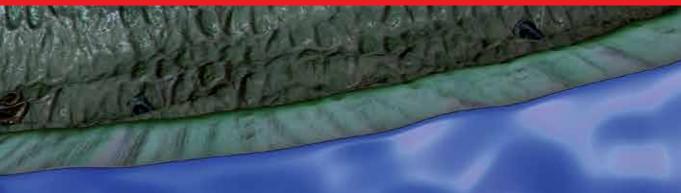


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Biogenic Amines in Neurotransmission and Human Disease

Edited by Ahmet Uçar





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Meet the editor



Prior to working at a major university of health sciences, Associate Prof. Ahmet Uçar received his degrees in pediatrics and then in pediatric endocrinology, achieving high honors at the national exams. He has been actively working in the field of pediatric endocrinology and diabetes, and he has contributed significantly to the definition of the characteristics of pubertal variants in children. He successfully completed a thesis on Turner syndrome,

which has dramatically increased his interest in and dedication to Turner syndrome ever since. He is an active member of the Turner Syndrome Study Group in Turkey. He is also a member of the Endocrine Society and the European Society of Paediatric Endocrinology.

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Preface

Remarkable advances have been achieved in the elucidation of the mechanisms of involvement of biogenic amines in neurotransmission and human disease. In this book, we briefly review neurotransmission, novel therapeutic options in neurodegeneration, thrombotic tendencies in catecholamine excess states, and the pharmacological potential of plant herbs in dealing with catecholamine associated stress. In the introductory chapter, I summarize the physiology of neurotransmission and describe how biogenic amine excess is associated with human disease, with the example of pheochromocytomas and paragangliomas. In the second chapter, Dr Rodriguez-Moreno describes the latest knowledge on kainate receptors and how they modulate glutamate release in the cerebellum, also presenting some insight on how aberrations in the described pathways relate to human disease. In the third chapter, Prof. Martin-Aragon describes the recent advances in the therapeutic management of neurodegenerative diseases, also giving some insight into homeostatic plasticity, which refers to the capacity of neurons to regulate their own excitability relative to network activity, a compensatory adjustment that occurs over the timescale of days. Therapies to modulate neuronal plasticity to halt neurodegeneration and their related clinical findings have shown much progress in the last decade, with enhanced understanding of neurobiology and molecular genetics that now allow genome editing that has the potential to halt the progress of neurodegeneration. In the fourth chapter, Prof. Nobuyuki presents information on the potential effects of plant herbs in alleviating the stress induced by catecholamine signaling. In the last chapter, the mechanisms regarding thrombotic tendencies in catecholamine excess are covered by Dr Rajan. The last two chapters succinctly describe how catecholamine excess states are associated with derangements in multiple systems in the human body.

I hope this book will serve as a point of reference and source of inspiration for researchers and clinicians interested in addressing issues regarding biogenic amine excess- or deficiency- related diseases in humans.

Ahmet Uçar, MD Associate Professor, University of Health Sciences, Şişli Hamidiye Etfal Education and Research Hospital, Istanbul, Turkey

Chapter 1

Introductory Chapter: Biogenic Amines in Neurotransmission and Human Disease from the Endocrinologist's Perspective

Ahmet Uçar

1. A synopsis of the normal physiology of neurotransmission in man

There are about 10¹¹ neurons and 10¹⁴ synaptic connections in the human brain. The neural circuitry is continuously sculpted in response to experience, modified as we learn and store memories, and irreversibly altered by the gradual loss of neurons and connections as we age [1].

Neuronal signals are transmitted from cell to cell at synapses. When an action potential arrives at the presynaptic site, the depolarization of the membrane opens voltage-gated calcium channels that are clustered in the presynaptic membrane. Calcium influx triggers the release of neurotransmitters which are stored in membrane-enclosed synaptic vesicles and released by exocytosis. The neurotransmitter provokes an electrical change in the postsynaptic cell by binding to and opening transmitter-gated ion channels. After the neurotransmitter is secreted into the synaptic cleft, it is rapidly removed: it is either destroyed by specific enzymes in the synaptic cleft or taken up by the presynaptic nerve terminal or by surrounding glial cells. Reuptake is mediated by a variety of Na⁺-dependent neurotransmitter symporters. The cycling of neurotransmitters allows cells to keep up with the high rates of release [1–3].

The chemical or electrical synapses can be excitatory or inhibitory. Excitatory neurotransmitters open cation channels, causing an influx of Na⁺ and also Ca⁺ in many cases, which reduces the threshold to fire an action potential. Inhibitory neurotransmitters open Cl⁻ or K⁺ channels, thereby making it difficult to depolarize the cell membrane. Depending on the secretion milieu, the type of the receptors they bind to, and the ionic conditions that they encounter, transmitters may be either inhibitory or excitatory. For example, acetylcholine may have inhibitory or excitatory effects depending on the type of the receptor it binds to. Usually glutamate and serotonin are excitatory, whereas γ -aminobutyric acid and glycine are inhibitory [1–4]. All neurotransmitter receptors fall into one of these classes based on their signaling mechanisms:

1. Ionotropic receptors—ion channels present at fast chemical synapses. Example: acetylcholine receptor of skeletal muscle cells is a transmitter-gated ion channel which is opened transiently by acetylcholine released from the nerve terminal at a neuromuscular junction. 2. Metabotropic receptors—also called G protein-coupled receptors. Signaling via is these receptors is somewhat slower, more complex, and longer lasting in its effects. For example: hormones such as parathyroid hormone operate via G protein-coupled receptors [1].

Biogenic amines such as dopamine, norepinephrine, serotonin, and histamine manifest their effects either directly or indirectly via their specified receptors [1].

2. Biogenic amines in human disease

Any dysfunction that occurs in the aforementioned physiological setups in the organism may be associated with a disease state, mostly with neurological consequences, and therapy is aimed at inducing the recovery of the dysfunctional pathway via drugs or, most recently, the gene therapy, which is elaborating groundbreaking results [5].

Disorders of neurotransmission in the human are either due to the lack of synthesis of the neurotransmitter, disordered synapse due to external insults such as trauma, autoimmunity, or impaired or enhanced receptor-ligand interaction at the postsynaptic area. The list of the neurogenerative diseases, the inborn errors of metabolism associated with dysfunctional neurotransmission, is not exhaustive and is discussed elsewhere [6].

From the endocrinologist's perspective, excess of biogenic amines includes many rare endocrine neoplasias such as pheochromocytomas (PHEO) and paragangliomas (PGL). The carcinoid syndrome, a constellation of clinical symptoms of biogenic amine excess particularly of serotonin, is much rarer in clinical practice, and it is usually observed in the context of clinically manifest neuroendocrine tumors, such as midgut carcinoids. In the pediatric patient, these syndromes of *amine excess* are usually associated with hereditary syndromes of endocrine neoplasia, such as von Hippel-Lindau disease and multiple endocrine neoplasia (MEN) type 2 syndrome, whereas sporadic cases are more common in the adult [7]. The diagnosis of *amine excess* states is challenging since single measurement of the culprit molecules via standard laboratory tests is often inconclusive and full of caveats that mandate repeat measurements of the suspected amines and/or their metabolites. Below is a brief description of catecholamine excess related tumors as an example to how *biogenic amine excess* may relate to human disease.

Pheochromocytomas and paragangliomas (PGL) are uncommon neuroendocrine tumors of neural crest origin, with the former being of adrenal medulla origin and the latter originating from sympathetic and parasympathetic system. All functional PHEO/PGL produce and metabolize catecholamines and contain chromaffin tissue, which refers to the brownish black color due to oxidation of catecholamines after staining with chromium salts. Pheochromocytomas comprise the majority of the tumors in the pediatric patient [7].

Dopamine, norepinephrine, and epinephrine are biogenic amines collectively referred to as catecholamines. These neurotransmitters and hormones are involved in regulation of numerous physiological processes and development of neuropsychiatric and cardiovascular diseases. They are synthesized from tyrosine, which is converted to 3,4-dihydroxyphenylalanine (DOPA) by the enzyme tyrosine hydroxylase, the rate-limiting enzyme in catecholamine synthesis. Decarboxylation and hydroxylation of dopa yield dopamine and norepinephrine, respectively. Norepinephrine is then converted to epinephrine via the enzyme phenylethanolamine N-methyltransferase.

The complex actions of norepinephrine and epinephrine are mediated by the G protein-coupled alpha- and beta-adrenergic receptors, whereas dopamine binds to a different class of metabotropic receptors.

Introductory Chapter: Biogenic Amines in Neurotransmission and Human Disease... DOI: http://dx.doi.org/10.5772/intechopen.89244

The clinical presentation PHEO/PGL is variable. Owing to the neuroendocrine origin, they might co-secrete other hormones that result ectopic hormone excess, such as gigantism due to growth hormone-releasing hormone, hypercalcemia due to parathyroid hormone-related peptide, or secretory diarrhea due to vasoactive intestinal peptide. The clinical symptoms and signs of functional PHEO/PGL also depend on differences in catecholamine secretion and release as individual patient sensitivities to catecholamines. Signs of catecholamine excess include hypertension, headaches, palpitations, diaphoresis (less common in children), orthostatic hypotension, pallor, tremor, and anxiety. Depending on the location of the tumor, nonspecific signs and symptoms include blurred vision, abdominal pain, diarrhea, or behavioral problems/decline in school performance [7].

The diagnosis of excess catecholamine secretion due to PHEO/PGL has been facilitated by the development of assays sensitive to diagnose these entities; measurements of fractionated metanephrine and normetanephrine in plasma and urine are considered the primary diagnostic tests in the initial evaluation of suspected PHEO/PGL. An elevation of metanephrines greater than fourfold above the reference range is considered indicative of a catecholamine secreting tumor [8]. It should be noted that some PGL can be nonfunctional and may be diagnosed incidentally or due to clinical signs and symptoms owing to their anatomical position, as is the case in some paragangliomas of the head neck which arise from parasympathetic ganglia. Catecholamine-secreting tumors may be noradrenergic as in tumors associated with von Hippel-Lindau disease and familial PGL or adrenergic as seen in tumors that arise sporadically or in the context of MEN type 2 and neurofibromatosis type 1. Dopamine-secreting tumors are very rare and typically extra-adrenal succinate dehydrogenase-mediated paragangliomas. Measurement of methoxytyramine may be of help in this context, but the test is not widely available [9]. Chromogranin A is also another effective tumor marker that may correlate with tumor size and malignant potential.

Radiographic studies and in selected cases, molecular genetic testing should be planned once the diagnosis of catecholamine secreting tumor is highly considered. Further information on management of PHEO/PGL is discussed elsewhere [7].

The expanding knowledge on synthesis and function of biogenic amines may pave the path for enhanced medical treatment of disease states of owing to *biogenic amine excess*. Recent studies have also shown that biogenic amines have additional important roles as signaling molecules mediating the function of the "microbiotabrain-gut" axis [10]. The potential role of biogenic amines in this axis is an exciting new area of medicine that awaits for further studies.

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Chapter 2

Kainate Receptors Modulating Glutamate Release in the Cerebellum

Pilar Losada-Ruiz, Rafael Falcón-Moya and Antonio Rodríguez-Moreno

Abstract

Glutamate receptors of the kainate type (Kainate receptors, KARs), are mediators of ionotropic postsynaptic synaptic transmission, as well as presynaptic modulators of neurotransmitter release where they show both ionotropic and metabotropic actions regulating glutamate and γ -aminobutiric acid (GABA) release. The mechanisms underlying these modulatory roles are starting to be understood at some brain regions. Here we review the KARs roles and mechanisms involved in the modulation of glutamate release in the cerebellum at parallel fibers (PF)-Purkinje Cells (PuC) synapses. KARs activation mediate a biphasic effect on glutamate release at this synapse, with low kainate (KA) concentrations mediating a facilitation of glutamate release and higher KA concentrations mediating a depression of glutamate release. KA-mediated facilitation is prevented by antagonizing KARs, by inhibition of PKA or stimulation of adenylyl cyclase (AC), by blocking Ca²⁺ permeant KARs, by depleting intracellular Ca²⁺ stores and by blocking calmodulin. Thus, at cerebellar parallel fiber-Purkinje cell synapses, presynaptic KARs mediate glutamate release facilitation through Ca²⁺-calmodulin dependent activation of adenylyl cyclase/cAMP/protein kinase A signaling. KAR-mediated depression of glutamate release involves the AC/cAMP/PKA pathway as for facilitation but not Ca²⁺-calmodulin, being in this case AC activated by a Gi/o protein to mediate a depression of glutamate release.

Keywords: cerebellum, KARs, glutamate release, presynaptic, PKA, adenylate cyclase, Ca²⁺ calmodulin

1. Introduction

Glutamate is the most abundant excitatory neurotransmitter in the central nervous system (CNS) of mammals. Glutamate mediates its actions by activating glutamate receptors. These receptors participate in normal synaptic transmission at different synapses, in plasticity processes as long-term potentiation (LTP) and long-term depression (LTD) that are considered the cellular and molecular correlation of memory and learning processes and in synaptogenesis and neuronal maturation and, additionally, failure in the functioning of this system can be the origin of some types of epilepsy and may contribute to the development of CNS disorders such as

Alzheimer's disease, Huntington's Korea, amyotrophic lateral sclerosis, Parkinson's disease, hypoglycemia, or cerebral ischemia [1–3].

Glutamate receptors are classically divided into two large families: ionotropic and metabotropic. Ionotropic glutamate receptors (iGluRs) participate in rapid neurotransmission in the nervous system; these ionotropic receptors are classified into three types depending on the agonist that activates them with higher affinity: N-methyl-D-aspartic acid (NMDA) receptors (NMDARs), α -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA) receptors (AMPARs), and kainate receptors (KARs). These receptors form a channel with different selectivity depending on their subunit composition, all of them being permeable to Na⁺ and K⁺ and, additionally, NMDARs are permeable to Ca²⁺ been some AMPARs and KARs also permeable to Ca²⁺ depending on subunit composition. They are integral membrane proteins, formed by four subunits (tetramers), being homomers or heteromers [1, 2].

Metabotropic glutamate receptors (mGluRs), which participate also in neurotransmission, are coupled to G proteins and are divided into eight types (mGluR 1–8) and three groups of receptors: group I mGluRs includes mGluR1 and mGluR5 receptors. These receptors are positively coupled to phospholipase C (PLC), which facilitates the conversion of inositol diphosphate (PIP2) to diacylglycerol (DAG) and inositol triphosphate (IP3). DAG activates protein kinase C (PKC) that phosphorylates different substrates and IP3 causes numerous intracellular effects, including the facilitation of Ca²⁺ release from intracellular stores. Group II mGluRs includes mGluR2 and mGluR3 receptors. These receptors are negatively coupled to adenylate cyclase-mediated AMPc formation, and group III mGluRs includes mGluR4, mGluR6, mGluR7, and mGluR8 receptors. These receptors are negatively coupled to the formation of AMPc mediated by adenylate cyclase [4].

1.1 Kainate receptors

Kainate (KA) is a potent neurotoxin derived from the alga *Digenea simplex*. The word "Kainic" is derived from the Japanese "Kaininso" ("Makuri"), which means "the ghost of the sea," and it is an agonist for both KARs and AMPARs (in the same way that the AMPARs agonist AMPA may activate KARs). Kainate is classically known for its potent epileptogenic actions [5, 6].

KA (and other agonists) activates KARs that are tetramers that resulted from different combinations of five subunits called GluK1, GluK2, GluK3, GluK4, and GluK5 (formerly known as GluR5, GluR6, GluR7 and KA1, and KA2, respectively). Of these subunits, GluK1 and GluK3 may form homomeric or heteromeric functional receptors, while GluK4 and GluK5 may only participate in functional receptors when associated with any of the GluK1, GluK2, or GluK3 subunits, but they do not combine with subunits of AMPARs [1, 7, 8].

KARs have been described in different invertebrates such as nematodes and flies [9, 10] and in different species of vertebrates such as amphibia, fish, and birds [11–13] in addition to mammals. In mammals, KARs have been observed virtually throughout the entire nervous system, although their subcellular location has not been yet fully determined. KARs are widely distributed throughout the CNS and found in the main cells and interneurons of the hippocampus, lateral amygdala, dorsal root ganglia, bipolar cells of the retina, cerebral cortex, and the cerebellum [14, 15].

The lack of knowledge about these receptors compared to other glutamatergic receptors (AMPARs or NMDARs) has been due to the lack of good agonist and

antagonist for receptors with particular subunit compositions and to the absence of specific antibodies for the different subunits of KARs, being therefore a significant limitation when exploring the distribution of these receptors. However, by using in situ hybridization techniques, it has been observed that the cells that present a significant expression of the kainate-type subunits GluK1, GluK2, GluK3, and GluK5 are distributed throughout the CNS, including nucleus striatum, hippocampus, cortex, and cerebellum [16]. Likewise, there is a high expression of the GluK4 subunit in the CA3 region of the hippocampus, as well as in the granular neurons of the dentate gyrus. The messenger of the GluK5 subunit, on the other hand, appears more abundantly and more extensively than that of the GluK4 subunit or those of the other subunits [15]. Because the in situ hybridization technique is informative and cannot reveal the subcellular distribution of a specific subunit, and because of pharmacological limitations, there is still much to know about the subcellular location and physiology of these receptors.

Kainate-type glutamate receptors are well established mediators of canonical, ionotropic postsynaptic synaptic transmission and, presynaptically, have a modulatory role in regulating neurotransmitter release. In the latter regard, KARs have been shown to have a noncanonical metabotropic capacity, whereby they affect the control of both glutamate and GABA release, for review see [15, 17–22]. At some excitatory glutamatergic synapses, KARs' activation can actually effect biphasic modulation, where low agonist concentrations facilitate glutamate release, while high concentrations decrease the release of the neurotransmitter, for review see [17, 18]. Mechanistic details of how this is achieved are subject of investigation and, indeed, the subcellular location of KARs responsible for presynaptic modulation remains contentious. Different roles of KARs in plasticity have also been described either in LTP or LTD, see [23] for a review of the role of KARs in plasticity.

