
Distribution of the Endocannabinoid System in the Central Nervous System

Sherry Shu-Jung Hu and Ken Mackie

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S.S.-J. Hu

Department of Psychology, National Cheng Kung University, Tainan 70101, Taiwan

e-mail: shujunghu@gmail.com

K. Mackie (✉)

Department of Psychological and Brain Sciences, Indiana University, 47405 Bloomington, IN, USA

Gill Center for Biomolecular Science, Indiana University, 47405 Bloomington, IN, USA

e-mail: kmackie@indiana.edu

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Abstract

The endocannabinoid system consists of endogenous cannabinoids (endocannabinoids), the enzymes that synthesize and degrade endocannabinoids, and the receptors that transduce the effects of endocannabinoids. Much of what we know about the function of endocannabinoids comes from studies that combine localization of endocannabinoid system components with physiological

or behavioral approaches. This review will focus on the localization of the best-known components of the endocannabinoid system for which the strongest anatomical evidence exists.

Keywords

CB₁ cannabinoid receptor • Immunocytochemistry • In situ hybridization

Abbreviations

2-AG	2-Arachidonoylglycerol
ABHD4/6/12	α/β -hydrolase domain 4/6/12
AEA or anandamide	<i>N</i> -arachidonylethanolamine
BLA	Basolateral amygdala
CB ₁ and CB ₂	Cannabinoid receptor 1 and 2
CRIP1a	Cannabinoid receptor interacting protein 1a
DAGL α/β	Diacylglycerol lipase α/β
DRG	Dorsal root ganglion
FAAH	Fatty acid amide hydrolase
GDE1	Glycerophosphodiesterase 1
M1	Muscarinic cholinergic receptor
MAGL	Monoacylglycerol lipase
mGluR5	Metabotropic glutamate receptor 5
MSNs	Striatal medium spiny neurons
NAAA	<i>N</i> -acylethanolamine-hydrolyzing acid amidase
NAPE-PLD	<i>N</i> -acyl phosphatidylethanolamine phospholipase D
PAG	Periaqueductal gray
SN	Substantia nigra
TH	Tyrosine hydroxylase
VTA	Ventral tegmental area

1 Introduction

1.1 Overview

Studies of the distribution of the protein components of the endocannabinoid system (ECS) are motivated by the notion that we can gain important insights into the function of the ECS by understanding the location of its component enzymes and receptors. This review has been written with that concept in mind, with an emphasis on studies that integrate function and location. Because of space limitations, this review will focus on components of the ECS in the CNS with the highest quality localization data. Thus, the emphasis will be on the cannabinoid

CB₁ receptors and on the enzymes *N*-acyl phosphatidylethanolamine phospholipase D (NAPE-PLD) (Di Marzo et al. 1994; Okamoto et al. 2004), fatty acid amide hydrolase (FAAH) (Cravatt et al. 2001), diacylglycerol lipase (DAGL) (Bisogno et al. 2003; Gao et al. 2010), monoacylglycerol lipase (MAGL) (Dinh et al. 2002, 2004), and α/β -hydrolase domain 6 (ABHD6) (Blankman et al. 2007; Marris et al. 2010). The interesting topic of neuronal expression of CB₂ receptors is complicated by the high inducibility of CB₂ in pathological conditions, the low levels of CB₂ compared to CB₁, the presence of CB₂ in microglia and endothelial cells, and nonspecific antibodies. Nonetheless, CB₂ may be present on a limited population of neurons with functional consequences, for example: (Viscomi et al. 2009; den Boon et al. 2012; Zhang et al. 2014). The interested reader can refer to the literature for additional details and consideration (Marsicano and Kuner 2008; Atwood and Mackie 2010). We will also focus on ECS component localization in the mature brain in this review. The ECS and its components are subject to dynamic regulation in the developing CNS, a topic that has been well covered in recent reviews (Harkany et al. 2008; Maccarrone et al. 2014).

A variety of techniques are available to determine protein localization in tissues. These include autoradiography (labeled ligands or GTP γ S), *in situ* hybridization, and antibody-based techniques. Each of these techniques gives complementary information, which taken together, can enhance our understanding of ECS function—autoradiography requires high affinity binding of a probe to the protein, *in situ* hybridization detects mRNA (often useful for identifying cell types synthesizing a protein), and antibody-based techniques detect a (protein) epitope resembling the epitope that antibody was raised against. As antibody-based techniques are most commonly used for ECS localization, it is important to be aware of some of their caveats. The issue of spurious results from antibody studies has received considerable attention in the neuroscience community (Saper 2005; Rhodes and Trimmer 2006; Michel et al. 2009; Manning et al. 2012), though numerous examples of poor practice continue to be published. “Best practices” have been discussed and implemented by several journals (Saper 2005; Rhodes and Trimmer 2006; Manning et al. 2012) and are summarized in Table 1. As much as is possible, studies that have adhered to those practices will be emphasized in this review.

1.2 Cells Expressing Components of the ECS

While the ECS in neurons has received the most attention due to the prominent effects of endogenous and exogenous cannabinoids on neuronal function, it is important to appreciate that glial cells are a major component of the ECS, sometimes acting independently of neurons and sometimes in concert. Strong evidence supports the presence of CB₁ receptors in some astrocytes and microglia (Rodriguez et al. 2001; Stella 2010; Bosier et al. 2013), and these cells as well as oligodendrocytes are prodigious synthesizers and degraders of endocannabinoids (Walter et al. 2002; Stella 2009); however, their complement of enzymes vary somewhat from those in neurons (Marris et al. 2010). In addition, the CNS

Table 1 Antibody controls for immunocytochemistry

Approach	Strength
Block with immunizing protein	Weak (block with immunizing protein is a necessary, but not sufficient condition to establish antibody specificity)
Lack of staining with preimmune serum	Weak (immune response can induce the expression of many serum proteins that can interact with extraneous epitopes)
Lack of staining in knockout (KO)	Strong (requires knockout; need to understand how the KO was made, particularly if the exon which the antibody was raised against remains; knockout studies should be conducted in parallel (identical tissue processing, incubation times, etc.) with experiments performed with tissue that expresses the genetically deleted protein)
Identical staining with antibody directed against independent epitopes	Strong (for alternatively spliced proteins need to ascertain that the appropriate exons are expressed)
Lack of staining in knockdown	Strong (need independent verification of knockdown; stronger if knockdown from limited population of cells so “controls” are adjacent)
Detection of protein in transfected cells	Medium (demonstrates antibody can detect protein, but not necessarily in tissue; stronger if conducted in a mixed population of cells (expressing and non-expressing) and target protein is epitope-tagged to allow its unequivocal detection; difficult to apply if target protein is present in the cell line used)
Detection of appropriate band on western blot	Can be helpful correlation (however, the conformation of protein in fixed tissue and denatured gel is quite different; many examples where antibody won't detect a specific band in Western blots and works for immunocytochemistry and vice versa)
Correlation with <i>in situ</i> hybridization	Useful at the level of cell populations (assumes mRNA is translated to protein)
Correlation with function	Helpful (e.g., CB ₁ -mediated responses and CB ₁ receptors detected; however can lead to circular reasoning)

vasculature also participates in endocannabinoid signaling (Gebremedhin et al. 1999; Schley et al. 2009; Zhang et al. 2009; Dowie et al. 2014).

1.3 Subcellular Localization of CB₁ Cannabinoid Receptors

Due to their prominent effects on presynaptic calcium channels and synaptic transmission, it is not surprising that high levels of CB₁ receptors are found on some presynaptic terminals and preterminal axon segments (Katona et al. 1999; Nyiri et al. 2005a). The ability of endocannabinoids to suppress spiking in low threshold spiking cortical interneurons and some pyramidal cells suggests that CB₁ receptors are also located on neuron somata (Marinelli et al. 2009). Within neurons,

CB₁ receptors are sometimes associated with specialized structures (e.g., multi-lamellar bodies (Katona et al. 1999)) that may be involved in their trafficking. A recent though not uncontroversial finding (Benard et al. 2012; Hebert-Chatelain et al. 2014a, b; vs. Morozov et al. 2013) is that CB₁ receptors are associated with some mitochondria, including mitochondria found in astrocytes, where they contribute to energy balance and may play a role in synaptic plasticity.

2 Retina

2.1 Receptors

Using the immunocytochemical approach, CB₁ receptors are found in subsets of amacrine cells and horizontal cells and densely expressed in the inner plexiform layer (Straiker et al. 1999a; Yazulla et al. 1999). CB₁ are also present in rod and cone photoreceptor terminals in a wide range of vertebrate retinas, including human (Straiker et al. 1999a, b; Hu et al. 2010), as well as in rod bipolar cells in rat retina (Yazulla et al. 1999).

