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Unveiling the role of adenosine A_{2A} receptors on neurogenesis induced by activation of cannabinoid receptors in rat SVZ and DG stem/progenitor cell cultures

Dissertação apresentada à Universidade de Coimbra para cumprimento dos requisitos necessários à obtenção do grau de Mestre em Biologia Celular e Molecular, realizada sob a orientação científica da Doutora Sara Xapelli (Instituto de Medicina Molecular, Faculdade de Medicina da Universidade de Lisboa) e do Professor Doutor Carlos Duarte (Departamento de Ciências da Vida, Faculdade de Ciências e Tecnologia, Universidade de Coimbra)

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“Ever tried. Ever failed. No matter. Try again. Fail again. Fail better.”

(Samuel Beckett)

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Abbreviations

Δ9-THC Δ9-tetrahydrocannabinol

2-AG 2-arachidonoylglycerol

AEA Arachidonylethanolamide

ARs Adenosine receptors

A₁Rs, A_{2A}Rs, A_{2B}Rs, A₃Rs Adenosine A₁, A_{2A}, A_{2B} and A₃ receptors

BDNF Brain-derived neurotrophic factor

CA3 Cornu Ammonis 3 area

CBRs Cannabinoid receptors

CB₁Rs, CB₂Rs Cannabinoid receptors type 1 and type 2

CNS Central nervous system

CSF Cerebrospinal fluid

DG Dentate gyrus

EC Entorhinal cortex

ECS Endocannabinoid system

EGF Epidermal growth factor

FGF-2 Fibroblast growth factor-2

GABA Gamma-aminobutyric acid

GCs Granule cells

GCL Granular cell layer

GDNF Glial-derived neurotrophic factor

GL Glomerular layer

GPCRs G protein-coupled receptors

HPA Hypothalamic-pituitary-adrenal axis

IGCs Immature granule cells

IPCs Intermediate progenitor cells

LV Lateral ventricle

MAPKs Mitogen-activated protein kinases

NPCs Neural progenitor cells

NSPCs Neural stem/progenitor cells

OB Olfactory bulb

PLC Phospholipase C

RMS Rostral migratory stream

SVZ Subventricular zone

SGZ Subgranular zone

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Abstract

Over the past years, the idea that neurogenesis, the process by which new functional neurons are produced, does not occur in the adult matured brain has considerably changed. In fact, neurogenesis actively occurs mainly in two distinct neurogenic niches, the subventricular zone (SVZ), along the lateral walls of the lateral ventricles, and the subgranular zone (SGZ), in the dentate gyrus (DG) of the hippocampus, where newly formed neurons, derived from neural stem/progenitor cells (NSPCs), have a role in olfaction/odor discrimination and in learning/memory, respectively.

Adenosine, which is a widespread neuromodulator in the brain, greatly influences synaptic neurotransmission and is involved in many physiological and neuropathological processes. Moreover, the endocannabinoid system, which activates primarily type 1 and 2 cannabinoid receptors (CB₁R and CB₂R), is a key regulator of synaptic function, especially by inhibiting neurotransmitter release. Furthermore, it has been shown that CB₁R and CB₂R activation modulates neurogenesis by promoting neural stem cell proliferation, differentiation and maturation. Importantly, an interaction between A_{2A} receptors (A_{2A}Rs) and CB₁Rs was reported and shown to allow the control of key modulatory effects on neuronal function and transmission. However, to date, no study has directly evaluated the effects of the crosstalk between A_{2A}Rs and CBRs on neurogenesis.

Therefore, in this work, the putative role of A_{2A}Rs on neurogenesis induced by the activation of CB₁Rs and CB₂Rs was investigated by looking at different stages of the neurogenic process.

Results show that CB₁R, CB₂R or A_{2A}R activation did not change the SVZ cell-fate. Moreover, activation of CB₂R or A_{2A}R did not induce SVZ cell proliferation. However, CB₁R activation promoted cell proliferation, an effect that was blocked by an A_{2A}R selective antagonist. Interestingly, activation of either CB₁R or CB₂R promoted an increase in SVZ neuronal differentiation that is impaired by A_{2A}R blockade, although A_{2A}R activation *per se* had no effect on neuronal differentiation.

Concerning the DG neurogenic niche it was observed that while CB₁R activation had no effect on the cell fate of DG cells, activation of CB₂Rs or A_{2A}Rs promoted self-renewing divisions. Furthermore, A_{2A}R selective antagonist blocked the effect mediated by activation of CB₂Rs in the self-renewal of DG cells. Moreover, A_{2A}R-mediated effect on self-renewal was blocked either by CB₁R or CB₂R selective antagonists. Moreover, although activation of CB₁R, CB₂R or A_{2A}R *per se* did not induce DG cell proliferation, an increase in the number of proliferating cells was observed upon CB₁R or CB₂R co-activation with A_{2A}Rs. Lastly, CB₁R, CB₂R and A_{2A}R activation promoted DG neuronal differentiation. Interestingly, the effect

mediated by CB₁Rs or CB₂Rs was blocked by an A_{2A}R selective antagonist, while the effect mediated by A_{2A}Rs was impaired in the presence of CB₁R or CB₂R selective antagonists.

Taken together, these results suggest that a possible crosstalk between the adenosinergic and endocannabinoid systems may exist, either by a structural interaction (formation of heterodimers) or by crosstalk at downstream signaling, that ultimately contributes to the control of neurogenesis.

Keywords: Neurogenesis, Neurogenic niches, Adenosine A_{2A} receptors, Cannabinoid receptors

Resumo

A ideia de que neurogênese, o processo pelo qual são produzidos novos neurónios funcionais, não ocorre no cérebro adulto foi alterada consideravelmente ao longo dos anos. Efetivamente, a neurogênese ocorre em dois nichos neurogênicos distintos, a zona subventricular (SVZ), que percorre a parede lateral dos ventrículos laterais, e a zona subgranular (SGZ, no giro dentado (DG) do hipocampo, onde neurónios recém-formados, descendentes de células estaminais/progenitoras neurais, apresentam papéis no olfato/discriminação de odores e em aprendizagem/memória, respetivamente.

A adenosina, um neuro-modulador amplamente presente no cérebro, tem um papel preponderante na transmissão sináptica e está envolvida na regulação de muitos processos fisiológicos e neuropatológicos. O sistema endocanabinóide, onde são principalmente ativados os recetores de canabinóides do tipo 1 e 2 (CB₁R e CB₂R), é um regulador-chave da função sináptica, especialmente pela inibição da libertação de neurotransmissores. Além disso, processos essenciais na neurogênese como proliferação, diferenciação e maturação neuronal são modulados pela ativação de recetores de canabinóides. Importaneamente, foi demonstrada a interação entre recetores A_{2A} de adenosina (A_{2A}Rs) e CB₁Rs que é especialmente importante no controlo da função neuronal e transmissão sináptica. Contudo, até à data, nenhum estudo avaliou diretamente os efeitos desta possível interação entre os recetores A_{2A} de adenosina e recetores de canabinóides na neurogênese.

Assim, neste trabalho, foi investigado o possível papel dos A_{2A}Rs na neurogênese induzida pela ativação dos recetores CB₁Rs e CB₂Rs, olhando para diferentes estádios do processo neurogénico.

Os resultados mostram que ativação dos A_{2A}Rs, CB₁Rs ou CB₂Rs não altera a divisão celular no sentido de promover um aumento ou diminuição da capacidade de auto-renovação na SVZ. Além disso, ativação dos A_{2A}Rs e CB₂Rs não promove proliferação das células SVZ. Contudo, a ativação dos CB₁Rs promove um aumento na proliferação das células SVZ, um efeito bloqueado pela presença de um antagonista dos A_{2A}Rs. Interessantemente, a ativação de ambos CB₁Rs ou CB₂R induz um aumento na diferenciação neuronal das células SVZ e este efeito é comprometido pelo bloqueio dos A_{2A}Rs, apesar de *per se* a ativação dos A_{2A}Rs não ter nenhum efeito na diferenciação neuronal.

Em relação ao nicho neurogénico DG, foi observado que, apesar da ativação dos CB₁Rs não promover nenhum efeito no destino celular das células derivadas do DG, ativação dos CB₂R e A_{2A}Rs promove a perpetuação do *pool* estaminal das células de DG. O efeito promovido pelos CB₂R é bloqueado pela presença de antagonista seletivo de A_{2A}Rs e semelhantemente, o efeito promovido pelos A_{2A}Rs é bloqueado pela presença de antagonistas seletivos para CB₁R ou CB₂R. Adicionalmente, apesar da ativação dos CB₁Rs,

CB₂Rs ou A_{2A}Rs *per se* não produzem qualquer efeito na proliferação das células DG, um aumento no número de células em proliferação é visto quando há co-ativação de CB₁Rs ou CB₂Rs com A_{2A}Rs. Por fim, a ativação dos CB₁Rs, CB₂Rs e A_{2A}Rs promove um aumento na diferenciação neuronal das células DG. Interessantemente, o efeito promovido pelos CB₁R ou CB₂R é bloqueado pela presença de um antagonista seletivo dos A_{2A}Rs assim como o efeito mediado pelos A_{2A}Rs é bloqueado pela presença de antagonistas seletivos para CB₁R ou CB₂R.

Em resumo, os resultados levam a acreditar que possa existir uma interação entre os sistemas adenosinérgico e endocanabinoide, provavelmente a nível estrutural (com a formação de heterómeros) ou interação ao nível das vias de sinalização, o que contribuirá em última instância para o controlo da neurogénese.

Palavras-chave: Neurogénese, Nichos neurogénicos, Recetores A_{2A} para adenosina, Recetores para canabinóides

CHAPTER 1

Introduction & Aims

1.1. The nervous system & brain, neurogenesis, adenosine and cannabinoids

In this thesis four different subjects of high complexity will be highlighted: (a) neurogenesis, which is an intricate process occurring throughout adulthood in two main neurogenic niches; (b) adenosine, which is ubiquitously present in all cells and is a neuromodulator enrolled in several physiological functions, with receptors distributed throughout all brain areas; (c) the endocannabinoid system, composed of cannabinoid receptors, endogenous ligands and proteins responsible for synthesis, reuptake and degradation, which inhibits neurotransmission and plays an important role in synaptic function and many physiological functions and, finally, (d) the putative crosstalk between adenosine A_{2A} receptors and cannabinoid receptors.

1.2. Neurogenesis

Neurogenesis, generally defined as the process by which new functional neurons are generated from neural stem/progenitor cells was first believed to be a process exclusively occurring during pre-natal development¹⁻³. The adult mature brain was supposedly incapable of producing new neurons. This dogmatic view began to change in the 60's because of Smart's and Altman's pioneering studies with [³H]-thymidine autoradiography⁴⁻⁶. They showed, for the first time, evidence for the generation of new dentate granule neurons in postnatal rat hippocampus. Subsequently, the first reports suggesting the existence of postnatal neurogenesis occurred in the late 70's and 80's^{2,7}. Later, technical advances revolutionized this field of research with the introduction of bromodeoxyuridine (BrdU), a synthetic thymidine analogue used as an S-phase marker of the cell cycle. This labelling technique allowed, along with other immunocytochemical and histological methods, the demonstration and validation that a great portion of newly generated cells in the adult brain were indeed neurons^{2,8-11}. Therefore, neural stem/progenitor cells are capable of dividing themselves and then differentiating into newborn neurons that can migrate to pre-existing circuitries¹⁰. Rapid and significant progress in the field has led to the overall acceptance of neurogenesis being one of the many phenomena occurring in the adult brain that contributes to brain plasticity, synaptic function and circuit dynamics^{10,12}.

Neurogenesis also occurs in the complex and highly evolved human brain, where new neurons continue to be added in a series of highly regulated and coordinated physiological mechanisms and by activity^{9,13,14}. However, this postnatal addition of newborn neurons slowly decreases with age^{15,16}. Importantly, this continual addition of neurons over a lifetime implies substantial structural changes and constant remodeling of brain circuits, which implicates a

tremendous impact in central nervous system (CNS) functioning, including, for instance, short- and long-term learning and memory^{10,12}.

In fact, neurogenesis is a wide spread phenomenon across many mammalian species (including rodents, rabbits, monkeys and humans)^{2,17,18} with a high degree of evolutionary conservation, suggesting that it is a fundamental biological mechanism – neurons generated and maintained in these specific regions during adulthood contribute to normal brain function and plasticity^{11,16,18–20}. This functional contribution is not just a restorative mechanism but also a way to respond to constant and dynamic challenges applied by environmental stimuli or internal variations, representing a unique feature worth the exceptional effort of brain continuously reshaping itself¹⁴.

Neural stem/progenitor cells (NSPCs) are present in the adult nervous system and have the ability to self-renew their own pool through cell proliferation and/or to generate cells from the neural lineage (neurons, astrocytes and oligodendrocytes)^{21,22}. Moreover, NSPCs pass through sequential developmental stages that show structurally and functionally distinct cellular properties and dynamics in order to generate the newborn neurons^{9,23}. Specifically, at the level of spatial organization, the presence or absence of specific cell constituents seems to be crucial for the self-renewal of NSPCs and neuronal differentiation¹³.

In the adult brain, the neurogenic process occurs, mainly in two restricted areas, the subventricular zone (SVZ) lining the lateral ventricles and the subgranular zone (SGZ) in the dentate gyrus (DG) of the hippocampus. NSPCs are abundant in these areas which have high rates of proliferation physiologically^{14,24–27}. New neurons generated at these sites then migrate toward their final destinations, where they differentiate into mature neurons and are integrated into the neuronal circuitry^{20,28,29}.

Adult neurogenesis is affected by several factors, such as physical activity, environmental enrichment, stress and aging and by various pathological conditions^{30,31}. Extrinsic and intrinsic elements largely regulate and influence the rate of proliferation, maturation, survival and integration of newborn neurons, including niche factors/receptors, cytoplasmic factors, transcriptional factors, and epigenetic regulators. Predominantly, neurotransmitters, neural peptides, adhesion molecules, growth factors, neurotrophins, cytokines, cell-cycle regulators, transcription factors and other key components are the major players involved in the regulation of adult neurogenesis^{2,22,25–27}. Epigenetically, many mechanisms perform diverse roles in regulating adult neurogenesis, such as DNA methylation, histone modifications, chromatin remodeling, and non-coding RNAs^{23,25,32,33}. In response to injury or pathological stimuli, such as ischemic stroke, neurodegenerative diseases (Parkinson's, Huntington's, Alzheimer's diseases) and brain trauma, the newborn neurons that are generated following the loss of neurons, migrate to atypical or lesioned areas (reactive neurogenesis)^{2,9,20,24,27,29}.

1.2.1. Neurogenic niches

A neurogenic niche is, by definition, a stem cell rich microenvironment that anatomically and functionally controls their development *in vivo*. Additionally, these restricted areas of the mammalian brain have unique structural architectures that allow specific properties at the molecular, cellular and circuitry levels thus contributing to constitutive generation of new functional neurons throughout life^{9,25,27}.

The major cellular components of adult neurogenic niches are NSPCs, endothelial cells, astrocytes, microglia, mature neurons, and intermediate neural precursors. They play an important role regulating proliferation and fate specification of adult neural precursors as well as in neuronal migration, maturation, targeting and synaptic integration^{9,25,26,30}.

In fact, SVZ and SGZ are regions rich in NSPCs that originate neuroblasts that migrate and reach their final target where they mature and integrate the preexisting neuronal network²⁹.

Moreover, recent studies have shown the generation of matured neuronal cells in other non-traditional neurogenic brain regions³⁴. In fact, several studies reported the occurrence of adult neurogenesis in the striatum, neocortex, piriform cortex (part of the olfactory cortex) and within the limbic system, primarily at the amygdala and hypothalamus¹⁷. Particularly in adult humans, the striatum is a well-developed structure in which there is a higher expression of genes associated with neuronal migration as compared to other brain regions^{15,18}. This neurogenesis in non-traditional regions is predominantly enhanced after certain physiological or pathological conditions^{35,36}.

1.2.1.1. Subventricular zone (SVZ)

SVZ is a thin cell layer located alongside the lateral walls of the lateral ventricles (LV) of the adult brain²⁰. This region exhibits a highly organized cytoarchitecture^{9,37}. The SVZ high density of NSPCs present in the dorsolateral and rostral sides of the ventricles is correlated with the rostrocaudal gradient of proliferative activity^{30,38}.

SVZ is, predominantly, constituted of four cell types, different in their morphological nature and ultrastructure (Figure 1)^{20,30,37,39–41}:

- Type E cells: nondividing ependymal cells, which are multiciliated cells facing the lumen of the ventricle and that are in close contact with the cerebrospinal fluid (CSF);

- Type B cells: astrocytic-like cells, which constitute the neural stem foundation and are responsible for the dividing/proliferative feature of SVZ; these astrocyte-like type B cells can be subdivided into two subtypes based on differences in their location and morphology:

 - type B1 cells are generally closely associated with ependymal cells;

-type B2 cells are more frequently located close to the underlying striatal parenchyma;

Type B cells can also be grouped into 3 main domains according to their position in the niche and the surrounding cells (proximal, intermediate and distal)^{28,42};

-Type C cells: transit-amplifying cells which are the immediate progeny of B cells, characterized by being highly proliferative intermediate progenitor cells (IPCs) that usually are close to blood vessels and form clusters interspaced among chains throughout the SVZ;

-Type A cells: young migrating neurons, called neuroblasts, that migrate through sheathed chains of astrocytes towards the olfactory bulb.

In summary, the type B1 astrocyte-like cells of radial glia origin (with unique functional characteristics between those of astrocytes and radial glia) are a quiescent stem cell-like subpopulation⁴³. B1 cells divide at a continuous and slow rate and are responsible for the long-term maintenance of the stemness/multipotency state of the niche. These cells generate a different pool composed of rapidly proliferating progenitor cells – type C transit amplifying cells –, committed to the neuronal lineage. These type C cells will, in turn, rapidly divide and generate type A cells^{20,30,38,44}.

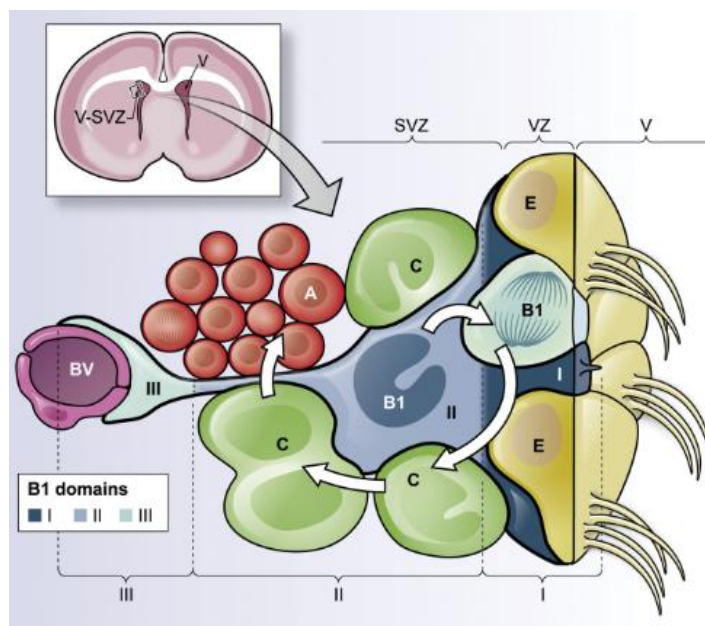


Figure 1 – Cellular organization of the SVZ niche. Cross-section showing the location of the SVZ niche; enlarged representative scheme of SVZ niche and how the process of neurogenesis occurs: NSPCs (type B1 cells, blue), which are surrounded by ependymal cells (type E cells, yellow), give rise to transit amplifying cells (type C cells, green), that divide and generate neuroblasts (type A cells, red); B1 cells domains are also represented as I, II and III according to their position in the niche (adapted from ²⁸Fuentealba et al, 2012).

Therefore, the process of adult SVZ neurogenesis can be simplified into the following linear, but multifaceted, sequence of events: 1) glial fibrillary acidic protein (GFAP)-expressing B-type astrocytes function as NSPCs and 2) produce transiently amplifying progenitors (type C cells) that 3) differentiate into neuroblasts (type A cells), which then 4) migrate anteriorly to the olfactory bulb (OB) via the rostral migratory stream (RMS)^{25,28,45,46}. In the RMS, neuroblasts generated in the SVZ migrate tangentially to the OB by forming elongated cell aggregates, known as “chains”, surrounded by a sheath of astrocytes called the glial tube²⁰. Once reaching the core of the olfactory bulb, immature neurons detach from the RMS and migrate radially along blood vessels toward glomeruli where they differentiate into different subtypes of olfactory interneurons²⁷. This terminal differentiation occurs in different layers of the OB, namely the granular cell layer (GCL), located in the deeper layer of the OB, and glomerular layer (GL), located in the most superficial layer. In these layers the neuroblasts will differentiate into two types of olfactory neurons, the granular cells (the majority) and the periglomerular cells (a small percentage)⁴⁷. Given the fact that the majority of interneurons are either gamma-aminobutyric acid (GABA)ergic or dopaminergic, the newly born granular and periglomerular neurons are synaptically integrated into the existing circuitry, exhibiting initially GABA and then glutamate receptors becoming sensitive to stimuli from the olfactory nerve layer of the OB^{26,48}.

As previously mentioned numerous factors are involved in the regulation of the neurogenic process. In fact, in the SVZ molecules mediating cell-to-cell and cell-to-substrate interactions, chemoattraction and chemorepulsion mechanisms, extracellular matrix remodeling and other key elements modulate neurogenesis^{25,28,30,40}. These factors are implicated in the control of different stages of cell development, namely in the maintenance, proliferation, differentiation, migration, survival, maturation and integration of functional new neurons in the given circuitry^{28,47}. Specifically, these factors include context-dependent release of neurotransmitters, neurotrophic/growth factors, morphogens, transcription factors and other extrinsic factors, like epigenetic regulators derived from sensory, motor or social stimuli (extensively reviewed in Lledo et al., 2006)^{26,27,37,40}.

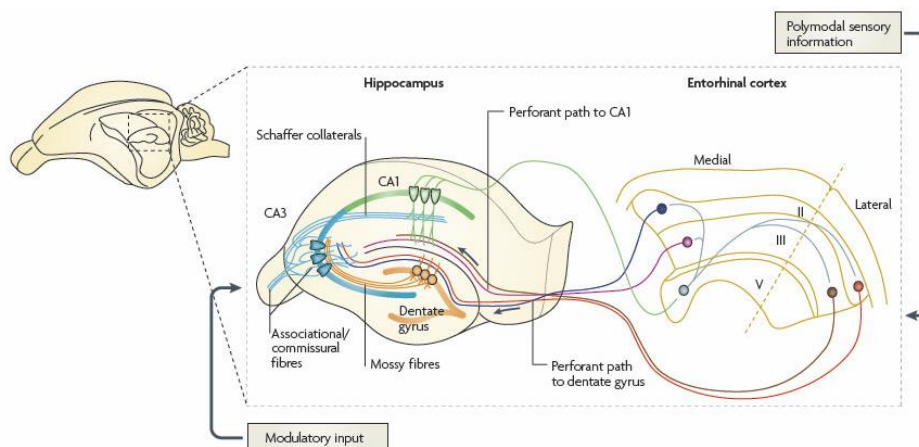
Therefore, the extracellular matrix, endothelial cells and vascular contacts associated with the adult SVZ, have also a direct impact on adult neurogenesis by constituting potential sources of signals regulating progenitor cell behavior and proliferative patterns²⁶. The SVZ area is extensively established on extracellular matrix which provides a platform for the presentation of components that affect proliferation and progenitor activity within the neurogenic niche^{9,37}. Notably, blood vessels-released factors and endothelial-derived factors may act as neurogenic signals that interact with NSPCs and stimulate stem cell self-renewal and transit-amplifying cells generation and proliferation^{26,37,47}. Furthermore, blood vessels may also be important to serve as a scaffold for neuronal migration and neuronal integration³⁷ and ependymal-derived factors can provide instructive cues to sustain new neuronal production

and to redirect niche responses to locally reintegrate neurons in damaged tissues (whenever in situations of brain injury or pathological conditions)^{46,49}.

Taking into consideration that the OB is the first processing center of odor information, the addition of new interneurons into this region is likely responsible for the plasticity of the olfactory system, which is especially important for rodents²⁰. The real functional significance is still unclear and relatively debatable but evidences point out that survival and proper synaptic integration of these newly-born neurons into the OB is highly correlated with odor acquisition and discrimination and short- and long-term odor memory^{3,26,39,47}. Additionally, SVZ neurogenesis may contribute to the regulation of pheromone-related events, such as mating and social recognition²⁵. Recent studies also suggest that SVZ-derived NSPCs may have non-neurogenic functions, such as regulating immune trafficking and maintaining CNS homeostasis^{42,50,51}.

1.2.1.2. Subgranular zone (SGZ)

The hippocampus, the brain region associated with learning and the memory formation, is composed of 3 distinct regions, the DG, the Cornu Ammonis 3 area (CA3) and the CA1 area, which form the trisynaptic hippocampal circuit (or loop)^{19,52,53}. The degree of complexity within the hippocampus is enormous but, to simplify the trisynaptic loop, this circuit can be fairly comprehended as: existence of synapses between the entorhinal cortex (EC) and the dendrites of the granule cells of the DG (1), between the mossy fibers (the axons of the granule cells) and the pyramidal neurons of CA3 (2) and between the axons of the CA3 pyramidal neurons and the pyramidal neurons of CA1 (3), which then again project out to the subiculum and the EC (Figure 2)^{19,54–56}. SGZ is located at the interface between the granule cell layer (GCL) and the hilar layer of the hippocampal DG^{20,30}. Unlike SVZ, the SGZ neurogenic niche is structurally less recognizable and NSPCs are not in direct contact with CSF^{9,28}.

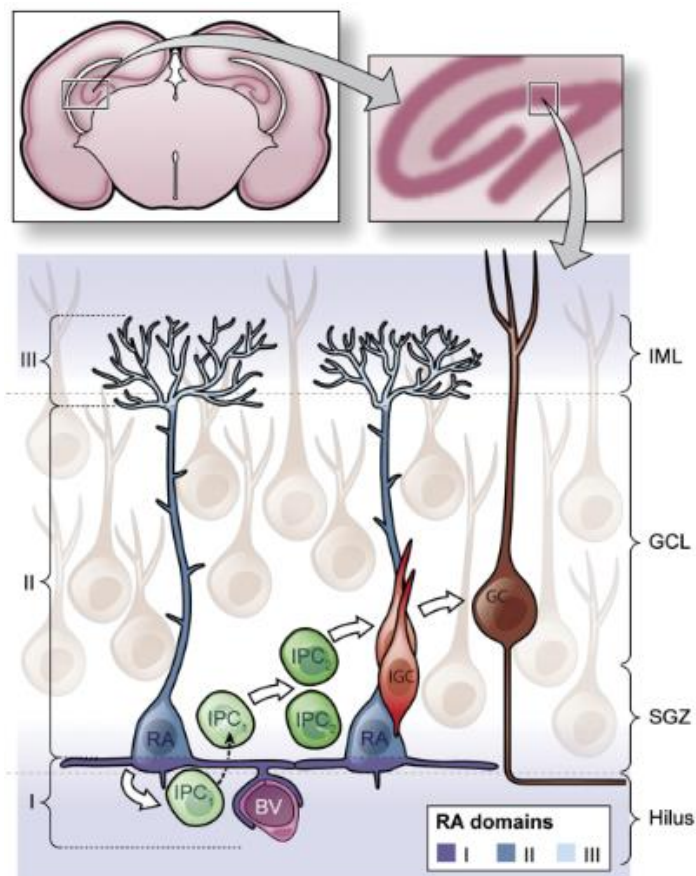


◀ **Figure 2 – Neuroanatomy of the hippocampus.** The trisynaptic arrangement of the hippocampus is due to the existence of 3 distinct zones: CA1, CA3 and DG. Excitatory projections connect the entorhinal cortex with the granule cells of the DG through the perforant path, which in turn connects, through mossy fibers, with the pyramidal cells of the CA3 area. The CA3 area connects through Schaffer collaterals with the CA1 area. Entorhinal cortex also connects with the CA1 area (adapted from ⁵⁶Neves et al., 2008).