As other glutamate receptors, KARs are directly or indirectly involved in different diseases, alterations of the nervous system and neurodegeneration and cell death processes. As previously indicated, KA is a potent neurotoxin that directly induces epilepsy and is used as a temporal lobe epilepsy model [5, 6]. Several lines of research indicate that KA directly activating KARs is involved in excitatory and inhibitory imbalances associated with epilepsy. The use of animal models for epilepsy through the use of KA injections has allowed to reproduce in great detail the symptoms observed in humans. The majority of studies of KARs' involvement in epilepsy have studied acute KA-induced seizures [24–27]. The best demonstrations of a mechanism for KARs' involvement in acute epilepsy come from studies of inhibition of GABA release by the activation of presynaptic KA receptors at interneuron-CA1 hippocampal synapses [24, 28, 29]. In chronic epilepsy, a role of KARs has been demonstrated at hippocampal mossy fibers making aberrant synapses onto granule cells of dentate gyrus expressing high number of KARs [30-32] reviewed in [6, 33]. In humans, genetic studies of members of a family affected by idiopathic juvenile absence epilepsy found elevated levels of Grik1 polymorphisms [34], and in TLE patients, GluK1 subunit containing KARs' increased levels have also been found [35]. In clinical studies, NS1209 (an AMPA/KARs' antagonist) has been found to decrease epileptic symptoms [36].

Different studies of neurotoxicity clearly indicate that KARs might be important targets for neuroprotection in neurons and glial cells. The mechanism by which KARs produce excitotoxicity and neuronal cell death is not well understood mainly because of the limitations in appropriate pharmacological tools. Toxicity of KARs involved in multiple sclerosis has also been found onto oligodendrocytes and myelin related to [37, 38], and damage has also been found at axonal levels, where AMPA/KARs' antagonists prevent it [39]. Interestingly, KARs have also been involved in pain. They are present at dorsal roots activating nociceptors, actually there are clinical trials using KARs' antagonists to try to prevent pain showing some levels of analgesia [36]. Additionally, KARs have been involved in ischemia [40, 41], migraine pain [36], Alzheimer's disease [42], Parkinson's disease [43, 44], Huntington's Chorea [45–47], Schizophrenia [48, 49], depression [50], bipolar disorder [51, 52], mental retardation [53], and autism [54, 55] as reviewed in [56]. In general, antagonists of KARs containing particular subunits might be good targets to ameliorate symptoms or treat different CNS diseases and alterations.

2. KARs in the cerebellum

As indicated above, KARs are expressed in the cerebellar cortex [57–59]. As known, the cerebellum participates in the modulation of movement by modifying the activity patterns of motor neurons. Structurally, the cerebellum is composed of the laminar cerebellar cortex and the deep cerebellar nuclei and has five types of cells: Purkinje, stellate, basket, Golgi, and granule cells. Purkinje cells (PuC) are aligned in front of each other. Their dendritic trees form two 2-dimensional layers through which parallel fibers from the mossy fibers located in the granular layer pass. These parallel fibers (PF) establish excitatory synapses between granular cells and the spines of the PCs dendrites as well as the climbing fibers (CF, originating from the inferior olivary nucleus) with the nearby dendrites and the cellular soma. The parallel fibers pass orthogonally through the dendritic tree of the Purkinje neuron. Up to 200,000 PF form a synapse with a single PuC. Each PuC receives up to 500 synapses of CF, all originated from a single CF. Both basket cells and stellate cells provide an inhibitory (GABAergic) entry to the PuC, with cells in the basket synapse to the initial segment of the PuC axon, and stellate cells to the dendrites [60, 61].

Presynaptic KARs participate in plasticity in the cerebellum where PF synapses onto PuCs mediate a form of LTD that is affected by the paired activation of CFs [62], **Table 1**; of these two types of fibers (PF and CF synapsing onto the same cells (PC), only PF have presynaptic KARs [62], similar to other brain regions as somatosensory and visual cortices in which fibers containing and noncontaining presynaptic ionotropic glutamate receptors synapse onto the same postsynaptic cell and induce LTD [63–69]. The exact role and action mechanism of KARs mediating LTD in the cerebellum are not well known yet and await further investigation.

The proper cerebellum development depends on a precise coordinated sequence of postnatal events, some of which are mediated by glutamate receptors. For example, NMDA receptors have been implicated in the migration of granular cells [70] and in the synaptic pruning of climbing fibers [71]. Although it has recently been shown that KARs are involved in synaptic transmission, little is known about their role in development. However, the expression of kainate-type glutamate receptor subunits in immature granule cells of the outer germinal layer of the developing cerebellum suggests that KARs may also have a role in neuronal maturation. Throughout the maturation process of the cerebellum, the quantity, composition, and function of KARs vary. Initially, cerebellar granular cells have a minimal amount of AMPARs in the postnatal period compared to KARs, which are predominant in immature granule cells. Different studies have shown that KARs composed of subunits GluK1, GluK2, and GluK5 predominate, and over the period of development, an increase in the number of KARs is observed and once the adult stage is reached, the number of KARs containing GluK1 subunits suffers a reduction in their expression in the granular layer, while the GluK2 and GluK5 remain constant, in contrast to AMPARs that increase their number, constituting a very notable majority compared to KARs.

KARs' activation	High concentrations of kainate	Depression of glutamatergic synaptic transmission	Delaney and Jahr [86]
	Low concentrations of kainate	Facilitation of glutamatergic synaptic synaptic transmission	Falcón-Moya et al. [80]
	Ionic imbalance	Calcification	Korf and Postema [78]
	Increase in Ca ²⁺	Neurodegeneration	
	Nodular cerebellum lesion Putrescine increase Histological damage	Ataxia	Maiti et al. [72] de Vera et al. [73] Yamaguchi et al. [74] Andoh et al. [75]
Parallel fibers paired with postsynaptic depolarization	Presynaptic KARs' activation	Long-term depression	Crépel [62]
Increase of GluR6 and GluK2 receptors	Reduction of GABAergic activity	Schizophrenia	Harrison et al. [76] Bullock et al. [77]

Table 1.

KARs' actions in the cerebellum.

All of these findings suggest that KARs have an important role in the development process of the cerebellum. Some indications suggest that GluK1-containing KARs participate in cerebellar development in the beginning of the differentiation of granular cells.

Additionally, KARs have been involved in some brain alterations in the cerebellum and a direct relationship exists between KA injection and cerebellar ataxia. Thus, the cerebellum is an important target to study functions of KARs and its possible role causing ataxia [72–75]. Furthermore, in patients with schizophrenia, an increase in KARs containing GluR6 and K2 subunits is observed, which would mediate a reduction in GABAergic transmission [76, 77]. In neurodegeneration, KARs may have a role in calcification of the brain tissue as it has been found that local application of KA in some areas of the cerebellum produces changes in different ion levels, highly increasing Ca²⁺ levels for more than 8 weeks, which mediate calcification [78] (**Table 1**). KARs have been described as producing increases in intracellular calcium [79, 80] and seems to signal increasing intracellular calcium without putting the cell at risk due to excitotoxicity, due to its low conductance in contrast to AMPARs. Due to the lack of knowledge on the subject, further exploration is necessary to determine the KARs' role in cerebellum development and cerebellar alterations.

2.1 KARs modulating glutamate release in the cerebellum: a biphasic effect

KARs are known to be expressed in the cerebellar cortex in the axons of cerebellar granule cells that form PF and make excitatory synapses with PuC [58]. Messenger RNA transcripts encoding for different KAR subunits and functional expression of KAR subtypes have been reported [81–84]. Biophysical studies with single-channel recording have shown GluK1 activity [85], suggesting these KARs are Ca²⁺ permeable. A biphasic action of KARs, activated by the agonist domoate, has been shown previously at PF-PuC synapse, with low agonist concentrations, facilitating synaptic transmission and higher concentrations depressing synaptic transmission [86] in agreement with what has been found in the hippocampus [87–89], cortex [90], amygdala [91], and the thalamus [92]. EPSC trial-to-trial fluctuation analysis, failure rates, as well as paired-pulse ratios have shown that these facilitatory and depressive actions of KARs in the cerebellum are mediated by presynaptic KARs [80]. However, the precise mechanism of action by which KARs mediate potentiation (and depression) of synaptic transmission at PF-PuC synapses has remained elusive until very recently [80] (**Table 1**).

2.1.1 Action mechanism for KARs-mediated facilitation of glutamate release at cerebellar PF-PuC synapses

We have recently demonstrated that the effect of the KARs' activation in this synapse requires protein kinase A (PKA) activation, since the inhibition of this protein by cAMP-Rp suppresses the effect of KA in glutamate release [80], in agreement with previous studies in hippocampus and cortex [87–89]. This congruence between mechanisms at different synapses has also been seen through the inhibition of PKA using H-89, which eliminates KARs-mediated facilitation of glutamate release. Similarly, the direct activation of AC (adenylyl cyclase) using forskolin caused an elimination of facilitation when KARs were activated by KA (with NMDARs and AMPARs blocked). These data indicate that a signaling mediated by AC/cAMP/PKA supports the facilitation of the modulation of synaptic transmission/glutamate release in these cerebellar synapses (**Figures 1–3**).

As observed in other synapses, Ca^{2+} seems to play a fundamental role in facilitating glutamate release at PF-PuC synapses. By blocking calcium-permeable KARs by the selective inhibitor philanthotoxin, KAR-mediated synaptic facilitation of glutamate release was prevented, indicating that there is a strict requirement for external Ca^{2+} entry through KARs to support the facilitation effect observed on glutamate release, indicating that KARs mediating the facilitation of glutamate release are calcium permeable [80].

Additionally, as has been reported at hippocampal synapses, the depletion of intracellular Ca^{2+} stores by a treatment with thapsigargin (a noncompetitive calcium inhibitor of the sarcoplasmic reticulum ATPase) eliminates the facilitation of glutamate release mediated by KARs' activation. The same result was found when selectively inhibiting Ca^{2+} -induced calcium release by using ryanodine, indicating that the entry of Ca^{2+} via KARs induces a mobilization of Ca^{2+} from the intraterminal Ca^{2+} reserves to mediate the increase in glutamate release observed [80].

Furthermore, it has been observed that the facilitation of glutamate release mediated by the activation of KARs is sensitive to calmodulin inhibitors. Previous studies showed that the increase of cytosolic calcium levels activates Ca²⁺ dependent on AC present in the terminals of parallel fibers. Through treatment with the calmodulin inhibitors, W-7 and calmidazolium, it has been recently shown [80] that the inhibition of calcium-calmodulin function prevents KAR-mediated presynaptic facilitation of glutamate release in cerebellar slices, supporting the hypothesis that after KAR activation and cytosolic elevation of Ca²⁺, a calmodulin-dependent calcium coupling activates AC, which subsequently activates the AC/cAMP/PKA pathway, thus promoting synaptic facilitation through an increase in neurotransmitter release at PF-PuC synapses [80].

2.1.2 Action mechanism for KARs-mediated depression of glutamate release at cerebellar PF-PuC synapses

Recently, in the same study discussed in the previous section [80], a transient synaptic depression of glutamate release with high concentrations of KA $(3 \mu M)$

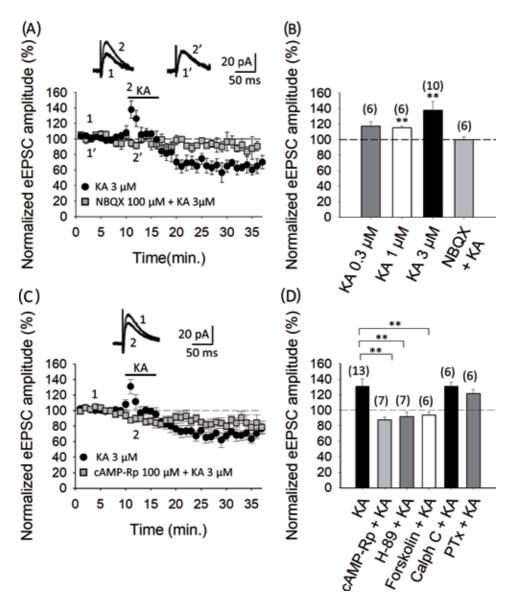


Figure 1.

KÅR-mediated facilitation of glutamate release involving activation of adenylyl cyclase (AC) and downstream protein kinase A (PKA) at PF-PuC synapses of the cerebellum. (A) Time course of KA ($_3 \mu$ M) effect on eEPSCs amplitude in the absence (circles) and presence of NBQX (squares). Insets show traces before and after 4 min of KA perfusion in the absence (1, 2) and in the presence of 10 μ M NBQX (1', 2'). (B) Quantification of modulation observed in (A) and dose-response curve. (C) Time course of the effect of KA on eEPSC amplitude in cAMP-Rp-treated slices. (D) Inhibition of PKA by cAMP-Rp (100 μ M) or H-89 (2 μ M), and activation of AC by forskolin (30 μ M) prevented the facilitatory action of KA. Inhibition of PKC with calphostin C (1 μ M) has no effect on the KA enhancement of the eEPSC amplitude. The facilitatory effect of KA is not affected in slices treated with pertussis toxin. Modified from [64].

was observed as reported for other different brain areas including thalamus, cortex, hippocampus, and amygdala [89–92]. This depression of glutamate release was prevented in the presence of cAMP-RP (which inhibits the activation of PKA), but was not affected by any other experimental modification discussed above with respect to the facilitation of glutamate release. This fact may indicate that the synaptic depression is probably related to an AC/cAMP/PKA signaling pathway (as for facilitation of glutamate release), but without the coupling of Ca²⁺ to the AC. Therefore,

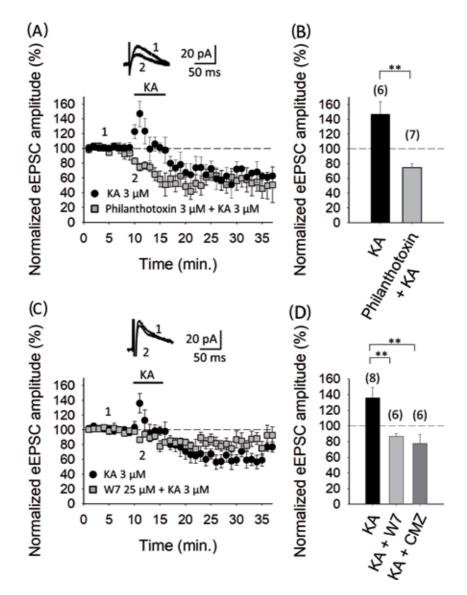


Figure 2.

Facilitation of glutamate release mediated by KAR activation requires an increase of Ca^{2+} in the cytosol and Ca^{2+} calmodulin at cerebellar PF-PuC synapses. (A) Time course of KA (3 μ M) effect on eEPSCs amplitude in control condition (circles) and in slices treated with philanthotoxin (squares). (B) Quantification of modulation observed in (A). (C) Time course of KA (3 μ M) effect on eEPSCs amplitude in control condition (circles) and in the slices treated with 25 μ M W-7 (squares). (D) Quantification of modulation observed in (A) and in the presence of 1 μ M CMZ. Modified from [64].

KA receptors have alternative mechanisms for facilitating and depressing glutamate release at PF-Pu synapses (**Figures 1–3**). In previous studies, investigating mossy fiber-CA3 hippocampal synapses [88, 89, 93, 94], as well as the amygdala [37] and cortex [36], a similar mechanism has been observed additionally involving the activation of a G-protein for the depressive effect that may well be also the case for these cerebellar synapses.

Although the presynaptic function of KARs facilitating glutamate release implies an increase in AC/cAMP/PKA signaling induced by the calcium calmodulin complex, KARs appear to be negatively associated with this pathway to carry out synaptic transmission of depression. Previous studies at hippocampal MF-CA3

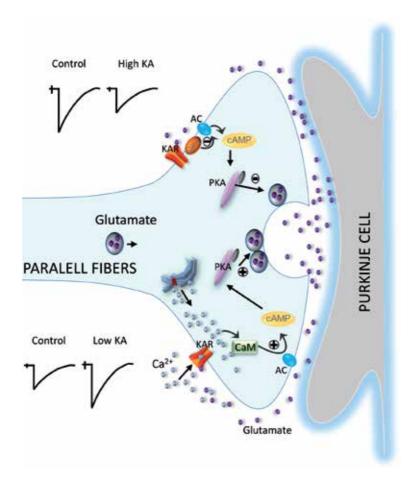


Figure 3.

KAR-mediated modulation of glutamate release in the cerebellum. Actions of KARs depressing or facilitating glutamate release at the PF-PuC synapse. KAR activation by high concentrations of KA (>3 μ M) depresses glutamate release at PF-PuC synapses, an effect that involves G_{i/o} protein and the adenylate cyclase/cAMP/ protein kinase A (AC/cAMP/PKA) pathway. KAR activation by low concentrations of kainate (<0.3 μ M) only facilitates glutamate release following activation of a Ca²⁺-calmodulin/AC/cAMP/PKA pathway.

synapses and thalamocortical synapses, as well as at PF-PuC synapses, have reported that the depression of glutamate release mediated by presynaptic KARs occurs through a negative coupling to the AC/cAMP/PKA pathway, being actually evoked by the action of a PTx-sensitive protein G [80, 88, 92]. Despite the hypotheses discussed above, it is also possible that these observed mechanisms reflect the presence of two different types of KARs, a clear objective being to clarify this hypothesis in future studies.

3. Conclusions

Regarding the role and mechanisms of KARs in the modulation of glutamate release in the cerebellum, new and recent data indicate that the KARs effecting facilitation of glutamate release and synaptic transmission show a mandatory dependence on adenylyl cyclase (AC) and cAMP-mediated protein kinase A (PKA) activity. Furthermore, the KAR-mediated facilitation of transmission is contingent on both external Ca²⁺ permeation into the cytosol through KAR and repletion of

intracellular Ca²⁺ stores. Finally, a major sensitivity of facilitation to calmodulin inhibition suggests that KARs are coupled through a Ca²⁺-calmodulin/AC/cAMP/ PKA pathway at PF-PuC synapses in the cerebellum. KARs seem to use the inhibition of the AC/cAMP/PKA pathway to mediate a depression of glutamate release at the same synapses, but the activation of the AC does not involve calcium calmodulin and seems to be directly activated by a PTX-sensitive G protein.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Chapter 3

Homeostatic Plasticity and Therapeutic Approaches in Neurodegeneration

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Abstract

The synapses transmit signals between neurons in an ever-changing fashion. Changes of synaptic transmission arise from numerous mechanisms known as synaptic plasticity. The importance and complexity of the synapse has fueled research into the molecular mechanisms underlying synaptogenesis, synaptic transmission, and plasticity. Particularly, homeostatic plasticity refers to the local changes in synaptic activation to generate local synaptic adaptations and network-wide changes in activity to generate adjustments between excitation and inhibition. This review chapter will focus on synaptic phenomena and mechanisms that are likely to contribute to network homeostasis. In addition, it will be discussed a putative modulation of the signaling mechanisms serving a homeostatic function as a viable therapeutic approach for disease modification in neurological and neurodegenerative disorders. To sum up, the main role of the following players in homeostatic plasticity will be analyzed, based on what a growing body of evidence has suggested recently: BDNF-mediated TrkB system activation; adenosine modulation system; nitric oxide/soluble GC/cGMP signaling; astrocyte involvement—astroglial CB1 receptors; the microtubule-associated neuronal protein Tau; the signaling pathway of the Wnt protein family; extracellular vesicles in the intercellular communication; and estrogen involvement in non-reproductive functions.

