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The effects of neuroinflammation induced by systemic lipopolysaccharides on the hippocampi of aged Sprague-Dawley rats

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Submitted in fulfilment of the requirements for the degree

Master of Science (MSc Human Physiology)

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Abstract

Dementia affects a significant number of South Africans. In 2015, an estimated 186, 000 South Africans were recorded as being patients of this irreversible condition. Alzheimer's disease (AD) accounts for 60% to 80% of reported dementia cases. It is estimated that more than 46.8 million people are affected by AD, worldwide. According to the 2015 World Alzheimer's Report, an estimated 186,000 South Africans struggled with dementia in 2015, and this number is anticipated to increase to 275 000 by 2030. In the past decades, two hallmarks termed amyloidosis and taupathy have received major acknowledgement in neurodegenerative studies. Although this knowledge has tremendously contributed towards the understanding of the molecular mechanisms underlying AD, therapeutic treatments that reverse the disease are yet to be discovered. Therefore, exploring the aetiology of the disease using a popular rodentmodel of inflammation can help provide some missing links required for long-term therapeutic strategies. Previous research has validated the use of lipopolysaccharide (LPS) in rodent models to replicate characteristics of AD and examine the inflammatory pathways and molecules involved. Therefore, in this study, the hippocampal region of male Sprague-Dawley rats was examined for AD-like (cognitive and histological) pathologies in response to repeated exposure to LPS. Subjects were assorted into one control group and three experimental groups, and the results of the experimental group were compared against the control group. LPS sourced from Escherichia coli 055:B5 was administered through repeated subcutaneous (SC) injection to induce a chronic systemic inflammatory response. Cognitive assesssments were conducted using a series of three behavioural experiments commonly used by researchers. This included the Y-maze, novel object recognition (NOR), and open-field tests. To quantitatively determine the effects of LPS-induced neuroinflammation on hippocampal neuroglia-astrocytes and microglia, biochemical assays involving ELISA and confocal microscopy were performed. To identify astrocytes and microglia, anti-glial fibrillary protein (GFAP) and anti-ionized calciumbinding adaptor molecule 1 (Iba1) fluorescent markers were used to stain hippocampal sections and view by microscopy. Previous findings have conveyed the benefits of honey as an anti-inflammatory agent against infectious pathogens. Therefore, during the experimental period of this study, Manuka honey was introduced to the subjects by oral gavage. The effects of honey as a "mopping agent" were identified by comparison of the experimental groups against the control group. Therefore the aim of this study was to investigate the effects of neuroinflammation induced by systemic lipopolysaccharide from *Escherichia coli* 055:B5 on the hippocampi of Sprague-Dawley rats. Cognitive assessments revealed that LPS exposure, over a 10-day period, did not significantly impair short-term spatial working memory, learning capacity and spontaneous memory, and anxiety, and locomotor activity. Confocal microscopy showed that LPS significantly increased the quantity of microglia detected by Iba1 antibody. This suggests that LPS exposure induced neuroinflammation in the hippocampal region, however, the nature of the inflammatory response, physiological or pathological, was unclear.

Keywords: Alzheimer's disease (AD), lipopolysaccharide (LPS), inflammation, *Escherichia coli*, hippocampus, microglia, astrocytes, honey, novel object recognition (NOR) test, open-field test, y-maze test, subcutaneous

Declaration of Originality and Authorship

I, the undersigned, hereby declare that the thesis titled

"The effects of neuroinflammation induced by systemic lipopolysaccharides on the hippocampi of aged Sprague-Dawley rats"

Is my original work and has not ever been presented for academic award to this or any other tertiary institution for another degree. I have read and understand the University of Pretoria's policy on plagiarism.

Herewith, I submit this thesis to the University of Pretoria for the degree Master of Science in Human Physiology.

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List of Abbreviations

Αβ	Amyloid beta
AD	Alzheimer's disease
ANS	Autonomic nervous system
APP	Amyloid precursor protein
АроЕ	Apolipoprotein E
BBB	Blood brain barrier
BSA	Bovine serum albumin
CA	Cornu Ammonis
CAS	Central animal services
C1q	Complement component
	1q
CD	Cluster of differentiation
CNS	Central nervous system
CSF	Cebrospinal fluid
CSFs	Colony-stimulating factors
СТ	Computerized tomography
Cdk5	Cyclin-dependent protein
	kinase 5
COX-2	Cyclooxygenase-2
CVO	Circumventricular organs
DAMPs	Damage-associated
	molecular pattern
DG	Dentate gyrus
ELISA	Enzyme-linked
	immunosorbent assay
GALT	Gastrointestinal-associated
	lymphoid tissue
GFAP	Glial fibrillary acidic protein
GIT	Gastrointestinal tract
GSK3β	Glycogen synthase kinase
	3β

H&E	Hematoxylin and eosin
lba-1	lonized calcium-binding
	adapter molecule
IF	Immunofluorescence
IHC	Immunohistochemistry
IL	Interleukin
iNOS	Inducible nitric oxide
	synthase
IFN	Interferon
LBP	LPS-binding protein
LPS	Lipopolysaccharide
MAC	Membrane attack complex
MAPs	Microtubule-associated
	proteins
MTL	Medial temporal lobe
MRI	Magnetic resonance
	imaging
NFT	Neurofibrillary tangle
NO	Nitric oxide
NORT	Novel object recognition
	test
NVU	Neurovascular unit
PAMPs	Pathogen-assosciated
	molecular patterns
PBS	Phosphate buffered saline
PET	Positron emission
	tomography
PNS	Peripheral nervous system
PRR	Pattern recognition
	receptor
ROS	Reactive oxygen species
TBS	Tris buffered saline

TIR	Translocated intimin
	receptor
TLR	Toll-liike receptor
TNF-α	Tumor necrosis factor α
OM	Outer membrane



Chapter 1: Introduction

1.1 Background

Dementia is characterized as a steady and sequential decline in cognitive ability with far-reaching effects that restrict independent occupational and social functioning.¹ It is best described as a syndrome rather than a distinct illness, which has been recognized as the main cause of disability and dependency amongst older individuals.² It is caused by a multitude of conditions or injuries that interfere with the brain's activity and performance, leaving patients with impaired memory, communication, and self-care skills.¹

Depending on the disease subtype, symptoms can manifest in various forms among different individuals.³ However, clinically, these symptoms are broadly categorized into four impairment groups, namely memory, cognitive, behavioural, and physical impairments.³ Individuals with mild dementia often display impaired episodic memory, spatial navigation, performance in social and work environments, and difficulties remembering recently learned information such as peoples' names and specific details of personal objects (agnosia).^{3,4} This is partially due to cortical damage that occurs along pathways that descend in the hippocampus and defects in the anterior-temporal system.⁴ In the moderate stage of dementia, individuals tend to display erratic changes in mood and sleeping patterns, difficulties in language and communication (dysphasia), and problems performing manual tasks like dressing themselves (apraxia).³ As the disease progresses to the severe stage; individuals become physically impaired and more dependent on caregivers for assistance with personal care as they become frail and confused.¹

