# D<sub>1</sub> Dopamine Receptor Activation of Multiple Transcription Factor Genes in Rat Striatum

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Abstract: Recent studies have shown that dopamine receptor agonists induce expression of Fos-like immunoreactivity in rat striatal neurons. The protooncogene c-fos belongs to a family of immediate early genes that are rapidly induced in fibroblasts by growth factors. In light of previous findings that several immediate early gene mRNAs that encode proven or putative transcription factors are differentially regulated by neuronal stimulation in vivo, we have examined the effect of dopaminergic agents on mRNA levels of several such genes using in situ hybridization and northern blot analysis. d-Amphetamine (2.5-10 mg/kg i.p.) causes a rapid but transient dose-dependent increase in zif268 and jun-B mRNA levels in striatum that was abolished by striatal 6-hydroxydopamine lesions or by pretreatment with the specific D<sub>1</sub> receptor antagonist SCH-23390 but not by specific D<sub>2</sub> receptor antagonists. Apomorphine, a dopamine agonist that acts at both D<sub>1</sub> and D<sub>2</sub> receptors, and SKF-38393, a specific D<sub>1</sub> receptor agonist, produce similar mRNA changes in rats pretreated with either 6-hydroxydopamine or reserpine, whereas LY-171,555, a specific D<sub>2</sub> receptor agonist, has no effect. Direct dopamine agonist effects on these immediate early gene mRNA levels are also blocked by D<sub>1</sub> but not by D<sub>2</sub> antagonists. We observed similar, although less robust, changes in c-fos and fos-B mRNA levels. These results demonstrate that striatal D<sub>1</sub> dopamine receptors are coupled to activation of multiple transcription factor genes, including zif268 and jun-B as well as members of the fos family. Key Words: zif268—jun-B—Immediate early gene—In situ hybridization—Dopamine—Amphetamine—SCH-23390. Cole A. J. et al. D<sub>1</sub> dopamine receptor activation of multiple transcription factor genes in rat striatum. J. Neurochem. 58, 1420-1426 (1992).

Cell surface receptor stimulation by growth factors or neurotransmitters can elicit the rapid activation of transcription factor genes thought to mediate long-term effects of these signaling agents (Greenberg and Ziff, 1984; Lau and Nathans, 1987; Milbrandt, 1987; Sheng and Greenberg, 1990). In recent studies, we and others have demonstrated that synaptic stimulation of Nmethyl-D-aspartate receptors elicits activation of the immediate early gene zif268 (Christy et al., 1988) [also termed NGF1A (Milbrandt, 1987), egr-1 (Sukhatme et al., 1988), and Krox-24 (Lemaire et al., 1988)], a transcription factor containing several zinc finger DNA binding motifs, but not c-fos or c-jun, demonstrating that this group of genes is differentially regulated in vivo (Cole et al., 1989; Wisden et al., 1990). Because several groups have reported that c-Fos-like immunoreactivity in striatum is regulated by D<sub>1</sub> dopamine receptor stimulation (Robertson et al., 1989a,b; Graybiel et al., 1990; Young et al., 1991), we wondered whether other transcription factor genes are also coupled to this neurotransmitter system.

We have used in situ hybridization and northern blot analysis to examine the responses of mRNAs encoding several immediate early genes, including zif268, jun-B (Ryder et al., 1988), and fos-B (Zerial et al., 1989) as well as c-fos, in brain following systemic administration of dopaminergic agents. A preliminary report of these results has been published in abstract form (Cole et al., 1990b). Induction of jun-B and c-fos mRNAs by indirect dopamine agonist treatment has also been reported in abstract form (Iadarola et al., 1990; Moratalla et al., 1990).