Keywords: adenosine, astrocytes, BDNF, estrogens, extracellular vesicles, neurodegeneration, nitric oxide, synaptic plasticity, Tau, TrkB system, Wnt proteins

1. Introduction

The neurotransmission may be defined as the set of biochemical and physiochemical signals which establishes neuronal communication. Changes of synaptic transmission arise from numerous mechanisms known as synaptic plasticity. Synaptic plasticity is crucial for regulating synaptic transmission or electrical signal transduction to neuronal networks, for sharing essential information among neurons and for maintaining homeostasis in the body.

Synaptic plasticity in the mature nervous system includes structural and morphological modifications constituting the cellular response to the changes in neuronal activity that are thought to be responsible for learning and memory [1].

The individual synaptic connections are constantly removed or recreated depending on the neuronal environment. This plasticity is highly regulated by signals from other cells of the nervous system, mainly the astrocytes whose function is the maintenance of the neurons in a determined position [2]. This modeling ability is achieved by genetic, molecular, and cellular mechanisms that influence synaptic connections and neuronal circuits.

Various neurotransmitter receptors are functionally associated with protein kinases and other G-proteins that modulate cascades of molecules which in turn maintain essential cellular functions [3]. Modifications of the MAPK- and cAMP-related signaling pathways may affect intracellular Ca²⁺ levels, neurotransmitter receptors, transcription factors, and the cross-link between signaling pathways, among other biological functions which are essential for neuroplasticity [4].

At the structural level, synaptic plasticity entails incorporation or disassociation of α -amino-3-hydroxy-5-methyl-isoxazole-4-propionic acid (AMPA) receptors from the postsynaptic membrane and the growth or constriction of the dendritic spines where most excitatory synapses are placed [5]. At the functional level, synaptic plasticity is considered as the long-term potentiation (LTP) or long-term depression (LTD) of synaptic strength, with modifications in conductance through AMPA receptors in the postsynaptic membrane. During the term of plasticity, N-methyl-D-aspartate (NMDA) receptor activation leads to Ca²⁺ ions passage through the postsynaptic membrane to trigger intracellular signaling cascades. These events set off gene transcription, trafficking of AMPA receptor via action dynamics, cytoskeleton reorganization, and enlargement or removal of dendritic spines. The integrity of the synaptic structure, trafficking of AMPA receptor, and dendritic spine dynamics are all crucial for generating lasting synaptic plasticity modifications [5].

The AMPA receptors and NMDA receptors are the ones which synergize at postsynaptic terminals to facilitate different forms of synaptic plasticity. Constant activation of AMPA receptors by a series of impulses arriving at presynaptic terminals leads to depolarization of the presynaptic membrane, which removes the Mg²⁺ ions that are obstructed at NMDA receptors [6]. Thereon, the simultaneous excitation of pre- and postsynaptic neurons speeds up the gating of NMDA channels and reinforces the synapse. This feature is crucial in NMDA channels for being specifically associated with synaptic plasticity and its high permeability to Ca²⁺ ions. Therefore, the second messenger Ca²⁺ modulates a set of signaling pathways and the responses that collectively lead to synaptic modification [6].

Furthermore, local changes in synaptic activation to generate local synaptic adaptations and network-wide changes in activity result in adjustments between excitation and inhibition. These mechanisms are likely to contribute to network homeostasis. Therefore, this review chapter will focus on synaptic phenomena involved in homeostatic plasticity. In addition, it will be discussed a putative modulation of the signaling mechanisms serving a homeostatic function as a viable therapeutic approach for disease modification in neurological and neurodegenerative disorders.

2. BDNF-mediated TrkB system activation

Synaptic plasticity at the molecular level can be driven by increased expression of plasticity-related genes, such as brain-derived neurotrophic factor (BDNF), calcium/calmodulin kinase II (CaMKII), and cAMP response element binding (CREB) protein, as well as by an augmented expression of both AMPA and NMDA receptors on surface [7].

The escalation of the synapse is a mode of homeostatic plasticity in which a prolonged enhancement in neuronal activity leads to a compensatory decline in excitatory transmission that is often mediated by a drop in levels of the synaptic AMPA receptors. The blockage of sustained activity results in the opposite effect, that is, an enhancement in excitatory transmission that is often mediated by an augmentation in levels of the synaptic AMPA receptors [8]. The neurotrophin BDNF and its signaling partners are the main regulators of synaptic plasticity. BDNF may serve as a real mediator rather than simply a modulator of synaptic plasticity and synaptic communication. As BDNF is synthesized and released in an activity-dependent manner [9], BDNF levels could serve as signals for changes in neuronal activity and thereby mediate synaptic scaling. BDNF and neurotransmitter signaling cascades can work together in close temporal association to induce immediate and guided effects on synaptic plasticity. What is more, specifically interfering with BDNF-related signaling is a crucial strategy for neuronal and functionally restorative treatments for neurological and psychiatric disorders [10].

The main functions of BDNF are mediated by its interaction with the tropomyosin-related kinase B (TrkB) receptor [11] whose intracellular signaling cascade activation can affect synaptic transmission and synaptic contact formation [12]. TrkB receptors are typically localized within vesicles inside the cell and translocate to the plasma membrane through neuronal activity [13]. The early effects of BDNF are a result of the modification (e.g., protein phosphorylation) of components that are already present at the synapse, while the long-term effects originate from the modification of translational activity at the synapse and changes in transcription. Stimulation of high-frequency inducing LTP leads to enhancement of BDNF production [14]. Furthermore, BDNF increases neurotransmitter release and promotes synaptic transmission and LTP [15]. Thus, it is assumable that the effects exerted by BDNF on synaptic plasticity are TrkB mediated.

A recent study has investigated the features of neural network formation and functions in primary hippocampal cultures in the context of chronic BDNF application and TrkB receptor blockage [16]. It has been demonstrated that the blockage of TrkB receptors affected the structures of synapses and even mitochondria, which were not regarded as capable of participating in synaptic modulation for a long period. Nevertheless, it has been shown that TrkB-mediated signaling can affect both the ultrastructural and functional parameters of brain mitochondria, even in normal oxygen and nutrient supply conditions. Chronic TrkB receptor blockade leads to destructive ultrastructural changes in mitochondria, whereas the functional activity of organelles remains intact apparently. Long-term application of BDNF enhances the enzymatic activity of mitochondria, though this modification is not related to changes in ultrastructure of organelles. These findings might demonstrate that TrkB-mediated mitochondrial regulation is associated with functional changes of the enzymatic apparatus of the respiratory chain but not with structural reshaping of organelles. Chronic BDNF application increased the basal oxygen consumption rate via activating respiratory chain complex II [17]. Under oxygen stress conditions, BDNF increases the adaptive potential by impacting the functional parameters of the mitochondrial apparatus. Further research should address whether the TrkB-mediated pathway for influencing mitochondria is generalized (i.e., carried out through nuclear genes) or directed toward an isolated organelle.

A genetically determined high level of BDNF can provide significant adaptive potential to the nervous system, and its neuroprotective effects are induced via the TrkB receptor system [18]. To date, it has been found that BDNF-mediated TrkB system activation accounts for the formation of more complex functionally active neural networks with a great level of efficiency in synaptic transmission. Thus, the TrkB signaling system can be involved in higher cognitive functions.

Recent progress into the etiology of neurological and metabolic diseases has been made with genome-wide association studies. BDNF/TrkB signaling contribution to synaptic development and plasticity, neurite outgrowth, and dendritic spine formation is a possible anatomical correlate of hyperconnectivity in autism-spectrum disorder (ASD) [19] or of hypoconnectivity in monogenic obesity [20].

Growing evidence suggests that BDNF upregulation plays a key role in the increased trophic effects in ASD. Studies on functional image have shown that, in most young ASD patients, cortical regions appear hyperlinked, and cortical thickness and brain size are enhanced [19]. These findings indicate that developing ASD brains may occur in a modified neurotrophic environment as some ASD patients and animal models have shown enhanced levels of BDNF.

Besides that, genome association studies now reveal different candidate obesity genes, most of which are strongly expressed or known to act in the CNS, highlighting the role of the brain in propensity to obesity. In this regard, BDNF is involved in energy metabolism and eating behavior as partial BDNF deficiencies in mouse models cause hyperphagia and obesity. The first case described by disruption of the BDNF gene (11p13) in humans was that of an 8-year-old obese girl, who presented an abnormality in chromosome 11 that altered the BDNF gene in one of the chromosomal breakpoints [21]. This patient with severe hyperphagia and obesity also presented a complex neurobehavioral phenotype, including impaired cognitive function and memory and a distinctive hyperactive behavior.

3. Adenosine modulation system

Adenosine is a ubiquitous molecule that is directly involved in the key processes sustaining cellular viability and adaptability, this is it, the energy charge, redox control, DNA and RNA, and epigenetic control. The role of adenosine in the brain is of great interest since the adenosine receptors are far more abundant in the brain than in any other organ or cell type in mammals [22]. Though there are four plasma membrane metabotropic receptors such as adenosine A1, A2A, A2B, and A3 receptors [23], A1 and A2A receptors are the responsible ones for the effects of adenosine in the brain. A1 receptors (A1R) are the most abundant and widely distributed, while A2A receptors (A2AR) are more abundant in the basal ganglia and in synapses throughout the rest of the brain. Both A1R and A2AR are mostly located in synapses, particularly in glutamatergic synapses, although both receptors are also present in other synapses [24].

Adenosine acts on A1R at a pre-synaptical level decreasing calcium influx and glutamate release, at a post-synaptical level decreasing the activation of ionotropic glutamate receptors and of voltage-sensitive calcium channels as well as hyperpolarizing dendrites through a control of potassium channels. The ability of A1R to control high-frequency-induced synaptic plasticity contrasts with that of A2AR. Thus, A2AR are capable of enhancing the evoked release of glutamate in different brain areas and the function of ionotropic glutamate receptors [25]. Besides, A2AR behave as fine-tuners of other neuromodulation systems, since A2AR activation is a requirement for the synaptic effects of growth factors or neuropeptides. What is more, A2AR activation diminishes the efficiency of presynaptic inhibitory systems. Thus, A2AR commute pre-synaptic modulation from inhibitory to facilitatory. Another significant feature of A2AR is that they have also been found in astrocytes and microglia cells, controlling Na⁺/K⁺-ATPase, the uptake of glutamate, the production of pro-inflammatory cytokines, and the effects potentially contributing to the selective A2AR-mediated control of synaptic plasticity [26].

Different sources of extracellular adenosine activate each of these adenosine receptors in order to control synaptic transmission by A1R and synaptic plasticity by A2AR [27]. The predominant role of adenosine under basal conditions, in excitatory synapses, is an A1R-mediated inhibition of synaptic transmission. Then, an endogenous A1R-mediated inhibitory tonus takes place under most experimental conditions. In contrast, A2AR are only recruited upon higher frequencies of nerve stimulation triggering plastic changes of synaptic efficiency (LTP) [28]. This selectivity is consequence of the adenosine formed by synaptic ecto-nucleotidases upon release of ATP from nerve terminals; high-frequency stimulations are needed to trigger an excessive release of ATP [29] that significantly results in the production of extracellular adenosine near A2AR. These A2AR then enhance the release of glutamate and the activation of NMDA receptors, effectively reinforcing the implementation of LTP [30].

Furthermore, the particular activation of A2AR upon higher frequencies of nerve stimulation favors the solution of the increased feedback inhibitory synaptic mechanisms with increasing frequencies of synaptic recruitment. The higher the stimulation frequency, the greater the extracellular levels of adenosine and of cannabinoids since supramaximal activation of pre-synaptic A1R or of cannabinoid CB1R can block synaptic transmission. The neuromodulator adenosine, by operating high affinity adenosine receptors, restrains transmission via A1 receptor (A1R) or CB1R [31].

In spite of the selective involvement of A2AR that allows simultaneously mitigating pre-synaptic inhibition and strengthening synaptic facilitation at the single synapse level, A1R and A2AR actually collaborate to encode the relevance of information at the level of brain circuits. The implementation of a potentiated transmission in a given synapse will elicit in parallel a process of hetero-synaptic depression involving synaptic strengthening of A1R function in surrounding synapses (with respect to this potentiated synapse). The increased recruitment of a synapse set off the activation of the astrocytic syncytium. Within the domain covered by this syncytium, there will be a greater astrocytic (i.e. non-synaptic) release of ATP [32], which will be degraded by ecto-nucleotidases degrading ATP into adenosine, which is channeled into A1R, further depressing the activity of neighboring synapses (where A2AR are not engaged).

In conclusion, a predominant role of neuronal A2AR in the control of neuronal damage is supported by several studies [33]. Dysfunction and damage of synapses may be triggered by the A2AR that are located synaptically. This synapse impairment is regarded as one of the earliest alterations found in different neurodegenerative disorders ranging from Alzheimer's disease (AD) to depression. Experimentally, it has been shown that blockade of A2AR prevents memory damage in AD models. Therefore, the question arises whether A2AR antagonists might have a therapeutic potential. To give some answer, a triple transgenic AD model $(3 \times \text{-Tg-AD})$ has been used with a defined onset of memory dysfunction occurring at the age of 4 months. After the onset of memory deficits in $3 \times$ -Tg-AD mice, a treatment of 3 weeks with a selective A2AR antagonist has shown to normalize the up-regulation of hippocampal A2AR, restore hippocampal-dependent reference memory, and decrease hippocampal synaptic plasticity and global and glutamatergic synaptic markers as well. These findings may point to a therapeuticlike ability of A2AR antagonists to recover synaptic and memory dysfunction that occur in early AD [34]. It may be assumed a role for A2AR in the early steps of neurodegeneration, whereby the up-regulation of A2AR that occurs at the beginning in synapses upon noxious brain insults [27] would set off an aberrant overenforcement of synaptic plasticity that would interrupt synaptic function and favor the likelihood of excitotoxic destruction of the synapse. A2AR would facilitate synaptic plasticity under physiological conditions, while their up-regulation would

lead to a synaptic toxicity that occurs early during the course of neuropsychiatric illnesses. Moreover, the up-regulation and role of A2AR in the astrocyte and microglial control suggest an involvement of A2AR in brain damage, which might also include the control of blood flow and endothelial permeability by these receptors. This entails that multiple cellular sites of action of A2AR would participate in the ongoing process engaged in neurodegeneration [33].

4. Nitric oxide/soluble GC/cGMP signaling in synaptic plasticity

Nitric oxide (NO) plays roles in maintaining synaptic plasticity and in helping to restore plasticity in the neuronal architecture in the CNS. NO is regarded as a chemical transmitter which has essential functions in the mammalian central as well as peripheral nervous system [35]. NO is the second mediator that can activate NMDA receptors. NMDA receptor activation persistently enhances the activity of neuronal nitric oxide synthase (nNOS) in the neuronal cytoplasm. It then catalyzes the generation of endogenous NO from L-arginine followed by the enhanced release of NO from neurons. Activation of these receptors by glutamate stimulates the calcium influx into cells and the generation of NO by NOS, which rapidly stimulates guanylate cyclase and increases cGMP synthesis [36]. Other glutamate receptors, such as AMPA, can also produce NO; this pathway modulates the release of glutamate and dopamine. Nevertheless, AMPA receptor trafficking, expression, and S-nitrosylation activity are maintained by NO. Specifically, the N-ethylmaleimidesensitive factor (NSF, an ATPase) is high in neurons which binds with GluR2 and the NSF-GluR2 interaction has shown to be important to maintain AMPA-mediated transmission at the synapse [37]. Physiologically, synaptic NSF is S-nitrosylated by NO from neuronal source in the mouse brain. Activation of NMDA receptors increases the NSF-GluR2 interaction, as well as the surface insertion of GluR2. NMDA receptors stimulate NO generation, which enhances NSF S-nitrosylation, stimulates its association with GluR2, and increases the surface expression of GluR2-containing AMPA receptors [38].

Additionally, NO is associated with the storage, uptake, and release of mediators, such as acetylcholine, noradrenaline, GABA, taurine, and a glycine. NO can stimulate its own extrasynaptic receptors, which are located some distance from sites of NO synthesis. In addition, nNOS-containing neurons actively participate in the rostral path of neuroblast migration, which involves new synaptic connections and influences neurogenesis [39]. Astrocyte migration is also regulated by the release of NO under the actions of inducible nitric oxide synthase (iNOS). NO is also recognized as critical for the formation of synapses and the growth of nerve fibers [40].

Deficits in synaptic plasticity are increasingly recognized as causes of memory loss in AD [41]. NO is produced by NOS through NMDAR-mediated calcium input. NO signaling comes into play in neurodegenerative diseases via the generation of reactive nitrogen species and cGMP signaling cascades. NO also exerts neuroprotective effects, as shown in AD mouse models, by reducing cell loss and Tau pathology [42]. In AD models, NO has shown to be altered through various mechanisms. For example, the NMDAR-mediated calcium entry that activates NOS is enhanced by abnormal ryanodine receptor-(RyR-) mediated calcium-induced calcium release. NOS and RyR protein levels are also increased in both AD mouse models and human AD brains. In AD mice, at the presynaptic level, these conditions increasing NO levels take place alongside excessive hippocampal synaptic depression. These deficits occur when homeostasis is placed at risk, such as in the presence of reduced RyR-calcium release. Consequently, the hippocampal network and cognitive function are not normal, though they appear to be so [43].