The SGZ niche is mainly represented by four different (regarding morphology and antigen properties) types of cells (Figure 3)^{22,26,28,42,57,58}:

-Type I cells, radial astrocytes, which are predominantly present and highly polarized cells with astrocytic properties and contribute to maintain niche stemness. These astrocytes, similarly to SVZ type B cells, can also be grouped into 3 main domains according to their position and neighboring cells (proximal, intermediate and distal)²⁸;

-Type II cells, intermediate progenitor cells (IPCs), which lack glial features and are thought to be the most active cells in terms of proliferation, and by giving rise to progenitor cells can be subdivided into “type-II a” or “type-II b” cells, according to the presence or absence of doublecortin, respectively;



◀ **Figure 3 – Cellular organization of the SGZ niche.** Cross-section showing the location of the SGZ niche in the hippocampus; enlarged representative scheme of SGZ niche and how the process of neurogenesis occurs: NSPCs, represented as radial astrocytes (RA, type I cells, blue), give rise to transit amplifying cells which are intermediate progenitor cells (IPCs, type II cells, green), that divide asymmetrically and generate neuroblasts, immature granule cells (IGCs, type III cells, red); RA domains are also represented as I, II and III according to their position in the niche (adapted from ²⁸Fuentealba et al, 2012).

-Type III cells, immature granule cells (IGCs) or neuroblasts, which are immediate precursors of granule cells and can be subdivided into immature (D_1) or more differentiated neuroblasts (D_2);

-Mature granule cells (GCs), which express specific neuronal markers and are functionally integrated into the circuit.

In summary, the process of adult SGZ neurogenesis can be simplified into the following sequence of events: 1) type I radial astrocytes function as NSPCs and 2) generate type II intermediate progenitor cells that progressively 3) differentiate into type III immature granule cells (IGCs), which then 4) migrate a short distance to the granule cell layer (GCL)^{20,25,27,59}. This migration to the GCL promotes the local maturation of newly generated immature cells and integration into the neural network as functional granule neurons of the DG. Compared to mature granule cells, newborn neurons exhibit hyperexcitability and enhanced synaptic plasticity during specific developmental stages. As they differentiate and progressively become more specialized granule cells, they develop axonal projections to CA3 area and the dendritic arborization becomes increasingly more complex and extends deeper into the molecular cell layer^{2,20,22,27}. After a prolonged maturation phase, adult-born neurons exhibit similar basic electrophysiological properties as mature neurons, such as firing behavior and the kinetics of excitatory and inhibitory inputs^{25,54,59}. Unlike SVZ, where NSPCs give rise to two distinct interneuron lineages in the OB, adult NSPCs in the SGZ of the hippocampus predominantly give rise to functional glutamatergic excitatory neurons that integrate DG circuitry^{28,60}.

Nonetheless, similarly to SVZ, the SGZ microenvironment is intimately associated with other cells, namely endothelial cells, glial cells and other neuronal types, that structurally and biochemically support and regulate the niche and contribute significantly to adult neurogenesis²⁸. Particularly, the vasculature, especially capillaries, plays an important role in regulating the proliferation of adult NSPCs in SGZ⁹.

Diverse niche components, signaling pathways and external stimuli differentially modulate NSPCs behavior and the course of adult hippocampal neurogenesis. These factors are involved in the control of stem cell behavior, cell fate determination, maintenance of

stemness, as well as the regulation of all different stages of the neurogenic process (differentiation, maturation, migration and integration)^{27,28,40}. They include hormones, neurotransmitters, growth factors, transcription factors and other activity/plasticity-related players in response to environmental stimuli (extensively reviewed in Lledo et al., 2006)^{25-27,40,61}.

In addition, the hippocampus, specifically the SGZ of the DG, is targeted by many inputs and neurotransmitter systems⁴⁸ which causes SGZ neurogenesis to be particularly sensitive to the surrounding neuronal activity. In fact, neurons at the neurogenic site may provide spatiotemporal regulation of adult neurogenesis in response to hippocampal-dependent neuronal activity⁹. SGZ neurogenesis is also known to be influenced by neuropathological states, such as epilepsy, where newborn neurons are incorrectly reorganized into aberrant synapses/network connections^{2,25}.

Importantly, adult hippocampal neurogenesis may contribute to a diverse number of functions. It is believed that immature granule cells, by being electrophysiologically more excitable, have more potential to modulate activity-dependent plasticity of the DG circuit processing^{55,59,62}. This continuous rewiring of the brain is particularly important in cognition, learning and memory formation^{26,55,63}. The addition of new neurons in the adult SGZ is critical during specific time windows and occurs to reorganize the circuit networks in order to process information in hippocampus-dependent learning activities, contributing to memory formation^{9,17}. Specifically, tasks like spatial-navigation learning, long-term spatial memory retention, and spatial pattern discrimination may be dynamically handled by the addition of new neurons to the hippocampal circuitry^{25,52,63}.

Stage specific markers can be used to understand how the entire neurogenic process is displayed. Developmental stages like cell fate, differentiation and neuronal maturation can be precisely detected because newly born neurons express a series of transient markers, such as GFAP, Sex determining region Y-box 2 (Sox2), Nestin, Doublecortin (DCX) and Neuronal nuclei (NeuN). Developmental stages are likely to reflect a continuum rather than discrete steps. Therefore, the entire neurodevelopmental progression of neurogenesis, from precursor cells to mature neurons, can be followed and dissected using these specific markers, as illustrated in Figure 4.

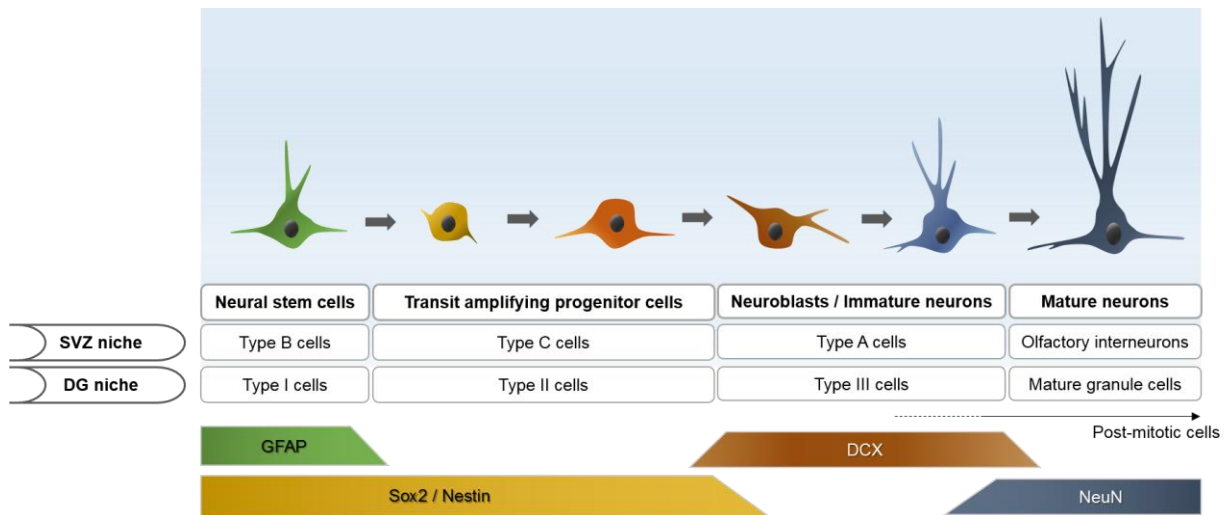


Figure 4 – Adult neurogenesis in the SVZ and DG niches and stage-specific markers. Summary of the 4 main developmental stages in adult neurogenesis both in SVZ and DG niches: neural stem cells divide into transit amplifying progenitor cells which give rise to neuroblasts/immature neurons that develop into mature neurons. GFAP: glial fibrillary acidic protein; Sox2: sex determining region Y-box 2; Nestin; DCX: doublecortin; NeuN: neuronal nuclei.

Overall, the SVZ and SGZ neurogenic niches exhibit key similarities and differences that seem important to mention. Overall, **similarities** between both neurogenic regions are displayed regarding niche composition (highly specialized with an heterogeneous population of cells), signaling pathways maintaining precursor pools, temporal sequence of new neurons' integration, critical periods of survival (the same regulatory factors act on specific time windows of stage development) and enhanced plasticity, and contribution to circuit dynamics. On the contrary, these regions show **differences** in specific features, specifically in niche organization, SGZ is a compacted region overflowed with different neuronal inputs and neurotransmitters which promote continuous interplay and plasticity, whereas SVZ is a rich vascular microenvironment and is not interconnected with such dense neuronal network, in neuronal subtype differentiation, and in migration of newborn neurons^{25,29,42,64,65}.

1.2.1.3. Non-traditional neurogenic regions

The presence of “local” progenitor cells in various brain regions outside the “standard” neurogenic niches, has been shown^{24,66,67}. These areas, where it still remains controversial whether adult neurogenesis indeed occurs, include the neocortex, piriform cortex, striatum, cerebellum, hypothalamus, amygdala and substantia nigra^{15,66,68–72}.

According to Bonfanti & Peretto, 2011, we can include neurogenesis in the mammalian CNS into 2 distinct groups: a 'complete' neurogenesis, greatly limited to SVZ and SGZ neurogenic sites; and a rather 'incomplete' neurogenesis, generally occurring in the parenchyma of the adult brain (non-neurogenic regions)⁷³. This "incomplete" definition of non-neurogenic regions results from several characteristics: parenchymal regions lack a well-defined functional and structural niche organization; the source and nature of the NSPCs vary from region to region; these regions are not directly related with germinal layers. Importantly, neurogenesis is interrupted at various intermediate levels in these non-neurogenic regions, possibly because they are more sensitive to modulation by external regulatory mechanisms and for not being well established^{35,69}. In addition, Bonfanti & Nacher, 2012, suggested, that immature neurons stay in the CNS for indeterminate time and might be recruited into preexisting circuits under certain stimuli, representing a new form of plasticity between synaptic remodeling and adult neurogenesis³⁶.

Although many studies exist concerning neurogenesis in the adult neocortex, amygdala and striatum^{15,74-77}, the most studied non-traditional neurogenic region, to date, is the hypothalamus. Recent evidences suggest that the adult hypothalamus may, indeed, be a neurogenic niche because of its localization in the brain (adjacent to the third ventricle) and its neurogenic potential, with two neurogenic regions: hypothalamic ventricular zone and hypothalamic proliferating zone^{34,69,71,78}. Given the critical role of hypothalamic neural circuitry in maintaining physiological homeostasis via the HPA axis, functional integration of newborn neurons may result in atypical effects in physiology and behavior^{67,79}. Possibly may serve as a compensatory mechanism contributing to the plastic control of energy balance and flexibility to adapt to metabolic challenges⁸⁰⁻⁸².

1.3. Adenosine and adenosine receptors

The purinergic neurotransmission was introduced in 1972 after it was shown that adenosine 5'-triphosphate (ATP) was a transmitter in non-adrenergic, non-cholinergic inhibitory nerves in the guinea-pig *taenia coli*⁸³⁻⁸⁵. ATP is one of the most important molecules for human biology because it acts as an energy exchange coin for almost every metabolic reaction⁸⁶. Particularly in the CNS, ATP acts both as a fast excitatory neurotransmitter and as a neuromodulator and has powerful long-term effects in the developing brain and in disease⁸³.

Adenosine, formed from the catabolism of ATP, is a ubiquitous purine ribonucleoside and signaling molecule/messenger with important functions at maintaining energetic homeostasis and other physiological processes for mammals, particularly in excitable tissues like the heart and the brain⁸⁶⁻⁸⁸.

Generally, many of its actions involve reducing the activity of these excitable tissues or increase the delivery of metabolic substrates, thus, regulating metabolic dynamics⁸⁶. In fact, adenosine can induce vasodilation in most vascular areas, reduce blood pressure or heart rate and regulate activity in the sympathetic nervous system⁸⁷.

Particularly in the brain, which highly expresses adenosine receptors, purinergic fast transmission is involved in a multitude of physiological processes, including regulation of sleep, arousal, mood, motivation and neuroprotection^{83,86,89,90}. Also, it is involved in the control of innate and adaptive physiological systems, being related with most neuropathological disorders of the CNS, ranging from neurodegenerative diseases, such as Alzheimer's (AD), Parkinson's (PD) and Huntington's (HD) diseases to multiple sclerosis (MS), epilepsy, schizophrenia, cerebral ischemia and mood disorders (depression, anxiety), as well as brain cancer (glioma)^{83,84,86,89,91-94}.

The role of adenosine in the CNS can be seen as an action on two distinct frontlines: first, as a neuromodulator, interfering with neuronal circuitry dynamics and, second, as a homeostatic modulator, coordinating metabolic activity^{89,95,96}.

Extracellular adenosine modulates neuronal activity and many other physiological mechanisms operating via metabotropic adenosine receptors (ARs). This modulation happens through the activation of a seven transmembrane G-protein-coupled receptor (GPCR) that lead to changes in the intracellular levels of second messengers and ion channels (Ca^{2+} and K^{+} channels), thus influencing pre- and post-synaptically the entire neuronal network^{92,97}. ARs are present throughout all brain areas and can be divided into four subtypes (A_1 , A_{2A} , A_{2B} and A_3), based on their unique pharmacological profile, tissue distribution and transducing signaling pathways^{92,98-102}. These four receptors can be divided according to their affinity to adenosine in high affinity (A_1 and A_{2A} - $K_m < 30$ nM) and low affinity (A_{2B} - K_m 1-20 μM) receptor subtypes; the affinity of A_3 receptor is species-dependent, and is low in rodents and high in humans^{88,103,104}. These receptors are also known as P1 receptors, when named according to the terminology of purinoceptors: P1 - adenosine-sensitive and P2 - ATP-sensitive^{92,97,105}.

Typically, ARs can also be divided into subcategories according to their associated signaling pathways: A_1 and A_3 receptors are coupled to G_i proteins, having an inhibitory effect on adenylate cyclase, therefore decreasing cyclic adenosine monophosphate (cAMP) levels whereas A_{2A} and A_{2B} receptors are coupled to G_s proteins, having an stimulatory effect on adenylate cyclase, thus increasing cAMP levels (Figure 5)^{85,106}. Furthermore, ARs can be involved in the activation of other pathways, namely phospholipase C (PLC), Ca^{2+} signaling and mitogen-activated protein kinases (MAPKs)^{88,100,106}.

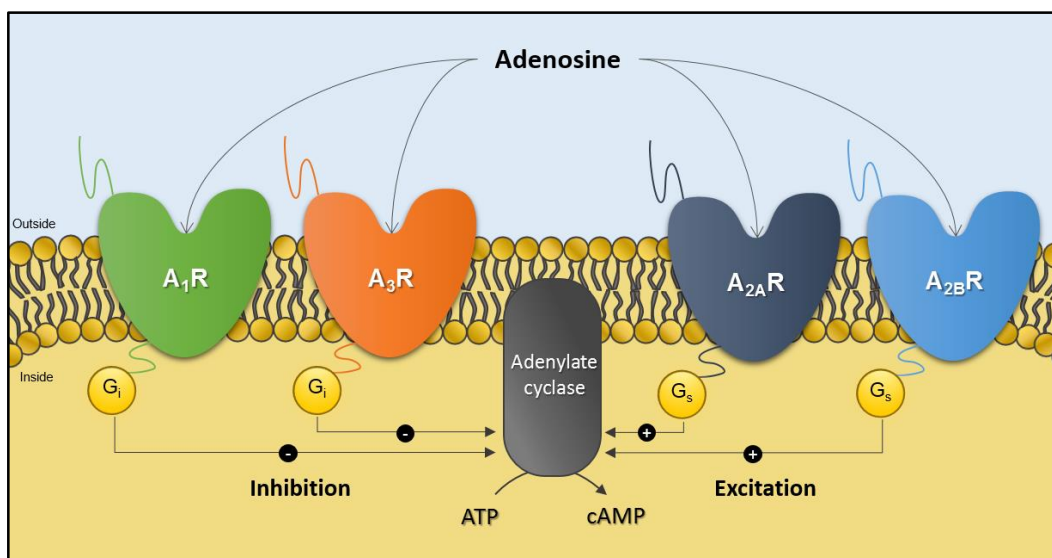
The distribution of ARs throughout the entire CNS is not homogeneous. A_1 Rs are most abundant in the neocortex, limbic system, dorsal horn of spinal cord, basal ganglia and cerebellum whereas A_{2A} Rs are highly expressed in the basal ganglia and olfactory bulb.

Subcellularly, A₁Rs are predominantly located in axons whereas A_{2A}Rs have a broader localization, whether dendritically in basal ganglia neurons or pre-synaptically in cortical neurons⁹⁷. A₁Rs and A_{2A}Rs are also located in astrocytes, microglia and oligodendrocytes and A_{2A}Rs in blood vessels¹⁰¹. In opposition, A_{2B}Rs and A₃Rs are expressed in low levels in the brain, being the expression of the latter moderate in the human cerebellum and hippocampus (extensively reviewed in the book ‘Adenosine Receptors in Health and Disease’ chapter Adenosine Receptors and the Central Nervous System¹⁰⁴, 2009).

Several key physiological and pathological effects of ARs have been described: e.g. A₁Rs are involved in the regulation of sleep and inhibition of neurotransmitter release; A_{2A}Rs are involved in the control of wakefulness and locomotion and play a role in neurodegeneration; A_{2B}Rs are linked with cardiac preconditioning and pro-inflammatory (acute injury) and anti-inflammatory (some chronic disease states) responses; A₃Rs mediate inflammatory responses and are involved in chronic neuropathic pain relief^{97,100,104,107}.

Although in a general sense many of adenosine effects are inhibitory, it is commonly accepted that activation of A₁R is translated into a neuroprotective function: by acting on A₁Rs it promotes the decrease of glutamate release (avoiding glutamate excitotoxicity conditions) and hyperpolarizes neurons, improving brain repair mechanisms⁹⁵. On the other hand, both stimulation or blockage of A_{2A}Rs under specific conditions was found to promote brain protection^{101,108,109}.

In particular, A_{2A}Rs modulatory role of neuronal activity seems to be especially important because of their numerous actions in the CNS, ability to “fine-tune” the functioning of other neurotransmitter modulatory systems and neuroprotective contribution on several brain diseases^{101,110,111}.



◀ **Figure 5 – Adenosine receptors signaling.** Adenosine activates four types of receptors: A₁R & A₃R, inhibitory receptors of adenylate cyclase; A_{2A}R & A_{2B}R, excitatory receptors of adenylate cyclase (re-illustrated from ¹⁰³Ham & Evans, 2012).

1.3.1. A_{2A} receptor

Adenosine A_{2A} receptor (A_{2A}R) neuromodulatory role can specifically comprehend the regulation of sleep-wake cycle, neuronal death, motor activity and psychiatric behaviors, inflammation, blood flow, angiogenesis and oxygen consumption^{95,101,107}. In addition to crucial roles in physiologic mechanisms, A_{2A}Rs are also involved in pathophysiological conditions of the CNS^{90,106,112,113}.

Structurally, A_{2A}Rs are organized, like every other GPCR, in seven transmembrane α -helices (7TM, helices 1–7) followed by one short membrane-associated helix, three extracellular loops (ECL1-3), three intracellular loops (ICL1-3), an extracellular amino-terminus (N-terminus) and a cytosolic carboxyl terminus (C-terminus). Particularly, their helical core, which constitutes the binding pocket, together with the extracellular loops and the four disulfide bridges of the extracellular N-terminal domain are especially important in ligand recognition and binding^{114–116}.

Regarding downstream signaling and second messengers (Figure 6), A_{2A}R interaction with the trimeric G-protein α -s/ β / γ causes the exchange of GDP to GTP, which is transduced into stimulation of adenylyl cyclase, increasing cAMP levels. A_{2A}R activation can, therefore, regulate MAPK activity, Ca²⁺ and K⁺ levels and other signaling pathways in a cAMP-dependent or -independent manner. In detail, MAPK activity is enhanced upon A_{2A}R activation by activating extracellular signal-regulated protein kinases 1/2 (ERK1/2) pathway. Furthermore, ERK1/2 pathway activation involves the activation of protein kinase A (PKA) which in turn phosphorylates and stimulates cAMP responsive binding element (CREB). Multiple PKA-related downstream mechanisms can also be triggered, like the Cdc42, which enhances PKC activity, ultimately, promoting cell survival. Moreover, A_{2A}R activation inhibits Ca²⁺ influx (PKA-dependent) and inhibits voltage-dependent Na⁺ channels (cAMP-dependent)^{97,100,101,117–121}.

Along with the complexity of A_{2A}R signaling, there is interplay between A_{2A}R receptors and receptors for other neurotransmitters and/or neuromodulators that enhances exponentially A_{2A}Rs-associated complexity and allows the modulation and “fine-tuning” of several systems. It can be detected either by A_{2A}Rs forming heteromers with other receptors or by targeting common intracellular transducing cascades^{92,110,111,121}. This interaction with other receptors (homo-, heteromerization or receptor mosaics), which can be direct or via intermediary adapter

proteins, may have an effect on communication networks intrinsic to the receptor complex, downstream signaling, ligand sensitivity/functionality, and compartmentalization of a given receptor. This interaction with other receptors and their signal transduction pathways may have an impact at the functional level and, ultimately, on phenotypical manifestation and behavioral output and responses^{122–126}.

Notably, strong evidences show that receptor-receptor interaction of A_{2A}R can occur with other G protein-coupled receptors (GPCRs), such as metabotropic glutamate receptor mGluR5¹²⁷, dopamine D₂ receptors¹²⁸ and cannabinoid CB₁ receptors (CB₁Rs)¹²⁹.

Interaction with other receptors has also been shown, namely with A₁R, with ionotropic receptors, like NMDA receptors or nicotinic cholinergic receptors (nAChR) and with receptors for neurotrophic factors, namely receptors for brain-derived neurotrophic factor – BDNF (TrkB) and glial-derived neurotrophic factors – GDNF (Ret and/or GFRα1) (extensively reviewed in the book ‘Adenosine Receptors in Health and Disease’ chapter Adenosine Receptors and the Central Nervous System¹⁰⁴, 2009).

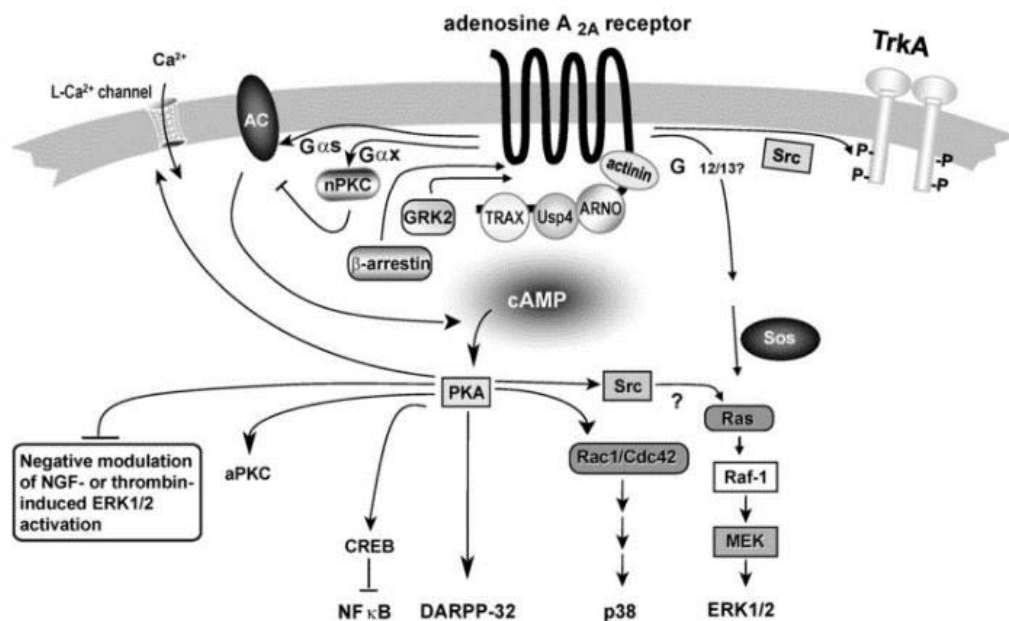


Figure 6 – A_{2A} receptor signaling pathways. A_{2A}R is a G protein alpha s (Gs) receptor. When A_{2A}R is activated, it activates adenylate cyclase which leads to production of cAMP. This induces the activation of cAMP-dependent kinase (PKA) pathway which triggers a sequence of downstream signaling events. Other kinases (e.g. nPKC) and G protein-independent mechanisms (e.g. coexistence and transactivation of TrkA receptors via Src pathway) are also illustrated as being part of A_{2A}R signaling (adapted from ¹²¹Fredholm et al., 2007).

The effects of agonists and antagonists of A_{2A}Rs (see Table 1) have been widely studied through crystallography and computational approaches in order to understand their potential as therapeutic modulators. Most of the A_{2A}Rs agonists are simply derivatives of adenosine modified at the 5'-position of the ribose and N⁶ positions of the purine. The most known A_{2A}R-selective agonist is CGS21680. On the contrary, A_{2A}Rs antagonists can be classically divided in xanthenes or non-xanthine derivatives. Natural occurring xanthenes like caffeine or theophylline generally have great affinity for ARs, with the highest affinity being at the A_{2A}R, constituting the most known typical antagonists for A_{2A}Rs⁹².