#### MATERIALS AND METHODS

### Animal preparation

Male Sprague–Dawley rats (weighing 150–200 g) were used for all experiments. d-Amphetamine sulfate, R(-)-apomor-

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Abbreviations used: MECS, maximal electroconvulsive seizures; SDS, sodium dodecyl sulfate; SSC, saline-sodium citrate.

phine hydrochloride, R(+)-SCH-23390 hydrochloride, S(-)-SCH-23388 hydrochloride, S(-)-eticlopride hydrochloride, (-)-quinpirole hydrochloride (LY-171,555), cocaine hydrochloride, and mianserin hydrochloride were dissolved in distilled water. Haloperidol, pimozide, R(+)-SKF-38393 hydrochloride, and reserpine were dissolved in dimethyl sulfoxide. Drugs were delivered by intraperitoneal injection in volumes of 1  $\mu$ l/g of body weight. At appropriate intervals, animals were killed by decapitation, and the brains were dissected and rapidly frozen. For blocking experiments, antagonists were given 15 min before agonists. For some experiments animals were pretreated with reserpine (5 mg/kg i.p.) 18 and 3 h before administration of the dopaminergic agonist to be tested.

For 6-hydroxydopamine experiments, animals were pretreated with desipramine (25 mg/kg i.p.) 30 min before intracranial injection (Iversen and Kelly, 1975). Under chloral hydrate anesthesia (400 mg/kg i.p.) animals were placed in a stereotaxic frame. Eight micrograms of 6-hydroxydopamine in 2  $\mu$ l of 0.9% NaCl with ascorbic acid (1 mg/ml) was injected into the caudate nucleus using a Hamilton syringe positioned at coordinates 0.2 mm anterior and 3.0 mm lateral to bregma and lowered to 4.7 mm below the dura (Javoy et al., 1975; Paxinos and Watson, 1982). Animals were allowed to recover for 7–10 days before experiments.

Maximal electroconvulsive seizures (MECS) were produced in male Sprague–Dawley rats (weighing 175–225 g) by standard techniques (Swinyard, 1972). Following attachment of saline-soaked earclips, electroshock consisting of a 1-s, 100-Hz, 85-mA stimulus of 0.5-ms square-wave pulses was delivered using a UGO Basile ECT unit (model 7801). At these settings, all animals had generalized tonic–clonic convulsions with hindlimb extension.

#### DNA and RNA probes

Murine zif268, rat NGFI-A, human c-fos, and mouse fos-B cDNA sequences were cloned in Bluescript M13+ (Stratagene, La Jolla, CA, U.S.A.). The murine jun-B cDNA sequence was cloned in p-Gem2 (Promega Biotech, Madison, WI, U.S.A.). For RNA probes, plasmids containing full-length or nearly full-length cDNAs for each of the immediate early genes were linearized with appropriate restriction enzymes to produce either antisense or sense transcripts. <sup>35</sup>S-Labeled RNA sense or antisense probes were prepared from cDNA

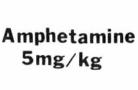
sequences using T7, T3, or SP6 RNA polymerase (Promega Biotech) and uridine 5'-[ $^{35}$ S]( $\alpha$ -thio)triphosphate (New England Nuclear) essentially as described (Melton et al., 1984). For northern blot analysis, inserts were excised from plasmids, gel-purified, and labeled with [ $\alpha$ - $^{32}$ P]deoxycytidine 5'-triphosphate using random primer-initiated synthesis with the Klenow fragment of DNA polymerase (Feinberg and Vogelstein, 1983). Mouse zif268 and rat NGFI-A probes were used interchangeably in these studies and gave similar results.