In a study on $3 \times$ -Tg-AD mice, the presynaptic terminals have shown to be the primary site of NO regulation, with increased evoked and spontaneous vesicle release, as determined by paired-pulse facilitation assays and spontaneous vesiclerelease properties [44]. Moreover, NO modifies the magnitude of vesicular release by transforming spare vesicles into easily releasable vesicles [45]. In addition, NO can increase the opening of RyR channels, possibly by means of S-nitrosylation. The opposite interactions between augmented RyR-calcium signaling and enhanced nNOS expression in AD neurons can sustain an increased NO production or synthesis and also strengthen the presynaptic gain. At the postsynaptic level, the neuroprotective characteristics of NO become evident by inhibiting the excessive NMDAR-induced calcium influx and excitotoxicity via S-nitrosylation of the NR2A subunit of the NMDA receptor. At the same time, apoptosis is decreased through the S-nitrosylation of caspase-3, -8, and -9. The enhanced nNOS activity and NO levels in AD brains might be neuroprotective, as proven by the selectively preserved NOS-positive neurons in AD. Hence, continuous increases in NO exert harmful effects, such as oxidative stress, the shredder loss of synaptic function, and apoptosis [46].

All in all, the NO up-regulation or down-regulation may result in neuropsychiatric conditions, and its improvement may restore synaptic plasticity and neuronal function. Understanding the specific molecular mechanisms maintaining these effects can give some light as to identify ways to treat these neuropsychiatric conditions [47]. Particularly, pathological deficits in NO signaling have been reported in corticostriatal circuits in Huntington's disease (HD). Studies indicate that deficits in cortical and striatal nNOS activity and nitrergic transmission may contribute to the progression of corticostriatal pathway dysfunction observed in HD. In this pathology, medium-sized spiny neurons (MSNs) projecting to the external globus pallidus appear to preferentially degenerate as a consequence of accumulation of the abnormal huntingtin protein. What is interesting is that cortical stimulation increases striatal NOS activity by means of a NMDA and dopamine D1 receptor-dependent mechanism. Thus, studies on nNOS knockout mice have proven that striatal MSNs are significantly less sensitive to cortical drive when compared with wild-type controls, indicating that NO signaling plays a critical role in maintaining corticostriatal transmission [48]. Therefore, deficiencies in nNOS activity and NO signaling reported in HD could be associated with decreased excitatory corticostriatal transmission and motor dysfunction. From this concluding remark, it is likely that HD patients might benefit from pharmacotherapies aimed to favor nitrergic signaling and corticostriatal transmission.

Conversely, the protective effect of NMDA antagonists in conditions of in vivo cerebral ischemia supports that influx of Ca²⁺ through NMDA-receptor channels constitute the main driver of glutamate neurotoxicity, a mechanism being of pathological importance. Studies on the role of NO in NMDA-mediated neurodegeneration have given some light with the use of mice lacking individual NOSs subjected to ischemia. For instance, it has been found that nNOS contributed to the early damage, whereas endothelial NOS (eNOS) was protective, reflecting its relevance in cerebral blood flow and, likely, in inhibiting both leucocyte adhesion to the endothelium and platelet aggregation [49]. Reduced synthesis and availability of eNO may contribute to the development of dementia by favoring the onset and progression of atherosclerosis, vasoconstriction, and impaired cerebral blood flow regulation. Furthermore, the inducible isoform of NOS (iNOS) that starts to be expressed in the following days to an ischemic event appeared to provide additional damage. iNOS is expressed in macrophages and glial cells in response to proinflammatory cytokines or endotoxin. In the brain, widespread expression of iNOS is pathologic and has been observed in neurological diseases, such as multiple

sclerosis, stroke, and AD. Increased NO production and iNOS expression have been observed after experimental and clinical traumatic brain injury (TBI). iNOS knockout mice showed reduction in infarct volume and motor deficits after focal ischemia compared with wild-type mice. In an exploratory phase II study trial, TBI patients treated with the most selective inhibitor of human iNOS reported to date showed significant improvement in clinical outcome [50]. These studies suggested a neuroprotective effect of iNOS inhibition after TBI.

5. Astrocyte involvement—astroglial CB1 receptors

Astrocytes are the most abundant glial cell population in the brain [51]. One of the most intriguing features of astrocytes is the control of synaptic plasticity and memory functions. Astrocytes contribute in shaping different types of synaptic plasticity that challenges the generation of experimental models enabling the distinction between the neuronal contribution and the astroglial contribution in the control of a given function [52].

The finding of a large number of astrocytes and increased expression of the α 7 nAChRs subunit on these cells in hippocampus and temporal cortex of sporadic AD patients have suggested an involvement of the astrocytic α 7 nAChRs in the metabolism of A β and a relationship with the amyloid cascade [53]. It has been proven that A β , at physiological levels, controls synaptic activity and acts as a positive or negative regulator post-synaptically. In fact, A β either activates or inhibits α 7 nAChRs in a dose-dependent manner. In the healthy brain, A β enhances spontaneous astrocyte calcium transients, regulating neuron-glia signaling in a nAChR-dependent manner [54]. On the other hand, A β accumulation increases astrocytic α 7 nAChRs expression and subsequently increases intracellular calcium released from intracellular stores contributing to a number of inflammatory cascades.

The nAChRs from astrocytes can be activated by the transmitters released from presynaptic terminals or neurotransmitters located in the interstitial space [52]. For instance, the nAChRs agonist acetylcholine elevates intracellular calcium in astrocytes, stimulating the release of glutamate gliotransmitter and consequently the NMDA receptor-dependent currents in neurons. It results in an increase of the efficient synaptic transmission and strength. This mechanism explains the Aβinduced hippocampal LTP. Then, A β - α 7 nAChRs interaction increases astrocytic activity to induce glutamate gliotransmission which contributes to synaptic plasticity and cognition [55]. However, pathological concentrations of A β , as occurred in AD, induces LTD as A β oligomers-induced glutamate release from astrocytes causes excitotoxic damage and synaptic loss in neurons. In essence, pathological levels of A β cause α 7 nAChR-induced gliotransmission dysregulation leading to increased excitability within the hippocampal neural network [56]. What is more, endogenous stimulation of α 7 nAChRs on astrocytes causes expression of further AMPA receptors post-synaptically at glutamatergic synapses of neurons. Consequently, the ratio of AMPA-evoked synaptic currents increases.

There is evidence that GABA is abundantly produced and released by activated astrocytes. A non-physiological increase in tonic GABA release from reactive astrocytes in the hippocampus may be directly responsible for the memory impairment in AD. It has been shown that alterations of kynurenic acid, an astrocyte-derived antagonist of the α 7 nAChRs, are associated with the impairment of the cognitive function in AD patients. In fact, kynurenic acid at low concentrations inhibits GABAergic neurotransmission triggered by α 7 nAChRs activation [57]. Therefore, prevention of GABAergic neurotransmission by α 7 nAChRs inhibition through kynurenic acid may protect against AD.

The α 7-specific agonists decrease IL-6 production in cell lines derived from astrocytes. Conversely, internalization of these receptors in astrocytes by binding to α 7specific antibodies could stimulate IL-6 production. The α 7 nAChRs internalization might provoke neuroinflammation within the brain by inducing IL-6 production in astrocytes. Furthermore, evidence shows that treatment with different α 7 nAChRs agonists prevent neuroinflammation in AD-affected brains and suppress elevated levels of astrocyte-related pro-inflammatory cytokines, such as TNF- α [58].

Microglia play different roles in the healthy brain which include neuronal synapse monitoring, phagocytosis of neuronal debris, and cell migration as well as proliferation. Microglia show a typical resting phenotype in the healthy CNS, becoming activated in response to excessive neuronal activity during neurochemical or electrically stimulations. It has been shown that increased activation of astrocytic $\alpha 4\beta 2$ and $\alpha 7$ nAChRs drives the expression of glial cell-derived neurotrophic factor (GDNF) [59]. Astrocyte-derived GDNF binds to GDNF family receptor alpha1 (GFR α 1) and activates intracellular signaling pathways that lead to inhibition of microglial activation and protection of neurons from neurodegeneration [60].

The astroglial type-1-cannabinoid (CB1) receptor signaling recently emerged as the mediator of several forms of synaptic plasticity associated to important cognitive functions [61]. The presence of CB1 and type-2 (CB2) receptors in astrocytes was first described in human and rodent primary cultures. Particularly, CB1 receptors are mainly located at presynaptic terminals of different types of neurons, inhibiting neurotransmitter release. Although the signaling of CB1 receptors in astrocytes has been partially investigated, it has been suggested that they are able to couple to different intracellular signaling pathways in this type of cells, therefore providing high adaptability to endocannabinoid-mediated responses in these cells. The first endocannabinoid identified was arachidonoyl-ethanolamide, known as anandamide [62]. Arachidonoyl-ethanolamide mimics the cannabinoid "tetrad" effects (i.e., hypolocomotion, antinociception, catalepsy, and hypothermia) in rodents. The second discovered endocannabinoid, 2-arachidonoyl-glycerol (2-AG) is considered as the main effector of most of the CB1 receptor-mediated endogenous regulation of synaptic transmission. Endocannabinoids' precursors are located in cell membranes and their transformation into actual endocannabinoids is thought to mainly occur in an activity-dependent manner. Endocannabinoids are likely produced, mobilized, and degraded by astrocytes *in vivo*, suggesting the presence of an astroglial autocrine and/or paracrine endocannabinoid signaling.

Astrocytic processes express transporters GLT-1 and GLAST for glutamate, participating in the rapid removal of glutamate released into the synaptic cleft, which is essential for both the termination of synaptic transmission and maintenance of physiological neuronal excitability. Since increased stimulation of glutamate receptors is highly toxic for neurons, glutamate uptake by astrocytes is crucial for protecting neurons against excitotoxicity. In this regard, it has been observed that pharmacological blockade of CB1 receptors reduces epileptiform discharges of hippocampal slices through a direct control on astrocytic calcium signaling, suggesting that over-activation of astroglial CB1 receptors might promote the maintenance of brain excitotoxicity [61].

Intriguingly, whereas CB1 receptors on glutamatergic neurons have been found to reduce glutamate release and excitoxicity, astroglial CB1 receptors have been shown to promote astroglial-dependent activation of glutamatergic transmission [63]. These potentially opposite effects at neurons or astroglial cells might indicate the complexity in the regulation of excitatory output by the endocannabinoid system.

However, arachidonoyl-ethanolamide has shown to reverse AMPA-induced neurotoxicity and down-regulation of GLT-1 and GLAST (glutamate transporters) mRNA in cultures of mouse astrocyte and in the spinal cord of a mouse model of multiple sclerosis via CB1 receptor [64]. Thus, the role of astroglial CB1 receptors in the regulation of excitatory amino acids in the context of excitotoxic processes might depend on the conditions of the brain circuits involved and the specific activation of the endocannabinoid system at different sites. It might be hypothesized that astroglial CB1 receptors might exert a bidirectional control of excitatory transmission and excitotoxicity [61].

The role of astroglial CB1 receptors in the modulation of excitatory transmission in the hippocampus was first reported by Navarrete and Araque [65]. They found that hippocampal astrocytes express functional CB1 receptors that increase intracellular Ca^{2+} levels in astrocytes following electrical stimulation of adjacent neurons. CB1 receptor-dependent increase of intracellular Ca²⁺ levels in astrocytes requires activation of PLC, suggesting that the interaction of CB1 receptors with the $G\alpha q$ protein subunit may be favored in astrocytes as compared to neurons. Astroglial CB1 receptors activation is necessary for NMDA receptor-mediated neuronal excitability evoked by astrocyte Ca²⁺ elevations, suggesting their importance for astrocyte-neuron excitatory signaling. However, these results do not characterize neither the endogenous NMDA receptor agonist implicated in this process nor the cellular origin and identity of the released eCBs that act at astroglial CB1 receptors in these conditions. Astroglial CB1 receptors mediate heterosynaptic potentiation at excitatory CA3-CA1 synapses in a Group I mGluR-dependent fashion. Importantly, this phenomenon coexists with depolarization-induced suppression of excitation (DSE, short-term inhibition of excitatory transmission) mediated by neuronal CB1 receptors at homosynaptic connections [66].

Finally, astroglial CB1 receptors mediate spike timing-dependent LTD (tLTD) in the neocortex. tLTD is induced by paired stimulations of the pre- and postsynaptic elements, leading to a reduction of presynaptic release of glutamate. CB1 receptordependent tLTD has been also described at cortico-striatal synapses and may imply astroglial CB1 signaling as well. This form of synaptic plasticity has been associated to the neuronal coding of sensory experiences *in vivo*. Interestingly, one of the major consequences of exogenous cannabinoids administration, including THC, is an alteration of perception, but little is known about the mechanisms underlying these effects [67]. Therefore, it is possible that the overstimulation of CB1 receptors impairs sensory functions by a deregulation of tLTD.

6. The microtubule-associated neuronal protein Tau

The neuronal cytoskeleton is the major intracellular structure that determines the morphology of neurons and plays a critical role in the development of the nervous system, neuronal plasticity, and neurodegenerative diseases [68]. In particular, changes in microtubule (MT) dynamics are involved not only in axon formation and axonal sprouting but also in mediating structural and functional changes of dendritic spines, which represent the major site of excitatory postsynaptic input [69].

MT dynamics are regulated by several factors affecting the assembly state of this polymer. MT-associated proteins (MAPs) promote MT nucleation and elongation in a compartment-specific manner and are also subject to regulation by posttranslational modification [69]. Tau is of particular importance since it becomes enriched in the axonal compartment during neuronal development and redistributes to the somato-dendritic compartment in a state of hyperphosphorylation during neurode-generative pathologies such as AD and other tauopathies [70]. Strikingly, acute or chronic knockdown of Tau does not appear to affect MT stability and organization or the overall structure of a neuron to a major extent. This feature raises the

question whether Tau possesses activities beyond its direct role in regulating axonal MT polymerization, which is altered in AD and other tauopathies. In this regard, Tau is an intrinsically disordered protein, which interacts with different partners. Tau stands out for its structural plasticity allowing it to react quickly in response to changes in their environment by posttranslational modifications [71]. Then, Tau may affect various signaling and regulatory processes dependent on its modification, thereby regulating the function and plasticity of neurons during learning, memory, and degenerative processes.

In order to make a functional assessment of Tau role with regard to neuronal activity and network properties, several electrophysiological studies have been performed using Tau knock-out (KO) mice or Tau knockdown experiments. Complete absence of Tau or a decrease in the amount of Tau on neuronal activity has been studied by recordings from individual cortical neurons as well as from intact neuronal circuits in acutely isolated tissue slices. The focus in most studies was placed on the influence of Tau on LTP or LTD generation in the hippocampus both being established experimental paradigms of synaptic plasticity underlying learning and memory formation. Cantero et al. [72] has studied the involvement of Tau in shaping electrophysiological properties of neural oscillations in different neocortical regions and hippocampus during spontaneous exploratory motor behavior of 4 month old mice. They observed a significant slowdown of the hippocampal theta rhythm, which is known to play a crucial role in learning and memory function. Furthermore, they found decreased levels of gamma long-range synchronization in the KO animals, while power and peak frequency in the gamma band oscillations (30-80 Hz) recorded from neocortex and hippocampus remained unchanged. As a result, it was hypothesized that the lack of physiologically phosphorylated Tau during early stages of development may influence the maturation of parvalbumincontaining interneurons affecting the spatiotemporal structure of long-range gamma synchronization between hippocampus and neocortex.

It was recently demonstrated that Tau exerts an important mediatory role in regulating the interaction between protein kinase C binding protein 1 and GluA2, a molecular mechanism fundamental to AMPA receptor internalization [73]. Biochemical and electrophysiological assays have allowed to show that specific phosphorylation at S396 of Tau is related to and required for LTD. In addition, the absence of Tau led to the reversion of deficiencies in spatial memory, pointing out a relevant role of Tau in hippocampal LTD. Thus, Tau seems to be decisively involved in the maintenance of physiological synaptic transmission and synaptic plasticity.

For many years, in terms of location, Tau has been considered to be an exclusively intracellular protein. However, Tau can be released from neurons and it has been found in the cerebrospinal fluid of AD patients. This feature may affect network functions by directly influencing the activity of multiple neurons. Stimulation of neuronal activity can induce Tau release from healthy cortical neurons, suggesting that Tau secretion is a physiological process [74]. In vivo microdialysis has proven that increasing neuronal activity raises the level of extracellular Tau. The elimination rate of Tau from the extracellular space in the brain is slow suggesting that Tau can be present for long times outside of neurons and influence other cells. Extracellular Tau has a reduced phosphorylation state and is enriched in Cterminally truncated forms. Besides, it has been demonstrated that a Tau fragment resulting from the cleavage through the effector caspase-3 exhibits enhanced secretion [75]. It is not yet clear in which form Tau is present in the extracellular space and how it potentially influences other neurons. Extracellular Tau might affect network functions by acting on other cells through receptor-mediated mechanisms. Alternatively, extracellular Tau might be taken up by other neurons and modulate their signaling cascades. Several studies have reported uptake of Tau aggregates and oligomers, which may cause network damage by propagation of protein abnormalities. For instance, it has been suggested a prion-like spreading of a pathogenic misfolded Tau in a deterministic manner to distinct brain regions [76]. However, it is not clear whether or not extracellular soluble Tau is internalized by neurons at physiological conditions. Application of exogenous Tau from cerebral cortex of AD brain evoked a strong LTP inhibition at CA3-CA1 synapses in hippocampal brain slices being elicited by theta burst stimulation. Moreover, a short exposure to oligomers, but not monomers, of extracellular recombinant human Tau causes a concentration-dependent impairment of LTP in the CA3-CA1 pathway and memory formation [77]. These effects were reproduced both by Tau oligomers from AD human specimens, as well as those produced in mice overexpressing the human Tau. *In vitro* conditions has evidenced that oligomerized Tau passes the cell membrane and is internalized by neurons. However, it is unclear whether extracellular Tau acts in these experiments through receptor-mediated mechanisms or after internalization.

With all of it, it appears clear that Tau exerts systemic effects and modulates network functions both in the CNS and the peripheral nervous system. Consequently, changes in the Tau level will produce various side effects. Though most available data are still controversial at length, general trends begin to become apparent from several experimental approaches. The course of motor impairments as a consequence of reduced Tau levels at advanced age might be caused by an impact on the peripheral motor system. Mild cognitive and memory deficits and a disruption of physiological synaptic transmission and synaptic plasticity have been repeatedly reported. Even posttranslational modifications of Tau possessing pathologic effect may serve protective functions for synapses, which would be impaired in circumstances of Tau depletion. Experiments with hibernating animals have rendered evidence for a protective function of Tau phosphorylation, providing a striking model for extreme plasticity, as numerous synaptic contacts are lost during hibernation and regenerated in a brief period of time after animal awakening. In these animals, a physiologically adaptive process associated with synaptic plasticity seems to be supported by a reversible hyperphosphorylation of soluble Tau [78].