Table 1 – A_{2A}R main agonists/antagonists. Affinity for each AR type is represented by K_i values of agonists/antagonists (resulted from binding assays using either recombinant human ARs or rat ARs or from cAMP functional assays) (adapted from ¹⁰⁶Jacobson and Gao, 2006)

Ligand	A ₁ R K _i value (nM)	A _{2A} R K _i value (nM)	A _{2B} R K _i value (nM)	A ₃ R K _i value (nM)
A_{2A}R agonists				
NECA	14	20	140	25
CGS21680	289	27	>10000	67
DPMA	168	153	>10000	106
Binodenson	48000	270	430000	903
ATL-146e	77	0.5	N.D.	45
CV-3146	>10000	290	>10000	>10000
A_{2A}R antagonists				
KW6002	2,830	36	1800	>3000
CSC	28,000	54	N.D.	N.D.
SCH58261	725	5.0	1,110	1,200
SCH442416	1,110	0.048	>10000	>10000
ZM241385	774	1.6	75	743
VER 6947	17	1.1	112	1470
VER 7835	170	1.7	141	1931
'Schering compound'	82	0.8	N.D.	N.D.

N.D., not determined or not disclosed. Other A_{2A}R ligands may be considered, namely partial agonists, synthetic ligands or other derivatives, but are not represented. For references and for further details on the structures of the compounds listed in this table, see ⁹²Cristalli et al, 2009.

Importantly, recent evidences show the involvement of A_{2A}Rs in the regulation of specific stages of neurogenesis, such as cell proliferation and neuritogenesis. Particularly:

- ♦ Impaired neuritogenesis caused by p53 blockage was rescued by activation of the A_{2A}R (designated the A_{2A} rescue effect) via KIF2A, a kinesin family member¹³⁰;

- ◆ Long-term administration of low dose of caffeine, an antagonist of A_{2A}Rs, was shown to inhibit hippocampal neurogenesis and hippocampus-dependent learning and memory¹³¹;
- ◆ Also, caffeine was shown to alter, by a 7-day administration, the proliferation of adult hippocampal neuronal precursors in mice in a dose dependent manner – moderate doses lead to a decrease in proliferation whereas supra-physiological doses increase proliferation of neural precursors¹³²;
- ◆ Sleep deprivation-induced decline of neuronal proliferation and differentiation was rescued by the 48h treatment with caffeine¹³³;
- ◆ Unpublished data from our lab shows that A_{2A}R activation induces neurogenesis from DG neurosphere cultures.

1.4. Endocannabinoids and cannabinoid receptors

Endocannabinoids and their receptor signaling system (ECS - endocannabinoid system), are highly common across vertebrates and invertebrate species¹³⁴. In fact, the ECS is an important intrinsic mechanism of the human biology that is known to affect both CNS and peripheral processes¹³⁵.

Specifically, it consists of a family of lipid signaling molecules referred as endocannabinoids (eCBs), their receptor(s), bioactive intermediaries, downstream signaling pathways, uptake mechanism and specific metabolic synthesizing and degrading enzymes¹³⁶.

The two main characterized eCBs are anandamide (arachidonylethanolamide, AEA) and 2-arachidonoylglycerol (2-AG), which are widely distributed in the brain and specifically target cannabinoid receptors^{135,137}. 2-AG concentration is 200-fold higher than that of AEA in the brain tissue¹³⁴. Other eCBs have also been identified, such as 2-arachidonoylglyceryl ether (noladin ether), N-arachidonoyl-dopamine (NADA) and O-arachidonoyl-ethanolamine (virodhamine)^{138,139}.

Cannabinoid receptors (CBRs) constitute specific binding sites for endo- and exocannabinoids¹⁴⁰. CB₁ and CB₂ receptors, the most described CB receptors, belong to the superfamily of GPCRs and are physiologically different, exhibiting only 48% similarity in their amino acid sequences and 68% in the transmembrane domains¹³⁵. Their tissue distribution and downstream signaling mechanisms are distinct, which suggests their physiological divergent importance in mammalian biology^{138,141,142}.

CB₁ receptor is considered as the neuronal receptor whereas CB₂ receptor is considered as the receptor of the immune system^{122,138,141}. Although these two receptors are the best characterized to date, other putative CBRs can be modulated by eCBs, CBR agonists

and/or antagonists: GPR55 activity can be modulated by certain eCBs and phytocannabinoids but its pharmacology remains puzzling; 5 transient receptor potential (TRP) cation channels are also activated by eCBs, specifically TRPV1 (transient receptor potential vanilloid 1); peroxisome proliferator activated receptors (PPARs) have been shown to weakly bind eCBs as well as phytocannabinoids and synthetic cannabinoids^{138,143,144}. All the more, despite CB₁R is the most associated to the actions of endo- and exocannabinoids in the brain, increasing evidence shows that CB₂Rs, TRPV1Rs are present in the brain¹⁴⁵.

In view of that, the main eCBs 2-AG and AEA target CB receptors differently: 2-AG acts as a full agonist at CB₁R and CB₂R and AEA acts as partial agonist for CB₁, CB₂, GPR55 and TRPV1 receptors^{138,144,146–148} (see Table 2 for CB₁R and CB₂R agonists and antagonists).

Exocannabinoids are, in general, referred as the group of exogenous substances that are structurally related to Δ^9 -tetrahydrocannabinol (Δ^9 -THC) – the main psychoactive cannabinoid found in cannabis – and that bind to CBRs^{135,138,149}. These usually include plant-derived cannabinoids (phytocannabinoids) and synthetic cannabinoids¹³⁹.

Table 2 – CB₁R and CB₂R main agonists/antagonists. Affinity for either CB₁R or CB₂R is represented by K_i values of selective agonists and antagonists (resulted from *in vitro* binding assays using [³H]CP55940, [³H]HU243 or [³H]BAY-38-7271 for CB₁R-and CB₂R-specific binding sites) (adapted from ¹⁵⁰Pertwee, 2008)

Ligand	CB ₁ R K _i value (nM)	CB ₂ R K _i value (nM)
CB₁R selective agonists		
ACEA (arachidonyl-2'-chloroethylamide)	1.4, 5.29	195, >2000
O-1812	3.4	3870
ACPA (arachidonylcyclopropylamide)	2.2	750
2-arachidonylglycerol ether (noladin ether)	21.2	>3000
R-m-AEA (R-(+)-methanandamide)	17.9 to 28.3	815 to 868
CB₂R selective agonists		
AM1241	280	3.4
JWH-133	677	3.4
L-759633	1043, 15850	6.4, 20
L-759656	529 to >20 000	11.8 to 57
JWH-015	383	13.8
HU-308	>10000	22.7
Agonists without significant selectivity for CB₁R and CB₂R		
HU-210	0.06 to 0.73	0.17 to 0.52
CP55940	0.5 to 5.0	0.69 to 2.8

R-(+)-WIN55212	1.89 to 123	0.28 to 16.2
Nabilone	1.84	2.19
(-)-Δ^9-THC ((-)-<i>trans</i>-Δ^9-tetrahydrocannabinol)	5.05 to 80.3	3.13 to 75.3
(-)-Δ^8-THC ((-)-<i>trans</i>-Δ^8-tetrahydrocannabinol)	44, 47.6	39.3, 44
Anandamide	61 to 543	279 to 1940
2-AG (2-arachidonoylglycerol)	58.3, 472	145, 1400

CB₁R selective antagonists/inverse agonists

SR141716A	1.8 to 12.3	514 to 13 200
AM281	12	4200
AM251	7.49	2290

CB₂R selective antagonists/inverse agonists

SR144528	50.3 to > 10 000	0.28 to 5.6
AM630	5152	31.2

Other CB₁R and CB₂R ligands may be considered, namely synthetic ligands or other derivatives, but are not represented. For references and for further details on the structures of the compounds listed in this table, see ¹⁵¹Pertwee, 2005.

eCBs are not stored in vesicles, like most neurotransmitters, but rather are synthesized and released in postsynaptic neurons upon demand ('on-demand' model), by neural stimulation, (depolarization followed by increasing Ca²⁺ levels). eCBs activate mainly CB₁Rs localized in presynaptic neurons and are reuptaken (travel backwards across the synapse through a endocannabinoid membrane transporter – EMT) and degraded by specific hydrolyzing enzymes^{135,143,152–154}. They serve as fast retrograde messengers that actively suppress neurotransmitter release in presynaptic terminals in a phasic or long-lasting manner (depending on the type of stimulus or the action of factors that control their levels) in both excitatory or inhibitory neurons throughout the CNS¹⁴³. The resulting effect of eCBs is an inhibition of presynaptic terminals called 'depolarization-induced suppression of inhibition' (DSI) or 'excitation' (DSE) depending if it occurs at GABA or glutamate synapses, respectively^{152,155}. Whether this presynaptic inhibition is mainly due to the action of AEA or 2-AG retrograde signaling it still remains unclear^{135,152}. However, there is also evidence suggesting that eCBs can also act in a non-retrograde manner, modulating postsynaptic activity direct (autocrine CBR activation) or indirectly (via gliotransmission to astrocytes that can modulate pre- and post-synaptic terminals), revealing that it is an evolved multidimensional process^{143,155}.

Synaptic eCB signaling is highly complex and involves many metabolic pathways, activation of CB receptors and crosstalk of intrinsic mechanisms (summarized in Figure 7)¹⁴⁵. eCBs spatiotemporal controlled synthesis is performed through the coordinated expression of

◀ **Figure 7 – Synaptic eCB signaling.** Main pathways associated with synaptic eCB signaling. Both pre- and postsynaptic signaling cascades are involved (adapted from ¹⁴⁵Melis et al., 2014).

1.4.1. CB₁ receptor

CB₁ receptor (CB₁R) is among the most abundant GPCRs in the CNS (equivalent densities as for GABA and glutamate-gated ion channels)^{135,165}. It is also present in numerous peripheral tissues (although with a lower expression) like the heart, spleen, reproductive tissues, endocrine glands and intestinal tracts, etc^{138,166}. In the CNS, there is a wide distribution of CB₁Rs (see Table 3). The high expression of CB₁Rs in motor/sensory regions correlates with the prominent effects observed by stimulation with cannabinoids, namely in motor coordination and perception, pain modulation, cognition, memory and learning and regulation of emotional states^{124,135,156,167}. Additionally, CB₁Rs are mainly confined at presynaptic central and peripheral nerves, and the majority is expressed on GABAergic terminals and, in less concentrations, in glutamatergic terminals^{138,155}.

Importantly, these receptors are also expressed at low levels by astrocytes, oligodendrocytes, and neural stem cells, highlighting the importance of this receptor as a neuromodulator in the CNS internetworks^{168,169}.

Structurally, the CB₁R is organized, like other GPCRs, in seven α -helical transmembrane domains (7TMs) with an extracellular glycosylated amino-terminus and a cytosolic carboxyl-terminus and intra- and extracellular loops. Particularly, the cytoplasmatic region – the carboxyl-terminus – is responsible for G protein-binding, desensitization and cellular trafficking of the receptor and the extracellular loop is involved in ligand binding and receptor localization^{170–172}.

Both AEA and 2-AG seem to bind to CB₁Rs as full agonists (or nearly full agonists), triggering specific downstream signaling pathways¹⁷³. Predominantly, by coupling to G_{i/o} proteins, CB₁R regulates, through several signaling machineries, the activity of many proteins, ion channels, enzymes, second messengers and kinases (Figure 8)¹⁶⁹. Specifically, CB₁R activation:

- inhibits adenylyl cyclase, which decreases cAMP levels and downregulates the cAMP/ PKA pathway¹²⁴;
- activates all 3 families of functional MAPKs – p44/42 MAPK, p38 kinase and JNK-terminal kinase and also PI3K pathway¹⁷⁴;
- regulates voltage-gated Ca²⁺ channels (VGCCs)¹⁴⁸;
- inhibits presynaptic Ca²⁺ influx and inward rectifying K⁺ channels (responsible for K⁺ influx)¹⁵³;

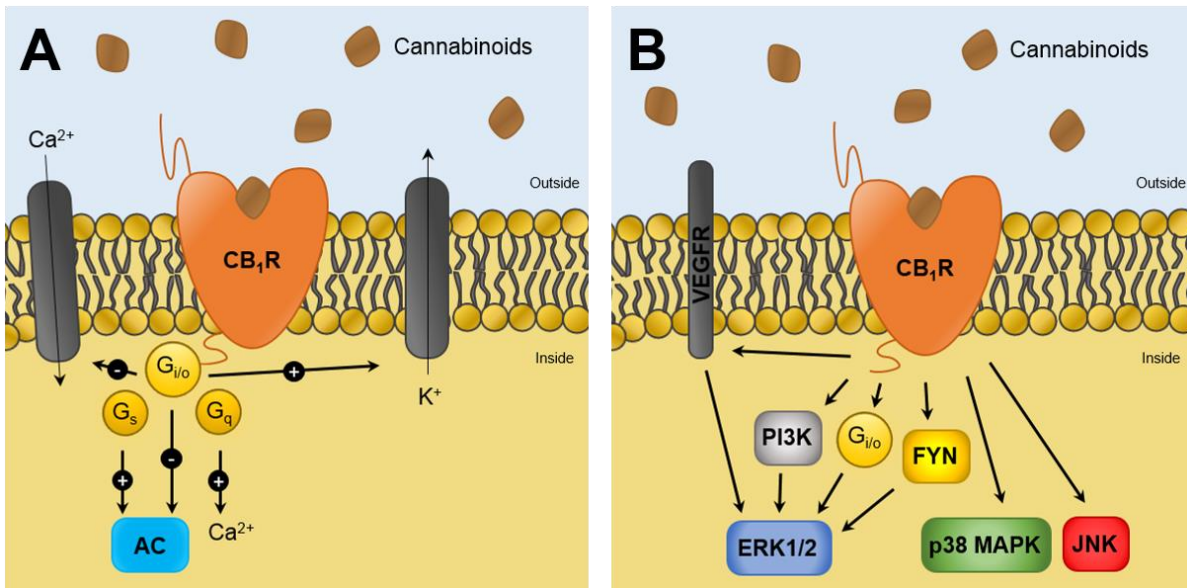


Figure 8 – CB₁R main signaling pathways. **A.** Activation of CB₁R activates G_{i/o} proteins inhibiting adenylate cyclase and cAMP levels and modulates ion channels. **B.** CB₁R activation can also activate different MAPK pathways (PI3K, ERK, p38 MAPK, JNK pathways) (re-illustrated from ¹⁷⁴Turu & Hunyady, 2010).

CB₁Rs can also modulate the release of several neuromodulators including serotonin, acetylcholine, dopamine, opioids, norepinephrine (among others) by interacting with other GPCRs, either by intracellular crosstalk of transduction signaling or by forming heteromers^{124,143,148,174}.

Evidence shows that heterodimerization occurs between CB₁R and other CB receptors, opiate δ receptors, dopamine D₁ and D₂ receptors and, with high relevance for this work, with adenosine A_{2A} receptors promoting alterations at the functional and signal transduction levels^{129,148,174–178}.

Importantly, in the brain, CB₁ and CB₂ receptors can form heteromers, demonstrating a bidirectional cross-antagonism phenomenon in which CB₁R antagonists have the ability to block the effect of CB₂Rs agonists and vice-versa¹⁷⁹.

Cannabinoid-based therapy, although particularly important due to the abundance of CB₁Rs throughout multiple brain areas and their complex interactions with other neuromodulatory systems, still remains controversial because the use of CB₁R agonists may lead to psychoactive side effects and craving consequences, as well as problems such as dosage concentrations and short window of beneficial actions may arise^{174,180,181}.

Table 3 – CB₁R expression in the mammalian CNS (adapted from ¹⁸²Svízenská et al., 2008)

Cannabinoid receptor	Localization	Intensity
CB ₁ R	– Telencephalon: Layers II, III, IV of the somatosensory cortex, layer II of the cingulate cortex, layers II and IV of the entorhinal cortex, layer III of the piriform cortex, association cortical regions of the frontal lobe; Molecular layer of the dentate area, CA1, CA2 and CA3 fields of Ammon's horn, subicular complex; Ependymal and subependymal zones of the olfactory bulb, anterior olfactory nuclei, olfactory part of the anterior commissure; Amygdala; Internal segment of the globus pallidus, caudate nucleus and putamen; Striatonigral pathway; Entopeduncular nucleus; – Brainstem: Substantia nigra (SN) <i>pars reticulata</i> ; Periaqueductal gray area (PAG); Gray matter around 4th ventricle; Spinal trigeminal tract and nucleus; – Cerebellum: Molecular layer – Spinal cord: Dorsal horn and lamina X – Dorsal root ganglia (DRG): Medium and large-sized neurons	Dense
	– Telencephalon: Layer V of the somatosensory cortex, temporal association cortex, secondary somatosensory and motor cortex, visual and auditory cortex; Polymorphic layer of the dentate area; Basal forebrain and septum; External segment of the globus pallidus, ventral pallidum, claustrum, and stria terminalis – Diencephalon: Anterior, mediodorsal, medioventral and intralaminar thalamic nucleus; Habenular nucleus; Lateral and paraventricular nucleus of the hypothalamus, infundibular stem – Brainstem: Solitary tract nucleus; Ambiguous nucleus; Inferior olive – Spinal cord: Deep dorsal horn; Thoracic intermediolateral nucleus	Moderate
	– Telencephalon: Primary motor and somatosensory, visual and auditory cortex; Granule cell layer of the dentate gyrus; Olfactory tubercle; Ventral pallidum; Nucleus Accumbens – Diencephalon: Sensory and motor thalamic nuclei; Subthalamic nucleus – Brainstem: Ventral tegmental area; SN pars compacta	Low

1.4.1.2. CB₁R role in neurogenesis

Cannabinoids, besides their neuromodulatory role, constitute a group of signaling cues that regulate neurogenesis at the levels of NSPCs proliferation, differentiation and migration^{136,139,141,183}. Importantly, CB₁R signaling influences the identity and cell features by regulating neuronal differentiation because its expression is not only increased with progressive differentiated stage but also because it is associated with proliferative and/or pro-survival cascades that allow the regulation of cell-cycle^{139,184,185}.

Growing evidences show that eCBs and CBRs have a huge impact on the regulation of the neurogenic process^{142,186,187}. Notably, CB₁R contribution to neurogenesis has been shown^{139,141,168} (for extensive review see Prenderville et al.¹⁸⁷). Specifically:

- ♦ Evidences shows that CB₁R knockout (KO) in mice results in impaired neurogenesis, suggesting a regulatory role of CB₁R in neurogenesis¹⁸⁸;

- ◆ The synthetic cannabinoid WIN-55,212-2, in addition to the selective FAAH inhibitor, URB597, have been shown to promote neurosphere generation, while WIN-55,212-2, URB597 and eCBs (both AEA and 2-AG) increase the number of BrdU (a marker for proliferation)-positive NPCs from dissociated neurospheres¹⁶⁸;
- ◆ CB₁R agonist ACEA was shown to promote murine neural precursor differentiation toward a neuronal lineage via CB₁R-dependent mechanism, suggesting that CB₁R activation may represent a pro-neuronal differentiation signal¹⁸⁹;
- ◆ Both CB₁R (ACEA) and CB₂R (JWH-056) agonists have been shown to stimulate the proliferation of primary murine cortical neurospheres¹⁹⁰;
- ◆ CB₁R activation (with R-m-AEA, a CB₁R agonist) was demonstrated to induce proliferation, self-renewal and neuronal differentiation in mouse neonatal SVZ cell cultures¹⁹¹;
- ◆ Treatment with CB₁R antagonist AM251 abrogated an exercise-induced increase of cell proliferation in the hippocampus, suggesting that endogenous cannabinoid signaling is required for this increase in cell proliferation¹⁹²;
- ◆ eCBs via CB₁R activation exerted a modulatory role on NPC proliferation and differentiation, under an excitotoxicity-induced neurogenesis context¹⁹³;
- ◆ AEA exposure was shown to affect murine NPCs cell fate determination; AEA treatment promoted an increase in glial differentiation followed by an increase in neuronal differentiation rates¹⁹⁴.

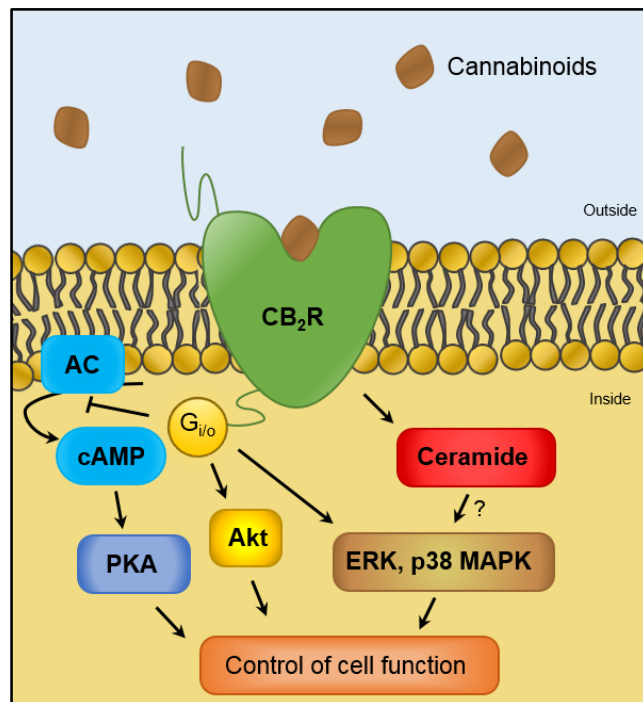
1.4.2. CB₂ receptor

CB₂ receptor (CB₂R) is also part of the receptor superfamily of GPCR and its localization is strikingly different from that of CB₁R¹⁶⁷. It is mainly present in cells of the immune and hematopoietic systems and other peripheral tissues^{124,180,195}. Specifically, it is present in specific tissues of immune cell production and regulation, the spleen, tonsils, thymus¹⁶⁵. In humans, CB₂R have important functions in the immune system by modulating the release of cytokines, molecules responsible for the regulation of immune function and inflammatory responses^{195,196}. The localization of these receptors in immune tissues suggests their role in key immunomodulatory function, particularly in the brain, where they are involved in immune surveillance through microglia (which are morphologically, phenotypically and functionally related to macrophages)^{124,148,196,197}. Nevertheless, and most importantly, CB₂R were also identified in the adult CNS particularly in glial cells, including microglia and astrocytes, neural and oligodendroglial progenitors, and neuronal subpopulations of certain brain structures^{195,198,199} (see Table 4 for CB₂R expression in the CNS). The multifocal existence of

functional CB₂Rs at CNS synapses implies their neuro/immunomodulatory action in brain's neurobiological processes associated with the stated regions like the control of pain, brain reward, cognition, emotion and others²⁰⁰.

As a GPCR, this receptor shares the same structural features as the CB₁R, characterized by seven transmembrane spanning domains, extracellular N-terminus, involved in ligand binding, 3 extracellular loops (EC1-3), 3 intracellular loops (IC1-3) and an intracellular C-terminal domain, which is involved in signal transduction and coupling to G proteins^{195,201}.

Different eCBs activate CB₂R signaling mechanisms differently. 2-AG was demonstrated to be a selective agonist whereas AEA was shown not to significantly bind to CB₂Rs¹⁹⁸. Similarly to CB₁R signaling, CB₂Rs can modulate adenylyl cyclase and MAP kinase activity, through their ability to couple to G_{i/o} proteins^{124,153}. This coupling triggers primarily the activation of canonical/classical intracellular responses that lead to the inhibition of adenylyl cyclase and subsequently to an impairment of cAMP/PKA short- and long-term effects. Since PKA modulates the expression of CREB, which is a transcription factor involved in the regulation of a variety of 'pro-survival-proliferation-differentiation' genes, the overall result of PKA inhibition is reduced cell maintenance. Additionally, there is stimulation of several MAPK cascades, namely ERK1/2 and p38 MAPK cascades and also PI3/Akt pathway, linked to pro-survival effects^{124,153,195,198}. However, in contrast to CB₁R, CB₂R stimulation is believed not to modulate ion channel function (Ca²⁺ and K⁺ channels), but has an impact in intracellular stores of Ca²⁺ from InsP3-sensitive stores by activation of PLC (Figure 9)^{124,195}.



◀ **Figure 9 – CB₂ receptor main signaling pathways.** CB₂R activation exerts an effect on adenylate cyclase and consequently on cAMP levels and protein kinase A (PKA) pathway; other pathways can also be activated, namely MAPK cascades (ERK, p38 MAPK) and Akt pathway (re-illustrated from ²⁰²Fernández-Ruiz et al., 2007).

Like CB₁R, cross-talk between CB₂R and other GPCRs is known to occur, but the molecular and cellular basis for the interactions, the extent to which they occur and the impact on CNS function is still not fully understood^{179,195}.

Overall, the neuroprotective role of CB₂R comes from their ability to predominantly mediate anti-inflammatory and immunomodulatory actions. This represents a crucial feature to specifically target the neuroinflammatory component of acute brain injuries (brain trauma or cerebral ischemia) or some neurodegenerative diseases, such as multiple sclerosis, Alzheimer's disease or amyotrophic lateral sclerosis, and consequently delaying brain damage^{149,181,198}.

All together, these evidences suggest that CB₂R play an important role in neuro-immunomodulatory responses and, because they do not have, unlike CB₁R, any psychoactive effect, they represent perfect targets for the development of new therapies based on CB₂R-agonist properties and actions^{148,149}.

Table 4 – CB₂R expression in the mammalian CNS (adapted from ¹⁸²Svízenská et al., 2008)

Cannabinoid receptor	Localization	Intensity
CB ₂ R	– Telencephalon: – Neurons of the layers III and V of the orbital, visual, auditory, motor and piriform cortex; Island of Calleja; Pyramidal neurons of the hippocampal CA2 and CA3 areas; Anterior olfactory nucleus; Striatum; Amygdala; – Diencephalon: Ventral and lateral posterior, posterior, and paracentral thalamic nuclei; Retina – Brainstem: Dorsal cochlear nucleus; Facial nucleus – Cerebellum: Purkinje cell bodies; Cerebellar granule cells – DRG: Neurons of a neonatal rat	Dense
	– Diencephalon: Geniculate body nuclei – Brainstem: SN <i>pars reticulata</i> – neurons larger than 20 µm; PAG; Inferior colliculus, interpeduncular, paratrochlear, and red nuclei; Paralemniscal nucleus, dorsal nucleus of lateral lemniscos; Pontine nuclei; Paratrochlear nucleus, medial and lateral vestibular nuclei; Parvocellular reticular nucleus; The spinal trigeminal tract nucleus – Cerebellum: Dendrites of Purkinje cells in the molecular layer	Moderate
	– Diencephalon: Paraventricular and mediodorsal thalamic nuclei; Ventromedial and arcuate hypothalamic nuclei	Low

1.4.2.1. CB₂R role in neurogenesis

Gathering evidences show the implication of the CB₂R in processes related to the control of proliferation, differentiation and survival of neural cells^{198,202,203}. Fairly recent evidences show that CB₂Rs have a role in neurogenesis by promoting NSPCs proliferation or by reestablishing neurogenic properties (for extensive review see Prenderville et al.¹⁸⁷). Specifically:

- ◆ Evidences show that CB₂R-KO reduces the self-renewal of murine embryonic cortical NPCs, while both HU-308 and JWH-133 (CB₂R agonists) increase both primary neurosphere generation and neural progenitor self-renewal *in vitro*²⁰⁴;
- ◆ Activation of CB₂Rs was shown to promote mouse NPCs proliferation and an increase in neurosphere formation through a mechanism dependent on the phosphoinositide-3 kinase/Akt pathway²⁰⁵;
- ◆ Treatment with a CBR agonist (WIN55,212-2) or with a CB₂R-selective agonist (JWH-133) in an *in vivo* study showed an increased NPCs proliferation on the SVZ of mice, via an autocrine DAGL-CB₂R signalling arrangement, with this effect being more pronounced in aged mice²⁰⁶;
- ◆ The administration of HU-308 (a CB₂R selective agonist) was shown to induce an increase in proliferation of NSPCs via PI3K/Akt/mTORC1-dependent signaling both *in vitro* and *in vivo*²⁰⁷;
- ◆ More recently, CB₂R agonist AM1241 has been shown to promote the proliferation/differentiation of primary normal human NSPCs (hNSCs) in the presence of the HIV-1 neurotoxic glycoprotein Gp120; Gp120-induced DNA fragmentation was reduced by the administration of AM1241 as well as astroglyosis and gliogenesis, which suggests a neuroprotective role of CB₂Rs against impaired neurogenesis²⁰⁸;

Accordingly, the fact that eCBs and exocannabinoids regulate NSPCs proliferation and differentiation constitutes a potential mechanism for the treatment of adult brain disorders, especially because 1) the ECS is activated to counteract/alleviate neuronal damage and neuroinflammation and 2) some symptoms associated with adult brain disorders appear to be correlated with dysregulation of eCBs^{135,141,156,169}.