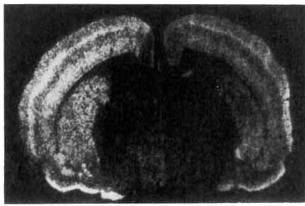
#### Northern blots

Total RNA was isolated from rat hippocampus or striatum using the acid guanidinium-phenol extraction procedure of Chomczynski and Sacchi (1987). RNA was separated by electrophoresis through a formaldehyde-containing 1% agarose gel, transferred to nitrocellulose membranes, and probed essentially as described by Linzer and Nathans (1983). In brief, filters were prehybridized in a sealed bag containing 50% (vol/vol) formamide, 4× saline-sodium citrate (SSC)  $(1 \times SSC \text{ contains } 150 \text{ m} M \text{ NaCl and } 15 \text{ m} M \text{ sodium citrate},$ adjusted to pH 7.0), 4× Denhardt's solution [1× Denhardt's contains Ficoll, polyvinylpyrrolidone, and bovine serum albumin each at 0.02% (wt/vol)], 0.1% (wt/vol) sodium dodecyl sulfate (SDS), 0.1% (wt/vol) sodium pyrophosphate, and 100 μg/ml of sheared salmon sperm DNA. <sup>32</sup>P-Labeled probe was added to a final concentration of  $2 \times 10^6$  cpm/ml, and filters were hybridized overnight at 52°C. Filters were washed in four changes of 2× SSC with 0.1% (wt/vol) SDS at room temperature and then in four changes of 0.1× SSC with 0.1% SDS at 50°C, air-dried, and exposed to x-ray film (Kodak XAR) at -70°C with an intensifying screen for up to 1 week.

#### In situ hybridization

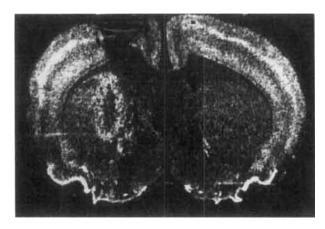
In situ hybridization was performed as described previously (Cole et al., 1990a). Frozen  $10-\mu$ m-thick sections were mounted on gelatin-coated glass slides and stored desiccated at  $-20^{\circ}$ C. Before hybridization, sections were fixed with depolymerized 4% (wt/vol) paraformaldehyde freshly prepared in phosphate-buffered saline, acetylated with 0.5% (vol/vol) acetic anhydride in 0.1 M triethanolamine dissolved in 0.9% (wt/vol) NaCl adjusted to pH 8.0, and delipidated. Hybridization was performed in a moist chamber under unsealed dimethyldichlorosilane (Atomergic Chemetals)-treated coverslips at  $56^{\circ}$ C using  $5 \times 10^{5}$ – $1 \times 10^{6}$  cpm of probe for each





Control

zif/268



**FIG. 2.** Effect of apomorphine on *zif268* mRNA levels in 6-hydroxydopamine-lesioned striatum. Ten days after intrastriatal injection of 8  $\mu$ g of 6-hydroxydopamine into the left caudate nucleus, the *animal* was treated with apomorphine (2 mg/kg i.p.). Thirty minutes after treatment the animal was killed, and the brain was processed for in situ hybridization with a <sup>35</sup>S-labeled *zif268* antisense riboprobe.

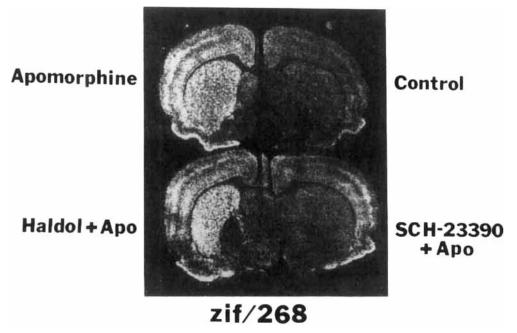
slide in 50  $\mu$ l of hybridization buffer containing 50% (vol/vol) formamide, 10 mM dithiothreitol, 600 mM NaCl, 1× Denhardt's solution, 1 mM EDTA, 0.2 mg/ml of yeast tRNA, and 10% (vol/vol) dextran. After overnight incubation, slides were washed in 2× SSC, treated with 10  $\mu$ g/ml of RNase A (Worthington) in 2× SSC at 30°C, air-dried, and apposed to autoradiography film (Kodak XAR) at room temperature for 1–5 days.

To ensure specificity of in situ hybridization, studies were done using control and experimental brains comounted in the same tissue block to minimize differences in handling of tissue sections, including freeze-thaw cycles and section thickness. No hybridization of sense strand probes was detected. <sup>35</sup>S-Labeled riboprobe complementary to the 3' noncoding region of *zi/268* gave results identical to those obtained using full-length antisense probes. Unless otherwise noted all experiments reported include five or more animals for each condition.

