthy et al., 1993) and mouse (Gregor et al., 1993); it remains possible that glial 5HT uptake arises from the expression of an additional SERT, generated by alternative splicing and lacking the CT-2 epitope, or that the CT-2 epitope is modified or masked in glia. Lesch and colleagues (Lesch et al., 1993a) have reported the presence of SERT mRNAs, using reverse transcription/polymerase chain reaction (RT-PCR) in regions associated with serotonergic terminals but not cell bodies. The clear ability of RT-PCR to amplify rare mRNAs of insufficient endogenous levels to contribute to functional protein pools (Chelly et al., 1989) suggests caution in attributing functional significance to these data. An alternative explanation for the in vitro glial findings is that SERTs may be expressed by glia early in CNS development but not in the adult, such that primary glial cultures derived from neonatal brain retain transport activity in vitro that eventually they lose in vivo. Finally, expression of 5HT uptake may be induced by culture conditions with little relevance to the pattern of expression in situ (Kimelberg et al., 1992).

In summary, we have achieved the first direct visualization of SERT proteins expressed in transfected and native membranes using SERT peptide and fusion protein-directed antibodies. SERT protein is distributed in a manner consistent with the known localization of 5HT uptake, SERT antagonist binding, and 5HT immunoreactivity. Differential N-glycosylation gives rise to variant SERT proteins in the CNS and periphery. The similar size of endogenous SERT proteins to purified antidepressant-binding proteins, the ability of single cDNAs to induce antidepressant-sensitive 5HT uptake, and the correspondence of SERT anatomical distribution with <sup>3</sup>H-antidepressant autoradiographic localization argue strongly that a single gene product comprises the functional unit of SERTs in vivo. Finally, the conservation of the SERT epitopes utilized for antibody production permits recognition of SERT proteins in both rodent and human preparations, providing an important opportunity to investigate alterations in SERT structure and expression in neurologic and psychiatric disorders.

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