

Review Article

Role of the Central Ascending Neurotransmitter Systems in the Psychostimulant Effects of Caffeine

Sergi Ferré*

National Institute on Drug Abuse, IRP, NIH, DHHS, Baltimore, MD, USA

Abstract. Caffeine is the most consumed psychoactive drug in the world. It is a non-selective adenosine receptor antagonist that in the brain targets mainly adenosine A₁ and A_{2A} receptors. The same as classical psychostimulants, caffeine produces motor-activating, reinforcing and arousing effects. This depends on the ability of caffeine to counteract multiple effects of adenosine in the central ascending neurotransmitter systems. Motor and reinforcing effects depend on the ability of caffeine to release pre- and postsynaptic brakes that adenosine imposes on the ascending dopaminergic system. By targeting A₁–A_{2A} receptor heteromers in striatal glutamatergic terminals and A₁ receptors in striatal dopaminergic terminals (presynaptic brake), caffeine induces glutamate-dependent and glutamate-independent release of dopamine. These presynaptic effects of caffeine are potentiated by the release of the postsynaptic brake imposed by antagonistic interactions in the striatal A_{2A}–D₂ and A₁–D₁ receptor heteromers. Arousing effects of caffeine depend on the blockade of multiple inhibitory mechanisms that adenosine, as an endogenous sleep-promoting substance, exerts on the multiply interconnected ascending arousal systems. Those mechanisms include a direct A₁-receptor mediated modulation of the corticopetal basal forebrain system and an indirect A_{2A}-receptor mediated modulation of the hypothalamic histaminergic and orexinergic systems.

Keywords: Adenosine A₁ receptor, adenosine A_{2A} receptor, ascending arousal systems, caffeine, dopaminergic system, local modules, receptor heteromers

INTRODUCTION

In the experimental animal as well as in humans, caffeine produces the same qualitative pharmacological effects as classical psychostimulants, such as cocaine and amphetamine: an increase in motor activity, arousal and reinforcing effects [1–3]. There has been some resistance to the claim that caffeine exerts reinforcing effects, in spite of the fact that its considerable worldwide consumption provides compelling circumstantial evidence. Nevertheless, there is enough unam-

biguous experimental evidence indicating that caffeine functions as a reinforcer under certain conditions both in laboratory animals and humans [2,4]. However, it is important to point out that caffeine has a weaker reinforcing efficacy than classical psychostimulants [2,4]. In fact, caffeine users often fulfill the criteria for drug dependence, but not for drug abuse, established by the Diagnostic and Statistical Manual of Mental Disorders (DSM IV; 4th edition) [2]. Furthermore, in humans, caffeine produces similar subjective stimuli to classical psychostimulants [2] and, in the experimental animal, caffeine and classical psychostimulants produce similar discriminative-stimulus effects [5–7].

Caffeine is a non-selective competitive adenosine receptor antagonist and produces its psychostimulant effects by counteracting the tonic effects of endogenous

*Correspondence to: Sergi Ferré, National Institute on Drug Abuse, IRP, NIH, DHHS, 251 Bayview Blvd, Baltimore, MD 21224, USA. Tel.: +1 443 740 2647; Fax: +1 443 740 2816; E-mail: sferre@ntra.nida.nih.gov.

adenosine on central adenosine receptors. This depends largely on the ability of adenosine to modulate the function of multiple central ascending neurotransmitter systems, which are involved in motor activation and reward (dopaminergic systems) and arousal (cholinergic, norepinephrine, histaminergic, orexinergic systems). Among the four cloned adenosine receptors (A_1 , A_{2A} , A_{2B} and A_3 receptors), A_1 and A_{2A} receptors are the ones predominantly expressed in the brain. Caffeine has similar *in vitro* affinities for A_1 , A_{2A} and A_{2B} receptors and much lower affinity for A_3 receptors [8]. A_1 and A_{2A} receptors are the preferential targets for caffeine in the brain, since physiological extracellular levels of adenosine are sufficient to occupy and, therefore, stimulate A_1 and A_{2A} receptors. On the other hand, A_{2B} receptors have a lower affinity for adenosine and are only activated by high pathological extracellular levels of adenosine [7,8]. A_1 receptors are widely expressed in the brain, including the striatum, while A_{2A} receptors are highly concentrated in the striatum [8,9]. The striatal localization of both receptors seems to underlie the motor-activating and reinforcing effects of caffeine, which depend on the adenosine-mediated modulation of the ascending dopaminergic system. On the other hand, A_1 receptors localized in the basal forebrain and A_{2A} receptors localized in the hypothalamus are believed to be mostly responsible for the arousing properties of caffeine, which depend on the adenosine-mediated modulation of the ascending arousal systems.

CAFFEINE AND THE ASCENDING DOPAMINERGIC SYSTEM

An important amount of experimental evidence supports a key role of dopamine in the psychostimulant effects of caffeine in animals and humans. For instance, dopamine depletion or blockade of dopamine receptors significantly impairs the motor and discriminative stimulus effects of caffeine [10,11]. Most probably the same basic dopamine-mediated mechanisms are involved in the motor-activating and reinforcing effects of caffeine, as it happens with classical psychostimulants [12,13]. The key to understand these mechanisms of action is understanding how adenosine modulates dopaminergic neurotransmission in the brain.

