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## Neurobehavioral Evidence for Changes in Dopamine System Activity During Adolescence

Dustin Wahlstrom<sup>1,3</sup>, Tonya White<sup>2,3</sup>, and Monica Luciana<sup>1,3,\*</sup>

<sup>1</sup>Department of Psychology, University of Minnesota, Minneapolis MN 55455 USA

<sup>2</sup>Department of Psychiatry, University of Minnesota, Minneapolis MN 55455 USA

<sup>3</sup>Center for Neurobehavioral Development University of Minnesota, Minneapolis MN 55455 USA

### Abstract

Human adolescence has been characterized by increases in risk-taking, emotional lability, and deficient patterns of behavioral regulation. These behaviors have often been attributed to changes in brain structure that occur during this developmental period, notably alterations in gray and white matter that impact synaptic architecture in frontal, limbic, and striatal regions. In this review, we provide a rationale for considering that these behaviors may be due to changes in dopamine system activity, particularly overactivity, during adolescence relative to either childhood or adulthood. This rationale relies on animal data due to limitations in assessing neurochemical activity more directly in juveniles. Accordingly, we also present a strategy that incorporates molecular genetic techniques to infer the status of the underlying tone of the dopamine system across developmental groups. Implications for the understanding of adolescent behavioral development are discussed.

### Keywords

Dopamine; COMT; Brain Development; Adolescence; Working Memory

### Introduction

Adolescence is a transitional period between childhood and adulthood characterized by behavioral, hormonal, and neurochemical changes designed to prepare organisms for independent survival (Casey et al., 2008; Doremus-Fitzwater et al., 2010; Spear, 2003). Although these transitions are largely positive, adolescence can be conceptualized as a period of vulnerability. Risk-taking increases during this time (Steinberg, 2008), as does vulnerability to psychiatric disorders (Kessler et al., 2005; Paus, Keshaven & Giedd, 2008). Theories abound to explain these patterns, most of which suggest that adolescent behavior patterns are attributable to changes in brain maturation during this period of the lifespan (Casey et al., 2008; Fareri et al., 2008; Steinberg, 2008). Many frameworks focus on one of two neural substrates, considering one or both. These include (a) the prefrontal cortex (PFC), which is structurally and functionally under-developed, leading to deficiencies in behavioral regulation,

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\*Corresponding author: Department of Psychology, 75 East River Road, University of Minnesota, Minneapolis MN 55455; phone: 612-626-0757; fax: 612-626-2079; lucia003@umn.edu

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as well as (b) limbic and striatal structures, which may be characterized by relatively heightened patterns of activation, conferring a state of motivational “over-drive”. Simplistically put, it is hypothesized that “go” signals are strong, while regulatory ‘stop’ or monitoring signals are weak. These processes may be inter-related in that sub-cortical signaling can be excessive due to a lack of overarching cortical control. Accordingly, it is unknown whether adolescents’ behavior patterns are due to deficiencies in the PFC’s structural and functional maturation, to a subcortical system that is in over-drive and could not be controlled even in the presence of an adequately functioning PFC, or to some combination of both factors. It should also be mentioned at the outset that not all models adhere to the formulation that motivational circuits are in a state of over-drive; some propose that the opposite is true (Comings & Bloom, 2000; Volkow et al., 2007). Although the brain is undergoing structural refinement during adolescence, adolescents’ difficulties with behavioral regulation may be exacerbated by other neurodynamic processes.

We have offered a unifying account of adolescent behavior that explains it in terms of the development of the dopamine (DA) system (Wahlstrom et al., 2010). Briefly, our perspective builds upon the theory that DA underlies a behavioral activation system that modulates incentive-motivated approach behavior (Depue & Collins, 1999). This system promotes reward-seeking through activity in limbic, striatal, and frontal networks, facilitating an individual’s ability to translate positive motivations into adaptive actions. The adaptive pursuit of positive incentives is critical to independent future-directed behavior. Our perspective is that activity in this system increases during adolescence to meet the demands associated with the transition to independent living. The increase occurs via a tonic increase in DA availability and impacts both subcortical (limbic and striatal) and cortical (prefrontal) circuits.

Operating within the context of continued structural brain development, heightened DA activity within this system may result in an apparent over-activation of incentive motivation in the absence of reliable levels of behavioral control. Unfortunately, the assessment of neurochemistry in human adolescents has proven elusive due to methodological limitations and human subjects concerns. The purpose of the current review is to describe developmental changes in the DA system through adolescence and to suggest a genetic model for how the integrity of the system can be assessed non-invasively in healthy adolescents. In addressing this aim, we will (i) provide an overview of brain development during adolescence; (ii) provide a brief overview of the DA system from a neurophysiological perspective, (iii) review the development of the DA system, presenting data from the animal literature; this review represents the major portion of this paper and is intended to provide a rationale for why this system should be scrutinized in humans for its role in adolescent behavior; finally, we will (iv) present a molecular genetic strategy for the indirect assessment of human neurochemical development. We have chosen DA, the COMT Val<sup>158</sup>Met single nucleotide polymorphism, and working memory to illustrate this strategy. Although working memory does not necessarily reflect the risk taking behaviors that are of primary interest in adolescents, it was chosen because there are well-established literatures regarding inter-relationships between it, COMT, and dopamine, allowing for more straightforward developmental inferences. However, similar models could be derived for other behavioral domains, particularly those associated with the processing of motivational signals, and other neurochemical systems.

## An Overview of Behavioral and Brain Development in Adolescence

Much of the literature on adolescent brain development focuses on the development of prefrontal systems, a topic that can be considered from both behavioral and brain imaging perspectives. In terms of behavioral approaches, executive functions that have been linked to prefrontal substrates are evident late in infancy, coincident with independent locomotion (Bell & Fox, 1992; Diamond, 1990a,b) and show a steady course of improvement through childhood

(Luciana & Nelson, 2002; Kerr & Zelazo, 2004; Levin et al., 1991; Ridderinkoff et al., 1997; Welsh & Pennington, 1991). These functions primary include rudimentary working memory skills, behavioral approach in the face of interference, and flexibility in shifting responses in response to feedback contingencies. With respect to all of these domains, studies conducted over the past several decades have generally concluded that adolescence is a period of continued PFC development. Adolescence is characterized by the continued maturation of a wide variety of PFC-mediated behaviors including planning (Asato et al., 2006; Luciana et al., 2009), concept formation (Chelune & Baer, 1986), working memory (Conklin et al., 2007; DeLuca et al., 2003; Luna et al., 2004; Luciana et al., 2005), inhibitory control (Hooper et al., 2004; Luna et al., 2010), delay discounting (Olson et al., 2007; Steinberg et al., 2008) and motivated decision-making as measured by gambling tasks (Crone & van der Molen, 2004; Hooper et al., 2004; Overman, 2004; Cauffman et al., 2009). When the age of maturation is reached varies by task demand (Luciana et al., 2005; Steinberg et al., 2009) and may depend on the amount or complexity of information that has to be processed within a given context (Luciana et al., 2005). Depending on the task, performance improves into the late teens and perhaps early twenties (DeLuca et al., 2003; Luciana et al., 2009; Steinberg et al., 2009). An exhaustive review of this literature is outside the scope of the current review, but this sampling of studies is representative of the literature as a whole in terms of the finding that executive skills continue to develop during adolescence.

These maturational trajectories are generally interpreted as reflecting the structural maturation of various tissue compartments that comprise the prefrontal cortex and associated networks. For instance, human neuroimaging studies indicate decreases in cortical gray matter during adolescence (Giedd et al., 1999; Giedd, 2004; Gogtay et al., 2004; Gogtay & Thompson, 2009; Jernigan, Trauner, Hesselink, & Tallal, 1991; Luders et al., 2005; Thompson et al., 2005; Sowell et al., 2003). Collectively, these studies indicate that the total volume of gray matter increases across the cortex prior to puberty (generally inferred by the age of the participant sample), reaching a peak somewhere in the early-to-mid pubertal period after which there is a post-pubertal decline. The post-pubertal changes in gray matter are non-linear and vary with respect to brain region. For instance, a recent longitudinal study, using time-lapsed images of participants studied repeatedly over time, indicates that the primary sensorimotor cortices and the frontal and occipital poles mature first. The remainder of the cortex develops in a parietal to frontal direction with the superior temporal cortex maturing last (Gogtay et al., 2004). Gray matter volumes may peak earlier in males versus females in most cortical regions (Lenroot & Giedd, 2010). The decline in gray matter that occurs after these peak values are reached is generally attributed to the pruning (elimination) of synaptic contacts.

As this elimination is occurring, there are concomitant increases in white matter volume and density (Bartzokis et al., 2008; Giedd et al., 1999; Paus, 2001; Pfefferbaum et al., 1994). These age-related increases appear to be more linear as compared to the non-linear changes in gray matter volumes. Indeed, white matter volumes accelerate through adolescence and through early-to-middle adulthood. As with gray matter, there is a suggestion of sexual dimorphism, with males showing steeper increases in white matter volumes across adolescence as compared to females (Paus, 2010). In addition to the assessment of white matter volumes, age-related increases in the directional organization of white matter, as assessed through diffusion tensor imaging (DTI), have been observed (Ashtari et al., 2007; Barnea-Goraly, 2005; Ben Bashat et al., 2005; Bonekamp et al., 2007; Eluvathingal, et al., 2007; Klingberg et al., 1999; Hasan et al., 2007; Lebel et al., 2008; Li & Noseworthy, 2002; Muetzel et al., 2008; Qui et al., 2008; Schmithorst & Yuan, 2009; Schneider, et al., 2004; Snook et al., 2005, 2007; Zhang et al., 2005). DTI measures the relative diffusion of water within voxels that have been characterized on the basis of automated tissue parcellations as white matter. Based on the observed diffusion patterns, inferences are made regarding the structural properties of the white matter within which the water molecules are located, given that water cannot diffuse in an unobstructed

manner in the presence of barriers to diffusion (Basser et al., 1994). Directionally oriented fibers represent one such barrier. Overall, these studies indicate that with increasing age, diffusion decreases in a manner that is directionally sensitive. These decreases have been attributed to increases in myelination or axonal caliber (Paus, 2010), supporting the notion that functional networks are being sculpted in a manner that increases information processing efficiency. Moreover, they indicate that with increasing development, frontal systems become increasingly capable from a functional neuroanatomical perspective of assuming greater regulation over subcortical, primary sensory, and motor processing (Eluvathingal et al., 2007).