The prevention and treatment of dementia is a global health issue. A cost of illness comparison study showed that in 2015 the global cost of dementia was 818 billion USD.⁵ According to a census conducted in 2011, approximately 2.2 million South Africans are suffering from some form of dementia, wherein a large proportion of this number is a result of the dementia-HIV/AIDS complex.^{6,7}

Chapter 2: Literature Review

2.1 Introduction to Alzheimer's Disease-type dementia

Serving as the most common cause of dementia, Alzheimer's disease (AD) is an irreversible and steadily advancing neurodegenerative condition characterized by cognitive and behavioural deterioration.⁸ The term "Alzheimer's disease" was authored by Emil Kraepelin in 1910.² This was in honour of his colleague Alois Alzheimer, a German psychiatrist and neuroscientist who was the first to note the most distinctive neuropathological brain change; neurofibrillary tangles (NFTs) in the brain of a female patient following her hospitalization for psychiatric related symptoms.²

Alzheimer's disease is known to only affect humans, with a disposition that exists across all racial groups.⁵ Due to a higher life expectancy and decreasing oestrogen levels with age-as a result of menopause- women are more vulnerable to developing AD than males.⁶ While the disease is often paralleled with aging, researchers have realized that it can arise at any age.⁷ The detail for this occurrence is however still unknown. Currently, there is no cure for AD; therefore, comprehensive understanding of the disease and the development of therapeutic targets are of particular interest to researchers.

2.2 Pathological hallmarks of Alzheimer's disease

The molecular premise underlying the pathology of AD is controversial.⁷ Nonetheless, two prominent pathological hallmarks have been widely recognized and accepted. First, AD is characterized by the extracellular deposition and aggregation of amyloid- β (A β) proteins, in the form of senileclumps known as plaques and second, the intracellular expression of aggregated tau proteins, in the form of hyperphosphorylated filaments known as NFTs.⁸

2.2.1) Amyloid hypothesis

Aβ peptides are derived from a type-1 transmembrane glycoprotein known as amyloid precursor protein (APP).⁹ APP is predominantly found in the synapses of neurons and various tissue structures including the brain, heart, thymus.⁹ APP processing is explicitly controlled by neuronal activity.¹⁰ Neurons that express high levels of APP favour the amyloidogenic pathway, in that they facilitate the process of neurotoxic Aβ

peptides formation.¹⁰ Therefore, Aβ accumulation in brain regions with high neuronal activity gives rise to AD vulnerability.¹⁰ In the amyloidogenic pathway, APP is enzymatically cleaved by β-secretase and γ-secretase to yield a Aβ peptides of 37 to 49 amino acids in length into the extracellular space, cerebrospinal fluid, and plasma.¹¹ These newly formed Aβ peptides exist in various forms; monomers, oligomers, fibrils, and plagues.¹² Essentially, A β monomers aggregate to form soluble oligomers that may spread across the brain and deposit as fibrils and plaques.¹² Aβ fibrils are insoluble, degradation resistant fibres made from aggregated oligomers and Aß plaques are formed by large, unbranched and insoluble aggregated fibrils.¹² Aß peptides with 40 and 42 amino acids are the most prevalent, where un-mutated A β_{42} is the most pathogenic isoform.¹³ Because of their additional hydrophobic residues, Aβ₄₂ monomers have rapid aggregation kinetics which makes them more prone to aggregation- hence they are closely linked to AD.¹⁴ However, earlier studies indicate that both A β_{40} and A β_{42} (A β_{42} :A β_{40} ratio) peptides influence each other's aggregation and toxic properties, such that the aggregation of A β_{42} is restricted by the presence of A β_{40} , whilst the presence of A β_{42} monomers promotes A β_{40} aggregation.^{15,16,17,18,19,20,21} Also, it has been reported that the accumulation of insoluble AB42 is a major contributor to AD pathogenesis.¹²

In AD, the amount of Aβ peptides produced remains constant but the clearance rate is significantly lowered.²² This is because excess native Aβ monomers are subject to abnormal and highly specific conformational changes that result in fibrillar structures represented as plaques.¹¹ Amyloid plaques can induce cellular damage and death via apoptosis or necrosis using receptor-ligand interactions or the disturbance of plasma membranes if the fibril is large enough.^{23,24} Examples of this cellular damage is observed when Aβ plaques interact with microglial receptors and through Aβ-driven lipid peroxidation, they cause inflammation and increases the output of reactive oxygen species (ROS).^{25,26} These pathways give value to the effects of amyloid deposition (amyloidosis) and its association with AD. To summarize, the amyloid hypothesis suggests that plaque formation prompts a series of events including the formation of histopathological lesions, NFTs, synaptic dysfunction and toxicity, neuronal death, microglial and astrocyte activation and mitochondrial dysfunction; all of which trigger neuroinflammation within the cerebral cortex, hippocampus and amygdala.²⁷

2.2.2) Tau hypothesis

Tau protein, a member of the microtubule-associated proteins (MAPs), is a heat-stable hydrophilic protein found in the axon of a neuron.²⁸ Tau protein is encoded by the MAPT gene and due to alternative splicing of its different mRNAs species; six notable tau isoforms are produced in the human brain.²⁹ The role of tau protein is to promote microtubule polymerization and stabilization.³⁰ Most importantly, tau serves to maintain neuronal morphology and regulate axonal transport.³¹ It has also been associated with defective oxidative phosphorylation and apoptotic activity that results in mitochondrial fragmentation.³¹ Unregulated mitochondrial dynamics promote an over-production of ROS that are capable of inducing post-translational modifications in proteins across a cell, thereby implicating tau protein in neurodegenerative diseases.³¹

In tauopathies like AD, changes in the structure, function, and quantity of tau protein occur.³¹ Patients with AD display a higher quantity of tau protein in all forms; normal, phosphorylated or aggregated.³² Aggregates appear as NFTs and their degree of phosphorylation is almost four folds greater than in the normal human brain.^{32,33,34} Aging is also an important risk factor for tauopathies because it coincides with a reduced tau protein turnover that results in its accumulation.³⁵ Similarly to Aβ peptides, excess tau proteins are subject to post-translational alterations such as phosphorylation, aggregation, glycosylation and glycation, all of which raise proteotoxicity.³⁵ Due to its phosphoprotein nature, the biological activity of tau protein is largely controlled by its degree of phosphorylation.³¹ In AD, tau is hyperphosphorylated by several proteins such as glycogen synthase kinase 3ß (GSK3β) and cyclin-dependent protein kinase 5 (Cdk5).³⁶ This abnormal and irreversible hyperphosphorylation causes an involuntary delay in neuronal activity and resultant progression in neurodegeneration.³⁷ Hyperphosphorylation, which precedes the formation of NFTs, decreases tau's affinity for microtubules thus increasing the quantity of unbound protein to abnormal levels.³⁶ Unbound tau proteins create nonfibrillary deposits which undergo further structural modifications where they transform from an unfolded pre-tangle to a more defined β -pleated sheet that self-assembles into NFTs typically found in the stoma of neurons.³⁶