2.2 Synthetic Enzymes

While the presence of synthetic enzymes for *N*-arachidonylethanolamine (anandamide or AEA) has not yet been examined in retina, two isoforms of the major synthetic enzyme for 2-arachidonoylglycerol (2-AG), diacylglycerol lipase- α and - β (DAGL α/β) (Bisogno et al. 2003; Gao et al. 2010), were both found in the mouse retina. DAGL α appears in the two synaptic layers, the outer plexiform layer and inner plexiform layer, whereas DAGL β immunoreactivity is limited to retinal blood vessels. Furthermore, DAGL α is present in postsynaptic type 1 OFF cone bipolar cells juxtaposed to CB₁-containing cone photoreceptor terminals (Hu et al. 2010). These findings suggest that retrograde 2-AG signaling exists at type 1 OFF bipolar cell-cone photoreceptor synapses, consisting of presynaptic CB₁ receptors and postsynaptic DAGL α .

2.3 Degradative Enzymes

Both degradative enzymes for anandamide and related *N*-acylethanolamines (NAEs), fatty acid amide hydrolase (FAAH) (Cravatt et al. 2001) and *N*-acylethanolamine-hydrolyzing acid amidase (NAAA) (Guo et al. 2005; Tsuboi et al. 2005), are present in retina. FAAH immunoreactivity was first detected in large ganglion cells, in large dopaminergic amacrine cells, in the dendrites of star-burst amacrine cells, and in the somata of horizontal cells of the rat retina (Yazulla et al. 1999). In the mouse retina, FAAH is widely expressed in the inner segments, outer nuclear layer, ganglion cell layer, and in the axon terminals of photoreceptors in the outer plexiform layer, and also co-localizes with CB₁ in subpopulation of amacrine cells

in the inner nuclear layer and in cells in the ganglion cell layer (Hu et al. 2010). The most notable difference between mouse and rat FAAH staining is the absence of staining in horizontal cells of the mouse, possibly a function of species difference. Finally, NAAA immunoreactivity is limited to retinal pigment epithelium in the mouse retina (Hu et al. 2010).

Among five candidate degradative enzymes for 2-AG, monoacylglycerol lipase (MAGL) and α/β -hydrolase domain 6 (ABHD6) (Blankman et al. 2007) were found in the mouse retina. Similar to DAGL α , MAGL staining was found in the outer and inner plexiform layers and additionally in the ganglion cell layer (Hu et al. 2010). While MAGL is not co-localized with DAGL α in the inner plexiform layer, it is distal to DAGL α in the outer plexiform layer. MAGL is present in photoreceptor terminals, including the rod spherules and cone pedicles (Hu et al. 2010), where it is well positioned to break down 2-AG after retrograde release onto both rod and cone terminals.

On the other hand, ABHD6 is widely distributed in the inner plexiform layer, inner nuclear layer, and ganglion cell layer. ABHD6 is localized to the calbindin- and GAD67-positive amacrine cells in the inner nuclear layer and to the dendrites of ganglion or displaced amacrine cells in the proximal inner plexiform layer (Hu et al. 2010). The postsynaptic staining of ABHD6 suggests its potential role in the breakdown of extrasynaptic 2-AG that has diffused beyond its intended target synapses.

3 Cerebral Cortex

3.1 Neocortex

3.1.1 Receptors

CB₁ receptors are densely expressed in all regions of the cortex, with high levels found in cingulate gyrus, frontal cortex, secondary somatosensory, and motor cortex, reviewed in (Mackie 2005). An immunocytochemical study found a heterogeneous distribution of CB₁-immunoreactive axons across neocortex in macaque monkeys and humans. Most neocortical association regions, such as the prefrontal and cingulate cortex, contain a higher density of CB₁-immunoreactive axons compared to the primary motor and somatosensory cortices (Eggan and Lewis 2007). Furthermore, in many cortical regions, CB₁-immunoreactivity displays a distinctly laminar pattern of expression, corresponding to the cytoarchitectonic boundaries. Although the distribution of CB₁ immunoreactivity across monkey neocortical regions is broadly similar to that observed in the rat using immunocytochemical (Egertova and Elphick 2000; Hajos et al. 2000; Katona et al. 2001; Bodor et al. 2005) and autoradiographic (Herkenham et al. 1991) approaches, there are several species differences in the laminar distribution of CB₁-immunoreactive axons. For example, CB₁-immunoreactive axons were reported to be most densely expressed in layers 2–3 and 6 and least densely expressed in layer 4 of the rat frontal and cingulate cortex (Egertova and Elphick 2000). In contrast, the highest density of CB₁-immunoreactive axons is localized in layer 4 of the same regions in monkey

(Eggen and Lewis 2007). In the primary somatosensory cortex, CB₁-immunoreactive axons are most densely localized in layer 5A in rat (Bodor et al. 2005), whereas a relatively similar density of CB₁-immunoreactive axons exists in layers 2–3 and 5–6, and a sparse axonal labeling presents in layer 4 of monkey (Eggen and Lewis 2007). Despite these differences, the laminar distribution of CB₁-immunoreactivity is quite similar within primates. For example, both autoradiographic (Glass et al. 1997) and immunocytochemical (Eggen and Lewis 2007) methods yield similar laminar distribution of CB₁ receptors in human and monkey neocortex, respectively.

In forebrain, high levels of CB₁ receptors have been primarily found on large cholecystokinin (CKK)-containing basket interneurons, with lesser levels found in non-CKK-expressing neurons (Marsicano and Lutz 1999). For example, CB₁ immunoreactivity is absent in nonadapting multipolar interneurons, such as the parvalbumin or bi-tufted adapting somatostatin-expressing interneurons (Tsou et al. 1999; Bodor et al. 2005). Despite original studies suggesting lack of expression of CB₁ immunoreactivity in principal glutamatergic neurons (Tsou et al. 1998a; Freund et al. 2003), more sensitive *in situ* hybridization studies revealed low but detectable levels of CB₁ mRNA in the great majority of glutamatergic neurons in many cortical regions including neocortex (Monory et al. 2006). Moreover, a single-cell real-time polymerase chain reaction (qPCR) study revealed that at least 50% of neocortical glutamatergic pyramidal neurons contain CB₁ mRNA (Hill et al. 2007), which is consistent with functional data showing that the CB₁ receptor agonist WIN-55212-2 decreased the intracortical electrical stimulation-evoked excitatory postsynaptic currents (EPSCs) in a CB₁ antagonist-dependent fashion (Hill et al. 2007). In summary, CB₁ receptors are densely expressed in multiple cortical regions. While CB₁ expression is highest on CCK-positive interneurons, functionally important CB₁ receptors are present on multiple neuron populations.

3.1.2 Synthetic Enzymes

Studies examining the protein and mRNA distribution of the anandamide synthesizing enzymes, NAPE-PLD, in neocortex have found it to be widespread, but at relatively low levels (Egertova et al. 2008). On the other hand, *in situ* hybridization revealed that moderate levels of DAGL α and DAGL β mRNA are expressed in mouse cerebral cortex (Yoshida et al. 2006). DAGL α is typically dendritic, as shown for a cultured cortical neuron in Fig. 1.

3.1.3 Degradative Enzymes

FAAH-immunoreactive neuronal somata and dendrites are present throughout the transitional and neocortical regions of the mouse cerebral cortex and are often surrounded by CB₁-immunoreactive fibers. Except for layer 1, FAAH immunostaining is evident in all cortical layers, especially in the large cells in layer 5 (Egertova et al. 2003).

In situ hybridization revealed that high levels of MAGL mRNA exist throughout the rat brain cortex, especially in layers 4, deep 5, and 6 (Dinh et al. 2002).