The striatum and the striatal spine module

The ascending dopaminergic systems originate in the mesencephalon (substantia nigra and ventral tegmen-

tal area) and innervate the striatum (caudate-putamen, nucleus accumbens and olfactory tubercle), cortex (mostly prefrontal), amygdala, and hippocampus [14]. These systems are involved in motor activation and learning and expression of goal-directed behaviors. The striatum is the main input structure of basal ganglia and contains the highest innervation of dopamine and the highest density of dopamine receptors in the brain [14]. In the striatum, the GABAergic striatal efferent neuron, also called the medium spiny neuron (MSN), constitutes more than 95% of the striatal neuronal population [14]. There are two subtypes of MSNs, which selectively express one of two peptides, enkephalin or dynorphin. Enkephalinergic MSNs predominantly express dopamine D_2 and A_{2A} receptors, while dynorphinergic MSNs predominantly express dopamine D_1 receptors and adenosine receptors of the A_1 subtype [15,16] (Fig. 1). The MSN receives two main afferents: glutamatergic afferents from cortical, thalamic, and limbic areas and the dopaminergic afferents from the mesencephalon. Both sets of afferents converge on the dendritic spine of the GABAergic efferent neuron. Glutamatergic and dopaminergic terminals make preferential synaptic contacts with the head and the neck of the dendritic spine, respectively [14,16] (Fig. 1). The dendritic spine, the glutamatergic terminal, the dopaminergic terminal and astroglial processes that wrap the glutamatergic synapse constitute the most common local module in the striatum, which we recently named striatal spine module (Fig. 1). We define "local module" as the minimal portion of one or more neurons and/or one or more glial cells that operates as an independent integrative unit [16]. The segregation of dopamine and adenosine receptors in the two subtypes of medium spiny neurons implies the existence of at least two subtypes of striatal spine module (Fig. 1).

Adenosine is a key modulator of dopaminergic and glutamatergic neurotransmission in the striatal local module. Recent studies indicate that astroglia plays a fundamental role in the formation of extracellular adenosine which can influence synaptic transmission. Astrocytes express glutamate receptors (mostly metabotropic) and ATP receptors which, when activated, induce astrocytes to release glutamate and ATP [17,18]. Astroglial-released ATP is then converted to adenosine in the extracellular space by means of ectonucleotidases [19]. Furthermore, there also seems to be a neurotransmitter-like formation of adenosine, a synaptic pool of adenosine. Adenosine can be produced from ATP co-released with glutamate, which

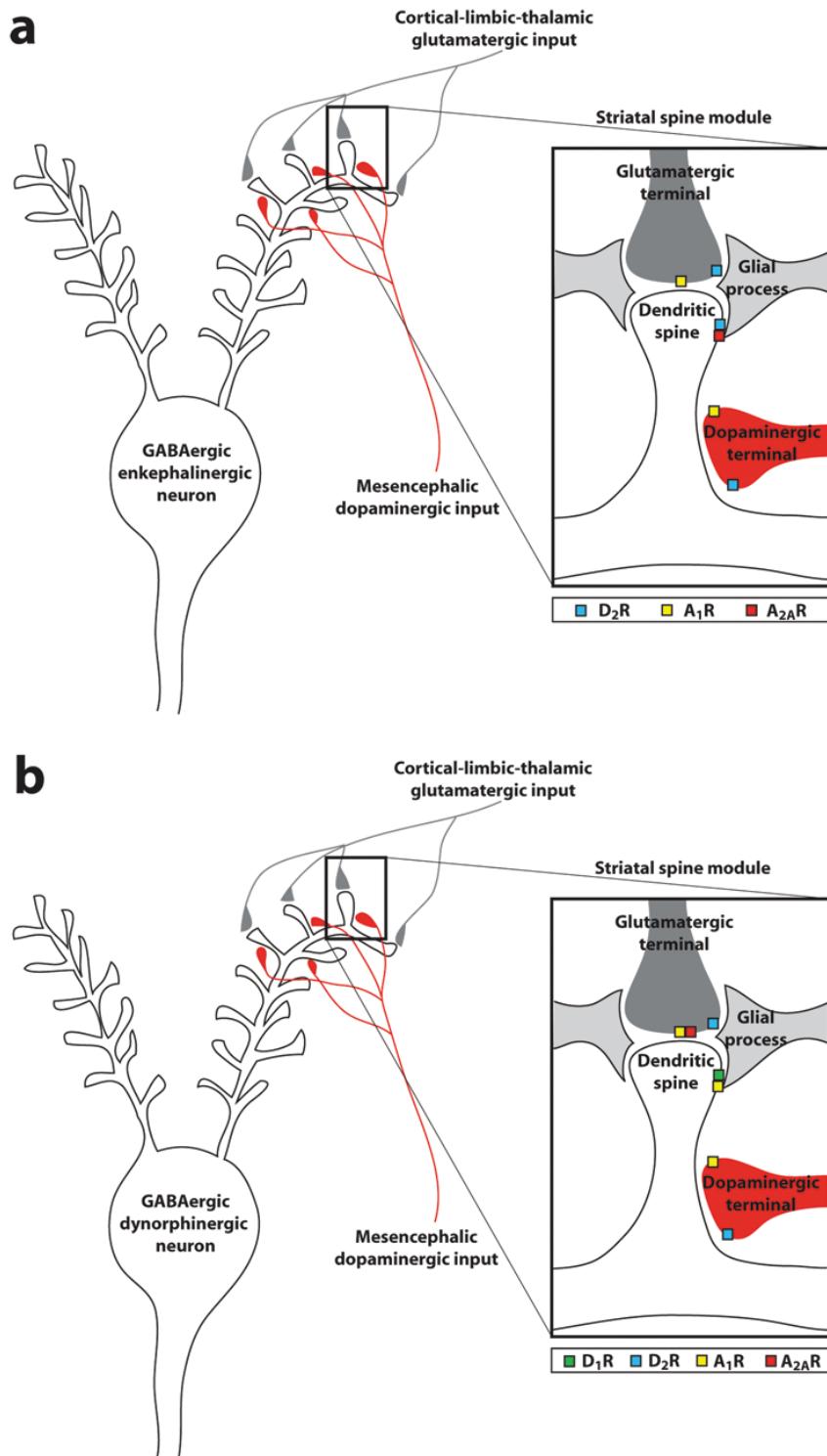


Fig. 1. Caffeine targets in the striatal spine modules of the GABAergic enkephalinergic neuron (a) and GABAergic dynorphinergic neuron (b). Adenosine A_{2A}-dopamine D₂ receptor heteromers are localized in the dendritic spines of the GABAergic enkephalinergic neuron and adenosine A₁-dopamine D₁ receptor heteromers are localized in the dendritic spines of the GABAergic dynorphinergic neurons. Presynaptic A_{2A} receptors form heteromers with A₁ receptors and are localized in the striatal glutamatergic terminals that make contact with the dendritic spines of the GABAergic dynorphinergic neuron. A₁ receptors are also localized in a fraction of dopaminergic terminals (modified from ref. [16]).