#### RESULTS

In initial experiments, we used in situ hybridization to examine the effects of amphetamine on levels of zif268 mRNA in brain. In animals killed 30 min after administration of amphetamine (2.5-10 mg/kg i.p.), we found an increase in zif268 mRNA levels in the striatum, as well as the overlying cortex (Fig. 1). The rise in zif268 mRNA level is apparent within 15 min of injection, and levels return to near baseline within 3 h. Amphetamine-induced increases in zif268 mRNA levels are seen throughout the caudate and putamen but entirely spare the globus pallidus. Similar results were obtained when we examined responses of jun-B mRNA, whereas less robust increases in levels of c-fos and fos-B mRNAs were observed. Increases in zif268 and jun-B mRNA levels were also seen in striatum 30 min after cocaine treatment (10 mg/kg i.p.).

To determine whether the response to amphetamine requires preserved dopaminergic innervation of the striatum, we used intrastriatal 6-hydroxydopamine to lesion dopaminergic afferents selectively. Ten days after intracerebral injections, animals were given amphetamine (10 mg/kg i.p.) or apomorphine (2 mg/kg). Am-



**FIG. 3.** Effect of dopamine receptor antagonists on apomorphine-induced *zif268* mRNA levels. All animals were pretreated with reserpine (see Materials and Methods). Animals were killed 30 min after saline injection (control), 30 min after apomorphine (2 mg/kg), or 30 min after apomorphine (Apo) following pretreatment with the indicated dopamine antagonist [haldoperidol (Haldol), 1 mg/kg; or SCH-23390, 0.5 mg/kg].

phetamine treatment induced vigorous rotation ipsilateral to the lesion, whereas apomorphine treatment induced rotation contralateral to the lesion, consistent with the establishment of functional receptor supersensitivity. Animals were killed 30 min after agonist treatment. In animals so treated, the zif268, c-fos, and jun-B mRNA response to amphetamine is abolished on the injected side but not on the contralateral side (n = 4). In contrast, treatment with apomorphine causes an increase in each of these mRNA levels in the lesioned striatum but not on the contralateral side (n = 5) (Fig. 2). These results indicate that amphetamine requires preserved presynaptic fibers for its action on this group of immediate early genes and further suggests that the direct dopamine agonist apomorphine can elicit a detectable immediate early gene mRNA response only in denervated striatum. Apomorphine's inability to mimic amphetamine's effect in unlesioned striatum was confirmed in naive animals at doses up to 5 mg/kg and at time points from 15 min to 4 h.

To assess the role of dopamine receptors in mediating amphetamine's effect, we examined whether it was blocked by selective dopamine receptor antagonists. The D<sub>2</sub> dopamine receptor antagonist haloperidol (0.1– 3 mg/kg) or the D<sub>1</sub> receptor antagonist SCH-23390 (0.01-1.0 mg/kg) (Hyttel, 1983) was administered 15 min before amphetamine. We found that whereas haloperidol does not block amphetamine's activation of zif268 or jun-B, SCH-23390 completely prevents these responses at 0.1 mg/kg but not at 0.02 mg/kg. SCH-23390's effect was not shared by its enantiomer SCH-23388 (1 mg/kg i.p.; n = 3), which lacks  $D_1$  receptor antagonist properties (Billard et al., 1984), or by mianserin (1 mg/kg i.p.; n = 3), which like SCH-23390 possesses a high affinity for serotonin 5-HT<sub>1C</sub> and 5-HT<sub>2</sub> receptors (Bischoff et al., 1986; Hess et al., 1986).