NFTs are described as cytotoxic intraneuronal masses of insoluble cytoskeletal components comprised mainly of phosphorylated tau proteins capable of inducing neuronal death, thereby contributing to AD.³⁷ In the brain, a substantial amount of NFTs are localized in the transentorhinal and entorhinal layer.³⁷ They spread across the brain and facilitate the appearance of A β plaques in the hippocampus as AD progresses.³⁷ Findings by Binder, L and colleagues suggest that NFTs play a crucial role in the progression of AD by promoting a synergistic relationship between A β and tau proteins that intensifies neuronal loss.²²

In short, A β peptides and NFTs are deposited mainly into the hippocampus, amygdala and entorhinal cortex where they impair learning, memory, and behavioral circuits.¹² Jointly, A β plaques connect to their binding receptors and drive downstream pathways that trigger tau hyperphosphorylation, and ROS production.¹² This results in inflammatory reactions which induce excessive calcium influx, synaptic and mitochondrial dysfunction and neuronal death, ultimately giving rise to AD.¹²

2.3 Brain regions affected in Alzheimer's disease

Neuropathological lesions associated with AD occur decades before the onset of clinical symptoms.³⁸ Difficulties in detecting AD during its primary stages are common. This often results in intensified symptoms such as the complete loss of intellectual ability, and great financial responsibility, not excluding the emotional and physical distress experienced by the caregiver.³⁹ Currently, clinicians are limited to providing a provisional diagnosis which often occurs at the last stage of the disease, because even though the preclinical stage is accompanied by clinical signs and symptoms, these are often obscured or misinterpreted as symptoms of other cognitive illnesses or old age.⁴⁰ Nonetheless, an accurate diagnosis requires posthumous assessment.⁴¹

As a result, considerable efforts have been made to establish neuro-imaging techniques for early detection and slowing disease progression such as computed tomography (CT), magnetic resonance imaging (MRI) and positron emission tomography (PET).

Neuropathological lesions associated with AD are coupled with structural irregularities in the brain, particularly the medial temporal lobe (MTL) and hippocampus.⁴² Initial A β aggregates appear on most parts of the cortex even during typical aging, whereas tau

pathology begins in the transentorhinal region and progresses in an activity-dependent fashion across sensitive functional pathways.^{43,44,45}

The MTL is a network of anatomically similar structures consisting of the: parahippocampal gyrus [dentate gyrus (DG), subicular complex and Cornu Ammonis (CA) subfields] as well as the neighboring entorhinal cortex, perirhinal cortex, and parahippocampal cortex.^{46,47} The MTL is the first brain region to display neurodegeneration (atrophy).⁴⁸ Atrophy of the MTL is frequently observed on MRI scans of AD patients and is correlated to neural damage and disease severity.^{49,50} Reports by Morrison, JH⁵¹ indicate that the entorhinal-hippocampal circuit is impaired during the early stages of AD, followed by a steady dissociation of the MTL regions, resulting in an ultimate loss of communication between the surrounding neocortex. This may possibly highlight the pathway of cognitive decline during aging in the onset of AD. Also, MTL atrophy is linked to reduced performance in memory-related tasks.⁵²

Histopathological examinations have revealed that the hippocampus is influenced by Aβ plaques and NFTs in the primary stages of the disease, thus confirming its significance in diagnosis.⁵³ The hippocampus is a highly vulnerable continuation of the cerebral cortex distinguished as an S-shaped layer of tightly packed neurons embedded deep within the MTL to form part of the limbic lobe.⁵⁴ It is a highly complex structure with ductile properties that make it easily modifiable by neuropathological disorders.⁵⁵ The hippocampus plays a role in neurogenesis, spatial memory, and orientation, emotional behavior, modulation of hypothalamic functions and the major functions of learning, memory formation, and retention.^{56,57,58} In rats, learning and memory are inherently associated with the dorsal hippocampus, whilst emotional behavior is associated with the ventral hippocampus.⁵⁹

As outlined above, the hippocampus is a major target for amyloid and tau pathology. By use of functional connections, NFTs first invade the transentorhinal and anterolateral entorhinal cortex followed by the CA subfields and the neocortex.^{60,61} The spatiotemporal sequence of NFT spreading matches the progression of brain atrophy.⁶² Unlike A β , the degree of NFT expression strongly correlates with disease severity.⁶³ A β deposition occurs mainly in the hippocampus, neocortex, and cerebrovasculature (also known as cerebral amyloid angiopathy).⁶⁴

That said, over the past decade, a third characteristic has been implicated in the pathogenesis of AD that may possibly present a link between the aforementioned hallmarks.⁶⁵ The brain tissue of AD patients indicates the occurrence of a persistent inflammatory response, neuroinflammation, which arises as a consequence of disrupted equilibrium between pro-inflammatory and anti-inflammatory signals.⁶⁶ Neuroinflammation is progressively being acknowledged as a core feature of neurodegenerative diseases such as AD not merely because it is linked to neurodegeneration but it also promotes and intensifies both A β and tau protein abnormalities.^{65,67}

2.4 Dementia-associated inflammation

2.4.1) Overview of systemic inflammation

Generally, a regulated systemic inflammatory response is a localized and beneficial defense mechanism facilitated by the host immune system to remove harmful agents such as an infection, cellular or tissue damage, irritation or trauma through various molecular and cellular events which act to minimize and clear the harmful agent, followed by restoration and repair processes.⁶⁸

Within the context of microbial infection; systemic inflammation is triggered when cells of the innate immune system alongside various epithelial cells use their surfaceexposed pattern recognition receptors (PRRs) to detect either; a) pathogenassociated molecular patterns (PAMPs), which are microorganism-derived and evolutionarily preserved exogenous structures found on pathogens, or b) damageassociated molecular patterns (DAMPs), which are endogenous stress cues produced by cells to prompt spontaneous cell death.^{69,70} Together, PAMPs and DAMPs form biomolecules that bind to specific receptors of the PRR family (i.e. Toll-like receptors) to enhance autophagy- thereby preventing genome instability and necrosis.⁷¹ Autophagy is an essential lysosome-mediated process, aimed at maintaining homeostasis.⁷¹ It involves the removal of toxic protein aggregates and degradation of macromolecules, (i.e. carbohydrates, proteins) intracellular pathogens and organelles (i.e. mitochondria, endoplasmic reticulum).^{71,72}

At the circulatory level, inflammation is marked by changes in vascular permeability that function to increase blood flow to the site of injury, and stimulate leukocyte chemotaxis that is driven by the release of pro-inflammatory cytokines and chemokines.⁶⁸ Whilst at the tissue level, inflammation is marked by: redness and heat (caused by the increase in blood flow), swelling (caused by fluid accumulation), pain (caused by the release of inflammatory mediators which excite neuronal endings), and tissue immobility.⁷³ Despite the fact that the resultant inflammatory response is dependent on many factors of the initial stimulus, the processes share a common mechanism consisting of i) recognition of the harmful stimuli using cell surface PRRs; ii) stimulation of inflammatory pathways; iii) release of inflammatory mediators and the iv) mobilization of inflammatory cells.⁷⁴ Ultimately, inflammation that runs unregulated in time, space or intensity is a hallmark of several pathologies.⁷⁵

