The multilingual nature of dopamine neurons

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Abstract

The ability of dopamine (DA) neurons to release other transmitters in addition to DA itself has been increasingly recognized, hence the concept of their *multilingual* nature. A subset of DA neurons, mainly found in the ventral tegmental area, express VGLUT2, allowing them to package and release glutamate onto striatal spiny projection neurons and cholinergic interneurons. Some dopaminergic axon terminals release GABA. Glutamate release by DA neurons has a developmental role, facilitating axonal growth and survival, and may determine in part the critical contribution of the ventral striatum to psychostimulant-induced behavior. Vesicular glutamate coentry may have synergistic effects on vesicular DA filling. The multilingual transmission of DA neurons across multiple striatal domains and the increasing insight into the role of glutamate cotransmission in the ventral striatum highlight the importance of analyzing DA neuron transmission at the synaptic level.

Keywords

1 THE PARTICULAR NATURE OF DOPAMINE NEURONS

Dopamine (DA) neurons of the ventral midbrain, encompassing the substantia nigra (SN), and ventral tegmental area (VTA), number about 450,000 in humans and 10–25,000 in rodents (German et al., 1983; Oorschot, 1996). Despite their small number, they provide a massive innervation of the striatum and also innervate other cortical and subcortical structures (Gauthier et al., 1999; Prensa et al., 2009). Single DA neurons give rise to approximately half a meter of striatal axon in rat, which is so highly branched that a single axon can occupy about 3% of striatal volume (Matsuda et al., 2009). A single SN neuron may possess over 300,000 axon terminals (Arbuthnott and Wickens, 2007; see also Bolam and Pissadaki, 2012).

Ultrastructural studies show that in the striatum of the rat or the monkey, approximately 70% of the dopaminergic inputs forming synapses are established with dendritic shafts, about one-quarter with dendritic spines, and very few with neuronal cell bodies (Descarries et al., 1996; Freund et al., 1984; Smith et al., 1994). However, the fraction of dopaminergic axonal varicosities that contains synaptic membrane specializations is relatively low. Depending on the methods used, estimates range from 20% to 50% in the rat (Antonopoulos et al., 2002; Descarries et al., 1996; Groves et al., 1994; Pickel et al., 1981) and the incidence appears to be even lower in mice (Berube-Carriere et al., 2012). It has therefore been proposed that DA neurons have two types of axonal release sites: some that are synaptic and others, perhaps the majority, that are nonsynaptic or asynaptic (Beaudet and Descarries, 1978; Berube-Carriere et al., 2012; Bérubé-Carriere et al., 2006; Descarries et al., 1980, 2008). Three-dimensional reconstruction of electron micrographs indicates that synaptic vesicles in dopaminergic axon terminals are spread widely throughout the axon, including narrow regions (Gaugler et al., 2012). Release sites in DA neurons are clearly not yet as well characterized as those in the far more studied glutamatergic hippocampal and cholinergic neuromuscular synapses. In DA neurons, DA release may occur throughout the axon at heterogeneous sites that remain to be fully characterized at the molecular and functional levels. The use of fluorescent false neurotransmitters that act as DA analogs may allow visualization of these release sites, independent of their localization and configuration along axons (Rodriguez et al., 2013). In addition, since catecholamines such as DA can be oxidized, the study of DA exocytosis through amperometric techniques has provided important insights into some of the fundamental properties of DA release sites. For example, amperometric measurements of isolated dopaminergic vesicles provided estimates of approximately 30,000 molecules per vesicle (Omiatek et al., 2013), which is higher than the measured quantal size obtained by amperometry on vesicles fusing in intact axons (approximately 10,000 molecules), therefore supporting the hypothesis that

exocytosis can occur by transient fusion (Pereira and Sulzer, 2012; Staal et al., 2004), a concept that remains controversial at other synapses. The rapid kinetics of amperometric measurements have also revealed the duration of the exocytotic events (<100 μs) and provided evidence for multiple transient fusion events (Staal et al., 2004), a phenomenon that could not be resolved with comparatively filtered postsynaptic recordings at conventional ionotropic synapses. In addition, much data suggest that quantal size is a parameter that can be modulated by a wide range of factors including transmitter synthesis, electrochemical gradients, and growth factors (Pereira and Sulzer, 2012; Staal et al., 2004; Sulzer and Pothos, 2000).

Another apparently unusual property of DA neurons is that they have long been known to release DA from their somatodendritic compartment (Geffen et al., 1976; Groves et al., 1975; Korf et al., 1976; Nieoullon et al., 1977), a mechanism that may lead to the activation of somatodendritic autoreceptors and regulate the firing of these neurons and their afferent input. Somatodendritic neurotransmitter release also occurs in other cell types including serotonin neurons (Hery and Ternaux, 1981) and has been well described in the context of neuropeptide release in some systems (Bergquist and Ludwig, 2008). As clusters of synaptic vesicles are essentially absent in dendrites, the most likely candidates for release are tubular structures that express VMAT2 (Nirenberg et al., 1996). The molecular machinery allowing somatodendritic DA release is still incompletely characterized, although a subset of SNARE proteins different from those present in axon terminals appears to be involved (Bergquist et al., 2002; Chen and Rice, 2001; Fortin et al., 2006; Mendez et al., 2011; Witkovsky et al., 2009). Differences between measured somatodendritic and terminal DA release could also involve differences in regional diffusion and reuptake in the VTA versus the striatum (Ford et al., 2010).

## 2 DISCOVERY OF COTRANSMITTERS IN DA NEURONS

DA neurons figure prominently in the elucidation of neurotransmission and cotransmission in the CNS. DA, which was discovered in the brain in 1957 (Montagu, 1957), was actually the first central neurotransmitter molecule to be identified due to its activation of motor responses in mice and rabbits (Carlsson, 1959; Carlsson and Waldeck, 1958). With the burgeoning of immunocytochemistry, DA neurons were more readily identified by the expression of the rate-limiting enzyme in DA production, tyrosine hydroxylase (TH). In addition to their complex axonal and dendritic arborization, allowing multiple modes of DA release, DA neurons have been known for many years to have the potential to release several other neurotransmitters, including neuropeptides, glutamate, and GABA. The physiological significance and pathological implications of these forms of cotransmission in DA neurons are still in the midst of being examined.