Furthermore, a small amount of Tau has been localized and accumulated in the nuclear compartment, in both the soluble and chromatin-bound fractions. A putative role of Tau in both nucleolar organization and DNA damage protection and chromosome stability have been proposed, since wild-type Tau prevents DNA damage under oxidative stress or hyperthermic conditions. Recently, it has been observed that promoting Tau nuclear translocation and accumulation, by Tau overexpression or disengagement from MTs, enhances the expression of the vesicular glutamate transporter VGluT1 [79], a disease-fundamental gene directly involved in glutamatergic transmission. Significantly, the P301L mutation in Tau gene, linked to frontotemporal dementia FTDP-17 [80], disturbs this mechanism leading to a function loss. This fact demonstrates a direct physiological role of Tau on modulating gene expression, particularly the expression of the vesicular glutamate transporter VGluT1, which is known to be strongly upregulated in early phases of tauopathies. Thus, alterations of this mechanism may be at the basis of the onset of neurodegeneration.

Taken together, though most current approaches to target Tau are based on its actions on MTs and are aimed at counteracting the aggregation of its soluble pool, unlinked functions of Tau to its MT-binding properties have been suggested lately. Therefore, in view of the evidence supporting considerable modulatory and potentially protective functions of Tau at physiological conditions, the strategies aimed at modulating the amount and modification of Tau should be carefully taken into consideration when planning Tau-based therapies.

7. The signaling pathway of the Wnt protein family

In humans, Wnts are a family of secreted glycolipoproteins (19 members) that are evolutionarily conserved [81]. Wnts are essential for axon pathfinding, dendritic development, and the formation and function of synapses [82]. The contribution of Wnts to synapse development in different model systems has been supported widely [83].

Several Wnt proteins (Wnt7a, Wnt5a, and Wnt3a) that signal through different receptors promote pre-synaptic differentiation [84]. Besides, Wnts signaling promotes excitatory synapse formation and spine growth [85]. Unlike Wnt7a, Wnt5a promotes inhibitory post-synaptic assembly by increasing GABA_A receptor clustering and enhancing the amplitude of inhibitory postsynaptic currents [86]. Members of the Wnt family might regulate excitatory and inhibitory post-synaptic properties, differentially. Moreover, Wnts act bidirectionally to promote the assembly of both sides of the synapse [84].

Regulation of synaptic transmission by Wnt proteins has been demonstrated by electrophysiological recordings. Members of the Wnt family can modulate neurotransmission pre-synaptically and post-synaptically [84]. For instance, gain of function of Wnt7a has shown to promote transmitter release in cultured neurons. Post-synaptically, Wnt7a enhances synaptic strength by increasing the number of AMPA receptors at the post-synaptic membrane in hippocampal neurons and Wnt5a potentiates post-synaptic NMDAR-mediated currents through RoR2 receptors in hippocampal neurons.

Members of the Wnt family are regulated by neuronal activity. Thus, neuronal activity modulates the mRNA and protein levels of Wnts and their receptors in different model systems. Wnt blockade completely abolishes activity-mediated synapse formation in cultured neurons [87]. Blockade of endogenous Wnt proteins, with secreted frizzled-related proteins (Sfrps), severely impairs LTP [88]. Conversely, addition of Wnt proteins can facilitate LTP [89].

Research is been currently focused on identifying the mechanisms by which Wnts modulate LTP. In this regard, Wnt5a has been shown to regulate NMDARmediated synaptic transmission in acute hippocampal slices [89]. However, it does not affect endogenous synaptic AMPAR localization or dendritic spine size in hippocampal cultured neurons [88]. Then, it is suggested that Wnt5a may contribute to later stages of LTP.

Endogenous Wnt signaling is required for structural and functional plasticity during LTP as it has been demonstrated. A study shows that Wnt7a/b regulate the early stages of NMDAR-dependent LTP [88]. Acute blockade of endogenous Wnts with Sfrps attenuates LTP induced by stimulating acute hippocampal slices at high frequency or LTP induced by glycine application in hippocampal neurons. Changes in dendritic spine structure, augmented synaptic strength, and synaptic localization of AMPA receptors are inhibited in the presence of Sfrps during LTP. Intriguingly, gain of function via single-particle tracking and super-ecliptic pHluorin-tagged AMPA receptors in hippocampal neurons has proven that Wnt7a quickly enhances dendritic spine growth, recruitment of synaptic AMPA receptors, and synaptic strength, similar to the early phases of LTP [88].

A proper balance between canonical and non-canonical Wnt signaling might determine whether synapses are lost or only their function is affected [84]. A mouse model characterized by a Wnt deficiency in the adult brain has demonstrated that Wnt proteins constitute a requirement for synapse stability and synaptic plasticity [90]. Transgenic mice expressing the Wnt antagonist Dickkopf-1 (Dkk1) protein in hippocampus display a loss of excitatory synapses, altered LTP, enhanced LTD, and deficits in long-term memory [91]. Acute blockade of endogenous Wnts by Sfrps does not affect the number of excitatory synapses but inhibits LTP induction. This is not due to acute versus long-term exposure to Wnt antagonists, as Dkk1 also rapidly induces synapse loss in mature neurons [92]. Both antagonists have a different mechanism of action. While Sfrps inhibits Wnt function by means of binding to Wnt proteins, Dkk1 affects a canonical Wnt signaling pathway by blocking the low density lipoprotein receptor-related protein 6 (LRP6), an essential Wnt co-receptor.

Growing evidence suggests that deficient Wnt signaling affects synaptic integrity in the adult brain. In AD, particularly, decreased levels of Wnt signaling could weaken synaptic function resulting in the subsequent degeneration and loss of synapses characteristic of this condition. Expression of the endogenous Wnt antagonist Dkk1 is increased in the brain of AD patients. On other hand, Aß quickly induces Dkk1 expression and blockade of Dkk1 protects synapses from Aß [92]. Besides, a variant of LRP6 has been linked to late onset AD [93] and deletion of LRP6 magnifies pathology in an AD mouse model [94]. These findings along with those obtained from mice expressing Dkk1 [91] demonstrate that deficits in Wnt signaling affect synaptic integrity in the adult brain. In AD, diminished levels of Wnt signaling could mine synaptic function leading to the subsequent degeneration and synapse loss which is characteristic of this condition. Reactivation of Wnt signaling might restore connectivity after substantial synapse degeneration. In fact, the ceasing of Dkk1 expression in transgenic mice that express Dkk1 fully restores the structural and functional plasticity and hippocampal-dependent memory [91]. Together, these studies prove the regenerative capacity of neurons in the adult hippocampus to assemble synapses within functional circuits after degeneration.

8. Extracellular vesicles in the intercellular communication

In the CNS, extracellular vesicles (EVs) have emerged as key players in the intercellular communication that underlies physiological processes such as synaptic plasticity, maintenance of myelination, and neuronal and glial response to brain injury. EVs are secreted membrane-enclosed 'packages' that contain cytosolic proteins, membrane proteins, mRNAs, noncoding RNAs, and even DNA [95].

In order to modulate synaptic plasticity, neuronal EVs can transfer cargo to other neurons. It has been suggested an uptake of neuronal EVs by microglia to enhance microglial pruning of synapses. Moreover, increased astrocytic uptake of glutamate may result from internalization of neuronal EVs into astrocytes. In turn, EVs from astrocytes have been shown to contain cargo-mediating neuroprotection under neuronal stress. Oligodendrocyte-derived EVs increase neuronal firing mediate as well as neuronal stress resilience [96]. Moreover, oligodendrocyte-derived EVs may behave as auto-inhibitors in expansion of myelin. As for microglial EVs, they may modulate neuronal firing, and also propagate inflammation and destabilize synapses in a CNS injury setting. In terms of neurodegenerative diseases, EVs are involved in both the spreading and clearance of neurotoxic protein aggregates [97]. EVs have also been shown to cross the blood-brain barrier particularly under inflammatory conditions, enabling molecular crosstalk between brain cells and the periphery [98].

The coupling of neuronal EV release to synaptic activity could be functionally relevant for plasticity-associated processes. For instance, activity-dependent disposal or transfer of AMPA receptor components, which have been shown to flow throughout EVs, may constitute a mechanism for the adjustment of synaptic strength [99]. Moreover, it has been suggested that neuronal EVs act as transsynaptic carriers of signaling proteins involved in synaptic plasticity. In a series of

in vivo studies at the *Drosophila* neuromuscular junction, it has been shown that the Wnt-family signaling protein Wingless was protected driving toward EVs by the trafficking protein Evi/Wntless for trans-synaptic transport and hampering this trafficking impaired synaptic bouton formation [100]. In the same model system, synaptotagmin 4 has been found to drive on EVs from the presynaptic motor neuron to the postsynaptic muscle, and this directional transport was fundamental for activity-dependent growth of presynaptic structures.

Plasticity-associated local protein synthesis at synapses appears to be affected by the activity-dependent trafficking of specific RNAs into EVs. In one study, depolarization of differentiated neuroblasts has been found to be correlated with a depletion of specific miRNAs from neurites, associated with an enhancement of a specific subset of these miRNAs in EVs. Furthermore, neuronal EVs have been recently shown to package mRNA in association with the activity-regulated cytoskeleton-associated protein (Arc), described as a master regulator of synaptic plasticity [101]. Arc appeared to self-assemble into capsids as if it were a viral group-specific antigen (Gag) protein, encapsulating its own mRNA or other highly abundant mRNAs, and trafficking between cells by means of EVs. The blockage of Arc trafficking from presynaptic terminals to the postsynaptic muscle, at the Drosophila neuromuscular junction, caused aberrations in synapse maturation and activity-dependent plasticity [102]. Therefore, Arc capsid-containing EVs may constitute a retrovirus-like mechanism for trans-synaptic transfer of at least Arc mRNA during processes associated with synaptic plasticity. The biological origin and composition of Arc-containing EVs, as well as the scope of the EV-associated Arc transport in the mammalian brain, have yet to be studied in the future.

It is intriguing that neuronal EVs can also transfer cargo to other cells in the brain, modulating their behavior with a potential impact on synaptic activity. One study found that the uptake of neuronal miR-124a-carrying EVs into astrocytes was correlated with overexpression of excitatory amino acid transporter 2/glutamate transporter-1 (EAAT2/GLT1), which could modulate synaptic activity via an increased uptake of glutamate into perisynaptic astrocytes [103]. Moreover, EVs from differentiated PC12 cells seemed to promote microglial phagocytosis of degenerating neurites by means of increasing the microglial expression of complement component 3 (C3), a factor associated with synaptic pruning. The release of EVs from active neurons could thereby serve to the removal of less functional synapses during developmental or learning-associated remodeling of neuronal connections, although validation of this finding is warranted in at least primary neurons.

With regard to glial EVs, they have been described to provide neurons with support and feedback on synaptic activity. And so, for example, oligodendrocytes contain multivesicular endosomes at sites near the axonal surface, and secrete EVs particularly in response to glutamate. While microglia appeared to degrade oligodendrocyte-derived EVs, neuronal internalization of these EVs led to functional cargo retrieval along with an increase in neuronal firing rate and modified gene expression of several plasticity-related targets, such as VGF nerve growth factor inducible and BDNF [104]. Microglial microvesicles (MVs), for its part, released in response to ATP and increased excitatory neurotransmission through stimulation of neuronal sphingolipid metabolism and a consequent increase in the presynaptic release of neurotransmitters. The functionality of the MVs seemed to depend on a lipid or a surface component since sheared MVs retained the functionality [105]. In another study, microglial EVs have shown to carry the active endocannabinoid N-arachidonoylethanolamine on their surface, which bound to presynaptic type 1 cannabinoid receptors, and thereby decreased the release of the neurotransmitter GABA. Overall, glial EVs seem to influence both excitatory and

inhibitory neurotransmission, providing regulatory feedback particularly on presynaptic activity [95].

As remarkable, EVs have emerged as possible therapeutic agents in inflammation-mediated demyelinating diseases, such as multiple sclerosis (MS). Increasing evidence suggests that EVs display anti-inflammatory properties, reducing the number of activated inflammatory microglial cells, supporting oligodendrocytes and protecting neurons. A very recent study illustrates a putative therapeutically role for EVs. Intravenously administered EVs derived from mesenchymal stem cells (MSCs) from human adipose tissue might mediate recovery in Theiler's murine encephalomyelitis virus (TMEV)-induced demyelinating disease, a progressive model of MS [106]. Intravenous EV administration in SJL/J mice improved motor deficits, lowered brain atrophy, enhanced cell proliferation in the subventricular zone, and decreased inflammatory infiltrates in the spinal cord of animals infected with TMEV. In addition, EV treatment was capable of modulating neuroinflammation, given glial fibrillary acidic protein and Iba-1 staining were reduced in the brain, whereas the expression of myelin was increased. Modifications of morphology in microglial cells from the spinal cord may indicate that EVs also modulate the activation state of microglia. EV administration attenuates motor deficits through immunomodulatory actions, diminishing brain atrophy and promoting remyelination.

So far, there is no evidence regarding the therapeutic potential of EVs during the neurodegenerative phase of MS. Further studies are necessary to establish EV delivery as a possible therapy for the neurodegenerative phase of MS.

9. Estrogen involvement in non-reproductive functions

Estrogens from neural origin are involved in a variety of non-reproductive functions including regulation of neurogenesis, neuronal development, synaptic transmission, and plasticity in brain regions not directly related with the control of reproduction [107]. Estrogens regulate the expression of genes for various specific subtypes of dopamine (DA) receptors and 5-HT receptors in a region-specific manner to contribute to behavioral responses to changes of the internal and external environment.

DA neurotransmission of the ventral tegmental area (VTA)-nucleus accumbens (NAc) pathway associated with motivation is modulated by estrogens, which results in functional differences of the mesolimbic dopaminergic pathway and explains numerous sex differences in reward processing and related behavior described in human and animal studies [108]. Generally, increased circulating levels of estrogens in rodents, either during their estrous cycles or by estrogen treatment, have shown to contribute to elevated dopaminergic signaling. Estrogens influence manifold aspects of dopaminergic neurotransmission both presynaptically and postsynaptically, including: (a) DA synthesis, release, and degradation; (b) presynaptic and postsynaptic receptors; and (c) DA transporters that uptake DA from the synapse to terminate DA neurotransmission. Nuclear and membrane-associated estrogen receptors (ERs) may constitute the mechanisms by which estrogens affect the dopaminergic system. Precisely, ERs are distributed in dopaminergic pathways involved in reward, what is pertinent to the effects of estrogens on the CNS reward system [109]. In male mice, dopaminergic projections from the VTA to the ventral caudate express the ER β isoform, while dopaminergic projections to the dorsal caudate do not. Dopaminergic projections to the basolateral amygdala also express $ER\beta$. Estrogens also act rapidly through membrane-associated ERs on dopaminergic neurons in the striatum to affect DA release. Estrogens increase the activity of the

tyrosine hydroxylase (TH) and thus DA synthesis in the NAc by means of both nuclear ERs and membrane ER; induce presynaptic DA release in the striatum; and decrease DA turnover in the NAc and reduce degradation of DA so that DA remains longer at the synapse [108].

Estrogens upregulate the expression and activity of tryptophan hydroxylase (TPH) to increase 5-HT biosynthesis. Estrogen administration has been found to increase the TPH mRNA level [110]. In ovariectomized (OVX) guinea pigs, estrogen treatment alone increases the protein expression of TPH at the dorsal raphe nucleus (DRN). Surprisingly, while treatment with estrogen alone increases levels of TPH mRNA and TPH protein, it does not modify 5-HT content; rather, the treatment combining estrogen plus progesterone enhances 5-HT levels at both the DRN and the projected medial basal hypothalamus in guinea pigs [111]. Thus, there is a disparity in the sex hormone regulation of TPH mRNA and TPH protein levels versus 5-HT production in estrogen-treated animals, likely due to activity of TPH that is regulated by means of phosphorylation by protein kinase A (PKA) [112]. Since only phosphorylated TPH possesses catalytic activity, progesterone treatment might increase PKA expression and activity of TPH. After loss of ovarian function in many women, there may be an adjustment in the serotonergic neurons so that TPH protein levels recover. This also may indicate that the decline in TPH expression is a potential point of vulnerability in pre- and postmenopausal women who experience mental disorders such as depression related to the decreased level of 5-HT.

Estrogens regulate the 5-HT receptors, 5-HT2A and 2C, which display classic features of G protein-coupled receptors. The 5-HT2A receptor has been associated with suicide and depression, and its mRNAs are found in brain areas fundamental for the mood control, mental state, and cognition [113]. In humans, 5-HT2A receptor mRNA is found in the cortex and hippocampus, but not in the DRN, striatum, substantia nigra, or cerebellum [114]. Estrogens increase 5-HT2A receptor mRNA and binding site densities in the male rat brain. Healthy men have shown significantly higher levels of 5-HT2A receptor binding capacity than healthy women in the frontal and cingulate cortices, according to determinations by PET and radiotracer 18F-labeled altanserin [115]. Human studies suggest that estrogens increase 5-HT2A binding in higher forebrain regions.

The NAc receives major inputs from the amygdala and projects to the cortex and hypothalamus. These regions are fundamental for the cognitive function, emotion, mental state, mood, and neuroendocrine control. Thus, increase in 5-HT2A receptor densities by estrogen stimulation in these stated regions would control behavior and mood. A single administration of estradiol to OVX rats induces a significant increase in 5-HT2A receptor labeling in the NAc, the amygdala, DRN, anterior frontal, anterior cingulate, and primary olfactory cortex [116]. Moreover, concomitantly with the spontaneous estrogen-induced LH surge, 5-HT2A receptor densities increase compared to diestrous females or males in the frontal and cingulate cortex, olfactory tubercle and NAc. OVX produces a reduction of 5-HT2A receptor mRNA and protein levels, and long-term estrogen replacement reverses this effect in the frontal cortex [117]. The receptor 5-HT2C, a relevant contributor of many psychiatric and neurological disorders, is the most prominent 5-HT receptor subtype in the rat brain. 5-HT2C mRNA and protein are found in discrete regions of the rat brain. In the primate hypothalamus, dense populations of 5-HT2C mRNA-labeled cells are found in the anterior hypothalamus, periventricular nuclei, ventromedial nucleus, dorsal hypothalamic area, lateral hypothalamus, arcuate nucleus, and infundibular nucleus [118].

