



Università di Foggia

Dipartimento di Studi umanistici. Lettere, Beni Culturali, Scienze della
Formazione

Corso di Dottorato in “Neuroscience and Education”

Ciclo XXXVI

Giuseppe Costantino

Unravelling neuroinflammatory pathways by using in vitro and in vivo models

PhD Thesis

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Anno Accademico 2022/2023

Preface

My work during my PhD course in “Neuroscience” at the University of Foggia is divided in two different parts. During the first period I carried out an *in vitro* study, pointing my attention on the analysis of the role of metabotropic glutamate receptors (mGlu) belonging to class II, mGlu2 and mGlu3 receptors, on the function of the blood brain barrier (BBB) under inflammatory conditions. To this purpose, we used cell populations that are components of the BBB of human origin. More specifically, we focused on the role of microglia in the modulation of BBB function. This part of the research was carried out at the University of Catania, in the Department of Biomedical and Biotechnological Sciences, laboratory of Pharmacology under the supervision of Prof. MA Sortino and SF Spampinato. The results obtained will be the object of the first part of this thesis.

During the second part of my PhD I moved to *in vivo* studies, being involved in a project in which we analyzed the complex mechanisms underlying ASD behavioural dysfunctions, by using the BTBR strain, an idiopathic model of ASD. In this context, we firstly investigated neurochemical and biomolecular alterations underlying such behavioural impairments and, subsequently effects in brain regions crucially involved in ASD, and after we evaluated the effect of a possible preventive treatment, on the core ASD symptoms and on the modulation of neuroinflammatory components of ASD by using β -carotene administration during pregnancy.

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1. *In Vitro* studies

1.1 Introduction

1.1.1 Blood Brain Barrier (BBB)

The blood-brain barrier (BBB) is a highly specialized structure with the main role of preserving the central nervous system (CNS). The BBB is composed of specialized endothelial cells that, together with other components of the neurovascular unit (NVU), astrocytes, pericytes, neurons and basement membrane, preserve the correct environment in the CNS, limiting the entry of potentially damaging blood-borne agents, metabolites, drugs and immune cells (Zenaro et al., 2017).

Changes in the BBB have been documented in different CNS pathological states. These include brain tumors (Wolburg et al., 2012), brain trauma (Chodobski et al., 2011), ischemia (Gursoy-Ozdemir et al., 2012) and epilepsy (Fabene et al., 2013). Furthermore, various conditions related to neurodegenerative disorders such as Alzheimer's disease (AD), Parkinson's disease (Bartels et al., 2008) and amyotrophic lateral sclerosis (ALS) (Garbuzova-Davis et al., 2011) are characterized by structural and functional changes of the BBB.

The endothelial cells are the primary cellular component of the BBB forming a continuous endothelium barrier. They are organized as a continuous intercellular layer forming tight junctions at cell-cell contacts that lack fenestration. For these reasons molecules and solutes movements through the BBB are selective (Garcia et al., 2014; Garg et al., 2008). The entry of water-soluble solutes, such as glucose or amino acids, is guaranteed by the presence of specific transporters that regulate their access, while their expulsion from the CNS is warranted by the presence of efflux pumps. Many BBB transporters (such as those for glucose) are facilitated diffusion systems, which allow molecules following their concentration gradient. Others, however, prevent the entry of exogenous substances into the CNS; among these, one of the most relevant is the P glycoprotein. Changing the activity of these transporters may be a possible approach in some pathologies.

The BBB is in close contact with some components of the immune system and reduces the entry of peripheral cellular components into the CNS. Endothelial cells in physiological conditions express a low level of adhesion molecules for leukocytes, which reduces the access of immune cells into the CNS, while preserving a certain level of immuno-surveillance. This is guaranteed by the few immune cells that pass the barrier but remain in the perivascular space, separated from the CNS by the presence of glia (Beurel et al., 2014). In pathological conditions, the expression of adhesion molecules increases and the connections between endothelial cells decrease, facilitating the entry of immune cells into the CNS. It is therefore clear that a completely efficient barrier is essential to preserve the CNS.

1.1.2 Endothelial cells: Tight Junctions and Adherens Junctions

Endothelial cells interact among each other through two important types of junctions: the tight junctions (TJs) and the adherens junctions (AJs), that regulate the paracellular permeability. TJs and AJs are involved in maintaining junctional integrity, limiting the permeability only to small lipophilic molecules and gases. The TJs constitute the "diffusion barrier" between adjacent endothelial cells and therefore have the task of regulating the flow of ions and mainly hydrophilic molecules through the paracellular space. In physiological conditions, molecules that have a molecular weight (MW) greater than 180 kDa cannot pass through TJs (Mitic & Anderson, 1998). If observed with the electron microscope, the TJs appear as a set of linear, long and parallel fibrils, which surround the cell (Pinto da Silva & Kachar, 1982). The assembly of these fibrils, important for the adequate functioning of TJs, is a very dynamic phenomenon, and seems to increase when endothelial cells are subjected to harmful stimuli (Gumbiner, 1993). Preservation of TJs is regulated by specific transmembrane proteins and among these, claudins, occludin and junctional adhesion molecules (JAMs) are particularly important. They are associated with cytoplasmic proteins (ZO-1, ZO-2 and ZO-3) connecting the TJs to the actin of the cytoskeleton and to the AJs. Alteration of specific phosphorylation of amino acid residues and their

translocation from the cell membrane lead to a loss of junctional integrity and a consequent increase in paracellular permeability. Furthermore, functional modifications of these proteins are observed in case of alteration of specific phosphorylation of amino acid residues (such as serine, threonine and tyrosine) (Sandoval & Witt, 2008). One of the most important AJs that contributes to the impermeability of the BBB is represented by the protein VE-cadherin.

Occludin (MW: 60 kDa) is present at the level of TJs in its phosphorylated form (Sakakibara A. et al., 1997). It binds to zonula occludens proteins (ZO-1, ZO-2 and ZO-3) and therefore interacts with the actin cytoskeleton (Fanning et al., 1998; Furuse et al., 1994; Haskins et al., 1998). This event has been shown to be important for maintaining the functionality and properties of the barrier (Balda et al., 1996). Accordingly, the presence of occludin at the membrane level is correlated with an increase in electrical resistance and a reduction in paracellular permeability (Balda et al., 1996).

Claudins (MW 20 to 27 kDa) are a family of membrane proteins that are among the major structural components of TJs. In the cerebral endothelium, the most represented are claudin-3 and claudin-5 (Morita et al., 1999; Wolburg et al., 2012), that are recognized to give an important contribution in the formation of TJs and in maintaining the integrity of the barrier. Claudins support the efficiency of TJs thanks to their ability to homodimerize and heterodimerize the two cis- and trans-forms (Piehl et al., 2010; Piontek et al., 2008; J. Zhang et al., 2010). Claudin-5 plays an important role in the organization of the barrier, which derives from its ability to associate with other claudins of neighboring endothelial cells (Greene et al., 2019).

The ZO-1 (zonula occludens-1) protein is a protein of the MAGUK (membrane-associated guanylate kinase) family together with ZO-2 and ZO-3. In TJs it acts as a scaffold for the organization of transmembrane proteins, thus connecting them to the actin cytoskeleton. It is also able to recruit various signal transduction pathways (González-Mariscal et al., 2000). ZOs share three main regions among

themselves: an SH3 domain, which is responsible for binding to signaling proteins and cytoskeletal elements, a guanylate kinase, which has the task of catalyzing the ATP-dependent transformation of GMP (guanosine-mono-phosphate) in GDP (guanosine-di-phosphate), and finally a PDZ domain, which binds the carboxy-terminal regions of transmembrane proteins, controlling their correct distribution in space (Zlokovic, 2008b). The localization and interactions of these proteins are essential for the assembly of claudin, occludin and JAMs filaments in TJs and for the anchoring of the entire complex to the actin cytoskeleton (Fanning et al., 2007; Utepbergenov et al., 2006). ZO-1 is a phosphoprotein and its phosphorylation state influences its localization and interactions with other TJs.

The adherence junctions have an important role in the immunosurveillance. The quote of leukocyte infiltration depends on the expression of AJs, in particular VE-cadherin, and adhesion molecules, such as ICAM-1, VCAM-1 and P- and E-selectins. In physiological conditions, ECs at the BBB express very low levels of these leukocyte adhesion molecules (Daneman, Zhou, Agalliu, et al., 2010; Wettschureck et al., 2019). Their expression is regulated at transcriptional levels, and can be induced by soluble inflammatory mediators, such as cytokines and chemokines (Wettschureck et al., 2019). When highly expressed, adhesion molecules and AJs may be more easily engaged by their leukocyte counterpart, leading to a signaling cascade that terminates with the transendothelial migration and leukocyte access into the CNS.

Endothelial cells are supported by astrocytes and pericytes that, releasing trophic factors, induce specific BBB properties (Hill et al., 2014). The components of the NVU are assisted by two extracellular matrix structures, recognized as basement membranes (BMs) (named also “basal lamina”). The inner vascular BM surrounds the endothelial cells, separating them from pericytes, while the outer parenchymal BM is interposed between the pericytes and the astrocytes endfeet (Balabanov & Dore-Duffy, 1998; del Zoppo et al., 2006; Sorokin, 2010). The BMs play an important role in the maintenance of the close connection between the elements of NVU (Yurchenco & Patton, 2009), constitute a source of growth

factors and adhesion receptors and contribute to the regulation of the BBB permeability (Daneman & Prat, 2015; Zlokovic, 2008a).

1.1.3 Pericytes

Pericytes are contractile cells that are responsible for the production of the extracellular matrix and proteins that contribute to the regulation of BBB homeostasis (Bohannon et al., 2021; Gautam et al., 2020). They contribute to the stability of the BBB through the release of trophic factors that help maintain vascular stability and regulate the diameter of the capillaries, thus also indirectly controlling blood flow (Winkler et al., 2011). The interaction between pericytes and endothelial cells contributes to the formation of TJs and counteracts transcytosis and the expression of cell adhesion molecules. It has been recently reported that the claudin-5, the glucose transporter Glut-1 and the efflux proteins (P-glycoprotein) were still expressed in animals increased BBB leakage, increased ICAM-1 expression and focal leukocyte extravasation (Mäe et al., 2021). Accordingly, others reported that pericytes control TJs alignment and transcytosis across the BBB (Bell et al., 2010; Daneman, Zhou, Agalliu, et al., 2010; Quaegebeur et al., 2010). Pericytes seem to be also involved in the regulation of the rapid control of neurovascular coupling. Thank to their contractile activity, they can actively relax or contract to modify the cerebral blood flow (CBF) in response to neuronal activity [29,30]. Pericytes are essential in BBB development and in the maintenance of brain microcirculation, as they directly contribute to the formation of the BM (Bergers & Song, 2005) and secrete angiogenic-promoting factors (such as VEGF, angiopoietin-1, TGF- β , PDGF-BB) that stabilize BBB functions (Ribatti et al., 2011). A lack of pericytes leads to hyperplasia, abnormal vascular morphogenesis (Hellström et al., 2001) and an incomplete function of the BBB during embryogenesis, as they inhibit the expression of proteins that increase vascular permeability, such as Angiopoietin 2 and the plasmalemma-vesicle-associated protein (Plvap), and leukocyte adhesion molecules on ECs (Daneman, Zhou, Kebede, et al., 2010). There are important interactions between pericytes and astrocytes, the absence of pericytes

influences astrocyte polarization, so affecting astrocytic endfeet/endothelial interaction (Geranmayeh et al., 2019). On their side, astrocytes facilitate pericyte migration, proliferation and contact with the endothelial layer (Nakagawa et al., 2009; Y. Yao et al., 2014).

1.1.4 Astrocytes

Astrocytes have an important role in supporting neuronal activity, regulate the balance of extracellular K⁺, and control the recovery and recycling of neurotransmitters. They support the activity of the BBB, surrounding the endothelial cells with their peduncles and releasing trophic factors (Capaldo & Nusrat, 2009; Garg et al., 2008). Astrocyte interaction with endothelium is important for the functions of the BBB. While pericytes are more important in the early stages of BBB property induction, astrocytes are more involved in the maturation and maintenance of barrier properties (Obermeier et al., 2016; Spampinato et al., 2019). Some studies have tried to explore how the depletion of astrocytes may affect BBB properties in animal models. The laser ablation of a single astrocyte may be not sufficient to induce BBB damage (Kubotera et al., 2019), while a complete astrocyte ablation in adult mice induced an early and sustained BBB dysfunction, consequent to alterations in TJs functions (Heithoff et al., 2021). Astrocytes regulate many functions of the barrier, modulate TJs, induce the expression and location of transporters such as P-glycoprotein and GLUT1 glucose transporter and influence the expression of specific enzymes (Cheslow & Alvarez, 2016). The barrier functions are modulated by factors released by astrocytes, such as angiopoietin 1, fibroblast growth factor (FGF), and neurotrophic factor derived from glial cells (GDNF), useful for promoting vascular stabilization (Broux et al., 2015). Astrocytes are highly secretory cells, and their secretome include many molecules (Dowell et al., 2009; Harada et al., 2015). For example, the morphogen sonic hedgehog (Shh) and retinoic acid strengthen endothelial junctional tightness (S. Liu et al., 2019; Mizze et al., 2013). Astrocytes are the main producer of Angiotensinogen (Stornetta et al., 1988), that is a central player in the regulation of blood pressure, although its role in the

reinforcing or weakening of BBB integrity is controversial: it can both reduce (Wosik et al., 2007) and increase barrier permeability (Takane et al., 2017). Astrocytes produce also the main vascular trophic factor, VEGF-A, that promotes angiogenesis and endothelial cell proliferation, differentiation, and survival during brain development (Esser et al., 1998). On the other hand, high levels of VEGF-A may induce BBB disruption in pathological conditions (Spampinato et al., 2019). In general, factors produced by astrocytes may both induce and disrupt barrier properties, depending on the disease context, age, and surrounding microenvironment. Astrocytes are in direct physical contact with endothelial cells; astrocytic terminal processes, called endfeet, cover the external surface of the endothelium and surround the vasculature (Abbott, 2002). Endfeet are thought to cover up to 99% of the cerebrovascular surface (Mathiisen et al., 2010), participate in the regulation of angiogenesis and the formation of cell-to-cell junctions (Cabezas et al., 2014). The importance of astrocyte support in BBB function varies during different pathological conditions of the CNS. In fact, when glial cells are involved, astrocytes can lose their functions or modify their secreting activity to produce inflammatory cytokines. All these events contribute to a change in the interaction between glia and neurons which leads to a modification of the functionality of the endothelial barrier. Finally, the interactions between endothelial and immune cells can also modify the activity of astrocytes, for example reactive astrocytes in inflammatory conditions can form their own junctions to limit leukocyte infiltration into the CNS, supplying the BBB barrier when endothelial functions are compromised (Horng et al., 2017).

1.1.5 Microglia and Neuroinflammation

Neuroinflammation is an important event that influences the severity and progression of neurodegenerative and psychiatric disorders, such as Alzheimer's disease (AD), Parkinson's disease (PD), Huntington's disease (HD), multiple sclerosis (MS), stroke, and others (Lyman et al., 2014). Numerous cytokines such as interleukin-23 (IL-23), IL-12 (Bennett, 2013), IL-1 β , and IL-6 are involved in neurodegenerative processes. Several studies in the past years have

demonstrated that different cell types such as oligodendrocytes, microglia, astrocytes (Domingues et al., 2016), both local and circulating lymphocytes, different dendritic cell subsets (Colton, 2013), and endothelial cells (Combes et al., 2012) are involved in neuroinflammation.

More recent evidence suggests a role for microglia in the control of BBB function. Microglia cells, the resident immune cells in the CNS, pathological mediators or traumatic events may be quickly activated in response to CNS insult. The nature of microglial responses can be different: they can remove cellular debris in an attempt to contribute to cellular repair or can accelerate the induction of the neuroinflammatory response (G. I. Caruso et al., 2021; Merlo et al., 2022; Merlo, Luaces, et al., 2020).

Microglia cells were described for the first time by the neuroanatomist Pio Del Rio-Hortega in 1919, who also noticed the mesodermal origin and phagocytic capability of these cells (Sierra et al., 2016). In the last few years, various studies in mice confirmed that microglia are mesodermal-derived cells, originating from erythromyeloid progenitors of the yolk-sac, which begin to colonize the developing CNS before the formation of the cerebrovascular network, from embryonic day 8.5 until the complete development of the BBB (Ginhoux et al., 2010; Gomez Perdiguero et al., 2015). In a similar way, microglial precursors in humans infiltrate the CNS primordium starting at around 4.5 gestational weeks (Menassa & Gomez-Nicola, 2018). The differentiation of progenitors in microglia is induced by two transcription factors, Pu.1 and Irf8, and their expression distinguishes microglia from other tissue-resident macrophages (Hagemeyer et al., 2016; Kierdorf et al., 2013). In the adult brain, microglia maintenance depends on the constant activation on their surface of Colony-stimulating Factor-1 Receptor (CSF-1R) by interleukin (IL)-34 and CSF-1, which are continuously released by neurons and astrocytes, respectively. Consequent to the pharmacological blockade of CSF-1R there is a rapid microglia depletion (Chitu et al., 2016; Elmore et al., 2014). Microglia are long-living and self-renewing cells that, in inflammatory conditions, can proliferate quickly through a selected clonal

expansion (Tay et al., 2017). When there is a BBB damage in CNS diseases, circulating monocytes can infiltrate the CNS and acquire a microglial-like phenotype (Ginhoux et al., 2013). In physiological condition, in brain and spinal cord, microglial cells take on a ramified morphology with long and motile processes that continuously scan the surrounding environment (Nimmerjahn et al., 2005). Beyond immune surveillance, microglia are involved in several and different functions in the CNS: microglia activity is in fact essential for a correct synaptic pruning during development (Paolicelli et al., 2011), synaptic plasticity (George et al., 2016; Schafer et al., 2013; Sipe et al., 2016), neuronal programmed cell death and neurogenesis in adults (Bachstetter et al., 2011; Sierra et al., 2010) as well as learning and memory processes (Parkhurst et al., 2013).

After an environmental negative stimulus or injury, microglia acquire an ameboid morphology, with much shorter ramifications and larger cell bodies (Morrison et al., 2017). In several pathological conditions, microglia clear the CNS through phagocytosis of protein aggregates, e.g., amyloid- β (Hu et al., 2023; Mandrekar et al., 2009), pathogens (Cockram et al., 2019), immune cells that have crossed the BBB and damaged or stressed-but-viable neurons in a process named “phagoptosis” (G. C. Brown & Neher, 2014). Activated microglial cells are capable to release a multitude of pro-inflammatory cytokines (IL-1 α and IL-1 β , Tumor Necrosis Factor (TNF)- α), chemokines (CCL2, CCL5, CXCL1, MIP-1), proteases (e.g., matrix metalloproteases, MMPs) and reactive oxygen species (ROS), which, while participating to the immunological response to injury, can also compromise BBB integrity if their secretion is protracted for a long time. Similarly, microglial cells are also able to secrete molecules that promote tissue healing and reduce inflammation (e.g., VEGF, TGF β -1, IL-10) (L.-R. Liu et al., 2020). Consequently, over the years, it has become increasingly clear that the behavior of microglia shows a high phenotypic plasticity and that microglia can modify the activity over time during disease. Starting from the early 2000s, microglial functional status has been classified as pro- or anti-inflammatory, named M1 and M2, respectively

(Paolicelli et al., 2022). This classification derives from *in vitro* studies on T-helper cells that, after the administration of the appropriate stimuli, acquire a pro- (Th1) or anti- (Th2) inflammatory activity (Paolicelli et al., 2022). In a similar way, microglia have been divided into these two categories, based on the expression of specific markers that indirectly implied a detrimental (M1) or protective (M2) phenotype (Paolicelli et al., 2022). Nowadays, the use of single-cell-transcriptomic and proteomic techniques demonstrated that the M1/M2 definition is not completely suitable for microglia, whose dynamic behavior is strongly dependent on the pathological and environmental conditions (Paolicelli et al., 2022).

1.1.6 Microglia role in BBB

As highlighted previously, the influence of microglial cells on physiological BBB functions has been only recently explored. During embryogenesis, microglial cells seem to be associated with the developing blood vessels in the retina and during neurogenesis in the cortex and contribute to vascular development (Penna et al., 2021). The absence of microglia in PU.1^{-/-} mice or pharmacological depletion with the CSF-1R inhibitor PLX5622 cause a choroidal vascular atrophy, the disorganization and dysfunction of retinal pigment epithelial cells and an altered expression of angiogenic growth factors (Dixon et al., 2021; Fantin et al., 2010). Confirming this, *Csf1op/op* mice, which produce a mutated inactive form of CSF-1, show an early postnatal defective retinal vasculature, with altered arterial–venous patterning and reduced ramification (Y. Kubota et al., 2009).

Microglia as part of the NVU interact with endothelial cells of the BBB, and although do not seem to play an essential role in BBB formation and stability, they modulate BBB functions both in physiological and pathological conditions (da Fonseca et al., 2014). In the adult CNS, microglia establish transient and dynamic physical contact with the neuro-vasculature; accordingly, it has been recently described a specific microglial population, called capillary-associated microglia (CAM), that interacts with microvessels in a continuous way (Bisht et al., 2021). However, the existence of a “juxtavasculature” microglial population had

been described much earlier (Grossmann et al., 2002). CAMs represent about 30% of the whole microglial population in adult mice and are localized preferentially along the vessels where astrocyte endfeet coverage is absent, thus suggesting the need for further investigations on their direct contribution to the regulation of microvessel diameter and, as a consequence, of CBF (Bisht et al., 2021; Mondo et al., 2020). It was shown that microglia depletion through the CSF-1R inhibitor PLX3397 results in an increased capillary diameter by 15% (Bisht et al., 2021), and a recent study by Csaszar and colleagues demonstrated that microglia make simultaneous contact with vascular elements (ECs, mural cells and astrocytes) and neurons to regulate neurovascular responses (Császár et al., 2022). Moreover, it has been demonstrated *in vitro* that the presence of microglia in co-culture with ECs promotes barrier properties in the endothelium, inducing the expression of the protein of TJs, like ZO-1 and occludin (Mehrabadi et al., 2017).

When neuroinflammation underlies the pathological condition, both astrocytes and microglia affect BBB stability.

Recruitment and migration of immune cells through the BBB is in fact facilitated by cytokines and chemokines released by reactive astrocytes (Spampinato, Merlo, et al., 2022), whereas microglial cells may promote BBB breakdown through the release and production of inflammatory cytokines, ROS and NO (Mahad & Ransohoff, 2003) that induce endothelial ICAM-1 expression (Huber et al., 2006) and destabilize the BBB by damaging supporting astrocytes (Haruwaka et al., 2019). In addition, reactive microglia contribute to the leaky BBB observed in these diseases through downregulation of TJs such as Claudin-5, occludin, and ZO-1 (Kleinberger et al., 2017; Obermeier et al., 2013).

Interestingly, however, both astrocytes and microglia can stabilize endothelial cells at the BBB, releasing trophic factors and molecules like IL-4, IL-13 and arginase implied in angiogenesis and barrier tightness (Z. Chen & Trapp, 2016; Spampinato et al., 2019). In particular, the involvement of microglia in increased

TJs stability in early phases of inflammatory conditions has been demonstrated (Haruwaka et al., 2019). Furthermore, microglia depletion can facilitate BBB permeability (Halder & Milner, 2019). Instead, during a sustained inflammation microglia can induce BBB damage (Haruwaka et al., 2019). In fact, the increased cyclooxygenase 2 activity in reactive microglia can facilitate the release of inflammatory cytokines (TNF, IL1, IL6) and the activation of pathways that contribute to BBB destabilization (Akundi et al., 2005; Ronaldson & Davis, 2020; Y. Yang & Rosenberg, 2011). Endothelium-derived factors including nitric oxide are known to suppress microglial activation and exert anti-inflammatory actions suggesting a bidirectional control between endothelium and microglia (Katusic & Austin, 2016). From this it follows that interacting with the cellular constituents of the BBB, microglia may play a dual role, initially protecting BBB integrity, but enabling its enhanced permeability during prolonged inflammation (Haruwaka et al., 2019).

A recent study investigated the role of microglia in a BBB model *in vitro*. Microglial cells were co-cultured with astrocytes and endothelial cells and the modulation of S1P receptor after an inflammatory stimulus was evaluated. S1P receptors were stimulated with BAF-312 (siponimod) a drug commonly used for MS. The drug facilitated microglial migration towards endothelial/astrocyte co-cultures, through the activity of the metalloprotease 2 (MMP2). Microglia actively cooperated with astrocytes in the maintenance of endothelial barrier stability: in the triple co-culture, selective treatment of microglial cells with BAF-312 significantly prevented cytokines' effects on the endothelial barrier, suggesting a potential additional effect of the drug in its pharmacological efficacy when used in the treatment of multiple sclerosis (Spampinato, Costantino, et al., 2022).

1.1.7 Microglia in a Chronic Inflammatory Condition: Alzheimer's Disease

Alzheimer's disease (AD) is the principal cause of memory loss and cognitive decline in the world. The pathogenic markers of the disease are A β plaques and hyperphosphorylated tau tangles, which lead to neuronal degeneration, synaptic loss and gliosis. The main target in AD are neurons. The progressive cognitive decline and memory impairment are caused by the reduction in synaptic functions and neuronal degeneration. A β have a direct action on neurons, where it modifies the activity of NMDA glutamate receptors, NA⁺/K⁺ ATPases, insulin receptors and integrins (Abbott, 2002; DiChiara et al., 2017; Texidó et al., 2011; Zhao et al., 2008) and can also modify neuronal membrane structure, altering its permeability and excitability (Merlo, Spampinato, et al., 2020). Amyloid- β (A β) has long been shown to be critical in Alzheimer's disease pathophysiology. Microglia contributes to the earliest responses to A β buildup, by direct interaction through multiple receptors. Microglial cells operate A β clearance and trigger inflammatory/regenerative processes that take place in the long years of silent disease progression that precede symptomatic appearance. But in time and with aging, the fine balance between pro- and anti-inflammatory activity of microglia deranges, negatively impacting its A β -clearing ability. Furthermore, in recent years, microglial activation has proven to be much more complex than the mere dichotomic pro/anti-inflammatory polarization previously accepted. Microglia can display a wide spectrum of phenotypes, which can even be mixed. On these bases, it is evident that while pharmacological intervention aiding microglia to prolong its ability to cope with A β buildup could be extremely relevant, its feasibility is hampered by such high complexity, which still needs to be completely understood (Merlo, Spampinato, et al., 2020). Furthermore, A β oligomers can be internalized in neurons, directly contributing to the generation of ROS (G. Caruso et al., 2019; Cheignon et al., 2018). In addition, also BBB impairment has been proposed as one of the principal elements in the pathogenesis of AD. Cerebrovascular diseases that cause an altered vascular perfusion, cerebral hemorrhage and hypoxia, can be also common comorbidities

in AD, often they constitute the first event (first hit) conducting to progressive A β accumulation (second hit) (Brkic et al., 2015; Sagare et al., 2012).