The widespread identification of neuropeptides in neurons otherwise using small-molecule, classical neurotransmitters established the idea of cotransmission in the

CNS (Hökfelt et al., 1984). Following the discovery and characterization of the distribution of neuropeptides in the nervous system in the 1980s, it was soon reported that DA neurons of the hypothalamic arcuate nucleus contain neuropeptide Y (Everitt et al., 1986; Harfstrand, 1986), while those of the SN and VTA did not (Murakami et al., 1989). Neurotensin (NT), vasoactive intestinal polypeptide, and cholecystokinin were however reported to be present in subsets of hypothalamic and mesencephalic DA neurons, but with phylogenetic limitations in the case of NT (Hökfelt et al., 1980, 1984; Kalivas et al., 1985; Seroogy et al., 1988; Studler et al., 1988). Neuropeptides are loaded in vesicles at the Golgi apparatus and are thus not released by recycling small synaptic vesicles. Very little is presently known about the regulation of neuropeptide release from DA neurons as well as its physiological and pathological significance.

While the modulatory actions of DA in the CNS were recognized quite early, physiological evidence later suggested that DA neurons also had direct excitatory actions (Wilson et al., 1982). The immunocytochemical visualization of glutamate in DA neurons provided a potential basis for these observations (Ottersen and Storm-Mathisen, 1984). The identification of the glutamate synthetic enzyme glutaminase in DA neurons (Kaneko et al., 1990) further suggested that DA neurons might use glutamate as an excitatory cotransmitter. However, later studies demonstrated that glutaminase is not an ideal and selective marker for glutamatergic neurons, as it is present at varying levels in many neurons and glial cells (Marquez et al., 2009). A new chapter in our understanding of the signaling and connectivity of DA neurons was opened with the report in 1998 of the capacity of rat DA neurons in culture to release glutamate at some of their synaptic contacts (Joyce and Rayport, 2000; Sulzer et al., 1998) (Fig. 1), in a manner similar to serotonin neurons, which had been shown to do the same in 1994 (Johnson, 1994). This capacity of DA neurons to mediate glutamate release was shown *in vitro* to be regulated by D2 autoreceptors and growth factors such as GDNF, suggesting that it can be regulated by some of the same signals that control DA release (Bourque and Trudeau, 2000; Congar et al., 2002; Joyce and Rayport, 2000; Sulzer et al., 1998).

Over the last decade, the identification of the vesicular neurotransmitter transporters has permitted the demonstration that classical neurotransmitters may be cotransmitters (El Mestikawy et al., 2011; Hnasko and Edwards, 2012). Following the cloning of the vesicular glutamate transporters at the turn of the millennium (see El Mestikawy et al., 2011; Fremeau et al., 2004; Omote et al., 2011, for reviews), the ability of DA neurons to release glutamate was explained by demonstrating that they selectively express VGLUT2, one of the three vesicular glutamate transporters (Dal Bo et al., 2004). *In situ* hybridization studies mapped the distribution of VGLUT2-expressing DA neurons (Dal Bo et al., 2008; Kawano et al., 2006; Yamaguchi et al., 2007), as will be discussed in more detail later. Single-cell RT-PCR has also been used to demonstrate expression of VGLUT2 mRNA in TH-positive DA neurons (Birgner et al., 2010; Dal Bo et al., 2004; Fortin et al., 2012; Mendez et al., 2008). The proportion of DA neurons containing VGLUT2 mRNA appear higher in single-cell RT-PCR studies than in *in situ* hybridization studies probably because
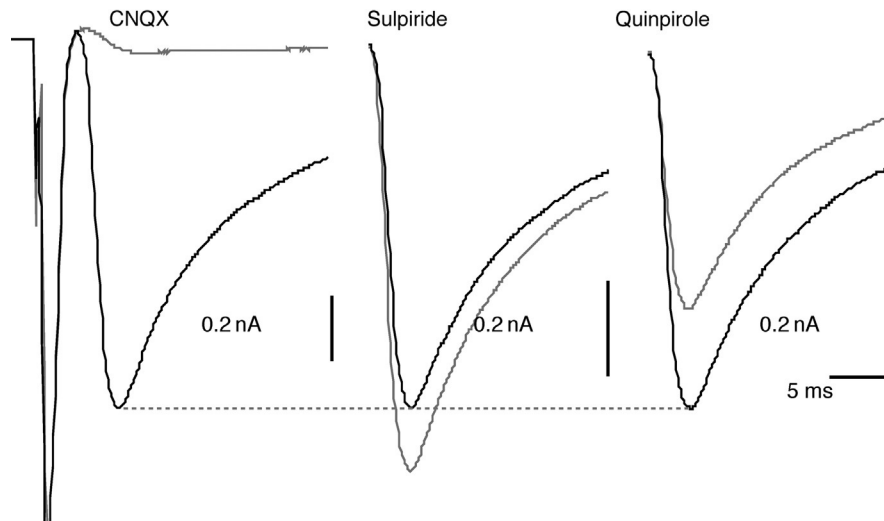
FIGURE 1

Glutamate-mediated synaptic currents recorded in isolated dopamine neurons *in vitro*. In an isolated neuron, subsequently shown to be TH-positive, a large autaptic EPSC was recorded under voltage clamp. This was almost completely blocked by CNQX (EPSC was 4% of control; traces shown are averages of 10 stimulations; traces during drug application are shown in *gray*). The reversible D2 antagonist sulpiride enhanced the EPSC (117%; shown here and in subsequent traces without the initiating action current), whereas the D2 agonist quinpirole markedly attenuated the autaptic EPSC (76%). This suggests that concomitant DA release modulates the glutamate-mediated EPSC.

in situ hybridization does not visualize the low copy numbers of VGLUT2 mRNA found in many TH-positive DA neurons (Li et al., 2013). Interestingly, the expression of VGLUT2 by DA neurons appears to be tightly controlled, as it is upregulated in isolated neurons forming autapses *in vitro* (Dal Bo et al., 2004; Mendez et al., 2008) and downregulated by contact with GABAergic neurons, both *in vitro* and *in vivo* (Mendez et al., 2008). Pathological conditions such as partial 6-OHDA neurotoxic lesions also upregulate the gene in surviving DA neurons (Dal Bo et al., 2008).

