



SHORT COMMUNICATION

$[^3\text{H}]$ WIN 35,428 [2 β -Carbomethoxy-3 β -(4-fluorophenyl)tropane] Binding to Rat Brain Membranes

COMPARING DOPAMINE CELL BODY AREAS WITH NERVE TERMINAL REGIONS

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ABSTRACT. Potential differences between somatodendritic and axonal dopamine transporters were examined by comparing the binding constants of $[^3\text{H}]$ WIN 35,428 [2 β -carbomethoxy-3 β -(4-fluorophenyl)tropane] binding to membranes prepared from the rat ventral mesencephalon, containing A9 and A10 dopamine cell bodies, and from the nucleus accumbens. Saturation analysis of $[^3\text{H}]$ WIN 35,428 binding, in the presence of compounds to occlude norepinephrine and serotonin transporters, was performed by both the “unlabeled” method (varying unlabeled ligand) and “labeled” method (varying radioligand). The density of binding was substantially lower in the ventral mesencephalon than in the nucleus accumbens, but the binding affinity was only slightly different. Likewise, the differences between the two regions in the inhibitory potency of cocaine and GBR 12909 [1-(2-(di(4-fluorophenyl)-methoxy)-ethyl)-4-(3-phenylpropyl)piperazine] were not substantial. The results suggest that somatodendritic and axonal dopamine transporters in the ventral mesencephalon and nucleus accumbens are not very different as far as their binding domains for uptake blockers such as cocaine and GBR 12909 are concerned. *BIOCHEM PHARMACOL* 51;4:563–566, 1996.

KEY WORDS. dopamine transporter; WIN 35,428 binding; rat ventral mesencephalon; rat nucleus accumbens; rat striatum

Central dopaminergic neurons release dopamine not only from their axon terminals, but also from their dendrites, and the released dopamine is inactivated by active reuptake into the dopamine neuron [1–3]. In consonance, autoradiographic studies with high-affinity radioligands show the presence of the dopamine transporter not only in dopaminergic nerve terminals but also in cell body areas [4, 5]. However, it is not clear whether axonal and somatodendritic transporters have the same properties. Thus, $[^3\text{H}]$ dopamine uptake into slices of rat ventral tegmentum is inhibited by various compounds with the same potency as that observed for the inhibition of uptake into NACC§ slices [2]; however, caution is needed when interpreting these results because more recent observations indicate the potential for $[^3\text{H}]$ dopamine to be taken up into serotonin terminals [3, 6]. The binding of $[^3\text{H}]$ WIN 35,428 in monkey substantia nigra, measured by autoradiography in coronal brain sections, is inhibited by various blockers with a rank order

similar to that in slices from caudate nucleus or putamen [7]; for the caudate-putamen, the IC_{50} values for various blockers in sections correlate closely with values obtained with membrane preparations [8]. However, detailed comparative information on the affinity of compounds for somatodendritic and axonal dopamine transporters is not available in the literature. In the present study, membranes were prepared from the VM, containing A9 and A10 dopamine cell bodies, and from the NACC, and the binding of $[^3\text{H}]$ WIN 35,428 was studied in the presence of citalopram and desipramine to occlude serotonin and norepinephrine transporters. Saturation analysis was conducted by varying either the nonradioactive WIN 35,428 (“unlabeled” method) or the $[^3\text{H}]$ WIN 35,428 (“labeled” method), and the inhibitory potencies of cocaine and GBR 12909 were determined.

MATERIALS AND METHODS

Male Sprague–Dawley rats (Harland Sprague–Dawley, Indianapolis, IN) between 8 and 10 weeks of age (225–300 g) were decapitated. For each experiment, the following brain regions were dissected rapidly as described by us previously [3]: the NACC and VM (from six rats, average total tissue weight: 110 mg for NACC and 117 mg for VM) and striatum (from one

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§ Abbreviations: NACC, nucleus accumbens; VM, ventral mesencephalon; GBR 12909, 1-(2-(di(4-fluorophenyl)-methoxy)-ethyl)-4-(3-phenylpropyl)piperazine; and WIN 35,428, 2 β -carbomethoxy-3 β -(4-fluorophenyl)tropane.