Estrogens regulate 5-HT auto-inhibition via the 5-HT1A auto-receptor. The 5-HT1A auto-receptor suppresses 5-HT synthesis by inhibiting firing of

serotonergic neurons at the DRN and median raphe and inhibiting 5-HT release in the hippocampus [119]. Therefore, the blockage of 5-HT1A receptors results in an antidepressant-like activity. Estrogen treatment lowers the 5-HT1A mRNA level in the dentate gyrus, CA2 region of the hippocampus, and DRN of OVX rats and in the DRN and median raphe of castrated rhesus monkeys, according to measurement by in situ hybridization. The5-HT1A auto-receptor is linked to an inhibitory G protein of the Gi/o/zfamily [120].

Estrogen treatment reduces 5-HT uptake to presynaptic cells by means of decreasing gene expression, translation, protein phosphorylation, trafficking, and stability of the serotonin transporter (SERT) [121]. Estrogens reduce SERT mRNA signal in the DRN of castrated rhesus monkeys treated with estrogen compared to the castrated control group and decrease [³H]-paroxetine binding, a selective indicator of 5-HT reuptake sites, in the hippocampus of estrogen-treated rats. Acute estrogen administration decreases SERT mRNA levels and 5-HT1A mRNA levels and binding. Thus, estrogen replacement therapy in postmenopausal women might decrease expression of the SERT gene and SERT protein [107]; thus 5-HT may remain longer in the extracellular space to continue neurotransmission. However, such argument does not support the observations that humans with depression have lower levels of 5-HT reuptake sites than healthy humans [122]. It might be that there is a mitigated 5-HT release and reduced levels of 5-HT in humans with depression to begin with, and a less 5-HT reuptake [107].

As we described, estrogens enhance 5-HT signaling via increasing 5-HT biosynthesis, upregulating expression and binding of receptors 5-HT2A and 5-HT2c, downregulating activity and expression of the 5-HT1A auto-receptor, decreasing expression of SERT gene and SERT protein allowing 5-HT to remain in the extracellular space for a longer period of time to continue neurotransmission, and decreasing 5-HT metabolism [107]. Consequently, the hypothesis of estrogen protection [123] claims that adjunctive estrogen treatment is beneficial at improving both symptoms of schizophrenia and the cognitive function, most notably in women. So far, the majority of studies on hormone replacement therapy for mood disorders have shown a positive response. Thus, estrogen has been demonstrated to improve mood in multiple mental illnesses, including major depressive disorder. Most recently, adjunctive estradiol treatment has been suggested as a promising treatment for women with comorbid depression and schizophrenia. In this direction, exploration of the impact of estrogen on serotonergic and other neurochemical pathways plus brain circuitry may guide not only the development of new hormone treatments but also assist in understanding the etiology of premenstrual dysphoric disorder, postnatal and perimenopausal depression, and even schizophrenia in women.

On the other hand, estrogens have shown that they do not always improve learning and memory. As an experimental example, acute administration of high doses of 17 β -estradiol and progesterone can impair performance in the memory test of Morris water maze. And most importantly, there is coherence among human studies since high endogenous levels of 17 β -estradiol are associated with poor performance on spatial tasks and cognitive function (Montreal Cognitive Assessment scale) and high dose of exogenous 17 β -estradiol impairs recognition memory. It is fundamental to assume that estradiol exerts a curvilinear influence on hippocampal-dependent performance, with low and high levels impairing versus a medium dose improving performance on a variety of tasks [124]. Similar dose response has been observed in the rapid effects of estrogens on dendritic spines, therefore, studies on the effects of estrogens on cellular morphology should take dose response into account. Furthermore, long-term exposure to estrogens can similarly result in dose-dependent responses on learning and memory, with low

levels of 17 β -estradiol reinforcing spatial working memory and high levels of estradiol hampering spatial working and reference memory. Studies have also shown that, whereas there is a dose-dependent facilitation in contextual fear conditioning by 17 β - and 17 α -estradiol, estrone results in dose-dependent impairments in contextual fear conditioning. Thus, it is essential to consider that the dose, frequency of administrations, and type of estrogen(s) utilized along with the type of memory task performed and the time point estrogens are administered (during acquisition or retrieval), are critical to the learning outcomes [124].

10. Conclusion

This review has concentrated on some cellular processes and signaling pathways acting during homeostatic molecular remodeling and their potential involvement in the maladaptive plasticity occurring in multiple neuropathologic conditions such as neurodegeneration and neuropsychiatric disorders.

There seem to be independent mechanisms for regulating presynaptic and postsynaptic strength, and there is evidence for independent homeostatic mechanisms operating on global and local spatial scales.

Involvement of different pharmacological approaches to compensate neurotransmission imbalance are under study, which may be considered as potential therapeutic approaches in neuropathologic conditions.

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Conflict of interest

Nothing declared.

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[124] Sheppard PAS, Choleris E, Galea LAM. Structural plasticity of the hippocampus in response to estrogens in female rodents. Molecular Brain. 2019; **12**:22. DOI: 10.1186/s13041-019-0442-7 **Chapter 4**

The Pharmacological Effects of Herbs on Catecholamine Signaling

Nobuyuki Yanagihara, Xiaoja Li, Yumiko Toyohira, Noriaki Satoh, Hui Shao, Yasuhiro Nozaki, Shin Ishikane, Fumi Takahashi, Ryo Okada, Hideyuki Kobayashi, Masato Tsutsui and Taizo Kita

Abstract

Herbs have many biologically and pharmacologically active compounds such as flavonoids and stilbenes. They have been used in remedies for various disorders. Here we review the effects of herbs on catecholamine synthesis and secretion in cultured bovine adrenal medullary cells. Ikarisoside A (1.0–100 µM), a flavonol glycoside, inhibited the catecholamine secretion induced by acetylcholine (0.3 mM). This inhibition was associated with the suppression of ²²Na⁺ and ⁴⁵Ca²⁺ influx induced by acetylcholine. The ethanol extract (0.0003-0.005%) of matsufushi (extract of pine nodules) inhibited the catecholamine secretion induced by acetylcholine. SJ-2, one of the stilbene compounds isolated from matsufushi, inhibited acetylcholine-induced catecholamine secretion. Matsufushi extract and SJ-2 reversibly inhibited acetylcholine-induced Na⁺ currents in *Xenopus* oocytes expressed with α3β4nicotinic acetylcholine receptors. Sweet tea is the processed leaves of Hydrangea *macrophylla*. The extract of sweet tea (0.3–1.0 mg/ml) suppressed catecholamine secretion induced by acetylcholine (0.3 mM). Moreover, sweet tea (0.1–1.0 mg/ml), ikarisoside A (1.0–100 μM), and matsufushi (0.001–0.003%) or SJ-2 (10–30 μM) inhibited acetylcholine-induced ¹⁴C-catecholamine synthesis from ¹⁴C-tyrosine. These findings indicate that ikarisoside A, matsufushi (or SJ-2), and sweet tea inhibit the catecholamine secretion and synthesis induced by acetylcholine in cultured bovine adrenal medullary cells and probably in sympathetic neurons.

Keywords: adrenal medullary cells, catecholamine secretion, ikarisoside A, matsufushi, sweet tea

1. Introduction

Since herbs have many biologically and pharmacologically active compounds such as flavonoids and stilbenes, they have been used in remedies for various disorders. A high dietary intake of herbs has become a focus of research because of herbs' potential to reduce the risks of diseases such as hypertension, coronary heart disease, diabetes, and cancers [1, 2]. Flavonoids are a group of plant secondary metabolites with variable phenolic structures, and they are found in plants, fruits, vegetables, roots, stems, flowers, wine, and tea [3, 4]. Over 5000 individual flavonoids have been reported [5], and 6 principal groups of flavonoids (flavones, flavonols, flavanones, flavanols, isoflavones, and anthocyanidins) are relatively common in human diets [1]. Polyphenol stilbenes have attracted scientific attention. For example, resveratrol (*trans-3*,4',5-trihydroxy- stilbene) is a natural phytoestrogen found in grapes, berries, and red wine [6, 7] that was reported to be implicated in the beneficial effect of red wine, i.e., the lower incidence of coronary artery disease in certain populations such as the French and the Greeks, despite diets rich in saturated fat and a rate of high smoking, which has been dubbed the "French paradox" [8].

In the human body, the most abundant catecholamines are adrenaline, noradrenaline, and dopamine, all of which are produced from phenylalanine and/or tyrosine. Catecholamines are biosynthesized mainly in the adrenal medulla, the postganglionic fibers of the sympathetic nervous system, and the central nervous system [2, 9, 10]. Catecholamines play very important roles in aspects of the cardiovascular system such as heart rate and blood pressure, blood glucose levels, and the general functions of the central and peripheral sympathetic nervous system [9].

Adrenal medullary cells derived from embryonic neural crests are functionally homologous to sympathetic postganglionic neurons [2, 10]. Our research demonstrated that in cultured bovine adrenal medullary cells, at least three distinct types of ionic channels participate in catecholamine secretion, including nicotinic acetylcholine receptor (nAChR)-ion channels, voltage-dependent Na⁺ channels, and voltage-dependent Ca²⁺ channels [2, 11]. In these cells, the Na⁺ influx induced by acetylcholine (ACh) via nAChR-ion channels or by veratridine via voltage-dependent Na⁺ channels is a prerequisite for Ca²⁺ influx via the activation of voltage-dependent Ca²⁺ channels and subsequent catecholamine secretion; in contrast, high K⁺ directly gates voltage-dependent Ca²⁺ channels to increase the Ca²⁺ influx without increasing the ²²Na⁺ influx [10, 11] (**Figure 1**). ACh-induced Ca²⁺ influx is also a prerequisite for the stimulation of catecholamine synthesis associated with the activation of tyrosine hydroxylase [2, 12–15]. The mechanisms underlying the stimulation of catecholamine synthesis and secretion mediated by these ion channels in adrenal medullary cells are thought to be similar to those of

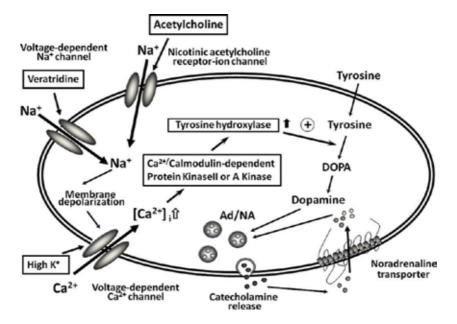


Figure 1.

The mechanism underlying the regulation of catecholamine synthesis, secretion, and reuptake in bovine adrenal medullary cells.

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noradrenaline in the sympathetic neurons and brain noradrenergic neurons [2]. Thus, adrenal medullary cells have provided a good model for the detailed analysis of antipsychotic [16], cardiovascular [17], and analgesic [18] drugs that act on catecholamine synthesis, secretion, and reuptake [2].

We have demonstrated the effects of several flavonoids and polyphenol stilbenes on catecholamine synthesis and secretion. For example, the treatment of bovine adrenal medullary cells with daidzein (an isoflavone derived from soy beans) stimulated basal catecholamine synthesis but inhibited the catecholamine synthesis and secretion induced by ACh [2, 19]. Genistein (another isoflavone in soy beans) but not daidzein stimulated the function of noradrenaline transporter in a human neuroblastoma cell line, SK-N-SH cells [2, 20]. Nobiletin (a compound of polymethoxy flavone in citrus fruits) stimulated the basal synthesis and secretion of catecholamines, but it suppressed both the ACh-induced synthesis of catecholamines and ACh-induced secretion of catecholamines [2, 21]. Resveratrol also inhibited the catecholamine synthesis and secretion induced by ACh [2, 22].

The present review summarizes our recent and current studies of the pharmacological effects of herbs and their components, i.e., ikarisoside A (a flavonol glycoside); matsufushi (extract of pine nodules), one of matsufushi's stilbene components (SJ-2); and sweet tea on the catecholamine signaling induced by ACh in cultured bovine adrenal medullary cells and on ACh-induced Na⁺ current in *Xenopus* oocytes expressing α 3 β 4 nAChRs.

2. Inhibitory effects of ikarisoside A, but not its aglycon, on the catecholamine secretion and synthesis induced by ACh

Ikarisoside A is a natural flavonol glycoside derived from plants of the genus *Epimedium*, which have been used in traditional Chinese medicine as tonics, antirheumatics, and aphrodisiacs [10, 23, 24], and is used as a tonic supplement in Japan. Ikarisoside A has antioxidant and anti-inflammatory effects [23] and anti-osteoporosis effects [10, 25]. Icariin, another flavonoid in the genus *Epimedium*, has an anti-stress effect in the forced swimming test in mice [26].

We observed that ikarisoside A (1–100 μ M) concentration dependently inhibited the secretion of catecholamines induced by ACh (0.3 m) (**Figure 2A**), but not the secretion of catecholamines induced by veratridine and 56 mK⁺ [10]. Ikarisoside A also suppressed the ²²Na⁺ influx and ⁴⁵Ca²⁺ influx induced by ACh in a concentrationdependent manner similar to that of catecholamine secretion (**Figure 2B, C**) [10]. Ikarisoside A is a flavonol glycoside with one rhamnose at the 3 position in the chemical structure. The aglycon of ikarisoside A is 3,5,7-trihydroxy-2-(4-hydroxyphenyl)-8-(3-methylbut-2-enyl)-4*H*-chromen-4one (**Figure 3A**).

It is interesting to note that the aglycon of ikarisoside A had little effect on catecholamine secretion induced by ACh (0.3 m) (**Figure 3B**), suggesting that the rhamnose moiety at the 3 position of ikarisoside A is essential to inhibit the function of nAChRion channels [10]. Ikarisoside A (1.0–100 or 10–100 μ M) inhibited ACh (0.3 mM)induced ¹⁴C-catecholamine synthesis from ¹⁴C-tyrosine and tyrosine hydroxylase activity [10].

3. Inhibitory effects of matsufushi and its stilbene component, SJ-2, on the catecholamine synthesis and secretion induced by ACh

Pine nodules of *Pinus tabulaeformis* or *Pinus massoniana* are formed by pine bark proliferation at places on the trunk or limbs that have undergone damage, either

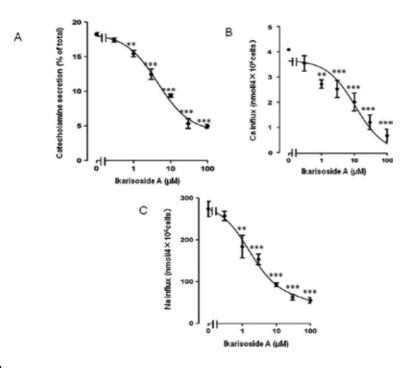


Figure 2.

Effects of ikarisoside A on ACh-induced catecholamine secretion (A), ${}^{45}Ca^{2+}$ influx (B), and ${}^{22}Na^+$ influx (C). (A) Cultured bovine adrenal medullary cells (10^6 /well) were stimulated with ACh (300μ M) in the presence or absence of ikarisoside A ($0.3-100 \mu$ M) for 10 min at 37° C. Catecholamine secretion is expressed as a percentage of the total catecholamines in the cells. (B and C) Cells (4×10^6 /well) are stimulated with ACh (300μ M) and 1.5μ Ci of ${}^{45}CaCl_2$ (B) or ${}^{22}NaCl$ (C) in the presence or absence of ikarisoside A ($0.3-100 \mu$ M) for 5 min at 37° C. ${}^{45}Ca^{2+}$ influx and ${}^{22}Na^+$ influx were expressed as nmol/4 × 10^6 cells. Data are means ± SEM from three separate experiments carried out in triplicate. P < 0.01 and P < 0.00 vs. ACh alone (by one-way ANOVA with Dunnett's multiple comparison post hoc test) (cited from [25]).

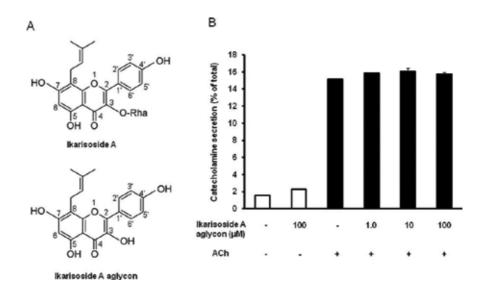


Figure 3.

Structures of ikarisoside A and its aglycon (A) and the effect of aglycon of ikarisoside A on the catecholamine secretion induced by ACh (0.3 mM) (B). (A) Chemical structures of ikarisoside A and its aglycon (3,5,7-trihydroxy-2-(4-hydroxyphenyl)-8- (3-methylbut-2-enyl)-4H-chromen-4-one). (B) Cells (10^{6} / well) were incubated with or without aglycon of ikarisoside A (1–100 μ M) and ACh (300 μ M) for 10 min at 37°C. Catecholamine secretion is expressed as a percentage of the total. Data are means ± SEM from three separate experiments carried out in triplicate (cited from [25]).

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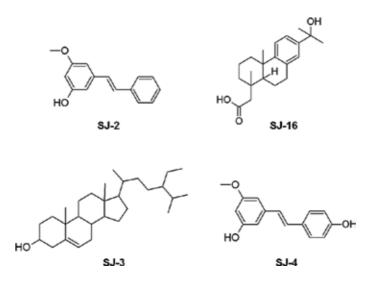


Figure 4.

Chemical structures of SJ-2, SJ-3, SJ-4, and SJ-16 (cited from [28]).

by pests or physical injury [27]. The effective curative components in pine nodule extract (matsufushi) have been used as an analgesic for joint pain, rheumatism, neuralgia, dysmenorrhea, and other complaints in traditional Chinese medicine [27–29]. Matsufushi is used as a healthy supplement in Japan.