Usually a compromised BBB, associated with increased permeability, can be considered as a biomarker of the aging process (Fang et al., 2023). In AD patients, it has been demonstrated that the damage of the BBB is already present in the early stages of the disease: imaging studies and postmortem tissues had shown the leakage of blood-borne proteins and immune cells through the BBB in various brain areas, such as the hippocampus, entorhinal cortex, and prefrontal cortex (Nation et al., 2019; Sweeney et al., 2019; van de Haar et al., 2016). Confirming this, brain ECs derived from induced pluripotent stem cells of AD patients showed altered properties, including a dysregulated expression of TJ proteins and an increased barrier permeability (Oikari et al., 2020). Often in AD postmortem brain through Immunohistochemistry, is observed an increased interaction between neuronal and non-neuronal cells, in particular among neurons, ECs, astrocytes and microglia (Kirabali et al., 2020). In AD, the modification occurring at the BBB exasperate A β accumulation in the brain. In ECs of the BBB there are expressed specific transporters that regulate the movement of molecules between the brain and the blood (Ramanathan et al., 2015). Among these transporters, the Receptor for Advanced Glycated End products (RAGE) and the low-density lipoprotein receptor-related protein 1 (LRP-1), are responsible of bind and transport A β (Alemi et al., 2016). In AD their activity is dysregulated (Deane et al., 2004; Deane & Zlokovic, 2007), that lead to A β accumulation in the brain (Ramanathan et al., 2015). The expression of glucose transporter Glut-1 is also severely compromised in microvessels in the brain of AD patients and mouse models (Hooijmans et al., 2007; Merlini et al., 2011), inducing early endothelial dysfunction and barrier leakage (Winkler et al., 2015). The neurovascular uncoupling, described in the disease, conduces to increased permeability to blood-borne solutes and to immune cells, that cause a CNS inflammatory response (Huang et al., 2023). Furthermore, A β may directly modify the BBB functions described: in *in vitro* BBB models, ECs exposed to A β oligomers

undergo to a TJs redistribution (Kook et al., 2012; Wan et al., 2015) and increase the expression of the adhesion molecules ICAM-1 and VCAM-1 (Gonzalez-Velasquez et al., 2008; Marco & Skaper, 2006; Nagababu et al., 2009). Various human postmortem studies indicate that the levels of occludin, ZO-1 and claudin-5 are reduced in the cerebral blood vessels of AD patients (Carrano et al., 2012; Keaney et al., 2015), although exist conflicting results on animal models (Lochhead et al., 2020). The weakening of BBB junctions, which may be directly correlated not just with the decreased levels of the proteins but also with the loss of the balance among all junctional proteins (Schlingmann et al., 2016), can facilitate the access of plasma proteins that get worse inflammation and neurotoxicity (Chodobski et al., 2011). Also, the endothelial overexpression of adhesion molecules such as ICAM-1 and VCAM-1 facilitates leukocyte transmigration, conducting to the aggravation of the inflammatory response. Supporting this, preventing the access of neutrophils to the CNS has been associated with improving of cognitive performance and reduced AD-related pathology in AD mouse models (Zenaro et al., 2015).

BBB properties are strongly influenced by endothelial interaction with the other component of the NVU, in particular with glial cells. In AD, glial cells may both directly and indirectly participate in neuronal loss and in the setup of neuroinflammation, that also influences BBB properties.

Microglia have a time-dependent role in BBB function during development of AD (Gullotta et al., 2023). In the early stages of AD, microglial cells surround and incorporate A β aggregates to limit their deposition (Condello et al., 2015, 2018) and participate actively in the disassembly and digestion of senile plaques (Keren-Shaul et al., 2017). Using genetically modified mouse it has been demonstrated that the complement factors C1q and C3 released by microglia are involved in A β homeostasis (Hong et al., 2016); confirming this, C3R ablation in APP mice have as result a reduction of A β accumulation (Czirr et al., 2017). Also, the triggering receptor expressed on myeloid cells 2 (TREM2) expressed by microglia is involved in A β degradation (P. Yuan et al., 2016).

At later stages A β is accumulated in the brain and, by binding to CD33 receptor on microglia, can reduce his clearance, accordingly, knocking out CD33 in APP/PS1 mice attenuated A β plaque deposition and cognitive decline (Griciuc et al., 2013; Perea et al., 2020). Activation of microglia for long periods have as a consequence the secretion of pro-inflammatory cytokines and the promotion of neurodegeneration (Merlo, Spampinato, et al., 2020), which can be, in fact, the consequence of A β -induced reactivity on glial cells. An important aspect of the microglia role in AD is an abnormal phagocytosis that has been described in AD (Hong et al., 2016). Microglial phagocytosis is regulated by various mechanisms, including the CX3CR1/CX3CL1 interaction and the complement. Alterations in the complement pathway are described already in the early stages of AD (Hong et al., 2016). A β in fact can directly bind and activate the complement system (Webster et al., 1995) inducing the microglia release of the C1p complement factor, which increases the astrocytic release of C3, conducting to synaptic phagocytosis (Hong et al., 2016; Huffels et al., 2023). A constant microglial activation causes an upregulation of inflammatory mediators and cytokines, conducting to an important neuroinflammation and neuronal death (Merlo, Spampinato, et al., 2020; Sajja et al., 2016). A β can directly bind to the microglial TLR4 receptor (Okun et al., 2009) and increase the stimulation of the NLRP3 inflammasome (Hanslik & Ulland, 2020), leading to the activation of caspase-1 and the consequent cleavage and secretion of IL-1 β and IL-18 (Freeman et al., 2017; Gustin et al., 2015).

The crosstalk between microglia and astrocytes is modified in AD (L.-R. Liu et al., 2020). A β induces the microglial release of complement factors and cytokines, which affect astrocytic response. The C1q complement component, released by reactive microglia, induces the astrocytic release of the C3 complement factor (Liddel et al., 2017), which is responsible for neuronal damage, and affects EC functions at the BBB (Propson et al., 2021). In AD animal models, has been showed that microglia release TNF α and TGF- β that modulate astrocytic calcium-ion-dependent signaling and glutamate release (Domercq et al., 2006),

responsible for the beginning of excitotoxicity (Rossi et al., 2005). Microglia-derived IL-1 β stimulates astrocytes to release VEGF, which lead to the degradation of TJ proteins and BBB permeability, in rats, IL-1 β , released by microglia, induced the astrocytes to release VEGF (Q.-S. Wang et al., 2020), while in *in vitro* studies, IL-1 β , derived from co-cultured microglia, suppressed the astrocytic release of Shh (Y. Wang et al., 2014), a factor essential for endothelial properties at the BBB. Finally, microglia-derived TNF α and IL-1 β suppress the expression of astrocytic gap junctions, inducing an altered astrocytic communication, thus facilitating the reactive gliosis observed in AD (Même et al., 2006).

In AD patients and mice models endfeet of astrocytes surrounding A β plaques bear structural alterations (Saavedra et al., 2022). Astrocytes are involved in the generation and clearance of A β (Iliff et al., 2012; Ries & Sastre, 2016; Saavedra et al., 2022), the accumulation of A β in the brain activate the NF-kB pathway in astrocytes, that lead to the complement activation and the release of inflammatory cytokines, chemokines and factors that modify neuron–glia communication and synaptic transmission contributing to the progression of neuroinflammation and BBB damage (Frost & Li, 2017; Garwood et al., 2017; Lian et al., 2015; Liddelow et al., 2017). *In vitro*, astrocytes exposed to A β together with inflammatory cytokines, release VEGF, that contribute to the degradation of the junctional protein claudin-5 and endothelial barrier permeability (Spampinato et al., 2017).

Astrocytes may also have a protective role depending on the disease's phase, they participate in the elimination of neuritic dystrophies in AD mouse models and in AD patients (Gomez-Arboledas et al., 2018) and release factors, such as TGF- β , that increase microglial phagocytic activity and reduce neuronal damage (Gomez-Arboledas et al., 2018; Iram et al., 2016). Astrocytes can also support BBB properties in AD. In response to exogenous trigger stimulus, such as A β accumulation, astrocytes activate heat-shock response, such as Dnajb1, Hspa1a and Hspa1b (Lau et al., 2020), in this way they maintain BBB functions (Thüringer

& Garrido, 2019). The upregulation of the gene Nr4a1 prevents astrocyte inflammatory response, by reducing NF- κ B activity, and contributes to the maintenance of BBB integrity (Pan et al., 2021). Astrocytes modulate also the access in the perivascular space of the leukocytes (Hornig et al., 2017). They can increase the release of protease inhibitors, such as Pcsk1n, that protect the endothelial barrier from the degradative enzymes released by infiltrating leukocytes (Pan et al., 2021). In *in vitro* model astrocytes, after a short A β exposure, regulate the endothelial expression of the adhesion molecule ICAM-1, contrasting leukocyte transmigration (Spampinato et al., 2017). ON the other hand, when the barrier is compromised, astrocytes may directly interact with infiltrated leukocytes, regulating their functions, lymphocytes exposed to A β in the presence of astrocytes showed a Th2 phenotype, increasing the release of BDNF and IL-4 that reduced endothelial barrier damage (Spampinato et al., 2020).

Regarding the interaction with neurons, microglia also have a dual role in the modulation of BBB properties in AD. Microglia activated by a constant inflammatory stimulus, can contribute to BBB leakage (McKim et al., 2018; Zenaro et al., 2017; H. Zhou et al., 2006); on the other hand, in the early phases of the insult, in AD, such as in other CNS pathological conditions, microglia can support the BBB (Haruwaka et al., 2019; Spampinato, Costantino, et al., 2022). In an *in vitro* model, the presence of microglia reduced endothelial permeability induced by an inflammatory stimulus (Spampinato, Costantino, et al., 2022). Accordingly, also in a mouse model, reactive microglia migrated toward the endothelial barrier, promoting the expression of the endothelial junctional protein, claudin-5, thus supporting ECs in reducing BBB leakiness (Haruwaka et al., 2019). Finally, the neurovascular coupling mediated by microglia through the purinergic receptor P2Y₁₂R is involved in AD, thus affecting the cerebrovascular response to neuronal activity, and reducing endothelial barrier properties (Császár et al., 2022; Kenkhuis et al., 2022).

After A β exposure, microglia can also cause a pericyte loss (Huang et al., 2023; Takata et al., 2021). In AD, also the pericyte dysfunction participate in BBB disruption, and a pericyte loss has been described in both AD patients and mice models (Sagare et al., 2013; Sengillo et al., 2013), caused by A β toxicity (Alcendor, 2020). In AD patients, the diameter of small capillaries, in particular in those in the proximity of senile plaques, are restricted and irregular (Kitaguchi et al., 2007), a recent study proposed the involvement of contracted pericytes as the cause of this phenomenon (Nortley et al., 2019). The loss of pericytes may also induce the expression of VEGF, contributing to barrier weakness (Hellström et al., 2001). At the end, A β -activated microglia may upregulate MMP9 released by pericyte (Takata et al., 2011) which, is one of the principal causes of TJs degradation and BBB disruption.

1.1.8 Glutamate Receptors Classification

Glutamate is the most important excitatory neurotransmitter of the mammalian central nervous system (CNS), it has a central role in memory, synaptic plasticity and neuronal development. Glutamate has also a negative role, because a glutamate overstimulation is implicated in neurodegeneration (Ferraguti et al., 2008; Nakanishi & Masu, 1994). Two types of glutamate receptors are described: ionotropic and metabotropic receptors. Ionotropic glutamate receptors are N-Methyl-d-Aspartate (NMDA), α -Amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA) and kainite receptors, all ligand-gated ion channels that promote a rapid excitatory neurotransmission (Dingledine et al., 1999) Metabotropic glutamate receptors (mGluRs) belong to family C G protein-coupled receptors (GPCRs) that, after stimulation, promote G α - $\beta\gamma$ uncoupling, conducting to G α -mediated increase in intracellular second messenger levels, $\beta\gamma$ regulation of ion channels, and furthermore the stimulation of G protein-independent pathways (Gerber et al., 2007; J.-P. Pin et al., 2003). mGluRs may form dimers composed of two subunits cross-linked by a disulphide bridge and dimer formation is important for mGluRs function (Kniazeff et al., 2011). Up to now, eight mGluR subtypes have been identified and characterized and are divided into three

subgroups based on sequence homology and cell signaling activation (Conn & Pin, 1997; J. P. Pin & Duvoisin, 1995). Group I mGluRs are composed by mGluR1 and mGluR5, these receptors are coupled to $G\alpha_{q/11}$, promote the release of Ca^{2+} from intracellular stores (Abdul-Ghani et al., 1996; Dhami & Ferguson, 2006), and are functionally linked to polyphosphoinositide hydrolysis and negatively coupled with KC channels (Abdul-Ghani et al., 1996; Nicoletti et al., 2011). Group II mGluRs are composed by mGluR2 and mGluR3. Finally, group III mGluRs comprises mGluR4, mGluR6, mGluR7 and mGluR8 (Gerber et al., 2007). Both Group II and Group III mGluRs negatively regulate adenylyl cyclase via $G\alpha_i$ but can also activate MAP kinase and PI-3-kinase pathways (Iacovelli et al., 2002; Nicoletti et al., 2011; Niswender & Conn, 2010). Both groups are principally localized presynaptically, and work as autoreceptors to inhibit glutamate or gamma-aminobutyric acid (GABA) release (Schoepp, 2001). Group I mGluRs are mostly located at postsynaptic elements in a perisynaptic zone surrounding the ionotropic receptors, where they modulate neuronal excitability (Lujan et al., 1996; Shigemoto et al., 1997). mGlu receptor subtypes form homo- and heterodimers among themselves (Kammermeier, 2012; Vafabakhsh et al., 2015; Yin et al., 2014), but mGlu receptors dimerize with also other receptors coupled to Gq such as 5-HT_{2A}, β 1-adrenergic, and GABAB receptors (J.-P. Pin & Bettler, 2016). Finally, evidence exists of functional interactions between mGlu receptors and other membrane G protein-coupled receptors such as estrogen receptors in neurons (Spampinato et al., 2012).

All component of CNS, neurons, astrocytes, microglia, oligodendrocytes, endothelial cells, and other circulating immune cells have a role in response to both acute and subacute injury and in chronic neurodegeneration. Some multifunctional treatment strategies can involve also targeting mGluRs, which are expressed in several cell types commonly distributed throughout the CNS (Ferraguti & Shigemoto, 2006). Glial cells express both, ionotropic glutamate receptors and mGluRs, and also glutamate transporters. The different and heterogeneous locations of mGluRs in the CNS constitute an interesting

opportunity to investigate selectivity different drug's target depending by receptor subtypes. A good number of studies have demonstrated that mGluRs are expressed in lymphocytes and in antigen-presenting cells, like dendritic cells, microglia, and macrophages (Fallarino et al., 2010; Pacheco et al., 2006). Furthermore, mGluRs regulate cell migration, glutamate release, and the induction of the inflammatory phenotype in microglia (Barker-Haliski & White, 2015). mGluRs can also modulate immune responses, the kind of the modulation depending on the specific subtype of mGluRs that is involved. (Boldyrev et al., 2005).

The Intracellular signaling stimulated by mGlu receptors has been predominantly studied in neuronal cells, (Gerber et al., 2007). Group I mGlu receptors activate MAP kinase signal that have an important role in protein synthesis-dependent neuronal plasticity (Gerber et al., 2007; Hellyer et al., 2017). Following mGlu receptors activation, the MAPK cascade have several translation and transcription factors as target, that have been well described (Gerber et al., 2007). Group I mGlu receptors dependent phosphorylation of JNKs increases transcription mediated by activator protein- 1 (L. Yang et al., 2006), meanwhile activation of p38 regulates NF- κ B (O'Riordan et al., 2006). Usually G-protein coupled receptor can activate canonical pathway mediated by the G_{α} subunit but also non-canonical pathways (e.g., MAPK activation) mediated by scaffolding proteins such as β -arrestin (Hathaway et al., 2015; Iacovelli et al., 2014). The recruitment of β -arrestin-dependent signaling pathways occurs in response to G-protein coupled Receptor Kinase (GRK)-dependent phosphorylation and is strictly ligand-dependent (Hellyer et al., 2017). The expression of mGlu receptors is highly regulated during the development. mRNA levels for mGlu1, mGlu2, and mGlu4 receptors are low at birth and increase during postnatal development (Luján et al., 2005). For mGlu5, the expression of the shorter mGlu5a receptor isoform is higher in prenatal stages, and mainly localized in cortex, hippocampus and subventricular zone, where it colocalizes with neural progenitors (Boer et al.,

2010), astrocytes and microglia. However, in mature brain, mGlu5b receptor is the main isoform expressed (Luján et al., 2005; Romano et al., 2002).

In glial cells, mGlu1, mGlu3, and mGlu5 receptors are found in astrocytes while mGlu2, mGlu3, and mGlu5 receptors are expressed in microglial cells. In oligodendrocytes, mGlu1 and mGlu4 are highly expressed (Ribeiro et al., 2017), while mGlu5a and mGlu2/3 receptors are present in early developmental stages and downregulated in mature MBPC oligodendrocytes (Deng et al., 2004; Luyt et al., 2003; Spampinato et al., 2015).

In glial cells mGlu receptors regulate cell proliferation (Ciccarelli et al., 1997), the release of growth factors, cytokines (Aronica et al., 2005; Ciccarelli et al., 1999), and neurotransmitters including glutamate, ATP and adenosine, which propagate Ca²⁺ signaling between astrocytes and other glial cells (Hamilton et al., 2010). Glial mGlu receptors can also modulate the activity and the expression of glutamate transporters, participating in this way to the regulation of synaptic function (Aronica, Gorter, Jansen, et al., 2003; Vermeiren et al., 2005).

1.1.9 Group II mGluRs

mGluRs constitute a class of membrane receptors associated with G proteins. Three different receptor classes are distinguished based on their structure, their binding to G proteins and their pharmacological profile. In particular, group 2 is made up of the mGluR2 and mGluR3 receptors, highly similar from a structural point of view, capable of stimulating the Gi protein, but with different distribution on the various cell types and capable of stimulating different, sometimes opposite, responses (Spampinato et al., 2018). Both have been studied as a relevant pharmacological target in neurodegenerative disorders (Bruno et al., 2017).

Regarding the cerebral localization of the Group II receptors, mGluR2 has been identified in only a few brain regions, like the olfactory bulb and cerebellar cortex. As cellular distribution, mGluR2 is mainly concentrated in neurons, more present in the pre-terminal region of axons, far from the sites of

neurotransmitter release (Tamaru et al., 2001). The presynaptic mGluR2/mGluR3 can be activated by an excess of glutamate coming from an over-release of synaptic glutamate or from the glutamate released from astrocytes via the cystine–glutamate membrane antiporter (Kalivas, 2009). Modifications of the expression and activity of the cysteine–glutamate antiporter may influence the function of mGluR2 and mGluR3 in brain areas involved in drug dependence (Crupi et al., 2019; Schmidt & Pierce, 2010). Presynaptic mGluR2/mGluR3 have the main role of reducing the release of neurotransmitters. Both receptor types play a role in the modulation of synaptic plasticity, particularly in stimulating LTD of excitatory synaptic transmission (Altinbilek & Manahan-Vaughan, 2009; Grueter & Winder, 2005; Nicholls et al., 2006).

In the case of mGluR3 distribution in the CNS, it is primarily expressed in the olfactory tubercle, dentate gyrus, cerebral cortex, nucleus accumbens, lateral septal nucleus, striatum, amygdaloid nuclei, cerebellar cortex, and substantia nigra pars reticulata (Petralia et al., 1996; Tamaru et al., 2001; Tanabe et al., 1993). Expression of mGluR3 is also reported presynaptically, postsynaptically, and on glial cells (Ferraguti & Shigemoto, 2006; Ohishi et al., 1993).

Group II mGlu receptors, have been studied as a possible pharmacological target in neurodegenerative disorders (Bruno et al., 2017).

Group II mGluRs were studied in Parkinson Disease, where it has been shown that the selective mGlu2/3 receptor agonist LY379268 improved rotarod performance in animal models of the disease but did not modify akinesia in 6-OHDA-lesioned rats and might even worsen motor symptoms (Johnson et al., 2009). Hence, Group II mGluR agonists were suggested as a possible treatment for neuropsychiatric symptoms associated with PD (J. S. Han et al., 2006).

The stress-related psychiatric conditions, among which major depressive disorder (MDD), anxiety, and drug abuse are considered a relevant problem concerning health worldwide (Cryan & Holmes, 2005). The L-glutamate system represents the primary excitatory neurotransmitter system involved in the circuits of

emotion and cognition and has a central role in the pathophysiology and persistence of disorders related to mental health. Clinical neuroimaging studies have reported a consistent volumetric alteration in brain areas where glutamatergic neurons predominate, among which there are amygdala, hippocampus, and several cortical regions (Lorenzetti et al., 2009). The expression of mGlu2Rs in the hippocampus has been strongly correlated with the mechanisms underlying resilience (or non-resilience) to stress, which have a central role in the pathophysiology of MDD and other stress-related disorders (McEwen et al., 2015).

The potential use of mGluR2/3 agonists in the treatment of schizophrenia was heavily tested in pre-clinical studies using different animal models (Kryszkowski & Boczek, 2021). In a phencyclidine (PCP)-induced model, LY379268 reversed certain behavioral phenotypes and pre-pulse inhibition (Cartmell et al., 2000) to a degree comparable with atypical antipsychotic clozapine. Other studies carried out on mGluR3 and mGluR2 knockout mice demonstrated that mGluR2, but not mGluR3, mediated the effects of LY379268 in experimental models predictive of antipsychotic activity (Woolley et al., 2008). LY379268 also showed promising effects in other schizophrenia models based on pharmacological blockage of NMDA receptor (ketamine and MK-801) (Sokolenko et al., 2019).

The first drug targeting mGluR2/3 receptors that was clinically tested in humans for the treatment of schizophrenia was LY-2140023, based on the assumption that it could normalize hyperactive cortical pyramidal neurons in the thalamus, prefrontal cortex and limbic system (Patil et al., 2007). In subsequent years more selective mGluR2/3 PAMS were tested in clinical trials and they can still be considered promising drugs in schizophrenia treatment to be used alone or in combination with other antipsychotics (Kryszkowski & Boczek, 2021).

Among studies carried out on mGluR2/3 in NVU, it has been shown that under conditions of diabetes complicated by depression, hippocampal neurovascular units were damaged, leading to decreased barrier function and also to other

disfunctions such as elevated Glutamate levels, upregulation of vesicular glutamate transporter 3 (VGLUT-3) and metabotropic glutamate receptor 2/3 (mGluR2/3) expression, downregulation excitatory amino acid transporter 1 (EAAT-1) expression, and finally alterations in the balance of key proteins associated with the extracellular signal-regulated kinase (ERK)/glial cell-derived neurotrophic factor (GDNF)/PI3K signaling pathway. Consequently the results of the study suggested that the Glu/mGluR2/3/PI3K pathway, induced by glucocorticoid receptor activation in the hippocampal neurovascular unit, may be associated with diabetes-related depression. (J. Liu et al., 2021).

As mentioned above, both receptors are expressed in glial cells, astrocytes express only mGlu3 receptor that is present at all developmental stages (Sun et al., 2013), while microglial cells express both mGlu2 and mGlu3 receptors (J. J. Geurts, 2003).

1.1.10 mGluR2-3 role in Astrocytes

Stimulation of Group II mGluRs increases nerve growth factor (NGF) and BDNF production by glial cells (Cicarelli et al., 1997; Matarredona et al., 2001). Elevated Glu levels stimulated the over-expression of the Glu receptor mGluR2/3 (Losonczy et al., 2003) that can also be increased in astrocytes in response to FGF and EGF (Aronica, Gorter, Ijlst-Keizers, et al., 2003) or after exposure to pro-inflammatory cytokines (TNF- α and IL-1 β), (Berger et al., 2012). mGlu3 receptor participates in the control of extracellular glutamate by increasing the expression of the transporter GLAST and GLT-1 (Aronica, Gorter, Ijlst-Keizers, et al., 2003; Gegelashvili et al., 2000; H. Yao et al., 2005; F. Zhou et al., 2006). On these bases, the use of mGlu3 receptor agonists and/or PAMs has been proposed in the treatment of ALS in which a defect of GLT-1 has been described (Battaglia et al., 2015; Rothstein et al., 1995).

In astrocytes, the stimulation of mGluR3 activate MAPK and PI3K pathways, increasing the production of neurotrophic factors (Bruno et al., 1998; Caraci et

al., 2011; Durand et al., 2017) that promote neuroprotection against different toxic insults (Ribeiro et al., 2017).

mGlu5 and mGlu3 receptor activation may control, either in cooperation or independently, various astrocyte functions, such as glutamate transporter activity, astrocyte–arteriolar and astrocyte–neuronal interactions (Bradley & Challiss, 2012). Among group I mGlu receptors, mGlu5 is the mostly expressed in astrocytes. Both mGluR3 and mGluR5 can have at the same time a positive and a negative influence on cell proliferation, and they are also both highly expressed in cultured oligodendrocyte progenitor cells (Aronica, Gorter, Ijst-Keizers, et al., 2003).

In addition, astrocytic mGlu3 receptors, through activation of MAPK and PI3K pathways, contribute to neuroprotection by increasing synthesis and secretion of neurotrophic factors (Bruno et al., 2017). These include TGF- β 1, that prevents both NMDA- and A β -induced toxicity on neurons (Bruno et al., 1998; Caraci et al., 2011; Corti et al., 2007) and GDNF, that is considered an important neurotrophic agent for nigral dopaminergic neurons and has shown neuroprotective and restorative activity in several preclinical models of parkinsonism (Ibáñez & Andressoo, 2017). Accordingly, the pharmacological activation of mGlu3 receptor in mice increases GDNF mRNA and protein levels in striatal neurons (Battaglia et al., 2009). GDNF has also a key role in the protection of neurons from excitotoxicity also in cultured spinal motor neurons (Battaglia et al., 2015). Thus, selective mGlu3 receptor stimulation may have a role in slowing neuronal degeneration in different conditions such as ALS (Battaglia et al., 2015) and PD (Bruno et al., 2017). The role of astrocytic mGlu3 receptors and the interaction astrocyte-neuron has been particularly claimed in this case. Accordingly, studies have shown that mGlu2/3 receptors agonists protect cortical neurons against excitotoxic death only in the presence of astrocytes (Bruno et al., 2017; Caraci et al., 2012). These data have been reinforced by studies carried out in cultured astrocytes from mGlu3 knockout mice (Battaglia et al., 2015; Caraci et al., 2011; Corti et al., 2007) demonstrating the key role of astrocytic mGlu3 receptor in the

mediation of the neuroprotective effects of mGlu2/3 receptor agonists. Activation of mGlu3 receptor activity also protects astrocytes from OGD (Ciccarelli et al., 2007) and nitric oxide damage, due to the reduction of cAMP content and consequent activation of PI3K/Akt pathway (Durand et al., 2010, 2013). It has also been suggested that mGlu3 receptor might represent a relevant pharmacological target to develop disease-modifying drugs in AD (Caraci, Nicoletti, et al., 2018). Even if no clear data are available in human AD brains, mGlu3 receptor expression seems reduced in several animal models of AD (Cha et al., 2001; Dewar et al., 1991; Durand et al., 2014; Knezevic & Mizrahi, 2018). When treated with the mGlu2/3 receptor agonist LY379268, astrocytes *in vitro* reduced neuronal A β toxicity through the release of neuroprotective factors such as TGF- β 1 (Caraci et al., 2011) and BDNF (Durand et al., 2017). This is in line with the observation that TGF- β 1 has anti-inflammatory and neuroprotective effects in experimental models of AD (J.-H. Chen et al., 2015), and a deficit of TGF- β 1 signaling has been found in the early phases of AD and appears to contribute to neuroinflammation and cognitive decline in AD (Caraci, Spampinato, et al., 2018). Activation of mGlu3 receptor can also have a relevance in other steps of AD pathogenesis by reducing A β production (Durand et al., 2014) or increasing A β clearance (Durand et al., 2017). Astrocytic mGlu3 receptors can stimulate the activity of α -secretase, the enzyme that cleaves APP downstream of the N-terminus domain of A β ₍₁₋₄₂₎ (Durand et al., 2014). When astrocytes were exposed to LY379268, they reduced the levels of β -secretase, while increasing the expression of sAPP α , with a consequent decrease of neurotoxic A β . It has been also demonstrated that LY379268 can increase A β uptake in astrocytes and microglia, promoting A β clearance from the extracellular space (Durand et al., 2017). The contribution of mGlu3 receptor seems not completely clear, because A β phagocytosis was not prevented by LY2389575, a selective mGlu3 receptor NAM, suggesting that the effects observed after LY379268 stimulation can also involve mGlu2 receptor activation (Durand et al., 2017).