By intervening on a pathophysiological context, cannabinoids may modulate adult neurogenesis and, ultimately, contribute for the treatment of a wide variety of disorders, such as anxiety, depression and neurodegenerative diseases like stroke, ALS, multiple sclerosis, Alzheimer's, Parkinson's and Huntington's disease^{156,180,181,209–213}.

1.5. Crosstalk between A_{2A} receptors and cannabinoid receptors in the CNS

A_{2A} and CB₁ receptors are highly expressed in the CNS, whereas CB₂Rs are expressed in low concentrations.

Notably, A_{2A}R crosstalk with CBRs (mainly CB₁R) is specifically significant because it modulates CB₁R actions which in turn has major repercussions on the motor depressant and rewarding effects of cannabinoids^{129,214}. Additionally, this interaction between A_{2A}Rs and CBRs is particularly important because it was shown to activate key modulatory effects on synaptic function and transmission^{92,215–217}.

Evidence for a structural and functional cross-talk between these receptors has been reported by several authors:

- ♦ In a human neuroblastoma cell line, CB₁R signaling was completely dependent on A_{2A}R activation. Moreover, on a behavioral perspective, blockage of A_{2A}Rs with selective antagonist ZM241385 counteracted the depreciating effects on motor control produced by the intrastriatal administration of a cannabinoid CB₁R agonist, suggesting that this was due to the formation of functional heteromeric complexes²¹⁴;
- ♦ A_{2A}R-CB₁R heterodimerization was shown to occur in living cells (human embryonic kidney cells, HEK-293T cells), as observed by bioluminescence resonance energy transfer (BRET) and fluorescence resonance energy transfer (FRET) – sequential BRET-FRET (SRET); additionally, the occurrence of A_{2A}R-D₂R, CB₁R-D₂R and A_{2A}R-CB₁R as well the existence of D₂R-A_{2A}R-CB₁R hetero-oligomers was also demonstrated^{175,177};
- ♦ The specific involvement of A_{2A}R in the addictive-related properties of cannabinoids was demonstrated by the reduction of THC-induced rewarding and aversive effects in mice lacking A_{2A}Rs compared to wild-type²¹⁸;
- ♦ A control of CB₁R function by A_{2A}Rs in glutamatergic terminals in the striatum was shown in which it was suggested an inhibitory presynaptic interaction between CB₁Rs and A_{2A}Rs²¹⁹;
- ♦ Detailed presynaptic interactions between A_{2A}R–CB₁R in glutamatergic nerve terminals of corticostriatal synapses in which presynaptic A_{2A}R activation dampened CB₁R-mediated inhibition of corticostriatal terminals was also demonstrated²²⁰;
- ♦ The activation of A_{2A}Rs was found to directly regulate the synaptic effects of CB₁Rs or indirectly, through activation of mGlu5Rs in the rodent striatum²²¹;

- ◆ Presynaptic CB₁Rs that interact with A_{2A}Rs were shown to be involved in the motor depressant and addictive effects of cannabinoids and that postsynaptic CB₁Rs interacting with A_{2A}Rs and D₂Rs are responsible for the cataleptogenic effects of cannabinoids¹²⁹;
- ◆ Blockade of CB₁Rs (either with antagonists or in CB₁R-KO mice) reduced the locomotor-activating effects of A_{2A}R antagonists, which, on the other hand, were able to increase endocannabinoid-dependent LTD²²²;
- ◆ Cannabidiol can induce robust neuroprotection (anti-inflammatory effects) mediated by CB₂Rs and adenosine receptors (mainly A_{2A}Rs) in an *in vitro* model of newborn hypoxic-ischemic brain²²³;
- ◆ Administration with an A_{2A}Rs-antagonist alters the reinforcing effects of cannabinoids (induced by CB₁R-agonists) by preferentially acting at presynaptic A_{2A}R sites²²⁴;
- ◆ A tight control of CB₁R function by presynaptic A_{2A}Rs in the inhibition of striatal glutamatergic terminals was described²¹⁹;
- ◆ Interestingly, the administration of A_{2A}Rs antagonists, in the presence of a sub-threshold dose of cocaine, was demonstrated to modulate synaptic activity and enhance locomotion and this enhanced activity required activation of CB₁Rs²²⁵;
- ◆ Receptor heteromers have also been demonstrated in native brain samples from parkinsonian rats and primates: in a rodent hemiparkinsonian model, the acute or chronic administration of L-DOPA was found to disrupt the A_{2A}-CB₁-D₂ receptor heteromers crosstalk²²⁶ and also L-DOPA treatment disrupts D₂R-A_{2A}R-CB₁R heteromers in the caudate nucleus of primates²²⁷;
- ◆ In a rodent model of Parkinson disease, single administration of A_{2A}R and CB₁R antagonists promoted dopaminergic survival whereas combined administration had a weakened effect, suggesting that the functional crosstalk between the adenosine and cannabinoid system may explain differences among single versus combined treatments²²⁸.

Overall, these studies suggest that the adenosinergic tone is important for cannabinoid-mediated effects and that A_{2A}Rs could exert a role on CBRs-mediated function via receptor heterodimerization²¹⁶.

Taking into account the differential effects of A_{2A} and CBRs agonists and antagonists in the brain it is possible to predict that this crosstalk may be involved in the regulation of the neurogenic process. It is also possible to infer that eCBs may play a role in regulating neurogenesis through interaction with A_{2A}Rs.

1.6. Main objective and specific aims

Growing evidences suggest that there is contribution of A_{2A} Rs and CB_1 Rs and CB_2 Rs to neurogenesis. Nevertheless, most of these studies only evaluate the individual effect of CBRs agonists and antagonists in neurogenesis whereas A_{2A} Rs role remains to be fully established. In fact, to date, the crosstalk between A_{2A} Rs and CBRs in neurogenesis has not been evaluated. Therefore, we proposed to evaluate this putative crosstalk based on 2 guiding criteria:

- Characterize the crosstalk of A_{2A} receptors with CB_1 Rs/ CB_2 Rs in both neurogenic niches (SVZ and SGZ/DG);
- Understand how the crosstalk between these receptors can affect the following neurogenic properties: cell-fate, proliferation and neuronal differentiation;

Considering the widespread brain distribution of A_{2A} Rs and the ECS, a better understanding of this possible interaction between CBRs and A_{2A} Rs, specifically at neurogenic niches, could contribute to the development of therapeutic alternatives to several brain disorders, either by potentiating endogenous repair mechanisms or preventing further brain damage. Consistently, both systems may, in the long run, represent promising pharmacological platforms for developing therapeutic proneurogenic compounds that could act upon these platforms and enhance neurogenesis.

CHAPTER 2

Material & Methods

2.1. Ethics Statement

All experiments were performed in accordance with the European Community (86/609/EEC; 2010/63/EU; 2012/707/EU) guidelines. The work was performed with biological material obtained from rat pups and subsequently maintained *in vitro*. The pups were handled according to standard and humanitarian procedures to reduce animal suffering.

2.2. SVZ and DG Cell Cultures

SVZ and DG neurospheres were prepared from early postnatal (P1-3) Sprague-Dawley rats. After sacrificing the animal and removing its brain, SVZ and DG fragments were dissected out from 450 μm -thick coronal brain slices, digested with 0.05% Trypsin-EDTA (Life Technologies, Carlsbad, CA, USA) in Hank's balanced saline solution (HBSS, Life Technologies), and mechanically dissociated with a P1000 pipette. The originated cell suspension was then diluted in serum-free medium (SFM), composed of Dulbecco's modified Eagle's medium/Ham's F-12 medium with glutaMAX (DMEM+GlutaMAX, Life Technologies) supplemented with 100 U/mL penicillin and 100 $\mu\text{g}/\text{mL}$ streptomycin (Pen/Strep; Life Technologies), 1% B27 (Life Technologies) and growth factors (for SVZ cells: 20 ng/mL epidermal growth factor (EGF; Life Technologies); for DG cells: 20 ng/mL epidermal growth factor (EGF; Life Technologies) and 10 ng/mL fibroblast growth factor-2 (FGF-2; Life Technologies) (*proliferative conditions*). SVZ cells were then plated on uncoated Petri dishes and allowed to develop for six days, whereas DG cells were allowed to develop for twelve days, both in a 95% air-5% CO₂ humidified atmosphere at 37 °C.

Six and twelve days after plating SVZ and DG cells, respectively, the resulting neurospheres were adhered for 24h onto glass coverslips coated with 0,1 mg/mL poly-D-lysine (PDL, Sigma-Aldrich, St. Louis, MO, USA) in SFM devoid of growth factors (*differentiative conditions*). One day after plating, the medium was renewed with or without (control) a range of pharmacological treatments for A_{2A}R, CB₁R and CB₂R ligands (illustrated in Figure 10).

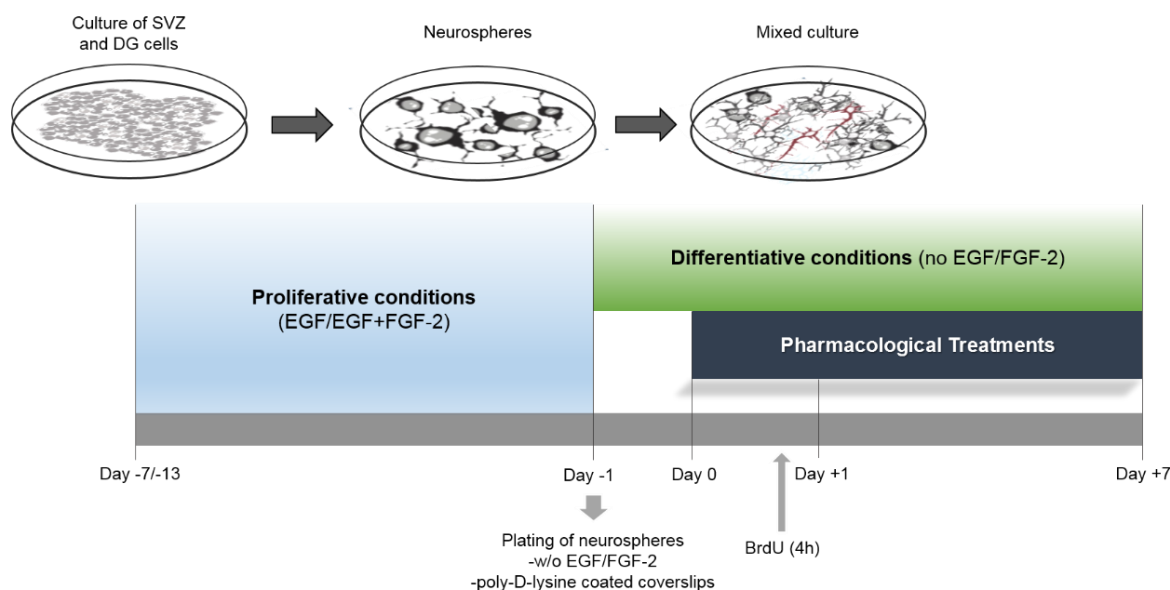


Figure 10 – SVZ and DG cell culture scheme. SVZ and DG cells were allowed to develop in proliferative conditions for 6 or 12 days, respectively. After that time period, the resulting neurospheres were plated onto PDL-coated coverslips in differentiative conditions without growing factors. Day 0 represents the day in which pharmacological treatments were executed.

2.3. Pharmacological Treatments

To study cell-fate, a Sox2 cell-pair assay was performed as described by Xapelli et al.¹⁹¹, where dissociated SVZ and DG cell suspensions obtained during the cell culture procedure were plated on poly-D-lysine coated glass coverslips at a density of 12800 cells/cm² and 19200 cells/cm², respectively. After seeding, SVZ and DG cells were grown, respectively, in SFM supplemented with 10 ng/mL EGF (low EGF) and in SFM supplemented with 10 ng/mL EGF and 5 ng/mL FGF-2 (low EGF/FGF-2). Moreover, plated cells were treated or not (control) with selective agonists and/or antagonists for A_{2A}Rs, CB₁Rs and CB₂Rs for 24h (Table 5). Thereafter, cells were processed for immunocytochemistry against Sox2, a marker of neural stem cells with the ability to self-renew.

To study cell proliferation, plated neurospheres were allowed to develop for 24h in the absence (control) or presence of the A_{2A}Rs, CB₁Rs and CB₂Rs ligands (Table 5). A thymidine analogue, 5-bromo-2'-deoxyuridine (BrdU, 10 μM, Sigma-Aldrich) was added for the last 4h of the culture session.

To study neuronal differentiation, neurospheres were allowed to develop for 7 days in the absence (control) or presence of the aforementioned ligands and an immunocytochemistry (ICC) for NeuN was performed.

Whenever cultures needed to be co-treated with a combination of drugs, SVZ or DG cells were primarily treated with CB₁R (ACEA) or CB₂R (HU-308) selective agonists for 30 minutes prior to A_{2A}R selective agonist (CGS21680) treatment and then grown for 24h in the case of cell-fate or proliferation studies, or 7 days, in the case of neuronal differentiation studies, in the presence of the ligands. Similarly, treatment with selective antagonists for CB₁Rs (AM251) and CB₂Rs (AM630) or A_{2A}Rs (ZM241385) was performed for 30 minutes prior to the treatment with CB₁Rs, CB₂Rs or A_{2A}Rs selective agonists and then co-incubated for further 24h in the case of cell-fate and proliferation studies, or 7 days in the case of neuronal differentiation studies.

Table 5 – Pharmacological treatments used.

Drug	Biological activity	Concentration used	Catalog number	Company
ACEA [N-(2-Chloroethyl)-5Z,8Z,11Z,14Z-eicosatetraenamide]	Cannabinoid CB ₁ receptor agonist	1 µM	1319	Tocris, Bristol, UK
HU-308 [4-[4-(1,1-Dimethylheptyl)-2,6-dimethoxyphenyl]-6,6-dimethylbicyclo[3.1.1]hept-2-ene-2-methanol]	Cannabinoid CB ₂ receptor agonist	1 µM	3088	
CGS21680 [4-[2-[[6-Amino-9-(N-ethyl-β-D-ribofuranuronamidoyl)-9H-purin-2-yl]amino]ethyl]benzenepropanoic acid hydrochloride]	Adenosine A _{2A} receptor agonist	30 nM	1063	
AM251 [N-(Piperidin-1-yl)-5-(4-iodophenyl)-1-(2,4-dichlorophenyl)-4-methyl-1H-pyrazole-3-carboxamide]	Cannabinoid CB ₁ receptor antagonist	1 µM	1117	
AM630 [6-Iodo-2-methyl-1-[2-(4-morpholinyl)ethyl]-1H-indol-3-yl](4-methoxyphenyl)methanone]	Cannabinoid CB ₂ receptor antagonist	1 µM	1120	
ZM241385 [4-(2-[7-Amino-2-(2-furyl)[1,2,4]triazolo[2,3-a][1,3,5]triazin-5-ylamino]ethyl)phenol]	Adenosine A _{2A} receptor antagonist	50 nM	1036	

2.4. Immunocytochemistry (ICC)

Cells were fixed for 30 minutes in 4% PFA in PBS, and permeabilized and blocked for non-specific binding sites for 1h30 with 0,5% Triton X-100 (Sigma-Aldrich) dissolved in PBS. Cells were then incubated overnight at 4°C with the primary antibodies (see Table 6) and 0.1% Triton X-100 and BSA 0.3% (w/v) in PBS, and for 1h at RT with the appropriate secondary antibodies in PBS as follows: donkey anti-rabbit Alexa Fluor 568, donkey anti-goat Alexa Fluor

568 or donkey anti-rat Alexa Fluor 488 (all 1:200 and all from Life Technologies). Nuclei were stained with Hoechst 33342 (6 µg/mL in PBS, Life Technologies). The final preparations were mounted using Mowiol fluorescent medium. Fluorescence images were recorded using an Axioskop 2 Plus fluorescent microscope (Carl Zeiss Inc., Göttingen, Germany).

2.5. Cell fate studies (Sox2 cell-pair assay)

SVZ and DG neurosphere-derived cells were stained for Sox2, a marker of NSPCs with the ability to self-renewal (Figure 11A). Cell pairs resulting from the division of a single NSPC were counted and categorized in 3 groups according to their Sox 2 expression: in both daughter cells (Sox 2 +/+), in only one of the daughter cell (Sox 2 +/-) and no expression (Sox2 -/-) (Figure 11B). Sox2 expression in the daughter cells characterizes the response of cells to the pharmacological treatment applied, ultimately reflecting the cell-fate of the pool of NSPCs, namely expansion (symmetrical self-renewal), maintenance (asymmetrical self-renewal) or extinction (symmetrical commitment).

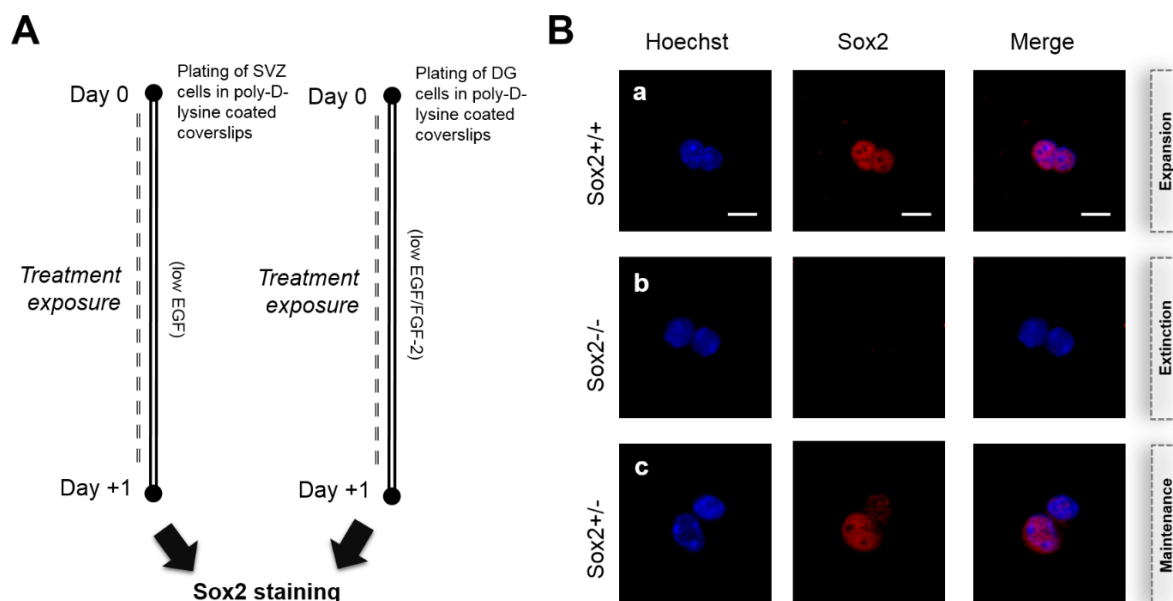


Figure 11 – Sox2 cell-pair assay. **A.** Schematic representation of the experimental protocol for studying SVZ and DG cell-fate. Day 0 represents the day of cultures in which cells were exposed to pharmacological treatments for the following 24h. **B.** Representative confocal images of cell pairs obtained following (a) the symmetrical division of a SVZ cell into two Sox2 + cells (Sox2 +/+), (b) the symmetrical terminal division into two Sox2- progenitors (Sox2 -/-) and (c) the asymmetrical division into a Sox2+ and a Sox2- progenitor (Sox2 +/-). Scale bars 20 µm. Sox2: sex determining region Y-box 2 (adapted from Xapelli et al.¹⁹¹).

2.6. Cell Proliferation Studies

To investigate the effect of the different pharmacological treatments on cell proliferation, SVZ and DG cells were exposed to 10 μ M 5-bromo-2'-deoxyuridine (BrdU) (Sigma-Aldrich), a synthetic thymidine analogue able to substitute thymidine in the DNA double chain synthesis occurring in dividing cells, for the last 4h of each specific pharmacological treatment (24h). Then, SVZ and DG cells were fixed in 4% PFA for 30 min and rinsed with PBS at RT. Subsequently, BrdU was unmasked by permeabilizing cells in PBS 1% Triton X-100 at RT for 30 min and DNA was denaturated in 1 M HCl for 40 min at 37°C. Following incubation in PBS with 0.5% Triton X-100 and 3% BSA to block nonspecific binding sites, cells were incubated overnight with the anti-BrdU antibody (see Table 6). After an additional rinse in PBS, nuclei counterstaining and mounting were performed as described previously.

Table 6 – Primary antibodies used for immunocytochemistry.

Antigen	Company	Catalog number	Host	Dilution
Sox2 (a marker of neural stem cells with the ability to self-renew)	Santa Cruz Biotechnology (Dallas, TX, USA)	sc-17320	Goat	1:100
BrdU (5-bromo-2'-deoxyuridine)	AbD Serotec, Bio-Rad Laboratories (Oxford, UK)	OBT00306	Rat	1:200
Neuronal Nuclei (NeuN) (mature neuronal marker)	Cell Signaling Technology (Danvers, MA, USA)	12943	Rabbit	1:200

2.7. Statistical Analysis

In all ICC experiments, measurements were performed at the border of SVZ and DG neurospheres, where migrating cells form a pseudo-monolayer of cells. In every independent experiment, each condition was measured in three different coverslips.

Percentages of Sox2 cell pairs were obtained from counting about 60 cell pairs in triplicate coverslips obtained from 3-6 independent cultures.

Percentages of BrdU and NeuN immunoreactive cells were calculated from cell counts in five independent microscopic fields in each triplicated coverslip with a 40x objective (approximately 200-300 cells per field).

Data are expressed as mean \pm standard error of the mean (SEM). Statistical significance was determined using one-way analysis of variance followed by Bonferroni's-multiple comparison test, with $p < 0.05$ considered to represent statistical significance. All graphs and statistical analysis were performed in Graph Pad Prism version 6.01 software (GraphPad Software, San Diego, California).

CHAPTER 3

Results & Discussion

3.1. Results Overview

The role of A_{2A}Rs on neurogenesis induced by activation of CB₁Rs/CB₂Rs in rat SVZ and DG stem/progenitor cell cultures was studied. In fact, using SVZ and DG neurospheres as a model, three distinct stages of neurogenesis were evaluated: cell-fate, proliferation and neuronal differentiation.

To have a clear picture of data, the results were displayed concerning SVZ and DG niches and then, within each niche, the three evaluated stages (cell-fate, proliferation and neuronal differentiation).

3.2. SVZ neurogenic niche

3.2.1.1. CB₁Rs or CB₂Rs or A_{2A}Rs activation induce no significant effects in SVZ cell-fate

In order to investigate the capacity of A_{2A}R, CB₁R and CB₂R ligands of modulating cell-fate of SVZ cells, a Sox2 cell-pair assay was performed as described by Xapelli et al.¹⁹¹, where SVZ cells were plated for 24h in medium complemented or not (control) with the ligands (Figure 12A).

It was observed that SVZ cells treated with CB₁R (ACEA, 1 μM) or CB₂R (HU-308, 1 μM) selective agonists or with A_{2A}Rs selective agonist (CGS21680, 30 nM) induced no significant changes in the percentages of either Sox2^{+/+} cell pairs (control: 50.93±0.84%; ACEA 1 μM: 53.76±3.00%; HU-308 1 μM: 53.19±3.64%; CGS21680 30 nM: 51.57±3.17%; N = 3) or Sox2^{-/-} cell pairs (control: 46.90±0.93%; ACEA 1 μM: 43.86±4.54%; HU-308 1 μM: 44.61±3.55%; CGS21680 30 nM: 47.66±3.06%; N = 3) when compared to control conditions (Figure 12B).

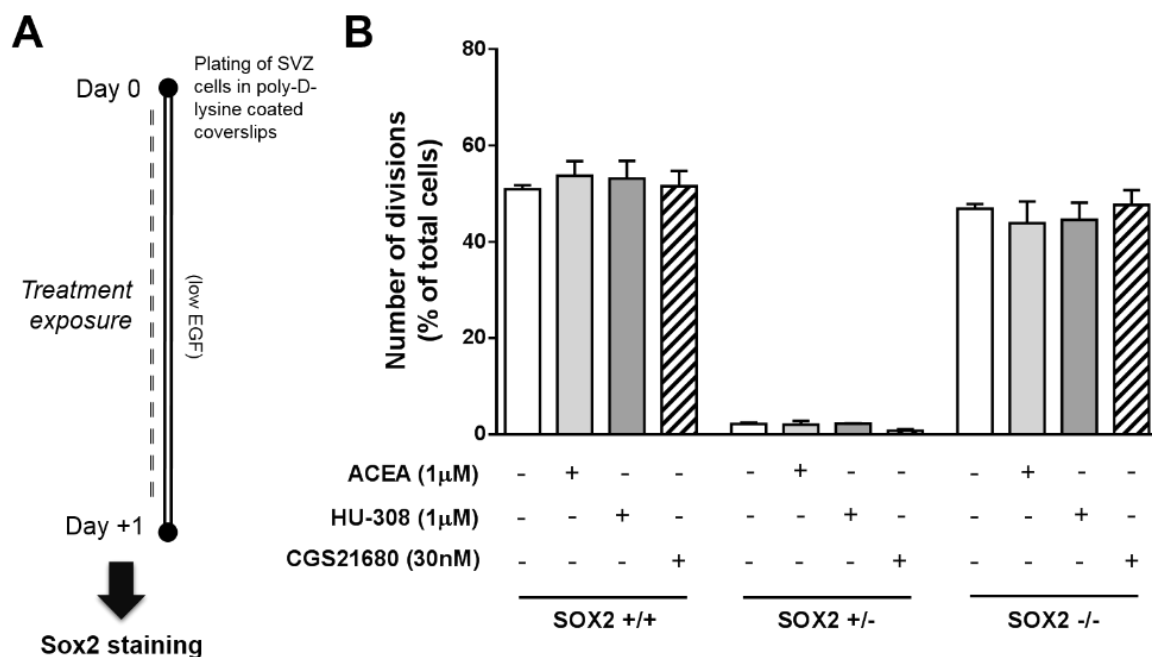


Figure 12 – CB₁R, CB₂R or A_{2A}R activation *per se* do not change the SVZ cell-fate. **A.** Schematic representation of the experimental protocol for studying SVZ cell-fate. Day 0 represents the day of cultures in which cells were exposed to pharmacological treatments for the following 24h. **B.** Bar graphs represent the number of each type of cell divisions counted (60 pairs counted *per* condition *per* experiment). Data are expressed as the percentage of total cell pairs and are represented as the mean ± SEM. N=3.