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rat, average tissue weight 75 mg) (sufficient for 30 binding assays or 10 conditions in triplicate per region). The tissues were homogenized in 2 mL of ice-cold 0.32 M sucrose with a Teflon-glass homogenizer. The pestle was then rinsed with 3 mL of sucrose solution, and this fluid was combined with the original homogenate, which was the starting material for the preparation of the P_2 fraction. All following steps in preparing and assaying P_2 preparations, and analyses of binding data, were as described previously except where indicated [9]. In the present experiments, P_2 pellets from six rats were homogenized with a Teflon-glass homogenizer in 5.5 mL of assay buffer (30 mM sodium phosphate, pH 7.4, at 25°, 0.32 M sucrose), giving on the average 120 μ g of protein/150 μ L of suspension. The binding assay mixtures contained in a final volume of 0.6 mL approximately 120 μ g of protein (delivered by the 150- μ L aliquot), 90 nM citalopram for blocking serotonin transporters, and 90 nM desipramine for blocking norepinephrine transporters. Saturation analysis by the unlabeled method involved adding increasing concentrations of unlabeled WIN 35,428 to assays containing a fixed concentration of [3 H]WIN 35,428 (2.2 nM for NACC or striatum and 4.5 nM for VM). Unlabeled WIN 35,428 ranged from 0.2 to 40 nM in a total of 8 points. Nonspecific binding was defined with 30 μ M cocaine. When the labeled method of saturation analysis was used, increasing concentrations of [3 H]WIN 35,428, ranging from 0.3 to 115 nM in a total of 7–9 points, were added along with

either 30 μ M cocaine or equivalent water volume. The specific activity of the batch of [3 H]WIN 35,428 used in this work (Lot No. 2824-220) was determined as described by us previously [10] by performing WIN 35,428 inhibition curves at four different radioligand concentrations, and turned out to be 48.2 dpm/fmol. Because the protein concentrations used in the present binding assays were higher (ca. 120 μ g/600 μ L) than used in our previous studies (ca. 25 μ g/200 μ L), control experiments were carried out with various protein concentrations, and it was found that [3 H]WIN 35,428 binding was linear with protein up to 150 μ g/600 μ L, the highest concentration tested.

RESULTS AND DISCUSSION

Saturable binding of [3 H]WIN 35,428 to membranes prepared from the NACC and the VM was best described by a one-site model. Figure 1 shows examples from the unlabeled experiments; data from the labeled analysis were similar. In a region such as the VM where [3 H]WIN 35,428 binding is low, saturation analysis is ideally performed with both the labeled and unlabeled method. The former allows a range of radioligand concentrations starting low, which results in increasing values for bound radioactivity as radioligand levels are increased; at the upper end, cost becomes the major limiting factor. The unlabeled method, on the other hand, allows the study of