2.5 Neuroinflammation

Neuroinflammation is a chronic and complex inflammatory reaction across the central nervous system (CNS); the brain or spinal cord.⁷⁶ It is mediated by biochemical and cellular reactions comprising the formation and release of pro-inflammatory cytokines such as; interleukins (IL)-1 β ,IL-6, IL-18, tumor necrosis factor α (TNF- α), interferons (IFNs), colony-stimulating factors (CSF), chemokines; CCL2, CCL5, CXCL1, ROS, nuclear factor kappa-light-chain-enhancer of activated B cells (NF- κ B), enzymes cyclooxygenase-2 (COX-2), inducible nitric oxide synthase (iNOS), coagulation factors like platelet-activating factor and secondary messengers; nitric oxide (NO) and prostaglandins.⁷⁶ Neurodegenerative conditions such as AD is linked and frequently preceded by neuroinflammation which plays an essential dual role.⁷⁷ This is partly due to the fact that A β accumulation induces a chronic neuroinflammatory state which participates in AD progression.^{78,79}

The degree and complexity of neuroinflammation are influenced by the setting, time span and course of the primary insult, given that each tissue displays diverse features of inflammation due to biochemical, immunological and physiological differences.^{76,80} Acute neuroinflammation, however, has advantages such as immune conditioning, neuro-immune communication, injury reduction and axonal and myelin restoration.^{77,76}

2.5.1) The complex foundation of neuroinflammation

Neuroinflammation participates in disease resolution and provides neuroprotection during an acute-trial reaction by restricting activation of the pathogenetic cascade but on the other hand, it may become deleterious under a chronic response, as seen in clinical conditions.^{81,82} Vicious or pathological neuroinflammation is linked to a disruption of the blood-brain barrier (BBB), microglial activation and proliferation, stimulated release of inflammatory mediators, astrocyte mobilization and abnormal neuronal signaling all of which may contribute to the degradation of the CNS microenvironment, consequently resulting in clinical implications.^{83,84} This phenomenon is depicted in **Figure 1**.⁸⁵ The CNS microenvironment possesses unique characteristics. It has a distinct chemical composition and neurotransmitter pool which promotes neural function, in addition to its poor protein content which reduces cell proliferation and oxidative stress.⁸⁶ This microenvironment is rarely subjected to systemic toxins that minimize neural impairment.⁸⁶ Moreover, it has the capability to restrict the movement of inflammatory mediators, thereby preventing local inflammation.⁸⁶



Figure 1: The complex inteaction between infection or injury, the induced inflammatory response, and subsequent clinical implications.

2.5.2) Blood-brain barrier

The BBB is a highly differentiated and stratified membrane made from non-fenestrated endothelial cells connected to each other by tight junctions and pericytes, which surrounds two continuous basement membranes and astroglial end-feet to collectively form the neurovascular unit (NVU).⁸⁶ At the molecular level, the BBB is embedded with receptors, transporter proteins, and exoenzymes which direct and regulate interactions along the barrier.⁸⁶ Taken as a whole, these components play a key role in the provision and maintenance of a homeostatic and immuno-protected CNS microenvironment by conserving the integrity of the BBB and limiting interactivity between the innate immune system and acquired immune system.^{87,83} In AD, systemic inflammation driven by microglia, macrophages, and immune cells is linked to CNS dysfunction, disrupted BBB, reduced bulk flow of cerebrospinal fluid (CSF) over the BBB and accelerated Aβ deposition in the hippocampus.^{88,89,90} At the same time, cerebrovascular deposition of A^β plaques in the CNS initiates pro-inflammatory and cytotoxic events whilst potentially playing a role in the degeneration of pericytes, endothelial cells, and smooth muscle cells, all of which promote extensive BBB permeability.^{91,92} BBB dysfunction also corresponds with higher levels of

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hyperphosphorylated perivascular tau protein close to vital hippocampal blood vessels.⁹³ Thus, both A β and tau pathology may advance BBB disruption to exacerbate neuroinflammation and neurodegeneration.⁹⁴ Reversibly so, neurodegeneration coexists with microgliosis and astrogliosis.^{95,96}

2.5.3) Glial cells of the CNS

Neuroglia (glia) are non-neuronal tissue cells of the nervous system isolated from the rest of the body by the BBB.⁹⁷ They serve as support structures for neurons whilst providing nourishment, maintaining homeostasis and form the myelin sheath.⁹⁷ Glia is broadly categorized into two classes; macroglia and microglia. Macroglia are further split into 7 different cell types, each playing a unique role (**Figure 2**).^{97,98,99,100,101}



Figure 2: Classification of neuroglial cells, with their main roles in the CNS and *peripheral nervous system (PNS).

Evidence that supports neuroinflammation as the link between A β deposition, tau abnormality and neurodegeneration arises from human and rodent studies where activated microglia, which is associated with disease progression, were observed near amyloid aggregates within the AD brain.^{102,103}

2.5.3.1) Microglia

Microglia are tissue-resident phagocytic cells of the innate immune system ubiquitously dispersed within the gray and white matter of the CNS.¹⁰⁴ They account for ±10%-15% of all glial cells and their amount generally stays constant from late postnatal development to old age due to concurrent proliferation and apoptosis.^{105,106} Microglia are involved in CNS function, evolution, homeostasis and disturbances.¹⁰⁵ They also participate in phagocytosis, synaptic maturation, pruning and protection, neurogenesis and the regulation of cognitive functions.^{105,107} From an immunological perspective of infected or injured tissue; microglia detect inflammatory signals, produce and release inflammatory mediators and participate in both the inflammatory response and resolution processes.¹⁰⁷ This primary function of immune surveillance is facilitated by a diverse range of receptors for neurotransmitters, cytokines, chemokines and toll-like receptors (TLRs).^{108,109}

2.5.3.2) Reactive microglia and its implications

In their inactivated resting state, as is the case in healthy adult brains, they are structurally referred to as ramified units with small stationary cell bodies and cell processes that constantly expand and contract; continuously scanning their assigned environment and relaying information straight to other microglia, nerve fibers, astrocytes, T-cells, myeloid progenitor cells and blood vessels.¹⁰⁵ Upon detection of a pathological stimuli (i.e. infection,neuronal degeneration or protein presence or aggregation) within the CNS, their impressive plasticity and regular state of motion enables them to rapidly respond by transitioning into an activated state (microglial activation) of altered structure and molecular composition characterized by contracted cell processes, enlarged cell bodies and migration towards the site of injury where they eradicate pathologens or damaged cells and execute inflammatory functions via innate immunity.¹⁰⁵ On a molecular scale, microglial activation involves cytoskeletal rearrangements that change the sequence of receptors displayed on their surfaces in addition to the amplified expression of potentially cytotoxic molecules such as cytokines, chemokines, prostaglandins and ROS.¹⁰⁷