3.2.1.2. Co-activation of CB₁R or CB₂R with A_{2A}R induces no significant changes on SVZ cell-fate

It was next assessed the effect of a co-administration of both CB₁R and CB₂R selective agonists with the A_{2A}R selective agonist on the SVZ cell-fate. SVZ cells were treated with either ACEA (1 µM) or HU-308 (1 µM) for 30 minutes prior to CGS21680 (30 nM) treatment and then grown for 24h in the presence of the ligands.

As expected, the exposure to both CB₁R selective agonist (ACEA, 1 µM) and A_{2A}R selective agonist (CGS21680, 30 nM) induced no significant changes in the percentages of either Sox2^{+/+} cell pairs (control: 50.93±0.84%; ACEA 1 µM+CGS21680 30 nM: 49.83±1.27% N = 3) or Sox2^{-/-} cell pairs (control: 46.90±0.93%; ACEA 1 µM+CGS21680 30 nM: 48.21±1.05%; N = 3) as compared to control conditions (Figure 13A). Similarly, the co-administration of CB₂R selective agonist (HU-308, 1 µM) and A_{2A}R selective agonist (CGS21680, 30 nM) induced no significant changes in the percentages of either Sox2^{+/+} cell

pairs (control: $50.93 \pm 0.84\%$; HU-308 1 μM +CGS21680 30 nM: $48.38 \pm 3.70\%$ N = 3) or Sox2^{-/-} cell pairs (control: $46.90 \pm 0.93\%$; HU-308 1 μM +CGS21680 30 nM: $50.04 \pm 3.66\%$; N = 3) when compared to control (Figure 13B).

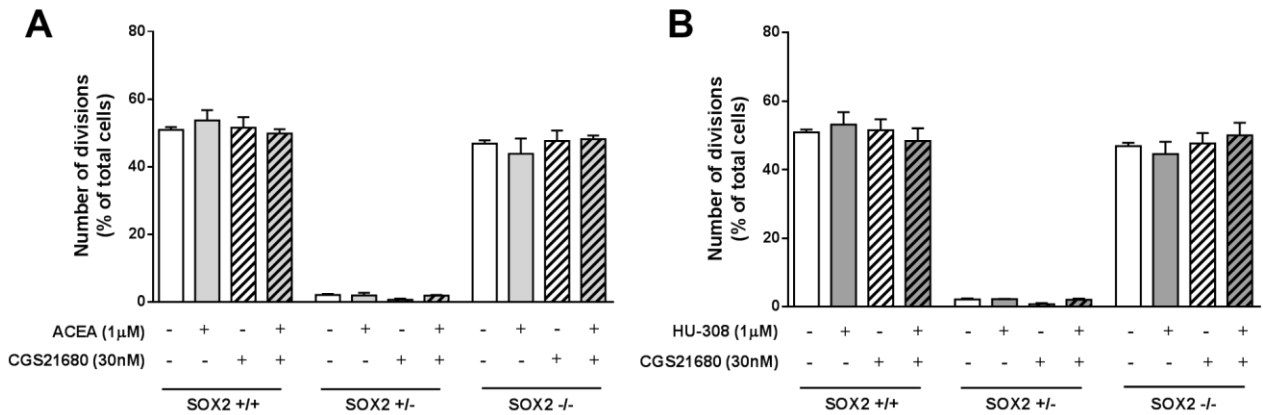


Figure 13 – Co-activation of CB₁Rs or CB₂Rs with A_{2A}Rs promotes no changes on SVZ cell-fate. Bar graphs represent the number of each type of cell divisions counted (60 pairs counted *per* condition *per* experiment) in cultures treated with CB₁Rs agonist ACEA (1 μM) and A_{2A}R agonist CGS21680 (30 nM) (**A**) or in cultures treated with HU-308 (1 μM) and A_{2A}R agonist CGS21680 (30 nM) (**B**). Data are expressed as the percentage of total cell pairs and are represented as the mean \pm SEM. N=3.

3.2.2.1. CB₁R agonist ACEA stimulates SVZ cell proliferation

Next it was investigated whether A_{2A}R, CB₁R and CB₂R ligands modulate SVZ cells proliferation. For that purpose, SVZ cells were treated with selective agonists for A_{2A}Rs, CB₁Rs and CB₂Rs for 24 days. BrdU, a thymidine analogue, was added during the last 4h of the culture to label SVZ cells that went through S-phase. After fixation, incorporated BrdU was immunolabeled and positive nuclei were counted (Figure 14A).

Treatment of SVZ cells with CB₁R agonist ACEA (1 μM) promoted a substantial increase in the number of BrdU-positive cells when compared to control cultures (control: $100.0 \pm 0.01\%$; ACEA 1 μM : $151.3 \pm 11.58\%$; N = 8, *** $p < 0.001$) whereas treatment with CB₂R agonist HU-308 (1 μM) and A_{2A}R selective agonist CGS21680 (30 nM) induced no significant alterations in the number of BrdU-positive cells when compared to control cultures (control: $100.0 \pm 0.01\%$; HU-308 1 μM : 99.75 ± 12.43 ; CGS21680 30 nM: $105.3 \pm 19.81\%$; N = 8) (Figure 14B, C).

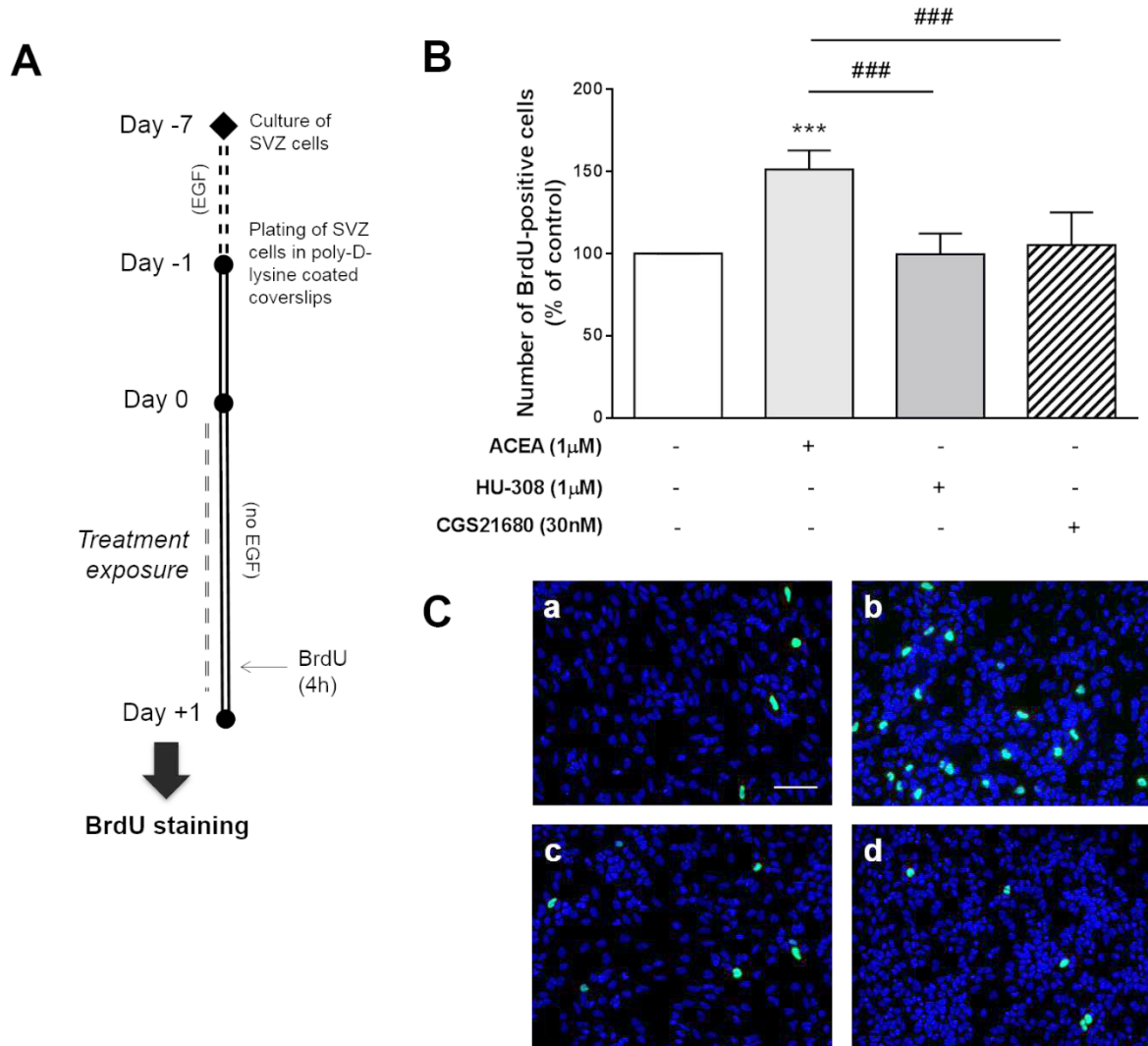


Figure 14 – CB₁R activation promotes SVZ cell proliferation while CB₂R and A_{2A}R activation does not. **A.** Schematic representation of the experimental protocol. Day 0 represents the day in which neurosphere-derived cells were exposed to pharmacological treatments for the following 24h. **B.** Bar graph depicts the number of BrdU-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean \pm SEM. Control was set to 100%. N=8. ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control; ###p<0.001, by ANOVA using Bonferroni's post test for comparison with ACEA. **C.** Representative fluorescent digital images of BrdU-positive cell nuclei (green) and Hoechst staining (blue nuclei), in control cultures (a) and in cultures exposed to ACEA 1 μ M (b), HU-308 1 μ M (c) and CGS21680 30 nM (d). Scale bar = 50 μ m.

3.2.2.2. CB₁R activation coupled with A_{2A}R activation preserves the increase on SVZ cell proliferation induced by CB₁R *per se*

In order to assess if co-incubation with the CB₁R or CB₂R selective agonists with the A_{2A}R selective agonist could induce any changes in SVZ cell proliferation, SVZ cells were treated with either ACEA (1 μM) or HU-308 (1 μM) for 30 minutes prior to CGS21680 (30 nM) treatment for 24h.

In the presence of both CB₁R selective agonist (ACEA, 1 μM) and A_{2A}R selective agonist (CGS21680, 30 nM), there was a significant increase in the number of BrdU-positive cells when compared to control cultures (control: 100.0±0.01%; ACEA 1 μM+CGS21680 30 nM: 163.8±27.63%; N = 8, ***p<0.001), while it was similar to ACEA exposure alone (ACEA 1 μM: 151.3±11.58%) (Figure 15). On the contrary, the co-administration of CB₂R selective agonist (HU-308, 1 μM) and A_{2A}R selective agonist (CGS21680, 30 nM) promoted, as expected, no alterations in the number of BrdU-positive cells as compared to control cultures (control: 100.0±0.01%; HU-308 1 μM+CGS21680 30 nM: 100.7±17.27%; N = 8) (Figure 16).

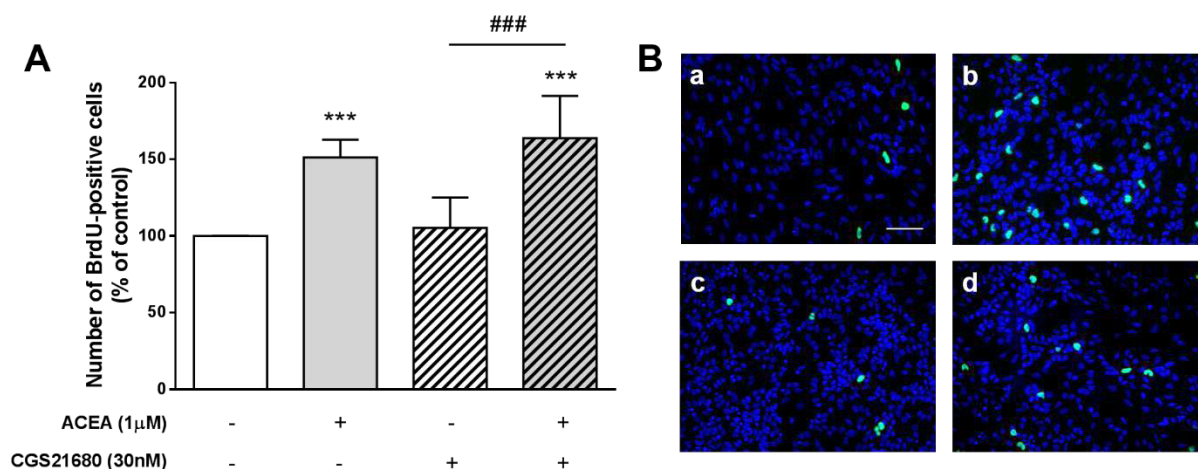


Figure 15 – Increased SVZ cell proliferation via CB₁R activation is maintained with A_{2A}R co-activation. **A.** Bar graph depicts the number of BrdU-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean ± SEM. Control was set to 100%. N=8. ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control; ###p<0.001, by ANOVA using Bonferroni's post test for comparison with CGS21680. **B.** Representative fluorescent digital images of BrdU-positive cell nuclei (green) and Hoechst staining (blue nuclei), in control cultures (a) and in cultures exposed to ACEA 1 μM (b), CGS21680 30 nM (c) and ACEA 1 μM+CGS21680 30 nM (d). Scale bar = 50 μm.

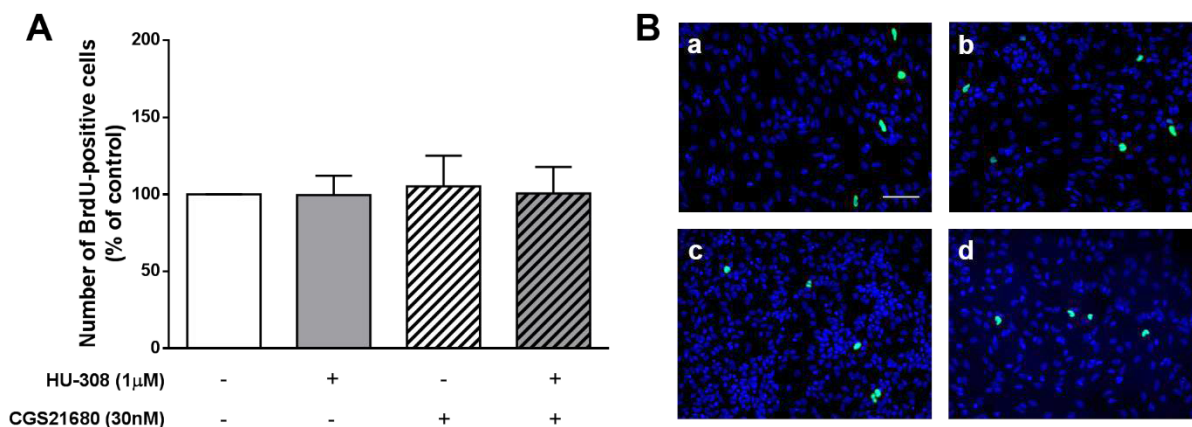


Figure 16 – SVZ cell proliferation is not affected by CB₂R and A_{2A}R co-activation. **A.** Bar graph depicts the number of BrdU-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean \pm SEM. Control was set to 100%. N=8. **B.** Representative fluorescent digital images of BrdU-positive cell nuclei (green) and Hoechst staining (blue nuclei), in control cultures (a) and in cultures exposed to HU-308 1 μ M (b), CGS21680 30 nM (c) and HU-308 1 μ M+CGS21680 30 nM (d). Scale bar = 50 μ m.

3.2.2.3. A_{2A}R is required for CB₁R-mediated stimulation of SVZ cell proliferation

I further wanted to study the putative role of A_{2A}Rs in CB₁R mediated increase in SVZ proliferation. For that purpose, SVZ cells were treated with A_{2A}R antagonist ZM241385 (50 nM) for 30 minutes prior to treatment with the CB₁R selective agonist ACEA (1 μ M) and then co-incubated for further 24h.

In fact, the increase in the number of BrdU-positive SVZ cells promoted by CB₁R activation was blocked by the presence of an A_{2A}R selective antagonist (ZM241385, 50 nM) (control: 100.0 \pm 0.01%; ACEA 1 μ M: 151.3 \pm 11.58%; ACEA 1 μ M+ZM241385 50 nM: 100.3 \pm 14.64%; ZM241385 50 nM: 96.13 \pm 10.80%; N= 4-8, ###p<0.001) (Figure 17).

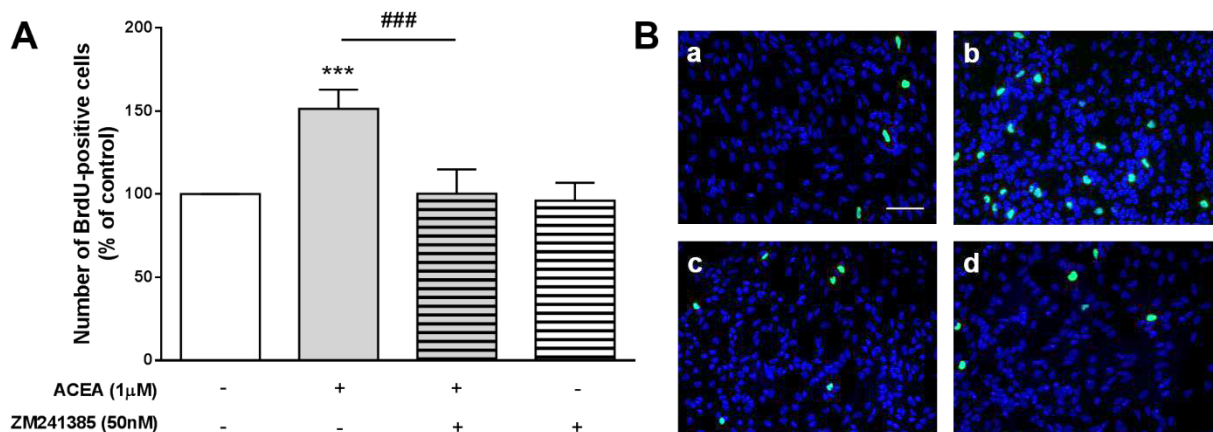


Figure 17 – A_{2A}R blockade impairs CB₁R-mediated increase in SVZ cell proliferation. **A.** Bar graph depicts the number of BrdU-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean \pm SEM. Control was set to 100%. N=4-8, ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control; ###p<0.001, by ANOVA using Bonferroni's post test for comparison with ACEA. **B.** Representative fluorescent digital images of BrdU-positive cell nuclei (green) and Hoechst staining (blue nuclei), in control cultures (a) and in cultures exposed to ACEA 1 μ M (b), ACEA 1 μ M+ZM241385 50 nM (c) and ZM241385 50 nM (d). Scale bar = 50 μ m.

3.2.3.1. CB₁R and CB₂R activation induces SVZ neuronal differentiation

Thereafter, the role of A_{2A}R, CB₁R and CB₂R ligands on SVZ neuronal differentiation was studied. For this, SVZ cells were treated with selective agonists for A_{2A}Rs, CB₁Rs and CB₂Rs for 7 days and then fixed and stained for NeuN (marker for mature neurons) (Figure 18A).

Treatment of SVZ cells with CB₁R (ACEA, 1 μ M) or with CB₂R (HU-308, 1 μ M) selective agonists induced an increase in the number of NeuN-positive cells when compared to control cultures (control: 100.0 \pm 0.02%; ACEA 1 μ M: 141.8 \pm 16.59%; HU-308 1 μ M: 128.6 \pm 12.36; N = 6, **p<0.01 and ***p<0.001) whereas treatment with A_{2A}R selective agonist (CGS21680, 30 nM) induced no significant changes (control: 100.0 \pm 0.02%; CGS21680: 111.6 \pm 11.87%; N = 6) (Figure 18B, C).

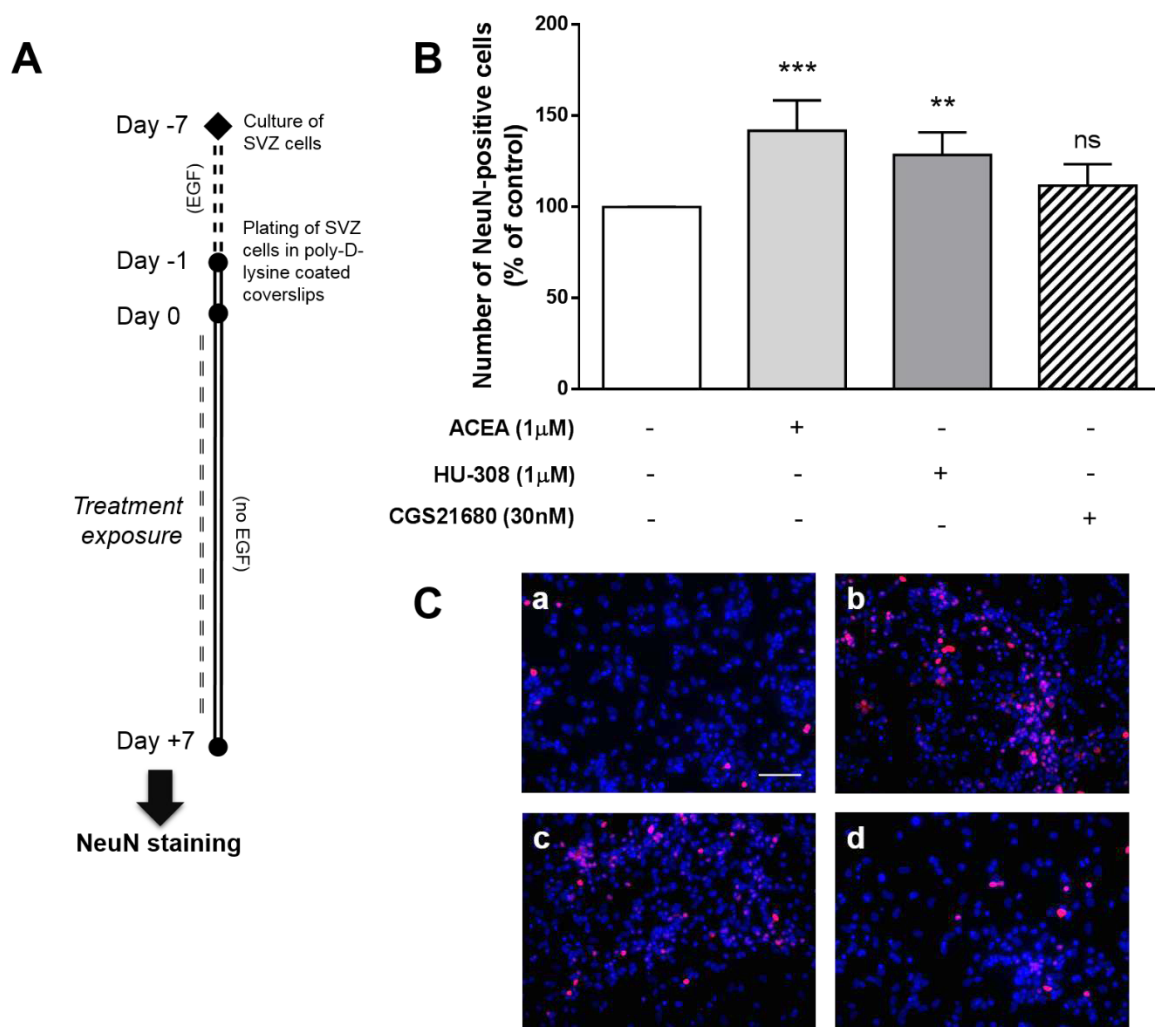


Figure 18 – CB₁R and CB₂R activation induces SVZ neuronal differentiation but not A_{2A}R activation. **A.** Schematic representation of the experimental protocol. Day 0 represents the day in which neurosphere-derived cells were exposed to pharmacological treatments for the following 7 days. **B.** Bar graph depicts the number of NeuN-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean \pm SEM. Control was set to 100%. N=6. **p<0.01 and ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control. ns: non-significant. NeuN: Neuronal Nuclei. **C.** Representative fluorescent digital images of NeuN-positive neurons (red) and Hoechst staining (blue nuclei), in control cultures (a) and in cultures exposed to ACEA 1 μ M (b), HU-308 1 μ M (c) and CGS21680 30 nM (d). Scale bar = 50 μ m.

3.2.3.2. SVZ neuronal differentiation promoted by CB₁R and CB₂R activation is not changed with A_{2A}R co-activation

In order to assess if a combined administration of both CB₁R and CB₂R selective agonists with the A_{2A}R selective agonist promoted any alteration on SVZ neuronal differentiation, SVZ cells were treated with either ACEA (1 μM) or HU-308 (1 μM) for 30 minutes prior to CGS21680 (30 nM) treatment and then grown for further 7 days in the presence of the ligands.

The results show that, in the presence of both CB₁R selective agonist (ACEA, 1 μM) and A_{2A}R selective agonist (CGS21680, 30 nM), there was a significant increase in the number of NeuN-positive cells when compared to control cultures (control: 100.0±0.02%; ACEA 1 μM+CGS21680 30 nM: 129.7±23.66%; N = 6, *p<0.05), while it was similar to ACEA exposure alone (ACEA 1 μM: 141.8±16.59%) (Figure 19). Also, the co-administration of CB₂R selective agonist (HU-308, 1 μM) and A_{2A}R selective agonist (CGS21680, 30 nM) induced an increase in the number of NeuN-positive cells as compared to control cultures (control: 100.0±0.02%; HU-308 1 μM+CGS21680 30 nM: 141.0±18.37%; N = 6, ***p<0.001), with a similar effect when comparing with HU-308 exposure alone (HU-308 1 μM: 128.6±12.36) (Figure 20).