is also metabolized to adenosine by means of ectonucleotidases [20]. In the striatal spine module, A_{2A} receptors are localized postsynaptically in the dendritic spine of enkephalinergic MSNs, colocalized with D₂ receptors and presynaptically in glutamatergic terminals to the dynorphinergic MSNs [3,9,15–17,20–23] (Fig. 1). Dopamine D₂ receptors are also localized in dopaminergic and glutamatergic terminals, where they do not seem to form heteromers with A_{2A} receptors [24]. A_{2A} receptors are also expressed in dopaminergic and GABAergic terminals, but with lower density than postsynaptic A_{2A} receptors in enkephalinergic MSN and presynaptic A_{2A} receptors in glutamatergic terminals, [25]. Thus, some studies suggest the existence of functional striatal A_{2A} receptors in the dopaminergic terminals, which do not directly modulate dopamine release, but that control dopamine release induced by glial cell line-derived neurotrophic factor (GDNF) [26,27]. Recent studies support the existence of A_{2A} receptors in the striatal collaterals of the enkephalinergic MSNs, activation of which facilitates GABA release [28]. Less clear is the significance of some studies suggesting that activation of striatal A_{2A} receptors in striatal GABA terminals inhibits GABA release [29]. Finally, functional A_{2A} receptors have also been suggested to be localized in nerve terminals from cholinergic interneurons (which should also be included in the striatal spine modules, mostly as asymmetric varicosities that release acetylcoline by volume transmission [16]) where they are co-localized with A₁ receptors [30]. Presynaptically, A₁ receptors are localized in glutamatergic terminals, colocalized with A_{2A} receptors in those terminals which contact dynorphinergic MSNs, and in a fraction of dopaminergic nerve terminals [21,22,31]. Postsynaptically, A₁ receptors are localized in the dendritic spine of dynorphinergic MSNs, colocalized with D₁ receptors [15,20] (Fig. 1). In order to understand the role of adenosine in the striatum, we have to understand the role that the different adenosine receptor subtypes play in the different elements of the striatal spine module.

Postsynaptic mechanisms: Adenosine-dopamine receptor heteromers

In the striatal spine, antagonistic interactions between A_{2A} and D₂ receptors modulate the function of the enkephalinergic MSN, and antagonistic interactions between A₁ and D₁ receptors modulate the function of the dynorphinergic MSN [15,16,20]. This gives the explanation at the neuronal level of an important

number of pharmacological findings indicating a selective modulation of A₁ and A_{2A} receptor ligands on D₁ and D₂ receptor agonists-induced behavioral effects, respectively. Thus, A₁ and A_{2A} receptor agonists counteract, respectively, the motor activating effects induced by dopamine D₁ and D₂ receptor agonists. Similarly, A₁ and A_{2A} receptor antagonists potentiate, respectively, the motor activating effects induced by dopamine D₁ and D₂ receptor agonists [15, 16,20]. The molecular mechanisms responsible for the selective antagonistic A₁-D₁ and A_{2A}-D₂ receptor interactions seem to depend on allosteric interactions between adenosine and dopamine receptors forming receptor heteromers. In fact, there is compelling evidence for the existence of A₁-D₁ and A_{2A}-D₂ receptor heteromers in artificial cell systems and in the striatum [16, 20,24,32]. In the A_{2A}-D₂ receptor heteromer, the stimulation of the A_{2A} receptor decreases the binding of dopamine to the D₂ receptor [33]. This allosteric interaction in the A_{2A}-D₂ receptor heteromer controls neuronal excitability and, consequently, neuronal firing and neurotransmitter (GABA) release by the enkephalinergic MSN [34,35]. This is related to the ability of D₂ receptors to suppress Ca²⁺ currents through L-type voltage-dependent calcium channels by a cAMP-protein kinase A (PKA)-independent and phospholipase C (PLC)-dependent signaling pathway [24,32]. Thus, stimulation of striatal A_{2A} receptor does not produce a significant effect on its own, but it strongly counteracts the depressant effects of D₂ receptor stimulation on neuronal firing and neurotransmitter release in the enkephalinergic MSN [34,35].

In addition to the allosteric interaction in the A_{2A}-D₂ receptor heteromer, a reciprocal antagonistic interaction between A_{2A} and D₂ receptors has been found at the second messenger level. In this case stimulation of D₂ receptors counteracts the activation of adenylyl-cyclase induced by stimulation of A_{2A} receptors [20,24,32]. Stimulation of A_{2A} receptor, through its coupling to G_{olf} proteins, can potentially stimulate adenylyl-cyclase, with consequent activation of cAMP-PKA signaling pathway and induction of the expression of different genes, such as *c-fos* and *preproenkephalin*, by activating the constitutive transcription factor cAMP response element binding protein (CREB) [20,24,32]. A_{2A} receptor-mediated activation of PKA can also induce phosphorylation of dopamine and cAMP-regulated phosphoprotein 32 kDa (DARPP-32) at threonine 34 (Thr₃₄) and a phosphatase 2-mediated dephosphorylation of DARPP-32 at Thr75 [20,24,32]. Furthermore, A_{2A} receptor-mediated activation of PKA