Diffusion tensor imaging studies of adolescents have suggested that increases in white matter integrity are associated with better performance on measures of general intelligence (Schmithorst, 2005; Shaw et al., 2006), delay discounting (Olson et al., 2009), interhemispheric transfer (Muetzel et al., 2008), working memory (Nagy et al., 2004; Olesen, et al., 2003), inhibitory control (Liston et al., 2006), and reading ability (Qiu et al., 2008). In a departure from this general pattern, Berns et al. (2009) reported that as risk-taking tendencies increase during adolescence, frontal lobe white matter integrity is *better* developed, a pattern that was interpreted as suggesting that risk-takers have more mature brains. Despite this inconsistency, these studies suggest regional associations between white matter development and behavior and also support, to a more limited extent, the notion that the structural integrity of broadly distributed functional networks supports behavioral development. This latter conclusion is more tentative, because studies have not always considered the extent to which behavior-white matter associations change with age. Because white matter integrity underlies connectivity between regions, these studies are particularly important in suggesting the manner in which functional networks mature across development.

In a more direct assessment of this functional capacity, several groups have used functional MRI to document developmental differences in brain activation patterns. These studies have found age-related differences in orbitofrontal cortex (OFC) and ventral striatum activation when individuals respond to rewards or when they make reward-relevant decisions, with the general pattern of heightened activity during adolescence (Van Leijenhorst, Zanolie, Van Meel, Westenberg, Rombouts & Crone, 2009; Ernst et al., 2005; Galvan et al., 2006; May et al., 2004). Additionally, adolescents exhibit unique patterns of amygdala activation when emotional facial expressions are viewed (Baird et al., 1999; Thomas et al., 2001). With respect to more purely cognitive functions, adolescents as compared to adults show inefficient patterns of neural activation during the performance of cognitive control tasks. Interpretation of this literature is complicated by disagreements regarding how more versus less focal patterns of BOLD activation should be interpreted (see Luna et al., 2010 for a comprehensive review).

However, based on this work, “dual system” models have been invoked to explain adolescent behavior (Casey et al., 2008; Fareri et al., 2008). These models advocate that there is poor regulatory control exerted by cortical structures, particularly the PFC, due to immaturity while at the same time there is unrestrained activity within limbic and striatal circuits. While it is compelling to attribute adolescents’ relative behavioral immaturities to deficiencies in the structural integrity of PFC circuits, it could be that neurochemical systems are also changing in a manner that impacts not only motivational circuits (see Spear, 2003 for review), but also cognitive skills (Goldman-Rakic, 1987a). Neurochemical changes would necessarily interact with structural ones in determining the course of behavioral development and may help to explain inconsistencies such as the data of Berns and colleagues (2009). In considering which neurochemical systems would best map onto the behavioral patterns observed in adolescence, we and others have suggested that the dopamine system is of interest given its known role in modulation of reward responding, reinforcement learning, and high-level cognition (Frank & Hutchinson, in press; Schultz et al., 2008; Spear, 2003; Wahlstrom et al., 2010). What, then,

do we know about the status of the DA system during childhood and adolescence? Much of the available knowledge is derived from rodent and non-human primate studies. Although one could question the merits of generalizing findings from the animal literature to humans, these findings represent virtually all that is known in a direct sense about the development of this system.

## Overview of the Dopamine System

### Synthesis Pathway and Inactivation

As a member of the catecholamine class of neurotransmitters, DA is defined by the presence of a catechol nucleus and ethylamine side-chain (Feldman, Meyer, & Quenzer, 1997). Its synthesis begins with the non-essential amino acid tyrosine and ends with the production of epinephrine. Tyrosine is converted to L-3,4-dihydroxyphenylalanine (L-DOPA) by tyrosine hydroxylase (TH), the rate-limiting enzyme in the synthetic pathway. L-DOPA is catalyzed by L-amino acid decarboxylase (AADC) into DA, which is converted to norepinephrine by dopamine  $\beta$ -hydroxylase (DBH) and then to epinephrine by phenylethanolamine *N*-methyltransferase (PNMT). Specific cell types express only those enzymes necessary to produce the end-product neurotransmitter. Thus, DA cells can be differentiated from those that produce other catecholamines by their lack of DBH.

Within the synaptic cleft, excess DA either diffuses or is regulated by reuptake or catabolic mechanisms. Reuptake is the process by which DA is taken back into the pre-synaptic terminal via transporter receptors. Some proportion is then repackaged into presynaptic vesicles for later release. Transporter knockout mice have dramatically slow rates of synaptic clearance, indicating the importance of reuptake for DA inactivation and of the DA transporter in regulating this process (Giros et al., 1996). DA transporters are abundant in subcortical regions but virtually absent in the PFC (Sesack et al., 1998). Accordingly, the PFC relies on other mechanisms, particularly enzymatic inactivation, for DA's regulation. Monoamine oxidase (MAO) and catechol *O*-methyltransferase (COMT) are the primary enzymes responsible for this inactivation, with MAO accounting for approximately 90% of DA catabolism and COMT accounting for the other 10% (Westerink, 1985). MAO catabolizes all monoamines and is expressed in one of two forms: MAO-A or MAO-B. Differences between the two are attributed to substrate selectivity and the fact that each is coded for by a unique gene. MAO-A is primarily responsible for the inactivation of DA in rats (Butcher et al., 1990), though similar findings have not been documented in humans. Conversely, COMT catalyzes catecholamines and appears to be most important in the PFC (Gogos et al., 1998), since COMT knockout mice exhibit increases in PFC DA despite unchanged striatal levels.

### Cell Bodies and Projections

DA is synthesized in several regions of the mammalian midbrain. Several major cell groups have been identified (Bjorklund & Dunnett, 2007), which are located mainly in mesencephalic and diencephalic regions. They can be divided into ascending, descending, and local neuron systems (Fuxe et al., 1974). The mesencephalic DA cell groups are composed largely of ascending projections; two of these projection systems are of primary interest in understanding DA's modulation of behavior. These systems originate in the midbrain substantia nigra and ventral tegmental area (VTA). The first system projects primarily to dorsal and ventral striatal regions. The second system projects to limbic structures, including the olfactory tubercle, nucleus accumbens, amygdala, hippocampus, and septum. These projections have been labeled collectively as the "mesolimbic" DA pathway (Moore and Bloom, 1978; Oades & Halliday, 1987; Simon, Scatton & LeMoal, 1980). The second system also projects to the cortex. Cortical dopamine projections are relatively sparse, with the exception of those that reach prefrontal and primary motor regions. These cortical projections are collectively labeled the

“mesocortical” DA pathway. The mesocortical DA system includes the transitional entorhinal, cingulate, and orbitofrontal cortices as well as dorsolateral PFC (Goldman-Rakic, 1987; Beckstead, Domesick, & Nauta, 1979). The prefrontal DA projection system is distinguishable from the mesolimbic pathway in terms of the basal firing rate and degree of burst activity of DA neurons, the amount of DA turnover, the sensitivity to DA agonists and antagonists, the absence of impulse- and nerve-terminal autoreceptors, and the development of tolerance following chronic neuroleptic administration (Bannon & Roth, 1983; Bjorklund & Dunnett, 2007; Tam & Roth, 1997).

## Receptors

Six subtypes of DA receptors have been identified and designated D<sub>1</sub> through D<sub>5</sub> (two isoforms of the D<sub>2</sub> subtype exist). All are G protein-coupled and have been categorized as belonging to one of two classes designated as D<sub>1</sub>-like (D<sub>1</sub> and D<sub>5</sub>) or D<sub>2</sub>-like (D<sub>2</sub>, D<sub>3</sub>, and D<sub>4</sub>). Autoreceptors, which are D<sub>2</sub>-like, have been identified on the presynaptic terminals of dopaminergic cells (see below). D<sub>1</sub>-like receptors can be differentiated from D<sub>2</sub>-like receptors because of their ability to stimulate adenylyl cyclase activity and increase cyclic adenosine monophosphate (cAMP), which results in an overall excitatory effect within the cell (Kebabian & Calne, 1979). Conversely, D<sub>2</sub>-like receptor activation either inhibits or has no effect on cAMP levels. D<sub>1</sub> and D<sub>2</sub> receptors are distributed throughout the brain. Autoradiographic techniques in rats have demonstrated D<sub>1</sub> and D<sub>2</sub> binding in the forebrain, striatum, nucleus accumbens (nAcc), olfactory tubercles, and substantia nigra (SN), with D<sub>1</sub> density higher in all areas assessed (Boyson, McGonigle, & Molinoff, 1986). Similar to rodents, D<sub>1</sub> and D<sub>2</sub> receptors in primates and humans are distributed throughout subcortical and cortical regions (Hall et al., 1994; Lidow et al., 1991; Madras et al., 1988). D<sub>1</sub> density appears to be higher compared to D<sub>2</sub> density in limbic and cortical regions (Hall et al., 1994), although the densities of both receptor types are greater in striatal and limbic regions as compared to the cortex (Meador-Woodruff et al., 1991). Cortically, DA receptor density is relatively greater in anterior versus posterior regions (Lidow et al., 1991). Autoreceptors are found on the dendrites, cell bodies, and terminals of DA cells. Autoreceptor stimulation inhibits DA synthesis and release. Autoreceptors have not been found on DA cells that terminate in the PFC or amygdala (Kilts et al., 1987).

## Genes that Regulate Dopamine Activity

There are genetic contributions to individual differences in DA signaling, quantified through the measurement of genetic polymorphisms (genetic variants appearing in at least 1 percent of the population). Some of the most prominent within the literature include the COMT<sup>108/158</sup>Val-Met single nucleotide polymorphism (SNP) (Grossman, Emanuel, & Budaf, 1992), dopamine transporter (DAT1) variable number of tandem repeats (VNTR) (Vandenberg et al., 1992), the Taq1A SNP, which appears to impact the D<sub>2</sub> receptor system (Neville et al., 2004; Noble, 2003), D<sub>3</sub> receptor Ser9Gly SNP (Crocq et al., 1992), and the MAO-A tandem repeat (Zhu et al., 1992). All influence DA signaling by altering the activity or effectiveness of the individual components they govern (Heinz et al., 2000; Lachman et al., 1996), and all have been linked to DA-mediated behaviors (Malhotra et al., 2002; Bellgrove et al., 2005), as well as psychiatric disorders such as ADHD, schizophrenia, and alcoholism (Brookes et al., 2006; Shifman et al., 2002). The proportion of phenotypic variance accounted for by genetic polymorphisms is typically low, which is unsurprising given the complexity of most phenotypes of interest to behavioral researchers. It is thus important to note within this context that interactions between genes may account for more variance than would be predicted by additive effects. For example, Reuter and colleagues (2005) demonstrated that an interaction between COMT Val<sup>158</sup>Met and DRD2 Taq1A polymorphisms accounted for 13% of the variance in performance on the Stroop interference task, more than what was accounted for by either SNP alone. Similar findings have been revealed with respect to disease risk, as Zappia

and colleagues (2004) have demonstrated that two polymorphisms impacting the deposition of  $\beta$ -amyloid significantly increased risk for Alzheimer's disease beyond what would be expected by additive effects. Other gene-gene interactive effects whereby the effect of one polymorphism is mediated by a second polymorphism (i.e., the effect of the first is evident in only in the presence of a certain allele in the second) are also emerging in the cognitive literature (Gosso et al., 2008; Reuter, Schmitz, Corr, & Hennig, 2006; Roffman et al., 2008; Stelzel et al., 2009; Tan et al., 2007).