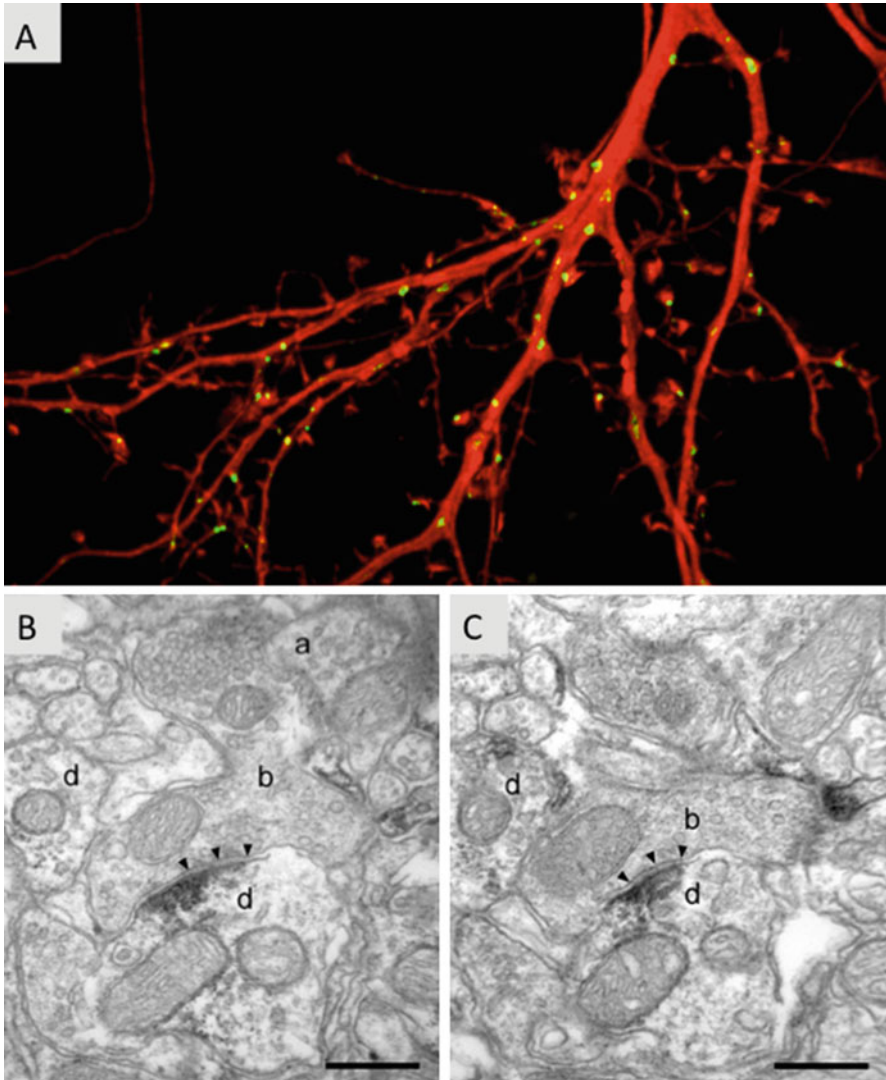


Fig. 1 Postsynaptic expression of diacylglycerol lipase alpha (DAGL α). (A) Dual immunofluorescent staining for DAGL α (green) and MAP2 (red). DAGL α as detected by a C-terminal antibody in a cultured neuron was found in close proximity to many dendritic spines. (B and C) Two consecutive ultrathin sections from mouse spinal cord demonstrate that the electron-dense reaction product representing DAGL α immunoreactivity (arrowheads) is present in the dendrite (d) close to the asymmetric postsynaptic density across from an excitatory terminal (b). Scale bar = 200 nm. Original figures provided by Barna Dudok and Istvan Katona (A) and Rita Nyilas and Istvan Katona (B, C)

In the mouse prefrontal cortex, ABHD6 immunoreactivity is predominantly localized to the postsynaptic dendritic spines, which is juxtaposed to the CB₁-positive presynaptic terminals (Marrs et al. 2010). Moreover, selective inhibition of ABHD6 allowed the induction of CB₁-mediated long-term depression by the subthreshold stimulation, suggesting this enzyme is a *bona fide* member of the endocannabinoid signaling system (Marrs et al. 2010).

3.2 Olfactory Areas (Olfactory Bulb, Piriform Cortex, Associated Regions)

3.2.1 Receptors

In the olfactory bulb, CB₁ receptors are highly expressed in the inner granular cell layer, followed by the inner plexiform layer, while less are expressed in the external plexiform layer, the glomerular layer, and the accessory olfactory bulb (Herkenham et al. 1991; Tsou et al. 1998a; Egertova and Elphick 2000). However, a detailed examination revealed that CB₁ receptor immunoreactivity is abundant in the periglomerular processes of GAD65-positive interneurons and the inner granular cell layer (A. Straiker, personal communication). Furthermore, CB₁ receptors are expressed uniformly by most neurons in the anterior olfactory nucleus and the anterior commissure, which connect the olfactory bulbs (Herkenham et al. 1991; Matsuda et al. 1993; Glass et al. 1997; Tsou et al. 1998a; Egertova and Elphick 2000). Moreover, most neurons in the piriform cortex contain CB₁ mRNA (Marsicano and Lutz 1999). Finally, CB₁ receptor immunoreactivity is present on dendritic processes in the olfactory epithelium of *Xenopus laevis* tadpoles, where it mediates cannabinoid modulation of odor-induced spike-associated currents in individual olfactory receptor neurons (Czesnik et al. 2007).

3.2.2 Synthetic Enzymes

NAPE-PLD mRNA expression has been detected in several olfactory areas. For example, NAPE-PLD mRNA is present in granule and periglomerular cells in the olfactory bulb and in neuronal cell body layers in the olfactory tubercle and the piriform cortex (Egertova et al. 2008). Interestingly, the immunostaining of NAPE-PLD is very intense in glomeruli of the accessory olfactory bulb and in the vomeronasal axons projecting into the accessory olfactory bulb (Egertova et al. 2008). Moderate expression of DAGL α and DAGL β mRNA was found in mouse olfactory bulb by in situ hybridization (Yoshida et al. 2006).

3.2.3 Degradative Enzymes

Despite the overall low levels of CB₁ receptor expression in the olfactory bulb, FAAH immunoreactivity is intense, especially in fibers of the olfactory nerve and in the olfactory glomeruli, as well as in the somata and dendrites of mitral cells (Egertova et al. 2003). Moreover, FAAH-immunoreactive neuronal somata are evident in the majority of cortical olfactory regions receiving direct input from the olfactory bulb, including the anterior olfactory nucleus, the piriform cortex,

the tenia tecta, and the indusium griseum (Egertova et al. 2003). Importantly, in all of these regions, FAAH-immunoreactive postsynaptic neuronal somata are surrounded by a complementary network of CB₁-immunoreactive fibers, which supports the hypothesis that anandamide and other acyl amides may function as transynaptic signaling molecules (Egertova et al. 2000).

3.3 Hippocampal Formation

3.3.1 Receptors

The hippocampus is highly involved in cognitive functions such as the spatial and declarative learning and memory, thereby drawing much attention as a site of action of endogenous and exogenous cannabinoids due to their effects on memory. Early autoradiographic studies found very high levels of CB₁ receptors in all subfields of the hippocampus as well as the dentate gyrus (Herkenham et al. 1991; Jansen et al. 1992). In situ hybridization studies revealed that most CB₁ receptor expression arose from a restricted subset of interneurons (Matsuda et al. 1990, 1993; Maillieux and Vanderhaeghen 1992). Immunocytochemical studies showed high levels of CB₁ receptors on large CKK-positive basket and Schaffer collateral-associated interneurons in the hippocampal pyramidal cell layer, as well as the molecular layer and at the base of the granule cell layer in the dentate gyrus (Katona et al. 1999; Marsicano and Lutz 1999; Tsou et al. 1999; Egertova and Elphick 2000) (e.g., Fig. 2a), while few or no CB₁ receptors were found on parvalbumin-positive interneurons. Agonist activation of CB₁ receptors on these interneurons was found to decrease power in theta, gamma, and ripple oscillations in the hippocampus (Robbe et al. 2006). Actions of cannabinoids such as these appear to be widespread in cortical and subcortical circuits (Sales-Carbonell et al. 2013) and may contribute to the effects of cannabinoids on memory (Chen et al. 2003; Freund et al. 2003; Klausberger et al. 2005; Robbe et al. 2006).

Building on earlier studies that detected CB₁ mRNA, but not protein in CA1 and CA3 pyramidal neurons, studies in which CB₁ receptors were selectively deleted from either GABAergic or glutamatergic neurons allowed the conclusive anatomical identification of low levels of CB₁ protein in glutamatergic hippocampal pyramidal neurons (Marsicano et al. 2003; Lutz 2004). Indeed, several groups independently identified CB₁ protein in glutamatergic pyramidal neurons in the CA1 and CA3 regions (Degroot et al. 2006; Katona et al. 2006) (examples of CB₁ expression in hippocampal inhibitory and excitatory terminals are shown in Fig. 2A–D). However, among glutamatergic neurons, the highest CB₁ levels are present in dentate gyrus mossy cells (Kawamura et al. 2006; Monory et al. 2006). Interestingly, many CB₁-positive hilar mossy cells also contain dopamine D₂ receptors, suggesting that this region might be involved in the interactions of these two neuromodulatory systems (Degroot et al. 2006). Moreover, CB₁ immunoreactivity was also identified in the majority of hippocampal cholinergic nerve terminals, where presynaptic CB₁ receptors control acetylcholine release in vitro (Gifford and Ashby 1996) and in vivo (Degroot et al. 2006). On the other hand, CB₁