Although the initial discoveries of glutamate release and VGLUT2 expression by DA neurons were made *in vitro*, multiple subsequent experiments validated these findings *in vivo* in intact mouse and rat brain. First, extracellular stimulation of dopaminergic cell bodies in the VTA was shown to induce glutamate-mediated synaptic responses in ventral striatal neurons (Chuhma et al., 2004) and in the prefrontal cortex (Lavin et al., 2005). Subsequently, conditional deletion (cKO) of the gene encoding VGLUT2 in DA neurons confirmed that such responses were largely mediated by glutamate release from DA neurons (Hnasko et al., 2010). With the advent of optogenetic techniques, the long-range projections of the major modulatory

neurotransmitter systems became synaptically accessible. It was shown that channelrhodopsin-2 (ChR2)-mediated photoactivation of dopaminergic axons in striatal brain slices prepared from DA transporter (DAT)-Cre mice (Zhuang et al., 2005) injected with floxed ChR2 viral vectors also induced glutamate-mediated synaptic currents in GABAergic spiny projection neurons (SPNs) of the nucleus accumbens (NAcc) (Stuber et al., 2010; Tecuapetla et al., 2010), with comparatively little response detected in the dorsal striatum (Tritsch et al., 2012). The discovery that such responses are preferentially detected in the NAcc is compatible with the localization of most double-phenotype DA neurons to the VTA and not the SN (Fortin et al., 2012; Yamaguchi et al., 2007, 2013). The reported glutamate corelease in the dorsal striatum (Tritsch et al., 2012) may be dependent on the extent of ChR2 expression, striatal subregion (e.g., medial vs. lateral), or the younger ages of the mice studied, since VGLUT2 expression is developmentally downregulated. In these optogenetic experiments, the magnitude of the evoked currents in SPNs was comparatively small, indicating that few ionotropic postsynaptic glutamate receptors may be activated by glutamate release from DA axons (Koos et al., 2011). This suggests that there may be functional roles of glutamate in mature DA neurons other than to drive action potentials in SPNs.

Indeed, more recently, DA neurons have been shown to make more robust glutamatergic connections to cholinergic interneurons (ChIs), specifically in the medial shell of the NAcc (Chuhma et al., 2014). In the dorsal striatum, DA neurons make discrete dopaminergic connections to ChIs, without glutamatergic or GABAergic components (Chuhma et al., 2014). This work suggests that not only do single CNS neurons use multiple neurotransmitters, but also DA neurons can exert diametrically opposite actions on different target neurons, exhibiting regional and postsynaptic cellular target heterogeneity. Further experiments will be needed to define whether single DA neurons projecting to SPNs or ChIs display differential expression of VGLUT2.

Recent work showing GABA-mediated synaptic currents in striatal SPNs (Tritsch et al., 2012) and in lateral habenula neurons (Stamatakis et al., 2013) elicited by optogenetic activation of dopaminergic axons suggests the existence of yet another possible cotransmitter in DA neurons. Surprisingly, this release was noncanonical in the neurons projecting to the striatum, due to an apparent lack of both the GABA synthetic enzyme GAD and the vesicular GABA transporter vGAT (Tritsch et al., 2012). Such a capacity has also been demonstrated in retinal DA neurons (Hirasawa et al., 2012) and in olfactory bulb DA neurons (Liu et al., 2013) and likely occurs in hypothalamic DA neurons, which show extensive GABA immunoreactivity (Schimchowitsch et al., 1991). Remarkably, DA neurons projecting to ChIs in the dorsal striatum do not appear to release GABA (Chuhma et al., 2014), highlighting further the cell-type specificity of DA neuron cotransmission. Further work will be required to delineate the physiological significance of such GABA release, although a role for the dopaminergic GABA projections to the lateral habenula in reward has already been proposed.

3 LOCALIZATION OF GLUTAMATERGIC AND MIXED PHENOTYPE DA NEURONS IN THE BRAIN

The DA neurons giving rise to the mesocorticolimbic system reside in the A10 region of the ventral midbrain, which comprises two major nuclei of the VTA, the parabrachial pigmented (PBP) nucleus and paranigral nucleus (PN), and three midline nuclei, the caudal linear nucleus (CLi), interfascicular nucleus (IF), and rostral linear nucleus (RLi) of the raphe (Swanson, 1982). Within the A10 region, DA neurons are interspersed with GABAergic and glutamatergic neurons, the latter expressing VGLUT2 (Kawano et al., 2006; Nair-Roberts et al., 2008; Yamaguchi et al., 2007). Some of these VGLUT2 neurons make local synapses (Dobi et al., 2010), countering the notion that all glutamatergic inputs to the A10 region are from extrinsic neurons. The relative prevalence of neurons expressing VGLUT2 mRNA in the PBP and PN nuclei of the VTA in the adult rat has been somewhat controversial, as some studies concluded that only very few VGLUT2 neurons were present in the PBP and PN in the adult rat (Kawano et al., 2006), and other studies reported a high prevalence of VGLUT2 neurons in these nuclei in the mature rat (Yamaguchi et al., 2007, 2011). As previously mentioned, such differences may involve the differential sensitivity of the *in situ* hybridization methods used for the cellular detection of the low level of VGLUT2 transcripts (Li et al., 2013).