This overview is presented primarily to convey the complexities involved in considering developmental changes within any given neurochemical system; in evaluating the nature of developmental changes, one must consider potential alterations in precursors, synthesis, projections, receptor molecules, and enzymatic processes that regulate synaptic activity.

### **Cross-species Considerations**

Although DA cell groupings and projections have been largely conserved across mammalian species, rodents and primates differ with respect to certain aspects of the system. This evidence is important to consider because later sections of this review will present findings that suggest the developmental trajectory of the DA system differs across species. For example, the primate brain is characterized by extensive cortical DA projections, which are either minor or absent in the rat (Berger, Gaspar, Verney, 1991). In addition, primates unlike rodents exhibit variations in DA concentrations across regions, with higher concentrations measured in the frontal cortex compared to more posterior regions (Goldman-Rakic & Brown, 1982; Kehr, Lindqvist, & Carlsson, 1976). Finally, DA afferents in the adult primate cortex, but not the rodent cortex, are distributed in a laminar specific manner (Berger et al., 1988).

## **Dopamine System Development From Fetal Life to Adulthood**

### **Prenatal and Early Postnatal Development**

In rats, which have a total gestation of approximately 21 days, the neurogenesis of TH-containing cells in the mesencephalon begins on embryonic day (E) 12 (Altman & Bayer, 1981). These cells migrate to the striatum by E15 (Voorn et al., 1986) and reach the sub-plate of the future PFC before birth (Kalsbeek et al., 1988). Initially, DA projections from the midbrain to the striatum are undifferentiated, but nigrostriatal and mesolimbic specificity is established sometime between E15 and birth (Hu et al., 2004), as are projections to the PFC and ACC (Berger, Gaspar, & Verney, 1991). Similar to rodents, the development of DA cells and the innervation of cortical regions in primates is largely complete by birth. However, the developmental changes that occur prenatally in non-human primates are protracted as compared to rodents, presumably to meet the needs of a more complicated neurochemical infrastructure as evidenced by the more widespread innervation and laminar-specific differences in density described above (Berger et al., 1988; Berger, Gaspar, Verney, 1991). For example, the origin of DA cells begins in the first third of gestation in primates (Berger, Alvarez, & Goldman-Rakic, 1983; Levitt & Rakic, 1982).

The developmental trajectory in humans is largely similar to that of non-human primates, where DA-containing cells reach the subplate by 13 weeks gestation and are distributed throughout the cerebral wall at 24 weeks (Verney et al., 1993; Zecevic & Verney, 1995), corresponding to the end of neuronal migration. Across species, the elaboration of DA projections takes place throughout the dorsal PFC and anterior cingulate cortex during the first few days of postnatal life (Berger et al., 1985; Kalsbeek et al., 1988).

The neurogenesis of TH-containing cells precedes that of dopamine receptors. In rats, D<sub>1</sub> and D<sub>2</sub> receptors have been stained as early as days E14 and E15 in the striatum and reach the frontal cortex by E18 (Caille et al., 1995; Jung & Bennett, 1996; Schambra et al., 1994). D<sub>2</sub>

receptors are more densely concentrated in the striatum compared to D<sub>1</sub> receptors in the prenatal brain (Jung & Bennett, 1996), and the density and affinity of both appears to increase with age (Jung & Bennett, 1996; Sales et al., 1989). Similar patterns in the general timing of DA receptor ontogeny have been demonstrated in primates. D<sub>1</sub> and D<sub>2</sub> receptors reach the marginal zone by E73 and begin to show regional differences across cortical regions by E107 (Lidow, 1995). The distribution of D<sub>1</sub> receptors reaches maturity by E128 in the PFC and somatosensory cortex but the 8<sup>th</sup> postnatal month in the motor cortex. D<sub>2</sub> receptors follow a similar pattern (Lidow, 1995). Human data are inconsistent given methodological difficulties in measuring receptor binding in post-mortem samples; however, evidence indicates that D<sub>1</sub> and D<sub>2</sub> receptors are present in both cortical and subcortical regions by gestational week 20 (Meng et al., 1999; Unis et al., 1998).

As would be expected given the proliferation of DA-containing cells in the prenatal brain, the activity of DA precursors can be measured early in development. TH has been identified as early as E11 in the rat brain (Teitelman et al., 1979) and gestational week 5 in humans (Pearson, Brandeis, & Goldstein, 1980). DBH and PNMT can be identified shortly after (Teitelman et al., 1979), as can COMT (Parvez et al., 1979) and MAO (Fizman et al., 1991). Precursors (e.g. TH) are identified prior to degradation enzymes (e.g. COMT and MAO), suggesting that the developmental time course mirrors the synthetic chain described in adults (Fizman et al., 1991).

The early onset of DA signaling may play a functional role in the developing brain. For example, DA may influence neuronal migration and differentiation given its modulation of growth cone activity and axonal outgrowth (Lankford, DeMello, & Klein, 1988). DA also impacts the morphology of cortical cells, since the dendritic arbors of cortical pyramidal cells are decreased in rats with VTA lesions (Kalsbeek, Matthijssen, & Uylings, 1989). Part of DA's role in these developmental processes might relate to nerve growth given that DA cells contain neuro-trophins such as brain-derived neurotrophic factor (BDNF) and neurotrophin-3 (NT-3) (Seroogy et al., 1994; Tucker, Meyer, & Barde, 2001).

Thus, the DA system appears to be well developed at birth. DA-containing neurons have reached all areas of the cortex, both D<sub>1</sub> and D<sub>2</sub> receptors can be identified across cortical regions, and the infrastructure required for DA transmission is intact. With respect to postnatal changes within the DA system, the primary events of interest are likely to comprise shifts within this predefined architecture (i.e. changes in receptor number; increases or decreases in synaptic density) as opposed to broader changes in neuroanatomical structure or function.

## Juvenile and Adolescent Development

**Innervation density and tissue concentrations**—Unlike the prenatal period, which is characterized by cross-species similarities in DA signaling, developmental changes in DA concentrations and innervation during childhood and into adolescence are distinct in primates and rats.

In rhesus monkeys, DA concentrations are evenly distributed throughout the cortex in early childhood but gradually shift in an anterior direction until adolescence, at which point the highest concentrations of DA can be found in the PFC (Goldman-Rakic & Brown, 1982). Concentrations decrease in the orbitofrontal cortex (OFC) and DLPFC between birth and 8 months but increase steadily into adolescence (e.g. age 2-3 years in the macaque); conversely, DA concentrations peak in the parietal cortex at age 5 months and do not exhibit any change after birth in the occipital cortex (Brown & Goldman, 1977; Goldman-Rakic & Brown, 1982). Adult monkeys were not included in the above samples; however, other work indicates that DA concentrations in the PFC, caudate, putamen, and nucleus accumbens of squirrel monkeys decrease significantly between 3 and 9 years, with additional decreases between 9



and 18 years (Irwin et al., 1994). Human findings are mixed, with some data indicating age-related decreases in DA between ages 9 and 30 in the striatum and others finding no changes with development (Haycock et al., 2003; Kalaria et al., 1993).

Changes in DA tissue concentrations are related to innervation density; thus, in primates, DA innervation of the frontal cortex peaks during adolescence. Cross-sectional comparisons of developing rhesus monkeys indicate that the total number of DA varicosities in Brodmann's areas 4 (i.e. primary motor cortex), 9 (i.e. dorsomedial cortex), and 46 (i.e. principal sulcus) increases from birth to 2-3 years and decreases afterwards (Rosenberg & Lewis, 1994; 1995). The axonal length of DA cells projecting to the cortex undergoes similar developmental changes, with adolescents exhibiting longer axons compared to infants, children, and adults. Although axonal length and varicosities peak during adolescence in all three areas, important differences occur between cortical layers. Adolescent-limited peaks in varicosities and axonal length in area 9 are found only in cortical layer III, whereas in area 46, varicosities also decrease between adolescence and adulthood in layer VI. Conversely, area 4 is characterized by an adolescent peak in axonal length in layers I/II and III, while the number of varicosities peak during adolescence in all three cortical layers (Rosenberg & Lewis, 1995). In contrast to these PFC regions, axonal length and varicosities peak at age 7 months in the rostral entorhinal cortex and then decline into adulthood (Erickson et al., 1998).

While region-wide DA innervation of the PFC appears to peak during adolescence in non-human primates, the density of connections from individual axons appears to undergo a more linear development. The density of catecholaminergic appositions on individual pyramidal cells in the PFC of rhesus monkeys increases gradually until 2 years and plateaus thereafter (Lambe, Krimer, & Goldman-Rakic, 2000). Conversely, the density of inputs to interneurons does not change with increasing age and is lower at all ages compared to inputs to pyramidal neurons. Similar to the work of Rosenberg and Lewis, these changes have been documented specifically in cortical layer III, although other layers were not specifically targeted.

A majority of DA terminals in layer III form synapses with pyramidal neurons, which have been implicated in the maintenance of working memory (Goldman-Rakic, 1998) and are the primary source of afferent projections to other cortical regions (Goldman-Rakic et al., 1989; Jones, 1984). DA modulates the excitability of pyramidal cells by lowering the voltage threshold required for cell firing (Henze et al., 2000); thus, it is possible that adolescence is characterized by increased excitability of the neural substrates responsible for cognitive functioning and cortico-cortical information processing. In addition, projections to layer III may originate from a different cell grouping as compared to deep cortical layers. In rodents, projections to layer III have been shown to stem from lateral A10/A9 cell groupings (originating from the VTA and substantia nigra), whereas deep layers appear to receive projections from the medial A10 (VTA) grouping (Berger, Gaspar, & Verney, 1991). No replications of this work are available.

Consistent with DA's role in modulating excitatory cortical circuits, DA release is thought to aid cognitive performance by increasing neural signal-to-noise ratios (Chambers, Taylor, & Potenza, 2003; Sawaguchi, Matsumura, & Kubota, 1990; Thurley et al., 2008). These effects stand in contrast to that of other monoamines, such as serotonin, which is hypothesized to offset the role of DA by exerting an inhibitory effect (Chambers, Taylor, & Potenza, 2003). Thus, it is important to note that the development of serotonin tissue concentrations and cortical innervation during adolescence differ from those of DA. For example, concentrations of serotonin shift gradually toward posterior regions between birth and adolescence, which is due largely to developmental increases in the parietal and occipital cortex while levels in the premotor and prefrontal cortices remain stable with increasing age (Goldman-Rakic & Brown, 1982). In addition, serotonergic appositions on pyramidal cells and interneurons remain

unchanged between childhood and adulthood (Lambe, Krimer, & Goldman-Rakic, 2000). Thus, while DA is found in higher concentrations compared to serotonin in the cerebral cortex during adulthood (Lewis et al., 1992), adolescence might be characterized by a temporally limited increase in the ratio of DA to serotonin, which may have functional implications with respect to the excitability of the PFC, perhaps biasing behaviors in the direction of “go” relative to “stop” signals.