can phosphorylate AMPA glutamate receptors, which play a crucial role in the initial plastic changes of glutamatergic synapses, which includes synaptic recruitment of AMPA receptors [20,24,32]. However, under basal conditions, stimulation of A_{2A} receptors can poorly activate cAMP-PKA signaling and increase gene expression, due to a strong tonic inhibitory effect of endogenous dopamine and D₂ receptor stimulation on adenylyl-cyclase [20,24,32]. D₂ receptor blockade is then necessary to reveal the effects of a tonic activation by endogenous adenosine on A_{2A} receptor-mediated PKA activation [20,24,32], which indicates that blockade of the A_{2A} receptor-mediated stimulation of cAMP-PKA signaling unlikely contributes to the acute behavioral effects of caffeine, as previously suggested [36]. It is most probable that the coexistence of two apparently incompatible A_{2A}-D₂ receptor interactions in the enkephalinergic MSN depends on the existence of a pool of A_{2A} and D₂ receptors forming heteromers, and a pool of A_{2A} and D₂ receptors forming homodimers. When forming homodimers, D₂ receptor couples preferentially to G_i, which allows its coupling to adenylyl cyclase. When forming heteromers with the A_{2A} receptor, the D₂ receptor cannot activate G_i, because a key epitope involved in Gi activation (an arginine-rich epitope localized in the amino-terminus of the third intracellular loop of the D₂ receptor) is also involved in A_{2A}-D₂ receptor heteromerization [37]. In this case, the D₂ receptor probably couples to G_{q/11} with the consequent activation of the PLC pathway [24]. This would be a similar situation to that recently described for the D₁-D₂ receptor heteromer [38].

The allosteric interaction in the A_{2A}-D₂ receptor heteromer seems to play a key role in the motor-activating effects of caffeine. In fact, this A_{2A}-D₂ receptor interaction has been given a lot of attention in the literature, more recently with the application of A_{2A} receptor antagonists as an adjuvant therapy for L-DOPA in Parkinson's disease (for recent review, see ref. [39]). But, A₁-D₁ receptor interactions are also of significant functional and pharmacological importance. Thus, as mentioned before, A₁ receptor antagonists potentiate the motor activating effects of D₁ receptor stimulation [40,41]. Similarly to what happens with A_{2A} and D₂ receptors, antagonistic interactions take place in the A₁-D₁ receptor heteromer and at the second messenger level. In this case, the interactions are not reciprocal, and stimulation of A₁ receptors inhibits both the binding of dopamine to the D₁ receptor [42,43] and the D₁ receptor-mediated activation of cAMP-PKA signaling pathway and the expression of genes, such as *c-fos* and *preprodynorphin* in the dynorphinergic MSN [43,44].

Presynaptic mechanisms: Dopamine release

In addition to the postsynaptic mechanisms related to the A_{2A}-D₂ and A₁-D₁ receptor interactions, adenosine also acts presynaptically in the striatal spine module, modulating glutamate and dopamine release. Different studies have repeatedly shown that A₁ and A_{2A} receptors exert opposite modulatory roles on extracellular levels of glutamate and dopamine in the striatum, with activation of A₁ receptors inhibiting and activation of A_{2A} receptors stimulating glutamate and dopamine release (reviewed in ref. [3]). More recently, we found that systemic or striatal administration of caffeine or an A₁ but not an A_{2A} receptor antagonist produces a significant increase in the extracellular concentrations of glutamate and dopamine in the ventral striatum, particularly in the most medial part, the medial shell of the nucleus accumbens [45,46]. It was hypothesized that dopamine release was mostly dependent on glutamate release induced by blockade of A₁ receptors localized in glutamatergic terminals and on stimulation of ionotropic glutamate receptors localized in dopaminergic terminals [46]. The ability of caffeine to release dopamine in the nucleus accumbens was questioned by another research group [47], but a recent study demonstrated the existence of subregional differences in the effect of A₁ receptor blockade in different parts of the nucleus accumbens and other striatal areas, most probably related to subregional differences in the level of tonic activation by endogenous adenosine [31]. Furthermore, glutamate-independent mechanisms were also found to be involved in A₁ receptor blockade-mediated striatal dopamine release, which depends on A₁ receptors localized in dopaminergic nerve terminals [31].

It has been demonstrated with electron-microscopy and immunocytochemical experiments that a high proportion of glutamatergic nerve terminals contain both A₁ and A_{2A} receptors [21]. Recent experiments indicate that A_{2A} receptors are in fact localized predominantly in glutamatergic terminals that establish contact with dynorphinergic MSNs, which constitute the so-called direct striatal efferent pathway [23]. In functional studies with striatal nerve terminal preparations, stimulation of the A₁ receptor was found to decrease while stimulation of A_{2A} receptors potentiated potassium-induced glutamate release [21]. Importantly, when both A₁ and A_{2A} receptors were stimulated, the response was not a counteractive effect, but the same as results from A_{2A} receptor stimulation, i.e., a potentiation of glutamate release [21]. Furthermore, in the same kind of preparations, low concentrations of

adenosine inhibited while high concentrations stimulated glutamate release [21]. This would agree with the reported higher affinity for adenosine of the A₁ compared to the A_{2A} receptor [48] and would provide a mechanism for a fine-tuned modulation of glutamate release by adenosine, with low concentrations inhibiting and high concentrations stimulating glutamatergic neurotransmission to the direct striatal efferent pathway. The mechanism by which A_{2A} receptor stimulation shuts down the effects of A₁ receptor stimulation seems to be related to an allosteric interaction in A₁-A_{2A} receptor heteromers. Thus, A₁ and A_{2A} receptors have been shown to form heteromers in transfected cells [21]. In membrane preparations from transfected cells and from rat striatum, stimulation of A_{2A} receptors decreases the affinity of A₁ receptor for agonist binding [21].