The visualization of VGLUT2 transcripts in the ventral mesencephalon and brain stem made possible a closer evaluation of the specific localization of DA neurons and other catecholaminergic neurons capable of glutamate cotransmission. Indeed, the glutamatergic signaling capacity of catecholaminergic neurons has received strong support following the demonstration of the coexpression of VGLUT2 mRNA and TH mRNA or protein in neurons of the brain stem (Stornetta et al., 2002) and A10 region (Dal Bo et al., 2008; Kawano et al., 2006; Yamaguchi et al., 2007, 2011) (Fig. 2). Although some of the anatomical studies reported VGLUT2–TH coexpressing neurons to be abundant in the VTA of immature rats, but very rare in the adult rat (Bérubé-Carrière et al., 2009), single-cell RT-PCR studies support the hypothesis of continued expression of VGLUT2 in adult DA neurons in the mouse (Berube-Carriere et al., 2012; Mendez et al., 2008). More recent anatomical and molecular biological studies in the adult rat also provide evidence that VGLUT2–TH coexpressing neurons are present at all levels of CLi, RLi, and IF and are interspersed with neurons expressing VGLUT2 but lacking TH (glutamate-only neurons) in the medial portions of PBP and PN (Li et al., 2013; Yamaguchi et al., 2013). VGLUT2-only and VGLUT2–TH coexpressing neurons each innervate the prefrontal cortex and the NAcc, indicating that in addition to the well-recognized mesocorticolimbic DAergic and GABAergic pathways, there exists a parallel mesocorticolimbic glutamatergic pathway comprising both glutamate-only and glutamate–DA neurons (Gorelova et al., 2012; Hnasko et al., 2012; Li et al., 2013; Yamaguchi et al., 2011).

The type of axon terminals established by VGLUT2–TH coexpressing neurons in their target areas is still relatively unclear. Although, *in vitro*, it has proved relatively
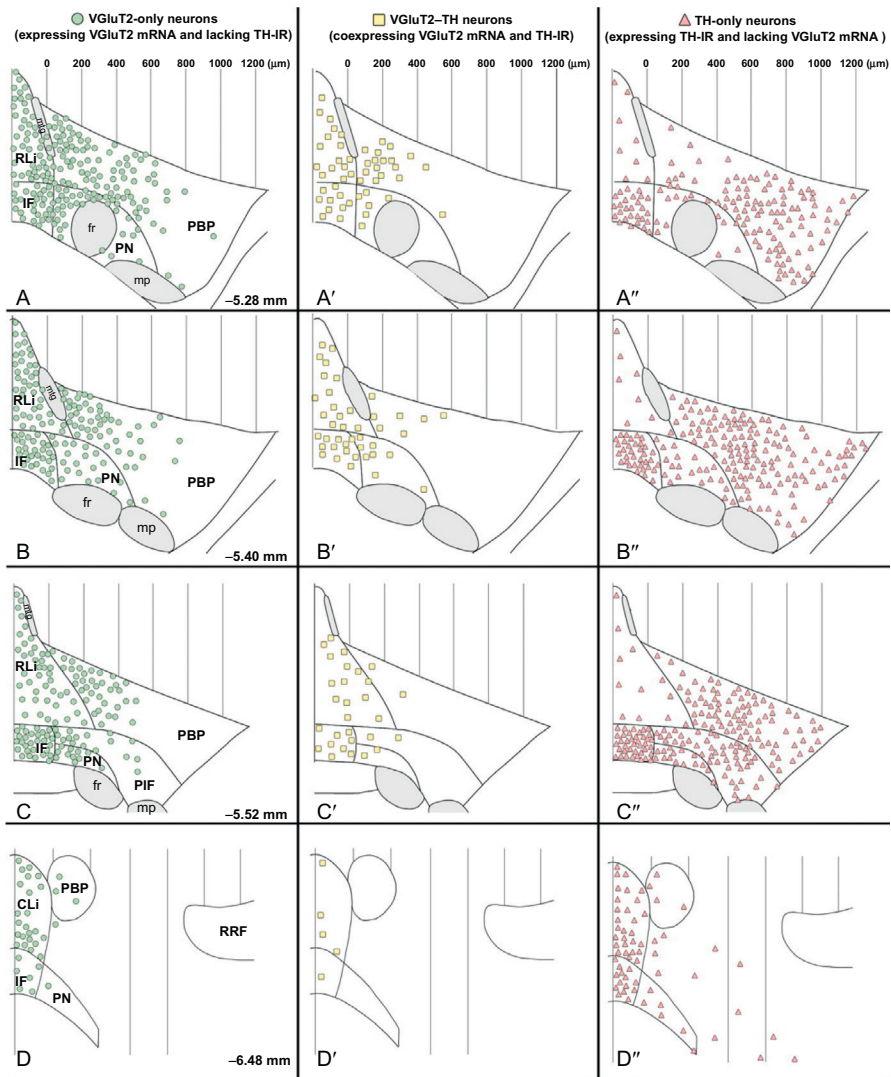
FIGURE 2

Summary diagram of the cellular heterogeneity within the A10 region. Differential distribution of VGLUT2-only neurons, VGLUT2–TH neurons, and TH-only neurons within each subdivision of the A10 region. (A–D) Distribution of VGLUT2-only neurons (neurons expressing VGLUT2 mRNA but lacking TH immunoreactivity). These VGLUT2-only neurons are present throughout the rostrocaudal levels of each subdivision of the A10 region with a lateromedial increasing gradient of concentration. VGLUT2-only neurons are infrequent in the most lateral region of the PBP, a region with a high concentration of TH-only neurons (A″, B″, and C″). (A′–D′) Distribution of VGLUT2–TH neurons (neurons coexpressing VGLUT2 mRNA and TH immunoreactivity). VGLUT2–TH neurons are restricted to the most