Unfortunately, the adolescent group used in the Rosenberg and Lewis studies contained only male rhesus monkeys, whereas the younger and older groups contained both males and females. Thus, it is possible that their findings were due to sex effects rather than age. While there were no sex differences within the other age groups studied (i.e. childhood and adulthood), adolescence may be unique due to hormonal changes.

Unlike primates, there is little evidence that the rodent cortex undergoes adolescent-limited increases in DA innervation; rather, it appears that innervation increases steadily until approximately postnatal day (P) 60, which corresponds to the early stages of adulthood in the rat (Berger et al., 1985; Kalsbeek et al., 1988). This pattern has been replicated in both the PFC and ACC. Findings regarding DA concentrations in the rodent brain are mixed. Several reports indicate slight age-dependent increases in DA tissue concentrations in the cortex (Giorgi et al., 1987; Nomura, Naitoh, & Seqawa, 1976) and striatum and midbrain (Ungethüm et al., 1996). However, others suggest that the available synaptic storage pool is increased in adolescence relative to both childhood and adulthood (Andersen, Dumont, & Teicher, 1997; Stamford, 1989). In addition, two major developmental milestones appear to be met during this time. First, the topographical distribution of DA innervation in the cortex attains an adult pattern (Berger et al., 1985). Second, neurons innervating the cortex develop with two distinct fiber structure characteristics: thick fibers that are also present in subcortical regions such as the hippocampus and thin fibers with multiple varicosities that form dense connectivity with (presumably) pyramidal cells in middle cortical layers (Kalsbeek et al., 1988). These maturations are achieved by postnatal days 30 and 35, respectively, which correspond to the early to mid stages of peri-adolescence.

Overall, the above evidence suggests that the primate cortex continues to undergo changes in DA innervation into adulthood, with rates increasing towards adolescence and decreasing shortly thereafter. This pattern is particularly evident in cortical layer III, indicating a specific pattern of development in the areas subserving intra-cortical information processing. Conversely, rat cortex is characterized by slight increases in DA innervation until adulthood, and adolescence does not appear to be unique with respect to these changes. The reasons for these cross-species differences are unknown. However, there are clear differences in adult level prefrontal structure and function between primates and rodents, with the former exhibiting both increased cerebral differentiation and more complex cognitive functioning (Preuss & Kaas, 1999; Uylings, Groenewegen, & Kolb, 2003). These differences, while not directly related to DA innervation during adolescence, may contribute to the species differences mentioned above.

**Receptor Density**—Unlike the prenatal period, where DA cell proliferation precedes the origin of receptors, D<sub>1</sub> and D<sub>2</sub> densities appear to peak in the cortex prior to adolescent-limited increases in cortical innervation and DA axonal growth. Both D<sub>1</sub> and D<sub>2</sub> receptors peak between 2 and 4 postnatal months in the developing rhesus monkey, with concurrent development occurring across prefrontal, motor, somatosensory, and visual cortices (Lidow, Goldman-Rakic, & Rakic, 1991; Lidow & Rakic, 1992). After peaking, receptor densities decrease slowly until early adulthood (e.g. 60 months), and at all time points D<sub>1</sub> receptors can be found at higher densities compared to D<sub>2</sub> receptors. Few studies have systematically assessed D<sub>1</sub> and D<sub>2</sub> receptor changes in humans, but similar to work in non-human primates,

they indicate decreases in D<sub>1</sub> and D<sub>2</sub> binding during adolescence. Seeman et al. (1987) demonstrated that D<sub>1</sub> and D<sub>2</sub> binding increases to age 4 and decreases throughout late childhood and adolescence. Sex differences were assessed, but none were found. Because the study relied on postmortem specimens, there were relatively few subjects between the ages of 5 and 20 years. As a result, conclusions regarding developmental changes in receptor density during adolescence were based on very few cases and should be considered tentative. Others have demonstrated decreases in D<sub>1</sub> density between infancy and adulthood in the caudate and putamen but also based on very few subjects (Montague et al., 1999).

Although changes in D<sub>1</sub> and D<sub>2</sub> densities occur simultaneously across cortical areas in non-human primates, the motor and visual cortices appear to be unique in that they display differential receptor changes across cortical layers that result in the reorganization of the laminar distributions of D<sub>1</sub> and D<sub>2</sub> densities (Lidow & Rakic, 1992). In the motor cortex, D<sub>1</sub> receptors are concentrated in layer V at birth but increase in layer III relative to other regions so that by 2-4 months of age, they are most densely concentrated there. Layers I, II, and III undergo the smallest rate of loss between 8 months and adulthood and as a result, they are the most densely concentrated areas in adults. D<sub>2</sub> receptors are most dense in layers I and II at birth, but within the first month of life, they increase significantly in layer V so that this area contains the highest density of receptors from that point forward. The laminar distribution in the visual cortex undergoes less substantial change compared to the motor cortex, with D<sub>2</sub> receptors shifting from layers II and III to V in the first month of life. In all regions, receptor affinity does not change after birth (Lidow, Goldman-Rakic, and Rakic, 1991), suggesting that the kinetic properties of DA receptors mature prenatally.

Whereas primate studies are relatively few, there is a large body of evidence assessing similar changes in the rodent. Although some variation across studies exists due to the fact that receptor concentrations are measured at different postnatal ages, most data indicate that in the striatum, D<sub>1</sub> receptors increase in number between birth and PD 35 to 40 and decrease thereafter (Andersen et al., 1997; Tarazi, Tomasini, & Baldessarini, 1999; Teicher, Andersen, & Hostetter Jr., 1995). This finding is most reliably demonstrated in the caudate and putamen (Gelbard et al., 1989; Giorgi et al., 1987; Tarazi, Tomasini, & Baldessarini, 1999; Teicher, Andersen, & Hostetter Jr., 1995), although other studies suggest that D<sub>1</sub> binding in the dorsal striatum increases to adult levels in periadolescence and then stabilizes, as opposed to peaking on approximately PD 35 (Leslie et al., 1991; Rao et al., 1991). These seemingly incoherent findings are likely due to differences in the age groups measured, as Rao et al. (1991) did not measure D<sub>1</sub> density between PD 30 and 60 and Leslie et al. (1991) did not measure past PD 42. The developmental trajectory of D<sub>1</sub> receptors in the nucleus accumbens is less clear. Some groups have found steady increases in D<sub>1</sub> density into adolescence (Leslie et al., 1991), others have reported a periadolescent-limited peak in density (Tarazi, Tomasini, & Baldessarini, 1999), and yet others suggest a periadolescent-limited peak followed by decline and then another peak at PD 80 (Andersen et al., 1997; Teicher, Andersen, & Hostetter Jr., 1995). The reasons for these cross-study differences are unclear but may be related to different binding agents, as well as differences in the age groups measured (similar to those mentioned above). For example, only Andersen et al. (1997) and Teicher, Andersen, & Hostetter Jr. (1995) measured D<sub>1</sub> density until day PD 80, whereas Leslie et al. (1991) measured until PD 42.

Interestingly, changes in D<sub>1</sub> receptor density in the frontal cortex of rats do not demonstrate the periadolescent overproduction exhibited in subcortical structures. For example, some have found that similar to primates, D<sub>1</sub> density peaks prior to periadolescence (e.g. PD 21) and decreases afterwards (Leslie, 1991). However, others have demonstrated that density increases steadily between birth, adolescence, and adulthood (Murrin & Zeng, 1990; Tarazi & Baldessarini, 2000), a pattern similar to the development of cortical innervation in the rodent. This overall developmental trajectory, characterized by subcortical peaks in D<sub>1</sub> density

occurring after cortical peaks, runs contrary to expectations given that the cortex is thought to be the last brain region to develop. This finding suggests that the cortical and subcortical circuits may develop independently. None of the above studies have demonstrated that D<sub>1</sub> densities peak in the cortex prior to the striatum.

D<sub>2</sub> receptor density follows a similar developmental pattern compared to its D<sub>1</sub> counterpart. Specifically, it appears to peak in the caudate and putamen between PD 28 and PD 40 depending on the study (Gelbard et al., 1989; Tarazi, Tomasini, & Baldessarini, 1998b; Teicher, Andersen, & Hostetter Jr., 1995), though others have demonstrated non-significant increases during that time compared to adulthood (Rao, Molinoff, & Joyce, 1991). In addition, D<sub>2</sub> density increases in the medial PFC until adulthood and does not exhibit a periadolescent-limited peak (Tarazi, Tomasini, & Baldessarini, 1998b). Contrary to observations in the striatum and cortex, similarities between D<sub>1</sub> and D<sub>2</sub> development have not been found in the nucleus accumbens (nAcc), where D<sub>2</sub> receptors do not exhibit density increases in later adulthood as do D<sub>1</sub> receptors (Andersen et al., 1997; Teicher, Andersen, & Hostetter Jr., 1995). In all studies directly comparing receptor subclasses, D<sub>1</sub> binding has been found to be higher compared to D<sub>2</sub> binding in the striatum and nAcc at all age groups studied (Andersen et al., 1997; Gelbard et al., 1989), and age interactions have been demonstrated whereby D<sub>1</sub> density increased faster and dropped off more rapidly around periadolescence (Gelbard et al., 1989). Interestingly, a pronounced medial-to-lateral gradient in D<sub>2</sub> density has been established in the striatum, with a specific increase in magnitude occurring during PD 40 (Teicher, Andersen, & Hostetter Jr., 1995). Similar findings have not been demonstrated within the D<sub>1</sub> family.

Sex differences in D<sub>1</sub> and D<sub>2</sub> receptor development are evident. Andersen et al. (1997) compared the changes in D<sub>1</sub> and D<sub>2</sub> receptor densities in the striatum and nAcc of rats aged between PD 25 and 100. They demonstrated that only males exhibited periadolescent-limited increases of D<sub>1</sub> and D<sub>2</sub> receptors in the striatum, although males and females exhibited similar densities in adulthood. A similar pattern was exhibited by D<sub>2</sub> receptors in the nAcc. Conversely, both males and females exhibited overproduction of D<sub>1</sub> receptors in the nAcc during periadolescence, although males also exhibited a peak at PD 100 and had higher D<sub>1</sub> density at all ages studied. This pattern is consistent with human MRI evidence that the striatum shrinks during adolescence in males but stays relatively the same size in females (Giedd et al., 1996). Interestingly, sex differences in receptor overproduction do not appear to be related to adolescent surges in gonadal hormones, as neither castration nor ovariectomy changed the developmental trajectory of D<sub>1</sub> or D<sub>2</sub> receptor density (Andersen et al., 2002). Other studies have indicated that the development of DA cells and receptors is similar in males and females (Lieb et al., 1996); however, rates of change during adolescence were not directly compared.