1.1.11 mGluR2-3 Role in Microglia

mGlu2 and mGlu3 receptors can differentially modulate microglial function. Activation of mGlu2 receptors causes neuroinflammation by stimulating the production of pro-inflammatory cytokines (Pinteaux-Jones et al., 2008; Taylor et al., 2002, 2005). In contrast, activation of mGlu3 receptors drives microglia to an anti-inflammatory function (Pinteaux-Jones et al., 2008). Group II stimulation with the agonist DCGIV also induces BDNF expression in microglia (Venero et al., 2002).

The role of Group II mGluRs in microglia has been investigated in AD models. In response to A β , microglial cells increase their glutamate release (Barger & Basile, 2001). The exposure of microglial cells to the active fragment A β ₍₂₅₋₃₅₎ induces also mGlu2 receptor activation, that can drive an increased neurotoxicity (Taylor et al., 2002, 2005). In fact, selective activation of mGlu2 receptors, promotes a pro-inflammatory and neurotoxic phenotype that releases TNF- α and FAS-L, and potentiates microglial reactivity in response to chromogranin-A, up-regulated in AD (Taylor et al., 2002, 2005).

It is well known that microglial activation plays a key role in the pathogenesis of MS (Strachan-Whaley et al., 2014). *In vitro*, exposure of microglial cells to myelin fragments induces microglia activation, increasing the release of glutamate and TNF- α , that cause neuronal death. In this context, activation of microglial mGlu2 receptor worsen myelin-evoked neurotoxicity, whereas activation of mGlu3 receptor is protective (Pinteaux-Jones et al., 2008).

A good neuroprotective effects of orthosteric mGlu2/3 receptor agonists have been observed in animal models of global and focal brain ischemia (Bond et al., 1998, 2000), probably as a consequence of the involvement of mGlu2 receptors expressed in neurons (Corti et al., 2007; Mastroiacovo et al., 2017; Motolese et al., 2015). However, the role of microglial mGlu2 receptor in stroke ischemia has not been completely clarified. mGlu2 and mGlu3 receptors are expressed by microglia in the ischemic penumbra, where apoptotic neuronal death progresses

slowly, making this area more open to therapeutic intervention and moreover microglial cells mediate neurotoxicity in the stroke penumbra (Kaushal & Schlichter, 2008). In experimental models of ischemia, it has been demonstrated that glutamate, released by “ischemic” neurons, activates microglia through group II mGlu receptors with the following activation of NF- κ B, induction of TNF- α , and subsequent neuronal death (Kaushal & Schlichter, 2008).

A recent study demonstrated that the microglial mGlu3 receptor is highly downregulated in a rat model of perinatal brain injury. Selective pharmacological activation of mGlu3 receptors during early postnatal life might mitigate neuroinflammation associated with perinatal adverse conditions (Zinni et al., 2021). In particular, after exposure to LPD (low protein diet) together IL-1 β , Grm3 gene expression results significantly downregulated in the developing microglia (Zinni et al., 2021). The study demonstrated that pharmacological blockade of mGlu3 receptors induced a pro-inflammatory phenotype in microglia sorted from rat pups. Finally microglia reactivity to inflammatory response induced by LPD/IL-1 β exposure was reduced by an mGlu3 receptor agonist (Zinni et al., 2021).

In mice, it has been demonstrated that the activation of mGlu3 receptors has a protective activity against ischemic brain damage and associated neuroinflammation (Mastroiacovo et al., 2021). In fact, in mGlu3^{-/-} mice, there is increased expression of several pro-inflammatory genes such as those encoding for IL-1 β , COX-2, TNF- α , CD86, and IL-6; these genes were more present in the peri-infarct region of mGlu3^{-/-} mice. On the other hand, following the medial cerebral artery occlusion, the expression of two genes associated with the anti-inflammatory phenotype of microglia (those encoding the mannose-1-phosphate receptor and the α -subunit of interleukin-4 receptor) and the gene encoding for the neuroprotective factor, GDNF, were enhanced in the peri-infarct region of wild-type mice, but not mGlu3^{-/-} mice (Mastroiacovo et al., 2021).

1.2 Aims of the study

The main aims of the first part of the study were to define the role of microglia in BBB function using an *in vitro* cellular model that allowed identifying the specific interplay among different cell types of the NVU and to assess the effect of pharmacological intervention. Among the various possible targets expressed in glial cells, group 2 metabotropic receptors were investigated. The specific aims of the study were:

- to evaluate whether, in an *in vitro* BBB model, modulation of mGlu2 and mGlu3 receptors plays a role in maintaining barrier properties following an inflammatory stimulus;
- to assess the role of microglia in modulating BBB properties and to evaluate whether this function can be affected by stimulation of mGlu2/3 receptors;
- to analyze the role of pericytes in the control of BBB function in inflammatory conditions.

1.3 Materials and Methods

1.3.1 Cells cultures of BBB *in vitro* model

To simulate the BBB *in vitro* model were used three adult human cell lines: TY-10 cells, brain microvascular endothelial cells; hAST, astrocytic cells; and HMC3 human microglial cell line (ATTC, LGC Standards, Manassas, VA, USA). Both TY-10 and hAst are transfected with a plasmid expressing temperature sensitive Simian virus-40 large T-antigen and the catalytic subunit of human telomerase, as previously described (Haruki et al., 2013). Both cell lines were developed at Yamaguchi University (Japan), in the labs of Dr. Sano and Kanda. TY-10 cells were grown in MCDB-131 media (Thermo Fisher Scientific, Milan, Italy), supplemented with EGM-2 SingleQuots (Lonza, Basel, Switzerland) and 20% heat inactivated FBS (Thermo Fisher Scientific). hAST were grown in astrocyte medium containing 3% heat-inactivated Fetal Bovine Serum (FBS), astrocyte growth supplement and

penicillin/ streptomycin solution, as provided with the Astrocyte media kit (Clinisciences, Nanterre, France). HMC3 was grown in Eagle's Minimum Essential Medium (EMEM, Sigma Aldrich) supplemented with 10% FBS (Thermofisher Scientific), Non-essential amino acids, sodium Pyruvate and penicillin (100 U/mL)/streptomycin (100 ug/mL). hAst and TY-10 were grown at 33 °C and 5% CO₂ for two days and then were transferred to 37 °C, where they exhibited growth arrest and differentiation. HMC3 were grown at 37 °C and 5% of CO₂ for 1 to 3 days. For the study, TY-10 were cultured alone or in co-culture with hAst or hAst and HMC3. For the endothelium/astrocytes were used specific insert with a polycarbonate membrane according to experiment protocol and for the triple-culture was also used a coverslip placed between the cell culture plate and the insert for the third cell type. Both hAst and TY10 were exposed to treatments after two days at 37 °C simultaneously at HMC3 for the triple-culture. All treatments were done in astrocyte medium supplemented with 2% FBS. Treatment with cytokines (TNF α , 1 ng/mL, and IFN γ , 1 ng/mL; T&I) was maintained either for 5 or 36 h, according to the experimental protocol. Endothelial monocultures, endothelial/astrocytes co-cultures, or endothelial/astrocytes/microglia co-cultures when required by experimental design, were pretreated with a negative allosteric modulator of mGluR2, (VU 6001966, 3 μ M, VU) for 20 min and with an agonist of mGluR2/3 (LY379268, 1 μ M, LY37) or an mGluR3 positive allosteric modulator LY2794193 (LY27, 5 μ M) for 40 min.

1.3.2 Western Blot

For western Blot analysis, Endothelial cell monocultures (5×10^5 cells for each well) were plated on 6 wells MW plates. When cocultured with endothelial cells, hAST (3×10^5 cells for each well) were plated on 0.4 μ m pores poly-carbonate membrane transwell inserts (Falcon) and when cocultured with endothelial cells and microglia, were plated on the downside of a glass coverslip (3×10^5 cells for each coverslip). In the triple culture, microglial cells were plated on the transwell insert (4×10^5 cells for each well). For double and triple culture, endothelial cells

and hAst were put together when the cells were passed to 37°C. For the triple-culture. Microglia was added in the culture before the treatments. This setting allowed the passage of soluble factors between the different cell types without direct physical contact, thus also facilitating their isolation for western blot investigations. All the treatments were performed in Astrocyte medium supplied with 2% FBS. After 36 hours of treatments, pellets were collected and lysed in RIPA lysis buffer (Sigma-Aldrich) supplemented with protease and phosphatase inhibitors for 5 hours at 4°C. Samples were sonicated and centrifuged at high speed for 5 min at 4 °C, and protein concentration was determined by a Bradford reagent (Sigma-Merck, Darmstadt, Germany), and absorbance was measured with a Varioskan™ Flash Multimode Reader. Sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) was performed by loading 40 ug of protein extracts per experiment on pre-cast “any-kDa” or 4–20% gradient gels (Bio-Rad, Hercules, CA, USA) followed by transfer to nitrocellulose membrane (Hybond ECL, Amersham Biosciences Europe GmbH, Milan, Italy) using a Transblot semidry transfer cell (Bio-Rad, Hercules, CA, USA). Membranes were blocked with a Blocker FL Fluorescent Blocking buffer (ThermoFisher Scientific) and incubated with primary antibodies: mouse anti-GAPDH (1:800; Millipore), rabbit anti-Flotillin-1 (1:700; Santa Cruz), and rabbit anti-Claudin-5 (1:300; Thermo Fisher Scientific) overnight at 4 °C. Membranes were then washed and exposed to secondary antibodies: anti-mouse AlexaFluor 488 Plus-conjugated secondary antibody (ThermoFisher Scientific) and anti-rabbit IRdye 800 secondary antibody (Licor) one hour at RT. The detection of specific bands was carried out using the iBright FL1500 Imaging System (ThermoFisher Scientific). Band intensity was analyzed using the ImageJ software, developed by the National Institutes of Health (NIH) and in the public domain.

1.3.3 Immunocytochemistry

For immunocytochemical analysis, endothelial cells will be fixed in ice cold acetone (10 minutes) and then in ice-cold methanol (15 minutes). Cells are incubated with primary antibody: rabbit anti-Claudin-5 (1:100; Thermo Fisher

Scientific) and rabbit anti-ZO-1 (1:100, Thermo Fisher Scientific), in the presence of 0.1% Triton X-100 at 4 °C overnight. The secondary antibody: anti-rabbit AlexaFluor 488 Plus-conjugated secondary antibody (ThermoFisher Scientific) is incubated for 1 hour at room temperature. Slides will be observed using an epifluorescent microscope (Zeiss Observer.Z1) equipped with the Apotome.2 acquisition system connected to a digital camera.

1.3.4 FITC-dextran permeability assay

For the test endothelial cells were plated on a 3 μM pores poly-carbonate membrane transwell insert (Corning) (15×10^4 cells for each well). Endothelial cells could be mono-cultured, co-cultured with hAst plated in 24 MW plate (9×10^4 cells for each well), cocultured with hAst and HMC3, in this case, HMC3 were plated on the well (15×10^4 cells for each well) and hAst were plated on a glass coverslip. Also in this experiment protocol, TY-10 and hAst were put together when cells were passed to 37 °C, and HMC3 before the treatments. After 36 hours of treatments inserts were equilibrated in the phenol red free DMEM media supplemented with 1% FBS. Solute permeability was assessed using 10 kDa FITC-conjugated dextran (1 mg/mL). Conjugated dextran was applied to the luminal compartment. One hundred microliters of sample will be collected from the abluminal compartment after 60 minutes and fluorescence of the sample measured at 485/520 nm (excitation/emission) using a Varioskan TM LUX multimode microplate reader (ThermoFisher Scientific).

1.3.5 Trans-endothelial electrical resistance analysis (TEER)

For TEER analysis were used the same insert and the same model of the permeability assay. The measurements were performed in real time, without causing damage to cells. The electrical resistance, in Ohm (Ω), was measured by means of an electrode which is positioned so that one end was in contact with the culture medium inside the insert and the other with the culture medium outside of it. The cells are subjected to various measurements at different times, before and during treatments. Various measurements were performed, and the average TEER value was calculated. For the TEER evaluation, was used the

specific device Millicell®-ERS (Electrical Resistance System) Voltometer (Millipore).

1.3.6 Quantitative Real-Time Polymerase Chain Reaction (PCR)

The PCR experiments were performed using the same model previously described in western blot analysis for the co-cultures and a microglial cells monoculture (5×10^5 cells for each well) plated on a 6 MW plate.

Total RNA was extracted from cell cultures using the RNeasy plus Mini Kit (Qiagen). One mg of RNA was used for cDNA synthesis, using the Superscript-VILO kit (Invitrogen) according to manufacturer's instruction. Quantitative real-time PCR was performed with Rotor Gene Q using QuantiNova SYBR Green PCR Kit (Qiagen). The melting curves obtained after each PCR amplification reaction confirm the specificity of the 2-[N-(3-dimethylaminopropyl)-N-propylamino]-4-[2,3-dihydro-3-methyl-(benzo-1,3-thiazol-2-yl)-methylidene]-1phenylquinolinium (SYBR Green assays). The following pairs of primers were used: human CCL2F (CCC CAG TCA CCT GCT GTT AT), human CCL2R (AGA TCT CCT TGG CCA CAA TG), human GRM2F (CCA GGA GCT GGG TCC CTT), human GRM2R (AAG TCT CCC TCC AGG GTC AG), human GRM3F (CGC TTT GCA CAA AAT GCA GC) and human GRM3R (AAC ACG TTG TAT CGC CCC AT) from Invitrogen (Thermo Fisher Scientific), and the following primeQuantitec primers (Qiagen) were used: human IL-6 (QT00083720), human BDNF (QT00235368), human IL-1 β (QT00021385) and human RPLP0 (QT00075012) that was used as an endogenous control. Expression fold changes were calculated by applying the $2^{-\Delta Ct}$ method.

1.3.7 Human Brain Pericytes (HBP) cell culture and treatment

Human brain pericytes (HBP) utilized in another part of the study (performed at "Research Center Blood-Brain Barrier Laboratory, LBHE", University of Artois, France) were originally isolated from a patient who had unexpectedly died from a heart attack (Shimizu et al., 2011). Study protocol for human tissue was approved by the ethics committee of the Medical Faculty (IRB#: H18-033-6), University of Yamaguchi Graduate School, and was conducted in accordance with

the Declaration of Helsinki, as amended in Somerset West in 1996. The protocol was approved by the French Ministry of Higher Education and Research (CODECOH Number DC2011-1321). All experiments were performed in accordance with the approved protocol. Briefly, these pericytes were transfected and immortalized using retro-virus vectors holding human temperature-sensitive SV40 T antigen (tsA58) and human telomerase (Htert). HBP were then amplified and cultured at 33 °C in high glucose (4.5 g/L) Dulbecco's modified Eagles' medium (DMEM/HG), supplemented with 10% non-heat-inactivated fetal calf serum (FCS, Sigma-Aldrich, Saint-Louis, MS, USA), 1% L-glutamine (Merck chemicals, Darmstadt, Germany) and 1% penicillin–streptomycin (Sciencell, Carlsbad, CA, USA). Before plating, well plates were coated with collagen I (Corning, NY, USA). Then, HBP were seeded in 12-well plates (for fluorescence experiments) or on a coverslip (for immunofluorescence experiments) at a density of 50,000 cells/well. Only brain pericytes at passage 11 were used for the experiments of this study. Cells were cultured at 37 °C in DMEM/HG supplemented with 1% L-glutamine (Merck KGaA, Darmstadt, Germany) and 1% penicillin–streptomycin (Sciencell) during 48 h to reach 80 to 90% of confluence. For the treatment performed, TNF- α (Sigma-Aldrich) was dissolved in DMEM/HG supplemented with 0.1% bovine albumin serum (BSA, Sigma-Aldrich) at a concentration of 10 ng/mL. When ready, HBP were rinsed once with warm DMEM/HG/0.1% BSA and then incubated in DMEM/HG/0.1%.BSA supplemented or not with 10 ng/mL of TNF- α .

1.3.8 Quantification of A β ₍₁₋₄₀₎ and A β ₍₁₋₄₂₎ fluorescence uptake experiments

For these experiments the HBP treated or not with TNF- α 10 ng/mL for 24h. Then cells were rinsed with warm DMEM/HG/0.1%.BSA and incubated with the solution of β -Amyloid HiLyte™ Fluor 488-labeled (1-40) (AS-60492-01, ANASPEC, Fremont CA, USA) or β -Amyloid (1-42) HiLyte™ Fluor 488-labeled (Cod: AS-60479-01, ANASPEC, Fremont CA, USA), both to a concentration of 100 nM in DMEM/HG/0.1%.BSA for 3 hours at 37 °C. After the treatment the cells were rinsed with Ringer Hepes (RH; 150 mM NaCl, 5.2 mM KCl, 2.2 mM CaCl₂, 0.2 mM

MgCl₂–6H₂O, 6 mM NaHCO₃, 5mM HEPES, 2.8mM glucose; pH: 7.4) for three times and lysed with RIPA buffer addition of protease inhibitors to assess the quantity fluorescence peptides were taken from cells. The lysates were collected, and an aliquot was taken in a dark 96-wells plate for measuring the fluorescence. The fluorescence quantification was quantified to values of λ : A β -40 (503-532) for A β -40 and (550-576) A β -42, excitation/emission with a spectrofluorometer (Synergy H1, Biotek, Colmar, France). The values of fluorescence obtained were normalized on the total amount of proteins measured with Bradford assay for each sample. Experiments were performed on at least three independent experiments in triplicate for each condition.

1.3.9 Immunofluorescence of A β ₍₁₋₄₀₎ and A β ₍₁₋₄₂₎ uptake

The HBP were treated as described above. First with TNF- α and then with A β -40 or A β -42 at the same concentrations and times. At the end of the incubation, cells were washed with RH buffer for three times and were treated briefly with Trypan Blue (0,2%) for 5 minutes used as autofluorescence quenching. After three washing with PBS the cells were fixed with Ice-cold Methanol for 1 minute, followed by three washing with PBS for 5 minutes. Lately the coverslips with the fixed cells were mounted on a microscope slide with a mounting solution with DAPI. Images were acquired using a Leica microscope (DMRD; Leica Microsystems, Wetzlar, Germany) and processed using the ImageJ software. Experiments were performed on at least three independent experiments in triplicate for each condition.

1.3.10 Statistical Analysis

All data are expressed as mean \pm SEM of 3–5 different experiments, each run in duplicate or in triplicate, as specified in the figure legends. Data were analyzed by one-way ANOVA, followed by the Newman–Keuls test for significance or by Unpaired Student's t-test two-sided. $p < 0.05$ was taken as the criterion for statistical significance.

1.4 Results

1.4.1 GRM2 and GRM3 expression

To evaluate the effects of the modulation of BBB's properties by mGluR2 and mGluR3, we first verified the expression of the two receptors in the three different cells lines with real time PCR analysis. Endothelial cells, astrocytes and microglia expressed both genes GRM2 and GRM3. (Tab.1).

	take off	amplification		take off	amplification
mGluR2_TY10	29,7	1,66	mGluR3_TY10	41,4	1,73
mGluR2_hAST	26,5	1,61	mGluR3_hAST	29,5	1,71
mGluR2_HMC3	25,5	1,59	mGluR3_HMC3	30,8	1,76

Tab. 1. mRNA expression levels for mGluR2 and mGluR3 in the cell lines utilized in the study.

1.4.2 mGlu3 receptor activation modifies endothelial function

After, to assess the effect of mGlu2/3 activation on the endothelium, a monoculture of endothelial cells grown alone was exposed to the inflammatory stimulus of TNF α (1 ng/mL) and IFN γ (1 ng/mL) (T&I) for 24-36 hours. This treatment caused impairment of barrier properties as measured by increased permeability to FITC-conjugated dextran; pretreatment with the mGluR2/3 agonist LY379268 (LY37, 1 μ M) contrasted the effect of T&I, significantly reducing permeability (Fig 1a). In contrast, the addition in the pretreatment of a mGluR2 NAM, VU6001976, did not significantly modify the effects of LY37 (Fig 1a). At 24 h there was also a decrease of Trans-endothelial electrical resistance (TEER) and pre-treatment with LY37 rescued TEER at basal values (Fig 1b). Endothelial permeability is dependent also on changes in the expression of endothelial tight junctions (TJs) like Claudin-5 and Zo-1. Expression of claudin-5 was significantly reduced by the inflammatory insult in the membrane fraction and mostly redistributed to the cytosolic compartment. Treatment with LY37 rescued claudin-5 localization to the membrane, whereas VU did not modify LY37 effect (Fig. 2a.) Furthermore, immunocytochemical analysis showed that both claudin-5

(Fig. 2b) and Zo-1 (Fig. 2c) were clearly reduced on the membrane and internalized in the cytosol and LY37 contrasted these effects.

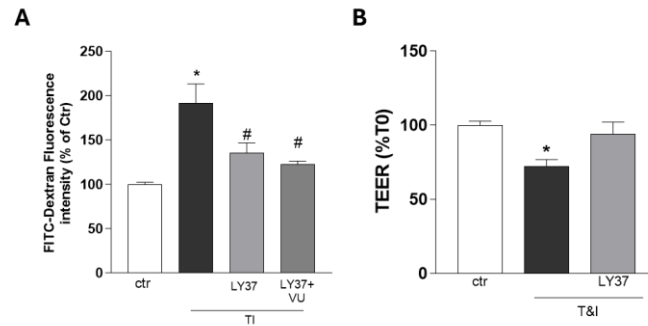


Fig. 1. Evaluation of Barrier function in Endothelial cells monolayer. **A)** FITC-Dextran Fluorescence Permeability assay on Endothelial cells monolayer after 36H of T&I treatment. One-way ANOVA followed by Newman–Keuls test. * $p < 0.05$ CTR vs T&I, # $p < 0.05$ T&I vs LY37, # $p < 0.05$ T&I vs LY37+VU. **B)** Measure of Trans Endothelial Electrical Resistance (TEER) on Endothelial cells monolayer after 24H of T&I treatment. One-way ANOVA followed by Newman–Keuls test. * $p < 0.05$ CTR vs T&I.

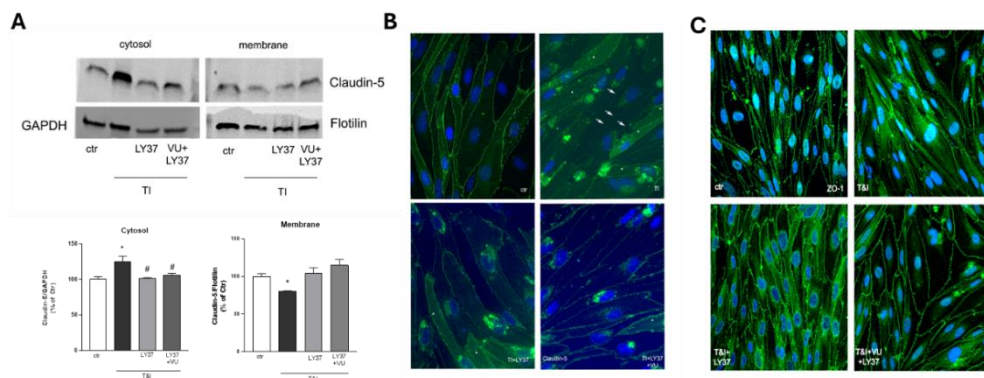


Fig. 2. Tight Junction (TJs) proteins expression in Endothelial cells monolayer. **A)** Claudin-5 expression levels in cytosolic and membrane compartment after 36H of T&I treatment. One-way ANOVA followed by Newman–Keuls test. * $p < 0.05$ CTR vs T&I, # $p < 0.05$ T&I vs LY37+T&I, # $p < 0.05$ T&I vs LY37+VU+T&I. **B)** Immunocytochemistry fluorescence image of claudin-5 expression after 36H of T&I treatment, Green: claudin-5, Blue: DAPI. **C)** Immunocytochemistry fluorescence image of claudin-5 expression after 36H of T&I treatment, Green: claudin-5, Blue: DAPI.

1.4.3 mGlu3 mediates the protective effect on BBB function in endothelium/astrocyte co-cultures

Exposure of the endothelium/astrocytes co-culture to T&I treatment (both at 1 ng/mL for 24-36 h) induced an enhancement of FITC-dextran fluorescence permeability measured at 36 h, LY37 prevented this effect, that was not modified by VU (Fig. 3a). T&I caused an increase of TEER already significant at 24 h, that was contrasted by pre-treatment with LY37 (Fig. 3b). Furthermore T&I caused also a reduction in the expression of the junction protein claudin-5, as detected by western blot analysis of the whole protein extract. LY37 prevented the effects of T&I on claudin-5 expression, that was not significantly modified when the NAM of the mGluR2 receptor VU was added (Fig. 4). We then selectively stimulated the mGluR3 receptor using the positive allosteric modulator LY2794193 (LY27, 5 μ M) and observed a significant effect on the increased permeability induced by T&I (Fig 5a). We also analyzed the inflammatory response on astrocytes grown in an endothelium/astrocytes co-culture following treatment with T&I and pretreatment with LY27. The latter completely prevented the gene expression of the pro-inflammatory cytokine IL-6, induced by a 5 h-exposure to T&I (Fig. 5b). In contrast, mGluR3 stimulation with LY27 increased the expression of transforming growth factor- β (TGF- β), that was not modified by the inflammatory challenge of T&I (Fig. 5c).