In the event of an exaggerated immune response and consequent overproduction of immune molecules: microglia lead to a significantly modified neural microenvironment,

consequently characterizing them as a hallmark of neuroinflammation and brain pathology.^{110,111} In AD, it is speculated that the force driving microglial activation and the resultant overproduction of inflammatory mediators is the expression of A β plaques; for the reason that some studies have indicated that activated microglia can engulf these plaques but because the microglia become so inflated and inefficient at binding A β that after a certain period they become incapable of processing the plaques.^{65,112} Microglia also have the ability to respond to APP and NFTs.¹¹³ During preclinical AD, the initial inflammatory reaction provides beneficial effects as it contributes towards the clearance of A β plaques, however, sustained inflammation and microglia-associated neurotoxins worsen AD pathology as they sustain reactive microgliosis which results in accumulated A β proteins and persistent cytokine signalling that recruits more microglia to the plaques, ultimately causing neuronal damage or death.^{112,114,115}

2.5.3.3) Astrocytes

Similarly to microglia, astrocytes are CNS-resident signal conductors of the innante immune system.¹¹⁶ They possess a variety of functional and structural features involved in several physiological activities.¹¹⁶ Astrocytes are the most abundant neuroglial cell.¹¹⁷ Their pathological reaction is known as reactive astrogliosis, a complex, multistep and disease-specific response marked by cellular, morphological and functional changes directed towards neuroprotection and the restoration of damaged neural tissue in response to CNS injuries.¹¹⁸ Astrogliosis is associated with important positive functions, however, under certain conditions it may lead to deleterious outcomes.¹¹⁹ Apart from supplying structural support to neurons, astrocytes are essential for the sustainability, homeostasis, and functioning of a healthy CNS.¹²⁰ Other functions include synaptogenesis, synaptic plasticity, neurotransmitter uptake and reuptake, receptor trafficking, gliotransmission, regulation of extracellular space composition and volume, maintenance of the BBB, inflammation and higher cognitive functions.^{120,121,122,123,124}

2.5.3.4) Astrogliosis and its implications

In light of their key role of facilitating CNS functions, it is unsurprising that any disruptions thereof promote the onset and progression of cerebral pathologies like

dementia.¹²⁵ Reactive astrocytes receive directive information from surrounding cells and deliver different messages to effecter cells that influence or initiate an appropriate response.¹¹⁹ Such astrocytes are marked by functional impairments, accumulation of hypertrophic astrocytes surrounding Aß plaques and amplified expression of glial fibrillary acidic protein (GFAP) in all four lobes of the brain and the CSF.¹²⁶ GFAP is a protein involved in cell communication, BBB function, mitosis, and repair processes, and its increased expression is associated with AD progression.^{126,127} Similarly to microglia when exposed to A^β plaques, astrocytes can initiate inflammation and the release of cytokines, chemokines and ROS, which exacerbates the progression of dementia.¹²⁸ Moreover, astrocytes produce and release gliotransmitters which facilitate synaptic plasticity and regulate learning and memory in distinct brain regions like the cerebral cortex and hippocampus.^{129,130} Any disturbances in gliotransmission (such as the presence of A β plaques) causes an increase in calcium (Ca²⁺) signaling, altered neural homeostasis, and synaptic transmission which also advances disease pathology.^{131,132} Not only do reactive astrocytes produce a variety of chemokines that permeate the BBB and attract inflammatory cells, they also produce molecules that dampen the effects of inflammatory cells.^{133,134}

In regions such as the cortex, hippocampus, and cerebellum, hypertrophic astrocytes bind and sheathe amyloid plaques while penetrating any unbound plaques resulting in their fragmentation, diffusion and varied morphology.¹³⁵ Extracellular molecules such as purines, growth factors, plasma proteins, $A\beta$ or steroid hormones invoked by endotoxins such as lipopolysaccharides (LPS) have the ability to initiate or regulate astrogliosis.¹¹⁹ When peripheral infections arise from immune system invaders like LPS, they produce PAMPs which infiltrate the BBB, and upon exposure to astrocytes, will induce and exaggerate their pro-inflammatory potential thereby steering astrogliosis towards cytotoxicity.^{136,137}

LPS is implicated in (neuro) inflammation, given that its systemic release is characterized by the activation of a strong inflammatory response and the infiltration of leukocytes from the periphery to the CNS resulting in neuroinflammation, mitochondrial dysfunction, and neurodegeneration.¹³⁸
2.6 Lipopolysaccharide

2.6.1) Structure of lipopolysaccharide

The membrane of Gram-negative bacteria (**figure 3**¹³⁹) such as *Escherichia coli*, is comprised of a cell envelope made from two membranes parted by peptidoglycan containing, hydrophilic cavity called the periplasm.¹⁴⁰ The outer membrane (OM) is the foremost layer of the envelope, composed of an asymmetric phospholipid bilayerinterlaced with proteins - serving as the inner leaflet, while LPS, serves as the outer leaflet.¹⁴¹



Figure 3: The membrane structure of gram-negative bacteria.

This bacterial endotoxin is a negatively charged, amphiphilic and heat-stable glycoconjugate that readily clusters into micelles or vesicles in aqueous solutions.¹⁴² It accounts for 10% - 15% of the overall quantity of molecules present in the OM and makes up 75% of the entire bacterial surface.¹⁴³ LPS is characterized by three distinct structural components: (i) Lipid A -the innermost hydrophobic component that is responsible for the toxicity of LPS- made from a disaccharide acylated with fatty acids; (ii) the core domain which joins lipid A to the O-antigen, which is made from a non-repetitive short-chain oligosaccharide of approximately 9 sugars; and the (iii) O-antigen, an unevenly distributed hydrophilic oligosaccharide of repetitive glycan units that extends toward the bacterial surface.¹⁴⁴ The formation and assembly of LPS components occurs in the cytoplasm, followed by translocation to the external

membrane.¹⁴³ The O-antigen exhibits a high level of inter- and intra-species variation, making it the key component governing strain specificity.¹⁴⁵

2.6.2) Function of lipopolysaccharide

LPS carries out two significant functions: firstly, it facilitates a defense reaction against adverse environmental conditions by providing a protective barrier to Gram-negative bacteria; consequently making it essential for bacterial resistance and viability.¹⁴⁶ Secondly, because of its quick recognition and sensing, LPS can initiate an immunological reaction within the host immune system leading to the removal of a bacterial infection.¹⁴⁷ In essence, it activates the complement system to enhance inflammation and inhibit the development of the membrane attack complex (MAC) on the pathogen cell surface.¹⁴³ However, this immunological reaction needs to be balanced or it may become detrimental to the host.¹⁴⁸ Rampant bacterial overgrowth causes an abundant release of unbound/free LPS into the circulatory system which activates monocytes and endothelial cells, inducing an exaggerated systemic immune response that involves an excess production of inflammatory mediators that cause tissue damage, septic shock or death.¹⁴⁸ In contrast, low amounts of LPS benefit the host by granting protection against local infections through the immediate activation of immune reactions such as the production and activation of dendritic cells, natural killer cells, B- and T-cell functions, initiation of fever and the complement cascade.¹³⁸