Matsufushi ethanol extract (0.0003–0.005%) concentration dependently inhibited the catecholamine secretion and ⁴⁵Ca²⁺ influx induced by ACh (0.3 mM) and veratridine (0.1 mM), but not 56 mM K⁺ in cultured bovine adrenal medullary cells [27]. Four compounds (SJ-2, SL-3, SJ-4, and SJ-16) were isolated from matsufushi extract (**Figure 4**). SJ-2, a phenol stilbene, and the mixture of four compounds (Mix4; SJ-2, SJ-3, SJ-4, and SJ-16), but not each of the other separate compounds, inhibited the catecholamine secretion (**Figure 5**) and ⁴⁵Ca²⁺ influx [27] induced by ACh (0.3 mM). In *Xenopus* oocytes

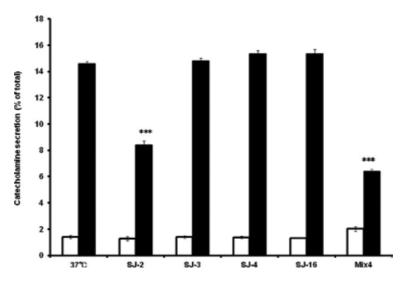


Figure 5.

Effects of SJ-2, SJ-3, SJ-4, SJ-16, and their mixture (Mix4) on catecholamine secretion induced by ACh in cultured bovine adrenal medullary cells. The cells (10^6 /well) were incubated with or without SJ-2 (10μ M), SJ-3 (10μ M), SJ-4 (10μ M), C-16 (10μ M), and their mixture (Mix 4) (10μ M) for 10 min at 37°C. Catecholamine secretion is expressed as a percentage of the total catecholamines in the cells. Data are means ± SEM from three separate experiments carried out in triplicate. P < 0.001 vs. ACh alone. Rha: rhamnose (cited from [28]).

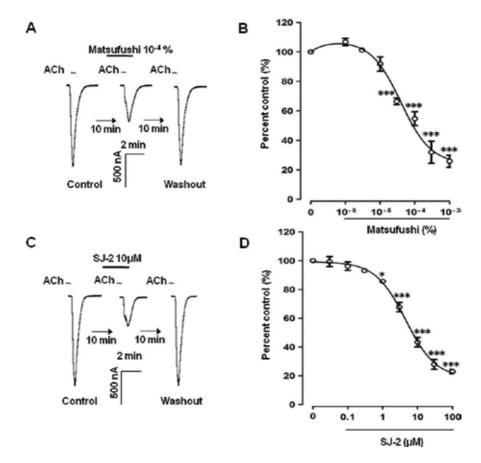


Figure 6.

Effects of ethanol extract of matsufushi and SJ-2 on inward currents induced by ACh in Xenopus oocytes expressing rat $\alpha_3\beta_4$ nAChRs. Representative traces from a single Xenopus oocyte are shown. The currents of matsufushi-treated (**A**) and SJ-2-treated (**C**) oocytes were recorded for 10 min after recording of the control currents, and the washout currents were obtained for 10 min after matsufushi extract and SJ-2 treatment. Matsufushi extract (0.0001%) and SJ-2 (10 µM) suppressed the currents induced by the EC50 (0.2 mM) of ACh, and the inhibitory effects were reversible. Concentration-response curve for the inhibitory effects of matsufushi extract (**B**) and SJ-2 (**D**) on ACh-induced currents. The peak current amplitude in the presence of matsufushi extract and SJ-2 was normalized to that of the control, and the effects are expressed as percentages of the control. Data are presented as means ± SEM from four separate experiments carried out in triplicate. P < 0.05 and $\stackrel{\text{w}}{=} P < 0.001$ w. the control (cited from [28]).

expressed with $\alpha 3\beta 4$ nAChRs, matsufushi extract and SJ-2 reversibly inhibited ACh (0.2 mM)-induced Na⁺ currents (**Figure 6A, C**). Matsufushi extract (0.00003–0.001%) (**Figure 6B**) and SJ-2 (1–100 μ M) (**Figure 6D**) significantly suppressed the Na⁺ current in a concentration-dependent manner [27]. In addition, matsufushi and SJ-2 suppressed the ACh (0.3 mM)-induced ¹⁴C-catecholamine synthesis from ¹⁴C-tyrosine and tyrosine hydroxylase activity [27]. These results suggest that matsufushi extract inhibits ACh-induced catecholamine synthesis and secretion mainly due to SJ-2 via the suppression of Na⁺ influx mediated through nAChR-ion channels [27].

4. Effects of the extract of sweet tea on catecholamine secretion and synthesis in adrenal medullary cells

Sweet tea is the processed leaves of *Hydrangea macrophylla* var. *thunbergii* Makino (hydrangeae dulcis folium), which is listed in the Japanese Pharmacopeia XV and used as a sweetening agent for diabetic patients. It also has antimicrobial and anti-allergic

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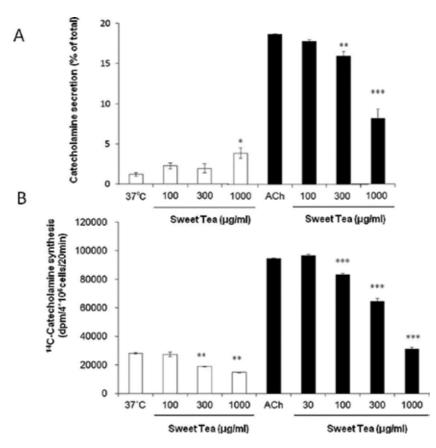


Figure 7.

Effects of sweet tea on ¹⁴C-catecholamine synthesis from [¹⁴C]tyrosine (**A**) and tyrosine hydroxylase activity (**B**) in the cells. (**A**) Cells (4×10^6 /dish) were incubated with L-[U-¹⁴C] tyrosine ($20 \ \mu$ M, $1 \ \mu$ Ci) in the presence or absence of sweet tea ($100-1000 \ \mu$ g/ml) and with or without $300-\mu$ M ACh at 37° C for 20 min. The ¹⁴C-catecholamines formed were measured. (**B**) Cells (10^6 /well) were incubated with L-[1-¹⁴C] tyrosine ($18 \ \mu$ M, $0.2 \ \mu$ Ci) in the presence or absence of sweet tea ($100-1000 \ \mu$ g/ml) and with or without $300-\mu$ M ACh at 37° C for 20 min. The ¹⁴C-catecholamines formed were measured. (**B**) Cells (10^6 /well) were incubated with L-[1-¹⁴C] tyrosine ($18 \ \mu$ M, $0.2 \ \mu$ Ci) in the presence or absence of sweet tea ($100-1000 \ \mu$ g/ml) and with or without $300 \ \mu$ M ACh at 37° C for 10 min, and tyrosine hydroxylase activity was measured. Data are means ± SEM from three separate experiments carried out in triplicate. P < 0.05, compared with 37° C (control), and "P < $0.01 \ and$ " P < $0.001 \ vs$. ACh alone in (**B**).

effects [30, 31]. Sweet tea is used as a healthy tea in Japan. There is, however, little evidence regarding sweet tea's effects on the sympathetic nervous system activity.

We investigated the effects of the extract of sweet tea on adrenal medullary cell function. A dry powder of sweet tea prepared from fermented leaves of hydrangeae dulcis folium was solubilized at 5.0 mg/ml and extracted at 90°C for 60 min. The extracted solution of sweet tea was used after centrifugation and filtration. The extract of sweet tea (1.0 mg/ml) slightly increased the basal secretion of catechol-amines (**Figure 7A**), whereas it suppressed the catecholamine secretion induced by ACh (0.3 mM) in a concentration-dependent manner (300–1000 µg/ml) (**Figure 7A**). In addition, the extract of sweet tea (300– or 100–1000 µg/ml) inhibited basal and ACh (0.3 mM)-induced ¹⁴C-catecholamine synthesis from ¹⁴C-tyrosine, respectively (**Figure 7B**). Sweet tea at concentrations of 3 mg/ml is usually used for drinking.

5. The insight of pharmacological potential of herbs in the catecholamine signaling induced by ACh in adrenal medulla

Adrenal medullary cells are derived from the embryonic neural crest and share many physiological and pharmacological properties with postganglionic sympathetic neurons [2]. The stimulation of AChRs in these cells increases the synthesis of catecholamines and causes the secretion of catecholamines into the systemic circulation [2, 11, 14]. In adrenal medullary cells, the Na⁺ influx induced by ACh via nAChR-ion channels is a prerequisite for Ca²⁺ influx via the activation of voltage-dependent Ca²⁺ channels and the subsequent catecholamine secretion and synthesis; in contrast, high K⁺ directly gates voltage-dependent Ca²⁺ channels to increase ⁴⁵Ca²⁺ influx [2, 11] (**Figure 1**).

As we noted, ikarisoside A and matsufushi (or SJ-2) inhibited the catecholamine secretion induced by ACh, but not the secretion induced by 56 mM K⁺ [10, 27]. In addition, ikarisoside A [10] and matsufushi [27] or SJ-2 [27] suppressed the Na⁺ current induced by ACh in *Xenopus* oocytes expressing $\alpha 3\beta 4$ nAChRs. These results suggest that the herbs and their components used as described herein inhibit the ACh-induced secretion and synthesis of catecholamines via a suppression of Na⁺ influx mediated through nAChRs in adrenal medullary cells [10, 27].

It is well known that catecholamines have important roles in the regulation of normal function in the central and peripheral sympathetic nervous systems as a neurotransmitter but also in the adrenal medulla as an endocrine hormone [10]. Strong and prolonged stress causes massive amounts of catecholamine release, which can lead to cardiovascular diseases (such as hypertension, coronary heart disease, heart failure, and atherosclerosis), and such stress also suppresses the immune system to induce some cancers [2, 9, 10]. Indeed, chronic heart failure is associated with the activation of the sympathetic nervous system as manifested by an increased circulating level of noradrenaline and increased regional activity of the sympathetic nervous system [2, 32]. It was reported that the stress hormone adrenaline stimulates β 2-adrenoceptors to activate the Gs-protein-dependent protein kinase A and the β -arrestin-1-mediated signaling pathway, which, in turn, suppresses p53 levels and triggers DNA damage [2, 33]. On the basis of these previous and present results, it appears that the herbs and their components such as ikarisoside A, matsufushi

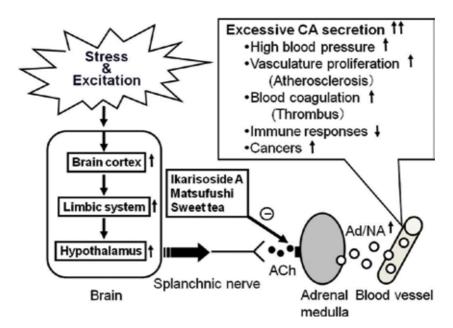


Figure 8.

Inhibitory mechanism of plant herbs (ikarisoside A, matsufushi, and sweet tea) on stress or excitation-induced excessive catecholamine secretion. Prolonged and strong stress or excitation stimulates the brain cortex, limbic system, and hypothalamus which evoke acetylcholine release from the splanchnic sympathetic nerves. Released acetylcholine induces a massive secretion of catecholamines from the adrenal medulla which may cause various deleterious symptoms or diseases such as high blood pressure (hypertension), vasculature proliferation (atherosclerosis), blood coagulation (thrombus), immune suppression, and cancers ([2] modified).

(or SJ-2), and sweet tea suppress the induction of a hyperactive catecholamine system induced by strong stress or emotional excitation (**Figure 8**).

6. Future perspective

Although the in vitro effects of the herbs and herb components described herein have been well clarified using cultured bovine adrenal medullary cells and *Xenopus* oocytes, the in vivo results are not yet clear. To confirm the pharmacological effects of these herbs on the catecholamine system, further in vivo studies of the effects of the administration of herbs to animals or humans are needed [2, 27]. We observed a disturbance of the autonomic nervous balance in women with climacteric symptoms measured by a power spectral analysis of heart rate variability [34]. Using this assay method, we will examine the effect of herbs on the autonomic nervous activity under some stress conditions.

7. Concluding remarks

We have reviewed the evidence that herbs and their components such as ikarisoside A, matsufushi (or SJ-2), and sweet tea inhibit the catecholamine synthesis and secretion induced by ACh in cultured bovine adrenal medullary cells and summarized them in **Table 1**. These findings may provide new insights into the pharmacological potentials of herbs on the hyperactive catecholamine system induced by stress.

Herb	Epimedium	Pine nodule	Sweet tea
Supplement	+	+	+
Compound	Ikarisoside A	SJ-2	ND
ACh-induced Na ⁺ current	Ļ	Ļ	ND
ACh-induced Ca ²⁺ influx	Ļ	Ļ	ND
Catecholamine secretion			
Basal	\rightarrow	\rightarrow	1
ACh-induced	Ļ	Ļ	Ļ
Catecholamine synthesis			
Basal	\rightarrow	\rightarrow	Ļ
ACh-induced	Ļ	Ļ	Ļ
Supplement +, available in Japan; ND, no	t determined.		

Table 1.

Summary of the effects of herbs and their components on catecholamine secretion and synthesis.

Conflict of interest

The authors have no conflict of interest to declare.

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Abbreviations

ACh acetylcholine nAChR nicotinic acetylcholine receptor

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Chapter 5

Thrombotic Tendencies in Excess Catecholamine States

Vivek K. Nambiar and Drisya Rajan Chalappurath

Abstract

Catecholamines are neurotransmitters distributed throughout the body including adrenal glands, chromaffin tissues and other tissues innervated by post ganglionic adrenergic neurons. The rate of release of medullary hormones is responsible for the control of serum catecholamines. Thrombogenicity of catecholamines are due various mechanisms including hypercoagulable states, endothelial damage, blood stasis and platelet aggregation. Oxidative stress generated by catecholamine excess causes coronary spasm, ultrastructural cell damage and arrhythmias. Elevated plasminogen activator inhibitor-1 during catecholamine excess causes hypercoagulability by hypofibrinosis. During stress, Catecholamines released in procoagulant environment causes vasoconstriction in adrenal veins resulting in venous thrombosis. Catecholamines generate moderately elevated levels of platelet count which enhances the risk of thrombosis. Hypercoagulability results in formation of coronary thrombus, rupture of atherosclerotic plaque and plaque progression due to gradual fibrinogen accumulation in the vessel walls. High levels of circulating catecholamines produce elevated levels thrombomodulin, the biomarkers of endothelial cell damage. In patients with hypertension in catecholamine excess, resistance to blood flow and damage to the integrity of blood vessels lead to atherosclerosis. A case report has been discussed which suggests an association of thrombotic tendency and catecholamine excess.

Keywords: catecholamines, thrombotic tendencies, atherosclerosis, hypercoagulation, platelet aggregation, endothelial damage, blood stasis

1. Introduction

Catecholamines are the class of neurotransmitters and hormones produced by the adrenergic nervous system, with a core structure of catechol and an amine group, regulating physiological processes [1]. Catechol refers to *ortho-dihydroxybenzene* and epinephrine and norepinephrine are the two major ethanolamine derivatives [2] occurring in the blood [3]. About 80% of epinephrine and 20% of norepinephrine are produced and released by the human adrenal medulla with a small amount of dopamine along with it [2].

1.1 Catecholamine biosynthesis

Catecholamines are synthesised and secreted from the chromaffin cells in adrenal medulla and sympathetic paraganglia [4]. Chromaffin granules produce catecholamines and fairly high concentration of adenosine triphosphate (ATP) and adenosine diphosphate (ADP) for active transport mechanism [2]. Syntheses of catecholamines occur at two levels [5]:

1. Sympathetic nervous fibre extremities for norepinephrine.

2. Chromaffin cells of adrenal medulla for both epinephrine and norepinephrine.

Tyrosine hydroxylase is the rate limiting enzyme of catecholamine synthesis. Tyrosine from dietary proteins is transported to the brain and tyrosine hydroxylase (in the dopaminergic and noradrenergic neurons of CNS and in the non-neuronal cells of gastrointestinal tract and kidneys) converts it to 3,4-dihydroxyphenylalanine (DOPA). In the presence of aromatic L-amino acid decarboxylase, DOPA is converted to dopamine (DA) which is transported to vesicular storage granules. It may be released in the dopaminergic neurons or converted to norepinephrine in the noradrenergic neurons by dopamine beta-hydroxylase [6]. Norepinephrine is converted to epinephrine by phenyl ethanolamine-N-methyltransferase (PNMT) [4].

Medullary hormones synthesised and stored in the chromaffin granules are a pool of ready to release hormones upon stimulation [7].

1.2 Catecholamine functions and receptors

Norepinephrine and epinephrine together contribute to an increase in cardiac output, heart-rate, respiration, blood pressure, blood flow to muscles, alertness and decrease in splanchnic and renal blood flow [2]. Norepinephrine alone causes a decrease in heart-rate indirectly as a result of reflex bradycardia because of vaso-constriction effects [2].

Functions of catecholamines are mediated and regulated by alpha, beta adrenoreceptors and dopamine receptors. Each receptor located at distinct locations serves different functions and actions.

Alpha-1 receptors mediate adrenergic vasoconstriction [8]. Alpha-2 receptor stimulation causes inhibition of norepinephrine secretion [6]. Three subtypes of α_2 receptors are α_{2A} , α_{2B} , and α_{2C} . Alpha-2A in the peripheral postsynaptic region contributes to vasoconstriction in different vascular beds [8]. Beta-1 receptor stimulation causes positive inotropic and chronotropic action [6]. Beta-2 receptors stimulate vasodilation [9]. Beta 3 causes lipolysis in fatty cells and increase renin secretion [9].

Two types of dopamine receptors are D1 and D2.Stimulation of D1 receptors in coronary, renal, mesenteric and cerebrovascular beds causes vasodilation, diuresis and natriuresis. D2 receptor stimulation in presynaptic nerve endings causes inhibition of norepinephrine release [6].

1.3 Catecholamine metabolism

Neuronal catecholamines get metabolised intraneuronally after leakage from its stores [1]. Catecholamine metabolism mainly occurs in the liver [2]. There are two major catecholamine metabolites (i) metanephrines (constitutes 15% of urinary catecholamine) and (ii) vanillylmandelic acid, VMA (major end-point of norepinephrine and epinephrine metabolism [1] and constitutes 85% of excreted catecholamine) [2].

1.4 Catecholamine inactivation

Two major enzymes involved in the inactivation of catecholamines are catecholo-methyl transferase (COMT) and monoamine oxidase (MAO).COMT adds methyl

group to hydroxyl at 3-position of catechol ring and MAO catalyses oxidative deamination reaction [2].

Norepinephrine is inactivated by two processes (i) reuptake into the nerve terminals by the active carrier mechanism associated with Na⁺-K⁺-ATPase of which some part is stored to granular vesicles and the rest is metabolised and (ii) extraneuronal accumulation to vascular smooth muscles [10].