easy to detect VGLUT2 protein in TH- or DAT-positive varicosities established by DA neurons (Berube-Carriere et al., 2012; Dal Bo et al., 2004; Mendez et al., 2008), *in vivo*, only sporadic VGLUT2–TH double-labeled varicosities have been detected in the target areas of these neurons, including the NAcc (Berube-Carriere et al., 2012; Fortin et al., 2012; Kawano et al., 2006). Moreover, ultrastructural evidence showed that VGLUT2–TH dual-labeled axon terminals are present in the NAcc of immature rats (Bérubé-Carrière et al., 2009), but absent either in the adult rat (Bérubé-Carrière et al., 2009; Moss et al., 2011) or at any age in mice (Berube-Carriere et al., 2012). Considering the relatively clear evidence for glutamate release in the NAcc in response to optogenetic stimulation of dopaminergic axons of adult mice (Chuhma et al., 2014; Stuber et al., 2010; Tecuapetla et al., 2010), the anatomical findings raise the intriguing hypothesis that double-phenotype DA neurons segregate their glutamatergic and dopaminergic terminals along the same axon in regions such as the NAcc, a possibility that has been considered by a number of groups (Berube-Carriere et al., 2012; Dal Bo et al., 2004; Descarries et al., 2008; El Mestikawy et al., 2011; Fortin et al., 2012; Hattori et al., 1991; Sulzer et al., 1998; S. Zhang and M. Morales, unpublished results) (Fig. 3).

## 4 SYNAPTIC CONNECTIVITY AND PLASTICITY OF GLUTAMATE RELEASE BY DA NEURONS

With global expression of optogenetic probes such as ChR2 in a genetically defined population of neurons and recording systematically from identified target neurons, it becomes possible to define key metrics of synaptic connectivity, including the incidence of connections and their strength. In the initial application of this *functional connectome* approach, restricted expression of ChR2 in striatal SPNs allowed measurements of functional connections of SPNs to principal intrinsic neurons and their projections to the pallidum and ventral midbrain (Chuhma et al., 2011); this revealed distinctive connectivity, with quantitative measures, and projection-specific DA modulation.

DA neuron terminals are the molecular target of psychostimulant action, so it would be anticipated that addictive doses or chronic administration of stimulants

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rostromedial aspects of the PBP and PN; however, they are present at all rostrocaudal and mediolateral levels of the RLi, CLi, and IF. (A″–D″) Distribution of TH-only neurons (neurons lacking expression of VGLUT2 mRNA but containing TH immunoreactivity). The TH-only neurons are present throughout the rostrocaudal levels of each subdivision of the A10 region, including the lateral aspects of the PBP. Each panel represents the average number of labeled neurons found in three sections, each section from a different rat. PIF, parainterfascicular nucleus; mtg, mammillotegmental tract; fr, fasciculus retroflexus; mp, mammillary peduncle; RRF, retrorubral field.

Taken from Yamaguchi et al. (2007).
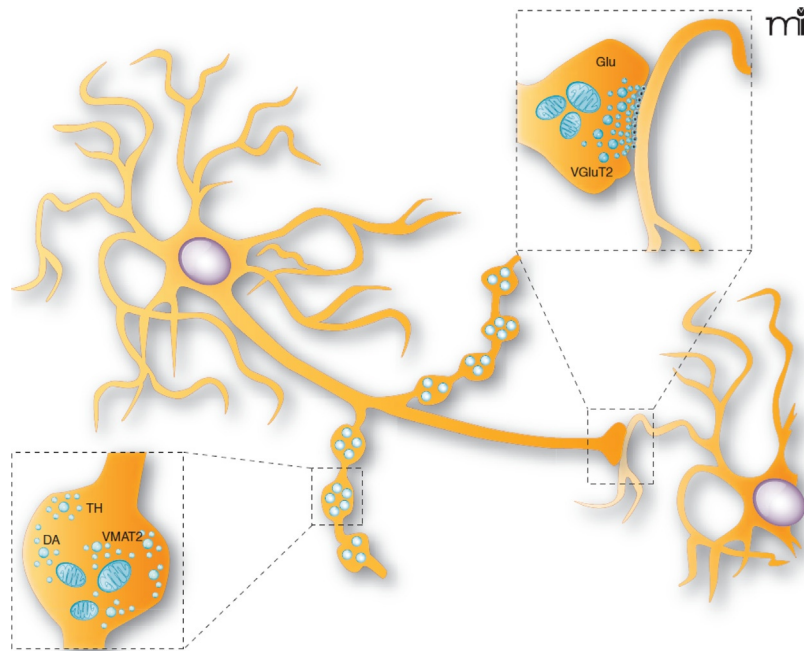
FIGURE 3

Schematic representation of the synaptic and nonsynaptic axon terminals established by dopamine neurons. DA neurons are known to establish two morphologically distinguishable axon terminals: some that are nonsynaptic and others that are synaptic. The nonsynaptic terminals (varicose-like structures) display no obvious pre- and postsynaptic specializations (see lower illustration showing a magnified view of a single nonsynaptic terminal), contain tyrosine hydroxylase (TH), and could be specialized for the release of DA. The synaptic terminals display a more classical active zone, postsynaptic density, and synaptic cleft and could be the site of VGLUT2 expression and of glutamate (Glu) release (see upper illustration showing a magnified view of a synaptic axon terminal). Although not shown, some of the terminals can also contain both TH and VGLUT2.

Taken from Trudeau and Gutierrez (2007).