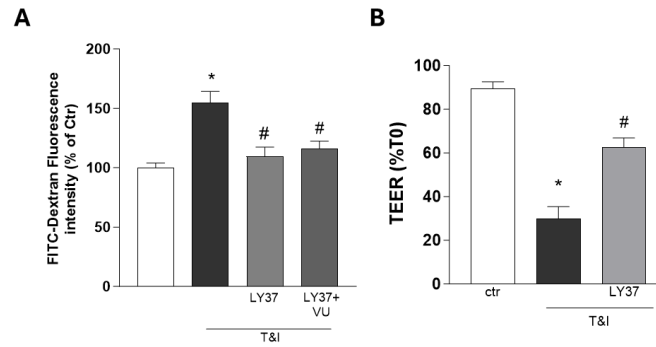


Fig. 3. Evaluation of Barrier function in Endothelial cells of Endothelium/Astrocytes co-cultures. **A)** FITC-Dextran Fluorescence Permeability assay on Endothelial cells from endothelium/astrocytes co-cultures after 36H of T&I treatment. One-way ANOVA followed by Newman–Keuls test. * $p < 0.05$ CTR vs T&I, # $p < 0.05$ T&I vs LY37+T&I, # $p < 0.05$ T&I vs LY37+VU+T&I. **B)** Measure of Trans Endothelial Electrical Resistance (TEER) on Endothelial cells from endothelium/astrocytes co-cultures after 24H of T&I treatment. One-way ANOVA followed by Newman–Keuls test. * $p < 0.05$ CTR vs T&I, # $p < 0.05$ T&I vs LY37+T&I.

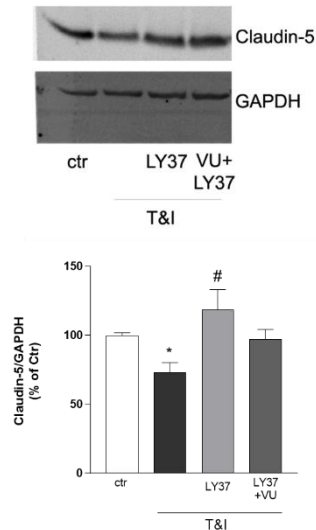


Fig. 4. Tight Junction (TJs) proteins expression in Endothelial cells from Endothelium/Astrocytes Co-cultures. Claudin-5 representative bands and expression levels in Endothelial cells from Endothelium/Astrocytes Co-cultures after 36H of T&I treatment. One-way ANOVA followed by Newman–Keuls test. * $p < 0.05$ CTR vs T&I, # $p < 0.05$ T&I vs LY37+T&I.

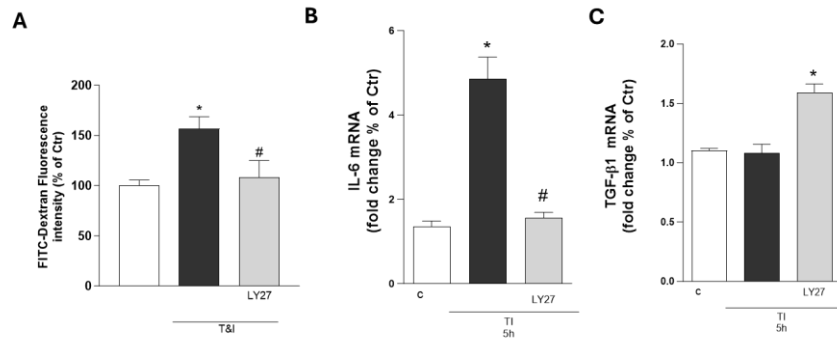


Fig. 5. Effects of selective mGluR3 stimulation on endothelial barrier properties and Astrocytes mRNA expression levels in Endothelium/Astrocytes Co-cultures. **A)** FITC-Dextran Fluorescence Permeability assay on Endothelial cells monolayer from Endothelium/astrocytes co-cultures after 36H of T&I treatment, One-way ANOVA followed by Newman–Keuls test. * $p < 0.05$ CTR vs T&I, # $p < 0.05$ T&I vs LY27+T&I. **B)** IL-6 mRNA expression levels in Astrocytes from Endothelium/astrocytes co-cultures after 5H of T&I treatment, One-way ANOVA followed by Newman–Keuls test. * $p < 0.05$ CTR vs T&I, # $p < 0.05$ T&I vs LY27+T&I. **C)** TGF-β1 mRNA expression levels in Astrocytes from Endothelium/astrocytes co-cultures after 5H of T&I treatment, One-way ANOVA followed by Newman–Keuls test. * $p < 0.05$ T&I vs LY27+T&I.

1.4.4 mGluR2 cooperates to protect BBB function in a triple endothelial-astrocyte-microglia co-culture

In the triple endothelial-astrocyte-microglia co-culture, the increased endothelial permeability to dextran induced by T&I, was reduced by LY37 (1 μ M) and this effect was contrasted by treatment with the mGlu2 NAM, VU (3 μ M); (Fig 6a). Similarly, T&I-reduced expression of claudin-5 was restored by pre-treatment with LY37, and this effect was again counteracted by VU (Fig. 6b).

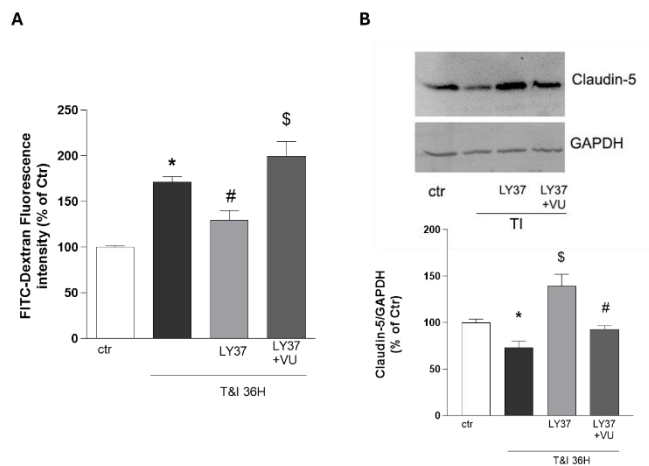


Fig. 6. Evaluation of Barrier function and TJs Protein expression in Endothelial cells of Endothelium/Astrocytes/Microglia triple co-cultures. A) FITC-Dextran Fluorescence Permeability assay on Endothelial cells from endothelium/astrocytes/microglia triple co-cultures after 36H of T&I treatment. One-way ANOVA followed by Newman–Keuls test. * $p < 0.05$ CTR vs T&I, # $p < 0.05$ T&I vs LY37+T&I, \$ $p < 0.05$ LY37+T&I vs LY37+VU+T&I. **B)** Claudin-5 representative bands and expression levels in Endothelial cells from Endothelium/Astrocytes/Microglia triple Co-cultures after 36H of T&I treatment. One-way ANOVA followed by Newman–Keuls test. * $p < 0.05$ CTR vs T&I, \$ $p < 0.05$ T&I vs LY37+T&I, # $p < 0.05$ LY37+T&I vs LY37+VU+T&I.

1.4.5 mGluR2 modulates microglia inflammatory response

To better understand the role of microglia, we therefore evaluated the expression of selected factors also in a microglial monoculture exposed to the same treatments via real time PCR. LY37 attenuated the expression of the inflammatory cytokines IL-6 and IL-1 β and the chemokine CCL2 in microglia cells activated by T&I (Fig. 7A-C) Blockade of mGluR2 with the NAM VU reduced these effects for IL-6 (Fig. 7A) and CCL2 (Fig. 7C) while prevented the effects of LY37 for IL-1 β (Fig. 7B). Treatment of microglia with LY37 also increased the expression of the neurotrophic factor BDNF, and this effect was blocked by VU (Fig. 7D).

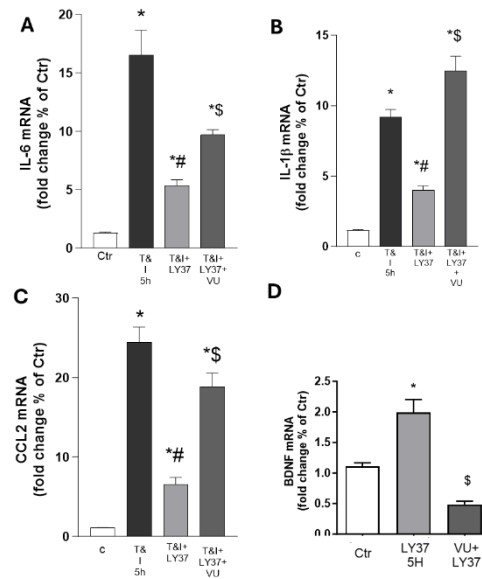


Fig. 7. Microglia mRNA expression levels in Microglia monoculture. **A)** IL-6 mRNA expression levels in Microglia after 5H of T&I treatment, One-way ANOVA followed by Newman–Keuls test. * $p < 0.05$ CTR vs T&I, # $p < 0.05$ T&I vs LY37+T&I, \$ $p < 0.05$ LY37+T&I vs LY37+VU+T&I **B)** IL-1 β mRNA expression levels in Microglia after 5H of T&I treatment, One-way ANOVA followed by Newman–Keuls test. * $p < 0.05$ CTR vs T&I, # $p < 0.05$ T&I vs LY37+T&I, \$ $p < 0.05$ LY37+T&I vs LY37+VU+T&I **C)** CCL2 mRNA expression levels in Microglia after 5H of T&I treatment, One-way ANOVA followed by Newman–Keuls test. * $p < 0.05$ CTR vs T&I, # $p < 0.05$ T&I vs LY37+T&I, \$ $p < 0.05$ LY37+T&I vs LY37+VU+T&I **D)** BDNF mRNA expression levels in Microglia after 5H of LY37 treatment, One-way ANOVA followed by Newman–Keuls test. * $p < 0.05$ CTR vs LY37, \$ $p < 0.05$ LY37 vs LY37+VU.

1.4.6 TNF- α treatment reduced the A β -peptides uptake on Human Brain Pericytes

To assess the function of brain pericytes in favoring transport and clearance at the BBB, we analyzed the uptake of the A β -peptides in cultured Human Brain Pericytes (HBP) utilizing fluorescent-conjugated forms of both A β (1-40) and A β (1-42) peptides. After TNF- α treatment, cells were incubated for 3 hours with one of the two peptides. At the end of incubation, the cells were washed to eliminate as much possible the external quote of A β -peptides and the uptake quotes were determined by quantifying fluorescence in the cell lysate or qualitatively with immunofluorescence on fixed cells. Regarding the fluorescence quantification, results showed that the TNF- α treatment was able to reduce fluorescence (normalized on the total protein amount), compared to the control cells, for both

$A\beta_{(1-40)}$ and $A\beta_{(1-42)}$ peptides and consequently the cells internalized less $A\beta$ -peptides (Fig. 8 A-B). For $A\beta_{(1-40)}$ data were confirmed also by immunofluorescence images where TNF- α treatment showed less internalization compared to control (Fig. 8C).

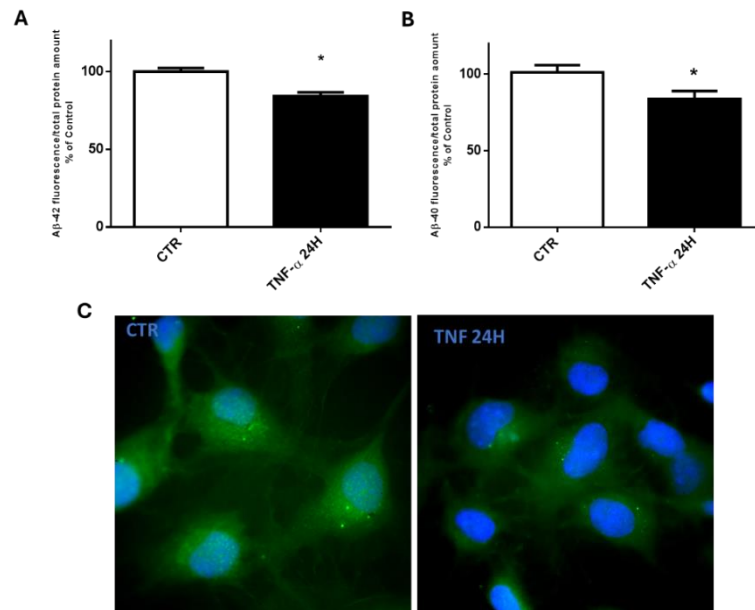


Fig. 8. $A\beta$ -peptides uptake on Human Brain Pericytes. A) $A\beta_{(1-42)}$ peptide fluorescence normalized on total protein amount after 3H of exposition to the peptide in human brain pericytes. Unpaired Student's t-test, $P < 0.05$ CTR vs TNF- α 24H. **B)** $A\beta_{(1-40)}$ peptide fluorescence normalized on total protein amount after 3H of exposition to the peptide in human brain pericytes. Unpaired Student's t-test, $P < 0.05$ CTR vs TNF- α 24H **C)** $A\beta_{(1-40)}$ peptide immunofluorescence after 3H of exposition to the peptide in human brain pericytes. $A\beta_{(1-40)}$: green, DAPI: blue.

Although these are only preliminary results, they confirm that an inflammatory condition can affect also the behavior of pericytes that reduce their ability to internalize $A\beta$ -peptides. These results suggest a main role also for this cellular component in BBB function. Further studies are needed to assess which BBB transporters, including LRP1, ABCA1, ABCB1 (P-gp), can be modified following treatment with TNF- α .

1.5 Discussion

In this study we investigated BBB properties in an in-vitro model, utilizing a single, double- and triple-co-culture system. The objective was to understand the influence of the mGluR2-3 receptors in the presence of an inflammatory insult, focusing in particular on the role of microglia cells. It is well known from literature that microglia can have a role in the modulation of BBB function in neuroinflammatory conditions that follow an acute event, such as stroke, but also participate in slow neurodegenerative conditions, such as Alzheimer's disease or multiple sclerosis (Gullotta et al., 2023; Spampinato, Costantino, et al., 2022). Attention has been paid to these conditions, taking into account the well-recognized role of neuroinflammation in the slow events that characterize neurodegenerative diseases. In response to an insult, the behavior of microglia can be different: they can remove cellular debris in an attempt to contribute to cellular repair or can accelerate the induction of the neuroinflammatory response (G. I. Caruso et al., 2021; Merlo et al., 2022; Merlo, Luaces, et al., 2020).

Regarding their role on BBB, microglia cells interact with endothelial cells of the barrier, and even though they do not seem to play an essential role in BBB formation and stability, they modulate BBB functions, both in physiological and pathological conditions (da Fonseca et al., 2014). In particular, in inflammatory conditions, microglia demonstrated to play a dual role. Microglial cells may promote BBB breakdown through the release and production of inflammatory cytokines, ROS and NO (Mahad & Ransohoff, 2003) that induce endothelial ICAM-1 expression (Huber et al., 2006) and weaken the BBB by damaging supporting astrocytes (Haruwaka et al., 2019). In addition, reactive microglia contribute to a leaking BBB that can be observed in neuroinflammatory conditions through downregulation of TJs such as Claudin-5, occludin, and ZO-1 (Kleinberger et al., 2017; Obermeier et al., 2013). On the other hand, the involvement of microglia in increasing TJs stability in early phases of inflammatory conditions has also been demonstrated (Haruwaka et al., 2019). Furthermore, endothelium-derived factors including nitric oxide are known to suppress microglial activation and

exert anti-inflammatory actions, suggesting a bidirectional control between endothelium and microglia (Katusic & Austin, 2016).

In this context, we investigated the role of mGluR2 and mGluR3, two receptors that are considered relevant pharmacological targets in neurodegenerative disorders (Bruno et al., 2017). Both receptors are expressed in glial cells: astrocytes express principally mGlu3 receptor that is present at all developmental stages (Sun et al., 2013). Alternatively, microglial cells express both mGlu2 and mGlu3 receptors (J. J. G. Geurts, 2003). The protective role of mGluR3 in astrocytes is well established (Spampinato et al., 2018) and it is largely known that stimulation of mGluR3 activates MAPK and PI3K pathways, increasing the production of neurotrophic factors (Bruno et al., 1998; Caraci et al., 2011; Durand et al., 2017). In contrast, the role of these receptors on microglia appears more uncertain. Activation of mGlu2 receptors causes neuroinflammation by stimulating the production of pro-inflammatory cytokines (Pinteaux-Jones et al., 2008; Taylor et al., 2002, 2005), whereas activation of mGlu3 receptors drives microglia to an anti-inflammatory function (Pinteaux-Jones et al., 2008). More recent evidence suggests that selective pharmacological activation of mGlu3 receptors in microglia during early postnatal life might mitigate neuroinflammation associated with perinatal adverse conditions (Zinni et al., 2021). In mice, it has also been demonstrated that the activation of mGlu3 receptors has a protective activity against ischemic brain damage and associated neuroinflammation (Mastroiacovo et al., 2021).

We examined the endothelial barrier function in the three different culture conditions. Starting from monoculture, endothelial cells were exposed to the inflammatory stimulus of TNF α and IFN γ (T&I), a treatment that caused a profound impairment of barrier properties, as demonstrated by increased membrane permeability and reduced endothelial electrical properties. Our results also showed that treatment with T&I caused a translocation of claudin-5 from membrane to cytosol. Regarding the modulation of mGluR2/3, results showed that pre-treatment with LY37 was able to prevent the effects of

inflammatory cytokines on permeability, TEER and claudin expressions, while the addition of VU did not modify these effects. These results suggest that mGluR3 is the receptor subtype more involved in the modulation of endothelial properties.

Similarly, in the endothelium/astrocyte co-cultures, T&I impaired the BBB properties, as measured by increased permeability, reduction in TEER and of total claudin-5 expression. Stimulation of mGlu2/3 receptors again prevented these effects and the lack of changes observed in the presence of blockade of mGlu2 receptor led us to hypothesize that in endothelium/astrocytes co-culture, the effect of LY37 was more linked to the activity of the mGlu3 receptor which, as is known from the literature (Spampinato et al., 2018), is highly expressed at astrocytic level. To analyze in more details this aspect, we utilized also the mGluR3 positive allosteric modulator LY2794193. As expected, in the endothelium-astrocyte co-cultures, LY27 prevented the increase of permeability, and the gene expression of the pro-inflammatory cytokine IL-6, both caused by T&Y, and increased the expression of transforming growth factor- β (TGF- β). This last result is in line with what is reported in literature where mGluR3 stimulation led to an increase of TGF- β in astrocytes (Caraci et al., 2011). Therefore, also in the endothelium/astrocytes co-culture, mGluR3 has a major role in the modulation of BBB properties.

Interesting results were obtained when we analyzed microglia influence on BBB properties in the triple endothelial-astrocyte-microglia co-culture. In this case, in fact, the effects of LY37, that were similar to those described above for the endothelial monoculture and the endothelial/astrocyte co-culture, were counteracted by VU that caused blockade of mGlu2 receptor. So, in contrast to what observed before, these results suggest that mGluR2 has a role in the protection of barrier function in the triple endothelial-astrocyte-microglia co-culture. To investigate in more details the specific role of microglia and the selective involvement of mGlu2 and mGlu3 receptors, we decided to use a monolayer of microglia cells challenged with T&I to assess expression of inflammatory cytokines and chemokines. The results obtained showed that LY37

attenuated the microglial inflammatory activation by T&I. All these effects were modified in the presence of blockade of mGluR2, confirming a role for mGlu2 receptor in this anti-inflammatory microglial response. Stimulation of mGlu2 and mGlu3 also increased the expression of the neurotrophic factor BDNF, an effect already reported in literature (Venero et al., 2002). These data are in apparent contrast with mGluR2 mediating a pro-inflammatory effect. Data from the literature report in fact that activation of 2 mGlu receptors stimulated the production of pro-inflammatory cytokines (Pinteaux-Jones et al., 2008; Taylor et al., 2002, 2005). One important aspect of mGlu2 and mGlu3 behavior is their capability to form homo- and heterodimers (Kammermeier, 2012; Vafabakhsh et al., 2015; Yin et al., 2014), a phenomenon particularly important for mGlu receptor function (Kniazeff et al., 2011). Although it is hard to conclude from the present data, we cannot rule out that the results we observed might be influenced by a change in the homo- and hetero-dimers proportion on microglial cells. The poor availability of very selective mGlu2 and mGlu3 receptor agonists/antagonists did not allow a clear identification of the specific role of each receptor subtype.

In conclusion, our results showed that mGluR3 and mGluR2 protect BBB functions in the presence of an inflammatory insult. In particular, the role of mGluR3 is prevalent in endothelial cells and astrocytes while mGluR2 appears to be involved in microglia. The effects observed are direct on the endothelial component where TEER, expression of tight junctions and consequently permeability are modified, but seem to be also indirect, through modulation of the inflammatory response on astrocytes and microglia.

2. *In Vivo* studies

2.1 Introduction

2.1.1 Animal models of Autism Spectrum Disorders

The Autism Spectrum Disorders (ASD) are a group of neurodevelopmental disorders generally diagnosed by three core behavioural symptoms, *i.e.* stereotyped repertoire, communication impairments and social dysfunctions (Hodges et al., 2020). However, besides the symptoms listed above there is an extensive range of additional symptoms, including cognitive dysfunctions, restricted interests, hyperactivity and impulsivity, that are often overlapping with different neuropsychiatric disorders, such as schizophrenia, attention-deficit/hyperactivity disorder, anxiety and obsessive-compulsive disorder (Guo & Commons, 2017; Murray, 2010).

ASD incidence has increased intensely during the last decade, by reaching the 1% estimation from the World Health Organization, that only accounts for approximately 16% of the global pediatric population; while European and US reports estimated around 1.4-2.5% in children of 8 years old on average based on population studies (Hodges et al., 2020; Sacco et al., 2023; Scattoni et al., 2023; Zeidan et al., 2022). The etiopathogenesis underlying the development of this type of diseases is very complex and still largely unknown, at the base genetic and environmental factors are hypothesized as the most involved. Although there is indication for a solid influence of genetic factors, with 60–90% concordance in monozygotic twins and 4–10% in dizygotic twins, (Muhle et al., 2004) with a study suggesting an incidence near to 30% for the seconds (Hallmayer et al., 2011), ASD seems to be polygenic, with hundreds of contributing loci (Bncur, 2011), among which the majority seems to be *de novo* mutations and copy number variants (Huguet et al., 2013; Iossifov et al., 2014; Jeste & Geschwind, 2014). ASD is more frequently diagnosed in boys, with the sex ratio of 4.3:1 (Newschaffer et al., 2007) for classic autism and 11:1 in Asperger Syndrome (Gillberg et al., 2006), thus suggesting that some of the genes involved are X-

linked. Moreover, only 10–15% of ASD cases can be directly associated with monogenic disorders, such as Rett syndrome, Fragile X syndrome or tuberous sclerosis complex (Chévere-Torres et al., 2012; Hagerman et al., 2011; Zoghbi, 2005), while most cases are described as idiopathic.

Recently, multiple factors are emerging as possibly implicated in this pathological picture, such as immune system, diet, metabolic system, and gastrointestinal system.(Napolitano et al., 2022; Samsam et al., 2014). In addition, an overgrowth of cortical gray matter and decreased migration of neurons during early development with consequences in heterotopias, thinning of the corpus callosum and malformations of the ventricular system have been described in ASD patients (Courchesne et al., 2011; Ecker et al., 2012; Wegiel et al., 2010). However, the presence of such alterations in the population of ASD patients is not enough to consider them main causes of the behavioural phenotype related to ASD. Other proposed impairments include imbalance in excitatory and inhibitory neurotransmissions, atypical formation of dendritic spines, dysfunctions in secondary messenger systems, extracellular matrix and synaptic protein and neuroinflammation during fetal development and later in life (Persico & Bourgeron, 2006). Furthermore, around 30% of autistic individuals also suffer from other mental health conditions, such as epilepsy (Tuchman & Rapin, 2002), anxiety disorders (van Steensel et al., 2011), obsessive compulsive disorders or attention deficit hyperactivity disorder (Leyfer et al., 2006).

Considering that, studies involving humans are frequently influenced by willful variables and biases and some differing results and inconsistencies in clinical evaluations have also been noted. In this regard, animal models can be really helpful to longitudinally study and understand behavioural alterations mimicking human symptoms in a translational way and to investigate the underpinning neurobiological mechanisms (Grosso, Galvano, et al., 2014; Grosso, Pajak, et al., 2014). To this purpose, the BTBR T+ Itpr3tf/J (BTBR) mice are an inbred mouse strain that well reproduces the principal behavioural deficits of ASD (Blanchard et al., 2012; Guo & Commons, 2017). Among the different ASD mice models, the

BTBR mouse strain shows strong and evident deficits in reciprocal social interactions, altered ultrasonic vocalization and repetitive stereotyped repertoire (Scattoni et al., 2013; Wöhr et al., 2011). Furthermore, BTBR mice show cognitive and emotional deficits typical of the psychiatric comorbidities of ASD (Chao et al., 2018; Chao, Marron Fernandez de Velasco, et al., 2020).

Regarding the social behaviour, in the three-chamber test for social approach, BTBR males failed to show a preference for the social chamber (Meyza et al., 2013). In addition, in the Visible burrow system (VBS) test, BTBR mice showed reductions in all interactive behaviours, including approach, follow and grooming another mouse, while they showed increases in non-interactive behaviours, like self-grooming and being alone (Bove et al., 2018; Pobbe et al., 2010). In social proximity test (Defensor et al., 2011) performed involving same-strain pairs, BTBR male mice showed decreased nose tip-to-nose tip, and upright behaviours, therefore suggesting that BTBR mice have an aversion for reciprocal frontal (face-to-face) orientations. In the Social conditioned place preference test, BTBR mice conditioned for 10 days didn't show a preference for the social chamber during testing, while a preference for this chamber was retrieved in C57BL/6 (BL6) mice, hence highlighting that BTBR mice showed social motivation deficits (Pearson et al., 2012). Indeed, BL6 mice are commonly used as control in several studies involving sociability, due to their normo-social behaviour.

Moreover, BTBR mice also displayed communication impairments, by emitting an uncommon pattern of vocalizations. Indeed, BTBR adult mice emitted a smaller number of ultrasonic vocalizations (USVs) in male-to-male, male-to-female, and female-to-female encounters (Scattoni et al., 2011). Furthermore, BTBR male mice also deposited few odor marks in the presence of female urine (Roullet et al., 2011; Wöhr et al., 2011). In this regard, some studies suggested that elevated USVs in BTBR pups may resemble the high levels of inconsolable crying in infants that are later diagnosed with autism (Scattoni et al., 2008); while reduced USVs and olfactory marking in BTBR adults are paralleling to communication difficulties in autistic children (Roullet et al., 2011; Scattoni et al., 2011; Wöhr et al., 2011).