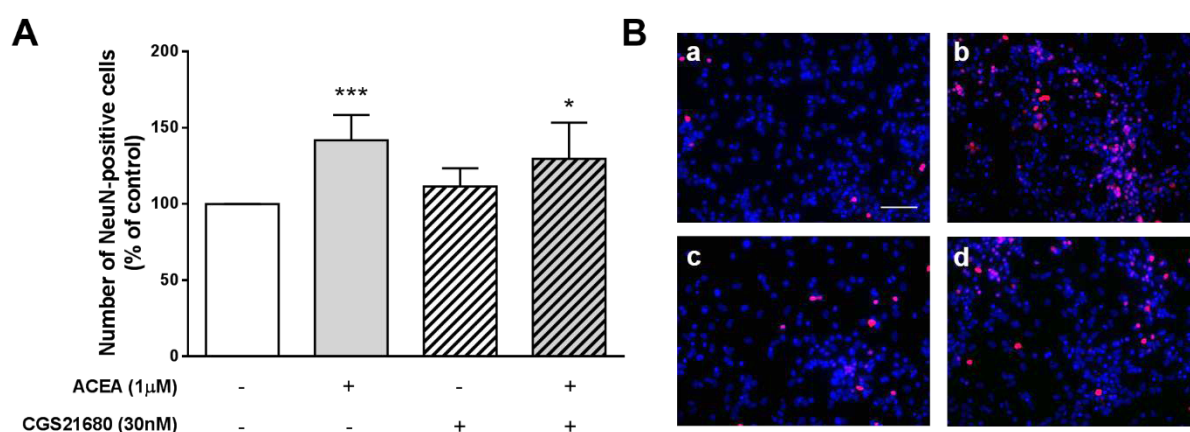


Figure 19 – Induced SVZ neuronal differentiation via CB₁R activation is not changed with A_{2A}R co-activation. **A.** Bar graph depicts the number of NeuN-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean ± SEM. Control was set to 100%. N=6. *p<0.05 and ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control. **B.** Representative fluorescent digital images of NeuN-positive neurons (red) and Hoechst staining (blue nuclei), in control cultures (a) and in cultures exposed to ACEA 1 μM (b), CGS21680 30 nM (c) and ACEA 1 μM+CGS21680 30 nM (d). Scale bar = 50 μm.

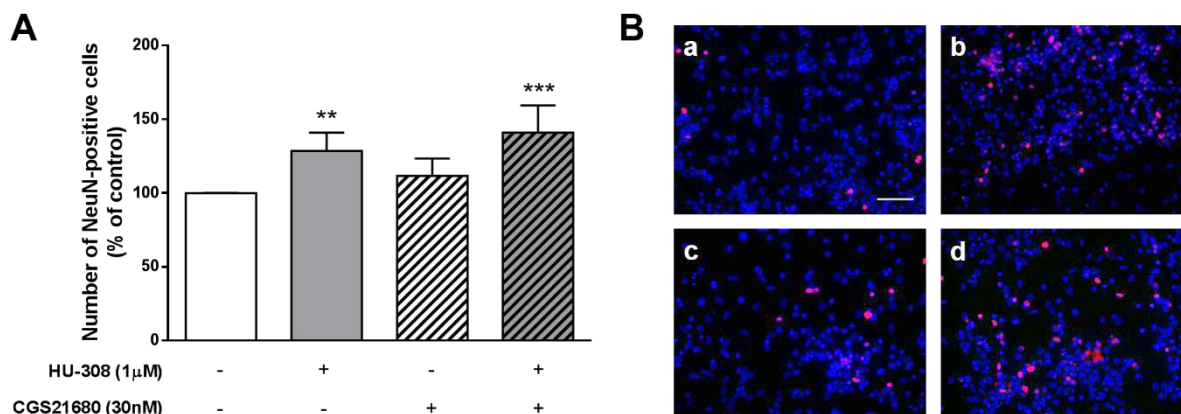


Figure 20 – Induced SVZ neuronal differentiation via CB₂R activation is not changed with A_{2A}R co-activation. **A.** Bar graph depicts the number of NeuN-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean ± SEM. Control was set to 100%. N=6. **p<0.01 and ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control. **B.** Representative fluorescent digital images of NeuN-positive neurons (red) and Hoechst staining (blue nuclei), in control cultures (a) and in cultures exposed to HU-308 1 μM (b), CGS21680 30 nM (c) and HU-308 1 μM+CGS21680 30 nM (d). Scale bar = 50 μm.

3.2.3.3. A_{2A}R activation is necessary for SVZ neuronal differentiation promoted by CB₁R and CB₂R

In line with the previous results we wanted to clarify the role of A_{2A}R in CB₁R and CB₂R activation-induced SVZ neuronal differentiation. For that purpose, SVZ cells were treated, as before, with A_{2A}R antagonist ZM241385 (50 nM) for 30 minutes prior to treatment with CB₁R or CB₂R selective agonists, ACEA (1 μM) or HU-308 (1 μM), respectively.

Interestingly, the results show that in the presence of an A_{2A}R selective antagonist (ZM241385, 50 nM), the increase in the number of NeuN-positive SVZ cells promoted by CB₁R activation is lost, being similar to control cultures (control: 100.0±0.02%; ACEA 1 μM: 141.8±16.59%; ACEA 1 μM+ZM241385 50 nM: 91.34±15.51%; ZM241385 50 nM: 106.4±8.53%; N = 6, ###p<0.001) (Figure 21A, C). In the same way, CB₂R activation coupled with A_{2A}R blockade induced no increase in the number of NeuN-positive SVZ cells, further suggesting the involvement of A_{2A}R on SVZ neuronal differentiation mediated by CB₂R activation (control: 100.0±0.02%; HU-308 1 μM: 128.6±12.36; HU-308 1 μM+ZM241385: 89.66±6.85%; ZM241385 50 nM: 106.4±8.53%; N = 6, ###p<0.001) (Figure 21B, D).

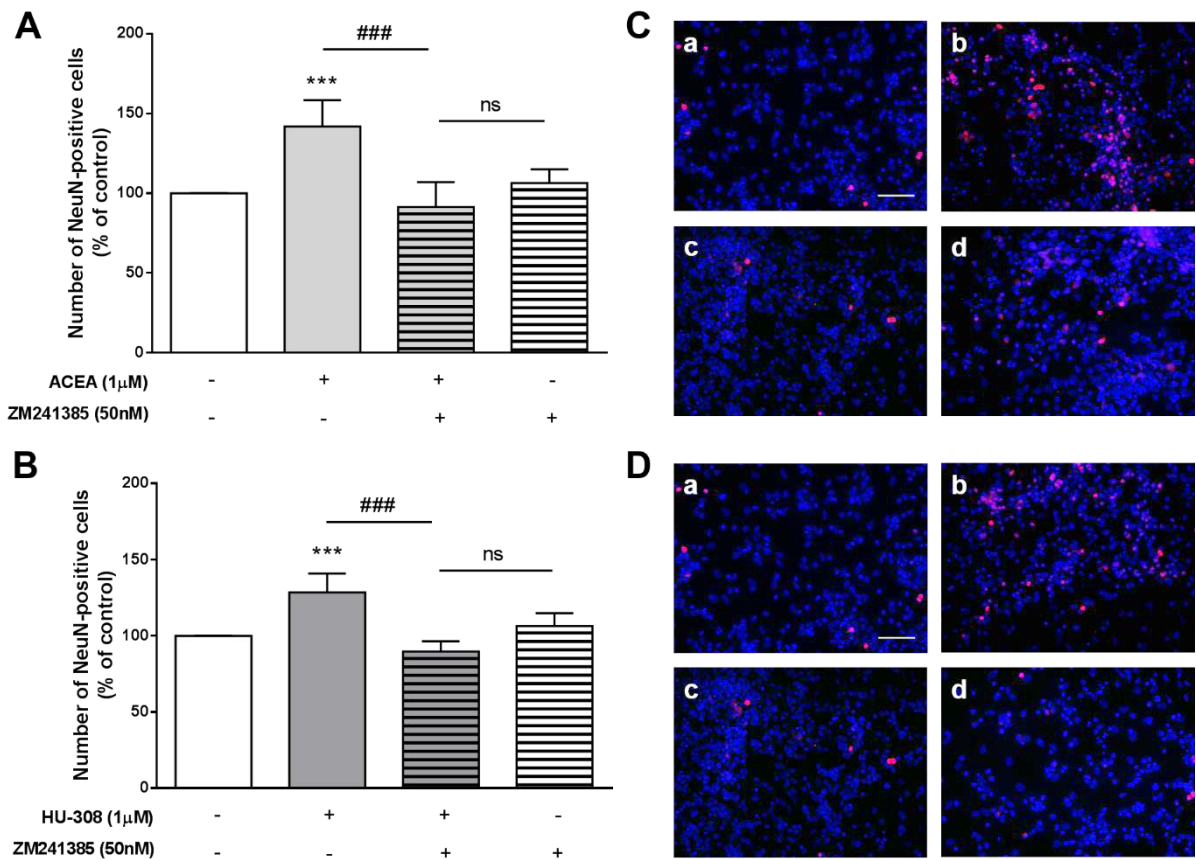


Figure 21 – A_{2A}R blockade impairs the CB₁R- and CB₂R-induced SVZ neuronal differentiation.

A, B. Bar graph depicts the number of NeuN-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean \pm SEM. Control was set to 100%. N=6. ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control; ###p<0.001, by ANOVA using Bonferroni's post test for comparison with ACEA and HU-308, respectively. ns: non-significant. **C, D.** Representative fluorescent digital images of NeuN-positive neurons (red) and Hoechst staining (blue nuclei), in control cultures (C,D-a) and in cultures exposed to ACEA 1 μ M (C-b), ACEA 1 μ M+ZM241385 50 nM (C-c) and ZM241385 50 nM (C-d); HU-308 1 μ M (D-b), HU-308 1 μ M+ZM241385 50 nM (D-c) and ZM241385 50 nM (D-d) Scale bar = 50 μ m.

3.3. DG neurogenic niche

3.3.1.1. CB₂R and A_{2A}R activation promotes self-renewal of DG cells

In order to investigate the ability of A_{2A}R, CB₁R and CB₂R ligands of modulating cell-fate of DG cells, a Sox2 cell-pair assay was performed as previously described (Figure 22A).

We observed that individual treatment of DG cells for 24h with CB₂R selective agonist HU-308 (1 μM) and A_{2A}R selective agonist CGS21680 (30 nM) induced a significant increase in the percentages of either Sox2^{+/+} cell pairs (control: 49.87±2.45%; HU-308 1 μM: 59.11±3.48%; CGS21680 30 nM: 57.66±4.03%; N = 6, **p<0.01 and ***p<0.001) with a concomitant decrease in the percentage of Sox2^{-/-} cell pairs (control: 48.54±2.20%; HU-308 1 μM: 39.24±3.02%; CGS21680 30 nM: 41.51±3.18%; N = 6, **p<0.01 and ***p<0.001) when compared to control (Figure 22B).

On the contrary, treatment with CB₁R agonist ACEA (1 μM) promoted no significant changes in the percentages of Sox2^{+/+} cell pairs (control: 49.87±2.45%; ACEA 1 μM: 49.53±1.43%; N = 6) and Sox2^{-/-} cell pairs (control: 48.54±2.20%; ACEA 1 μM: 49.24±1.22%; N = 6) when compared to control conditions (Figure 22B).

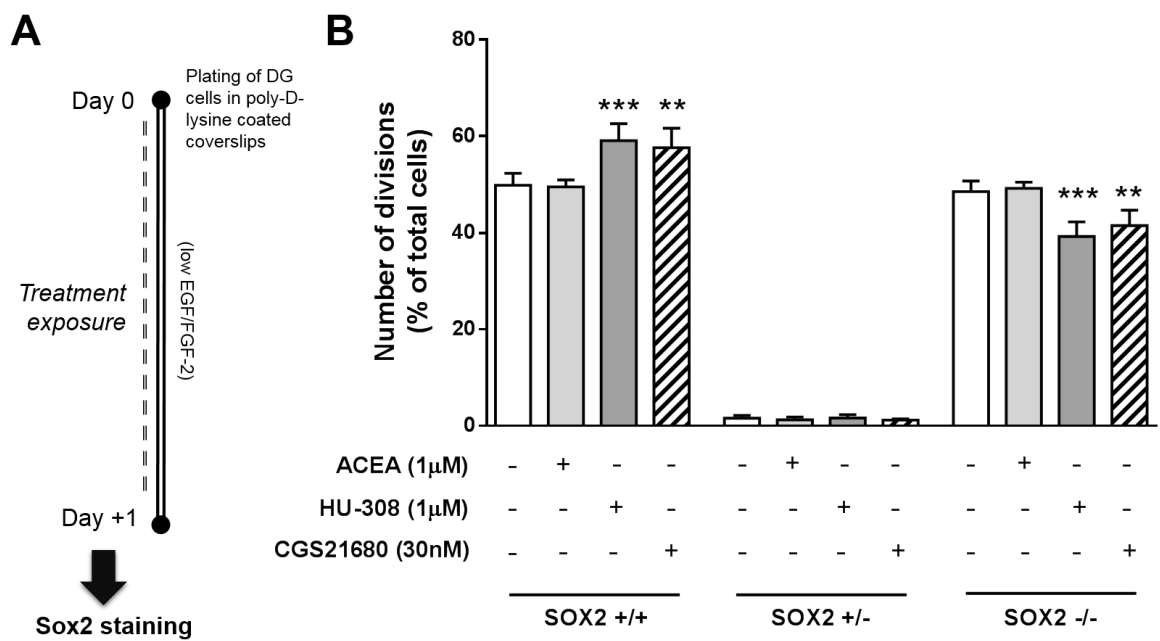


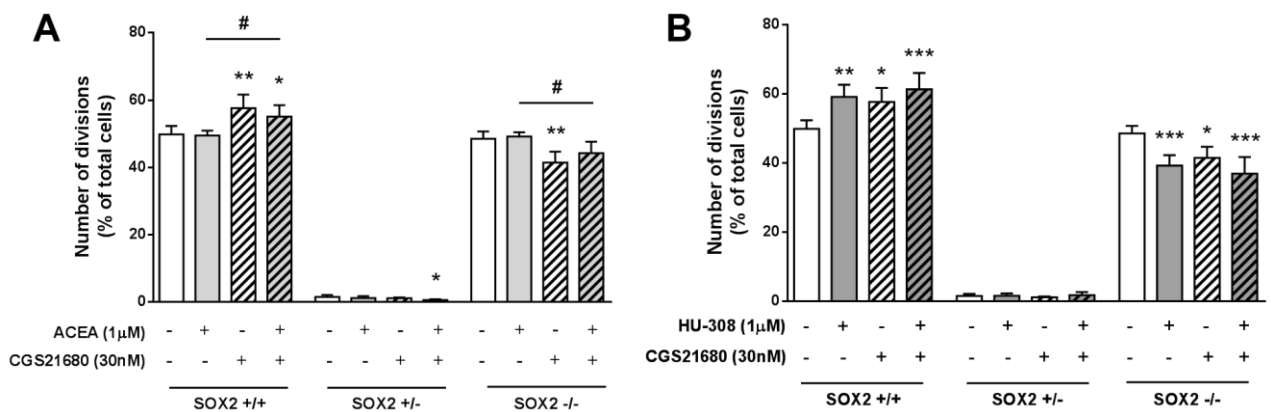
Figure 22 – CB₂R and A_{2A}R activation *per se* induce an increase in self-renew divisions on DG, whereas CB₁R activation does not. **A.** Schematic representation of the experimental protocol for studying DG cell-fate. Day 0 represents the day of cultures in which cells were exposed to pharmacological treatments for the following 24h. **B.** Bar graphs represent the number of each type of cell divisions counted (60 pairs counted *per condition per experiment*). Data are expressed as the percentage of total cell pairs and are represented as the mean ± SEM. N=6. **p<0.01 and ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control.

3.3.1.2. Self-renewal of DG cells is maintained when both CB₁R and CB₂R are co-activated with A_{2A}R

It was next assessed the role of a combined administration of both CB₁R and CB₂R selective agonists with the A_{2A}R selective agonist on DG cell-fate. Similarly to the SVZ treatment, DG cells were treated with either ACEA (1 μM) or HU-308 (1 μM) for 30 minutes prior to CGS21680 (30 nM) treatment and then grown for further 24h in the presence of the ligands. Considering that A_{2A}R activation had an effect on DG cell-fate *per se*, as previously presented, we expected this effect to be maintained when we combined ACEA or HU-308 with CGS21680 treatment, respectively.

In agreement with this hypothesis, the results demonstrate that, in the presence of both CB₁R selective agonist (ACEA, 1 μM) and A_{2A}R selective agonist (CGS21680, 30 nM), there is a significant increase in the percentage of Sox2^{+/+} cell pairs similar to CGS21680 exposure alone (control: 49.87±2.45%; CGS21680 30 nM: 57.66±4.03%; ACEA 1 μM+CGS21680 30 nM: 55.04±3.42%; N = 6, *p<0.05), as compared with control, with a tendency to decrease in the percentage of Sox2^{-/-} cell pairs (control: 48.54±2.20%; ACEA 1 μM+CGS21680 30 nM: 44.24±3.39%; N = 6) (Figure 23A).

In the same way, co-administration of CB₂R selective agonist (HU-308, 1 μM) and A_{2A}R selective agonist (CGS21680, 30 nM) induced a significant increase in the percentage of Sox2^{+/+} cell pairs (control: 49.87±2.45%; HU-308 1 μM: 59.11±3.48%; HU-308 1 μM+CGS21680 30 nM: 61.27±4.75% N = 6, ***p<0.001) with a concomitant decrease in the percentage of Sox2^{-/-} cell pairs (control: 48.54±2.20%; HU-308 1 μM: 39.24±3.02%; HU-308 1 μM+CGS21680 30 nM: 36.09±4.81%; N = 6, ***p<0.001) (Figure 23B).



◀ **Figure 23 – Co-activation of CB₁Rs or CB₂Rs with A_{2A}R promotes an increase in self-renew divisions on DG cell-fate.** Bar graphs represent the number of each type of cell divisions counted (60 pairs counted *per condition per experiment*) in cultures treated with CB₁Rs agonist ACEA (1 μM) and A_{2A}R agonist CGS21680 (30 nM) **(A)** or in cultures treated with HU-308 (1 μM) and A_{2A}R agonist CGS21680 (30 nM) **(B)**. Data are expressed as the percentage of total cell pairs and are represented as the mean ± SEM. N=6. *p<0.05, **p<0.01 and ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control. #p<0.05, by ANOVA using Bonferroni's post test for comparison with ACEA.

3.3.1.3. A_{2A}R activation is necessary for DG self-renewing promoted by CB₂R activation

In line with the previous results we wanted to clarify the role of A_{2A}Rs in DG cell-fate. Hence, in order to test this, DG cells were treated with A_{2A}R antagonist ZM241385 (50 nM) for 30 minutes prior to treatment with CB₁R or CB₂R selective agonists and then for further 24h in the presence of the ligands.

The results show that, as expected, in cultures treated with both CB₁R agonist (ACEA, 1 μM) and A_{2A}R antagonist (CGS21680, 30 nM), no changes were observed, when compared to control cultures, in the percentages of Sox2^{+/+} cell pairs (control: 49.87±2.45%; ACEA 1 μM: 49.53±1.43%; ACEA 1 μM+ZM241385 50 nM: 49.38±1.51%; ZM241385 50 nM: 50.38±3.67%; N = 6) or in the percentages of Sox2^{-/-} cell pairs (control: 48.54±2.20%; ACEA 1 μM: 49.24±1.22%; ACEA 1 μM+ZM241385 50 nM: 48.57±2.14%; ZM241385 50 nM: 50.94±1.40%; N = 6) when compared to control cultures (Figure 24A). Interestingly, in the presence of the A_{2A}R antagonist, the observed increase in self-renewal divisions of DG cells promoted by CB₂R activation was blocked, resulting in the decrease of the percentage of Sox2^{+/+} cell pairs (control: 49.87±2.45%; HU-308 1 μM: 59.11±3.48%; HU-308 1 μM+ZM241385: 51.80±1.99%; ZM241385 50 nM: 50.38±3.67%; N = 6, ####p<0.001) and in a concomitant increase in the Sox2^{-/-} cell pairs (control: 48.54±2.20%; HU-308 1 μM: 39.24±3.02%; HU-308 1 μM+ZM241385: 46.71±2.01%; ZM241385 50 nM: 50.94±1.40%; N = 6, ###p<0.01), when compared to HU-308 treatment, to similar values as control cultures (Figure 24B).

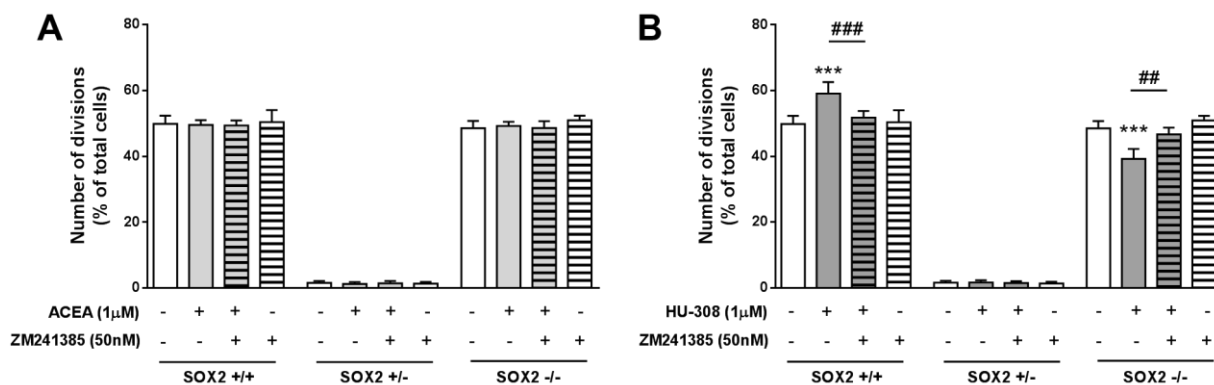


Figure 24 – A_{2A}R blockade impairs the CB₂R-mediated increase in self-renewal divisions of DG cells. Bar graphs represent the number of each type of cell divisions counted (60 pairs counted *per* condition *per* experiment) in cultures treated with CB₁R agonist ACEA (1 μ M) and A_{2A}R antagonist ZM241385 (50 nM) (**A**) or in cultures treated with HU-308 (1 μ M) and A_{2A}R antagonist ZM241385 (50 nM) (**B**). Data are expressed as the percentage of total cell pairs and are represented as the mean \pm SEM. N=6, ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control; ##p<0.01 and ###p<0.001, by ANOVA using Bonferroni's post test for comparison with HU-308.

3.3.1.4. DG self-renewing promoted by A_{2A}R activation is dependent on CB₁R or CB₂R activation

In addition, since both CB₂R and A_{2A}R promote an increase in DG self-renewing properties it was further tested if CB₁R and CB₂R are required for A_{2A}R role in DG cell-fate. To address this question, DG cells were treated with either CB₁R selective antagonist AM251 (1 μ M) or CB₂R selective antagonist AM630 (1 μ M) 30 minutes prior to treatment with A_{2A}R selective agonist CGS21680 (30 nM) and then incubated for further 24h in the presence of the ligands.

The results show that, the effect of A_{2A}R agonist CGS21680 (30 nM) in DG cell-fate is lost when cultures are co-treated with both A_{2A}R agonist (CGS21680, 30 nM) and CB₁R antagonist (AM251, 1 μ M), resulting in a decrease in the percentage of Sox2+/+ cell pairs (control: 49.87 \pm 2.45%; CGS21680 30 nM: 57.66 \pm 4.03%; CGS21680 30 nM+AM251 1 μ M: 50.19 \pm 2.66%; AM251 1 μ M: 49.06 \pm 1.25%; N = 6, ##p<0.01) and an increase in the percentages of Sox2-/- cell pairs (control: 48.54 \pm 2.20%; CGS21680 30 nM: 41.51 \pm 3.18%; CGS21680 30 nM+AM251 1 μ M: 48.62 \pm 2.60%; AM251 1 μ M: 49.58 \pm 2.06%; N = 6, ##p<0.01) when compared to CGS21680 treatment alone (Figure 25A).

Similarly, the same happened in cultures co-treated with both A_{2A}R agonist (CGS21680 30 nm) and CB₂R antagonist (AM630, 1 μM), where it is observed a decrease in the percentage of Sox2^{+/+} cell pairs (control: 49.87±2.45%; CGS21680 30 nM: 57.66±4.03%; CGS21680 30 nM+AM630 1 μM: 50.74±3.24%; AM630 1 μM: 46.48±2.47%; N = 6, #p<0.05) and an increase in the percentages of Sox2^{-/-} cell pairs (control: 48.54±2.20%; CGS21680 30 nM: 41.51±3.18%; CGS21680 30 nM+AM630 1 μM: 47.91±3.46%; AM630 1 μM: 52.54±2.58%; N = 6, #p<0.05) when compared to CGS21680 treatment alone (Figure 25B).

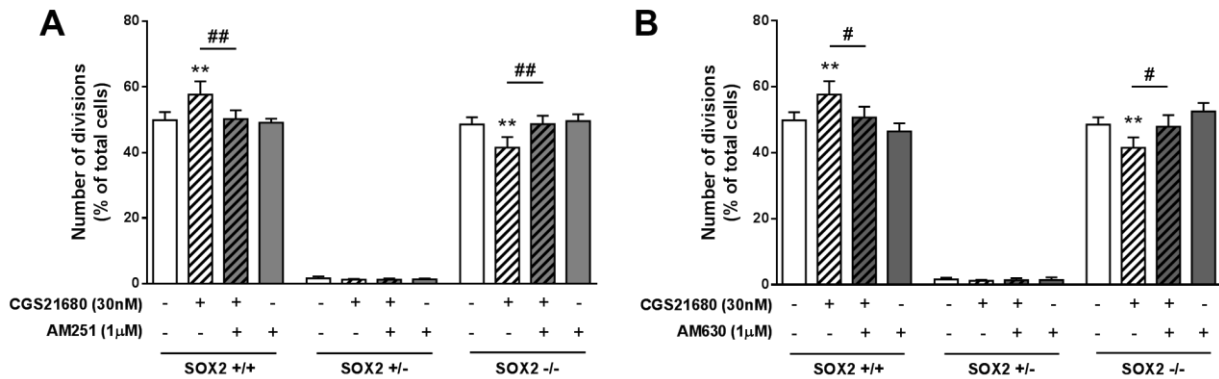


Figure 25 – CB₁R or CB₂R blockade impairs the A_{2A}R-induced self-renewing of DG cells. Bar graphs represent the number of each type of cell divisions counted (60 pairs counted *per condition per* experiment) in cultures treated with A_{2A}R agonist CGS21680 (30 nM) and CB₁R antagonist AM251 (1 μM) (**A**) or in cultures treated with A_{2A}R agonist CGS21680 (30 nM) and CB₂R antagonist AM630 (1 μM) (**B**). Data are expressed as the percentage of total cell pairs and are represented as the mean ± SEM. N=6, **p<0.01, by ANOVA using Bonferroni's post test for comparison with control; #p<0.05 and ##p<0.01, by ANOVA using Bonferroni's post test for comparison with CGS21680.

3.3.2.1. CB₁R, CB₂R or A_{2A}R activation induces no significant changes on DG proliferation

We next investigated whether A_{2A}R, CB₁R and CB₂R ligands could modulate DG cells proliferation. For that purpose, DG cells were treated with selective agonists for A_{2A}R, CB₁R and CB₂R for 24h, as similar to SVZ cells. Incorporated BrdU was immunolabeled and positive nuclei were counted (Figure 26A).