should affect connectivity. With comprehensive viral transfection of DA neurons with ChR2, a functional connectome analysis can be applied to drug-induced plasticity (Fig. 4). Surprisingly, a recent study revealed that repeated cocaine exposure had minimal impact on DA neuron connections to SPNs (Ishikawa et al., 2013). In contrast, a single dose of amphetamine had profound effects on DA neuron connections to striatal ChIs. A few hours after a single low dose of amphetamine, which induced hyperlocomotion, functional connectome measures showed a profound reduction in DA neuron glutamatergic connections to ChIs in the medial shell of the NAcc (Chuhma et al., 2014). Consistent with the lack of repeated cocaine exposure on connections to SPNs, no impact was seen of amphetamine exposure on DA neuron
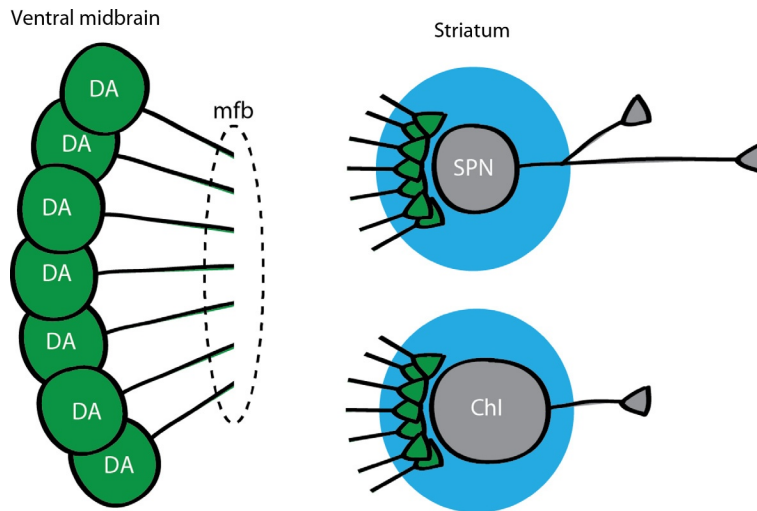
FIGURE 4

Functional connectome analysis of DA neuron connections in striatum. Mice are generated with ChR2 expressed selectively in DA neurons. DA neuron cell bodies are shown in the ventral midbrain (left) with their axons projecting into the median forebrain bundle (mfb) toward the striatum. In the striatum (right), spiny projection neurons (SPNs) and cholinergic interneurons (ChIs) are recorded and DA neuron inputs activated by wide-field photostimulation (larger circle around cells and terminals). With repeated measurements, the incidence and strength of DA neuron inputs to SPNs and ChIs are quantitated to determine functional connectivity.

connections to SPNs. Several hours after a single high dose of amphetamine, which induced stereotypy, dopaminergic connections to ChIs were significantly attenuated. This differential drug-induced plasticity in DA neuron connections shows further how DA neuron transmission can be heterogeneously affected on a dose- and region-dependent manner by drugs of abuse.

5 DEVELOPMENTAL ROLE OF GLUTAMATE RELEASE BY DA NEURONS

Although the developmental pattern of expression of the three VGLUTs has never been extensively examined, initial comparative evaluations of the expression of VGLUT1, VGLUT2, and VGLUT3 in the rat brain revealed that while at birth, expression of VGLUT1 or VGLUT3 was quite low, that of VGLUT2 was already quite high (Boulland et al., 2004; Schafer et al., 2002). Compatible with the finding of early expression of VGLUT2, *in situ* hybridization studies reported high expression of VGLUT2 in the embryonic ventral mesencephalon as early as E12 (Birgner et al., 2010) and E14 (Dal Bo et al., 2008). Single-cell RT-PCR data also support embryonic

and perinatal expression of VGLUT2 in mesencephalic DA neurons (Birgner et al., 2010; Dal Bo et al., 2008; Mendez et al., 2008). Considering these findings and the recent demonstration that glutamate can positively regulate the growth and axonal branching of DA neurons (Schmitz et al., 2009, 2013), a plausible hypothesis is that one of the functions of glutamate release by mesencephalic DA neurons is to promote the growth and connectivity of these neurons during development.

Arguing in favor of this hypothesis, a recent paper reported that cKO of the gene encoding VGLUT2 in DA neurons caused a number of morphological and functional alterations that are compatible with reduced axonal development (Berube-Carriere et al., 2012). The cKO was carried out by using a DAT-Cre mouse (Ekstrand et al., 2007; Zhuang et al., 2005), which, based on the known developmental expression pattern of DAT, should drive Cre recombinase expression and gene deletion approximately by embryonic days 13 or 14 (Fujita et al., 1993; Le et al., 1992; Perrone-Capano et al., 1994). The cKO mice showed reduced numbers of TH-positive neurons in the VTA and SN and a reduced density of TH-positive axon terminals in the NAcc (Fortin et al., 2012). Intriguingly, in a separate study, no change in TH protein or enzymatic activity was detected in NAcc from similar mice (Hnasko et al., 2010). Examined *in vitro*, DA neurons from these mice showed reduced axonal growth (Fortin et al., 2012), compatible with the possibility that glutamate release from growth cones contributes to the growth potential of these neurons. The presence of glutamate NMDA receptors on the growth cones of cultured DA neurons has been demonstrated previously, making the autocrine action of glutamate plausible (Schmitz et al., 2009). Further work will be required to determine whether glutamate release from growth cones induces growth through autocrine activation of receptors on the growth cone itself or through activation of glutamate receptors on target cells, such as astrocytes, that could provide retrograde trophic factor support.

The VGLUT2 cKO also led to reduced action potential-evoked DA release in the NAcc (Fortin et al., 2012; Hnasko et al., 2010).This decreased release could result from the reduced number of DA neurons and reduced axonal growth. Intriguingly, although the number of DA neurons was reduced both in the VTA and in the SN, DA release was only reduced in the NAcc and not in the dorsal striatum (Fortin et al., 2012). One possibility that remains to be evaluated is that the reduced number of DA neurons in the SN was due to loss of the small contingent of SN neurons that project to the NAcc (Lynd-Balta and Haber, 1994a,b; Prensa and Parent, 2001; Prensa et al., 2009). In a separate cKO mouse, potassium-evoked DA release was found to be reduced in both the NAcc and the dorsal striatum (Alsiö et al., 2011). An alternate explanation of the reduced DA in these cKO mice is that the loss of vesicular synergy (El Mestikawy et al., 2011; Gras et al., 2008) between VGLUT2 and VMAT2 leads to reduced packaging of DA inside vesicles, as recently suggested (Hnasko et al., 2010, 2012). As previously mentioned, evidence for the partial segregation of VGLUT2 in a distinct set of axon terminals in DA neurons (Berube-Carriere et al., 2012; Dal Bo et al., 2004; Fortin et al., 2012; Sulzer et al., 1998) may argue against contribution of vesicular synergy in the adult, at least at some

presynaptic sites. The reduced number of TH-positive DA neurons and terminals in VGLUT2 cKO mice could be due to the degeneration of DA neurons lacking the trophic signaling provided by glutamate. Further work will be required to test this and to consider alternate possibilities such as that the reduced number of neurons results from downregulation of the dopaminergic phenotype of these neurons.