2.6.3) Proteins involved in liopolysaccharide induced toll-like receptor 4 signalling cascade

LPS is easily identified by the innate immune system.¹⁴⁰ Contact with the immune system is initiated through recognition of the lipid A receptor, which prompts the involvement of the adaptive immune response through the O-antigen.¹⁴⁰ The innate immune system uses PRRs such as Toll-like receptor 4 (TLR4) and glycoprotein MD-2, as the first line of defense that forms a complex functioning to identify PAMPs (i.e. LPS) for the purpose of stimulating intracellular signaling pathways that activate the cellular expression of inflammatory mediators. ^{147,149} Cluster of differentiation 14 (CD14) is a glycoprotein anchored on the membrane of monocytes and macrophages (mCD14) or soluble in plasma (sCD14) which increases the potency of the immune response to LPS via stimulation of the TLR4 signal pathway. ¹⁵⁰

2.6.4) Mechanism of action of liopolysaccharide induced toll-like receptor 4 signalling cascade

LPS-binding protein (LBP), a lipid transferase present in the liver and released into plasma, will first detect the presence of endotoxin bacterial components and proceed to bind to the lipid A domain of LPS, dislodging it from the bacterial membrane.¹⁵⁰ LBP then shuttles LPS monomers to CD14 proteins and these CD14 proteins will finally deliver the LPS monomers to a TLR4/MD2 complex.¹⁵⁰ The receptor binding of LPS to the TLR4/MD2 complex activates the fusion of cytosolic translocated intimin receptor (TIR) domains of TLR4 causing the mobilization of downstream accessory proteins.¹⁵¹ This TLR4 signaling transduction pathway activates the expression of pro-inflammatory cytokines and IFNs via NF- κ B nuclear translocation and IFN regulatory factors, respectively.^{148,152} Importantly, TLR4 can be stimulated by LPS at a concentration as low as 0.1 ng/mL.¹⁵³ The signaling cascade is also characterized by the release of ROS, anti-inflammatory cytokines i.e. IL-1 receptor antagonist (IL-1Ra), IL-4, IL-10, IL-13, anti-microbial peptides, TNF- α , prostaglandin E2, IFN- γ , C-reactive protein, hydroxyl radicals, NO and O₂^{-.150,154} **Figure 4** represents the LPS induced TLR4 signalling cascade.



Figure 4: The LPS induced TLR4 signaling cascade that activates an innate immune response. Source: Supplementary Figure 1, Park, BS *et al* (2012).¹⁵⁵

LBP-deficient mice show a hypo-responsive effect to LPS- as demonstrated by reduced amounts of serum inflammatory mediators and increased mortality rates after infection with *Escherichia coli*.¹⁵⁶ In the gut, LBP is secreted into the lumen not only to detect bacterial invasion but to also regulate symbiont-derived LPS, making it fundamental for intestinal homeostasis.¹⁵⁷

2.7 Probable sites of lipopolysaccharide entry

2.7.1) Gut microbiome and its development

All mammals are colonized by plentiful and heterogeneous populations of microbial species that are central to the organization and function of the host; in that, they play a role in energy regulation, metabolism, intestinal epithelial health, immunological processes, and neurodevelopment.^{158,159,160} The human gastrointestinal tract (GIT) houses 10¹³ -10¹⁴ different species of symbiotic, commensal and pathogenic microorganisms consisting of bacteria, archaea, viruses, fungi, phages and protozoa that collectively form an ecological community known as the microbiome.^{161,158} The microbiome is an adaptive community susceptible to extensive changes in composition and efficacy in response to numerous factors such as diet, external environmental signals (i.e. geography and culture), pathological conditions and medical therapies (i.e. antibiotics).¹⁶² However, microbiome inhabitation is not limited to the gut, other body areas of microbial colonization include the skin and oral cavity.¹⁶³ Microbial flora in the neonatal GIT is inherited maternally during childbirth where the species of the flora is largely influenced by the mode of delivery.¹⁶³ In toddlers, the composition, heterogeneity, and efficacy parallel the healthy adult gut microbiome and is related to usual developmental milestones, up until late adulthood (~65 years) where there is a shift that favors Bacteroidetes and Clostridium cluster IV.^{164,165} Considering that almost 70%-80% of the immune system inhabits the mucosal lymphoid tissue of the GIT, it is unsurprising that growing evidence proposes that microbiota participate in the physiology, development, and pathology of their host organisms, thus implicating them in both health and disease states, where they develop and maintain host immunity.166,167

2.7.1.1) Limitations of the gastrointestinal tract

The intestinal lumen is sheathed by a continuous layer of epithelial cells that form a protective barrier and interaction surface for internal organs against microbial pathogens and antigens and the relay of information between the host and gut microbiota, respectively.¹⁶⁸ However, intestinal bacteria are capable of synthesizing and releasing a variety of substances that disrupt the integrity of the epithelial barrier by increasing its permeability and enabling the uncontrolled movement of gut microbiota into the lamina propria, where a majority of intestinal immune cells are localized within the gut-associated lymphoid tissues (GALT).^{169,170} Consequently, systemic inflammatory responses that compromise the integrity and functionality of the BBB are triggered.¹⁶⁹ When the BBB is compromised, microbes from the gut infiltrate the CNS environment to induce neuroinflammation which ultimately leads to neurodegeneration.¹⁷¹

2.7.1.2) Brain-Gut-Microbiome Axis

The gut-brain axis describes a neural, immune, endocrine and metabolic mechanism of communication between the microbiota, GIT and the brain in a bidirectional pathway by virtue of the autonomic nervous system (ANS) and circumventricular organs (CVOs).^{169,172} The ANS is at the epicenter of this interaction through its signaling divisions: the sympathetic (via the prevertebral ganglia) and parasympathetic (via the vagus nerve) nervous systems. Jointly, each division functions to detect gut microbiota and transmit information to the CNS where it is integrated, modified and a suitable reaction is implemented.^{173,174} Afferent fibers of the vagus nerve express TLR4s that are capable of detecting gut pathogens like LPS and sending signals to the brain to activate inflammatory pathways.¹⁶¹

Elderly individuals are faced with the possible development of a hyper-stimulated immune state that is characterized by chronic, low-grade inflammation called inflammaging.¹⁷⁵ Inflammaging is associated with a sustained inflammatory reaction of the GIT mucosa that is induced by an age-related shift in the microbiome composition, heterogeneity, and efficacy resulting in increased: permeability of the epithelial barrier, bacterial translocation into the bloodstream, pro-inflammatory

cytokines, BBB permeability, and neuroinflammation.¹⁷⁶ **Figure 5**¹⁷⁷ illustrates the gutbrain communication and the presumed involvement of gut microbiota in AD.