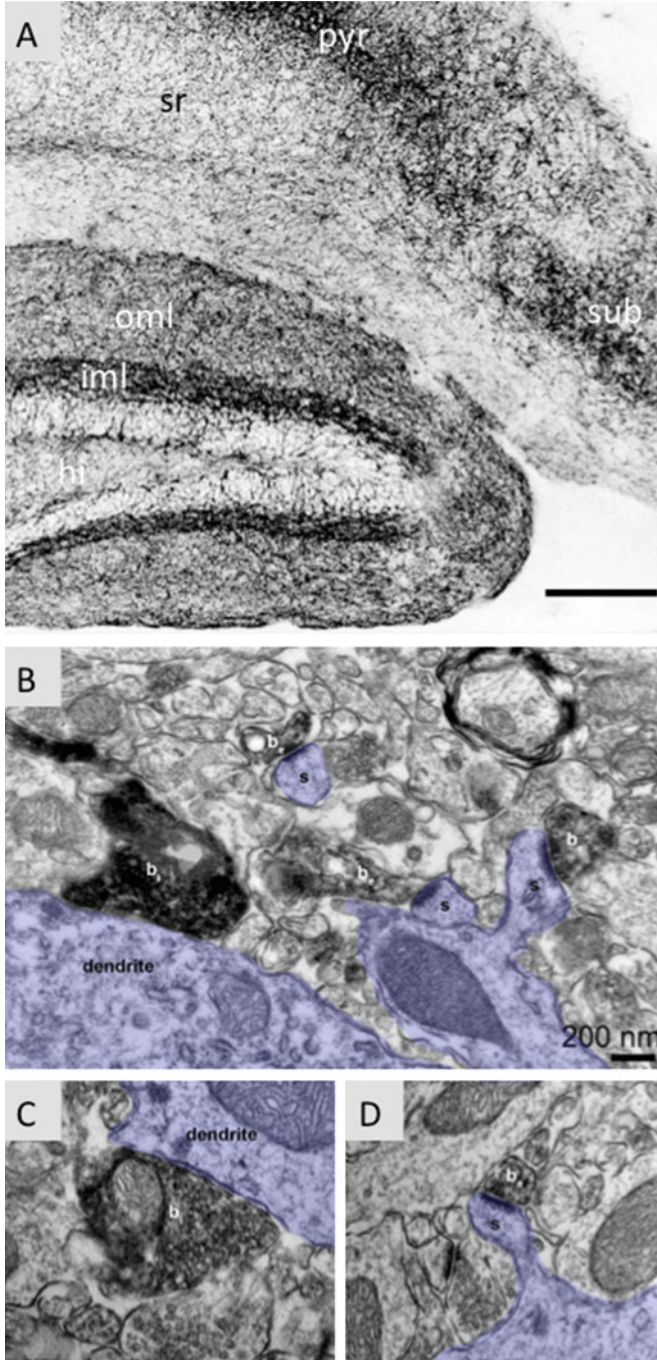


Fig. 2 CB₁ expression in rodent brain is primarily presynaptic. (A) CB₁ receptors in mouse hippocampal formation were detected by an antibody directed to its C-terminus. A dense meshwork of CB₁-expressing axons is evident. Levels are particularly high in the inner molecular layer

receptors appeared to be absent from granule cells of the dentate gyrus. Finally, low but detectable levels of CB₁ receptors were found in a subset of progenitor cells in the subgranular zone of the dentate gyrus, which regulate proliferation, survival, and differentiation of these adult progenitor cells (Aguado et al. 2005; Galve-Roperh et al. 2007).

3.3.2 Synthetic Enzymes

NAPE-PLD mRNA, as revealed by in situ hybridization, is most intensely expressed in the dentate gyrus granule cell layer, followed by the pyramidal cell layer of the hippocampus throughout all three fields (CA1-CA3) (Cristino et al. 2008; Egertova et al. 2008; Nyilas et al. 2008). Moreover, NAPE-PLD immunoreactivity was detected in granule cell axons (Egertova et al. 2008; Nyilas et al. 2008) as well as in many neurons of the hilus region in mouse dentate gyrus (Cristino et al. 2008; Nyilas et al. 2008). Therefore, in contrast to 2-AG's prominent role as a retrograde signaling messenger (Kano et al. 2009), anandamide and related NAEs generated by NAPE-PLD in axons may act as anterograde synaptic signaling molecules to regulate the activity of postsynaptic neurons (Egertova et al. 2008).

High levels of DAGL α mRNA and protein have been found in postsynaptic dendritic spines of the majority of hippocampal pyramidal neurons, including the spine head, neck, or both (Katona et al. 2006; Yoshida et al. 2006). A similar pattern for DAGL α was found in the postmortem human hippocampus, with highest levels in strata radiatum and oriens of the cornu ammonis and in the inner third of stratum moleculare of the dentate gyrus (Ludanyi et al. 2011). The juxtaposition of DAGL α -immunoreactive postsynaptic dendritic spines to CB₁-expressing presynaptic terminals at excitatory glutamatergic synapses highlights the prominence of 2-AG as a retrograde messenger at many synapses (Katona et al. 2006).

3.3.3 Degradative Enzymes

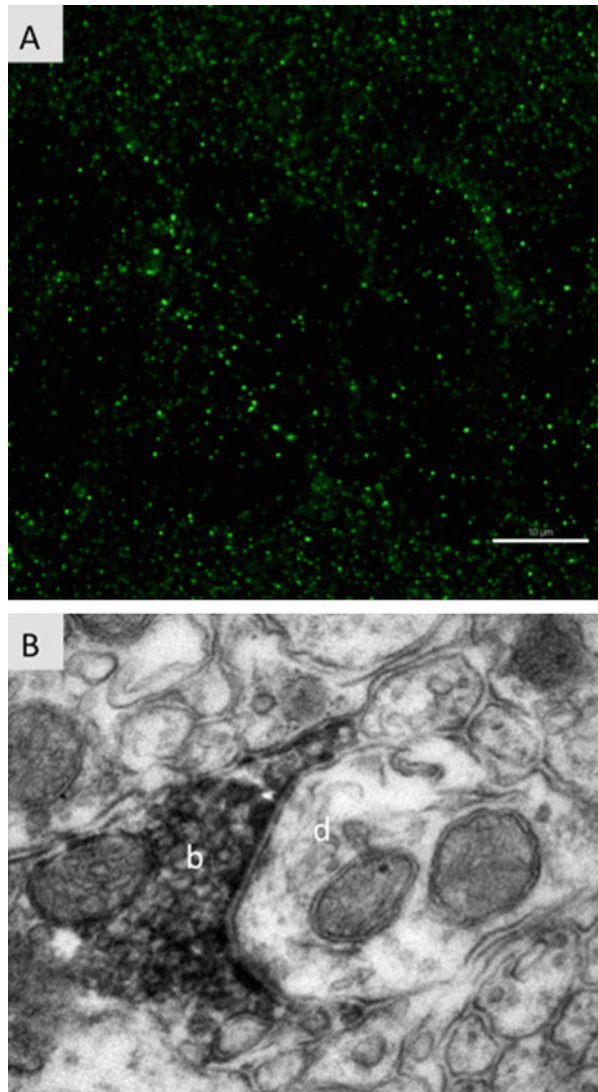
FAAH is highly expressed in the somata and proximal dendrites of the pyramidal cells, which are densely innervated by CB₁-positive axon terminals (Egertova et al. 1998, 2003; Tsou et al. 1998b; Gulyas et al. 2004). However, it is important to note that FAAH is more frequently present on the membrane surface of intracellular organelles (e.g., mitochondria, smooth endoplasmic reticulum) than on somatic or dendritic plasma membranes (Gulyas et al. 2004). In addition, FAAH



Fig. 2 (continued) of the dentate gyrus and the pyramidal neuron layer: *hi* hilus, *iml* inner molecular layer, *oml* outer molecular layer, *pyr* pyramidal cell layer, *sr* stratum radiatum, *sub* subiculum. **(B–D)** Expression of CB₁ receptors at the ultrastructural level in the mouse hippocampus assessed using a C-terminal CB₁ antibody. **(B)** Electron-dense reaction product is present in both inhibitory (*b_i*) and excitatory (*b_e*) boutons. Note that the excitatory boutons synapse onto spines (*s*) while the inhibitory boutons form synapses on a dendritic shaft. Dendritic structures are pseudocolored blue for ease of identification. **(C)** Higher magnification image showing a CB₁-positive inhibitory terminal synapsing onto the shaft of a dendrite. **(D)** Higher magnification image showing a CB₁-positive excitatory bouton forming an asymmetric synapse onto a dendritic spine. Scale bar = 200 nm in **B**. Original figures provided by Jim Wager-Miller **(A)** and Chris Henstridge and Istvan Katona **(B–D)**

immunostaining is also evident in the somata of mouse dentate gyrus granule cells (Egertova et al. 2003), which contrasts to the absence of FAAH immunoreactivity in granule cells of the rat dentate gyrus (Tsou et al. 1998b; Gulyas et al. 2004). An in situ hybridization study showed that MAGL mRNA is abundantly expressed in the CA3 field of rat hippocampus (Dinh et al. 2002). Additional immunocytochemical studies revealed that MAGL protein is particularly prominent in axon terminals of granule cells, CA3 pyramidal cells, and some interneurons of rat hippocampus (Gulyas et al. 2004), as well as in axon terminals of glutamatergic neurons in both rodent and human hippocampus (Yoshida et al. 2006; Ludanyi et al. 2011) (Fig. 3A, B).