Treatment of DG cells with CB₁R (ACEA, 1 μM) and CB₂R (HU-308, 1 μM) selective agonists and A_{2A}R selective agonist (CGS21680, 30 nM) induced no significant increase in the number BrdU-positive cells when compared to control cultures (control: 100.5±1.22%; ACEA

1 μM : $109.7 \pm 18.39\%$; HU-308 1 μM : $102.8 \pm 5.55\%$; CGS21680 30 nM: $110.7 \pm 18.98\%$; N = 5)
(Figure 26B, C).

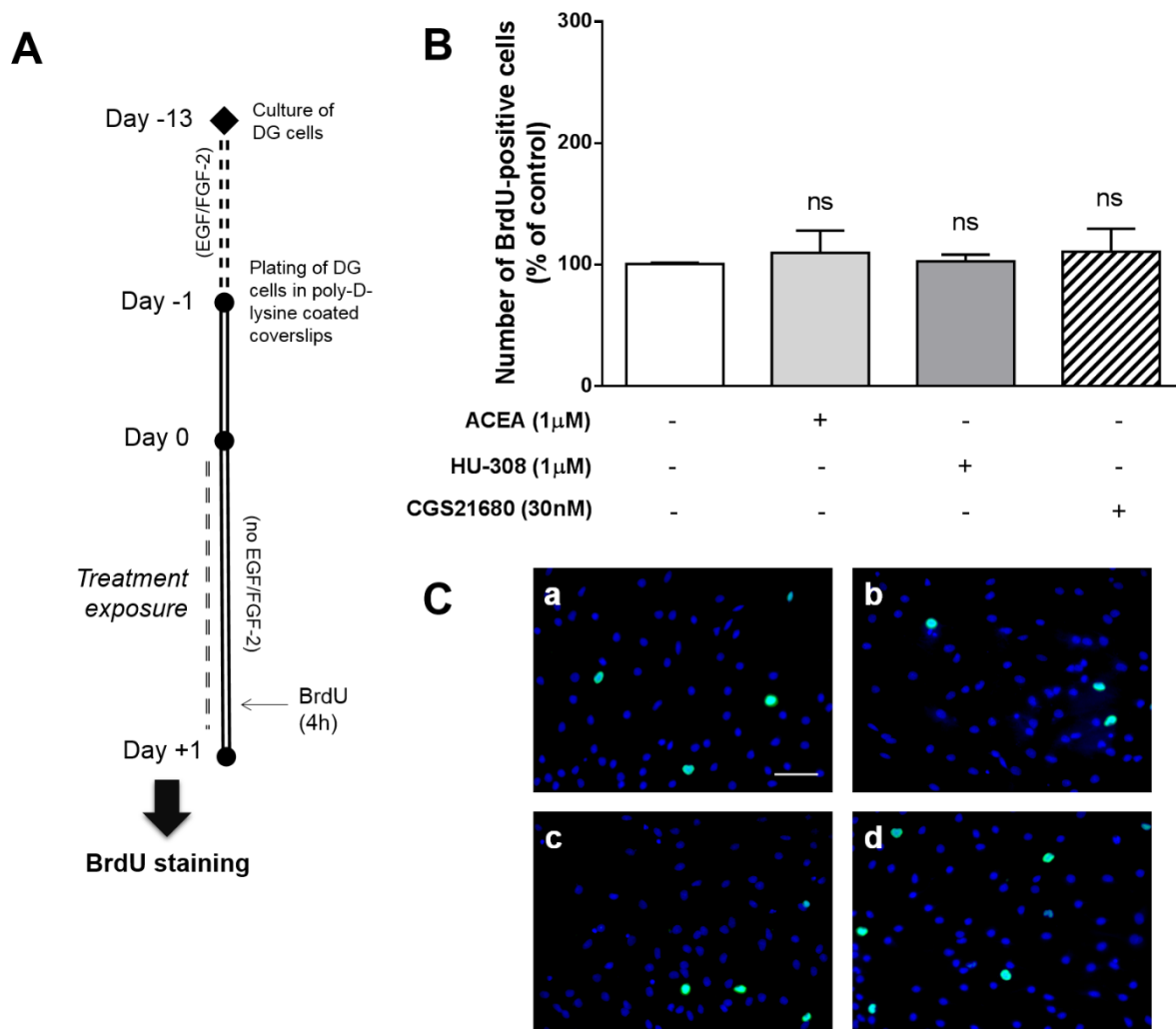


Figure 26 – CB₁R, CB₂R and A_{2A}R activation have no effect on DG cell proliferation.

A. Schematic representation of the experimental protocol. Day 0 represents the day in which neurosphere-derived cells were exposed to pharmacological treatments for the following 24h. **B.** Bar graph depicts the number of BrdU-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean \pm SEM. Control was set to 100%. N=5. **C.** Representative fluorescent digital images of BrdU-positive cell nuclei (green) and Hoechst staining (blue nuclei), in control cultures (a) and in cultures exposed to ACEA 1 μM (b), HU-308 1 μM (c) and CGS21680 30 nM (d). Scale bar = 50 μm .

3.3.2.2. Co-activation of CB₁Rs or CB₂Rs with A_{2A}R increases DG proliferation

To further understand the role of CB₁Rs, CB₂Rs and A_{2A}Rs in DG proliferation, it was assessed if a combined administration of either CB₁Rs or CB₂Rs selective agonists with the A_{2A}R selective agonist promoted any change in the number of BrdU-positive DG cells. Similar

to SVZ treatment, DG cells were treated with either ACEA or HU-308 for 30 minutes prior to CGS21680 treatment and allowed to grow for further 24h in the presence of the ligands.

Surprisingly, we observed that a significant increase in the number of BrdU-positive nuclei was obtained in cultures incubated with both CB₁R selective agonist (ACEA, 1 μM) and A_{2A}R selective agonist (CGS21680, 30 nM) when compared to control (control: 100.5±1.22%; ACEA 1 μM+CGS21680 30 nM: 204.3±43.98%; N = 5, ***p<0.001) (Figure 27). In the same way, the co-incubation of CB₂R selective agonist (HU-308, 1 μM) and A_{2A}R selective agonist (CGS21680, 30 nM) induced a significant increase in the number of BrdU-positive cells as compared to control cultures (control: 100.5±1.22% HU-308 1 μM+CGS21680 30 nM: 205.5±25.50%; N = 5, ***p<0.001) (Figure 28).

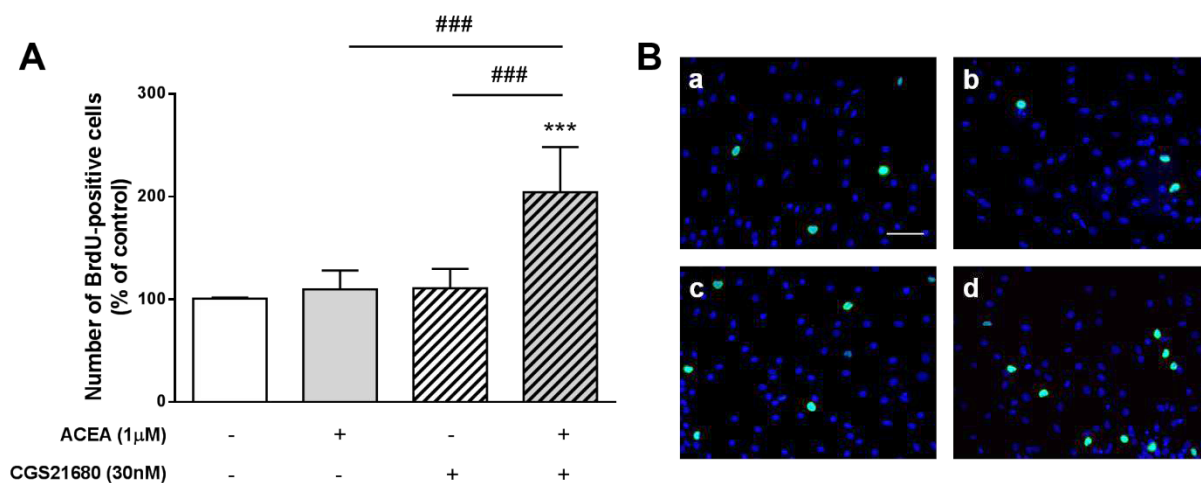


Figure 27 – CB₁R activation coupled with A_{2A}R activation induces DG proliferation.

A. Bar graph depicts the number of BrdU-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean ± SEM. Control was set to 100%. N=5. ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control; ###p<0.001, by ANOVA using Bonferroni's post test for comparison with ACEA+CGS21680. **B.** Representative fluorescent digital images of BrdU-positive neurons (green) and Hoechst staining (blue nuclei), in control cultures (a) and in cultures exposed to ACEA 1 μM (b), CGS21680 30 nM (c) and ACEA 1 μM+CGS21680 30 nM (d). Scale bar = 50 μm.

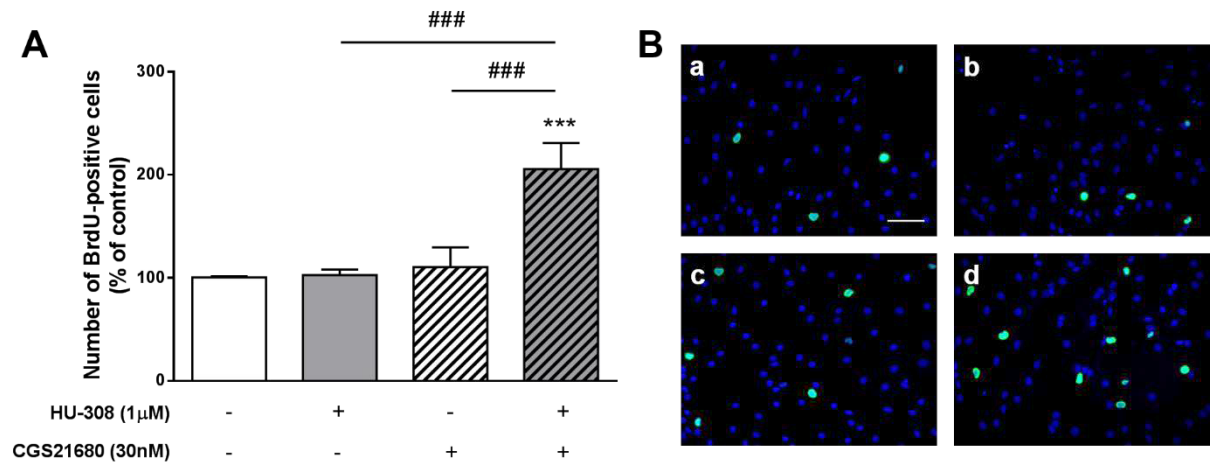


Figure 28 – CB₂R activation coupled with A_{2A}R activation induces DG proliferation.

A. Bar graph depicts the number of BrdU-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean \pm SEM. Control was set to 100%. N=5. ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control; ###p<0.001, by ANOVA using Bonferroni's post test for comparison with HU-308+CGS21680. **B.** Representative fluorescent digital images of BrdU-positive neurons (green) and Hoechst staining (blue nuclei), in control cultures (a) and in cultures exposed to HU-308 1 μ M (b), CGS21680 30 nM (c) and HU-308 1 μ M+CGS21680 30 nM (d). Scale bar = 50 μ m.

3.3.3.1. CB₁Rs, CB₂Rs or A_{2A}Rs activation promotes DG neuronal differentiation

To determine the role of A_{2A}R, CB₁R and CB₂R ligands on DG neuronal differentiation, DG cells were treated with selective agonists for A_{2A}Rs, CB₁Rs and CB₂Rs for 7 days. Cells were then stained for NeuN, a marker for mature neurons (Figure 29A).

Treatment of DG cells with CB₁R (ACEA, 1 μ M) and CB₂R (HU-308, 1 μ M) selective agonists and A_{2A}R selective agonist (CGS21680, 30 nM) induced a significant increase in the number of NeuN-positive cells when compared to control cultures (control: 99.99 \pm 0.02%; ACEA 1 μ M: 173.6 \pm 12.37%; HU-308 1 μ M: 151.9 \pm 8.16%; CGS21680 30 nM: 158.2 \pm 8.69%; N = 5, ***p<0.001) (Figure 29B, C).

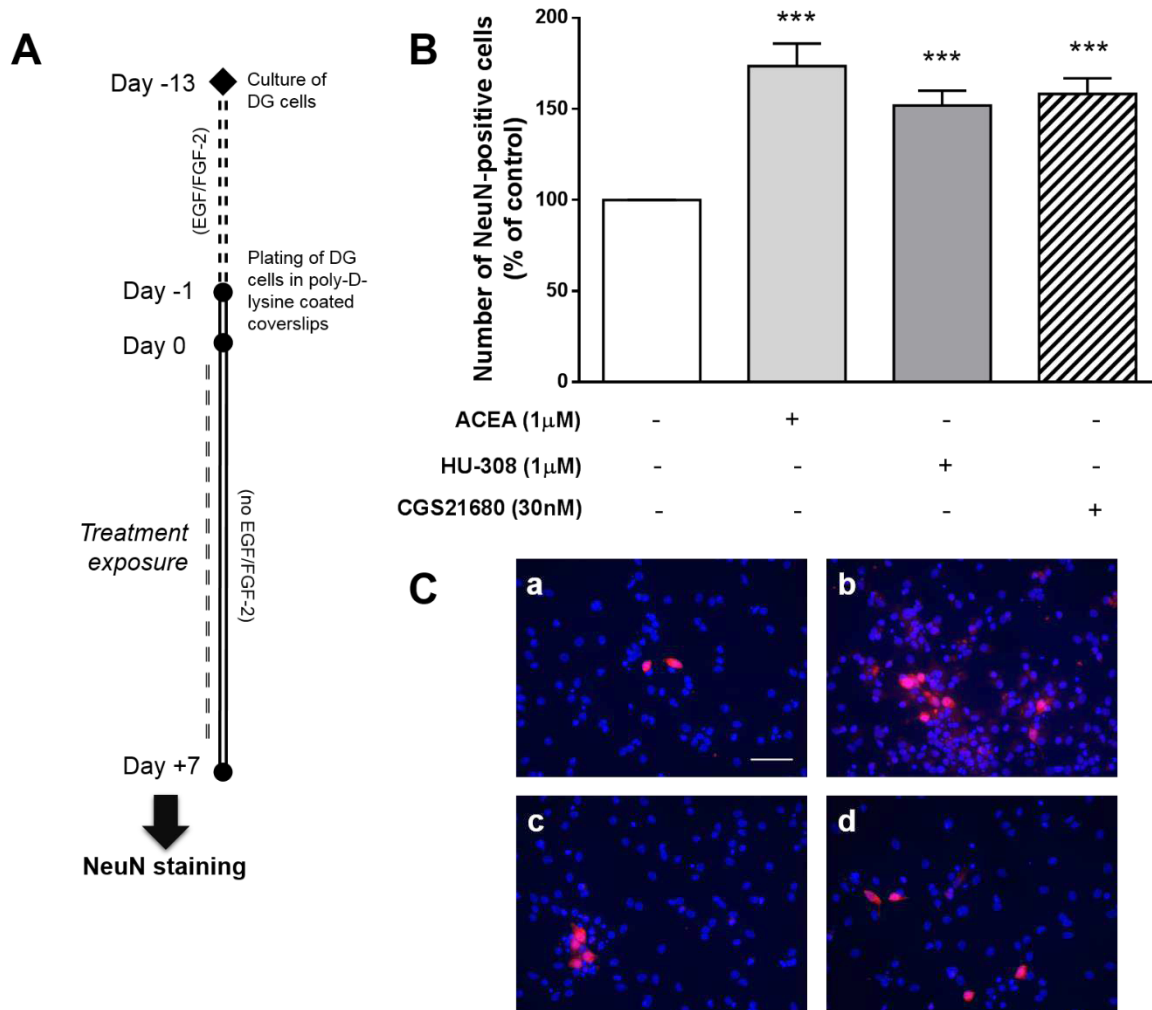


Figure 29 – CB₁R, CB₂R and A_{2A}R activation induces DG neuronal differentiation.

A. Schematic representation of the experimental protocol. Day 0 represents the day in which cells were exposed to pharmacological treatments for the following 7 days. **B.** Bar graph depicts the number of NeuN-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean \pm SEM. Control was set to 100%. N=5. ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control. NeuN: Neuronal Nuclei. **C.** Representative fluorescent digital images of NeuN-positive neurons (red) and Hoechst staining (blue nuclei), in control cultures (a) and in cultures exposed to ACEA 1 μ M (b), HU-308 1 μ M (c) and CGS21680 30 nM (d). Scale bar = 50 μ m.

3.3.3.2. Co-activation of CB₁R or CB₂R with A_{2A}R increases DG neuronal differentiation

To study whether co-incubation with CB₁R or CB₂R selective agonists with the A_{2A}R selective agonist could change neuronal differentiation, DG cells were treated with either ACEA (1 μM) or HU-308 (1 μM) for 30 minutes prior to CGS21680 (30 nM) treatment and then cells were grown for 7 days in the presence of the ligands.

It was observed that, in the presence of both CB₁R selective agonist (ACEA, 1 μM) and A_{2A}R selective agonist (CGS21680, 30 nM), there was an increase in the number of NeuN-positive cells when compared to control cultures (control: 99.99±0.02%; ACEA 1 μM+CGS21680 30 nM: 147.8±13.44%; N = 5, ***p<0.001), with no synergistic effect when comparing to ACEA (1 μM) and CGS21680 (30 nM) treatments alone (ACEA 1 μM: 173.6±12.37%; CGS21680 30 nM: 158.2±8.69%) (Figure 30). Similarly, the co-administration of CB₂R selective agonist (HU-308, 1 μM) and A_{2A}R selective agonist (CGS21680, 30 nM) induced an increase in the number of NeuN-positive cells as compared to control cultures (control: 99.99±0.02%; HU-308 1 μM+CGS21680 30 nM: 146.1±28.51%; N = 5, **p<0.01), also, with no synergistic effect when comparing to HU-308 (1 μM) and CGS21680 (30 nM) treatments alone (HU-308 1 μM: 151.9±8.16%; CGS21680 30 nM: 158.2±8.69%) (Figure 31).

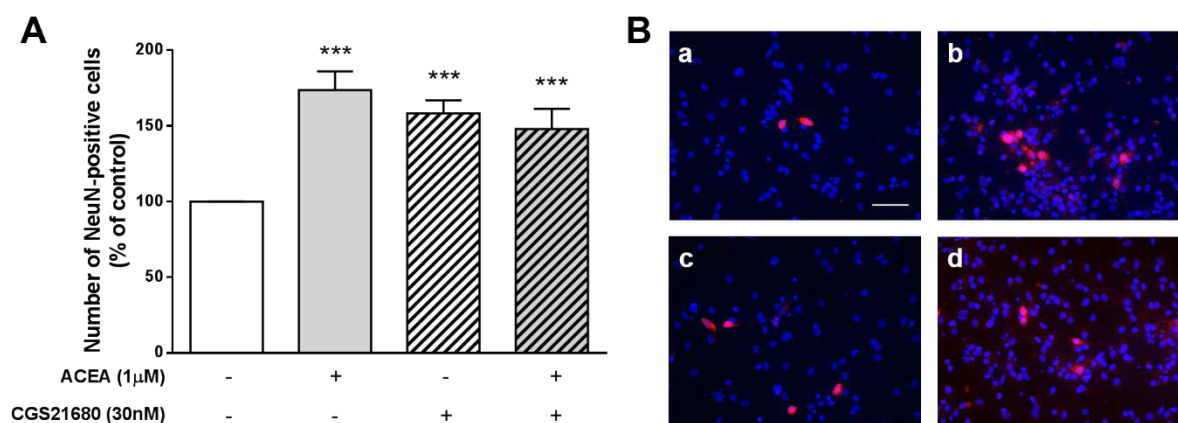


Figure 30 – CB₁R activation coupled with A_{2A}R activation induces DG neuronal differentiation. A. Bar graph depicts the number of NeuN-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean ± SEM. Control was set to 100%. N=5. ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control. **B.** Representative fluorescent digital images of NeuN-positive neurons (red) and Hoechst staining (blue nuclei), in control cultures (a) and in cultures exposed to ACEA 1 μM (b), CGS21680 30 nM (c) and ACEA 1 μM+CGS21680 30 nM (d). Scale bar = 50 μm.

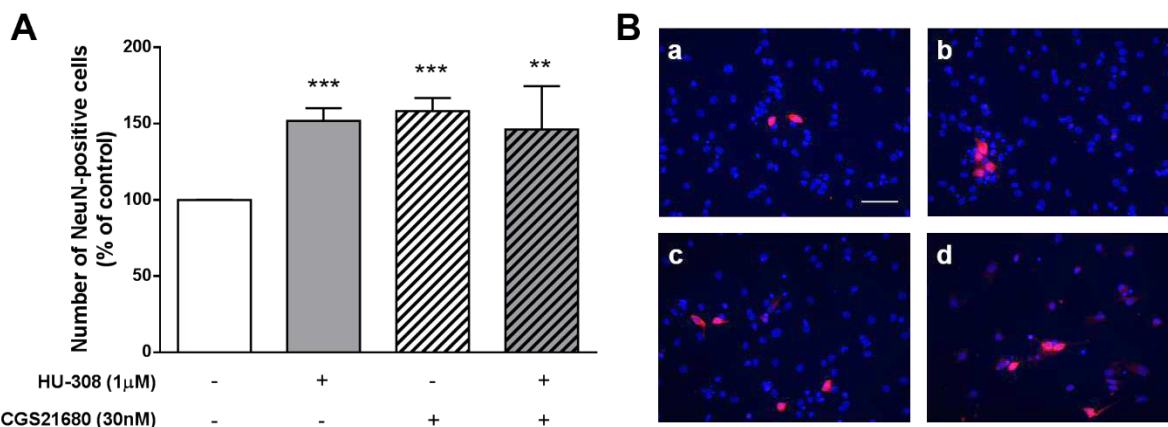


Figure 31 – CB₂R activation coupled with A_{2A}R activation induces DG neuronal differentiation.

A. Bar graph depicts the number of NeuN-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean \pm SEM. Control was set to 100%. N=6. **p<0.01 and ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control. **B.** Representative fluorescent digital images of NeuN-positive neurons (red) and Hoechst staining (blue nuclei), in control cultures (a) and in cultures exposed to HU-308 1 μM (b), CGS21680 30 nM (c) and HU-308 1 μM+CGS21680 30 nM (d). Scale bar = 50 μm.

3.3.3.3. A_{2A}R role in DG neuronal differentiation promoted by CB₁R and CB₂R

Considering that treatment with A_{2A}R, CB₁R and CB₂R selective agonists promoted DG neuronal differentiation, we further wanted to clarify the putative role of A_{2A}R in DG neuronal differentiation promoted by CB₁R and CB₂R activation. For that purpose, similar to SVZ cultures, DG cells were treated with A_{2A}R antagonist (ZM241385, 50 nM) for 30 minutes prior to treatment with CB₁R (ACEA, 1 μM) or CB₂R (HU-308, 1 μM) selective agonists, and then further co-incubated for 7 days.

The results show that the increase in the number of NeuN-positive DG cells promoted by CB₁R activation is lost in the presence of the A_{2A}R selective antagonist ZM241385 (50 nM) (control: 99.99 \pm 0.02%; ACEA 1 μM: 173.6 \pm 12.37%; ACEA 1 μM+ZM241385 50 nM: 117.9 \pm 10.42%; ZM241385 50 nM: 102.1 \pm 12.96%; N = 5, ###p<0.001) (Figure 32A, C). In the same way, the increase in the number of NeuN-positive DG cells promoted by CB₂R activation is partially blocked in the presence of an A_{2A}R selective antagonist ZM241385 (50 nM) (control: 99.99 \pm 0.02%; HU-308 1 μM: 151.9 \pm 8.16%; HU-308 1 μM+ZM241385: 117.4 \pm 8.09%; ZM241385: 102.1 \pm 12.96%; N = 5, *p<0.05 and ###p<0.001) (Figure 32B, D).

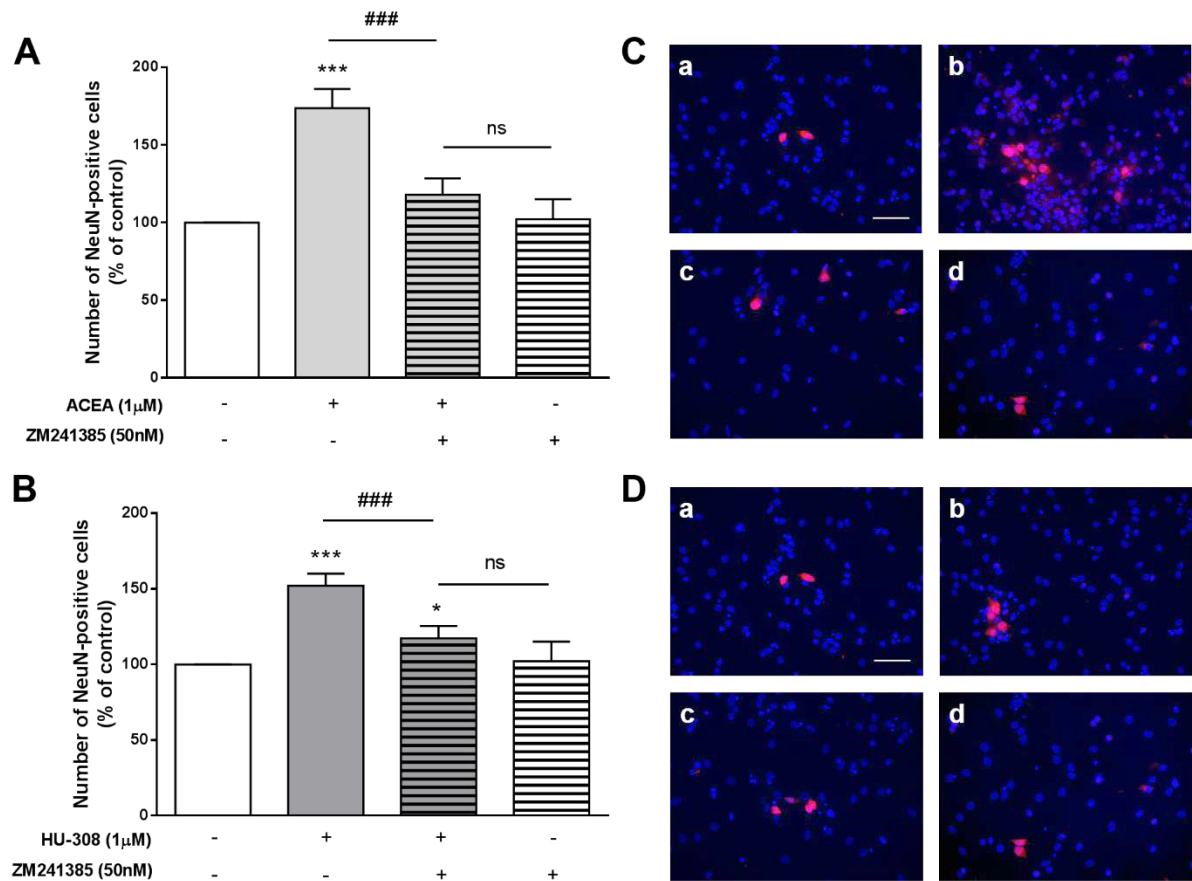


Figure 32 – A_{2A}R blockade impairs the CB₁R- and CB₂R-induced DG neuronal differentiation.