6 POSSIBLE CONTRIBUTION OF VGLUT2 IN DA NEURONS TO VESICULAR SYNERGY

Vesicular neurotransmitter loading is dependent on the transport of protons (H^+) into the vesicle lumen mediated by the vacuolar H^+-ATPase (vATPase), thereby establishing a H^+ electrochemical gradient ($\Delta\mu_{H+}$) composed of a chemical gradient (ΔpH) and membrane potential ($\Delta\Psi$). However, the vesicular accumulation of different transmitters by distinct transport mechanisms is differentially dependent on ΔpH versus $\Delta\Psi$ (Edwards, 2007). DA and other substrates of the VMAT2 rely heavily on ΔpH, while glutamate transport through the VGLUT is dependent most on $\Delta\Psi$ (Fig. 5A). Because the stoichiometry of the VMAT2 requires the exchange of two lumenal H^+ for one charged cytosolic DA^+, DA accumulation results in depletion of ΔpH relative to $\Delta\Psi$; efficient DA filling thus requires mechanisms to dissipate $\Delta\Psi$ (Johnson et al., 1981; Knoth et al., 1981). One such mechanism could involve glutamate; because glutamate is an anion, its entry through the VGLUT dissipates $\Delta\Psi$ leading to secondary activation of the vATPase and increased ΔpH (Fig. 5B) (Maycox et al., 1988; Tabb et al., 1992).

Because vesicular monoamine loading is principally dependent on ΔpH and because glutamate loading increases ΔpH, the potential for glutamate coentry to increase DA quantal content has been suggested (Fig. 5C). This mechanism requires localization of VGLUT2 and VMAT2 in the same synaptic vesicles. Consistent with this possibility, vesicular coimmunoisolation of VMAT2 with VGLUT2 has been demonstrated in rat NAcc (Hnasko et al., 2010). It was further shown that glutamate selectively increases uptake of serotonin into synaptic vesicles isolated from the NAcc of 3-week-old rats, providing biophysical evidence that vesicular glutamate coentry contributes to an increase in vesicular DA filling in a native vesicle preparation. Similar experiments have not yet been performed in synaptic vesicles prepared from mice or from adult rats. Along with a developmental impairment in the mesolimbic DA pathway, discussed in the preceding text, vesicular synergy provides an alternate explanation for the reduction in NAcc DA tissue content, evoked DA release, and psychostimulant-induced locomotion in cKO mice lacking VGLUT2 selectively in DA neurons (discussed in the succeeding text) (Alsiö et al., 2011; Birgner et al., 2010; Fortin et al., 2012; Hnasko et al., 2010). Importantly, a similar role for vesicular glutamate coentry promoting the loading of other cationic transmitters has been observed, suggesting this mechanism may be general (Amilhon et al., 2010; El Mestikawy et al., 2011; Gras et al., 2008; Hnasko and Edwards, 2012).
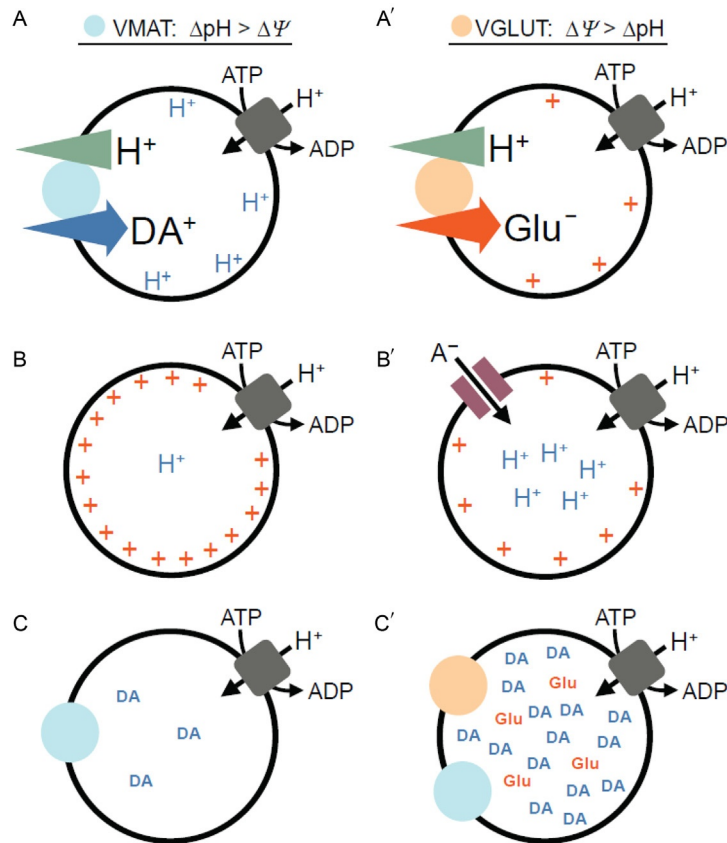
FIGURE 5

Vesicular coloading of glutamate with dopamine leads to synergistic effects on vesicle filling. Vesicular neurotransmitter loading is dependent on both the ΔpH and $\Delta\Psi$ established by the vATPase. However, due to both the stoichiometry of the transporters and the opposite charges of the substrates, (A) dopamine loading through VMAT is dependent mostly on ΔpH, while (A') glutamate uptake through VGLUT depends principally on $\Delta\Psi$. Although the efficient filling of vesicles with dopamine relies on ΔpH—in the absence of a counterion (B)—the transport of H$^+$ by the vATPase sets up a large $\Delta\Psi$ that impedes vesicle acidification. The generation of a large ΔpH therefore requires dissipation of $\Delta\Psi$, for example, by the (B') entry of a permeant anion. Chloride anions are abundant at the nerve terminal and serve this function. However, glutamate anions are also present, and their vesicular entry through VGLUT can facilitate acidification (i.e., increase ΔpH) of dopamine-containing synaptic vesicles. (C) This model predicts increased dopamine quantal content for a subset of (C') VGLUT2- and VMAT2-cocontaining vesicles. Abbreviations: ΔpH, vesicular proton gradient; $\Delta\Psi$, vesicular membrane potential; H$^+$, protons; A$^-$, anion; DA, dopamine; Glu, glutamate.