Fig. 3 Presynaptic localization of monoacylglycerol lipase (MAGL) detected with an antibody recognizing residues positioned in the middle of mouse MAGL. (A) Punctate expression of MAGL immunoreactivity in the mouse hippocampus pyramidal cell layer. Scale bar = 10 μm . (B) Electron-dense reaction product representing MAGL immunoreactivity is present in an inhibitory terminal (*b*) synapsing onto a dendritic shaft (*d*). Original figures provided by Stephen Woodhams and Istvan Katona



3.4 Cortical Subplate (Other Amygdala Nuclei)

3.4.1 Receptors

The amygdala is divided into a cortical component (e.g., the basolateral, lateral, and basomedial nuclei) and a striatal component (e.g., the central and medial nuclei) (Swanson and Petrovich 1998). This subdivision corresponds to the different structural organization and neurochemical properties of the two components. For example, while most principal neurons in the cortical subdivision of the amygdala are glutamatergic, the great majority of neurons in the striatal subdivision are GABAergic. Therefore, similar to cortex, high levels of CB₁ receptors are primarily expressed in the CCK-positive GABAergic basket cells (Marsicano and Lutz 1999; Katona et al. 2001). For example, in the basal (but not lateral) nucleus of the basolateral amygdala (BLA), high levels of CB₁ receptors are localized to presynaptic CCK-positive GABAergic terminals at invaginating synapses (Yoshida et al. 2011). On the other hand, as in cortex, low but functionally significant levels of CB₁ receptor are present in glutamatergic neurons in the cortical part of amygdala (Monory et al. 2006; Yoshida et al. 2011). Consistent with the above anatomical data, functional studies suggest that CB₁ receptors and endocannabinoids facilitated extinction of fear conditioning *via* inhibiting GABA release in the BLA (Marsicano et al. 2002). Finally, in contrast to earlier studies finding very weak CB₁ immunoreactivity within the central amygdala (Katona et al. 2001; Kamprath et al. 2011), a recent study using a highly sensitive CB₁ receptor antibody showed the presence of functional CB₁ receptors in the central amygdala (Ramikie et al. 2014).

3.4.2 Synthetic Enzymes

In the amygdaloid complex, NAPE-PLD mRNA is expressed in neurons of the cortical and medial amygdaloid nuclei, while less staining has been found in the basal and lateral nuclei (Egertova et al. 2008). At invaginating synapses, a unique type of perisomatic synapses in the basal nucleus of the BLA, DAGL α is recruited to somatic membrane of postsynaptic pyramidal neurons, juxtaposed to the CB₁-, MAGL-, and CCK-containing presynaptic terminals (Yoshida et al. 2011). In the central amygdala, DAGL α is localized to postsynaptic dendritic spine heads and dendritic shafts, juxtaposed to the CB₁-containing presynaptic terminals at glutamatergic synapses (Ramikie et al. 2014).

3.4.3 Degradative Enzymes

FAAH-immunoreactive neuronal somata are also present throughout the basolateral complex of the amygdala, which includes the lateral, basolateral (BLA), and basomedial nuclei. In all of these nuclei, the FAAH-immunoreactive somata are surrounded by CB₁-immunoreactive fibers (Egertova et al. 2003).

As mentioned above (Sect. 3.4.2), MAGL is co-expressed with CB₁ receptors in the presynaptic CCK-positive terminals at invaginating synapses in the basal nucleus of the BLA. Together with the postsynaptic DAGL α expression, this constitutes a molecular convergence for 2-AG-mediated retrograde signaling.

This contrasts with the flat perisomatic synapses made by parvalbumin-positive interneurons (Yoshida et al. 2011) where no such arrangement is present.

4 Subcortical Nuclei (Striatum, Basal Ganglia)

4.1 Striatum (Dorsal, Caudate)

4.1.1 Receptors

The subcortical nuclei with the highest level of CB₁ receptor expression are the basal ganglia. In situ hybridization studies showed that many striatal medium spiny neurons (MSNs) express CB₁ receptors, while adult pallidal and nigral neurons contain little or no CB₁ mRNA (Matsuda et al. 1993; Julian et al. 2003). Due to its axonal terminal localization, the high levels of pallidal and nigral CB₁ receptor binding and protein observed in autoradiographic and immunocytochemical studies mostly arose from GABAergic neurons projecting from the caudate putamen (Tsou et al. 1998a; Egertova and Elphick 2000). The staining for CB₁ in axons is denser in the globus pallidus than in the caudate putamen, while both show a gradient in the staining intensity increasing from medial to lateral (Tsou et al. 1998a; Egertova and Elphick 2000). Moreover, CB₁ receptors are present on both the striatonigral and striatopallidal projection pathways; thus, they are well positioned to modulate both the direct and indirect striatal output pathways (Hohmann and Herkenham 2000). In addition, a study utilizing a high-sensitivity CB₁ receptor antibody showed CB₁ protein to be intensely expressed on GABAergic axon terminals of striatal MSNs and parvalbumin-positive interneurons (Uchigashima et al. 2007). Finally, CB₁ protein was found on excitatory corticostriatal afferents (Gerdeman and Lovinger 2001; Rodriguez et al. 2001; Uchigashima et al. 2007), GABAergic aspiny interneurons (Hohmann and Herkenham 2000) and neurons in the subthalamic nucleus (Matsuda et al. 1993), with important functional implications (Kreitzer and Malenka 2007).

4.1.2 Synthetic Enzymes

The immunostaining of NAPE-PLD is present in the caudate putamen (Egertova et al. 2008). On the other hand, because 2-AG is synthesized by DAGL after membrane depolarization and G_q-coupled receptor activation, an immunocytochemical study was carried out to examine the subcellular distribution of DAGL α , metabotropic glutamate receptor 5 (mGluR5), and muscarinic cholinergic receptor 1 (M1) in mouse striatum (Uchigashima et al. 2007). Even though all three proteins were present on somatodendritic membranes of MSNs, only DAGL α and mGluR5 were present in spines and the perisynaptic region, while M1 receptors were absent in these domains (Uchigashima et al. 2007). This subcellular arrangement may account for the differential involvement of mGluR5 and M1 in endocannabinoid-mediated depolarization-induced suppression of inhibition and depolarization-induced suppression of excitation (Uchigashima et al. 2007).

4.1.3 Degradative Enzymes

In the mouse caudate putamen, FAAH-immunoreactive oligodendrocytes are present in fiber tracts (white matter) (Egertova et al. 2003). FAAH protein is also localized to the myelin sheath surrounding the unstained axons of large neurons (Egertova et al. 2003). This is consistent with the findings that FAAH mRNA is present in white matter of the rat brain (Thomas et al. 1997). However, the functional significance of FAAH expression in these non-neuronal cells is not yet understood. Moreover, although FAAH and CB₁ receptors are anatomically associated in many brain regions (Egertova et al. 1998), FAAH-expressing neurons are present in some brain areas such as thalamus and midbrain that express few or no CB₁ receptors (Egertova et al. 2003). Conversely, there is a population of striatal GABAergic MSNs where CB₁ receptors are present in their axonal terminals projecting to globus pallidus, entopeduncular nucleus, and substantia nigra, but FAAH-expressing neurons are absent (Egertova et al. 2003). In these situations, FAAH's biological role may be to degrade related acyl amides other than anandamide, for example, oleoylethanolamine and palmitoylethanolamine (Melis et al. 2008).

4.2 Striatum (Ventral, Accumbens)

4.2.1 Receptors

CB₁ receptors are expressed at low to moderate levels in the nucleus accumbens, with a pattern reminiscent of the striatum. CB₁ receptor protein is also localized on the terminals of the prefrontal glutamatergic efferents projecting to the nucleus accumbens as well as on the GABAergic axon terminals of accumbens MSNs and parvalbumin-positive interneurons (Robbe et al. 2001; Uchigashima et al. 2007). However, CB₁ receptors seem to be absent in the dopaminergic terminals projecting from the ventral tegmental area (VTA) to the accumbens. Therefore, cannabinoid stimulation of dopamine release in nucleus accumbens is likely mediated by inhibition of GABA release (either from within the nucleus accumbens or in the VTA) (Tanda et al. 1997; Szabo et al. 1999, 2002).