A, B. Bar graph depicts the number of NeuN-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean \pm SEM. Control was set to 100%. N=5. * p <0.05 and *** p <0.001, by ANOVA using Bonferroni's post test for comparison with control; ### p <0.001, by ANOVA using Bonferroni's post test for comparison with ACEA and HU-308, respectively. ns: non-significant. **C, D.** Representative fluorescent digital images of NeuN-positive neurons (red) and Hoechst staining (blue nuclei), in control cultures (C,D-a) and in cultures exposed to ACEA 1 μ M (C-b), ACEA 1 μ M+ZM241385 50 nM (C-c) and ZM241385 50 nM (C-d); HU-308 1 μ M (D-b), HU-308 1 μ M+ZM241385 50 nM (D-c) and ZM241385 50 nM (D-d) Scale bar = 50 μ m.

3.3.3.4. A_{2A}R-mediated DG neuronal differentiation is dependent on CB₁R_s and CB₂R_s

In line with the previous results, it was next investigated if CB₁R_s or CB₂R_s blockage coupled with A_{2A}R activation would influence DG neuronal differentiation. To address this question, DG cells were treated with either CB₁R (AM251, 1 μ M) or CB₂R (AM630, 1 μ M) selective antagonists 30 minutes prior to treatment with A_{2A}R selective agonist CGS21680 (30 nM), and co-incubated for further 7 days.

Remarkably, we found that, treatment with CB₁R antagonist AM251 (1 μM) blocked the effect mediated by A_{2A}R agonist CGS21680 (30 nM) on DG neuronal differentiation, resulting in a decrease number of NeuN-positive cells (control: 99.99±0.02%; CGS21680 30 nM: 158.2±8.69%; CGS21680 30 nM+AM251 1 μM: 91.44±16.67%; AM251 1 μM: 81.30±16.62%; N = 5, ###p<0.001) when compared to CGS21680 treatment (Figure 33A, C). In the same way, cultures co-treated with both A_{2A}R agonist (CGS21680, 30 nM) and CB₂R antagonist (AM630, 1 μM), it was observed a decrease in the number of NeuN-positive DG cells (control: 99.99±0.02%; CGS21680 30 nM: 158.2±8.69%; CGS21680 30 nM+AM630 1 μM: 102.0±14.43%; AM630 1 μM: 110.0±12.89%; N = 5, ###p<0.001) when comparing with CGS21680 treatment alone (Figure 33B, D).

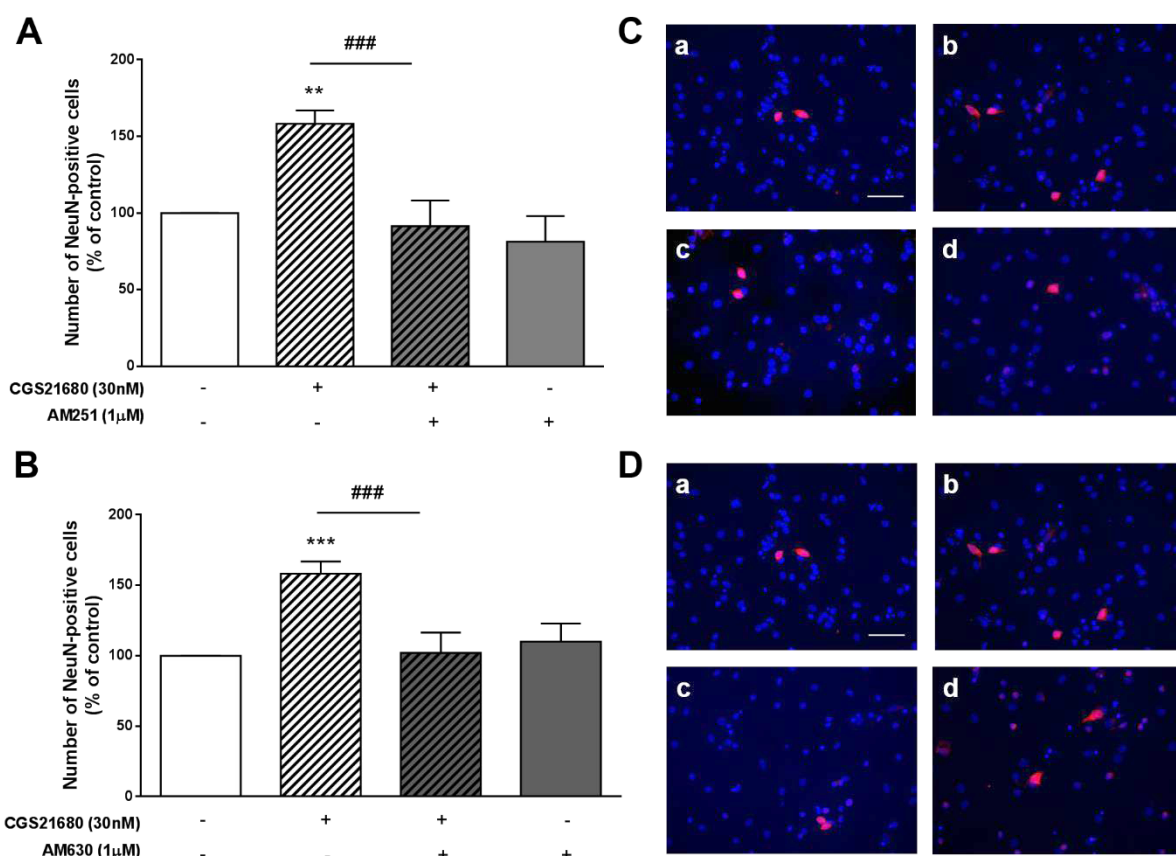


Figure 33 – CB₁R and CB₂R blockade impairs A_{2A}R-induced neuronal differentiation of DG cells. **A, B.** Bar graph depicts the number of NeuN-positive cells. Values were normalized to the control mean for each experiment and are represented as the mean ± SEM. Control was set to 100%. N=5. **p<0.01 and ***p<0.001, by ANOVA using Bonferroni's post test for comparison with control; ###p<0.001, by ANOVA using Bonferroni's post test for comparison with CGS21680. **C, D.** Representative fluorescent digital images of NeuN-positive neurons (red) and Hoechst staining (blue nuclei), in control cultures (C,D-a) and in cultures exposed to CGS21680 30 nM (C-b), CGS21680 30 nM+AM251 1 μM (C-c) and AM251 1 μM (C-d); CGS21680 30 nM (D-b), CGS21680 30 nM+AM630 1 μM (D-c) and AM630 1 μM (D-d) Scale bar = 50 μm.

3.4. Discussion

ECS and adenosinergic system are two major players in the CNS. Although several studies have been trying to understand the role of eCBs and adenosine in the regulation of many physiological processes, gaps still persist in our knowledge regarding endocannabinoid action and A_{2A}Rs role in neurogenesis.

Here, it was examined the effects of CB₁R, CB₂R and A_{2A}R activation on SVZ- and DG-derived cells, given that SVZ and DG represent the two main neurogenic niches of the adult brain. These neurogenic niches are packed with NSPCs that can be modulated by a wide array of factors. In fact, stem cell-based therapy is emerging as a thriving area for regenerative medicine with a lot of clinical applications. Nonetheless, this approach is severely hindered by the shortage of knowledge about the proneurogenic potential of certain compounds.

Previous studies have mainly focused their attention on the effects played by cannabinoids in neurogenesis, mostly regarding proliferation experiments, and not many studies have evaluated the adenosinergic action on neurogenesis. In fact, it was already shown that CB₁Rs, CB₂Rs and A_{2A}Rs are expressed in SVZ and DG tissue and neurosphere-derived cells. Particularly, NSPCs express a functional endocannabinoid system and are able to produce both AEA and 2-AG and, also, that are targeted by cannabinoids to promote neurosphere generation and NSPC proliferation^{168,204,205}. In the same way, SVZ-derived primary neurospheres were found to express A_{2A}Rs and it is well established that hippocampal neurons express the adenosinergic system²²⁹. Moreover, purinergic signaling was shown to regulate NSPCs expansion and neurogenesis²³⁰.

It was firstly observed that CB₁R, CB₂R or A_{2A}R agonist treatment had no effect on SVZ cell fate. Also, a combined activation of either CB₁R or CB₂R with A_{2A}R also had no effect on SVZ cell fate, resulting in no changes in the percentages of Sox2+/+ or Sox2-/- cell pairs. These results are not in complete accordance with past studies where it was seen that CB₁R activation altered fate specification to a more stem state either in SVZ-derived cells¹⁹¹ or in mice cortical NSPCs²³¹.

Moreover, it was observed that CB₁Rs activation promoted an increase in cell proliferation of SVZ cultures, an effect that persisted when cultures were co-incubated together with A_{2A}Rs selective agonist. In fact, it was previously shown an inhibition of NSPCs proliferation in both hippocampus and SVZ of CB₁R- or CB₂R-KO animals, or in animals treated with CB₁R or CB₂R-selective antagonists^{204,232}. Additionally, these data are in accordance with several data showing that CB₁R activation promotes SVZ cell proliferation^{184,188,191,233}. However, we found that CB₂R activation did not induce cell proliferation, although others have shown that CB₂R activation could promote proliferation in embryonic cell lines, in SVZ

neurosphere cultures and in the SVZ of young mice^{206,207}. Importantly, the effect mediated by CB₁Rs on proliferation was blocked by an A_{2A}R selective antagonist, further indicating a role for A_{2A}Rs on CB₁R-mediated stimulation of SVZ proliferation. In fact, Stafford and colleagues showed that A_{2A}R activation led to cell cycle arrest in primary SVZ-derived NSPCs, therefore inhibiting the proliferation, potentially playing a role in the initiation of neuronal differentiation²²⁹, which strongly correlates with the results previously described.

I have also found that SVZ neuronal differentiation significantly increases upon CB₁Rs or CB₂Rs activation. Regarding the role of CBRs in neuronal differentiation, the existing data also appears to be conflicting. In fact, the work done by Compagnucci et al.¹⁸⁹ and Xapelli et al.¹⁹¹ also shows that CB₁R activation is important to promote neuronal differentiation, while Aguado et al.²³⁴ and Gomez et al.²³⁵ show that endocannabinoid treatment promotes an astroglial and/or oligodendroglial differentiation rather than neuronal differentiation. Importantly, I observed that the effect mediated by CB₁R or CB₂R activation in neuronal differentiation remained unaltered when CB₁Rs or CB₂Rs activation was coupled with A_{2A}R agonist treatment. On the other hand, this effect was blocked by the presence of an A_{2A}R selective antagonist, although A_{2A}R activation *per se* did not change neuronal differentiation. These data suggests a pivotal role of A_{2A}Rs in SVZ neuronal differentiation promoted by CB₁Rs and CB₂Rs activation.

Regarding DG, it was observed that CB₂R and A_{2A}R activation induced self-renewing divisions, resulting in a significant increase in the percentages of Sox2+/+ daughter cell pairs. This is in accordance with the work of Palazuelos et al.²⁰⁴ where they show that in murine embryonic cortical NSPCs of CB₂R KO mice a reduced self-renewal rate was observed and that CB₂R activation increased neural progenitor self-renewal *in vitro*. Moreover, I also found that the increase in self-renewal observed in DG cells was maintained when CB₁Rs or CB₂Rs were co-activated with A_{2A}Rs. Importantly, CB₂R-induced increase in self-renewal was blocked with an A_{2A}R selective antagonist. In the same way, the effect mediated by A_{2A}Rs activation on self-renewal was blocked by the presence of CB₁R or CB₂Rs selective antagonists. These very interesting data further suggests evidences for a putative crosstalk between these two systems.

Most studies relating A_{2A}Rs role in neurogenesis generally focus their attention on the action of caffeine (a non-selective A₁R and A_{2A}R antagonist) in hippocampal-related neurogenesis. Several studies show that acute treatment with caffeine has beneficial effects on cell proliferation or overall neurogenesis in the hippocampus, whereas chronic administration has the opposite effect¹³¹⁻¹³³. On the other hand, the role of cannabinoids on hippocampal neurogenesis has been intensively studied to try to understand how eCBs can affect neurogenesis and subsequently shape learning and memory processes. Evidences

show that the number of proliferating cells in the DG is reduced in CBRs-deficient mice and that CBRs play major roles in NSPCs proliferation, morphogenesis and differentiation^{139,204}.

I observed no changes on DG proliferation when cultures were treated with CB₁R, CB₂R and A_{2A}Rs selective agonists. These results are not in complete accordance to past studies where it was seen that chronic administration with HU-210 (a drug that has a high affinity for both CB₁Rs and CB₂Rs) enhanced both proliferation and survival of cells in the rat DG²³² or stimulated BrdU incorporation and neurospheres formation²⁰⁵. Rubio-Araiz and co-workers also demonstrated that both CB₁R and CB₂R agonists, stimulate the proliferation of primary murine cortical neurospheres¹⁹⁰. Aguado and her colleagues showed the involvement of the CB₁R in NSPCs proliferation and neurogenesis induced by excitotoxic injury¹⁹³. Moreover, Palazuelos and his colleagues showed that chronic administration of the CB₂R selective agonist also exhibits proliferative-enhancing effects²⁰⁷. Nevertheless, when using combined treatments for DG cells using CB₁R and A_{2A}R selective agonists or CB₂R and A_{2A}R selective agonists a surprisingly significant increase on DG cell proliferation occurred, further indicating a possible synergistic effect between CBRs and A_{2A}Rs.

Moreover, DG neuronal differentiation is stimulated by CB₁Rs, CB₂Rs or A_{2A}R agonist treatment and this effect persists when DG cells are treated in a combined manner with the CB₁Rs or CB₂Rs and A_{2A}Rs selective agonists. These results are in accordance with Compagnucci and colleagues work in which AEA enhances cell differentiation toward a neuronal lineage via a CB₁R-dependent mechanism¹⁸⁹ and, also, with Wolf and collaborators work in which, in the absence of CB₁Rs, cell proliferation was increased and neuronal differentiation reduced²³⁶. Furthermore, Avraham and co-workers showed that administration of CB₂R agonist promoted the differentiation of human NSPCs²⁰⁸. Interestingly, a study from Lin and colleagues suggested that activation of the purinergic system may act as a proliferation signal for NSPCs, and thereby serve as a negative regulator for neuronal differentiation²³⁰.

Similarly to what happens in DG cell fate decision, CB₁R- or CB₂R-induced increase in neuronal differentiation was blocked with an A_{2A}R selective antagonist. Likewise, the effect mediated by A_{2A}R activation on neuronal differentiation was blocked with CB₁R or CB₂R antagonists. Since proliferation and differentiation are mutually excluding cellular processes¹³⁹, CB₁Rs/CB₂Rs and A_{2A}R signaling may be modulating these two processes in an entirely different fashion, being more active either in the cell cycle or in the differentiation stage. Thus, these data further supports the idea that A_{2A}Rs play a crucial part in the regulation of DG cell proliferation and on the DG neuronal differentiation promoted by CB₁Rs and CB₂Rs activation.

Divergent effects are observed throughout literature concerning the role of eCBs on neurogenesis which may partly be explained from the use of different pharmacologic approaches. These discrepancies can also be explained by differences in the study design (treatment schedule or time-points where measurements are performed) or the animal species

or gender used, which may induce misleading interpretations¹⁹¹. So, all these readout parameters may account for the differences seen in this study from the combined activation of CB₁Rs and CB₂Rs with A_{2A}Rs.

In fact, several studies resort to the use of non-selective agonists/antagonists for CBRs to evaluate the effects of eCBs on neurogenesis. For example, Abboussi and colleagues showed that chronic administration of the synthetic CB₁R/CB₂R non-selective agonist WIN55,212-2 to rats during adulthood had no effect on the number of immature neurons in the DG, however, administration during adolescence decreased the number of immature neurons²³⁷. Using the same compound, Marchalant and colleagues found that a low, continuous administration significantly increased neurogenesis in aged rats²³⁸. Moreover, Aguado and colleagues, through an *in vitro* approach, also showed that WIN55,212-2 promoted an increase in the number of neural progenitors and generation of neurospheres¹⁶⁸. Moreover, the use of SR141716A, a CB₁R and TRPV1 antagonist, has been shown to increase cell proliferation in the DG and the in lateral ventricles of mice by Jin and colleagues and this was reproducible in both wild type and CB₁R, but not TRPV1, KO mice¹⁸⁸. Overall, these findings demonstrate that compound selectivity is also really important when studying neurogenesis and that multiple receptors may be responsible for the cannabinoid-induced effects on neurogenesis, which may account for the complexity/variety of general results.

The adenosinergic signaling was shown to be important for some cannabinoid-mediated effects, either CB₁Rs or CB₂Rs^{219,223}. In particular, A_{2A}Rs were shown to be involved in the addictive-related behavior promoted by cannabinoid administration, having a role in the counteraction of the rewarding effects promoted by cannabinoids²¹⁸. In fact, A_{2A}Rs could play a role in CB₁Rs/CB₂Rs functions via receptor heterodimerization or signaling cascades cross-linking^{175,177,214}. Certainly, the occurrence of a putative crosstalk between A_{2A}Rs and CB₁Rs and/or CB₂Rs presents as a valid deduction, either by a structural interaction (formation of heterodimers), as others have suggested in other brain areas^{175,176}, or by crosstalk at downstream signaling^{104,179}. Considering that CB₁Rs, CB₂Rs and A_{2A}Rs act through signaling pathways, such as ERK1/2 and p38 MAPK, which are common, heterodimerization of receptors or signaling crossing can emerge as solid hypotheses to explain the differences observed in these experiments, among single versus combined treatments with CB₁Rs or CB₂Rs and A_{2A}Rs selective agonists.

Taking together the current data is, as far as I know, the first study showing a putative crosstalk between A_{2A}Rs and CB₁Rs and/or CB₂Rs in neurogenesis.

CHAPTER 4

Closing Remarks

4. Conclusion and Future Perspectives

Generally, NSPCs are spatially located within very particular regions of the brain (at SVZ and DG regions) sensitive to different stimulus and are endowed with quiescent pools of cells that keep continuously self-renewing. This makes them ideal therapeutical targets for stem cell-based therapies and regenerative medicine platforms in order to enhance brain repair in brain trauma or pathological situations.

There is considerable evidence from previous studies suggesting that eCBs can control a variety of physiological processes in the adult brain. However, the current available data about eCB action on neurogenesis is still quite controversial. Furthermore, the role of adenosine and especially of A_{2A} Rs is not known. In fact, the adenosinergic signaling was shown to be important for some cannabinoid-mediated effects and that A_{2A} Rs could play a role in CBRs functions via receptor heterodimerization or signaling cascades cross-linking.

Altogether, this work dissects the modulatory role of A_{2A} Rs on CBR-mediated SVZ and DG neurogenesis, suggesting a possible crosstalk between the cannabinoid and adenosinergic systems.

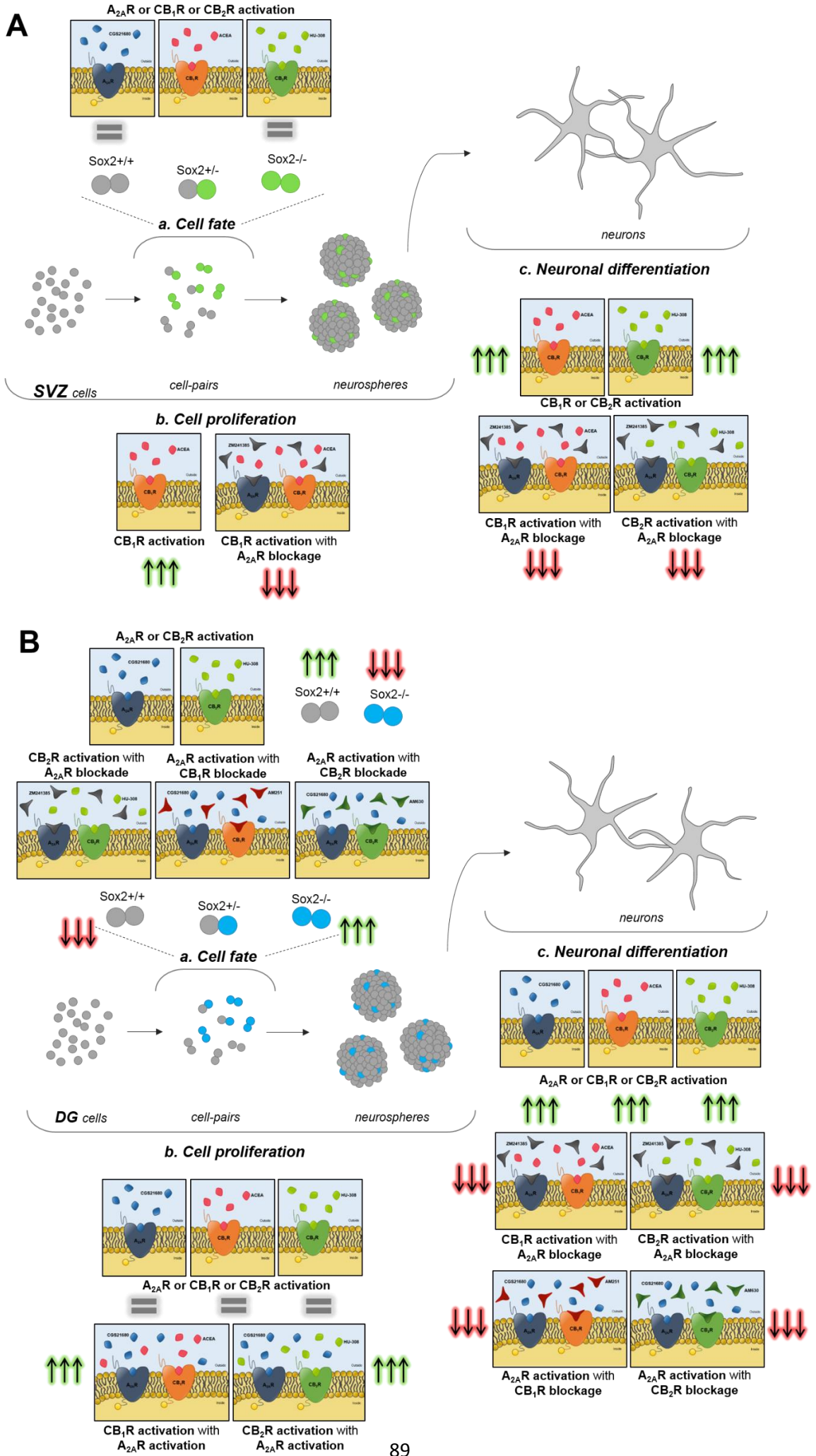
In summary, it was shown that treatment with CB_1 R, CB_2 R or A_{2A} R selective agonists induces no effect on the SVZ cell-fate. However, in the DG either CB_2 R or A_{2A} R activation could promote self-renewing divisions. Moreover, the effect on self-renewal mediated by CB_2 R activation in the DG was blocked with an A_{2A} R selective antagonist, while the effect mediated by A_{2A} R activation was blocked with either CB_1 R or CB_2 R antagonists. Additionally, CB_1 R activation was shown to promote SVZ cell proliferation and this effect was blocked by an A_{2A} R selective antagonist. Similarly, treatment with either CB_1 R or CB_2 R selective agonists promoted an increase in SVZ neuronal differentiation, an effect that was blocked by A_{2A} R blockade. Interestingly, while activation *per se* of CB_1 R, CB_2 R or A_{2A} R did not induce cell proliferation of DG cells, co-incubation of CB_1 R and A_{2A} R selective agonists or with CB_2 R and A_{2A} R selective agonists resulted in an increase in DG cell proliferation. Furthermore, CB_1 R, CB_2 R or A_{2A} R activation promoted DG neuronal differentiation. Finally, the effect mediated by CB_1 R or CB_2 R was inhibited in the presence of an A_{2A} R selective antagonist, while the effect mediated by A_{2A} R activation was blocked in the presence of CB_1 R or CB_2 R selective antagonists.

Taking all this into account, it is possible to speculate about the potential regulatory role of A_{2A} Rs in neurogenesis: cross-signaling interaction between A_{2A} Rs and CB_1 Rs or CB_2 Rs may complement the complexity of endocannabinoid signaling and influence some cell stage-specific features in order to promote neurogenesis. Taken together, a putative crosstalk

between the adenosinergic and cannabinergic systems may indeed exist, either by the formation of heterodimers or by crosstalk at downstream signaling.

Therefore, further analysis is ultimately necessary to define the importance of this putative interaction of A_{2A} Rs and CB_1 Rs and/or CB_2 Rs, either by forming heterodimers or by interacting at the signaling level, and to clearly define how it may influence the neurogenic process. Indeed, this modulation of A_{2A} Rs and CB_1 Rs and/or CB_2 Rs of neurogenesis may be occurring at distinct levels of action or at unique temporal time-windows, resulting in different roles depending on which neurogenic stage NSPCs are in (e.g. different modulatory actions on cell fate decisions compared to cell survival upon integration in the new circuitry). In fact, eCBs and A_{2A} Rs may be essential for mediating particular aspects of some neurogenic stages. The role of cannabinoids on neurogenesis is complex and the contribution of A_{2A} Rs action to the net-effect still remains unclear and requires additional investigation to elucidate their full proneurogenic potential. Indeed, further *in vivo* studies will be required to comprehensively understand the role of adenosine, in particular A_{2A} Rs, in the modulation of CBRs-mediated effects on neurogenesis.

Therefore, based on the established knowledge and my data, the main findings of this work can be illustrated as described in Figure 34.



◀ **Figure 34 – General schematic representation of the main findings of this work.** **A.** SVZ neurogenic niche: activation of CB₁Rs, CB₂Rs or A_{2A}Rs induced no changes in the cell fate (**a**); CB₁Rs activation promoted a significant increase in SVZ cell proliferation, an effect blocked by an A_{2A}R selective antagonist (**b**); similarly, activation of either CB₁Rs or CB₂Rs promoted a significant increase in SVZ neuronal differentiation, an effect abolished by A_{2A}R blockade (**c**). **B.** DG neurogenic niche: activation of CB₂Rs or A_{2A}Rs induced a significant increase in self-renewal divisions and the effect mediated by CB₂Rs was blocked by an A_{2A}R selective antagonist and, similarly, the effect mediated by A_{2A}Rs was blocked by CB₁R or CB₂R antagonists (**a**); co-activation of CB₁Rs and A_{2A}Rs or CB₂Rs and A_{2A}Rs promoted an increase on DG cell proliferation, although activation of CB₁Rs, CB₂Rs or A_{2A}Rs *per se* had no effect on cell proliferation (**b**); DG neuronal differentiation was significantly increased by the activation of CB₁Rs, CB₂Rs or A_{2A}Rs; the effect mediated by CB₁Rs or CB₂Rs was blocked by an A_{2A}R selective antagonist and, similarly, the effect mediated by A_{2A}Rs was blocked by CB₁R or CB₂R antagonists (**c**).

CHAPTER 5

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