In conjunction with the complex interaction between the microbiome, gut, and brain, it is meaningful to note that the gut microbiome is interrelated to the oral cavity microbiome by composition and heterogeneity.¹⁶³ Recent evidence suggests that there is an association between oral pathogens and changes in gut microbial composition in addition to inflammatory changes in brain tissue.^{163,178}



Figure 5:¹⁷⁷ Intestinal microbiota from the gut lumen is the source of a great number of bacterial factors such as bacterial amyloids (produced by *Escherichia coli*), LPS and other toxins, that are detected by TLRs of enteroendocrine cells (EECs) which are found on the epithelial membrane.^{179,180} (i)These factors are capable of disrupting the permeability of the epithelium to promote their movement into the bloodstream. (ii) In the blood lumen, these factors can penetrate the GALT where they activate immune cells to proliferate e.g. T-cells resulting in direct communication with the innate immune system.^{171,181} (iii) Regulatory T-cells circulating in the bloodstream survey the blood, lymphoid tissue, and CSF, this altered cellular environment stimulates T-cell brain permeation.¹⁸²(iv) Bacterial factors such as LPS activate inflammatory pathways that trigger increased cytokine levels and BBB permeability; they may also be detected by endothelial TLR4s of the brain thereby stimulating neuroinflammation and CNS pathologies.¹⁸³ (v) Microbial metabolites are capable of activating tight junction proteins, subsequently restoring BBB integrity, though, they are also capable of infiltrating the BBB, priming microglia and astrocytes, stimulating neuroinflammation, disrupting Aβ clearance and increasing neurotoxicity.^{184,185}

2.7.1.3) Oral cavity pathogens

The periodontium is a set of tissue structures composed of the gingival, periodontal ligaments, cementum and alveolar bone, that surround and support the teeth.¹⁸⁶ Periodontal disease (periodontitis) is a persistent inflammatory condition of the periodontium.¹⁸⁷ In its progressed state, periodontitis is presented by damage to the periodontal ligament and decay of the alveolar bone.¹⁸⁷ Risk factors for this disease include smoking, poor oral hygiene, hormonal changes in females, medications, stress and advancing age.¹⁸⁸

2.7.1.4) The interrelation between oral cavity flora and the gut microbiome

Growing evidence suggests an association between periodontitis and systemic diseases including AD.¹⁸⁹ It is unclear whether periodontal pathogens initiate the onset of systemic diseases or if systemic diseases trigger the accumulation of periodontal pathogens.¹⁸⁸ However, it is noted that the swallowed salivary bacterial content of patients with periodontitis is ~10¹² bacteria/day.¹⁹⁰ A study by Tiisanoja, A *et al* (2019).¹⁹¹ found a stronger association between tooth decay and AD in comparison to other oral diseases such as periodontitis and stomatitis. Nonetheless, research suggests that toxic bacterial molecules like LPS or peptidoglycan may originate from the ingestion of periodontal pathogens (i.e. *P. gingivalis* and *T.denticola*) causing either a change in the composition of the gut microbiome or the activation of an inflammatory response.¹⁹² Mice that are orally infected with *P.gingivalis* exhibit impaired learning and memory skills in addition to depression-associated behavior.^{193,194} Migrating periopathogens and pathogenic gut microbes in the bloodstream are detected by systemic and CNS immune cells.¹⁹⁵ If pathogens are abundant, inflammatory mediators are produced in surplus and released.¹⁹⁵ These mediators disrupt the integrity of the BBB ultimately leading to neuroinflammation and contributing towards AD development.^{195,163} The pathways resulting in pathological outcomes are illustrated in figure 6.



Figure 6:¹⁶³ The pathways linking gut microbiota, oral flora, and AD.

Pathogens from oral diseases cause periodontal inflammation which releases toxic molecules into the intestinal lumen causing infiltration of microbes into the bloodstream and signal transmission to the CNS that destabilizes the BBB, leading to chronic neuroinflammation and possibly AD.

2.8 Using the hippocampus to measure the degree of induced neuroinflammation

Though the pathophysiology of AD has been central to dementia research, the basic processes that facilitate disease pathogenesis and progression are yet to be understood.

Due to a close link between the vulnerability to amyloid and tau pathology, high astrocyte and microglial activity and its role in learning and memory decline, the hippocampus is a classic and valuable model for investigating the effects of dementia-related neuroinflammation.¹⁹⁶ For this study, three important components have been selected which may shed light on the effects of LPS induced neuroinflammation as well as the protective and therapeutic benefits of honey in a dementia-type animal model.

2.8.1) Neuroglial activation in neuroinflammation

Neuroinflammation is prevalent in neurodegenerative conditions. Characterized by three features; glial cell activation, the release of inflammatory mediators and permeation of peripheral immune cells into the brain tissue, its chronic state plays a major role in disease pathogenesis.¹⁹⁷ The neuron-astrocyte-microglia triad is essential for the normal functioning of the brain, any inefficiency in this interaction may trigger neurodegeneration.^{198,199} The surface membranes of astrocytes and microglia contain receptors that bind molecules secreted by neurons.^{200,201} This activates downstream intracellular pathways that induce the degradtion of apoptotic neurons and abnormal cells.²⁰² Also, because Aβ production occurs in several cells including neurons, its presence leads to an innate immune reaction that activates astrocytes and microglia to facilitate its own clearance from the brain.203,204 Characterized by hypertrophic cells and increased hippocampal GFAP expression, the degree of astrogliosis is linked to cognitive deterioration.²⁰⁵ Along with this, microglia is the predominant cell type to engulf fibrillary Aß proteins irrespective of the conditions but because this phagocytic process is dysfunctional in AD, pro-inflammatory mediators escalate and AB deposition rises to overwhelming and pathogenic levels.^{206,207} Marked by an upregulated state at the site of injury following neuronal damage-mediated inflammation, calcium-binding adapter molecule Iba1, is used as a marker for microglial activation and migration during the clearance of tau deposits.208,209,210

Promisingly, new studies exploring alternative dementia treatments suggest that honey may provide prophylactic effects against A β deposition and LPS-induced neuroinflammation on the hippocampus.²¹¹

2.8.2) Honey as a protective agent

Honey is a natural functional food produced by worker bees that has been used as a nutrient and traditional medicine since ancient times.²¹² It is a hypersaturated liquid composed mainly of fructose and glucose in addition to various bioactive compounds and antioxidants including; polyphenols, flavonoids, amino acids, minerals, vitamins B,C,E, enzymes and organic acids.²¹² In favor of its biological, pharmacological and physiological properties, many people resort to honey as an alternative to chemical treatments, also because it has fewer side effects.²¹³ Amongst a myriad of health benefits, shown in **figure 7**, honey is becoming more popular in neural disease research.



Figure 7: A summary of the documented therapeutic effects of honey.²¹⁴

Numerous studies have attempted to elucidate the exact biological pathways that render honey as a neuroprotective agent in dementia. An inquiry by Tonks *et al.* (2003) showed that exposing monocytes to Manuka honey increased the production of TNF- α and interleukins

IL-1β, IL-6 via a TLR4 dependent pathway.²¹⁵ Another study revealed that Manuka honey improves cellular viability while reducing apoptosis and neuroinflammation by lowering caspase 3, p-p38 and p-Erk1/2 protein levels in LPS-exposed macrophages.²¹⁶ The benefits of honey extend to the GIT as well as brain tissue and its respective functions. The high carbohydrate content in honey facilitates *bifidobacterium* proliferation in the GIT whilst the polyphenol content suppresses the growth of intestinal pathogens that incite gut leakiness and allow LPS entry into circulation.²¹⁷ Furthermore, Al-Rahbi *et al.*(2014) found that oral ingestion of honey increases the expression of neurons in the hippocampus in addition to brain-derived neurotrophic factor, which is related to enhanced spatial memory.²¹⁸

Considering the inefficiency and harsh side effects of synthetic remedies of dementia, it is valuable to explore the protective effects of functional foods such as honey.