4.2.2 Synthetic Enzymes

DAGL α immunostaining, similar to that of mGluR5, is intense in spines and the perisynaptic region of the somatodendritic surface of striatal MSNs (Uchigashima et al. 2007).

4.2.3 Degradative Enzymes

FAAH immunoreactivity is absent in the nucleus accumbens (Egertova et al. 2003). However, moderate amounts of MAGL mRNA were found in the nucleus accumbens shell, islands of Calleja, and pontine nuclei by *in situ* hybridization (Dinh et al. 2002).

4.3 Striatum Medial (Lateral Septum, Septohippocampal, etc.)

4.3.1 Receptors

Moderate levels of CB₁ protein and mRNA are present in the basal forebrain including the medial and lateral septum and the nucleus of the diagonal band (Herkenham et al. 1991; Mailloux and Vanderhaeghen 1992; Matsuda et al. 1993; Marsicano and Lutz 1999). In situ hybridization studies showed the presence of CB₁ mRNA in cholinergic territories, especially the medial septum in mice. An immunocytochemical study revealed that a dense network of CB₁-positive fibers is present in the tenia tecta, ventral pallidum, and substantia innominate, whereas a fine network of CB₁-positive fibers is localized to the medial septum, diagonal bands, and nucleus basalis (Harkany et al. 2003). In the same study, no CB₁ immunoreactivity was detected in the cell bodies of basal forebrain cholinergic cells; instead these cells contain high levels of FAAH (Harkany et al. 2003). This is consistent with CB₁ receptors being synthesized and rapidly transported to axon terminals. Indeed, a later study showed that CB₁ protein is expressed in at least one-third of cholinergic neuron somata when axonal protein transport was blocked by cholchicine (Nyiri et al. 2005b). In rat medial septum, at least two types of cholinergic cells were identified, one with large somata, expressing CB₁ and GABA_B receptors and projecting to the hippocampus, whereas the other had smaller somata and lacked these two receptors (Nyiri et al. 2005b). Therefore, endocannabinoid signaling is implicated in the function of a population of septohippocampal cholinergic neurons, including cognition as well as generation of hippocampal theta rhythms.

4.3.2 Synthetic Enzymes

The immunostaining of NAPE-PLD is localized to the lateral nucleus of the septum (Egertova et al. 2008).

4.3.3 Degradative Enzymes

Neuronal FAAH-immunoreactivity has been detected in the lateral septum and the triangular septal nucleus (Egertova et al. 2003), as well as in the basal forebrain cholinergic neurons (Harkany et al. 2003).

4.4 Striatum Caudal (Striatum-like Amygdala Nuclei, Central Amygdala, Bed Nucleus, Medial Amygdala, Etc.)

4.4.1 Receptors

In contrast to the cortical component of the amygdala, the striatal component of amygdala (e.g., central and medial nuclei) displays much lower levels of CB₁ receptors. CB₁ mRNA in the striatal amygdala was revealed by sensitive in situ hybridization with the absence of signal in the same region of CB₁ knockout mice (Marsicano and Lutz 1999).

5 Cerebellum and Associated Nuclei

5.1 Cerebellar Cortex

5.1.1 Receptors

The patterns of CB₁ receptor expression in the cerebellum are striking. While autoradiographic and immunocytochemical studies showed intense labeling of CB₁ protein in the molecular layer, *in situ* hybridization studies yielded robust CB₁ mRNA in the granule cell layer (Matsuda et al. 1990; Herkenham et al. 1991; Glass et al. 1997; Tsou et al. 1998a; Egertova and Elphick 2000). Purkinje neurons are devoid of CB₁ protein labeling, while the axon terminals of basket cells surrounding the Purkinje cell axon initial segment display extremely strong CB₁ labeling. Putting this together, CB₁ receptors are mainly expressed in the axon terminals of the climbing fibers, parallel fibers, and (some) basket cells, suggesting a prominent presynaptic localization of CB₁ receptors, mediating modulatory effects of (endo)cannabinoids at glutamatergic and GABAergic inputs onto Purkinje neurons. Consistent with this pattern of expression, several elegant electrophysiological studies demonstrated a role for endocannabinoid inhibition of glutamatergic and GABAergic neurotransmission onto Purkinje neurons (Kreitzer and Regehr 2001; Maejima et al. 2001; Diana et al. 2002; Brenowitz and Regehr 2003). However, additional electrophysiological studies supported the functional somatic expression of CB₁ receptors (Kreitzer et al. 2002).

5.1.2 Synthetic Enzymes

NAPE-PLD mRNA has been found in the granule cell layer and Purkinje cells, but not in the molecular layer or white matter (Egertova et al. 2008). However, NAPE-PLD immunoreactivity is localized in the pre- and post-synaptic areas of the Purkinje neurons and in the somata of the basket cells in the molecular layer (Cristino et al. 2008; Suarez et al. 2008).

DAGL α immunoreactivity is highly concentrated at the base of the spine neck of cerebellar Purkinje cells. In contrast to its distribution in hippocampal pyramidal cells, DAGL α is excluded from the main body of spine neck and head (Yoshida et al. 2006). However, there are no DAGL α -immunoreactive neurons in the granular layer and in any subdivisions of the inferior olive, suggesting that DAGL α is not present in parallel and climbing fibers (Suarez et al. 2008).

High levels of DAGL β mRNA have been found in the mouse cerebellar granular layer by *in situ* hybridization (Yoshida et al. 2006). However, DAGL β immunostaining in the cerebellar cortex is less intense than that of DAGL α . DAGL β protein is localized to cell bodies of Purkinje neurons and in the molecular layer. In contrast to DAGL α , DAGL β -containing neuropil of the molecular layer probably represents parallel and climbing fibers from the granular cells and inferior olive neurons, respectively (Suarez et al. 2008).

5.1.3 Degradative Enzymes

FAAH immunoreactivity is present in the somata and dendrites of the Purkinje cells, which are innervated by CB₁-positive axon terminals (Egertova et al. 1998, 2003; Tsou et al. 1998b; Gulyas et al. 2004). Weak FAAH immunostaining is also evident in the somata of granule cells (Egertova et al. 2003), which is consistent with the detection of FAAH mRNA in rat cerebellar granule cells (Thomas et al. 1997).

Both Northern blot and in situ hybridization analyses revealed that MAGL mRNA is present in rat cerebellum (Dinh et al. 2002), whereas MAGL immunoreactivity is localized to the axon terminals in the molecular layer but absent in the FAAH-positive Purkinje cell dendrites in rat cerebellum (Gulyas et al. 2004). Interestingly, a recent immunocytochemical study found that MAGL is heterogeneously expressed in mouse cerebellum, with highest levels in parallel fiber terminals, weak levels in Bergman glia, and complete absence in other synaptic terminals (Tanimura et al. 2012). Even though the expression of MAGL is limited to a subset of nerve terminals and astrocytes in the cerebellum, MAGL still regulates 2-AG retrograde signaling broadly at parallel fiber or climbing fiber to Purkinje cell synapses (Zhong et al. 2011; Tanimura et al. 2012).

5.2 Deep Cerebellar Nuclei (Fastigial, Interpos, Dentate Nucleus)

5.2.1 Receptors

All cerebellar nuclei (medial, lateral, and interposed nuclei) contain very weak CB₁-immunoreactivity throughout the neuropil. However, intense CB₁ immunostaining was found in the neuropil of the dorsal part of the principal nucleus of the inferior olive (Suarez et al. 2008).

5.2.2 Synthetic Enzymes

Most cerebellar nuclei showed intense neuropil NAPE-PLD immunoreactivity and a number of moderately NAPE-PLD-labeled neurons. On the other hand, both the posterior parvicellular part of the interposed cerebellar and lateral cerebellar nuclei have considerably fewer NAPE-PLD-labelled neurons and a less intense neuropil immunoreactivity (Suarez et al. 2008).

DAGL α immunoreactivity is absent in cell bodies of cerebellar and vestibular nuclei and in other regions with mossy fiber projections in the granular layer such as the pontine nuclei or the spinal cord (Suarez et al. 2008). In contrast, DAGL β immunoreactivity is associated mainly with cell bodies embedded in a network of fibers. As with NAPE-PLD immunostaining, the posterior parvicellular parts of the interposed cerebellar and lateral cerebellar nuclei contain fewer DAGL β -positive neurons when compared with the remaining cerebellar nuclei (Suarez et al. 2008).

5.2.3 Degradative Enzymes

The strong FAAH immunoreactivity observed in all cerebellar nuclei is related mainly to the presence of a dense meshwork of fibers, consisting of FAAH-positive

punctate labeling that contain immunoreactive somata (Egertova et al. 2003; Suarez et al. 2008). These cerebellar nuclei are devoid of CB₁ immunoreactivity and receive synaptic input mainly from Purkinje cell axons. In white matter surrounding the cerebellar nuclei, FAAH-immunoreactive oligodendrocytes are conspicuous (Egertova et al. 2003).