Regarding the repeated and stereotyped behaviours, numerous studies have shown that BTBR mice displayed elevated levels of general self-grooming (Amodeo et al., 2012; McFarlane et al., 2008; Pearson et al., 2011; Silverman, Tolu, et al., 2010). BTBR mice also showed high levels of bar biting (Pearson et al., 2011), and marble burying (Amodeo et al., 2012), that are two additional indicative parameters of stereotyped motor behaviour in laboratory rodents (Londei et al., 1998; Nevison et al., 1999). In the hole board test, BTBR mice showed a preference for a particular corner hole (Moy et al., 2008). The same was retrieved in the novel object test, where BTBR mice introduced in a box containing four novel objects in each corner showed increased preference for individual objects, by significantly spent more time exploring the first and second most investigated objects and reduced time exploring the smallest preferred object compared to controls. Moreover, BTBR mice showed more episodes of visiting the objects in a specific order (Pearson et al., 2011) due to the cognitive inflexibility characterized by resistance to change that has been correlated with stereotypies (Tanimura et al., 2008). Nonetheless, cognitive inflexibility manifestation in autism is highly variable (H. M. Geurts et al., 2009; Zandt et al., 2007), maybe because tests for resistance to change require that mice first have to learn an established routine and then to start a new routine. A relevant paradigm to study cognitive inflexibility is the Morris water maze test. This test is commonly used for studying spatial learning by measuring the time necessary to find a hidden, submerged platform, first during an acquisition phase, then in a new location in a reversal phase. In this test, BTBR mice had normal results in the acquisition phase, but showed reduced performance in finding the platform (M. Yang et al., 2012) and did not show a preference for the quadrant containing the platform (Moy et al., 2007) during the reversal phase. Furthermore, in the spatial reversal learning test (Amodeo et al., 2012), BTBR mice showed normal acquisition, but reduced results in reversal phase. However they performed normally in both phases when in one arm a reinforcement was continuously present (Amodeo et al., 2012; Moy et al., 2007), hence indicating that

diminished reversal learning and cognitive flexibility deficits in BTBR mice are dependent on the type of reinforcement and the percentage of reinforcement.

Regarding anxiety-related and stress-related behaviours, in the elevated plus maze tests and in the light/dark exploration test used for evaluating anxiety-like behaviours, results on BTBR mice have been contradictory. The elevated plus maze (EPM) is a test generally used for evaluating general anxiety by measuring the number of entries and duration of time spent in an open arm compared to a closed arm, with more open arm activity usually indicating less anxiety. In the EPM, the results reported are different and variables, indeed BTBR mice showed normal (Moy et al., 2007), high (M. Yang et al., 2009), and low (Pobbe et al., 2011) number of entries into the open arms, and showed, at the same time, normal (Benno et al., 2009; Moy et al., 2007; M. Yang et al., 2009) and low (Pobbe et al., 2011) open arm durations. While in the elevated zero maze, where the maze has a “zero” form, several studies reported that BTBR mice spent more time in the open quadrants (McFarlane et al., 2008; Pobbe et al., 2011). The light/dark exploration test is also utilized for evaluating anxiety-related behaviours and is based on the aversion of mice to intensely lighted areas. In this test, the BTBR mice did not showed difference compared to BL6 mice in the number of transitions between the light and dark compartments, the time spent in the dark chamber or in the latency to enter the dark chamber.

The mouse defense test battery is another test where is evaluated the anxiety-like behaviour, by using a hand-held anesthetized rat to evaluate behaviours in response to predator exposure in various contexts (Blanchard et al., 2003). In the MDTB, BTBR mice showed more vocalizations when the contact with the predator was forced, meanwhile BTBR mice also allowed the predator to approach within shorter distances before escaping in an approach/avoidance setting (Pobbe et al., 2011). Moreover, BTBR mice revealed increased locomotion, increased wall jumping and defensive standing behaviours after, but not before, the predator was introduced, suggesting increased predator-induced anxiety.

Regarding the stress-reactive behaviours, BTBR mice seem to respond differently depending on the stressor stimulus. Indeed, they showed normal reactivity to an acoustic alarm stimulus and tail flick, but they showed also a normal prepulse inhibition and low reactivity to a hot plate stimulus (Silverman, Yang, et al., 2010). Furthermore, whether the EPM test was preceded by tail suspension, BTBR mice showed increased anxiety-like behaviour (Benno et al., 2009). Lastly, in tests for evaluate depressive-like behaviours, BTBR mice showed low levels of immobility in the forced swimming test and the tail suspension test, suggesting a decreased tendency to depressive-like behaviour (Silverman, Yang, et al., 2010).

2.1.2 Neurochemical and biomolecular pathways underlying ASD symptoms

Considering their very specific behavioural profile, BTBR mice may represent a helpful model to better understand the neurobiological alterations that are the base of the complex heterogeneity of ASD symptoms, to identify possible biomarkers involved and, finally, to develop possible pharmacological treatments (Kas et al., 2019). Going into specifics, concerning repetitive and restricted behaviours, BTBR mice showed both “lower-order” motor stereotypies and “higher-order” cognitive stereotypies (Pearson et al., 2011), giving to this animal model the characteristics for being used to study the neurobiological pathways related to these behavioural dysfunctions. Pharmacological treatments to reduce repetitive behaviours are principally focused on impulsivity and irritability, by using typical and atypical antipsychotics, consequently they are utilized in limited cases. Nowadays, there are no drugs approved for an effective treatment of ASD repetitive symptoms (Aishworiya et al., 2022; McPheeters et al., 2011; Sharma & Shaw, 2012; Whitehouse & Lewis, 2015). Therefore, to discover more details on mechanisms related to stereotypies would offer a significant step forward to the development of novel pharmacological treatments for ASD repetitive repertoire. Although the pathophysiology of ASD is still not completely clear, a wide variety of imaging and post-mortem studies described the key role of corticostriatal and limbic circuits, by showing cortical, hippocampal and amygdalar anatomical

anomalies and functional alterations (Haznedar et al., 2000; Hollander et al., 2005; Kemper & Bauman, 1998; Langen et al., 2011; Zilbovicius et al., 2000).

Regarding neuroanatomic structure, BTBR mice show severe anomalies of the corpus callosum, often accompanied with a substantial reduction of the hippocampal commissure (HC) (Kusek et al., 2007; Wahlsten et al., 2003), a collapse of lateral ventricles (LV) (Meyza et al., 2012) with lateral displacement of the hippocampi (Mercier et al., 2011). The length of the subventricular zone (SVZ) is decreased in BTBR mice and both the concentration of laminin and heparan sulfate (HS) therein are diminished (Meyza et al., 2012). This is probably an effect of downregulation of exostosin-1 Ext1 expression in BTBR mice, because Ext1 is involved in synthesis and elongation of HS polysaccharides, that in the appropriately sulfated form is considered a guidance molecules crucial for the crossing of corpus callosum (CC) neuronal fibers across the midline during development (E14-E17) (Busse et al., 2007). Indeed the Ext1 KO and Hs6st1^{-/-} and Hs2st^{-/-} mice, missing sulfotransferases responsible for sulfation of the polysaccharide HS chain, showed a lack of CC, and altered guidance of other neuronal tracts in the brain (Conway et al., 2011; Inatani et al., 2003). Although the intact CC is a prerequisite for development of normal social behaviours in mice, there are evidence that in postnatal day (PND) 7 there was a lesion of CC in BL6, but mice social behaviour was not affected (M. Yang et al., 2009). Unfortunately, it is difficult to understand the effect of earlier disruption of CC, and also Ext1 KO, Hs6st1^{-/-} and Hs2st^{-/-} mice did not postnatally show any social behaviour deficits. Probably, it seems that the absence of the guidance factors influencing the CC genesis, more than the functioning of the CC itself, could be implicated in the behaviours of BTBR mice. Indeed, conditional knockouts of Ext1 (Ext1CKO, CaMKII-Cre2834; Ext1flox/flox, (Irie et al., 2012), in which Ext1 is disrupted from PND21, exhibited abnormal social interactions, improved repetitive behaviours and reduced ultrasonic vocalizations. The Ext1CKO animals also showed decreased neuronal activation to social stimuli in the amygdala and damaged excitatory synaptic transmission therein, suggesting

that HS is involved in correct functioning of glutamatergic synapses. As regarding BTBR mice, it has been reported decreased neurogenesis in the dentate gyrus of the hippocampus (Stephenson et al., 2011) and such characteristic has been linked to the autism-like behavioural phenotype (L. Wei et al., 2011), proving that youthful neurogenesis is crucial for development of normal social repertoire. It has also showed that BTBR mice also have reduced general levels of plasma sulfates (Corley et al., 2012), which is in accordance with clinical data from ASD patients (Adams et al., 2011) and could indicate that atypical sulfate metabolism or excretion may be affecting HS levels in these mice.

For their metabolism, BTBR mice were studied many years before the knowledge of their social behaviour impairment. Indeed, BTBR leptin *ob/ob* mice are susceptible to develop type-2 insulin resistant diabetes and were studied as an animal model for this disease. Even in the absence of leptin, *ob/ob* BTBR mice have been shown to have elevated plasma insulin levels compared to BL6 mice (Stoehr et al., 2000). Male BTBR mice are prone to develop abdominal obesity, hypertriglyceridemia and hypercholesterolemia, with a decreasing insulin-stimulated uptake of glucose to white adipose tissue (Flowers et al., 2007; Ranheim et al., 1997). Insulin resistance in BTBR males is correlated with increased expression of several genes in the adipose tissue, including cytoskeletal and focal adhesion molecules (*Actb*, *Acta2*, *Flna*, *Diap*, *Rhoa*), proteins involved in inflammation (*Pir's*, *Fos*, *Ptgs2*, *Syk*, *Cflar*, *H2ab1*) and phosphatases (*Ptprs*, *Pp2r4*, *Pp6c*) (Flowers et al., 2007)

Regarding the immune system, BTBR mice have a different immune response from those of BL6 mice on both humoral and cellular levels. In particular, they showed higher basal levels of plasma IgG, IgE, anti-brain antibodies (Abs), and also of brain tissue deposited IgG and IgE (Heo et al., 2011), together with higher expression of proinflammatory cytokines, such as IL-33, IL-18 and IL-1b. Moreover, the brain–blood barrier permeability measured by Evans blue was comparable in BTBR and BL6 mice and, the number of mast cells in the brain was significantly higher in BTBR mice, thus suggesting a constant neuroinflammation.

In addition, the microglia of BTBR differed from BL6 for the number of MHC class II-expressing cells. However, these immune differences could not be related to differences in NF- κ B signaling, the studies showed that the phosphorylation of NF- κ B p65 are not significantly altered in the cerebellum and cortex of BTBR mice and in a similar way both the expression and phosphorylation of NF- κ B p65 are not significantly altered in the cerebellum and cortex of autistic subjects as compared with the control subjects.(Malik et al., 2011; Manzardo et al., 2012).

From a genetical point of view, BTBR is an inbred strain developed originally at Columbia University, by crossing mice carrying the wildtype T (brachyury) gene with the stock carrying tufted (tf) mutation. Subsequently the strain was outcrossed to c129 mice and then maintained by inbreeding. Several studies aimed to investigate the genes involved in the behavioural differences, including social behaviour of these mice. Investigations performed in large scale compared to gene expression in 10 inbred mouse strains, including BTBR and BL6 mice, tried to correlate this expression with motor coordination in the rotarod test. The results indicated a correlation of motor activity with the expression of several transcripts in the cerebellum, hippocampus and amygdala, but did not focus on between strain differences and only provided a general statement regarding a cluster analysis in which BTBR mice were placed closest to BALB/cByJ mice, in terms of overall expression patterns (Nadler et al., 2006). In another study performed analyzing the entire genome on BTBR mice backcrossed to obtain an obese phenotype (ob/ob), 3 main loci that differentiated BTBR from BL6 mice and influence plasma glucose and/or insulin levels were identified; the first of these was situated on chromosome 2 and was related to insulin sensitivity and the other two were situated on chromosomes 16 and 19 and influenced quickly glucose and insulin levels (Clee et al., 2005). Analysis of 124 assumed autism candidate genes showed that there are four genes with nonsynonymous coding single nucleotide polymorphisms (SNPs) that differentiate BL6 from BTBR mice. Two of them, Smo (smoothened, signaling protein regulated by Sonic hedgehog) and Pkd1 (polycystic kidney disease 1) were common for other inbred strains.

Regarding the other two, Slc6a4 encoding serotonin transporter was different in BL6 mice, while Kmo encoding kynurenine 3-hydroxylase was different in BTBR mice. This gene is constituted by 3 different SNPs situated on chromosomes 9 and 13, among which the one situated on chromosome 13 is a cytosine/thymine substitution in a sequence that code for a transmembrane part of the protein. This part is a conserved domain identical in mice and humans (McFarlane et al., 2008) and may constitute a relevant biomarker, because Kynurenine 3-hydroxylase is involved in synthesis of kynurenic acid, a potent antagonist of glutamate and nicotinic receptors, involved in neuroprotection, dendritic spine formation and dopamine release (Alkondon et al., 2004; H.-Q. Wu et al., 2007; Yu et al., 2004). Further analysis showed that BTBR mice also carry a Disrupted in Schizophrenia (Disc1) mutation, a 25 bp deletion causing a frame shift and a premature termination of translation of the protein. It is located on chromosome 7 [<http://jaxmice.jax.org/strain/002282.html>].

Interestingly, a large study collected a wide range of transcriptomic and proteomic data indicating a different expression of a big variety of genes and proteins in the hippocampus and cortex of BTBR and BL6 mice. Among those genes and proteins, brain derived neurotrophic factor [BDNF], p21-activated kinase type1 [Pak1] and cortistatin [Cort] were downregulated in BTBR hippocampus and cortex (Daimon et al., 2015), while some members of MAPK signaling pathway, were both up and downregulated. (Daimon et al., 2015) These data were in line with previous studies that have described a BDNF deficiency in adult BTBR mice (Scattoni et al., 2013; Stephenson et al., 2011), MAPK signaling interruption (Faridar et al., 2014; Seese et al., 2014) and an increased p-ERK levels (Seese et al., 2014). Such high levels of p-ERK in the prefrontal cortex was correlated with deficits in juvenile sociability (Faridar et al., 2014) and in adult memory formation (Seese et al., 2014).

Furthermore, Wei et colleagues in a proteomic study showed that BTBR mice have down-regulated levels of the stable tubule only polypeptide protein (STOP) and myelin-related proteins, such as myelin basic protein (MBP) and myelin

associated glycoprotein (MAG) (H. Wei, Ma, Ding, et al., 2016). Accordingly, histopathological examination of BTBR brain tissue showed a reduction of myelin markers, such as CNPase and MBP and an increase in the oligodendrocyte precursor NG2 (Stephenson et al., 2011). Analysis of fetal brain proteins demonstrated reduced levels of glial fibrillary acidic protein (GFAP) and increased BDNF and MBP levels in BTBR compared to FVB/NJ mice. Concerning BDNF, the upregulation of its expression at an early developmental stage might be confirmed by clinical data showing increased plasma's level of BDNF in children with ASD (M. Wang et al., 2015).

An interesting study performed proteomic analysis in hippocampi and cortices in aged BTBR mice, by identifying elevated levels of some synaptic proteins, including spinophilin, Synapsin 1 (and p-Synapsin 1), PSD95 and NeuN in the hippocampus and spinophilin, p-Synapsin 1 and NeuN in the cortex (Jasien et al., 2014), together with reduced BDNF levels in these aged animals.

In addition, some genes related to ASD are known to modify the neural structure, function and connectivity, but also influencing neurotransmission and neurotropism (Gandhi & Lee, 2020; Rubenstein, 2010). Among the neurotrophins family, particularly BDNF and Nerve Growth Factor (NGF) play an essential role in neurodevelopmental processes which are atypical in ASD pathology (Theoharides et al., 2015). BDNF is crucial for the appropriate cerebral development and significantly associated to synaptic plasticity. It has been recently proposed as a possible diagnostic marker in children suffering from ASD since different line of evidence have reported its involvement in ASD onset and development (Barbosa et al., 2020; Crespi, 2019; Hellings et al., 2017). On the other hand NGF has a key role in the regulation of nerve-cell growth, survival and differentiation, being mainly expressed in the central nervous system, particularly in the cerebral cortex, hippocampus (HIP) and amygdala (AMY) (von Richthofen et al., 2003). Recently, few studies linked NGF and ASD, by describing an involvement of the NGF signaling pathway in the pathogenesis of ASD (Lu et al., 2013; Theoharides et al., 2015), also considering the central role of this neurotrophins

in the immune modulation and in the induction of the release of different neuropeptides and neurotransmitters (Minnone et al., 2017), such as glutamate and dopamine (DA) (Paredes et al., 2007). On the other side, catecholamines are notoriously known to increase NGF content (Furukawa et al., 1986), thus resulting in a constant crosstalk between these two pathways. Concerning this, catecholamines, such as DA and noradrenaline (NA) are known to play a fundamental role in the modulation of executive functions, attention, impulsivity, and emotional state, which are all processes disturbed in ASD (Aston-Jones, 2005; Sara, 2009). In this context, recent studies have reported a possible implication of DA neurotransmission and metabolism in ASD development (DiCarlo et al., 2019; Mandic-Maravic et al., 2021; Pavăl & Micluția, 2021), whereas several evidence demonstrated that neuropathological changes associated to the noradrenergic system often happen in ASD (Beversdorf, 2020; M. Kubota et al., 2020; Leshem et al., 2021). Moreover, an imbalance of the excitatory-inhibitory synaptic transmission has also been related to ASD pathology (Yizhar et al., 2011). In support of this, alterations of glutamate and gamma-aminobutyric acid (GABA) receptors expression have been described in *post-mortem* brains of ASD patients (Blatt & Fatemi, 2011) and decreased GABA levels have been detected in different brain regions of ASD children (Horder et al., 2018). In addition, brain regions with a massive cholinergic innervation, such as prefrontal cortex (PFC), hippocampus (HIPP) and amygdala (AMY), are linked to social cognition, thus resulting remarkably involved in ASD pathophysiology (Tanimizu et al., 2017). Indeed, lower basal levels of extracellular acetylcholine were described in BTBR prefrontal cortex (McTighe et al., 2013). In this regard, it has been showed that low acetylcholine (ACh) levels in mice presented reduced social interactions and that, by treating with acetylcholinesterase inhibitors such as Donepezil, this impairment might be contrasted (Kljakic et al., 2021) and both social approach and social novelty preference in the three chamber social approach task were improved (Karvat & Kimchi, 2014). Although there were no differences in acetylcholinesterase (AChE) expression in the forebrain of BTBR

and BL6 mice (Stephenson et al., 2011). Regarding drugs acting on Ach transmission, such as Oxotremorine, an antagonist of muscarinic receptor, it was able to reduced repetitive behaviours (grooming and marble burying) in BTBR mice (Amodeo, Yi, et al., 2014), while concerning nicotinic receptor agonist, nicotine administered in low doses enhanced sociability and in high doses reduced grooming (L. Wang et al., 2015).

As reported above, BTBR mice present impairments in monoaminergic (5HT, DA) and cholinergic transmission that have been related with the seriousness of symptoms in ASD patients (Cartier et al., 2015; Gangi et al., 2016; Muller et al., 2016). BTBR mice showed higher hippocampal 5HT_{1A} receptor presence (Gould et al., 2011, 2014), as well as lower [(3)H] cyanoimipramine and citalopram binding to the serotonin transporter SERT (Gould et al., 2011). Increased 8-OH-DPAT-stimulated (8-OH-DPAT is a full agonist of 5HT_{1A} receptor) GTPγS binding in the BTBR hippocampus indicated an elevated capacity of 5HT_{1A} receptors to stimulate G-proteins (Gould et al., 2011). At the same time, less serotonergic cells were detected in the hippocampus of BTBR compared to BL6 mice (Guo & Commons, 2017). On the other hand, this was compensated by improved number of serotonergic cells in brain areas projecting to the hippocampus, especially in the median raphe. Moreover, dopaminergic neurotransmission is also altered in the BTBR mice. In particular, D2 receptor function seems to be compromised and the result is a minor activation of the reward system in a functional magnetic resonance imaging (fMRI) scan after administration of a dopamine reuptake inhibitor. (Squillace et al., 2014). Several drugs acting on monoaminergic pathways have been investigated for their efficacy in ameliorating autism-like behaviours in the BTBR mouse (Meyza & Blanchard, 2017). For instance, risperidone, that act on both serotonergic and dopaminergic transmission had no effect on sociability (Chadman, 2011; Gould et al., 2011) and anxiety (Chadman, 2011), but it showed a reduction of repetitive behaviours (Gould et al., 2011) and an enhancement of reversal learning in the probabilistic learning paradigm (Amodeo, Jones, et al., 2014). The reduction in grooming was also observed after

intraperitoneal administration of a 5HT_{2A} antagonist, M100907 (Amodeo et al., 2016, 2017; Amodeo, Jones, et al., 2014). In opposite way, fluoxetine, a SERT blocker, increased social approach, but had no effect on the repetitive behaviour (Gould et al., 2011) and anxiety level (Chadman, 2011). Similar effects were described for buspirone, a 5HT_{1A} receptor partial agonist (Gould et al., 2011). Moreover, administration of tryptophan from diet was able to enhance social approach without influencing social novelty preference, it was also able to reduce the excessive grooming and the number of marbles buried in the BTBR mice (W. Q. Zhang et al., 2015). Furthermore, paracetamol, beyond its analgesic effect, also acts as 5HT agonist, and it was able to improve social approach in the three-chamber apparatus (Gould et al., 2012). In addition, different psychostimulant drugs that act on the dopaminergic circuitry, are able to affect the behaviour of BTBR mice, such as the D-amphetamine that increased the number but reduced the duration of social sniffing and reduced the excessive grooming in BTBR mice (Silverman et al., 2013), the while (-)-trans-9-tetrahydrocannabinol (THC) was able to reduce spontaneous, repetitive wheel running (S. Onaivi et al., 2011).

Regarding the serotonergic system, drug enhancing such neurotransmission, including citalopram (Crowley et al., 2005), fluoxetine and buspirone, demonstrated a reduction of depressive-like behaviour and an increase in sociability (Chadman, 2011; Gould et al., 2011). Moreover, In SERT and 5HT_{1a} receptors binding were observed some differences between BTBR and BL6 mice, but there was no difference in the 5HT_{2a} receptor binding to (Tanimura et al., 2008). Indeed, it has been demonstrated that risperidone a DA/5HT_{2a,c} antagonist/blocker had no effects on social behaviour (Chadman, 2011; Gould et al., 2011), although it decreased marble burying (Gould et al., 2011). In ASD patients' risperidone has been utilized to repress aggression, but had no effect in enhancing the social behaviour (West et al., 2009). Regarding glutamatergic neurotransmission, a decrease in repetitive self-grooming behaviours was detected in BTBR mice after administration of 2-methyl-6-(phenylethynyl)-pyridine (MPEP), a mGluR5 receptor antagonist (Burket et al., 2011), which might

suggest that alterations in NMDA signaling may cause excessive self-grooming in these mice. Supporting this, the administration of GRN-529, a GluR5 antagonist, was able to decrease repetitive behaviours in BTBR mice and increase different parameters related to social approach and social interactions (Silverman et al., 2012), while an improvement of GABA neurotransmission induced by diazepam moderated BTBR jump escape, vertical position and crawl under behaviours in the social proximity test (Defensor et al., 2011).

Going deeper to the Excitatory/Inhibitory (E/I) neurotransmission imbalance, it has been proposed as pivotal mechanism in autism-like behaviours (Uzunova et al., 2016). In this direction it was reported abnormalities in glutamate release in BTBR mice, specifically were reported decreased baseline and KCl-evoked glutamate release levels from the cortical synaptoneurosome in both young adult (8 weeks-old) and aged BTBR mice. (H. Wei et al., 2015; H. Wei, Ma, Liu, et al., 2016). Moreover, during the early postnatal period, in the CA3 region of BTBR mice hippocampi, an increase in GABAergic and a reduction in glutamatergic currents was observed (Cellot et al., 2016). Conversely, the frequency of spontaneous inhibitory post-synaptic current was significantly reduced, whereas the amplitude and the frequency of spontaneous excitatory post-synaptic current was increased in BTBR hippocampal slices of adult mice. Both currents were restabilized to a normal function by the treatment with clonazepam, a GABA-A receptor agonist (S. Han et al., 2014). In support of this hypothesis, different pharmacological studies reported that by acting on E/I balance was possible to reverse at least one characteristic of autism-relevant behaviours in the BTBR mouse (Meyza & Blanchard, 2017)

Regarding their hormonal situation, BTBR mice showed increased basal plasma level of corticosterone (Benno et al., 2009; Silverman, Yang, et al., 2010) Although it was not associated by an increase in CRF peptide levels or its mRNA expression after exposure to stressors (Silverman, Yang, et al., 2010), but they had higher levels of glucocorticoid receptors in the CA1 (but not CA2) field of the hippocampus, thus possibly indicating a modified negative feedback loop for the

hypothalamo-pituitary-adrenal axis (HPA axis) (Silverman, Yang, et al., 2010). The study also suggested that elevated basal corticosterone levels might be linked to insulin resistance previously observed in BTBR and to higher levels of plasmatic oxytocin compared to BL6 mice (Silverman, Yang, et al., 2010). Moreover, BTBR female mice also expressed increased circulating progesterone levels and enhanced 5 α pregnan-3 α -ol-20-one levels in both plasma and hypothalamus (Frye & Llaneza, 2010), suggesting that the higher basal CORT described in these mice may be a consequence of an reduced feedback of 5 α pregnan-3 α -ol-20-one on the HPA axis. In a similar way, BTBR male mice had higher levels of plasmatic testosterone (Flowers et al., 2007) that from one side may be responsible for their insulin resistance, since testosterone could inhibit the production of adiponectin (A. Xu et al., 2005), and on the other side might stimulate the expression of TNF- α in macrophages (Ashcroft & Mills, 2002), by modulating their immune reactivity.

Furthermore, it was demonstrated that an Oxytocin (OT) signaling deficiency (observed in oxytocin knockout mice, OTKO; oxytocin receptor knockout mice, OTRKO and CD38KO mice where the CD38 enzyme that is a multifunctional enzyme involved in calcium signaling and Nicotinamide Adenine Dinucleotide (NAD⁺) metabolism, regulates the Ca²⁺ dependent secretion of oxytocin, was associated to social behaviour impairments. Such deficiency could be counteracted by intranasal supplementation of oxytocin, or by a non-peptide agonists of OTRs or by OT release promoting factors (Modi & Young, 2012).

Oxytocin exerts also a central role to maintain E/I imbalance due to its capability to promote the formation of GABAergic synapses (Theodosis et al., 2006). Indeed, it was reported that mice deficient of oxytocin receptors had a reduced number of GABAergic synapses in hippocampus that led to an increased rate of glutamatergic over GABAergic synapses and an increased convulsion susceptibility, together with impaired social behaviour and cognitive flexibility (Sala et al., 2011). The administration of a single dose of oxytocin was able to enhance the social information maintenance and to decrease repetitive

behaviours in ASD patients (Hollander et al., 2007). In addition, intranasal OT administration had positive effects on empathic accuracy (Guastella et al., 2010) and trust (Andari et al., 2010). However, chronic intranasal oxytocin administration in BTBR mice did not show an improvement in juvenile reciprocal social interactions and it did not influence repetitive self-grooming, open-field exploratory activity or fear-conditioned learning and memory (Bales et al., 2014).