1.5 Catecholamine excess

The three distinct peripheral catecholamine systems are [11].

1. Sympathetic nervous system (SNS)

2. Adrenomedullary hormone system (AHS)

3. Dopamine autocrine/paracrine system (DOPA)

Sympathetic nervous system: the measure of norepinephrine release in the specific regions allows the clinical assessment of organ-specific sympathetic nervous tone and penetrating analysis of SNS in the pathophysiology of a disease state [12].

Adrenomedullary hormone system: the catecholamines stored in the chromaffin cells get released by innervations of preganglionic sympathetic fibres of splanchnic nerves and get dispersed after its entry to medulla [2].

Dopamine derived from plasma DOPA acts as an autocrine/paracrine substance. Human urine contains higher concentration of dopamine and its metabolites rather than norepinephrine and its metabolites as dopamine is generated in the nonnoradrenergic and non-adrenergic cells [13].

Catecholamines are distributed throughout the body including adrenal glands, chromaffin tissues and other tissues innervated by post ganglionic adrenergic neurons [3]. The rate of release of medullary hormones is responsible for the control of serum catecholamines [2].

The cellular uptake and metabolism of catecholamines in the extraneuronal tissues and the exogenous administration of catecholamines together contribute to 25% of total metabolism of catecholamines produced in the SNS and adrenergic chromaffin cells [1].

During stressful conditions, the level of enzymes catalysing biosynthesis of catecholamines increase and enzymes degrading the catecholamines decrease [14]. Stress-induced changes in the adrenergic receptors occur due to the combination of neuronal and extraneuronal catecholamine uptake system, level of cAMP, density of adrenoceptor subtypes and coupling of agonist receptor complex with different G-proteins [14].

During chronic stress, increased levels of catecholamines produce aminolutins and oxyradicals. Oxidative stress thus generated causes coronary spasm, ultrastructural cell damage and arrhythmias [14]. They also cause Ca²⁺overload and cardiac dysfunction by acting on beta-1-adrenoceptor signal transduction pathway [14].

1.6 Catecholamine excess in pheochromocytoma

Catecholamine excess is seen in certain types of tumour such as pheochromocytomas that are functional tumours of the chromaffin cells [15]. Norepinephrine secreting tumours are known to cause hypertension (in 80–90% of these patients [4]) and hypertensive crisis and their abrupt release may cause arrhythmias [6]. Of this nearly 40% of the patients suffer from sustained hypertension (norepinephrine secreting tumours), 45% from paroxysmal hypertension (epinephrine secreting tumours) and 5–15% from normotension (dopamine secreting tumours) [4].

Other vasoactive substances like neuropeptide Y, adrenomedullin and atrial natriuretic peptide contribute to hypertension in these patients [6]. Neuropeptide Y increase coronary and peripheral vascular resistance and potentiate norepinephrine-induced vasoconstriction [7].

Pheochromocytoma produces enormous amounts of catecholamines due to mutation of succinate dehydrogenase by oxidative phosphorylation deficit [6].Two major mechanisms in pheochromocytoma that leads to catecholamine excess are translocation of dopamine into storage granules and the presence of dopamine beta hydroxylase enzyme [1].

In extra-adrenal tumours, norepinephrine levels are high, in adrenal tumours both norepinephrine and epinephrine are in excess and in multiple endocrine neoplasia type 2 and neurofibromatosis type 1, epinephrine levels are high [6].

In non-functional/silent pheochromocytoma, metanephrine levels are elevated instead of catecholamines [6].

Complication of pheochromocytoma in hypertensive crisis are sustained hypertension by continuous elevated levels of catecholamine causing vasoconstriction and orthostatic hypertension due to reduced blood volume that resulting in vasoconstriction, postural tachycardia and postural hypotension [4].

Neurological complications are hypertensive encephalopathy, haemorrhage (due to paroxysmal hypertension) and acute ischemic stroke (due to postural hypertension) [4].

In 78% of pheochromocytoma patients, thrombus exists at the time of their tumour discovery and 75% of the secondary thrombi are located in inferior vena cava or heart or both [16]. Venous thromboembolism and acute arterial thrombi are well described in such patients [17] but evidence of stroke, aortic thrombi and myocardial infarction are fewer [18].

Several mechanisms explain thrombotic tendencies in pheochromocytomas by elevated catecholamines:

- Catecholamines generate moderately elevated levels of platelet count which enhance the risk of thrombosis [19].
- Platelet aggregation is increased in pheochromocytoma [18] and produce thrombi in low-flow areas causing coronary events [19].
- Contributing factors to thrombus formation are erythropoietin, pro-coagulation and serotonin [6].
- Catecholamine excess in other tumours like paragangliomas may cause ADPmediated platelet aggregation leading to thrombotic events [20].
- During stress, catecholamine release in procoagulant environment causes vasoconstriction in adrenal veins resulting in venous thrombosis [21].
- Loss of antithrombin III in glomerular diseases associated with hypertension may increase thrombotic risk [22].
- Pheochromocytoma causes thrombosis of caudal vena cava and aortic thromboembolism through Virchow's triad list [20, 22]:

- (i) Endothelial damage
- (ii) Blood stasis
- (iii) Hypercoagulability state

Mortality of pheochromocytoma patients occur due to either acute intramyocardial necrosis or diffuse patchy myocardial fibrosis [23].

2. Mechanisms associated with thrombus formation in catecholamine excess

Thrombus formation and dissolution are physiological processes with dynamic balance between procoagulation and anticoagulation mechanism [24]. Catecholamine excess contributes to long-term development of atherosclerosis and cardiovascular risk [25] due to their effect on carbohydrate and lipid metabolism [23].

3. Cascade of thrombus formation

Thrombotic process occurs by the activation of coagulation cascade in different phases [26].

- 1. Initiation phase—the extrinsic pathway of coagulation in which low amounts of active pro-coagulant factors are produced.
- 2. Amplification phase—level of coagulation factors increase.
- 3. Propagation phase—active coagulation factors are generated like thrombin which binds to phospholipid at the surface of activated platelets.
- 4. Stabilization phase—thrombin provides strength and stability to growing clot and thrombin activatable fibrinolysis inhibitor.

Injury to the vessel walls due to resistance, in catecholamine excess states, causes exposure of glycoprotein tissue factor (TF) to the blood. After exposure, TF bind to Factor VII/VIIa and forms tertiary complex in the presence of calcium to activate factor X to Xa, factor IX to IXa and factor VIIa. At the site of injury, platelet activation, conversion of fibrinogen to fibrin and thrombus formation occurs. Platelets are able to regulate their gene and protein expression for binding to sub endothelium [26].

Diseases affecting Virchow's triad can induce clot formation [27].

4. Hypercoagulability

Hypercoagulability is the tendency to produce thrombosis by inherited or acquired molecular defects. Thrombus is formed based on the number of predisposing factors and environmental stress [13].

Cancer is the second major cause of hypercoagulation with clinical thrombosis in 15% and thrombosis on autopsy in 50% patients [28].

Catecholamines increase plasminogen activator inhibitor-1 (PAI-1) mRNA by expression of PAI-1 genes in the cardiovascular system by neurological and neuroendocrine mechanisms via beta-adrenoreceptors [29]. PAI-1, a member of serineprotease inhibitor, inhibits both tissue type and urinary-type plasminogen activator. Elevated PAI-1 during catecholamine excess causes hypercoagulability by hypofibrinosis. High levels of PAI-1 act on adipokines in the atherogenic process and cause atherosclerotic lesions and atheromas [30]. Epinephrine regulates Beta-mediated tissue-type plasminogen activator to produce an increase in fibrinolytic capacity [25].

Antithrombin III (AT-III) inhibits the protein-splitting reaction, characteristic of haemostasis, by interfering with the activity of four of the serine proteases and antagonising thrombin [31]. Hypertension associated glomerular diseases in catecholamine excess may cause loss of AT-III by protein loss in the urine. Hypoalbuminemia and hypercholesterolemia in Proteinuria causes platelet hyperaggregability and aggravate hypercoagulability. Increased levels of fibrinogen and thromboxane in protein-losing nephropathy contribute to thrombosis [27].

Hypercoagulability result in formation of coronary thrombus, rupture of atherosclerotic plaque and plaque progression due to gradual fibrinogen accumulation in the vessel walls [24].

In patients with borderline hypertension, coagulation activation occurs before clinical manifestations of vascular damage appear [32].

5. Platelet aggregation

In thrombus formation, a high proportion of platelets translocates from their initial point of attachment to the injured vessel wall and forms a firm adhesion contact [31]. Catecholamine infusion induces platelet aggregation in ex-vivo and animal models [16]. They activate platelets by interacting with platelet α 2-adrenoreceptor in in-vitro studies. Stimulation of α 1-adrenoreceptor inhibits platelet aggregation by inactivating platelet response to epinephrine [25]. In-vivo adrenergic infusion causes platelet activation and an increase in platelet size, aggregation and releasing factors [24]. Elevated factor V: C, factor VIII: C, v Willebrand Factor: Ag, tissue-plasminogen activator (t-PA): Ag, contribute to risk of atherothrombotic events [25]. Redistribution of cardiac blood flow in catechol-amine infusion causes regional slowing of blood flow and enhanced tendency for intravascular platelet aggregation [23].

The anticoagulative effect of epinephrine is explained by a 50% decrease in the ratio of maximal coagulation activation and maximal fibrinolytic activation by decreased tumour necrosis factor (TNF) and interleukin (IL) 10 [33]. Epinephrine induce fibrinogen receptor exposure and fibrinogen binding to potentiate platelet aggregation. Epinephrine infusion causes short-term recruitment of platelets and functionally active Factor VIII from the spleen for thrombus formation. They discharge high molecular von Willebrand Factor (vWF) which increase the activity of platelet adhesion from its storage compartments. Elevated epinephrine levels increase platelet deposition on the atherosclerotic vessels with high local shear rate [25].

Norepinephrine activates platelet aggregation in-vivo by stimulating α 2-adrenoreceptor [24].

It causes platelet aggregation in two phases [16]:

- Initial reversible aggregation.
- Simulated release of adenosine diphosphate.

Norepinephrine initiates platelet thrombosis by directly damaging the endothelial walls of vessels and breaking its integrity [24]. In acute mental stress, 60% of variation in thrombin formation depends on amount of norepinephrine secreted and sensitivity of β 2-adrenergic receptor [19].

6. Clotting time

Adrenergic activation by epinephrine injection increases factor VIII: C and cause a decrease in clotting time. In-vivo adrenergic infusion stimulate β -2 adrenergic vascular receptors to release clotting factors like FVIII, v WF, and t-PA from endothelium to the circulation [24] resulting in thrombotic risk in both coronary artery disease and hypertension [25].

7. Endothelial damage

Endothelium maintains balance between prothrombotic and atherothrombotic tendencies [27]. Endothelial cells provide a non-thrombogenic surface that inhibits platelets and blood cells from adhering and activating coagulation cascade [34]. Antithrombotics like thrombomodulin, heparin sulfate, tissue factor pathway inhibitor and protein S are involved in reducing the activation of AT III and protein c to elevate antithrombotic tendencies. Synthesis and induction of prothrombotic mediators such as tissue factor and antifibrinolytic factors like PAI-1 take part in prothrombotic tendencies [27].

Disproportion in the tendencies causes endothelial damage where subendothelial proteins are exposed and favour platelet adhesion [27]. According to Gando et al. [35], endothelial damage can occur due to both exogenous and endogenous release of catecholamines. High levels of circulating catecholamines produce elevated levels thrombomodulin, the biomarkers of endothelial cell damage [36]. Due to loss of endothelial cell protection and expression of procoagulant and prothrombotic molecules, platelets adhere to the subendothelial surface and release ADP and thromboxane A2 causing imbalance in haemostasis and thrombosis [37].

8. Blood stasis

Blood stasis plays an important role in the formation of thrombus. Thrombus forms in regions of blood stasis like narrow artery due to atheromatous plaque, mechanical vasoconstriction, prolonged immobilisation and varicose vein. From the injured tissue, tissue thromboplastin produced causes initial thrombus formation followed by platelet activation by thrombin. Thrombin transforms fibrinogen to insoluble fibrin and release to stagnant blood [36].

In patients with hypertension in catecholamine excess, resistance to blood flow and damage to the integrity of blood vessels lead to atherosclerosis [27, 32]. Three mechanisms explain the overall increased vascular resistance [37]:

(i) Rarefaction of arterioles and capillaries.

- (ii) Reduced internal diameter of arterioles.
- (iii) Increase in arterial and arteriolar wall mass.

S. No.	Mechanism of thrombosis	Factors involved in thrombosis
1.	Hypercoagulability	1. Tissue plasminogen activator inhibitor-1 (t-PAI-1)
		2. Antithrombin-III (AT-III)
		3. Fibrinogen
		4. Thromboxane
2.	Platelet aggregation	1. Factor V: C
		2. Factor VIII: C
		3. v Willebrand factor (v WF)
		4. t-PAI-1
		5. Tumour necrosis factor (TNF)
		6. Interleukin 10 (IL-10)
3.	Clotting time	1. Factor VIII:C
	-	2. (vWF)
		3. t-PAI-1
4.	Endothelial damage	1. t-PAI-1
	Ũ	2. Thrombomodulin
		3. Thromboxane A2
5.	Blood stasis	1. Thromboplastin
		2. Thrombin

Table 1.

Summary of factors associated with thrombosis.

The significant role of sympathetic nervous system in elevated haemostatic activity in atherosclerosis and thrombus formation explains the role of catecholamines in cardiovascular diseases and associated arterial thrombus formation [25]. The factors associated with thrombosis in different mechanisms are summarised in **Table 1**.

9. Management of thrombosis

Hypertension associated with catecholamine excess plays an important role in the thrombus formation by mechanisms explained above. So treatment for hypertension is the primary objective in treatment and prevention of thrombotic tendencies. The most preferred antihypertensive is phenoxybenzamine which is an alpha and beta receptor antagonist given at 10 mg once a day with gradual dose titration to 30 mg three times a day. Appropriate anti-coagulation with heparin is vital for patients with risk of thrombosis to prevent devastating outcomes [19].

Removal of tumours producing excess catecholamines play a crucial role in alleviating symptoms associated with hypertension and thus thrombus formation. Due exaggerated fall in blood pressure after surgical removal of these tumours, adequate vascular volume replacement with oral fluids and salt is recommended [38].

Here in a case report has been discussed which suggests an association of thrombotic tendency and catecholamine excess.

10. Case report

A Case report on cerebral venous thrombosis due to pheochromocytoma in a patient with Von Hippel Lindau (VHL) mutation [39] explains an unusual connection between thrombosis and pheochromocytoma in a 15 year old boy with pheochromocytoma and associated hypertension. The patient was primarily presented with cerebral venous thrombosis and seizures. Initial examination revealed left

frontoparietal hematoma and superior sagittal and transverse sinus thrombus for which he was treated with heparin and antiepileptics. Follow-up showed elevated blood pressure and high urine catecholamine levels. The diagnosis of abdominal mass after evaluation for consistent elevated blood pressure turned out to be pheochromocytoma. He underwent para-aortic dissection and right adrenalectomy.

VHL disease is an autosomal, dominant inherited tumour syndrome which occurs as a result of a germline mutation in the VHL gene. Even though its association with pheochromocytoma is under-studied, VHL proteins are found to cause tumorigenesis in pheochromocytoma by induction of either gain of function, dominant negative effect or gene dosage effect in chromaffin cells [40].

Some mechanisms are established to explain the association of Catecholamine excess in pheochromocytoma and thrombus formation:

- 1. Tumours produce different chemicals with prothrombotic nature that cause hypercoagulability as a paraneoplastic feature [41]. The release of procoagulants, fibrinolytics and proteolytic factors induce thrombosis. Neoplastic vascular invasion damages the thromboresistant properties of vessel wall and tumour cell attachment to the endothelial surface leading to endothelial retraction and exposure of basement membrane. The damaged endothelial cells and matrix cause platelet adhesion and aggregation by release of procoagulatory and antagonistic substances [42].
- 2. Adrenal tumours release excess catecholamines and cause thrombophilia and clot formation [16].
- 3. In the case of VHL mutation, certain changes are produced in the Hypoxia inducible factor (HIF) resulting in polycythaemia and increased venous thrombosis [43]. HIF regulates both oxygen hemostasis and erythropoietin receptor expression [44, 45].

To associate hereditary disorders with thrombosis in excess catecholamine state is restricted because of unavailability of literatures explaining the same. Excess catecholamines themselves produce thrombosis even though an inherent mutation of VHL was found in the above patient as a cause of thrombosis. Hypertension associated with high catecholamine levels make a patient more prone to thrombosis than others with normal blood pressure. In patients with tumours such as pheochromocytoma and paraganglioma, there is high catecholamine levels which increase the risk of thrombotic tendencies compared to other patients with catecholamine excess alone.

11. Conclusion

Though there have been evidences of thrombus formation after infusion of catecholamines in animals and humans, available evidence of the exact mechanism of thrombotic tendencies is scanty. Catecholamine excess states are a diagnostic challenge due to heterogenous clinical presentation like fluctuating hypertension in young which may not attract enough clinical and laboratory attention. Thrombogenicity of catecholamines are due various mechanisms including hypercoagulable states, endothelial damage, blood stasis and platelet aggregation. Surgical removal of tumours associated with catecholamine excess provide a clinical approach in treating abnormalities of thrombus formation.

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Conflict of interest

There are no conflicts of interest.

Abbreviations

ADP ATP	adenosine diphosphate adenosine triphosphate
DOPA	3,4-dihydroxyphenylalanine
DA	dopamine
PNMT	phenyl ethanolamine-N-methyl transferase
COMT	catechol-O-methyl transferase
MAO	monoamine oxidase
SNS	sympathetic nervous system
ANS	autonomic nervous system
DOPA	dopamine autocrine/paracrine system
PAI-1	plasminogen activator inhibitor-1
AT-III	antithrombin III
t-PA	tissue plasminogen activator
TNF	tumour necrosis factor
IL	interleukin
v WF	von Willebrand factor
VHL	Von Hippel Lindau
HIF	hypoxia inducible factor

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Drawing on the expertise of experienced researchers in neurotransmission and catecholamines, this book provides a brief overview of the latest knowledge in the field. The book contains an introductory chapter that aims to explain the subsequent four chapters for researchers who are new to the field.

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