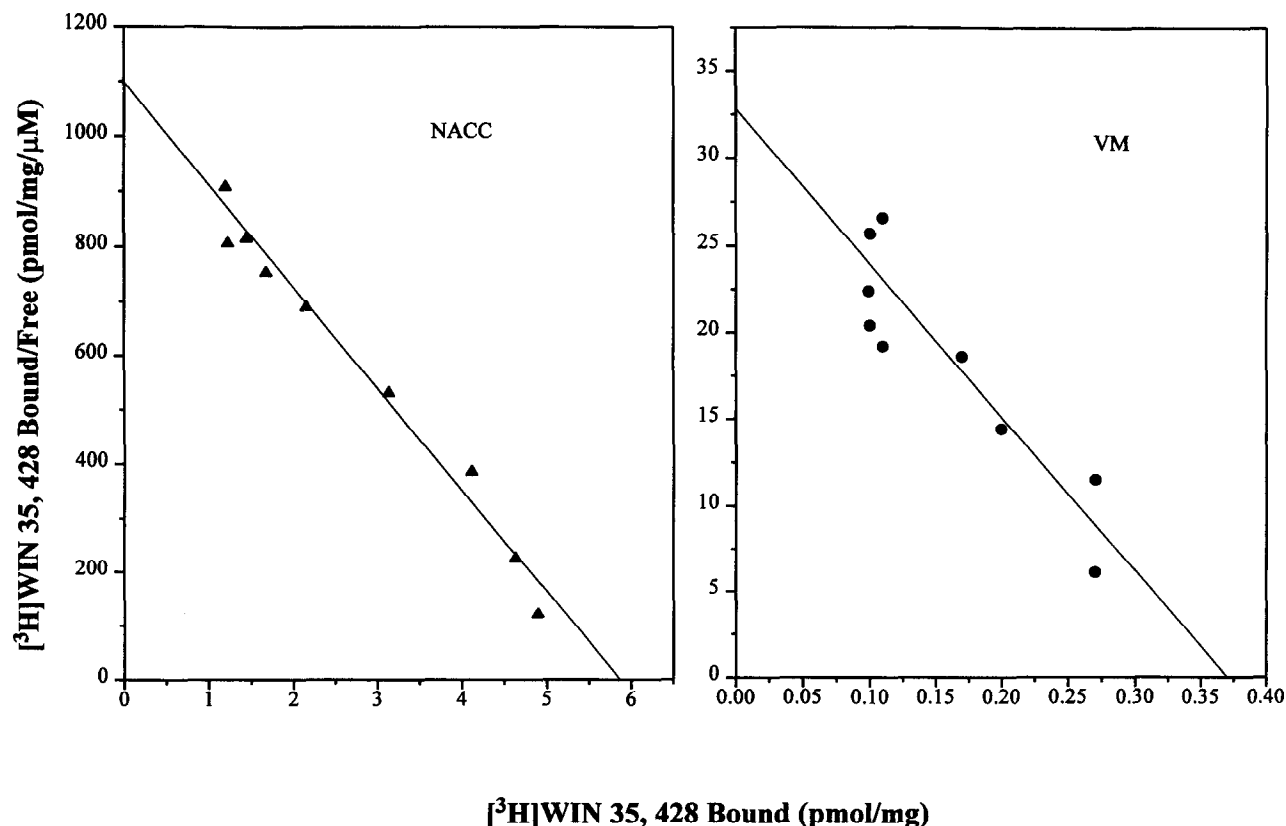


FIG. 1. Saturation analysis by the unlabeled method graphed as Scatchard plots for illustrative purposes. Straight lines represent the best fits chosen by the LIGAND program. A representative experiment is shown that was carried out four times for the NACC and three times for the VM (see Table 1 for averages of all unlabeled and labeled results together).

relatively higher concentrations of nonradioactive ligand but requires a substantial labeling at the fixed radioligand concentration chosen as the starting point (well below the K_d). Under the present conditions for the VM, within the constraints of available brain tissue, the lowest possible radioligand concentration was 0.3 nM with the labeled method (on the average 225 dpm specific binding per assay) and the fixed radioligand concentration was 4.5 nM with the unlabeled method (750 dpm). These low, but detectable, binding values probably underlie the greater variation in the binding data collected for the VM as compared with the NACC. Because the specific activity of the radioligand impacts on the estimation of the binding constants, especially with the labeled method [11], we addressed this experimentally for the batch of radioligand used in this study. We used the competition method for measuring specific activity, as described previously [10].

The results taken together indicated a slightly elevated K_d for [^3H]WIN 35,428 binding in the VM ($P < 0.05$, two-tailed t -test) (Table 1). Cocaine also appeared to have a slightly higher K_i in the VM as compared with the NACC ($P < 0.05$, one-tailed t -test); the Hill numbers were close to unity in both regions. The results obtained for GBR 12909 did not indicate a regional difference in the same direction as for the cocaine-like compounds; if indeed more observations would substantiate the tendency for GBR 12909 to be a stronger, rather than weaker, inhibitor in the VM, it would support the notion of different binding domains involved in GBR- and cocaine-like compounds. The Hill numbers for GBR 12909 tended to be somewhat higher than unity, but this did not reach statistical significance (one-sample, Student's t -test) (Table 1). Overall, the differences in potencies, between regions, were not substantial, suggesting that somatodendritic and axonal dopamine transporters have comparable affinities for cocaine- and GBR-related compounds. In this context, it is of interest that i.v. administered cocaine may be metabolized relatively faster in the substantia nigra of primate brain as compared with other brain regions including caudate-putamen and NACC [12].