To sum up, several molecular alterations, in terms of neurotrophins, catecholamines and aminoacids, have been described in both animal and human ASD studies, and, from an anatomical point of view, tendency towards overgrowth of PFC and AMY have been found in individuals with ASD compared to neurotypical controls (Bellani et al., 2013b, 2013a; Pardo & Eberhart, 2007). Furthermore, data from structural magnetic resonance imaging highlighted common cerebral abnormalities in ASD patients, that involved total brain volume, fronto-parieto-temporal and cerebellar regions (J. K. Lee et al., 2022).

2.1.3 The role of neuroinflammation in ASD

As widely reported in the first part of the study, neuroinflammation is a well-coordinated inflammatory response within the brain or spinal cord mediated by various groups of glial cells, particularly astrocytes and microglia, that acting releasing inflammatory mediators, namely cytokines, chemokines, reactive oxygen species, and secondary messengers, together with peripherally derived immune cells (DiSabato et al., 2016).

Inflammation and dysregulation of the immune system are clinical signs of ASD (Ashwood et al., 2011; Ashwood & Wakefield, 2006). Several studies theorized that chronic inflammation and neuroinflammation during the period of early brain development could trigger behavioural and cognitive losses, ultimately affecting the etiology of ASD (Ashwood et al., 2011; Liao et al., 2020; Sciara et al., 2020). In ASD children, higher plasma levels of pro-inflammatory cytokines, such as IL-1 β , IL-6, IL-17, IL-12p40, and TNF- α , were described (Inga Jácome et al., 2016). Indeed, elevated cytokine levels in plasma were correlated with

seriousness of ASD symptoms, in particular, increasing cytokine levels were related with reduced communications and unusual behaviours (Ashwood et al., 2011; Inga Jácome et al., 2016).

Chronic neuroinflammation might be implicated in the biogenesis of behavioural and cognitive impairments during the early brain development. Indeed, neuroinflammation plays a key role in developing and maintaining the dendritic spines in glutamatergic and GABAergic neurotransmission (Alabdali et al., 2014; El-Ansary & Al-Ayadhi, 2014). In this regard, cytokines can influence the length, location, and organization of dendritic spines on excitatory and inhibitory neurons and recruit and impact glial cell function around the neurons (Eroglu & Barres, 2010), with consequences on the development that may contribute to the ASD behavioural and cognitive symptoms.

It has been reported that acute or chronic maternal immune activation could modify proinflammatory cytokine levels in the fetal environment, by affecting neuroinflammation mechanisms, developing brain, and possibly led ASD development. (Toscano et al., 2021). Both Clinical and preclinical studies have suggested a link between maternal immune activation during pregnancy and the development of ASD (Atladóttir et al., 2009, 2010; A. S. Brown et al., 2014; Choi et al., 2016; B. K. Lee et al., 2015; Rudolph et al., 2018), by reporting a positive correlation between maternal autoimmunity and ASD development (Atladóttir et al., 2009). This association could be related to a mixture of a common genetic background and a possible prenatal mother's antibodies exposure or an alteration in the fetal environment during pregnancy (Atladóttir et al., 2009) that could cross the placenta and affect fetal brain development (Braunschweig et al., 2012; Dalton et al., 2003; Zimmerman et al., 2007).

In ASD patients, astrocytes reactivity was identified in several brain regions. A significant increase in GFAP was described in the *postmortem* superior frontal cortex, parietal cortex, prefrontal cortex, and cerebellum (Edmonson et al., 2014; Laurence & Fatemi, 2005). In addition, GFAP levels were also increased in fresh-

frozen brain tissue of the cerebellum, middle frontal gyrus, and anterior cingulate gyrus from ASD patients (Vargas et al., 2005).

Interestingly, children with ASD may also suffer of autoimmune disorders (Ormstad et al., 2018). It has been suggested that autism symptoms might originate from central nervous system imbalances due to chronic inflammatory reactions, with activation of microglial cells (Bjorklund et al., 2016; Vargas et al., 2005). In support of this hypothesis, ASD children had improved numbers of circulating monocytes, essential precursors for macrophages, dendritic, and microglial cell activation (Rodriguez & Kern, 2011; Sweeten et al., 2003). Microglial cells are the brain immune residence cells and it was described an increase in microglial density in cortices of ASD patients (Morgan et al., 2010; Tetreault et al., 2012; Vargas et al., 2005) and increase activated microglia in the striatum (Morgan et al., 2014), accompanied by no changes in the number of microglia cells in the amygdala (Morgan et al., 2014). Moreover, some methylated genes involved in microglial cell differentiation and synaptic pruning during brain development were affected in ASD (Nardone et al., 2014; The DDD Study et al., 2014). Confirming this, the progenies of the autoimmune ASD mice model showed hypermethylated DNA regions in numerous transcription factor subjects critical for early microglial development and immune activation, and in the cytokines IL-6, IL4, IL-8, and Jak-STAT, TNF, and mTOR signaling (Vogel Ciernia et al., 2018).

Furthermore, samples from cerebrospinal fluid, blood, and *postmortem* brain tissue from ASD patients showed increased level in pro-inflammatory cytokine, together with reduced levels of anti-inflammatory cytokines and activation of microglia with changes in the microglia-neuronal spatial organization (A. S. Lee et al., 2017; Morgan et al., 2010; Rodriguez & Kern, 2011). A major density of microglial cells were observed in the fronto-insular and visual cortex of the autopsy brains of ASD patients compared to control brains (Tetreault et al., 2012). In ASD patients, by using the microglia marker IBA-1 through optical fractionator analysis, it was described a major microglia density in the brain sections of the

dorsolateral prefrontal cortex, with evidently morphological alterations of microglia, including expansion of soma, retraction, and thickening of processes and extension of filopodia (Morgan et al., 2010). The same study had also showed a major microglial cell density in the gray-matter and an increased somal volume of microglia in the white-matter of ASD patients (Morgan et al., 2010), accompanied by signs of excessive microglia activation in amygdala (Morgan et al., 2014).

It usually happens that changes in microglia density can be accompanied by changes in microglia phenotype and morphology. In this regard, in ASD *postmortem* brain temporal cortex, six structurally and functionally microglial phenotypes were identified and quantified with IBA-1 immuno-staining. These results showed a minor density of ramified microglia and a major density of primed microglia in ASD subjects compared with controls (A. S. Lee et al., 2017). In another study using positron emission tomography and a radiotracer for microglia, an enhanced microglia activity in the cerebellum, fusiform gyri, anterior cingulate and orbitofrontal cortices was founded (Suzuki et al., 2013). In addition, a marked activation of microglia and astroglia was described in the cerebral cortex, white matter and in the cerebellum of autistic patients and also the cerebrospinal fluid reported an important increase in the proinflammatory cytokine MCP-1 (Vargas et al., 2005).

Furthermore, some microglial-specific markers, such as TREM2, DAP12, and CX3CR1, had a higher expression in the prefrontal cortex of autistic people compared to controls. In particular, the expression of TREM2 was the highest of all microglial markers in brain tissue from ASD patients, although it was lower in autistic than in controls when analyzed in the *postmortem* cerebellum (Edmonson et al., 2014). Another confirming result of neuroinflammatory profile of microglia activation in postmortem brain of ASD patients is the increased levels of proinflammatory cytokines (such as IL-1 β , IL-6, IL-8, INF- γ , and TNF- α) described in both brain and blood samples of autistic patients (X. Li et al., 2009; Vargas et al., 2005; H. Wei et al., 2011). In the middle frontal gyrus of ASD

patients an increase in TGF- β 1, increased MCP-1, IL-6, and IL-10 in the anterior cingulate gyrus were showed (Vargas et al., 2005). Whereas, regarding the dorsolateral prefrontal cortex of ASD patients, there were no changes in characteristic markers of microglial activation, such as IL-6, IL-1 β , and TNF- α (Chana et al., 2015). Elevated levels of TNF- α have been associated with altered pineal melatonin release and sleep dysfunction in ASD, since it was proposed that circadian dysregulation in ASD is strongly linked to amplified immune-inflammatory activity, that can lead to sleep disorders, cognitive and behavioural alterations (da Silveira Cruz-Machado et al., 2021). Coming back to the proinflammatory cytokines, IL-6 was increased in the frontal cortex and in the cerebellum (X. Li et al., 2009; H. Wei et al., 2011). Moreover, it was described an important increase in Th1 cytokine (IFN- γ), but not in Th2 cytokines (IL-4, IL-5, and IL-10) in brain tissues of ASD, thus indicating that ASD brain can excessively adapt its response through activation of the Th1 pathway compared to the activation of the Th2 pathway (X. Li et al., 2009). Both excitatory and inhibitory synapse formation and transmission were altered by increased levels of IL-6 in shape, length, and distribution of dendritic spines (H. Wei et al., 2012). These studies suggested an unusual microglial-specific gene expression in autistic brains, indicating as a consequence the microglial activation, that might play an important role in ASD development.

Regarding BTBR mice, in this strain they were demonstrated enhanced brain cytokine levels, such as IL-33, IL-18 and IL-1 β , an increased number of CD4+ T cells in the spleen, mesenteric lymph nodes and peripheral blood, and an increased proportion of microglial cells compared to controls (Heo et al., 2011). It has to be taken into account that when there is a Chronic neuro-inflammation, BBB allows the entrance to peripheral cytokines and immune cells in the CNS. Although the effects of peripheral cytokines on the CNS immune environment in ASD has to be fully explained yet, they were described some alteration in gene expression associated with BBB integrity and function (Fiorentino et al., 2016).

2.1.4 The role of oxytocin and microglia in ASD

As previously described, neuroinflammation plays an important role in mechanisms of ASD and other early brain developmental diseases. In addition, oxytocinergic system impairment appears to be crucially involved in ASD and other neurodevelopmental diseases development, particularly such disorders that implicate social behaviours deficits. Recently, possible novel functions of OT for the developing brain are progressively emerging. Indeed, OT seems to be able to modulate glial activity in neuroinflammatory states, in particular via the activation of astrocytes and microglia state (Knoop et al., 2022). A meta-analysis of 18 studies, comprising a total of 1422 patients, found lower levels of endogenous OT in the blood, plasma or saliva of children with ASD compared to neurotypical controls (Moerkerke et al., 2021). In this regard, changes in OT system during brain development were reported also in ASD animal model (Knoop et al., 2022).

Furthermore, studies on neuroinflammation, by using interleukin-1 β (IL-1 β) or lipopolysaccharide (LPS) administration to adult male rats, have demonstrated that these cytokines caused an increase in plasmatic OT levels, and a major activity of the immediate early gene c-FOS in all regions where OT was synthesized (Matsunaga et al., 2000). Similar increase in OT activity were described also in MG6 microglia cells following LPS stimulation (Maejima et al., 2022). Moreover, in a rat model of fetal growth restriction, increased pro-inflammatory cytokine levels and microglial activation were described, accompanied by a downregulation in hypothalamic OT levels in the neonatal pups (Mairesse et al., 2019). These results support a developmental and functional alteration of the OT system in inflammation-related neurodevelopment diseases. This hypothesis piloted some researchers to study whether the social deficits seen in ASD models, could be due to a lack in the hippocampal OT system (Bertoni et al., 2021; Meziane et al., 2015). To this purpose, several studies tested neonatal OT as a therapy in neuroinflammatory and neuroinflammation-related models. (Knoop et al., 2022).

Focusing on OT role on astrocyte and microglia state, the expression of OTR was described in several brain areas including hypothalamus, hippocampus, frontal cortex and amygdala (Baudon et al., 2022). Concerning astrocytes, electrophysiological experiments described a depolarization of the astrocyte membrane potential and an increase in calcium release after activation of the OT-receptor (OTR) (Kuo et al., 2009; P. Wang et al., 2017). In addition, OTR activation in astrocytes could possibly increase local astrocyte network activity (Wahis et al., 2021). Furthermore, the reactive astrocyte phenotype is associated with a decrease in the expression of OTR (Guttenplan et al., 2020; Hasel et al., 2021), indeed, it was reported that OT administration reduced astrocyte-specific GFAP immunoreactivity in the supraoptic nucleus (SON) of healthy adult male rat (P. Wang et al., 2017), and that such effect was related with the inhibition of ERK1/2 kinase and with the activation of protein kinase A (PKA), thus suggesting that the reduced effects of OT in astrocytes might involve the MAPK pathway (Baudon et al., 2022). Subsequently, a possible target for OT modulation of reactive Astrocytes was identified as TGF- β . A link between TGF- β and OT has been found in astrocytes purified from E16 rat embryos and cultured for 12 days, where TGF- β increased the binding of astrocyte OTR and astrocyte OTR mRNA levels (Mittaud et al., 2002). However, OT effects on astrocytes could aim to reduce their neurotoxic effects, since it was showed that OT application to astrocytes was able to cause the retraction of their processes from the synapse in healthy conditions in the rodent hypothalamus (Baudon et al., 2022). This OT-mediated changes in astrocyte cytoskeleton is probably due to an effect on microtubule/actin dynamics (Baudon et al., 2022; Y.-F. Wang & Hatton, 2007). These results might let hypothesize that in neuroinflammatory states, OT-mediated retraction of astrocyte processes from the synapse/gap-junction could form a spatial obstacle in order to avoid astrocytes neurotoxic effect (Baudon et al., 2022).

Regarding microglial response, several studies described that OT treatment was able to induce a shift in microglial phenotype and reactivity during the

inflammatory response (Knoop et al., 2022). Different *in vitro* studies showed that OT pre-treatment decreased the LPS-induced increase in the production of pro-inflammatory cytokines from primary microglia and macrophages (Szeto et al., 2017; L. Yuan et al., 2016). In addition, in OTR knock-out mice an improved microglial activation in the medial amygdala compared to WT mice was reported, as demonstrated by an increased Iba1-immunoreactivity, and a smaller presence of amoeboid-shaped cells (Miyazaki et al., 2016). In this regard, there are several possible mechanisms underlying the effect of OT on microglial reactivity during neuroinflammation. Firstly, a possible mechanism could be linked to the reduction of phagocytic activity via NADPH/ROS signaling, since the neurotoxicity mediated by activated microglia is extremely dependent on NADPH oxidase (Block et al., 2007). Supporting this, it was shown that the OT treatment in macrophages and THP-1 monocytes markedly reduced the production of NADPH oxidase-dependent superoxide (Szeto et al., 2008). Secondly, another possible mechanism is represented by the regulation of cytokines secretion via the NF- κ B/MAPK pathway. Indeed, the OT-induced reduction in pro-inflammatory cytokines expression in cultured BV-2 murine microglial cell line and primary microglia was related with decreased phosphorylated ERK1/2 and p38 MAPK protein levels (L. Yuan et al., 2016). Strictly related to MAPK protein, there is the NF- κ B protein, a transcription factor that regulates pro-inflammatory cytokine production in microglia. Concerning this transcription factor, in a neonatal maternal separation rat model, it was shown that OT pre-treatment was able to reverse the increase in GFAP signaling and TLR4/NF- κ B signaling compared to vehicles (S. Xu et al., 2018). Furthermore, in a mouse model of early stage Alzheimer's disease, in which there were an increased in neuroinflammatory levels, it was found that OT release inhibited microglial activation and cytokine secretion by blocking the ERK/p38 MAPK and COX2/iNOS NF- κ B signaling pathways (Ye et al., 2022). Taken together, these data suggest that the NF- κ B/MAPK pathway could be an important factor contributing to the neuroprotective effect of OT on microglia.

Although OT might be able to influence the reactivity of microglia in neuroinflammation, it has not been clarified yet whether it can influence the phenotype. The presence of OTR expression on microglia are not completely described, but, although not in the neonatal age, several studies described microglial OTR expression predominantly in *in vitro* experiments (Knoop et al., 2022). To report some examples, LPS stimulation of P5 primary microglia induced a reduction in OTR mRNA levels, as assessed with RNA sequencing (Guttenplan et al., 2020). Moreover, LPS stimulation of P1-P2 primary microglia triggered an increase in OTR mRNA and protein levels (L. Yuan et al., 2016). A hypothesis about the effect of OT on microglia could involve the modulation of neurotransmitter systems, such as serotonin. It has been demonstrated in mice that serotonin can modulate microglial reactivity via the microglial 5-HT_{2b} receptor, and that restricted knock-out of this gene in microglia caused a prolongation of the inflammatory response 4 and 24 h post LPS injection (Béchéde et al., 2021). Therefore, such modulation could be mediated by OT, by having a positive correlation with serotonin levels and ultimately cause the release of serotonin in several brain regions of the limbic system, including the amygdala (Dölen et al., 2013; Nishimura et al., 2022). OT also upregulates the availability of the 5-HT_{1A} receptor (Lefevre et al., 2017) and this effect is due to the presence of OTR expression on serotonergic cells (Yoshida et al., 2009). These data imply that the serotonergic system is another potential, indirect, pathway of effect between OT and microglial activity in neonatal neuroinflammation. In support of this hypothesis, a study showed only the consequences of 5-HTR_{2B} knockout on neuroinflammation without studying the possible protective effect via increased upregulation of this receptor. (Béchéde et al., 2021).

OT may also influence the crosstalk between astrocytes and microglia. In this regard, it has been reported that inflammation caused by LPS injections in adult mice increased TNF- α reactivity levels of both astrocytes and microglia, although pre-treatment with OT was able to reverse only the TNF- α immunoreactivity of microglia and did not affect astrocytes, thus suggesting that the anti-

inflammatory effect of OT may be independent of astrocytic reactivity in adults (L. Yuan et al., 2016). Therefore, it could be hypothesized that OT treatment during inflammatory states might affect the number of astrocytes, but not their phenotype. However, it has been showed that OT injections between P2 and P6 in healthy rats enhanced the expression of hippocampal GFAP mRNA and protein two months later, without reporting changes in CD68 levels, a marker of microglial phagocytic activity (Havránek et al., 2017), hence hypothesizing that the effects of OT on astrocytes and microglia are dependent on the presence of inflammation (Bordt et al., 2019). Furthermore, positive link between TGF- β and OT has been suggested (Mittaud et al., 2002), since TGF- β can also have a role on microglial cells by binding to the TGF- β receptor type II, which phosphorylates TGF- β receptor type I (Zöller et al., 2018). In addition, it was described that TGF- β secretion reduced microglial activation by decreasing microglial production of pro-inflammatory cytokines, NO and oxygen-free radicals (W. Liu et al., 2011). Moreover, an *in vitro* study reported that, in baseline conditions, astrocytes involved TGF- β 2 to exert an immune-regulatory control on microglia, which repressed the microglial response to low inflammatory stimuli (Baxter et al., 2021).

2.1.5 β -Carotene impact on neuroinflammation and microglia

β -carotene belongs to the category of carotenoids, plant pigments that are precursors of vitamin A (retinol). As it has been well shown, carotenoids exert many beneficial effects. Indeed, they are principally known for their antioxidant properties, being main scavengers of ROS (Saini et al., 2015). B-carotene exhibits different pharmacological effects, such as anti-inflammatory, antioxidant and anti-tumoral (Liebler et al., 1997; Y. Yang et al., 2021; Y. Zhang et al., 2016). Several *in vitro* and *in vivo* studies reported that carotenoids can be involved in a variety of processes associated to the immune inflammatory response. In this regard, it has been described that these compounds influence both cellular and humoral mechanisms of immunity (J. S. Park et al., 2010; Seyedzadeh et al., 2014). Concerning anti-oxidant properties, it has been shown that β -carotene was

able to prevent the upregulation of heme oxygenase 1 expression in human skin fibroblasts (FEK4) exposed to UV-A (El-Agamey et al., 2004; Jomova & Valko, 2013), although being less effective in preventing lipid peroxidation (El-Agamey et al., 2004). In addition, β -carotene was able to counteract the production of NO, prostaglandin E2 and superoxide dismutase, while downregulating the expression of iNOS/cox-/NADPH oxidase proteins and mRNA, associated by inhibition TNF- α , thus showing important anti-inflammatory properties (Hadad & Levy, 2012). Moreover, β -carotene treatment reduced the oxidation level in porcine intestinal epithelial cells induced by LPS, by reducing the expression of Caspase-3 and mitigating the inflammatory response through the inhibition of NF- κ B, JNK2/STAT3 and JNK2/p38MAPK signaling pathways in macrophages (R. Li et al., 2019). Interestingly, a study of network pharmacology, by using molecular docking analysis, identified potential targets and possible signal pathways of the anti-inflammatory effects of β -carotene, by showing a good affinity with TNF, IL1B, and Leptin (LEP) (S. Wu et al., 2023).

Furthermore, the protective role of β -carotene against oxidative stress and neuroinflammation was demonstrated in a rat model of spinal cord injury, where β -carotene reduced the production of pro-inflammatory cytokines, including TNF- α , IL-1 β , IL-18 and COX-2, and repressed the activation of astrocyte in the spinal cord, acting a strong inhibition of the NF- κ B pathway activation (L. Zhou et al., 2018). In addition, β -carotene also showed a protective effect against myeloperoxidase-mediated oxidative stress and inflammation in a rat ischemic brain injury model, by reducing NF- κ B and myeloperoxidase activity in the brain. Moreover, β -carotene was able to significantly block Bcl-2-associated X protein and caspase-3 expression, while upregulated B-cell lymphoma-2 expression, suggesting a neuroprotective role (Althurwi et al., 2022).

Several studies reported that antioxidant and anti-inflammatory properties of carotenoids might improve the efficiency of cognitive function, by increasing neuronal efficiency or stabilizing the lipid-protein bonds in neuronal membranes (Kaulmann & Bohn, 2014; Mohammadzadeh Honarvar et al., 2017; Sorrenti et al.,

2020; Sujak et al., 1999). In addition, other neuroprotective mechanisms of carotenoids, such as improvement of communication between gaps and modulation of the functional properties of synaptic membranes (Stahl & Sies, 2001; Sujak et al., 1999), have been described. Interestingly, carotenoids might play a protective role in the development of depression through various mechanisms (Rasmus & Kozłowska, 2023), including a possible allosteric activation of BDN (S.-J. Park et al., 2021). Indeed, concerning depressive context, low levels of BDNF probably influenced by pro-inflammatory cytokines, such as IL-6 and TNF- α , might have a key role in the onset of such diseases (Videbech & Ravnkilde, 2004). Although further studies are necessarily required, it has been hypothesized that carotenoid could promote the BDNF activation and, considering their antioxidant and anti-inflammatory effects, they might ultimately exert a promising antidepressant effect. (Rasmus & Kozłowska, 2023).

Recently, different studies revealed that patients with ASD have reduced levels of oxytocin and its regulatory ectoenzyme CD38. Since β -carotene is an inducer of the CD38 enzyme (Avraham et al., 2019), the reduced CD38 expression, and consequent reduced OT levels, observed in ASD might be counteracted by β -carotene treatment, suggesting its use as a new therapeutic strategy in the treatment of ASD (Ebstein et al., 2011, 2012; Riebold et al., 2011). All trans-retinoic acids are potent CD38 inducers, but they are also toxic and teratogenic. Hence, β -carotene, being not toxic and not teratogenic, could represent a good safe and preventive strategy for ASD.

2.2 Aims of the Study

In this study, we used the BTBR strain, an idiopathic model of ASD, to better understand the complex mechanisms underlying ASD behavioural dysfunctions and, subsequently, to develop a possible preventive strategy.

In the first part of the study, we focused on BTBR behavioral abnormalities resembling human ASD symptoms and we investigated neurochemical and

biomolecular alterations underlying such behavioural alterations in brain regions crucially involved in ASD, such as PFC and HIPV.

In the second part of the study, we aimed to assess the effect of a possible preventive treatment, by using β -carotene administration during pregnancy, on the core ASD symptoms, such as stereotyped repertoire and social impairments, occurring in BTBR offspring. In addition, we evaluated the effects of β -carotene treatment on the modulation of neuroinflammatory components of ASD.

2.3 Materials and Methods

2.3.1 Animals

The study was focused on male sex since ASD shows a male prevalence, being males three times more diagnosed than females (Loomes et al., 2017). In the first part of the study, a total of 20 (2-3 per litter, randomly distributed) 10-weeks-old male mice, C57/BL6J (BL6) (Envigo, San Pietro al Natisone, Italy) and BTBR (Charles-River Italia, Milan, Italy), were used in this experimental study. They were raised at constant room temperature ($22 \pm 1^\circ\text{C}$) with relative humidity ($55 \pm 5\%$), under a 12 h light/dark cycle (lights on from 7:00 AM to 7:00 PM) and free access to water and food. All experiments on animals and their care were carried out in accordance with the institutional guidelines of the Italian Ministry of Health (D.Lgs. n. 26/2014), the Guide for the Care and Use of Laboratory Animals: Eight Edition, the Guide for the Care and Use of Mammals in Neuroscience and Behavioural Research (National Research Council, 2004), the Directive 2010/63/EU of the European Parliament and of the Council of 22 September 2010 on the protection of animals used for scientific intents, as defined by Animal Research: Reporting of *In Vivo* Experiments (ARRIVE) guidelines 2.0. The experimental protocol was approved by the Italian Ministry of Health (protocol nr. B2EF8.24). Animals' health state was checked daily during experimental period. Moreover, to accomplish the 3R's principles, we performed the behavioural test battery on the same group of animals; in order to reduce as

much as possible, the number of animals used, and every procedure was performed with the aim to minimize their suffering.

2.3.2 β -Carotene Treatment

In the second part of the study was performed a treatment with β -carotene on BTBR mice. The treatment with β -carotene was performed through prenatal supplementation and it was carried out at different doses 8, 16, 32 and 64 mg/kg subcutaneously in seed oil in pregnant mothers twice a week during the three weeks of pregnancy on five female BTBR mice for each dosage and vehicle respectively.

2.3.3 Battery of behavioural tests

2.3.4 Marble burying test

During day 1, The marble burying test was performed in an unfamiliar standard polycarbonate cage (26 x 48 x 20 cm) filled with unscented bedding material distributed in an even layer that was 5 cm deep. Twenty-four marbles were positioned inside the cage without food and water. Each mouse was allowed to explore the cage for 30 min and then returned to its home cage. At the end of the test, an experimenter counted marbles buried, scoring only those buried by 2/3 or more of its surface.

2.3.5 Hole board test

During day 2, animals performed the Hole Board task. This test was carried out in a wooden box (40 x 40 x 35 cm) having 16 holes with a diameter of 3 cm on the ground and placed 5 cm from the floor, as described in literature (Souza et al., 2016). The test lasted 10 min and an automatic counter registered the number of times the animals poking the hole for a duration of at least 1 sec, reported as number of poking holes.

2.3.6 Open Field test

During day 3, the Open Field test was performed. The test was executed according the literature (Carola et al., 2002). Briefly, the animals were left to explore an open field arena (40 x 40 x 35 cm) for 5 minutes. ANY-maze tracking

software version 7 (Ugo Basile-Varese, Gemonio, Italy) recorded and analyzed the locomotory activity of each mouse by measuring the distance travelled, the duration and the frequency of freezing behaviour and the time spent in the center and in the wall of the arena. Between one test and another, a solution at 70% of ethanol was used to clean arena floor and avoid inter-assay bias.