D<sub>3</sub> and D<sub>4</sub> receptors have been less extensively studied. Available evidence suggests that D<sub>4</sub> and D<sub>2</sub> receptors follow similar developmental trajectories. D<sub>4</sub> density within the striatum increases until periadolescence and decreases thereafter; conversely, the forebrain undergoes steady increases in density until periadolescence that then plateau into adulthood (Tarazi, Tomasini, & Baldessarini, 1998b). Unfortunately, studies of D<sub>3</sub> development bypass periadolescence and exclude measurements between PD 21 and PD 90 (Demotes-Mainard et al., 1996; Stanwood et al., 1997). Despite these limitations, evidence indicates D<sub>3</sub> density in the striatum and nAcc increases between birth and PD 21 and increases again to PD 90. Whether a brief peak occurs during periadolescence is unknown.

**Transporter Density**—Findings regarding postnatal changes in DA transporters have been inconsistent. This is especially true in the SN and VTA, where different lines of evidence have indicated no developmental changes (Moll et al., 2000), steady increases between birth and adulthood (Galineau et al., 2004), and peaks in transporter density at PD 21 (Coulter, Happe, & Murrin, 1996). The reasons for these discrepancies are unclear, although different binding

agents were used in each study and the age groups studied are not totally comparable. For example, neither the Moll et al. (2000) nor the Galineau et al. (2004) studies examined transporter levels at age PD 21, which is the age at which density peaked according to the evidence of Coulter, Happe, and Murrin (1996). Findings in subcortical regions are more coherent, with converging lines of evidence indicating that transporter density increases into periadolescence and plateaus thereafter in the striatum, nAcc, thalamus, and bed nucleus of the stria terminalis (Coulter, Happe, & Murrin, 1996; Galineau et al., 2004; Tarazi, Tomasini, & Baldessarini, 1998a; but see Moll et al., 2000 for contrary findings). Studies of developmental changes in the rat cortex are sparse, likely due to the relative dearth of DA transporters in this area. Coulter, Happe, and Murrin (1996) found that frontal transporter density increased rapidly between PD 21 and adulthood, while Moll et al. (2000) were unable to yield interpretable results.

**Summary**—D<sub>1</sub> and D<sub>2</sub> receptors in nonhuman primate and human cortex appear to peak in early childhood and decline thereafter, reaching relatively stable levels by early adulthood. D<sub>1</sub> receptors are more densely concentrated at every developmental stage, and similar findings have been found in subcortical regions. Contrary to the results in primates, DA receptor densities in the rodent cortex appear to increase during adolescence, and D<sub>1</sub> and D<sub>2</sub> densities in the striatum peak during peri-adolescence.

**Precursors and Metabolites**—Evidence regarding changes in DA precursors and metabolites is inconsistent and may provide limited information about the development of DA signaling. This is especially true because metabolite levels are confounded by DA usage, which decreases with age (Leslie et al., 1991; Teicher et al., 1993), and regional innervation patterns (McGeer et al., 1967), which vary widely across developmental periods and species (Kalsbeek et al., 1988; Rosenberg & Lewis, 1994). For example, TH activity has been shown to decrease with age in human PFC (Weickert et al., 2006) and striatum (McGeer & McGeer, 1976), though others have shown brief increases in the striatum in early childhood (Haycock et al., 2003). Findings in other species have indicated the opposite to be true, as TH increases during childhood and adolescence in rat and rabbit midbrain, striatum, and cortex (McGeer et al., 1967; Ungethüm et al., 1996). Similar inconsistencies have been found with respect to AADC, where data demonstrate both decreases in the striatum after the age of 2 in humans (Haycock et al., 2003) and increases throughout the brain in pigs (Brust et al., 2004). HVA levels have been shown to decrease in the rat striatum (Ungethüm et al., 1996) and human cerebrospinal fluid (Seifert Jr., Foxx, & Butler, 1980), though others have indicated that it does not change with age in the human striatum (Haycock et al., 2003). COMT, on the other hand, is unique in that its activity has been shown to increase in both human PFC (Tunbridge et al., 2007) and porcine striatum (Brust et al., 2004) during adolescence. Overall, the factors underlying the discrepancies in these data are unknown but may relate to differences in the species and brain areas studied, as well as the underlying changes in regional DA innervation and usage, neither of which are typically measured simultaneously with presynaptic metabolites. At the very least, these data indicate that DA metabolites may vary with age.

## Integration: Evidence for Increased Dopamine Availability in Adolescence Compared to Adulthood

Despite some conflicting results, the available evidence, summarized in Table 1, suggests that both primates and rodents exhibit increases in DA signaling during adolescence, though differences exist with respect to the regions and aspects of the DA system affected. Rodents do not exhibit peaks in cortical DA innervation during adolescence, as evidence suggests that innervation undergoes monotonic increases lasting until early adulthood (Kalsbeek et al., 1988). However, cortical DA availability, and synaptic DA availability in the striatum, peak

during adolescence (Andersen, Dumont, & Teicher, 1997; Stamford, 1989). Similar to innervation patterns, receptor densities in the cortex increase throughout adolescence (Tarazi & Baldessarini, 2000). D<sub>1</sub> and D<sub>2</sub> densities appear to peak in subcortical structures such as the striatum and nAcc, as do the densities of D<sub>2</sub>-like D<sub>3</sub> and D<sub>4</sub> receptors (Andersen et al., 2002; Tarazi, Tomasini, & Baldessarini, 1998b). Thus, it appears that in rodents, adolescence is characterized by a subcortical increase in DA compared to childhood and adulthood. Conversely, DA availability in the cortex is lower compared to adulthood (but see Andersen, Dumont, & Teicher, 1997).

In primates, cortical and subcortical tissue concentrations of DA are increased during adolescence compared to childhood and adulthood (Goldman-Rakic & Brown, 1982; Irwin et al., 1994). In addition, DA innervation of the frontal cortex peaks during adolescence relative to childhood and adulthood, specifically in cortical layer III, which contains pyramidal cells responsible for cortico-cortical information processing (Rosenberg & Lewis, 1995). D<sub>1</sub> and D<sub>2</sub> receptor densities appear to be heightened during adolescence compared to adulthood in both cortical and subcortical regions, though peaks in receptor density occur in childhood (Lidow & Rakic, 1992; Seeman et al., 1987). Thus, cortical and subcortical regions undergo specific increases in DA concentrations and innervation during adolescence, with receptor levels decreasing from peaks achieved during childhood.

These patterns indicate that the DA system is patterning itself in a manner during adolescence that could support functional changes in behaviors that are DA-modulated. For example, the receptor evidence supports the notion that limbic and striatal DA systems may be in a state of overdrive during adolescence. Moreover, there is evidence in rodents of a cortical/subcortical distinction. This distinction is not as apparent in primates, but it could be the case that “overdose” thresholds differ between cortical and subcortical regions in primates based on autoreceptor and dopamine transporter characteristics. Accordingly, adolescent-limited increases in dopamine availability could account for the accelerated drive toward potential rewards that is a hallmark of adolescent behavior.

Of course, virtually all of this work relies on animal models. There are limited strategies for assessing DA system development in healthy human developmental samples. Postmortem samples have been the primary means through which changes in DA system characteristics have been inferred. Other potential strategies include the use of pharmacological probes to provoke reactions in targeted chemical systems and/or positron emission tomography (PET) to measure brain activation changes in the presence of radioactive ligands. Neither of these methods is easy to justify for non-clinical samples. Studies of clinical samples, such as children with attention-deficit-hyperactivity-disorder (ADHD) are potentially informative because of the use of psychostimulants to treat the disorder, however the assessment of psychostimulant effects in ADHD is complicated by the unknown influences of the disorder itself on observed patterns. Other conditions of interest, such as PKU, have been used as models to illustrate developmental aspects of PFC DA availability (Diamond, 1996; Luciana, Whitney, & Hanson, 2004), but this work is indirect.

Despite these logistical obstacles, several groups have used fMRI techniques that allow for assessment of changes in neural systems innervated by DA (which may reflect changes in DA signaling in these areas). For example, despite activating a similar network of neural structures, adolescents generate larger BOLD activation in the nAcc compared to adults and young children when presented with rewarding stimuli (Ernst et al., 2005; Galvan et al., 2006; May et al., 2004). In addition, either smaller (Esher et al., 2007) or less focally distributed (Galvan et al., 2006) activations in response to reward presentation have been demonstrated in the orbitofrontal cortex (OFC) and anterior cingulate cortex (ACC), indicating that increased subcortical activity might be accompanied by less-developed prefrontal control. As mentioned

previously, these data have led to theories of adolescent risk-taking whereby active ventral striatal responsivity in the context of poor prefrontal regulation accounts for an increased propensity to engage in risky activities (Casey, Getz, & Galvan, 2008). On the one hand, more focal activation in brain regions in response to task demands have been interpreted to reflect more efficient neural processing (see Mattay et al., 2003), whereas other data suggest that activation increases as a function of learning and practice (Karni et al., 1998). Furthermore, cross-sectional and longitudinal studies indicate that development is accompanied by more focal activation in brain areas necessary for task completion, further complicating interpretation (for review, see Casey et al., 2008).

While it is possible to speculate about how these fMRI findings might be related to neurotransmission, inferences regarding neurochemical development are impossible to make with any certainty. Increases in neurochemical availability could result in BOLD signal increases or decreases depending on the task employed, the region of interest, and the nature of the chemical system. Thus, the BOLD signal in and of itself is not necessarily informative regarding directional changes in neurochemical activity.

In the absence of more direct assessment strategies, we have been intrigued by the notion of using molecular genetic probes to index the underlying status of the DA system to study neurochemical development. We have hypothesized, based on the animal data, that adolescents have high functional levels of DA activity in the cortex (Wahlstrom et al., 2007; Wahlstrom et al., 2010). If true, then there might be age-related variation in the relationship between genes and cognition, when genes that code for specific aspects of DA transmission are examined. The COMT SNP is of interest, because it is thought to have some specificity to PFC function and acts on the DA system in the synapse, impacting how much released DA reaches postsynaptic receptors. If DA release changes in magnitude throughout development and if COMT activity does not change to parametrically match how much DA is released, then the optimal COMT genotype for the enhancement of prefrontally-guided cognition might change in concert with underlying changes in DA availability. This model will be expanded upon in the next section.