TABLE 1. Binding of [^3H]WIN 35,428, cocaine, and GBR 12909 to membranes prepared from NACC and VM

	NACC	VM
[^3H]WIN 35,428		
K_d (nM)	5.2 ± 0.2 (8)	$7.7 \pm 1.2^*$ (6)
B_{\max} (pmol/mg)	3.46 ± 0.44 (8)	$0.53 \pm 0.16^\dagger$ (6)
Cocaine		
K_i (nM)	47.4 ± 0.98 (3)	$66.4 \pm 7.82^\ddagger$ (3)
Hill	0.96 ± 0.03 (3)	0.99 ± 0.08 (3)
GBR 12909		
K_i (nM)	2.8 ± 0.36 (3)	1.9 ± 0.54 (2)
Hill	1.3 ± 0.29 (3)	1.2 ± 0.12 (2)

Values are means \pm SEM ($=0.5 \times$ range where $N = 2$). The number indicated between parentheses denotes the number of independent membrane preparations, with each preparation consisting of pooled tissue from six rats. The K_d and B_{\max} values from the unlabeled and labeled experiments were combined.

* $P < 0.05$ compared with the same variable in NACC (Student's t -test, two-tailed).

$^\dagger P = 0.0001$ compared with the same variable in NACC (Student's t -test, two-tailed).

$^\ddagger P < 0.05$ compared with the same variable in NACC (Student's t -test, one-tailed with alternative hypothesis that K_i is higher in VM).

The present results would suggest that such a diminished activity of cocaine is not compensated for by an increased affinity of cocaine towards somatodendritic dopamine transporters; rather, the slightly reduced affinity would accomplish the opposite. In the present *in vitro* experiments at 0–4°, cocaine is not likely to be metabolized as no metabolism of [^3H]cocaine was found with rodent brain membranes at 21° [13].

The B_{\max} of [^3H]WIN 35,428 binding sites was substantially (7-fold) lower in the VM than in the NACC (Table 1). Most likely, this is partly due to tissue surrounding the A9 and A10 somatodendritic regions that does not contain dopamine transporters, because autoradiographic studies show only an approximately 2-fold difference in dopamine transporter labeling between the ventral tegmental area/substantia nigra (combined in our dissection of the VM) and the NACC. Our attempts to measure [^3H]dopamine uptake into VM synaptosomes in the presence of 90 nM citalopram and desipramine were unsuccessful. The uptake was much lower than anticipated from the reduced density of [^3H]WIN 35,428 sites, indicating, for that region, a relatively lower affinity for dopamine, a lower dopamine uptake velocity, or an increased sensitivity to citalopram/desipramine. The addition of these inhibitors was necessary because [^3H]dopamine can be taken up into serotonergic terminals in dopamine cell body areas [3, 6], and dopamine has a high affinity for the norepinephrine transporter, in fact higher than norepinephrine itself as described for brain preparations by Coyle and Snyder [14] and the cloned norepinephrine transporter by Pacholczyk *et al.* [15].

In addition to the lack of differences in WIN 35,428 binding between cell body and terminal dopamine transporters observed in this study, the affinity of the transport complex for WIN 35,428 was similar in the two different terminal areas of the mesolimbic pathway (the NACC), and the nigrostriatal pathway (the striatum). Thus, the K_d of striatal [^3H]WIN 35,428 binding was 6.8 ± 1.5 nM (mean \pm SEM for three independent preparations, each derived from one rat) when measured under the present conditions, close to the value observed under the same experimental conditions in the NACC (Table 1). The B_{\max} was 9.7 ± 1.2 pmol/mg protein, as expected somewhat higher than that found for the NACC. Other reports have described generally similar properties of dopamine transporters in the striatum and NACC [16, 17]. All data taken together with the present results suggest that the dopamine transporter is similar in the VM, NACC, and striatum.

The observation of Einhorn *et al.* [18] that cocaine is much weaker in suppressing dopamine cell firing when administered iontophoretically into the ventral tegmental area than when injected intravenously, as well as our own findings showing that focal application of cocaine into the ventral tegmental area is less effective in enhancing extracellular dopamine than application into the NACC (manuscript in preparation), are probably not the result of a lower affinity of cocaine for somatodendritic as compared with axonal dopamine transporters. Rather, the present study suggests that other factors should be considered. One is the lower density of dopamine uptake

sites in the ventral tegmental area as compared with the NACC (see above), and another factor is the existence of an inhibitory NACC-ventral tegmental area feedback pathway [18]. The present study does not rule out the possibility that there are regionally dependent differences in monoamine transport functioning beyond the level of substrate, blocker, and Na⁺ recognition.

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