2.3.7 Social Interaction test

During day 4, the Social Interaction test was performed, as described in literature (Bove et al., 2022). The animals, after 2 days of individual housing, were left in the same large box used for the Open Field arena and let free to explore one plastic object, one paper cylinder, one ball and an unfamiliar stimulus mouse (same strain, sex and age of subject mouse) for 5 minutes. A camera recorded the test and a blind observer scored the frequency and duration of social behaviours from an investigative and affiliative point of view and also frequency and duration of non-social behaviour, considering the attitude towards objects, like sniffing, exploring and playing with the objects. (Kraeuter et al., 2019; Laviola et al., 2004).

2.3.8 Elevated Zero Maze test

During day 5, the Elevated Zero Maze test was performed, as described in literature (Tucker & McCabe, 2017). Precisely, a maze built in black acrylic in a circular track 10 cm wide, 105 cm in diameter and 72 cm high was used. Two opposed closed quadrants and two opposed open quadrants with black acrylic walls 28 cm high composed the maze. On the test day the animal was placed at a casually chosen boundary between an open and a closed zone, facing the closed area. After each trial, the maze was cleaned with a 70% ethanol solution. The test lasted 5 minutes and a blind observer evaluated the time spent in the open and in the closed corridors, expressed in seconds.

2.3.9 Post-mortem tissues analyses

After behavioural tests, the mice were sacrificed by cervical dislocation and brains were randomly divided for ex vivo analysis. PFC and HIPPO were removed

from brains, in accordance with the mouse brain atlas of Paxinos and Franklin, and frozen, stored at -80°C for subsequent analyses. To perform biomolecular studies, the tissues were homogenated and diluted 1:10 w/v in PBS buffer with 1:100 protease and phosphatase inhibitor (HALT inhibitors, Thermo Fisher Scientific, Cleveland, OH, USA) at 4 °C; while for neurochemical analyses, the samples were homogenated and diluted 1:10 w/v in perchloric acid 0.1M at 4 °C. In both cases, after dilution, a centrifuge at 10.000 x g at 4 °C for 10 min was carried out and the supernatants were analyzed.

2.2.10 Western blotting quantification

In this procedure, protein extracts were generated from fresh frozen PFC and HIPP tissues following samples homogenization with a Halt™ Protease and Phosphatase Inhibitor Single-Use Cocktail, EDTA-Free (Thermo Fisher Scientific, Cleveland, OH, USA). Subsequently, lysates were measured for total protein concentration using a Pierce™ BCA protein assay kit (Thermo Fisher Scientific, Cleveland, OH, USA) and the Multiskan™ FC Microplate spectrophotometer (Thermo Fisher Scientific, USA) at 570 nm. For SDS-PAGE the total amount of samples protein (40 µg) loaded in to the 4–15% Mini-PROTEAN™ TGX Stain-Free™ Protein Gels (Bio-Rad Laboratories Inc, Segrate (MI), Italy) for electrophoresis and then transferred on the Nitrocellulose membrane (Bio-Rad Laboratories Inc, Segrate (MI), Italy) by Pierce™ Power Blotter (Thermo Fisher Scientific, Cleveland, OH, USA). Then, the membranes were blocked in the 5% skimmed milk for 1 h at room temperature followed the incubation with rabbit monoclonal antibodies against NGF (ab52918; 1:1000, Abcam, Cambridge, UK), BDNF (ab226843; 1:1000, Abcam, Cambridge, UK), NF-κB (Ab16502; 1:1000, Abcam, Cambridge, UK), Occludin (Ab216327; 1:750, Abcam, Cambridge, UK), CD11-β (Ab133357; 1:1000, Abcam, Cambridge, UK), IBA-1 (Ab178846; 1:500, Abcam, Cambridge, UK), and mouse monoclonal antibodies against β-actin (ab8226; 1:1000, Abcam, Cambridge, UK), GFAP (Ab279290; 1:1000, Abcam, Cambridge, UK); overnight at 4°C. In add to this, the incubation (1 h at room temperature) with horseradish peroxidase-conjugated specific (Goat anti-rabbit (ab6721; 1:5000, Abcam, UK)

and goat anti-mouse (ab205719; 1:5000, Abcam, UK) secondary antibodies were used. Clarity™ Western ECL Substrate (Bio-Rad Laboratories Inc, Segrate (MI), Italy) was used for protein bands visualization. ChemiDoc™ XRS+ system (Bio-Rad Laboratories Inc, Segrate (MI), Italy) was used to detect chemiluminescence and ImageJ software (version 1.52a; National Institutes of Health, USA) was utilized to quantify the optical densities of the bands that were then normalized versus bands of β -actin.

2.3.11 Neurochemical quantifications

The neurochemical quantifications were carried out in PFC and HIPP of BTBR and BL6 animals. In particular DA, NA, 3,4-dihydroxyphenylacetic acid (DOPAC), homovanillic acid (HVA) and Kynurenine (KYN) were measured by using high-performance liquid chromatography coupled with an electrochemical detector (Ultimate ECD, Dionex Scientific, Milan, Italy). LC18 reverse phase column (Kinetex, 150mm x 3.0 mm, ODS 5 μ m; Phenomenex, Castel Maggiore-Bologna, Italy) was utilized to separate the catecholamines that were detected by a thin-layer amperometric cell (Dionex, ThermoScientific, Milan, Italy) with a 5-mm diameter glassy carbon electrode by using 400 or 750 mV as working potential vs. Pd. An aqueous buffer (pH 3.0) composed of 75 mM NaH₂PO₄, 1.7 mM octane sulfonic acid, 0.3 mM EDTA, acetonitrile 10%, was used as mobile phase and an isocratic pump (Shimadzu LC-10 AD, Kyoto, Japan) worked at 0.7 ml·min⁻¹ as flow rate. To perform the data acquisition and integration, Chromeleon software (version 6.80, Dionex, Thermo Scientific, San Donato Milanese, Italy) was utilized. Furthermore, GABA and Glutamate amounts were quantified by high-performance liquid chromatography with fluorescence detection, after derivatization with ophthalaldehyde/mercaptopropionic acid (emission length, 460 nm; excitation length, 340 nm) by using an OD column (Kinetex, 150mm x 3.0 mm, ODS 5 μ m; Phenomenex, Castel Maggiore-Bologna, Italy). The mobile phase used was a gradient phase of 50 mM sodium acetate buffer, pH 6.95, with methanol increasing linearly from 2 to 30% (v/v) over 40 minutes. A pump (JASCO, Tokyo, Japan) maintained the flow rate at 0.5 ml/min and the Borwin

software (version 1.50; Jasco) was used to analyze the results. Results, after being normalized for total area weight, were expressed as concentration/mg of tissue.

2.3.12 Acetylcholine (ACh) assay

The levels of ACh were quantified in tissue homogenized by using the commercially available kit (Catalog Number MAK435, Sigma-Aldrich, Milano, Italy), following the manufacturer's instructions. Tissue has been prepared by homogenization in cold 1× PBS and centrifugation (5 minutes at 14,000 × g). In the assay utilized, acetylcholine is hydrolyzed by acetylcholinesterase to choline which is oxidized by choline oxidase to betaine and H₂O₂. H₂O₂ reacts with a specific dye to form a colored product. The color intensity was analysed by the Multiskan™ FC Microplate spectrophotometer (Thermo Fisher Scientific, USA) at 570 nm and it is directly proportional to the acetylcholine concentration in the sample. Results were expressed in accordance to the linear detection range for the acetylcholine assay method (10-200 µM). Each sample analysis was performed in duplicate to avoid intra-assay variations.

2.3.13 Enzyme-Linked Immunosorbent Assays (ELISA)

Plasma samples were collected after centrifuge to 10000g at 4 °C of fresh blood samples and were stored at -80 °C until the analysis were performed. Commercially available ELISA kits were used for measurement of Oxytocin (EIAM-OXT-1, RayBiotech, Peachtree Corners, GA, USA), Prolactin (ELM-Prolactin-1, RayBiotech, Peachtree Corners, GA, USA) and Vasopressin (EIAM-VAS-1, RayBiotech, Peachtree Corners, GA, USA) levels, according to the manufacturer's instructions. In order to avoid intra-assay variations, all samples and standards were analyzed in duplicate.

2.3.14 Blindness of the study

For each test, scoring process and analysis, the experimenters were blind with respect to the experimental groups.

2.3.15 Statistical analyses

Sample size calculation has been performed a priori by using G power software. Statistical analyses were carried out by GraphPad Prism software (version 9.5.0; San Diego, CA, USA). In particular, Shapiro-Wilk test for normality and ROUT method to identify statistical outliers were performed for each group. Subsequently, Unpaired Student's t-test two-sided, with Welch's correction when needed, was used to analyze data that were expressed as mean \pm standard error of the mean (SEM). When the group were more than two, was performed One-way ANOVA followed by Tukey's multiple comparisons test.

2.4 Results

2.4.1 BTBR mice showed increased repetitive behaviour and novelty-induced hyperlocomotion.

To assess repetitive behaviours, the Hole Board test was performed on BTBR and BL6 mice. The results obtained showed that BTBR mice reported a significant increase in the number of poking holes respect to control mice (Fig. 1A). Moreover, the locomotor activity in a novel environment was explored in the Open Field paradigm. We observed an increase of the distance travelled in the arena (Fig. 1B), in BTBR mice compared with control animals. Additionally, BTBR mice also showed a reduction in freezing duration (Fig. 1D) and freezing frequency (Fig. 1C) compared with BL6 animals.

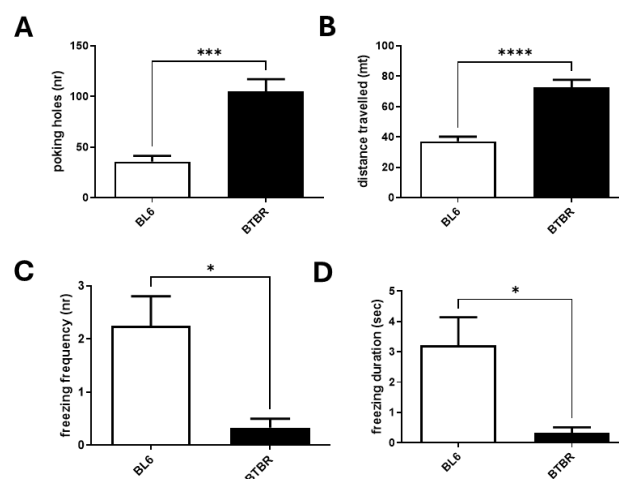


Fig. 1. Repetitive and stereotyped behaviours of BTBR mice. **A)** Number of poking holes in Hole Board task, Unpaired Student's t-test, $P < 0.001$ BTBR vs. BL6. **B)** Distance travelled in the Open Field paradigm, Unpaired Student's t-test, $P < 0.0001$ BTBR vs. BL6. **C)** Freezing duration in Open Field paradigm, Unpaired Student's t-test, $P < 0.05$ BL6 vs. BTBR **D)** Freezing frequency in Open Field paradigm, Unpaired Student's t-test, $P < 0.05$ BL6 vs. BTBR.

2.4.2 BTBR mice showed reduced social behaviours and increased non-social behaviours

The Social Interaction test was performed to explore social dysfunctions. During the test, BTBR mice demonstrated significantly less duration of social interaction (sec) compared to BL6 mice (Fig. 2A). Furthermore, the time employed (sec) performing non-social interactions in BTBR animals was increased against controls (Fig. 2B). Observing the social frequency time, measurements of BTBR mice were significantly decreased (Fig. 2C), while there was a significant improvement in non-social frequency vis-a-vis control (Fig. 2D).

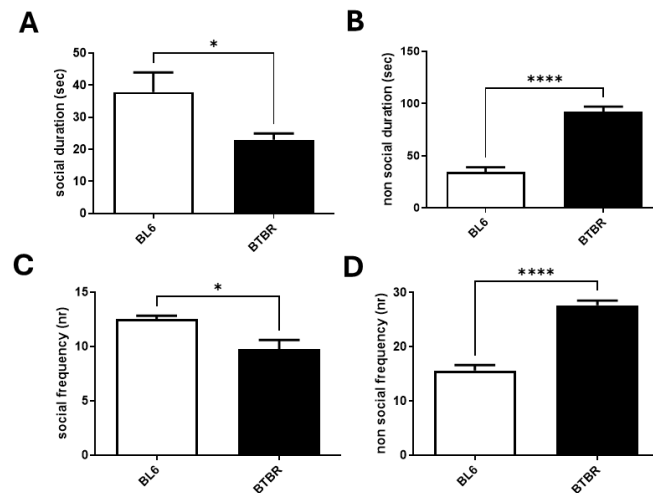


Fig. 2. Social-related behaviours of BTBR mice. **A)** Time spent (sec) in Social Interaction test, Unpaired Student's t-test, $P < 0.05$ BTBR vs. BL6. **B)** Time spent (sec) in non-social interactions, Unpaired Student's t-test, $P < 0.0001$ BTBR vs. BL6. **C)** Social frequency measurements in social interaction Test, Unpaired Student's t-test, $P < 0.05$ BL6 vs. BTBR. **D)** Non-social frequency, Unpaired Student's t-test, $P < 0.0001$ BTBR vs. BL6.

2.4.3 BTBR mice did not show anxiety-like behaviours

To evaluate the anxiety-like behaviours, were performed the Elevated Zero Maze and the Open Field tests. In contrast to control animals, BTBR animals showed an increase in the time spent exploring the open arms compared to controls (Fig. 3A). In addition, results from Open field test showed that the time spent in the center in BTBR animals was significantly augmented respect to control animals (Fig. 3B), whereas the time spent in the wall was a significantly reduced (Fig. 3C), respectively.

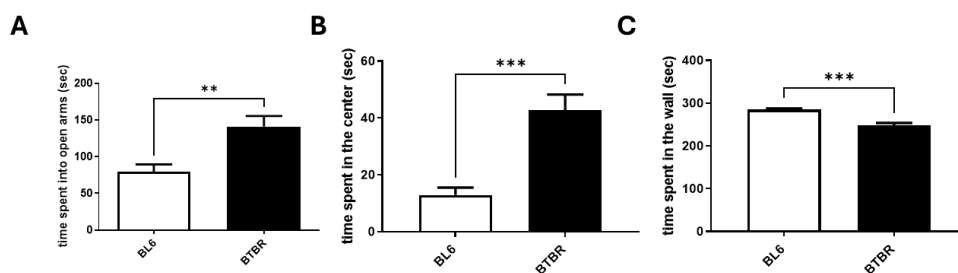


Fig. 3. Anxiety-related behaviours of BTBR mice. A) Time spent (sec) exploring the open corridors in Elevated Zero Maze, Unpaired Student's t-test, $P < 0.01$ BTBR vs. BL6. **B)** Time spent (sec) in the center in Open field test, Unpaired Student's t-test, $P < 0.001$ BTBR vs. BL6. **C)** Time spent (sec) in the wall in Open Field test, Unpaired Student's t-test, $P < 0.001$ BTBR vs. BL6).

2.4.4 BTBR mice showed reduced NA, ACh and GABA levels, and increased Kyn levels in PFC

The results report a significantly reduction of NA levels in the PFC of BTBR animals related to BL6 (Fig. 4A), similarly also ACh (Fig. 4E) and GABA (Fig. 5B) levels was reduced in the BTBR mice. Concerning Kyn, its levels were increased in BTBR mice respect to the control animals (Fig. 5C), whereas levels of DA (Fig. 4B), DOPAC (Fig. 4C), HVA (Fig. 4D), NGF (Fig. 6A), BDNF (Fig. 6B) and glutamate (Fig. 5A), were not significantly different between the two groups.

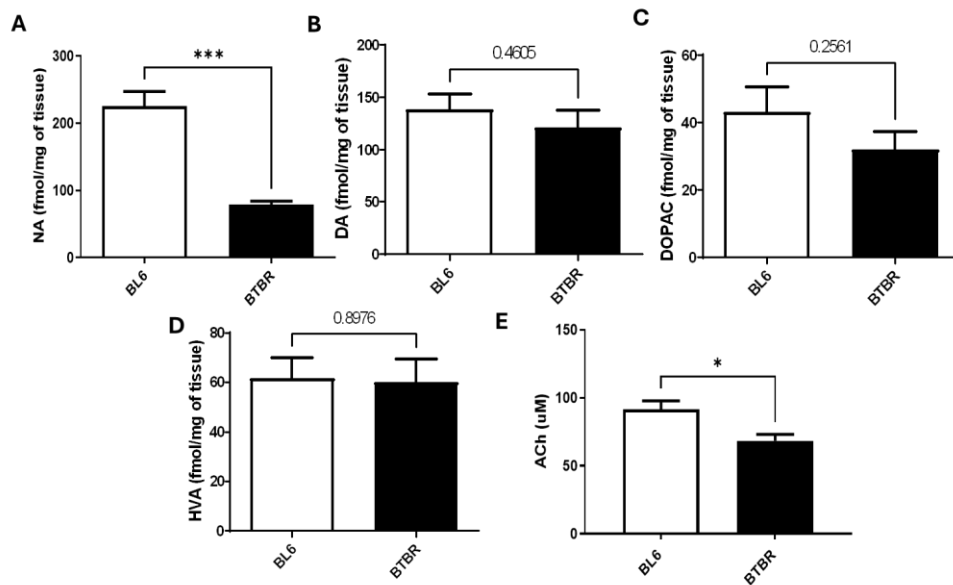


Fig. 4. Catecholamines and Acetylcholine levels in PFC of BTBR and BL6 mice. A) NA levels, Unpaired Student's t-test, $P < 0.0001$ BTBR vs. BL6. **B)** DA levels, Unpaired Student's t-test, n.s. **C)** DOPAC levels, Unpaired Student's t-test, n.s. **D)** HVA levels, Unpaired Student's t-test, n.s. **E)** Ach levels, Unpaired Student's t-test, $P < 0.05$ BTBR vs. BL6

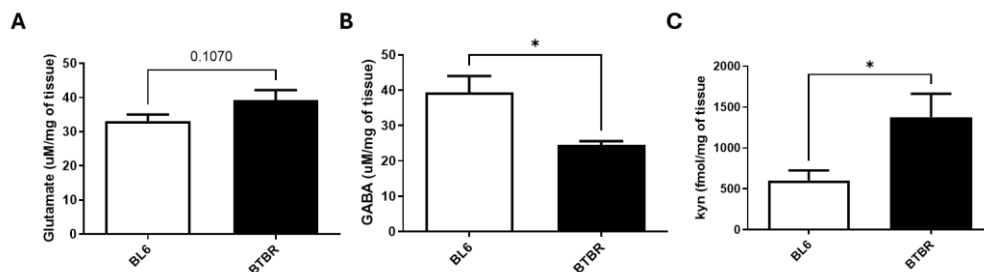


Fig. 5. Glutamate, GABA and Kyn levels in PFC of BTBR and BL6 mice. A) Glutamate levels, Unpaired Student's t-test, n.s. **B)** GABA levels, Unpaired Student's t-test, $P < 0.05$ BTBR vs. BL6. **C)** Kyn levels, Unpaired Student's t-test, $P < 0.05$ BTBR vs. BL6.

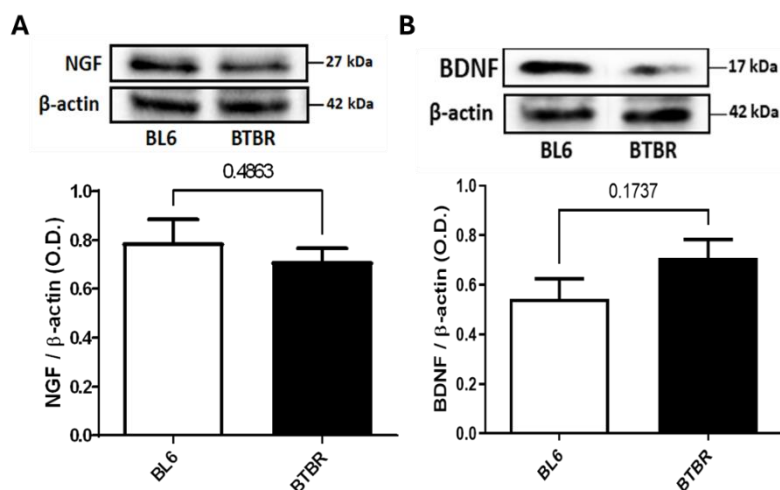


Fig. 6. NGF and BDNF expression levels in PFC of BTBR and BL6 mice. A) NGF representative bands and expression levels, Unpaired Student's t-test, n.s. **B)** BDNF representative bands and expression levels, Unpaired Student's t-test, n.s.

2.4.5 BTBR mice showed enhanced DOPAC and Kyn levels, and decreased NGF, BDNF and GABA levels in HIPP

In HIPP, the results did not show any difference in NA (Fig. 7A), DA (Fig. 7B), HVA (Fig. 7D), Ach (Fig. 7E) and glutamate (Fig. 8A) levels between BTBR and the control groups. Furthermore, analysis of DOPAC levels reported a significant increase in BTBR animals against controls (Fig. 7C). Whereas GABA levels were decreased (Fig. 8B). Regarding neurotrophins, both NGF and BDNF expression levels were reduced (Fig 9 A-B). Lately, Kyn levels showed an increase in BTBR mice respect to the control mice (Fig. 8C).

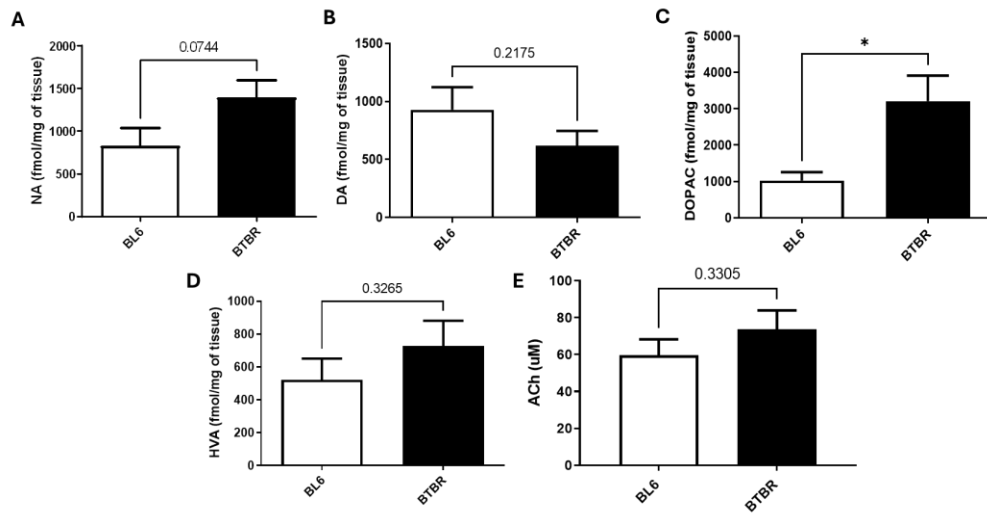


Fig. 7. Catecholamines and Acetylcholine levels in HIPP of BTBR and BL6 mice. A) NA levels, Unpaired Student's t-test, n.s. **B)** DA levels, Unpaired Student's t-test, n.s. **C)** DOPAC levels, Unpaired Student's t-test, $P < 0.05$ BTBR vs. BL6. **D)** HVA levels, Unpaired Student's t-test, n.s. **E)** Ach levels, Unpaired Student's t-test, n.s.

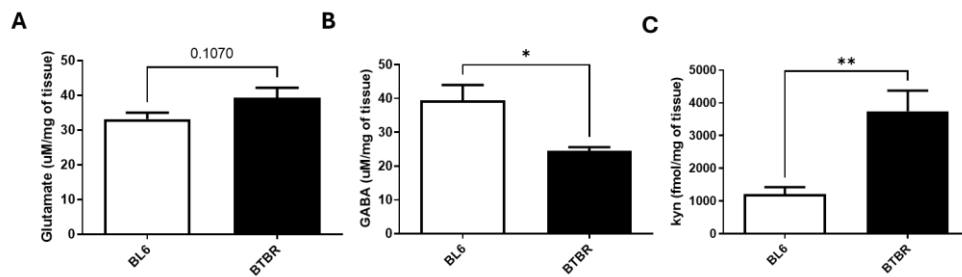


Fig. 8. Glutamate, GABA and Kyn levels in HIPP of BTBR and BL6 mice. A) Glutamate levels, Unpaired Student's t-test, n.s. **B)** GABA levels, Unpaired Student's t-test, $P < 0.05$ BTBR vs. BL6. **C)** Kyn levels, Unpaired Student's t-test, $P < 0.01$ BTBR vs. BL6.

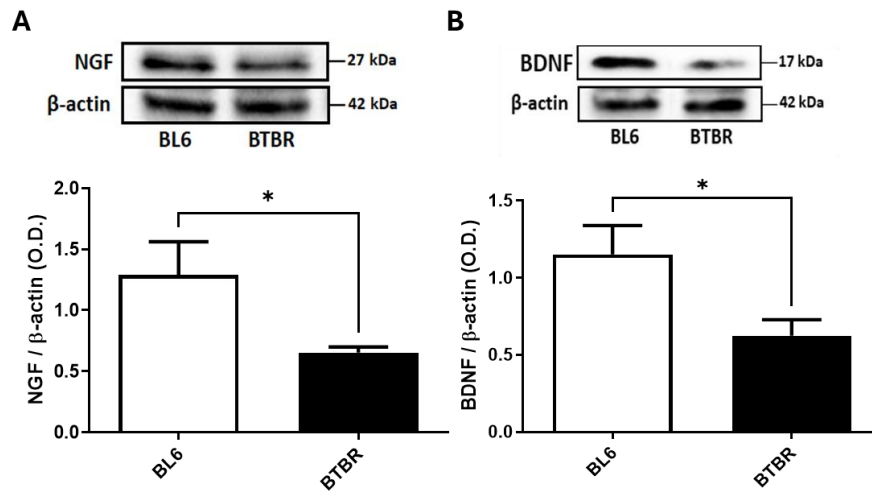


Fig. 9. NGF and BDNF expression levels in HIPP of BTBR and BL6 mice. **A)** NGF representative bands and expression levels, Unpaired Student's t-test, $P < 0.05$ BTBR vs. BL6. **B)** BDNF representative bands and expression levels, Unpaired Student's t-test, $P < 0.05$ BTBR vs. BL6.

BTBR mice showed decreased Oxytocin plasma levels.

Oxytocin levels were measured in plasma using an ELISA kit. The results demonstrated that control indicators of Oxytocin plasma levels in the BTTR group exceeded those of the control group.

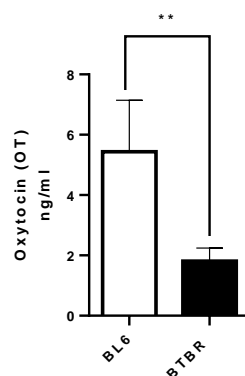


Fig. 10. Oxytocin levels in Plasma of BTBR and BL6 mice. Oxytocin plasma levels, Unpaired Student's t-test, $P < 0.01$ BTBR vs. BL6.