## COMT: Functions and Associations with Human Cognition

As described above, the COMT enzyme, along with monoamine oxidase (MAO), represents the first step in the catabolism of excess synaptic DA with some specificity to the PFC due to a lack of DA transporter activity in this region (Giros et al., 1996; Gogos et al., 1998; Huotari et al., 2002; Lewis et al., 2001; Napolitano, Cesura, & Da Prada, 1995; Sesack et al., 1998; Weinshilboum et al., 1999). The gene that codes the COMT enzyme resides on the q11 region of Chromosome 22 (Grossman et al., 1992), and the val<sup>158/108</sup>met polymorphism results from a single nucleotide polymorphism (SNP) on exon 4 caused by a G-to-A base-pair mutation (Lachmann et al., 1996). This mutation results in the substitution of valine (Val) for methianine (Met) on codon 158 of the membrane-bound enzyme and codon 108 of the soluble enzyme (Lundström et al., 1991). Thus, there are two alleles, Met and Val. Functionally, the Met allele confers four times lower enzymatic activity compared to Val, which results in increased levels of synaptic DA (Lachmann et al., 1996). In humans, individuals that are homozygous Met exhibit higher levels of PFC DA compared to individuals homozygous for Val. Heterozygotes may exhibit intermediate levels of DA, as the alleles are codominant (Lotta et al., 1995; Weinshilboum et al., 1977).

### COMT and Cognitive Functioning: Adult Studies

Due to DA's role in regulating prefrontal cognition, the COMT polymorphism was identified early as a possible contributor to individual variation in functions dependent on this region. A wealth of evidence suggests that COMT is associated with performance on working memory

(Goldberg et al., 2003) and executive set-shifting tasks such as the Wisconsin Card Sort (Malhotra et al., 2002), with the Met allele typically conferring relatively better performance. This effect has been demonstrated and replicated in healthy volunteers and schizophrenic patients (Bilder et al., 2002; Egan et al., 2001; Mattay et al., 2003; though for a recent meta-analysis that suggests the effect of COMT holds in healthy control samples only, see Barnett, Jones, Robbins, & Muller, 2007). Egan and colleagues (2001) have shown that COMT is related to changes in prefrontal physiology during the completion of cognitive tasks, as they demonstrated that Met/Met volunteers performed better on the Wisconsin Card Sort compared to Val/Met and Val/Val individuals while simultaneously generating smaller BOLD responses in the DLPFC, indicating more efficient neural processing. Better attentional function, as measured by the digit span forward task, has also been demonstrated in a sample of elderly Asian males who were free of cognitive impairment (Liu et al., 2008). Overall, it should be noted that the effect of COMT on cognitive variables is small, similar to other polymorphisms; for example, the effect size of COMT on Wisconsin Card Sort performance as estimated by Barnett and colleagues (2007) in healthy adult controls is Cohen's  $d = 0.22$ . Moreover, sample sizes are often small in individual studies, which may account for inconsistencies in the literature, particularly developmental studies.

### COMT and Cognition: Developmental Findings

The literature linking COMT to cognitive functioning in children and adolescents is relatively sparse; moreover, available findings are inconsistent. In a sample of 8 to 14 year-old children (mean age = 9), Diamond et al. (2004) demonstrated that the Met allele was associated with a test of inhibition previously shown to depend on prefrontal DA. Similarly, Barnett and colleagues (2007) found that being homozygous Met predicted optimal performance on measures of verbal IQ, working memory, and inhibition in a sample of 8 and 10 year-old children; however, COMT appeared to predict cognitive performance only in males, and heterozygotes outperformed both homozygote groups on a measure of sustained attention. COMT was not related to intelligence in a sample of healthy children in China (Zhang et al., 2007). Studies of children with ADHD have suggested that COMT is unrelated to performance on tests of executive functions such as working memory, set-shifting, planning, and response inhibition (Mills et al., 2004; Taerk et al., 2004), whereas others have found that being homozygous for the Met allele impairs sustained attention performance (Bellgrove et al., 2005).

The role of COMT in prefrontal cognition has been of considerable interest to those groups studying 22q11.2 deletion syndrome, which is caused by a hemizygous deletion in a small region of chromosome 22 that contains the COMT gene and results in debilitating cognitive impairments and a significant increase in the risk for schizophrenia (Bassett & Chow, 1999; Scambler, 2000). Individuals with 22q11.2 deletion are either hemizygous Met or hemizygous Val, and as a result, the effects of the Val<sup>158</sup>/Met polymorphism on behavior are presumably larger than in healthy controls. The Met allele has been linked to higher overall IQ and improved sustained attention in children with 22q11.2 (Shahi et al., 2006), though other work using samples of both children and young adults found no association between COMT and measures of IQ, verbal fluency, or response inhibition (Glaser et al., 2006). The reasons for the null findings in the latter study may have to do with the broad range of ages assessed (6 – 37 years) and the fact that the relationship between COMT and cognition changes with age in 22q11.2. Specifically, Golthelf et al. (2005) demonstrated that the Met allele predicts higher IQ in young adolescents (mean age 13 years) with 22q11.2; however, their cognitive abilities decline at a more rapid rate so that by 18 years, Val individuals exhibit higher IQs.

Noticeably missing from the developmental literature are studies assessing the link between COMT genotype and prefrontal cognition in adolescence, especially given the importance of



this time as a period of normative developmental transitions and increased risk for psychopathology. We have associated the COMT genotype with working memory and attention in a sample of adolescents and young adults (Wahlstrom et al., 2007). We administered a battery of working memory, attention, motor coordination, and motor speed measures to children and adolescence ages 9-17 ( $n = 70$ ) genotyped for COMT. Composites were created to represent the cognitive domains of working memory/attention and motor coordination/speed. These composites were created to maximize both theoretical coherence (e.g., similarity of function) and psychometric strength (e.g., high internal consistency) and to increase statistical power in reporting associations given the small sample size. We reported that the COMT Val<sup>158</sup>/Met genotype was significantly related to measures of working memory/attention, and motor coordination, all of which depend on prefrontal DA (Glickstein, DeSteno, Hof, & Schmauss, 2005; Luciana & Collins, 1997; Yang et al., 2003). Conversely, genotype was not related to a measure of motor speed, indicating specificity in the modulation of frontal behaviors, which is consistent with other studies of children (Diamond et al., 2004). The critical finding in our study concerned the optimal genotype for cognitive performance in this adolescent sample. Instead of finding Met-Met superiority of cognitive function as would have been predicted from the adult literature, we found that Val-Met heterozygotes exhibited superior performance on all cognitive constructs compared to both homozygote groups (see Figure 1: Wahlstrom et al., 2007). Specifically, heterozygotes outperformed the Met-Met group (but not the Val-Val group) on the working memory/attention composite, and heterozygotes outperformed both homozygote groups on the motor composite. At first glance, this pattern might be seen as consistent with the overall degree of inconsistency of findings relating the COMT genotype to cognition in developmental samples. However, another proposal is that these data are reflective of underlying transitions in DA availability that occur during adolescence as compared to other points in the lifespan. The rationale for this proposal is as follows.

First, in the adult literature, the optimal COMT genotype for cognitive performance varies based on state characteristics. Mattay and colleagues (2003) recruited a sample that varied in its representation of the COMT genotype. All participants completed a short battery of executive function measures (the *N*-back task and the Wisconsin Card Sort Task) under baseline conditions. All participants then completed two fMRI scans, during which the *N*-back task was administered. The fMRI scans were distinct in that one scan took place under placebo conditions and the other took place after participants ingested a small dose of *d*-amphetamine, which non-specifically increases DA activity. Under placebo conditions, Met/Met individuals exhibited better working memory performance compared to Val/Val individuals, and this improved performance was associated with smaller BOLD activations in the DLPFC, which the authors interpreted as an indication of greater PFC efficiency. However, after the administration of *d*-amphetamine, which presumably raised dopamine levels in the PFC, the Val/Val group improved their working memory performance in the context of decreased BOLD activations, indicating that less neural processing was necessary to obtain improved behavioral performance. The opposite pattern was evident in the Met/Met group, which exhibited less efficient neural processing and poorer behavioral performance in the *d*-amphetamine, relative to the placebo, condition. Thus, under conditions of increased DA release, the Met-Met group performed worse, while the Val-Val group performed better. The results were interpreted by the study's authors in accord with the inverted-U model that represents the DA-cognition association. That is, there is an optimal level of dopamine availability in relation to cognitive function (Williams & Goldman-Rakic, 1995). It was suggested that Met/Met individuals rested near the apex of this curve at baseline (as evidenced by better performance and more efficient neural processing under placebo conditions) and that Val/Val individuals rested near the lower left slope of the curve due to decreased DA levels. After increasing DA levels with *d*-amphetamine, Met/Met individuals' neural processing suffered due to excess levels of DA, which pushed them to the right-most downslope of the inverted-U, while the Val/Val group

showed improvements as they shifted from the lower left portion of the curve to a point close to its apex. Recently, animal models have been developed to illustrate similar relationships. Transgenic mice who over-express the Val polymorphism exhibit disrupted attentional set-shifting and show impairments in multiple forms of memory, including both working and recognition memory, as compared to mice with a null COMT mutation (Papaleo et al., 2008). Amphetamine administration improved the experimental animals' memory deficits but impaired memory in control animals.

Based on these findings, one interpretation is that the relationship between COMT alleles and cognitive functioning changes in a similar manner during adolescence due to the underlying development of the DA system (Wahlstrom et al, 2007). Specifically, in accord with the evidence that DA levels increase in the PFC during adolescence, it may be that adolescents who have the Met/Met genotype may experience excessive levels of DA in prefrontal circuits, and hence, poorer cognitive functioning as compared to Val/Met and even Val/Val individuals.

Figure 2 illustrates this hypothesis. In adulthood, characterized by mature levels of DA, Met/Met individuals rest near the apex of the inverted-U shaped function that describes the relationship between DA and cognition. Conversely, Val/Met and Val/Val individuals, who exhibit lower levels of synaptic DA due to more efficient COMT enzymes, would rest toward the left-most downslope of the curve, which is characterized by less optimal cognitive functioning. In adolescence, when increased levels of DA are experienced due to developmental changes, the relationship between COMT allele and cognitive functioning may presumably shift as seen in adults administered *d*-amphetamine. Specifically, increases in DA may interact with the Met/Met genotype to cause excess levels of DA in the PFC, shifting these individuals to the upper right portion of the inverted U that is characterized by inefficient cognitive functioning. Val/Met and Val/Val individuals would benefit from these increases in DA transmission, which would shift them closer to the apex of the curve and result in improvements in cognitive functioning relative to Met/Met volunteers.

In understanding this model, several points are worthy of emphasis. First, developmentally, it is expected that regardless of genotype, individuals will improve their levels of performance on cognitive tasks between adolescence and adulthood, as many factors in addition to the COMT genotype contribute to the development of executive function skills (e.g., structural brain development, increasing levels of experience, other components of the DA system, etc.) (Nagy, Westerberg, Klingberg, 2004). Second, it is assumed that all individuals, regardless of genotype, experience increases in functional DA availability during adolescence. And finally, although there may be changes in COMT enzyme activity throughout development (Turnbridge et al., 2007), these changes are stable within age groups. What is expected on the basis of this model is that there will be age by genotype interactions related to the differential impacts of individual COMT genotypes on cognition at each point in development and related to how the different genotypes respond in the course of development. That is, Met/Met individuals may make larger developmental gains between adolescence and adulthood, because they are moving from an inefficient area of the curve (due to excess DA) to the apex. Conversely, the developmental gains made by Val/Met and Val/Val individuals may be smaller, because as they move into adulthood, which is characterized by lower DA levels compared to adolescence, they shift down to the lower downslope of the curve as defined by their COMT genotype.