Cerebellar and functionally related vestibular nuclei have numerous MAGL-immunoreactive neurons, showing a perikaryal and dendritic Golgi-like labeling, similar to that of DAGL β (Suarez et al. 2008).

6 Brainstem

6.1 Diencephalon

6.1.1 Thalamus (All Nuclei, Including Reticular Thalamic Nucleus, Habenula)

Receptors

CB₁ receptor expression is very low in most areas of the thalamus, with the exception of strong labeling in the lateral habenular nucleus, the anterior dorsal thalamic nucleus, and the reticular thalamic nucleus (Herkenham et al. 1991; Mailloux and Vanderhaeghen 1992; Tsou et al. 1998a; Marsicano and Lutz 1999). Since CB₁ mRNA is quite abundant in the lateral habenular nucleus, which has extremely diverse projections, it is likely that (endo)cannabinoids, acting through CB₁ receptors, significantly affect the diverse functions of the lateral habenula (Herkenham and Nauta 1977; Herkenham et al. 1991).

Synthetic Enzymes

NAPE-PLD mRNA is found in several thalamic nuclei such as lateral posterior nuclei and the medial geniculate nucleus, albeit at quite low intensity in mice (Egertova et al. 2008; Nyilas et al. 2008). In contrast, the highest levels of NAPE-PLD activity, protein and mRNA, were identified by enzyme assay, western blotting, and qPCR in rat thalamus, among nine different brain regions examined (Morishita et al. 2005). In addition, in situ hybridization revealed that moderate levels of DAGL α mRNA are expressed in mouse thalamus (Yoshida et al. 2006).

Degradative Enzymes

FAAH immunoreactivity has been detected in neuronal somata in the majority of thalamic nuclei, including the anterodorsal, the anteroventral, the anteromedial, the ventroanterior, the paratenial, the mediodorsal, the reticulothalamic, the ventrolateral, the ventroposterior, the ventromedial, the posterior, the lateral geniculate, and the medial geniculate (Egertova et al. 2000, 2003). Interestingly, FAAH protein is much more abundant in these nuclei than the cannabinoid CB₁ receptor, suggesting its role may be to degrade acyl amides other than anandamide in these regions.

An *in situ* hybridization study revealed that MAGL mRNA is abundantly expressed in the anterior thalamus, particularly in the anterodorsal nucleus, whereas it is sparse in other thalamic nuclei (Dinh et al. 2002).

6.1.2 Hypothalamus (All Nuclei)

Receptors

High levels of CB₁ immunoreactivity have been found in the arcuate, paraventricular, ventromedial, dorsomedial nuclei, and the external zone of the median eminence (Wittmann et al. 2007), as well as in the infundibular stem and lateral hypothalamic area (Tsou et al. 1998a). Further analysis revealed that CB₁ immunoreactivity is detectable in the preterminals of approximately equal numbers of symmetric and asymmetric synapses, suggesting the occurrence of retrograde signaling by endocannabinoids in both excitatory and inhibitory hypothalamic neuronal networks (Wittmann et al. 2007). An *in situ* hybridization study suggests that CB₁ mRNA is primarily present on glutamatergic neurons in the hypothalamus (Marsicano and Lutz 1999). Despite the relatively low levels of CB₁ receptors in the hypothalamus, functional GTP γ S assays revealed these receptors are more efficiently coupled to G proteins than in many other regions (Breivogel and Childers 1998). Finally, a recent study showed that mice with viral-mediated knockdown of the CB₁ receptor gene (~60 % decrease of the mRNA level) in the hypothalamus, while maintained on a normocaloric standard diet, displayed decreased body weight gain over time, subsequent to increased energy expenditure and elevated β_3 -adren-ergic receptor expression in brown adipose tissues (Cardinal et al. 2012). This result suggests that hypothalamic CB₁ receptor signaling plays an important role in energy expenditure under basal conditions, contributing to the antiobesity effect of CB₁ receptor antagonism.

Synthetic Enzymes

The staining of NAPE-PLD mRNA is evident in cells of the ventromedial nucleus (Egertova et al. 2008).

6.1.3 Mesencephalon (Colliculi, VTA, PAG, SN, Raphe)

Receptors

A. Substantia Nigra

Both autoradiographic and immunocytochemical studies showed extremely high levels of CB₁ receptor protein in the substantia nigra (SN) pars reticulata (Herkenham et al. 1991; Egertova and Elphick 2000). In contrast, *in situ* hybridization studies showed very low amounts of CB₁ mRNA in the SN (Matsuda et al. 1993), suggesting the high levels of CB₁ protein are restricted to incoming axonal projections from other brain regions. For example, CB₁ receptor protein is restricted to the GABAergic axonal terminals from the putamen MSNs and the glutamatergic terminals from the subthalamic nucleus, which may be involved in

the control of locomotility by CB₁ activation in the SN (Mailleux and Vanderhaeghen 1992; Sanudo-Pena and Walker 1997; Sanudo-Pena et al. 1999a). On the other hand, very low levels of CB₁ receptors exist in sparse intrinsic nigral neurons, which may exert a direct control on dopaminergic transmission (Matsuda et al. 1993; Julian et al. 2003). Finally, in rat striatal nerve terminals, a low but significant percentage of CB₁-immunoreactivity is co-localized with tyrosine hydroxylase (TH), a marker for both noradrenergic and dopaminergic terminals (Kofalvi et al. 2005).

B. Ventral Tegmental Area

Both cannabinoids and endocannabinoids modulate the primary rewarding effects of many abused drugs, including exogenous cannabinoids, *via* regulation of drug-induced increases in dopaminergic neural activity in the VTA (Maldonado et al. 2006). Therefore, it is of interest to elucidate the expression and function of CB₁ receptors in the VTA. A sensitive CB₁ polyclonal antibody (Fukudome et al. 2004) revealed a dense neuropil labeling of CB₁ receptors in the VTA (Matyas et al. 2008). The CB₁ immunoreactivity is restricted to presynaptic axon terminals of symmetric synapses, which may belong to local intrinsic GABAergic neurons (Matyas et al. 2008). Moreover, CB₁-immunoreactivity is co-localized with vesicular glutamate transporter in presynaptic terminals near dopamine neuron dendrites in the VAT, indicating the presence of CB₁ receptors in glutamatergic terminals (Kortleven et al. 2011). Interestingly, co-localization of CB₁ receptor and TH has been revealed in several brain areas including VTA, thereby pointing to a possible direct influence of CB₁ receptor activation on dopaminergic neurons (Wenger et al. 2003). However, further studies are required to clarify this possibility.

C. Periaqueductal Gray

Low to moderate levels of CB₁ receptors have been found in the midbrain periaqueductal gray (PAG), where the ECS is involved in the control of pain sensation, including stress-induced analgesia (Walker et al. 1999; Hohmann et al. 2005; Gregg et al. 2012). In contrast to opiate receptors on GABAergic aqueductal neurons, CB₁ receptors are preferentially, but not exclusively, localized in the dorsal portion of the PAG (Tsou et al. 1998a; Azad et al. 2001). In addition, moderate levels of CB₁ mRNA and protein have been found in the reticular formation and raphe nucleus, the latter being the main neuronal source of serotonin in the brain (Glass et al. 1997; Haring et al. 2007), which might have functional implications in emotion/mood modulation.

Synthetic Enzymes

A. Ventral Tegmental Area

Moderate to high levels of DAGL α are expressed in most neurons of the VTA (Matyas et al. 2008). High-resolution electron microscopy further revealed that DAGL α is accumulated in postsynaptic plasma membrane of glutamatergic and GABAergic synapses, of both TH-positive and negative dendrites. The finding that

DAGL α is present in postsynaptic dendrites juxtaposed to presynaptic CB₁ receptors suggests that 2-AG-CB₁-mediated retrograde synaptic signaling may modulate the drug-reward circuitry at multiple types of synapses in the VTA.

B. Periaqueductal Gray

DAGL α protein is co-localized with mGluR5 within the same dendritic spine heads at postsynaptic excitatory synapses in rat dorsolateral PAG, which is involved in 2-AG-mediated stress-induced analgesia (Gregg et al. 2012).

Degradative Enzymes

A. Substantia Nigra

FAAH-immunoreactive neurons are not evident in the pars reticulata of the SN, which contains a very high concentration of CB₁ immunoreactivity in both mouse and rat (Egertova et al. 2000, 2003).

B. Other Nuclei

FAAH-immunoreactive neurons are present in the superior and inferior colliculus, the rhabdoid nucleus, and several mesencephalic raphe nuclei and are also intensely stained in the mesencephalic trigeminal nucleus (Egertova et al. 2003). FAAH-immunoreactive oligodendrocytes associated with fiber tracts are abundant in the midbrain. For example, FAAH immunostaining was found to be localized in the myelin sheath surrounding the unstained axons of the mesencephalic trigeminal tract (Egertova et al. 2003).