The question arises as to which is the contribution of A₁ and A_{2A} receptors and the many pre- and postsynaptic mechanisms in the psychostimulant effects of caffeine. There has been a long-running debate about the preferential involvement of A₁ and A_{2A} receptors in the motor-activating effects of caffeine (reviewed in refs. [3,22]). Particularly influential were the experiments with A_{2A} receptor knockout (A_{2A}KO) mice, which showed a lack of motor-activating effects of caffeine [49]. However, by comparing both quantitative and qualitative aspects of the motor activity induced by caffeine and selective A₁ and A_{2A} receptor antagonists, recent studies have clearly shown that caffeine, when administered acutely, shows a profile of a non-selective adenosine receptor antagonist with even a preferential A₁ receptor antagonism [50,51]. Importantly, chronic exposure to caffeine differentially modifies its motor effects dependent on A₁ and A_{2A} receptor blockade. Thus, chronic exposure to caffeine in the drinking water of rats results in partial tolerance to the motor effects of an additional acute administration of caffeine, and total cross-tolerance to the motor effects of an A₁ but not an A_{2A} receptor antagonist [50]. This indicates that tolerance to the effects of A₁ receptor blockade is mostly responsible for the tolerance to the motor-activating effects of caffeine, and that the residual motor-activating effects of caffeine in tolerant individuals might be largely because of A_{2A} receptor blockade [50]. Numerous experimental findings indicate that dopamine release in the medial striatal compartments is involved in invigoration of approach and in some aspects of incentive learning (for recent review, see ref. [13]). In relation to psychostimulants, dopamine release in the very medial striatal compartments seems to be involved in

both their motor-activating and reinforcing effects [13, 52]. Therefore, the pre- and postsynaptic dopaminergic mechanisms mentioned above (striatal dopamine release and adenosine-dopamine receptor-receptor interactions) taking place in the medial striatal compartments are most probably involved in the motor and reinforcing effects of caffeine. Importantly, chronic administration of caffeine in the drinking water completely counteracted the effects of caffeine or an A₁ receptor antagonist on dopamine and glutamate release, while the effect of an A_{2A} receptor antagonist was not modified [53]. Thus, these biochemical changes correlate with the studies on motor activity [50], strongly suggesting the involvement of presynaptic mechanisms in the psychostimulant effects of caffeine. In relation to the dopamine-releasing effects, at least two factors could explain the weaker reinforcing properties of caffeine as compared with other psychostimulants: its specific subregional effects [31] and the development of tolerance [50,53].

In summary, caffeine produces its motor and reinforcing effects by releasing the pre- and postsynaptic brakes that adenosine imposes on dopaminergic neurotransmission in the striatal spine module. By targeting A₁-A_{2A} receptor heteromers in glutamatergic terminals and A₁ receptors in dopaminergic terminals (presynaptic brake), caffeine induces glutamate-dependent and glutamate-independent release of dopamine. These presynaptic effects of caffeine are potentiated by the release of the postsynaptic brake imposed by antagonistic adenosine-dopamine receptor-receptor interactions in the A_{2A}-D₂ and A₁-D₁ receptor heteromers. However, it is important to point out the possible additional contribution coming from A₁ receptors localized in the nuclei of origin of the ascending dopaminergic systems [54,55].

CAFFEINE AND THE ASCENDING AROUSAL SYSTEMS

Arousal is a state of behavioral readiness in response to sensory stimulation, which is associated with cortical electroencephalographic (EEG) activation, which depends on the activation of ascending arousal systems localized in the pontomesencephalic tegmentum, basal forebrain and hypothalamus. The arousal-enhancing properties of caffeine are mostly dopamine-independent and they are related to its ability to antagonize the sleep-promoting effects of adenosine. To understand the mechanisms involved in caffeine-induced arousal we have to understand how adenosine modulates the function of the ascending arousal systems.

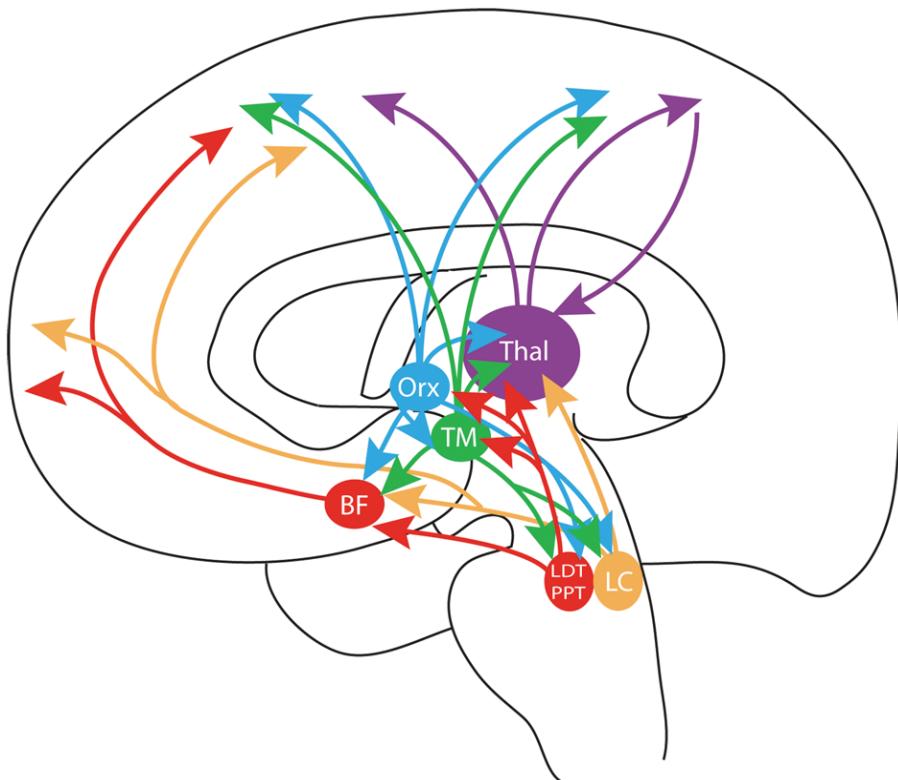


Fig. 2. Scheme of the highly interconnected multiple ascending arousal systems, which are all targets of caffeine (see text). BF: area of origin of the corticopetal basal forebrain system; LC: locus coeruleus; LDT and PPT: laterodorsal and pedunculopontine nuclei; TM: tuberomammillary nucleus; Orx: area of origin of the orexinergic system; Thal: thalamus.