2.4.6 β -Carotene treatment was able to reduce repetitive and stereotyped behaviours of BTBR mice

In order to evaluate the effect of β -Carotene administration on repetitive and stereotyped behaviours, the Marble Burying test, the Hole Board test and the Open Field test were performed in the offspring of treated and vehicle dams. The results obtained showed that the administration of β -carotene in pregnant mice at a dose of 16 mg/kg was able to significantly reduce the duration (Fig. 11A) and frequency of grooming (Fig. 11C), the number of buried marbles (Fig. 11B) and the number of poking holes (Fig. 11D) in male offspring compared to offspring of pregnant mice treated with vehicle (seed oil) alone. Regarding the other dosages, the administration of β -carotene at a dose of 32 mg/kg led to a significant reduction in the frequency of grooming (Fig. Fig. 11C) and the 8 mg/kg dose significantly reduced the number of poking holes (Fig. 11D). Therefore, supplementation with β -carotene at a dose of 16 mg/kg was the most effective in preventing repetitive behaviours in BTBR mice. In addition, in the Open Field test, the total distance travelled, an index of locomotor activity, was measured, the results showed that in male offspring treated with β -carotene at a dose of 16 mg/kg was able to significantly reduce the distance traveled (Fig. 11E) compared to controls and doses of 32 mg/kg and 64 mg/kg.

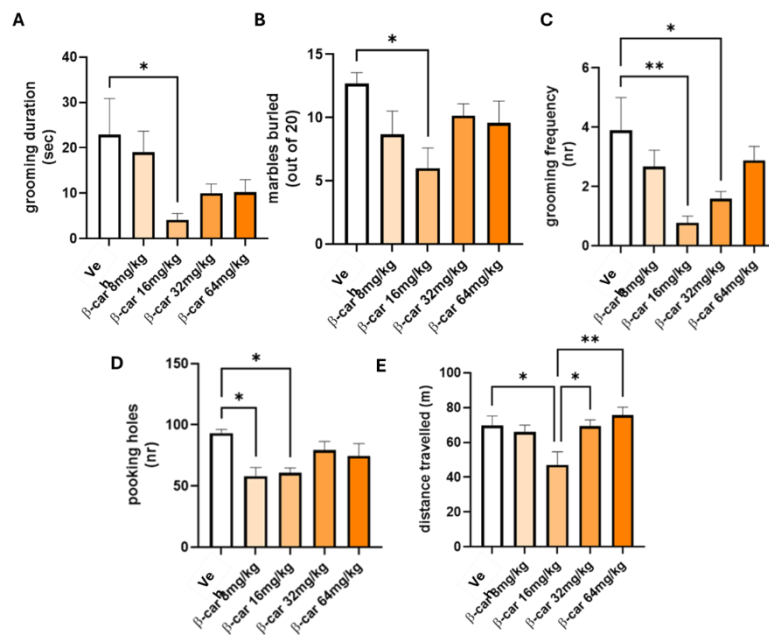


Fig. 11. Repetitive and stereotyped behaviours in BTBR dams treated with vehicle or β -carotene. **A)** Grooming duration, One-way ANOVA followed by Tukey's multiple comparisons test, $P < 0.05$ β -car 16 mg/kg vs VEH. **B)** Number of marbles buried, One-way ANOVA followed by Tukey's multiple comparisons test, $P < 0.05$ β -car 16 mg/kg vs VEH. **C)** Grooming Frequency, One-way ANOVA followed by Tukey's multiple comparisons test, $P < 0.01$ β -car 16 mg/kg vs VEH and One-way ANOVA followed by Tukey's multiple comparisons test, $P < 0.05$ β -car 32 mg/kg vs VEH. **D)** Number of Pooking holes in hole board test, One-way ANOVA followed by Tukey's multiple comparisons test, $P < 0.05$ β -car 8 mg/kg vs VEH and One-way ANOVA followed by Tukey's multiple comparisons test, $P < 0.05$ β -car 16 mg/kg vs VEH. **E)** Distance travelled in Open Field paradigm, One-way ANOVA followed by Tukey's multiple comparisons test, $P < 0.05$ β -car 16 mg/kg vs VEH.

2.4.7 β -Carotene Treatment enhanced social related behaviours of BTBR mice

To evaluate the social related behaviours the social interaction test was performed. The results obtained showed that supplementation with β -carotene at a dose of 16 mg/kg was able to increase the frequency of social interactions in male offspring compared to control mice and the other three doses (Fig. 12B). Regarding the duration of the non-social interactions with the three objects, β -carotene doses of 32 mg/kg and 64 mg/kg administered to pregnant mice were able to decrease this parameter in male offspring compared to control progeny (Fig. 12C).

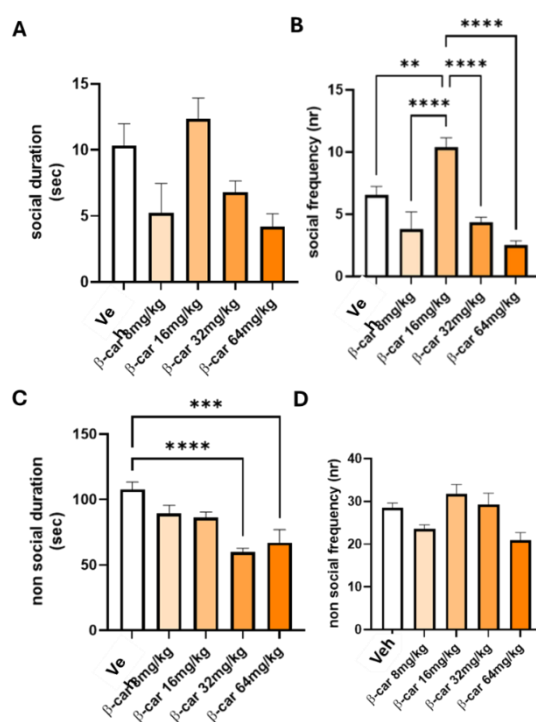


Fig. 12. BTBR dams treated with vehicle or β -carotene in the social interaction test. A) Time spent (sec) in social interactions, One-way ANOVA followed by Tukey's multiple comparisons test β -car 16 mg/kg vs VEH n.s. B) Frequency of social interactions, One-way ANOVA followed by Tukey's multiple comparisons test, **** $P < 0.0001$ β -car 16 mg/kg vs β -car 8 mg/kg, β -car 32 mg/kg and β -car 64 mg/kg; ** $P < 0.01$ β -car 16 mg/kg vs VEH. C) Time spent (sec) on non social interactions, One-way ANOVA followed by Tukey's multiple comparisons test, **** $P < 0.0001$ β -car 32 mg/kg vs CTRL; *** $P < 0.001$ β -car 64 mg/kg vs VEH. D) Frequency of non social interactions, One-way ANOVA followed by Tukey's multiple comparisons test β -car 16 mg/kg vs VEH n.s.

2.4.8 β -Carotene Treatment increased plasmatic Oxytocin levels BTBR mice

Based on behavioural results we choose the dose of 16 mg/kg to perform *ex-vivo* analysis. In particular we performed analysis of oxytocin, prolactin and Vasopressin on plasma samples. Among the factors examined the B-Carotene treated dams did not show differences in Prolactin and Vasopressin levels compared to the Vehicle treated dams (Fig. 13 B-C). At the same time, treatment with β -carotene produced an increase in oxytocin levels in the male offspring from β -carotene treated dams compared to the offspring from vehicle treated dams (Fig. 13A).

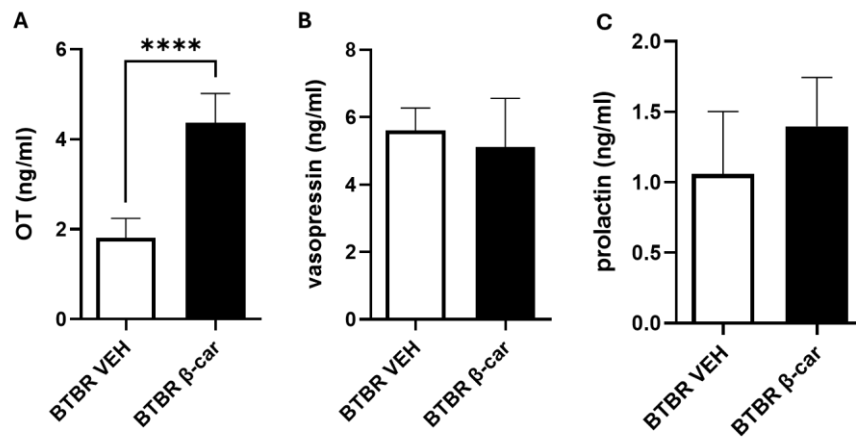


Fig. 13. Effects of β -carotene treatment on Oxytocin, Vasopressin and Prolactin levels of BTBR mice. A) Oxytocin plasmatic levels, Unpaired Student's t-test, $P < 0.0001$ β -car 16 mg/kg vs VEH. B) Vasopressin plasmatic levels, Unpaired Student's t-test n.s. C) Prolactin plasmatic levels, Unpaired Student's t-test n.s.

2.4.9 β -Carotene treatment decreased the Kynurenine levels in PFC of BTBR mice

To evaluate the effect of β -Carotene treatment on some markers related to BBB function and neuroinflammation in PFC, NF- κ B, BDNF, Kynurenine, Occludin, GFAP, CD11-b and IBA-1 were quantified. The results obtained did not show any substantial differences in the expression levels of NF- κ B (Fig. 14A) and BDNF (Fig. 14B) between offspring from β -Carotene treated dams and offspring of vehicle treated dams. Regarding instead the kynurenine in PFC the offspring of the β -carotene treated dams showed a significant decrease in the kynurenine levels compared to offspring of vehicle treated dams (Fig. 14C). Furthermore, the results obtained did not show any substantial differences in the expression levels of Occludin (Fig. 15A), GFAP (Fig. 15B), CD11-b (Fig. 15C) and IBA-1 (Fig. 15D) in the offspring of β -carotene from treated dams compared to the offspring from vehicle treated dams.

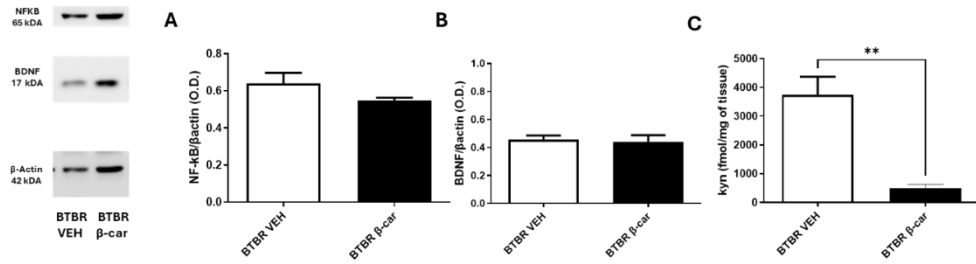


Fig. 14. Effects of β -carotene treatment on NF-kb and BDNF expression levels and on Kyn Levels in PFC of BTBR mice. A) NF-kb expression levels, Unpaired Student's t-test, n.s. B) BDNF expression levels, Unpaired Student's t-test, n.s. C) Kyn levels, Unpaired Student's t-test, $P < 0.01$ β -car 16 mg/kg vs VEH

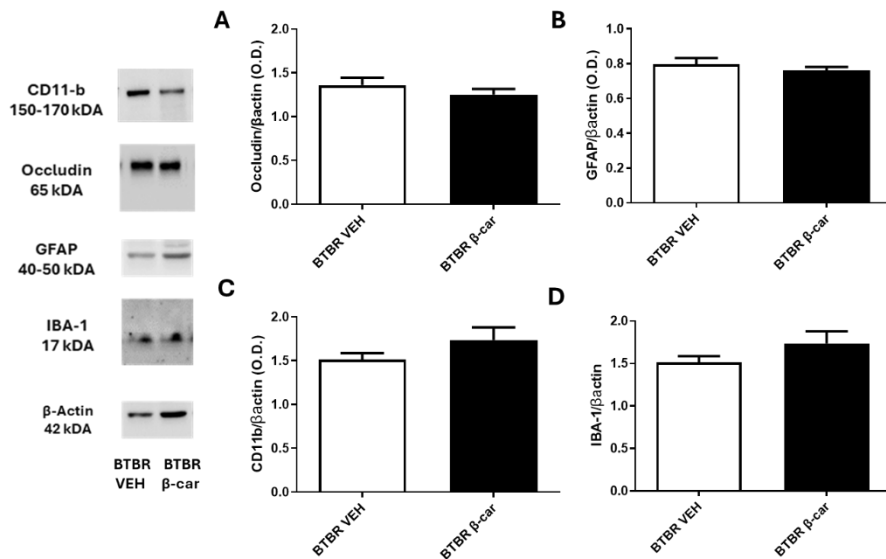


Fig. 15. Effects of β -carotene treatment on Occludin, GFAP, CD11-b and IBA-1 expression levels in PFC of BTBR mice. A) Occludin expression levels, Unpaired Student's t-test, n.s. B) GFAP expression levels, Unpaired Student's t-test, n.s. C) CD11-b expression levels, Unpaired Student's t-test, n.s. D) IBA-1 expression levels, Unpaired Student's t-test, n.s.

2.4.10 β -Carotene Treatment reduced Kynurenine and NF-kB levels and increased the BDNF, Occludin and IBA-1 levels in HIPP of BTBR mice

To evaluate the effect of β -Carotene treatment on some markers related to BBB function and neuroinflammation in HIPP, NF-kB, BDNF, Kynurenine, Occludin, GFAP, CD11- β and IBA-1 were quantified. The treatment effect showed a reduction in the expression levels of NF-kB (Fig. 16A) and an increase in the expression levels BDNF (Fig. 16B) in HIPP for the offspring from β -Carotene 16/mg treated dams compared to the offspring of the vehicle treated dams.

Moreover, the content of kynurenine in HIPP was significantly reduced in the offspring from β -Carotene Treated dams compared to the offspring from controls (Fig. 16C). Lastly the offspring from dams treated with B Carotene 16/mg showed increased levels in the expression of Occludin (Fig. 17A) and IBA-1 (Fig. 17D), while did not show any substantial differences in the expression levels of CD11- β and GFAP (Fig. 17 B-C) compared to offspring from controls.

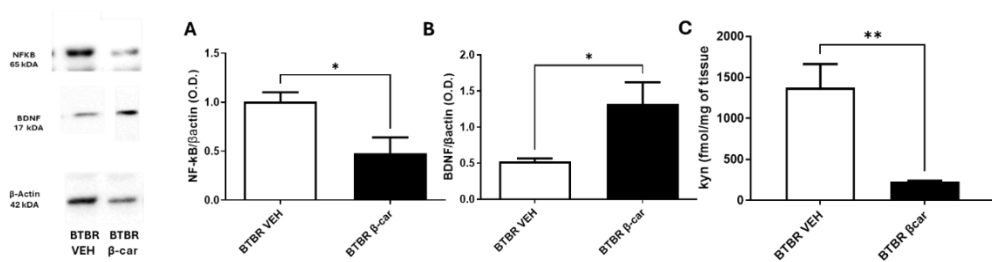


Fig. 16. Effects of β -carotene treatment on NF-kb and BDNF expression levels and on Kyn Levels in HIPP of BTBR mice. **A)** NF-kb expression levels, Unpaired Student's t-test, $P < 0.05$ β -car 16 mg/kg vs VEH. **B)** BDNF expression levels, Unpaired Student's t-test, $P < 0.05$ β -car 16 mg/kg vs VEH. **C)** Kyn levels, Unpaired Student's t-test, $P < 0.01$ β -car 16 mg/kg vs VEH.

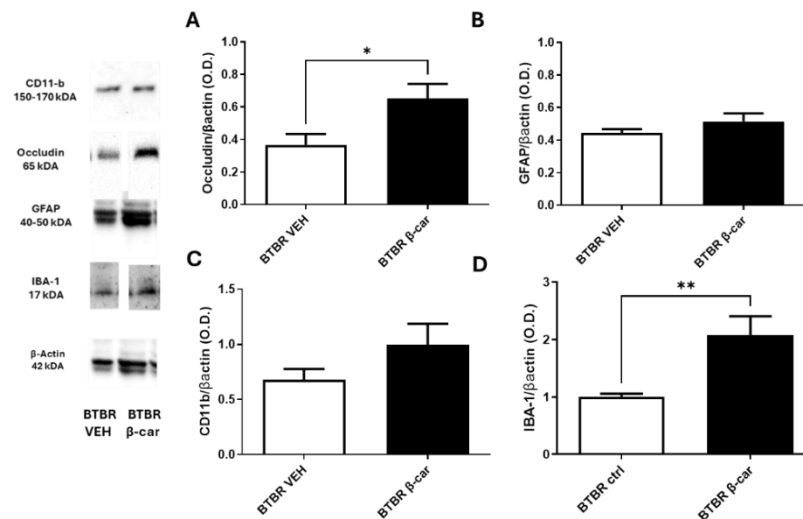


Fig. 17. Effects of β -carotene treatment on Occludin, GFAP, CD11- β and IBA-1 expression levels in HIPP of BTBR mice. **A)** Occludin expression levels, Unpaired Student's t-test, $P < 0.05$ β -car 16 mg/kg vs VEH. **B)** GFAP expression levels, Unpaired Student's t-test, n.s. **C)** CD11- β expression levels, Unpaired Student's t-test, n.s. **D)** IBA-1 expression levels, Unpaired Student's t-test, $P < 0.05$ β -car 16 mg/kg vs VEH.

2.5 Discussion

Although ASD are exclusively human diseases, mice models able to resemble the wide range of ASD behavioural phenotypes represent a useful instrument to understand the underlying neurobiological alterations (Bove et al., 2022).

The results obtained in this part of the thesis confirmed that the BTBR mice model of ASD reported stereotyped repertoire and social interaction deficits, by performing the Hole Board task and a modified version of the Social Interaction test, in which we evaluated social duration and frequency, together with frequency and time spent performing exploration of non-socially related objects. Interestingly, despite the well-identified reduction of social interactions in BTBR mice (Blanchard et al., 2012; Bove et al., 2022; Karvat & Kimchi, 2014; Nadeem et al., 2019), results also showed an increase of the time spent performing non-social exploration, together with frequency of this behaviour. Regarding this, the exploration of three objects non socially related might have similarities with the absence of interests in people and in developing relationships, an important behavioural feature of ASD patients (Davidson et al., 2015). Furthermore, results showed an increase in the distance travelled and a reduction in the duration and frequency of freezing in the Open Field test, both considered as indexes of hyperactivity, an important ASD collateral symptom (Aman & Langworthy, 2000). Behavioural results are in agreement with the previous literature on this animal model, in particular regarding repetitive behaviours (Brierley et al., 2021; McFarlane et al., 2008; Moy et al., 2007) and higher initial activity in the open field test (Silverman, Tolu, et al., 2010). Regarding the anxiety-related behaviours, BTBR mice showed a reduction of these behaviours in the Elevated Plus Maze and in the Open Field tests, as already reported in literature (Pobbe et al., 2011). Interestingly, spending more time in the center and in the open areas might be correlated to an increase in risk taking behaviours. In addition, the reduction in the anxiety-related behaviours could also be interpreted as an enhanced hyperactivity and impulsiveness, important traits typical of ASD patients (Rodrigues et al., 2021).

From a molecular point of view, different neuroanatomical abnormalities shared across BTBR mice and ASD subpopulations have been described (Chao, Pathak, et al., 2020; Ellegood et al., 2013). This study focused on two different brain areas (PFC and HIPP) that, together with AMY, are centrally involved in ASD and that are strictly interplaying among each other. In this study dopaminergic, noradrenergic, and excitatory-inhibitory neurotransmissions were investigated, together with kynurenine, BDNF and NGF levels, in both PFC and HIPP. Concerning DA, results did not show differences in PFC and HIPP, but there was an increase in the levels of its metabolite DOPAC in PFC. In this regard, several studies reported a dopaminergic modulation in widespread forebrain areas of ASD patients (Chadman et al., 2012; Garnier et al., 1986; Ghanizadeh & Moghimi-Sarani, 2013; Launay et al., 1987; Martineau et al., 1992; Nguyen et al., 2014; Squillace et al., 2014). Moreover, we found a decrease in cortical NA levels in BTBR mice, with no differences in the hippocampal area. In this regard, it has been reported that stereotyped behaviours, considered as derivatives of poor adaptive behaviours, are principally mediated by the noradrenergic system and that the administration of an alpha-2-receptor agonist might be helpful to improve ASD behavioural dysfunctions (Nanjappa et al., 2022). In line with our results, another model of ASD, the Engrailed-2 knock-out (EN2KO) mice, also reported NA neurotransmission alterations in the ventral hindbrain (Brielmaier et al., 2014). In addition, it has been shown that the administration of atomoxetine, a NA reuptake inhibitor, could be effective for treating hyperactivity in children with ASD (Arnold et al., 2006). In regards to ACh content, results showed a decrease in PFC according to neurochemical alterations in the cholinergic pathway observed in a postmortem study involving ASD patients (Brielmaier et al., 2014). Moreover, it has been reported that ACh increase could conduce to an improvement of cognitive deficits in ASD and other neuropsychiatric disorders (L. Wang et al., 2015). Interestingly, reduced ACh in the basal forebrain has been linked to decreased social interactions and social memory dysfunctions in mice (Kljakic et al., 2021).

In our results, we also found impairments of excitatory-inhibitory neurotransmissions, in particular, regarding PFC and HIPP, we only reported a decrease in GABA levels, while Glu levels remained unchanged. Therefore, BTBR animals demonstrated a perturbation of the excitatory-inhibitory balance, by showing disruptions in both PFC and HIPP. As regarding GABA, the reduction in its levels in HIPP has been associated with decreased sociability, especially considering that HIPP is a crucial component of the social brain (Paine et al., 2017). Regarding kynurenine levels, our results showed an increase both in PFC and HIPP, thus confirming the neuroinflammatory component of ASD (DiSabato et al., 2016). Accordingly, a tryptophan-kynurenine pathway dysregulation was reported in inflammation and immune activation contest (Q. Wang et al., 2015). Furthermore, our results showed a decrease in the expression levels of the neurotrophins NGF and BDNF in HIPP, a brain region in which such neurotrophins are mainly present (Dinçel et al., 2013). To this purpose, different studies reported an interplay between NA neurotransmission and NGF expression (Benitez et al., 2021; Counts & Mufson, 2010; Naoi & Maruyama, 2010). In particular, it has been shown that, in cellular models, monoamine oxidase inhibitors increased NGF expression (Naoi & Maruyama, 2010) and that NA can exert neuroprotective properties by inducing NGF expression (Counts & Mufson, 2010). In line with our results, hippocampal BDNF and NGF deficiency has been associated with ASD development (Dingsdale et al., 2022; C. Liu et al., 2023), however results are contrasting, since BDNF brain levels fluctuations have been reported in response to several known, such as brain areas and age, and unknown factors (Kasarpalkar et al., 2014).

Furthermore, our results showed a decrease in the oxytocin plasmatic levels of BTBR mice. Accordingly, it has been demonstrated that an impairment of OT signaling could be associated to social behaviour impairments (Modi & Young, 2012). Concerning the treatment used in this study, β carotene subchronic administration (at the dose of 16 mg/kg, 2 times/week for the whole pregnancy) was able to increase OT plasmatic levels but did not affect the levels of

vasopressin and prolactin in the offspring of BTBR mice after β -carotene or vehicle treatment. In line with this, it has been demonstrated that in patients with ASD there were reduced levels of oxytocin and its CD38 regulatory ectoenzyme. Interestingly, β -carotene is a potent inducer of the CD38 enzyme (Avraham et al., 2019). Considering the role of OT in the social behaviours, results obtained showed that supplementation with β -carotene during pregnancy was able to increase the frequency of social interactions in male BTBR offspring compared to vehicles. In particular, among the four doses tested (8, 16, 32 and 64 mg/Kg), the dose of 16/kg appeared to be the most effective. Moreover, such dose was also able to reduce repetitive and stereotyped behaviours of BTBR mice, in particular regarding grooming, numbers of marbles buried, numbers of head dipping in hole board task and distance travelled in open field test. In line with our results, pups treated with β -carotene injection reported a reduction of repetitive and stereotyped behaviours (Avraham et al., 2019). Behavioural results were corroborated by *ex vivo* analyses. Indeed, cortical and hippocampal neurochemical and biomolecular measurements between the BTBR vehicle and the BTBR treated with 16 mg/kg of β -carotene group were performed. By focusing on the possible role of β -carotene to counteract neuroinflammatory components of ASD, we quantified kynurenine, NF-Kb and BDNF in HIPP and PFC. Indeed, beyond its effects on the sociability and on the stereotyped repertoire, β carotene owns also anti-inflammatory and anti-oxidant effects (Liebler et al., 1997; Y. Yang et al., 2021; Y. Zhang et al., 2016). In this regard, our results showed that β carotene was able to reduce kynurenine levels in both HIPP and PFC and the NF-kB expression levels in HIPP, but not in PFC, thus supporting its role in contrasting neuroinflammation. In line with our results, different studies reported that β -carotene strongly inhibited the NF-kB pathway activation in a rat model of spinal cord injury (L. Zhou et al., 2018) and that the β -carotene treatment reduced NF-kB expression in a model of rat ischemic brain injury (Althurwi et al., 2022). Furthermore, the results obtained showed that β -carotene administration was able to enhance the expression levels of BDNF in

HIPP. Accordingly, recent studies on BDNF and carotenoid docking may indicate the possibility of an allosteric activation of BDNF by carotenoids (S.-J. Park et al., 2021).

Concerning BBB and glia-related markers, our results showed different interesting effects of β -carotene administration in HIPP. In particular, β -carotene treatment increased the expression levels of Occludin and IBA-1, while the expression levels of GFAP and CD11-b did not show any difference between β -carotene treated and vehicle-treated BTBR in HIPP. As regarding PFC, no difference was detected between the treated BTBR group compared to the vehicle one. The hippocampal effects could be due to the increase in OT levels induced by β carotene treatment. Indeed, OT can modulate the inflammatory response of microglia cells (Knoop et al., 2022) maybe acting through NF- κ B/MAPK pathway (Ye et al., 2022). Although it is not clear how OT influences the microglial phenotype, the increase we observed after β carotene treatment in microglial cell expression, via the IBA-1 marker, could be linked to the increase in OT induced by β -carotene administration. Moreover, enhanced the Occludin expression levels could be linked to a BBB strengthening following the less inflammatory conditions induced by β -carotene treatment.

In conclusion, the molecular characterization of the BTBR strain suggested that such idiopathic animal model might be useful to understand neurobiological and neurochemical correlates underlying ASD behavioural dysfunctions, highlighting the important role of neurotransmitter and neurotrophins alterations in a specific brain region, such as HIPP and PFC. Furthermore, our results showed also the promising preventive effect of β -carotene treatment during pregnancy to ameliorate ASD behavioural dysfunctions in offspring and also to counteract neuroinflammatory components of these neurodevelopmental pathologies.

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