Ideally, longitudinal designs would be implemented to test this model, and our laboratory is in the midst of data collection toward this end. This framework is consistent with a growing body of literature that indicates the relationship between COMT and cognition changes as a function of development. As mentioned above, Golthelf and colleagues (2005) demonstrated in a longitudinal design that individuals with 22q11.2 deletion syndrome hemizygous for Met exhibited more rapid cognitive decline compared to those hemizygous for Val. Given the

widespread impairments associated with 22q11.2 deletion, it is difficult to interpret these findings within the context of normative development, but their results do indicate that age modulates the effect of COMT genotype on cognition during the adolescent period.

Similar models have been advocated to explain age-related decrements in behaviors that are DA-modulated (Backman et al., 2006). In the context of healthy aging, several cognitive processes are known to decline, notably processing speed, working and episodic memory, and fluid reasoning skills (Verhaeghen & Salthouse, 1997). Similarly, there is evidence for age-related loss in functional dopamine availability, primarily observed in nigrostriatal regions due to the relative ease through which biomarkers of DA availability can be measured in the striatum during Positron Emission Tomography (PET) scans and SPECT imaging (Bäckman et al., 2006; Kaasinen et al., 2000; Suhara et al., 1991; Volkow et al., 1994; 2000; 2004). These studies, along with findings from human autopsy studies, indicate age-related declines in D<sub>1</sub> and D<sub>2</sub> receptor numbers, the DA transporter, and DA cells in the midbrain and striatum (Volkow et al., 1998). Consistent with our model, it has been suggested that these normative mechanisms may have greater functional consequences for individuals with lower levels of baseline DA availability (Backman et al., 2006), and indeed, individuals who carry the Val-Val COMT configuration appear to be more susceptible to these age-related changes. In a sample of adults ages 20-31 and 60-70 years, Nagal et al. (2008) demonstrated a COMT genotype by age interaction on Wisconsin Card Sorting Test (WCST) performance, whereby the functional consequence of the Val allele was increased in the oldest group. This finding would seem to reflect the interaction of the Val allele with age-related decreases in dopamine, which would theoretically shift Val homozygotes further down the inverted-U relating DA transmission and cognition. In contrast, Met-Met individuals, who have presumably higher levels of functionally available dopamine at the start, may be more invulnerable to age-related declines in the system. Accordingly, this model predicts individual differences in neuroplasticity on the basis of genotype, and indeed, we observe that adolescents who carry the Met-Met genotype have a more protracted rate of working memory development than individuals with the Val-Met and Val-Val genotypes (Wahlstrom et al., in press). Given that the aging work is the subject of another paper within this issue, we have not covered it in detail here but refer interested readers to the paper by Backman and colleagues.

Recently, some supportive evidence has emerged indicating that sex also modulates the effect of COMT on cognition, as elderly females appear to benefit from the Val allele when performing the Wisconsin Card Sort, whereas aging males show benefit from possessing the Met allele (Diamond, 2007). COMT activity is lower in females compared to males (Boudikova, Szumlanski, Maidak, & Weinshilboum, 1990); thus, females presumably exhibit higher DA levels in the PFC and would show no benefit from the Met allele, which pushes them into the excess range of the inverted-U and impairs their performance. Males, on the other hand, exhibit lower levels of baseline DA and do show improvement. How these sex effects interact with normative development is unclear, but these findings raise interesting questions about how hormonal state and perhaps pubertal development might impact some of these observed associations.

The evidence presented above suggests that the influence of COMT on cognition may depend on state factors such as age and individual differences, such as sex, both of which are known to impact the DA system that mediates the relationship between COMT and cognitive processes. Given that small samples and effect sizes characterize most COMT studies, results regarding its impact on cognitive performance in adults have been inconsistently replicated, and even meta-analyses have yielded inconsistent findings (Barnett et al., 2007; Barnett, Scoriels, & Munafò, 2008). Thus, the results discussed here are admittedly preliminary, and several replications focused on specific age groups (e.g., adolescents, elderly individuals, etc.) will be necessary. It is important to note within this context that the findings of Mattay and

colleagues (2003), which stimulated the conceptualization of this framework, have not been replicated.

Finally, research utilizing the COMT Val<sup>158</sup>/Met polymorphism is complicated by the fact that the precise mechanisms by which it impacts DA signaling are debated. Although it is typically assumed that COMT's relationship to prefrontal cognition is accounted for by generalized changes in DA in that region, others have proposed that Val and Met alleles differentially affect tonic vs. phasic aspects of striatal DA transmission. Specifically, the Met allele may be related to higher tonic levels of subcortical DA, and, by association, decreased subcortical phasic amplitude in response to neuronal firing and increased cortical D<sub>1</sub> transmission (Bilder et al., 2004). Functionally, these changes may result in stability of neural networks underlying cognition, which comes at the cost of decreased flexibility and would thus affect various phenotypes differentially. Interestingly, fMRI evidence supports these claims, with Met individuals exhibiting a heightened transient activation during the maintenance phase of a working memory task and Val individuals a heightened sustained response throughout the maintenance period (DeFrias et al., 2009). With respect to the data presented in this review, the tonic/phasic theory would suggest that Met is related to improved working memory (due to improved cognitive stability during maintenance phases of the task); albeit by different mechanisms than those outlined in this review (which suggest they are related to prefrontal dopamine transmission).

Our group is one of many to suggest that the relationship between behavior and genetic polymorphisms is state dependent (Diamond et al., 2007; Mattay et al., 2003; Tunbridge, Harrison, & Weinberger, 2007). Presumably, state changes are occurring in the neurochemical systems that are mediating the relationships between these polymorphisms and behaviors, but the factors causing these changes are numerous and may include development (Wahlstrom et al., 2007), interactions between gender and stress (Diamond, 2007), or interactions between age and psychopathology (Getholf et al., 2005). Furthermore, a number of recent studies have received considerable interest because they demonstrate that early life experiences modify the risk for future psychopathology conferred by genetic polymorphisms (Arseneault et al., 2002; Caspi et al., 2002). The mechanisms underlying these findings are unknown, but it is possible that environmental influences result in changes within the neurochemical systems acted upon by the polymorphisms of interest, resulting in different risk potentials. Moreover, this model suggests that genotypic groups may vary in the quality of their life experiences based on how an individual's "neurochemical tone" provokes him/her to select experiences. These selections, in turn, may impact the continued development of the neurochemical system. It is clear that further work is necessary to clarify how environmental influences interact with genetics and underlying neurochemical systems to impact current and future behavior.

Given that our findings suggest a role for development in modulating the influence of genetic polymorphisms on behavior, future studies attempting to link genes to behavioral phenotypes should consider age as an important variable in study design. The effect sizes of individual polymorphisms on behavior are often small (Barnett, Jones, Robbins, & Müller, 2007), and meta-analyses attempting to link COMT to cognition or psychopathology have sometimes resulted in null findings (Barnett et al., 2008; Munafó, Bowes, Clark, & Flint, 2005). Given these results, attempts to link individual genes to behavior may benefit from more homogeneous samples with respect to age, which could hypothetically increase the likelihood of detecting significant relationships. Similar consideration should be given to demographic variables such as gender, which have also been shown to modulate the underlying neurochemical systems acted on by candidate genes (Diamond, 2007).

## Conclusions

Human adolescence is recognized as a period of rapid cognitive development, increased risk-taking and inconsistencies in behavioral regulation. These patterns have been attributed to structural aspects of brain development but may also be impacted by neurochemical changes that interact with these structural refinements. The dopamine system is intriguing in this respect, because it has been linked to reinforcement learning and to higher-level cognitive processes salient to cognitive control. Yet, there are few (if any) studies that have directly documented changes in DA transmission in human adolescents, most likely because of the invasiveness of the techniques necessary for this type of measurement and the related human subjects concerns. Although DA's modulation of working memory and other behavioral processes is well-established, behavioral findings alone cannot directly assess the question of whether this system demonstrates unique properties during the adolescent period. The animal literature suggests that such properties may be evident and that the system should be a focus of study in humans. Although based primarily on findings in rodents and non-human primates, the available evidence indicates that the dopamine system, which includes factors such as cell proliferation, receptor density, and metabolite concentrations, undergoes widespread and dramatic changes. Furthermore, while the data are characterized by some cross-species differences and inconsistencies across research groups, the general trends point to a system that is overactive compared to both childhood and adulthood. The implications of these changes are not fully understood, but it is likely that specific developmental alterations of dopamine transmission play an important role in both normative developmental behaviors (i.e. preferences for novelty, increased interest in risky situations such as drugs and promiscuous sex) and the onset of psychiatric disorders. We have suggested a model through which the underlying state of dopaminergic tone can be inferred through developmental changes in gene-behavior associations.

Several limitations of this presentation should be mentioned. First, our primary goal is to provide a rationale for why neurochemistry should be directly assessed in human developmental samples. The behavioral evidence is compelling in suggesting that dopamine activity changes during this lifespan period. In reviewing the animal data pertinent to dopamine system development, it is important to recognize the limitations in generalizing the findings to humans. That said, the findings are compelling in suggesting that dopamine receptor densities change throughout adolescence and that there are cortical versus subcortical distinctions. Many researchers might find limbic/subcortical dopamine activity particularly compelling as a target of study, and we agree that dopamine overactivity in this circuitry may account for adolescent risk-taking and other problem behaviors (Wahlstrom et al., 2010). Yet, we have focused our molecular genetic presentation on the COMT enzyme and prefrontally-guided working memory functions. That decision was guided by our laboratory's available data. It should be emphasized that this model is intended to use the COMT genotype as an exemplar that may apply to other polymorphisms. It is unlikely, for instance, that the COMT genotype modulates reward salience or other incentive-motivated behaviors given that the enzyme is most active in dorsal prefrontal regions (Beyer & Stekee, 2000; Carr & White, 1987). It is likely that there are other candidate genes that control for processes that are expressed more strongly in limbic and striatal regions and that could be examined for age-by-genotype interactions. For instance, the Taq1 A allele, associated with D<sub>2</sub> dopamine receptor mechanisms (Neville et al., 2004), has been associated with novelty seeking as well as with reinforcement learning (Berman et al., 2002; Frank & Hutchinson, 2009). The D<sub>4</sub> dopamine receptor and the dopamine transporter genes are also likely candidates for analysis, because individual differences in SNPs associated with these genes have been linked to variations in personality traits, externalizing behaviors, and higher-order cognition (DeYoung et al., 2006; Durston et al., 2008; Munafò et al., 2008). Moreover, as the aging and animal developmental literatures suggest, there are lifespan changes in dopamine receptor and transporter

mechanisms that impact behavior. The primary point to be made here is that genotype-by-age interactions can be informative regarding the underlying state of this neurochemical system given that more direct assessments (via psychopharmacological probes, for example) are lacking. In our view and based on success in using pharmacological probes to index neurochemical reactivity in young adults (Luciana & Collins, 1997; Luciana et al., 2004; Vrshek-Schallhorn et al., 2006), this is a technology that could be applied safely to adolescent samples. However, such an approach requires justification in the context of the current scientific climate. Perhaps neuroimaging techniques could be combined with behavioral paradigms to index differences in brain activation across age groups and genotypes to provide a more compelling rationale for the use of pharmacological probes to assess the integrity of neurochemical systems in developmental groups.