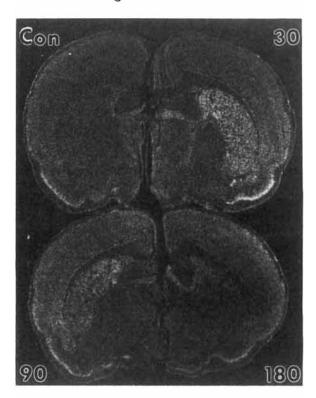
Because these results pointed to a role for D<sub>1</sub> dopamine receptors in activation of zif268 and jun-B, we wondered whether these responses could be potentiated by reserpine pretreatment, which has been found to enhance other responses to D<sub>1</sub> receptor stimulation (Vasse et al., 1990). Rats were pretreated with 5 mg/ kg i.p. of reserpine 18 and 3 h before administration of dopaminergic agents. Amphetamine (2.5-5 mg/kg) produced a robust activation of zif268 and jun-B, as did the nonselective dopaminergic agonist apomorphine (1-2 mg/kg i.p.) (Figs. 3 and 4). The rise in levels of these mRNAs elicited by both of these agents was blocked by SCH-23390 (0.5 mg/kg i.p.) but not by haloperidol (1 mg/kg i.p.), eticlopride (1 mg/kg i.p.), or pimozide (1 mg/kg i.p.), three chemically distinct  $D_2$  receptor antagonists (Fig. 3). We confirmed the findings of Miller (1990) and Dragunow et al. (1990) that  $D_2$  dopamine antagonists activate immediate early gene mRNA responses in naive animals, making it difficult to interpret the failure of these agents to block dopamine agonist effects. As shown in Fig. 5, after reserpine pretreatment the haloperidol effect on zif268

mRNA levels is abolished, confirming the notion that this  $D_2$  dopamine antagonist is inactive in blocking apomorphine's effect after reserpine pretreatment.

To examine further the effects of direct dopamine receptor agonists, we used reserpinized animals to check the effect of the selective  $D_1$  agonist SKF-38393 (1–20 mg/kg) (Anderson et al., 1985) and the selective  $D_2$  agonist LY-171,555 (0.5–2 mg/kg) (Nagahama et al., 1986). Whereas SKF-38393 produced an increase in zif268 and jun-B mRNAs in striatum that was blocked by SCH-23390, LY-171,555 had no effect. Similarly, in animals lesioned with unilateral 6-hydroxydopamine injections, LY-171,555 was inactive (n = 5). As reported above for apomorphine, neither of these agents induced a rise in levels of zif268 or jun-B mRNAs in the absence of reserpine pretreatment.

We used northern blot analysis to confirm the results obtained with in situ hybridization. As shown in Fig. 6, mRNA levels of *zif268*, *jun-B*, *c-fos*, and *fos-B* are elevated at 30 min after apomorphine, with lower levels apparent at 90 min. This profile fits well with that observed by in situ hybridization (Fig. 4). Furthermore, as suggested by in situ hybridization, northern blot

## jun-B



Reserpine + Apomorphine (2 mg/kg)

**FIG. 4.** Time course of *jun-B* mRNA response to apomorphine. Each hemisphere was obtained from a reserpine-treated rat at the indicated time after treatment with apomorphine (2 mg/kg i.p.). The control animal was reserpine-treated and killed 30 min after injection with saline.

SALINE

HALOPERIDOL

1 MG/KG

+

APOMORPHINE
2 MG/KG



HALOPERIDOL 2 MG/KG

Apomorphine 2 mg/kg

FIG. 5. Effect of reserpine pretreatment on  $D_2$  dopamine antagonist-induced increases in zif268 mRNA levels. All animals were pretreated with reserpine (see Materials and Methods). Animals were killed 45 min after the indicated antagonist and 30 min after the indicated agonist treatment. Each hemisphere is from a different animal.

**ZIF/268** 

analysis shows that whereas zif268 and jun-B levels are increased to values comparable to those observed in hippocampus following MECS, increases in c-fos and fos-B levels are smaller.