Although the colocalization of VGLUT2 and TH by dual immuno-EM in DA terminals of young (P15) rats (Dal Bo et al., 2008; Descarries et al., 2008) is consistent with the possibility that VGLUT2 and VMAT2 colocalize in vesicles in a subset of dopaminergic axon terminals in young rats, the lack of TH and VGLUT2 colocalization in young mice and in adult rats or mice (Berube-Carriere et al., 2012; Bérubé-Carrière et al., 2009; Moss et al., 2011) argue that vesicular synergy may be a developmentally or species-restricted phenomenon. Finally, it is possible that L-dopa synthesized by TH diffuses to neighboring axon terminals lacking TH or DAT where it is released after conversion to DA. Further experiments will be required to explore the possibility that vesicular colocalization and vesicular synergy may be restricted to subpopulations of DA neuron terminals, on the developmental stage, and on species.

7 DOES GLUTAMATE CORELEASE MEDIATE A REWARD-RELEVANT SIGNAL?

Several experimental approaches have demonstrated that phasic increases in mesencephalic DA neuron firing occur in response to unexpected rewards or to reward predictive cues; such observations have led to the influential hypothesis that phasic DA responses encode a reward-prediction error (Schultz, 2013; Schultz et al., 1997). The cellular and molecular mechanisms by which released DA shape learning remain less clear but involve neuromodulatory effects on excitatory inputs to the striatum and other regions. Although DA-mediated synaptic effects can sometimes be relatively fast (a few tens of ms) (Beckstead et al., 2004; Chuhma et al., 2014; Gantz et al., 2013; Schmitz et al., 2002), postsynaptic actions mediated by metabotropic DA receptors are nonetheless generally considered slower in onset and duration compared to the very rapid effects of ionotropic neurotransmitter receptors. The utility of DA to mediate a temporally precise postsynaptic reward-prediction error signal has therefore been questioned (Lapish et al., 2007; Seamans and Yang, 2004), and testing the potential for glutamate corelease as mediating a reward-relevant fast excitatory synaptic signal has driven much of the recent work on the possible functional roles of the glutamate cophenotype of DA neurons.

To define the functional consequences of glutamate release by DA neurons, several laboratories have employed a conditional Cre–Lox strategy to disrupt the gene encoding VGLUT2 (reviewed in Wallén-Mackenzie et al., 2010) and hence vesicular glutamate packaging and release, specifically in DA neurons. In each case, cKO mice carrying floxed *Slc17a6* (i.e., *VGLUT2*) alleles were crossed with mice expressing Cre recombinase under the control of *DAT* (*Slc6a3*) regulatory sequences (Alsiö et al., 2011; Birgner et al., 2010; Fortin et al., 2012; Hnasko et al., 2010). Although the precise mutant mouse lines and background strains have varied across laboratories, several key results have been independently corroborated and indicate that VGLUT2 expression in DA neurons shapes behavioral responses to psychostimulants and other reinforcers.

Acute behavioral responses to the psychostimulants amphetamine and cocaine, which elicit hyperlocomotion, were blunted in cKO mice (Birgner et al., 2010;

Fortin et al., 2012; Hnasko et al., 2010). Notably, when tested with higher doses of amphetamine, behavioral activation was still blunted, but the distance traveled was actually higher in cKO than control mice, suggesting that cKO mice are less sensitive to the stereotypy inducing effects of higher-dose amphetamine (Birgner et al., 2010). In a cocaine (and morphine)-conditioned place preference paradigm, a commonly used test for measuring the reinforcing properties of drugs of abuse, cKO mice did not display any deviations from controls (Hnasko et al., 2010; Å. Wallén-Mackenzie, unpublished). However, on operant self-administration tasks, cKO mice consumed higher levels of both sugar and cocaine and displayed a significant elevation in cue-induced drug seeking, a behavioral model of relapse (Alsiö et al., 2011). The cKO mice show no significant differences in cognition measured in the radial arm maze, anxiety measured in the elevated-plus test, or depression measured in the forced swimming test (Birgner et al., 2010; Fortin et al., 2012). However, in the rotarod test of motor coordination, one study found a decrease in performance (Fortin et al., 2012), while others did not (Birgner et al., 2010; Hnasko et al., 2010).

cKO mice thus show diminished behavioral responses to acute and repeated psychostimulant administration but display heightened reward seeking for both sugar and cocaine. These behavioral changes appear to be consistent with the reductions in DA stores and release seen in the cKO (Alsiö et al., 2011; Fortin et al., 2012; Hnasko et al., 2010) and may indicate that in the absence of VGLUT2, rewards produce a weaker DA response and are consequently more sought-after. An alternate possibility is that the changes in drug intake result indirectly from developmental alteration of the mesotelencephalic DA system due to embryonic deletion of the gene encoding VGLUT2, as suggested by biochemical assays revealing multiple adaptations including altered DA receptor levels (Alsiö et al., 2011). However, it remains to be definitively established whether these behavioral perturbations are due to the effects of VGLUT2 on DA packaging, direct alterations in glutamate signaling, developmental changes, or some combination thereof.

In conclusion, the work discussed in this chapter suggests that DA neurons establish a highly heterogeneous set of axonal release sites, allowing the release of DA, glutamate, GABA, or combinations thereof onto target neurons at different sites in the brain, adding further dimensions to the multilingual nature of DA neuron transmission. Further work will be required to extend our understanding of the fine localization and function of individual release sites established by individual DA neurons. While glutamate release by DA neurons contributes to the acute response of animals to natural rewards and psychostimulant drugs, additional work will be required to establish the specific mechanisms involved.

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