6.2 Hindbrain

6.2.1 Medulla (Area Postrema, Cochlear Nuclei, Nucleus of the Solitary Tract, Trigeminal Nuclei, Various Other Cranial Nerve Nuclei)

Receptors

Expression of CB₁ receptors, in contrast to the opioid receptors, in the medullary respiratory control centers is relatively low (Herkenham et al. 1991; Glass et al. 1997). This likely explains the low mortality caused by cannabinoid intoxication in humans and animals. However, relatively high levels of CB₁ receptors are present in the medullary nuclei associated with emesis and vagal control of gut motility, which may underlie the inhibition of emesis and gastrointestinal motility by cannabinoids (Krowicki et al. 1999; Van Sickle et al. 2001, 2003). For example, high to moderate levels of CB₁ receptors were found in the area postrema, the dorsal motor nucleus of the vagus, as well as the medial subnucleus and the subnucleus gelatinosus of the nucleus of the solitary tract (Van Sickle et al. 2001, 2003; Mackie 2005; Storr and Sharkey 2007).

In mouse dorsal cochlear nucleus, CB₁ receptors are highly expressed in glutamatergic terminals of the parallel fibers, at intermediate levels in glycinergic terminals, and completely absent in the auditory nerve inputs innervating to the same DCN principle neurons—fusiform and cartwheel cells (Zhao et al. 2009). Therefore, CB₁ receptors are well positioned to mediate short- and long-term plasticities exhibited at parallel fiber synapses, but not at auditory nerve inputs (Zhao et al. 2011; Zhao and Tzounopoulos 2011).

Synthetic Enzymes

Both DAGL α and DAGL β proteins were identified in the somata of fusiform and cartwheel cells of mouse DCN, while they were present only in the dendritic spines of cartwheel cells (Zhao et al. 2009). These findings suggest that the synthesis of 2-AG is more distant from parallel fiber synapses in fusiform than cartwheel cells.

7 Spinal Cord (Dorsal, Ventral, Dorsal Root Ganglion)

7.1 Receptors

Intrathecal application of cannabinoids has been found to suppress pain in various pain models (Smith and Martin 1992; Welch et al. 1995), which is consistent with their suppression of noxious stimulus-evoked neuronal firing (Hohmann and Herkenham 1998) and Fos protein expression in the spinal dorsal horn (Hohmann et al. 1999b). Moreover, cannabinoids inhibited glutamate release from afferents in lamina I of dorsal horn in a CB₁ receptor-dependent fashion (Jennings et al. 2001; Morisset and Urban 2001). Compatible with these functional studies, moderate levels of CB₁ receptors were found in the superficial layers of the dorsal horn, the dorsolateral funiculus, and lamina X, all regions of the spinal cord associated with analgesia (Farquhar-Smith et al. 2000; Nyilas et al. 2009).

However, only a small percentage of CB₁ receptors are localized at central terminals of primary afferent C fibers, with many more present on large, myelinated A β and A δ fibers, as well as postsynaptic interneurons (Hohmann and Herkenham 1998, 1999; Hohmann et al. 1999a; Farquhar-Smith et al. 2000). A dorsal rhizotomy produced time-dependent 50 % losses in cannabinoid binding densities in the dorsal horn since rhizotomy destroyed the terminals of both small- and large-diameter fibers (Hohmann et al. 1999a). However, a quantitative autoradiographic study showed a modest (16 %) decrease in cannabinoid binding sites in the superficial dorsal horn by neonatal capsaicin-mediated destruction of sensory C fibers (Hohmann and Herkenham 1998). Moreover, another study showed little decrease in CB₁ receptor immunoreactivity following dorsal rhizotomy or hemisection of the spinal cord, suggesting CB₁ receptors are primarily expressed on interneurons (Farquhar-Smith et al. 2000). Similarly, CB₁ receptors are only minimally co-localized with markers for C primary afferents both in the superficial dorsal horn and dorsal root ganglion (DRG) (Farquhar-Smith et al. 2000; Bridges et al. 2003). The above data suggest that the majority of CB₁ receptors are not

localized at the presynaptic terminals of nociceptive primary afferents, but rather may exist on postsynaptic interneurons (Farquhar-Smith et al. 2000; Salio et al. 2002), a CB₁ distribution which differs from the strong presynaptic axon terminal localization of CB₁ receptors in most other brain regions (Katona et al. 1999; Nyiri et al. 2005a). Indeed, research showed that CB₁ receptors localized at dorsal horn inhibitory postsynaptic interneurons mediate C-fiber-induced pain sensitization. However, it is important to emphasize the unexpected pro-nociceptive role of endocannabinoids is specific for C-fiber-mediated activity-dependent hyperalgesia, in contrast to the anti-nociceptive effect of these endogenous lipid molecules in models of inflammatory and neuropathic pain (Pernia-Andrade et al. 2009). Moreover, a conditional nociceptor-specific loss of CB₁ was found to reduce spinal CB₁-specific ligand binding by approximately 20 % only and did not substantially decrease CB₁-immunoreactivity in spinal laminae I and II (Agarwal et al. 2007). However, this conditional knockout of CB₁ in peripheral nociceptive neurons led to a substantial reduction of analgesia produced by local and systemic delivery of cannabinoids, suggesting that low levels of CB₁ expression did not necessarily mean lack of functional significance (Agarwal et al. 2007). Therefore, it is likely that the interplay between cannabinoid actions on peripheral primary afferents, interneurons, and descending pathways collectively contributes to the analgesic effects of CB₁ receptor activation in the spinal cord.

Interestingly, despite the earlier reports of minimal localization of CB₁ receptors in nociceptive DRG neurons (Hohmann and Herkenham 1999; Bridges et al. 2003), more recent studies suggest a much broader (40–80 %) distribution of CB₁ receptors in nociceptive neurons of DRG and trigeminal ganglia (Mitrirattanakul et al. 2006; Agarwal et al. 2007). While the possible reasons for these discrepancies have been well discussed elsewhere (Marsicano and Kuner 2008), it is also important to note that the expression and peripheral transport of CB₁ receptors in the DRG can be upregulated by peripheral inflammation (Amaya et al. 2006). Finally, some immunocytochemical evidence suggests CB₁ receptors are also present in the ventral horn (Tsou et al. 1998a; Sanudo-Pena et al. 1999b), a spinal cord area associated with movement.

7.2 Synthetic Enzymes

DAGL α mRNA is widely expressed in spinal dorsal horn neurons (Nyilas et al. 2009). Similar to CB₁ receptors, high levels DAGL α protein have been found to be localized at the superficial dorsal horn. High-resolution electron microscopy demonstrated a postsynaptic localization of DAGL α at nociceptive synapses, which is juxtaposed to the presynaptic CB₁-containing excitatory primary afferents (Nyilas et al. 2009) (Fig. 1B, C). Interestingly, postsynaptic DAGL α is co-localized with mGluR5, whose activation induces 2-AG biosynthesis (Nyilas et al. 2009).

7.3 Degradative Enzymes

FAAH has been found in the cell bodies of ventral horn neurons (Tsou et al. 1998b). The presence of FAAH and CB₁ receptors in circuits involved in spinal reflexes may underlie the antispastic effects of (endo)cannabinoids.

8 Summary, Concluding Thoughts, and Future Directions

As this brief review has shown, the components of the endocannabinoid system (ECS) are widespread throughout the CNS. Endocannabinoids have a major role as retrograde transmitters in many brain regions, although often with region-specific specialization as discussed above. Thus, in most brain regions, the highest levels of CB₁ receptors are found presynaptically, while endocannabinoid synthesizing enzymes are present postsynaptically. Interestingly, some endocannabinoid degrading enzymes are found presynaptically (e.g., MAGL) and others postsynaptically (e.g., ABHD6 and FAAH), suggesting specialized roles of each in endocannabinoid degradation (Gulyas et al. 2004; Blankman et al. 2007), and introducing an extra layer of complexity in endocannabinoid metabolism. A variation on the presynaptic localization of CB₁ is the somatic expression of CB₁, most commonly in some cortical and cerebellar neurons, where endocannabinoid signaling is cell autonomous. Future studies are needed to better define the distribution of some less-well studied (putative) ECS components, including α/β -hydrolase domain 4 (ABHD4) (Simon and Cravatt 2006), glycerophosphodiesterase 1 (GDE1) (Simon and Cravatt 2006, 2010), ABHD6 (Blankman et al. 2007; Marrs et al. 2010), α/β -hydrolase domain 12 (ABHD12) (Blankman et al. 2007), cannabinoid receptor interacting protein 1a (CRIP1a) (Niehaus et al. 2007), and DAGL β (Gao et al. 2010).

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