Multiply interconnected ascending arousal systems

Moruzzi and Magoun, in 1949, were the first to demonstrate that activation of the pontomesencephalic tegmentum, what they called the ‘ascending reticular activating system’, is associated with arousal [56]. We know now that the ascending reticular activating system is heterogeneous and contains interconnected ascending glutamatergic, cholinergic and noradrenergic pathways [57–59] (Fig. 2). A main characteristic of these chemically defined ascending systems, as well as of the corticopetal basal forebrain system and the hypothalamic ascending systems (see below), is their ability to modulate simultaneously large areas of the brain and synchronize high-frequency cortical activity during the wakefulness period. This characteristic is related to their morphological and functional properties. First, these systems have a relatively low number of cells of origin with widespread branching terminal processes. Second, a synchronization of activity takes place between their cells of origin [60]. Third, their terminals often contain varicosities without synaptic spe-

cializations [61] which release neurotransmitters in a “volume transmission” mode [62].

The physiological activation of the ascending reticular system depends on multiple rostral and caudal inputs carrying motor, limbic, visceral and sensorial multimodal information. It even receives an important cortical input from the prefrontal cortex [63,64]. The different involvement in cortico-subcortical activation induced by the different ascending glutamatergic, cholinergic and noradrenergic pathways is determined by their preferential target innervation, as well as the different inputs to their pontomesencephalic nuclei of origin [57–59]. Glutamatergic cells of the reticular formation project mostly to the thalamus, to the midline-intralaminar and reticular nuclei, which transmit the activity of the ascending reticular activating system to extensive areas of the cerebral cortex. Most cholinergic cells originate in the laterodorsal and pedunculopontine nuclei (LDT and PPT nuclei), which through a dorsal tegmental pathway also project heavily to the thalamus (Fig. 2). A ventral tegmental pathway forms the extrathalamic cholinergic relay to the cortex and in-

nerves the cells that are the origin of the corticopetal basal forebrain system (see below) (Fig. 2). It must be pointed out that PPT nucleus is not only cholinergic, but that also contains non-cholinergic cells with extensive projections [65]. In fact, the projection from the PPT nucleus is considered to be largely non-cholinergic and, most probably glutamatergic [66]. Furthermore, some studies indicate that nitric oxide (NO) is an important neuronal messenger of the cholinergic cells of the ascending reticular activating system [67]. Noradrenergic cells originate mostly in the locus coeruleus, just lateral to the LDT nucleus, and project in a diffuse manner to many brain areas, including the thalamus, amygdala, the corticopetal basal forebrain system, and the cerebral cortex [59] (Fig. 2).

The firing of cholinergic neurons of the LDT and PPT nuclei is temporally correlated with cortical activation. The activity of the cholinergic cells decreases during slow-wave sleep and increases again during fast-wave sleep [58]. The activity of brainstem cholinergic neurons during the waking state is not stimulus-specific. This is consistent with their multiple connections with adjacent ascending fiber systems in the reticular formation, as well as with their extensive afferent connections from subcortical limbic (nucleus accumbens) and motor brain areas (substantia nigra pars reticulata, globus pallidus). Animals with PPT nuclei lesions suffer from a global impairment in attention, rather than deficits of specific attention-related tasks [68]. Similarly to the brainstem cholinergic cells, noradrenergic cells of the nucleus coeruleus are also more active during the waking state and less active during slow wave-sleep. However, their activity ceases during fast-wave sleep [58]. During the waking state the activity of locus coeruleus cells is decreased when the animal is engaged in automatic, consumatory, non-goal directed motor activity, where it is mostly inattentive to extrapersonal environmental stimuli [69]. Furthermore, different from the brainstem cholinergic cells, the activity of the noradrenergic cells during the waking state is stimulus-specific. Thus, they increase their activity in response to novel or stressful “alerting” stimuli, with the concomitant general increase in cortical activity. This is related to the afferent connections of the nucleus coeruleus from nucleus paragigantocellularis and nucleus prepositus (both located in the rostral medulla) and with the amygdala, brain structures that convey information from emotionally arousing stimuli [70–72]. Animals with lesions of the ascending noradrenergic pathway experience difficulties in some tasks requiring attention, particularly those involving “alerting” stim-

uli [59]. Furthermore, there is experimental evidence suggesting that locus coeruleus also plays an important role in attending to non-stressful motivational stimuli [60]. Motivational stimuli most probably influence locus coeruleus by means of its afferent connections with amygdala and prefrontal cortex [71,73].

The corticopetal basal forebrain system is considered as the major extrathalamic relay of the ascending reticular activating system to the cortex [58,74,75]. This system represents a continuous stream of cells that extends from the medial septum, through the diagonal band into the substantia innominata, ventral pallidum and nucleus basalis of Meynert [57]. Although it was initially suggested that in both rodents and primates most neurons from the basal forebrain projecting to the cortex were cholinergic, it is now known that in rodents a roughly equal number of basal forebrain GABAergic and cholinergic cells project to cortical areas [76, 77]. Finally, there is also evidence for a glutamatergic component [78]. The corticopetal basal forebrain system innervates the entire telencephalic mantle, including the cerebral cortex, hippocampal formation and amygdala. Main inputs of this system come also from these areas, although the cortical input is restricted to the prefrontal cortex and seems to target preferentially non-cholinergic cells [76,79]. Also, important afferent connections originate in the ventral striatum (motor input), the hypothalamus, the ascending serotonergic and dopaminergic systems and the different components of the ascending reticular activating system [80], and the ascending hypothalamic arousal systems [81, 82] (Fig. 2). The corticopetal basal forebrain system plays a decisive role in cortical activation. Although the predominantly cortical excitatory role of acetylcholine is well established, GABA is basically an inhibitory neurotransmitter. Nevertheless, GABAergic cells of the corticopetal basal forebrain system induce cortical activation by disinhibition, by targeting the cortical system of GABAergic interneurons [83]. Activity of its cells correlates with cortical activation during waking and fast-wave sleep. During the waking state, the activity of the cells of the corticopetal basal forebrain system is stimulus-specific and specially associated with motivational stimuli [84–86] and motor activity [87]. This is most probably related, first, to the very marked glutamatergic afferent connections from the amygdala and prefrontal cortex, where motivational stimuli are represented [88]. Second, the motor activity-related changes in cell activity can be mediated by the direct connections from the ventral striatum and the indirect connections from the basal ganglia, through the