Such studies would be informative in resolving several issues. First, the precise relationship between dopamine activity and risk-taking behaviors is unclear. In addition, there is little knowledge of how changes in one facet of the dopaminergic system impact others during adolescence, how these changes are influenced by other neurochemical systems, and how these changes are influenced by or otherwise interact with structural brain changes, particularly those associated with white matter. The factors that regulate these changes are also unknown. Many may be under strong genetic control, but it could also be that differences in the onset or length of adolescent-typical events (i.e. dating, school, parenthood) have an influence on when dopaminergic changes come online or stabilize to adult levels. Unlike changes in brain structure, those that relate to neurochemistry may be more readily malleable. Accordingly, if adolescent behavior is to continue to be described as poorly regulated, as disordered in some respect, or in terms of the increased mortality that accompanies it, it should be viewed as a scientific and ethical imperative to broaden the scope of inquiry into the reasons for such behavior in humans and how we might successfully intervene.

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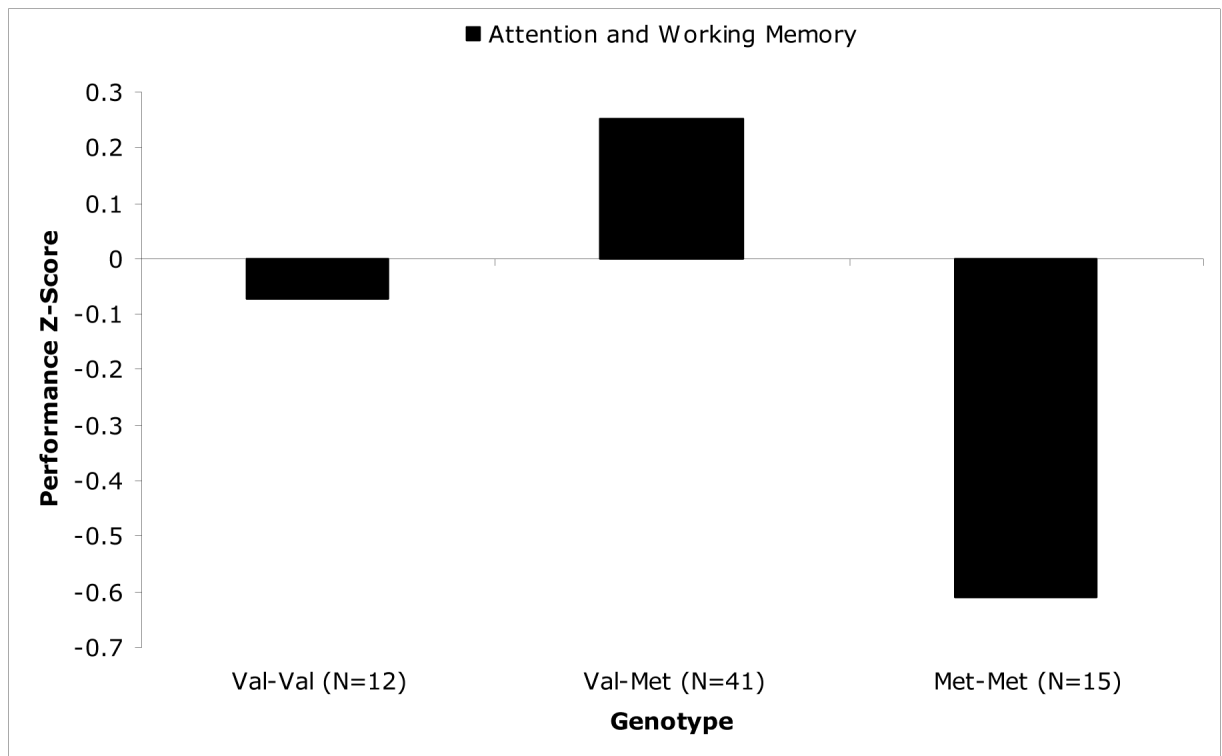
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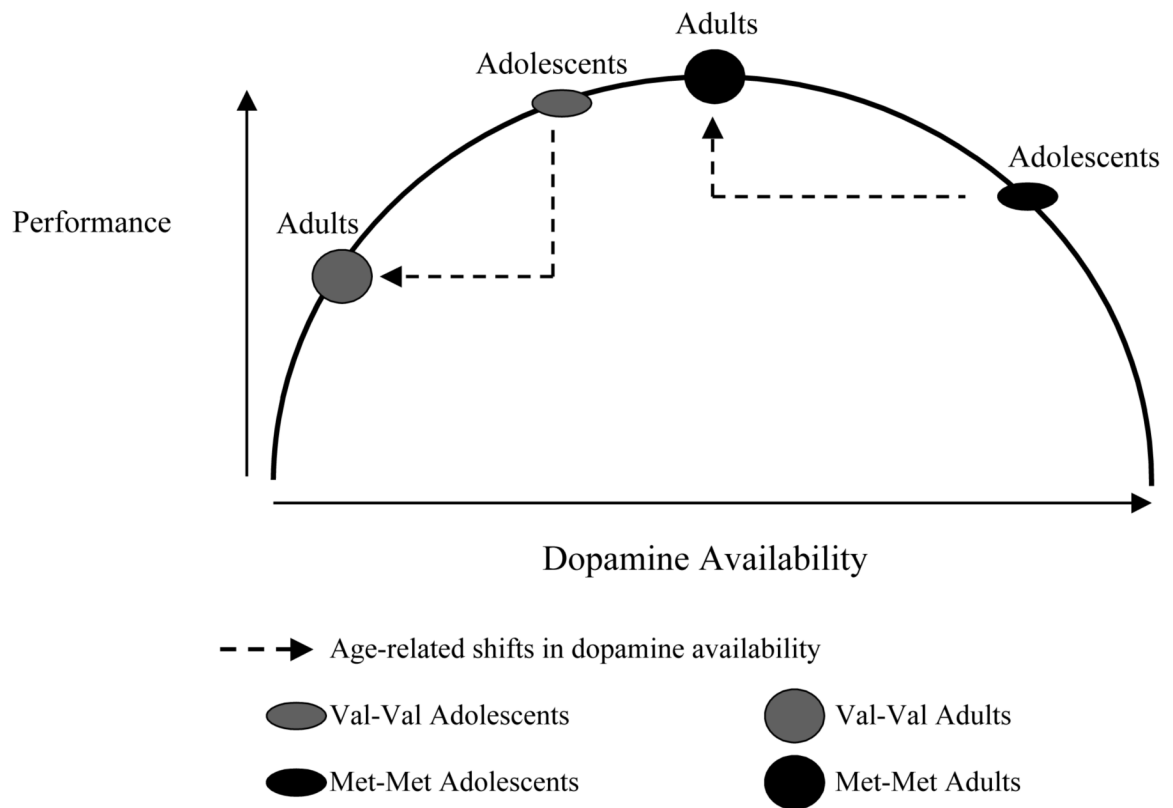
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**Figure 1.**

The COMT/Cognition Relationship in 9-17 year-old Healthy Adolescents

Adapted from Wahlstrom et al. (2007), this figure illustrates working memory and attentional performance in healthy adolescents as a function of COMT genotype. Working memory and attention were assessed through spatial delayed response, digit span, and spatial span tasks and scored as a standardized composite. The heterozygote Val/Met group performed better than either homozygote group.



**Figure 2.** Modeling the COMT-Cognition Relationship from Adolescence to Adulthood  
Changes in the relationship between COMT genotype and cognitive performance as a function of age. In adolescence, Val homozygotes are conferred functional benefits due to developmental increases in dopaminergic availability. Conversely, the performance of Met homozygotes suffers due to excess dopamine (as a result of genotype and age-related increases in dopaminergic availability), which moves them to the right of the inverted U-shaped function.

TABLE 1

Summary of major changes in dopaminergic signaling in primates and rodents. Highlighted boxes indicate evidence suggesting heightened dopamine activity compared to adulthood as discussed in the text

		Cortical	Subcortical	References
Primates	Tissue Concentrations	Peaks during adolescence	NA	Brown & Goldman, 1977; Goldman-Rakic & Brown, 1982
	Dopaminergic Innervation	Peaks in PFC layer III during adolescence	NA	Rosenberg & Lewis, 1994; 1995; Lambe et al., 2000
	D <sub>1</sub> -Type Density	Peaks in childhood. Elevated in adolescents compared to adults	Peaks in childhood. Elevated in adolescents compared to adults	Lidow et al., 1991; Lidow & Rakic, 1992; Montague et al., 1999; Seeman et al., 1987
	D <sub>2</sub> -Like Density	Peaks in childhood. Elevated in adolescents compared to adults	Peaks in childhood. Elevated in adolescents compared to adults	Lidow et al., 1991; Lidow & Rakic, 1992; Seeman et al., 1987
Rodents	Tissue Concentrations	Monotonic increases between childhood and adulthood, but adolescent peaks in dopamine synthesis	Monotonic increases between childhood and adulthood, but adolescent peaks in synaptic availability	Andersen, Dumont, & Teicher, 1997; Giorgi et al., 1987; Nomura, Naitoh, & Seqawa, 1976; Stamford, 1989; Ungethüm et al., 1996
	Dopaminergic Innervation	Monotonic increase between childhood and adulthood.	NA	Berger et al., 1985; Kalsbeek et al., 1988
	D <sub>1</sub> -Type Density	Monotonic increases between childhood and adulthood.	Peaks during periadolescence	Andersen et al., 1997; Gelbard et al., 1989; Giorgi et al., 1987; Rao et al., 1991; Tarazi et al., 1999; Tarazi & Baldessarini, 2000; Teicher et al., 1995
	D <sub>2</sub> -Like Density	Monotonic increases between childhood and adulthood.	Peaks during periadolescence	Andersen et al., 1997; Gelbard et al., 1989; Rao et al., 1991; Tarazi et al., 1998b; Teicher et al., 1995