#### **DISCUSSION**

Elevation of levels of several transcription factor mRNAs, including zif268 and jun-B, by either amphetamine or direct dopamine agonists is blocked by the selective D<sub>1</sub> receptor antagonist SCH-23390, indicating that dopamine D<sub>1</sub> receptor stimulation is required for the rapid and transient elevation of levels of these mRNAs in striatum. In contrast to the striking rise seen in zif268 and jun-B mRNA levels following activation of D<sub>1</sub> dopamine receptors, we see lesser changes in c-fos and fos-B mRNA levels. These findings extend several previous reports of increased Fos-like immunoreactivity (Robertson et al., 1989a,b; Young et al., 1991) and c-fos mRNA levels (Graybiel et al., 1990) to indicate that several transcription factor mRNAs are coupled to D<sub>1</sub> dopamine receptor activation in striatum.

 $D_1$  receptors are coupled to stimulation of adenylate cyclase (Kebabian and Calne, 1979), suggesting that, as demonstrated in cells in vitro, this second-messenger system may mediate the activation of selected immediate early genes (Cho et al., 1989). However, recent studies indicate that  $D_1$  dopamine receptors are also coupled to phosphoinositide turnover via activation of phospholipase C (Felder et al., 1989; Mahan et al., 1990). Accordingly, the intracellular second messenger system mediating the rise in immediate early gene levels mediated by  $D_1$  receptors remains unclear.

Our finding that the indirect dopamine agonists amphetamine and cocaine activate several immediate

early gene mRNAs in the naive animal, but the direct agonists appear to increase mRNA levels only in animals pretreated with reserpine or 6-hydroxydopamine, fits well with previous reports (Robertson et al., 1989a; Graybiel et al., 1990; Young et al., 1991) regarding c-

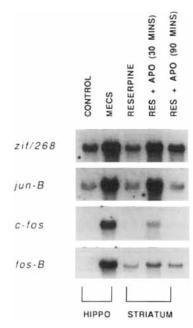


FIG. 6. Comparison of MECS- and apomorphine-induced immediate early gene responses. Total RNA was isolated from hippocampus (HIPPO; lanes 1 and 2) or striatum (lanes 3–5) after the following treatments: lane 1, no treatment (control); lane 2, 30 min after MECS; lane 3, reserpine (5 mg/kg) 18 and 3 h before killing; lane 4, reserpine (RES) pretreatment plus apomorphine (APO; 2 mg/kg) 30 min before killing; and lane 5, reserpine pretreatment plus apomorphine 90 min before killing. Fifteen micrograms of total RNA was loaded in each lane. Replicate filters were prepared and hybridized with the indicated immediate early gene cDNA probes.

Fos-like immunoreactivity following similar treatments. Potential explanations for this apparent disparity include the possibility that direct agonists act as partial agonists that become effective at eliciting this genomic response only after denervation or reserpine treatment, which have been shown previously to potentiate other D<sub>1</sub> receptor-mediated responses. Alternatively, indirect agonists may result in the release of other monoamines that facilitate activation of these genes.

Our observation that fos-B mRNA levels are elevated by amphetamine or direct agonist treatment fits nicely with the recent report of Young et al. (1991) that cocaine treatment induces an increase in several Fos-related antigens as assessed by western blot analysis. Recent studies have demonstrated that two forms of Fos-B generated by alternative splicing are induced in brain following seizure activity. These migrate on SDS-polyacrylamide gel electrophoresis with apparent molecular masses of 44 and 35 kDa (Nakabeppu and Nathans, 1991). Accordingly, it appears likely that the 44-and 35-kDa proteins induced by cocaine (Young et al., 1991) represent alternatively spliced forms of Fos-B.

The striking selective involvement of  $D_1$  receptors in activation of these transcription factor genes suggests that this receptor subtype may be particularly important in mediating long-term effects of dopamine on neuronal function. In this context, it is noteworthy that blockade of D<sub>1</sub> receptors prevents sensitization to amphetamine-induced locomotor effects (Vezina and Stewart, 1989). Furthermore, some of these transcription factor genes could be involved in the marked change in neuropeptide mRNA levels observed following treatment with D<sub>1</sub> receptor agonists (Gerfen et al., 1990). Taken together, these studies suggest that identification of the target genes regulated by these transcription factors in striatum will be critical to understanding long-term effects of amphetamine and cocaine, as well as agents used for dopamine replacement in parkinsonism.

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