ascending reticular activating system [74,87]. There is an increasing number of experimental data suggesting the existence of anatomical and functional segregated subsystems in the corticopetal basal forebrain, with distinct afferent-efferent connections such as the GABAergic subsystem, that selectively receives input from the prefrontal cortex. These subsystems should afford different roles in different attention processes [76,77].

Two more ascending arousal systems originate in the hypothalamic area: the histaminergic and orexinergic systems, which originate in the caudal and dorsolateral hypothalamus, respectively [81,82,89–91]. The same as the noradrenergic (and also the serotoninergic) system, histaminergic and orexinergic cells are active during the waking state, particularly active waking, and virtually inactive during slow and fast-wave sleep [92–94]. Histaminergic cells originate in the tuberomammillary (TM) nuclei of the hypothalamus and innervate diffusely most areas of the brain, including the thalamus and cortex [81,89]. TM nuclei receive a main GABAergic-galaninergic inhibitory input from the ventrolateral-preoptic (VLPO) area of the anterior hypothalamus, considered as a main sleep center. Furthermore, TM cells reciprocally connect with LDT and PPT nuclei and project to the origin of the corticopetal basal forebrain system and to the locus coeruleus [81,92] (Fig. 2).

Orexinergic neurons are localized in the perifornical lateral hypothalamic area, extending medially to the dorsomedial hypothalamic nucleus and sending projections to the midline hypothalamic nuclei, basal forebrain, and cortex, with a very similar distribution to that of the noradrenergic system, with which it is heavily interconnected [82,91]. Their most prominent projections include many hypothalamic areas (including TM nucleus), locus coeruleus (the densest extrahypothalamic projection), basal forebrain, dorsal raphe nucleus, LDT and PPT nuclei, and substantia nigra and ventral tegmental area [82]. The connections with the histaminergic system seem to be particularly relevant and, in fact, several studies strongly suggest that the arousal effects of orexin depend on the activation of the histaminergic system [95,96]. Orexin neurons also receive direct projections from the hypothalamus, including TM nucleus, VLPO area and the suprachiasmatic nucleus, which is critically involved in the generation of circadian rhythm [97] (Fig. 2).

The ascending dopaminergic and serotoninergic pathways, which also originate in the brainstem and innervate diffusely many brain areas including the cerebral cortex, are often included as part of the ascending

reticular activating system (see, for instance, ref. [59]). However, these systems are not ‘directly’ involved in arousal or attention. Nevertheless, the activity of serotonergic cells (but not dopaminergic cells) is state-dependent, and their firing is maximal during the waking state, decreases during slow-wave sleep and stops during fast-wave sleep. During the wakefulness period, the activity of the ascending serotonergic cells, which originate in the dorsal and median raphe nuclei, does not vary with any kind of sensorial stressful or non-stressful stimulation. On the other hand, just opposite to what happens with locus coeruleus cells, they increase their activity during automatic, consumatory, non-goal directed motor activity [98].

Adenosinergic modulation of the ascending arousal systems

An important amount of evidence indicates that adenosine is a mediator of sleepiness following prolonged wakefulness. Initial evidence came from pharmacological studies showing the sleep-inducing effects of systemic or intracerebral administration of adenosine and adenosine receptor agonists (reviewed in ref. [99]). These results, together with the fact that adenosine is a byproduct of energy metabolism and that energy restoration seems to be one of the main functions of sleep, led to the hypothesis that adenosine may serve as a homeostatic regulator of energy. Although it was initially thought that adenosine would accumulate in the extracellular space as a function of neuronal activity, either following its equilibrative transport or after metabolism from released ATP, recent studies indicate that sleep homeostasis depends largely upon gliotransmission. Thus, inhibiting gliotransmission, and therefore glial ATP release (ATP and glutamate are the main glial neurotransmitters), attenuated the sleep pressure following sleep deprivation and prevented the arousing effects of A₁ receptor antagonists [100].

After prolonged wakefulness in the cat, adenosine was found to accumulate in the basal forebrain and to a lesser extent in the cerebral cortex [101]. This accumulation was brain region-specific and adenosine was not found to increase in thalamus, preoptic hypothalamic area, PPT, or dorsal raphe [101]. The rise of adenosine levels inhibits the firing of the cells of origin of the corticopetal basal forebrain system, and experiments with selective ligands indicate that A₁ receptors are preferentially involved [99,102]. Nevertheless the activity of LDT/PPT cells is also under an inhibitory local tone by endogenous adenosine through the acti-

vation of A₁ receptors, probably localized both postsynaptically and presynaptically in glutamatergic terminals [103,104]. Furthermore, recent experiments indicate that stimulation and blockade of A₁ receptors in the prefrontal cortex decreases and increases arousal, respectively, and that this modulation is mediated by the descending connection from the prefrontal cortex to the PPT/LTD [105].

However, the sleep-generating effects of adenosine entail additional and not less important modulations of the other ascending arousal systems. As mentioned before, the hypothalamic VLPO area (and the adjacent median preoptic nucleus) is an important sleep center, and anatomical and physiological evidences suggest that the preoptic area neurons promote sleep via descending inhibitory modulation of the arousal system. The GABAergic-galaninergic cells of the VLPO project heavily to the histaminergic TM nuclei, and electrical stimulation of the VLPO area inhibits neuronal excitability of the histaminergic cells [106]. Physiologically, VLPO area is activated by endogenous sleep promoting substances that accumulate during the waking state, including adenosine and prostaglandin D₂ (PGD₂) [107,108]. Activation by adenosine is mediated by A_{2A} receptors localized in a subset of VLPO neurons [109], and the sleep-enhancing effects of PGD₂ also seem to require activation of VLPO sleep regulatory neurons by adenosine acting at A_{2A} receptors [109]. Apart from the connection to the TM nuclei, GABAergic cells from the VLPO also project to the locus ceruleus and the perifornical lateral hypothalamic area and several experimental findings suggest that deactivation of the histaminergic, noradrenergic and orexinergic arousal systems occurring at sleep onset and during the fast- and slow-wave sleep results from GABA-mediated inhibition originating in the preoptic hypothalamus [108].

In summary, it is now generally believed that the direct A₁-receptor mediated modulation of the corticopetal basal forebrain system and the indirect A_{2A}-receptor mediated modulation of the hypothalamic histaminergic and orexinergic systems are the main mechanisms by which adenosine exerts its sleep-promoting effects [102,108]. However, it is important to point out the possible additional contribution coming from A₁ receptors localized in the nuclei of origin of the histaminergic, orexinergic and noradrenergic arousal systems [110–112].

The same as for the motor-activating effects of caffeine, there has been a strong debate about the role of A₁ and A_{2A} receptors in its arousing effects. Sim-

ilarly to the studies on motor activation, A₁ receptors (in the basal forebrain) were initially the main candidates (reviewed in ref. [102]), but again experiments performed in A_{2A}KO mice strongly suggested a predominant role of A_{2A} receptors [113]. A_{2A}KO did not show slow-wave sleep rebound after sleep deprivation while A₁KO did. Furthermore, caffeine induced wakefulness in wild-type and A₁KO, but not in A_{2A}KO mice (reviewed in ref. [114]). However, recent studies using conditional CNS A₁KO mice question the validity of the apparently obvious interpretations of experiments with adenosine receptor KO mice [114], underscoring the limitations of the global absence of a gene during development. Thus, conditional CNS A₁KO mice showed a significant attenuation of the slow-wave EEG activity rebound response to restricted sleep and they also fail to maintain cognitive performance in a working memory task [114]. Nevertheless, recent experiments that measure c-Fos immunoreactivity to measure neuronal activation showed that behaviorally relevant doses of caffeine induce a remarkably restricted pattern of c-Fos expression in the ascending arousal systems. Those studies support a predominant activating effect of caffeine of the orexinergic, histaminergic and noradrenergic systems [115]. Surprisingly, caffeine induced very little or no effect on the cholinergic neurons of the basal forebrain or mesopontine tegmentum [115], which might call into question the generally accepted validity of c-Fos as a universal marker of neuronal activation. In summary, when considering all the results obtained from different studies, there seems to be evidence for a role of an A₁-receptor mediated modulation of the corticopetal basal forebrain system and a complex A₁ and A_{2A} receptor mediated modulation (direct and indirect, respectively) of the noradrenergic system and the hypothalamic histaminergic and orexinergic systems in the arousing effects of caffeine.

CONCLUSION

The study of the psychostimulant effects of caffeine implies the analysis of the multiple effects of endogenous adenosine on multiple ascending neurotransmitter systems. It is obvious that only an integrative view of those effects can allow us to understand the mechanisms of action of the most consumed psychoactive drug in the world. We have explicitly separated motor-activating (and reinforcing) and arousing properties by separating the dopaminergic systems from the ascending arousal systems. But even this separation can be

artifactual. One of the purposes of this review is to show the tight interconnectivity between all the central ascending neurotransmitter systems. As described above, some of the most important inputs to the areas of origin of the ascending arousal systems come from motor brain areas, from the basal ganglia, including the substantia nigra and ventral tegmental area. It was also mentioned that, in fact, the activity of the cells of the corticopetal basal forebrain system is not only associated with motivational stimuli but also with motor activity [87]. That means that some of the arousing effects of caffeine may be secondary to an increased motor activation. In this respect, a recent study in mice with a selective inactivation of striatal A_{2A} receptors showed that in these animals both motor-activating and arousing properties of caffeine were compromised [116]. An additional level of complexity arises when looking at a more molecular level, at the level of local modules, where we discover that the main targets of caffeine (A₁ and A_{2A} receptors) are localized in different parts of different neuronal and glial elements where they form different receptor heteromers. We seem to be quite advanced in our understanding of the modulatory role of adenosine in the striatal local modules (striatal spine module), but this is just one of the terminal fields of the ascending dopaminergic systems. To reach a complete understanding of the mechanisms behind the psychostimulant effects of caffeine we still need to find out about the role of adenosine, adenosine receptors, and adenosine receptor heteromers in local modules from all the other areas of origin and projection of the ascending dopaminergic and arousal systems. Furthermore, we are only starting to understand the mechanisms underlying some temporal properties of the psychostimulant effects of caffeine, such as sensitization and tolerance [22,50,117–119]. In addition to pharmacokinetic mechanisms [117], we need to establish which pharmacodynamic changes, such as modifications in the expression and function of adenosine and dopamine receptors and receptor heteromers [22,50,119], are involved. Some of these temporal effects, particularly sensitization, which is known to be a context-dependent process for classical psychostimulant [120], might be related to the ability of adenosine to modulate long-lasting, activity-dependent changes in synaptic efficacy at excitatory synapses, particularly in the striatum [20]. Nevertheless, in the meantime, caffeine shows to be a very good pharmacological tool to investigate the functions of the central ascending neurotransmitter systems.

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