2.8.3) Behavioural tests in dementia-type models

Animal models are used in a wide range of biomedical studies to investigate and gain insight into specific theories concerned with the physiological, cellular and molecular mechanisms underlying AD dementia etiology. Behavioral assays are also used to quantify cognitive impairments such as learning and memory in animal species. The main goal of these tests is to advance knowledge on already existing treatments while developing new and better effective strategies against diseases affecting humans.²⁵⁴

2.9 Aim and Objectives

With a comprehensive understanding of the implications of neuroinflammation on dementia and the mechanisms underlying its manifestation in the hippocampus, the following questions emerge;

- Doesa chronic systemic LPS exposure cause AD-like brain damage and symptoms? If so, how does it present itself?
- ii. If chronic systemic LPS does indeed cause AD-like brain damage and symptoms, are these effects evident in the behavioral pattern of Sprague Dawley rats?
- iii. Does oral ingestion of honey mop-up the deleterious effects of chronic LPS exposure, both histologically and behaviorally?

These questions give rise to the following aim and objectives:

Aim

The aim of this study is to investigate the effect of neuroinflammation induced by systemic lipopolysaccharide on the hippocampus of 10 weeks old male Sprague-Dawley rats.

In order to achieve the aim the objectives are to

Quantitative biochemical assays

- Assess the progression of amyloid β (Aβ) pathology by measuring the concentration of soluble Aβ₄₂ levels in the hippocampal tissue of male Sprague-Dawley rats using sandwich ELISA assay.
- Determine the number of astrocytes present in the hippocampal tissue of Sprague-Dawley rats by immunofluorescence staining with glial fibrillary acidic protein (GFAP) antibody using confocal microscopy and ImageJ.
- Identify astrocyte activity (astrogliosis) in the hippocampal tissue of Sprague-Dawley rats by measuring fluorescent intensity using a GFAP antibody using confocal microscopy and ImageJ.
- Identify microglial activity (microgliosis) in the hippocampal tissue of aged Sprague-Dawley rats by co-labeling tissue sections with Iba1 and CD68 antibodies and counting the number of Iba1 positive cells using confocal microscopy and ImageJ and measuring the fluorescent intensity of co-localized Iba1 and CD68 positive cells using ImageJ.
- Identify the protective effects of honey on the astrocyte and microglial expression/quantity in the hippocampal tissue of aged Sprague-Dawley rats using ImageJ.
- Identity the amyloidosis in the hippocampal tissue of aged Sprague-Dawley rats by measuring fluorescent intensity using Thioflavin-T (ThT) dye using confocal microscopy and ImageJ.

Behavioral analysis

- Assess the effects of systemically induced neuroinflammation on the short-term spatial working memory of Sprague-Dawley rats using the spatial recognition Y-maze test.
- Assess the effects of systemically induced neuroinflammation on the learning capacity and spontaneous memory of Sprague-Dawley rats using the novel object recognition test.
- Assess the effects of systemically induced neuroinflammation on the exploratory behavior, anxiety and locomotor activity of Sprague-Dawley rats using the open field test.

Chapter 3: Study Design, Sampling, and Statistics

3.1 Study Design

This study adopted a randomized intervention approach using an animal model. A total of forty Sprague-Dawley rats, housed at the University of the Witwatersrand Central Animal Services (CAS) unit was used. The subjects were randomly assigned into four sample groups, with each group containing ten subjects (n=10). The four groups and their treatment were assigned as indicated in **table 1**.

Table	1:	Categorization	of	the	experimental	groups	and	the	treatment	received	per	group.
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Sample group name	Treatment type	Treatment received	Category
PBS	PBS only	Daily SC injection of 0.1M PBS at a volume of 0.1ml/kg for 10 days.	Control group
PBS + H	PBS and Honey	Daily SC injection of 0.1M PBS at a volume of 0.1ml/kg for 10 days + 0.5 ml honey* per kg of rat via oral gavage from day 4 until day 10.	Experimental group 1
LPS	LPS	Daily SC injection of 0.1M LPS dissolved in 0.1M PBS at a volume of 0.1ml/kg for 10 days.	Experimental group 2
LPS + H	LPS and Honey	Daily SC injection of 0.1M LPS dissolved in 0.1M PBS at a volume of 0.1ml/kg for 10 days. + 0.5 ml honey* per kg of rat via oral gavage from day 4 until day 10.	Experimental group 3

*The Manuka honey was mixed 50% v/v with distilled water to enable a comfortable consistency for the rats.

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Prior to termination, behavioural tests using the spatial recognition Y-maze, novel object recognition (NOR) test and open field test were performed on all the animals. This was followed by quantitative sample analysis on the dorsal hippocampal region using ELISA and Immunostaining assays.

3.2 Ethical Considerations

Ethical approval was obtained from the University of Pretoria Health Sciences Research Ethics Committee as well as University of Pretoria Animal Ethics Committee under protocol number 181-2020 (addendum 1). Ethical approval was also obtained from the University of the Witwatersrand Animal Research Ethics Committee, Faculty of Health Sciences under protocol number 2019/07/44/C (addendum 2). Approval from the University of Pretoria, MSc Committee was granted (addendum 3).

The following animal ethical concerns were adhered to:

- All animal subjects were treated with sensitivity and respect;
- No animal subjects were subjected to intentions which were not directly concerned with the research project objectives and methodology;
- Professional standards were upheld in accordance with research training given prior;
- Responsibility for the proper care and use of the animal subjects;
- Integrity towards the animal subjects were promoted through honesty and fairness;

3.3 Sampling Population and Procedures

Sampling Population:

Ten-week-old Sprague Dawley rats, with an average weight of 250-300 grams were maintained at the University of the Witwatersrand, CAS housing facility, as shown in **figure 8 (A-D)**. In this study, the Sprague Dawley rat strain was favoured because of its clam temperament and ease of handling²¹⁹, which is significant for behavioural analyses. A male-only population was used in this study to limit the number of probable confounding variables caused by hormonal fluctuations in female populations. The subjects were housed conventionally with sizes laid down per the South African National Standards (SANS) 10386:2008 recommendations. Subjects were provided with standard irradiated "Epol" rat pellets and municipal water ad libitum. The subjects were housed in pairs per cage and

autoclaved pinewood shavings were used as a bedding material, along with white facial tissue paper for enrichment as photographically indicated in **figure 8 (E-F)**. The room temperature was maintained at 23°C (\pm 2°C) with a relative humidity of 50% (\pm 20%) and a 12-hour night/dark cycle throughout the study.



Figure 8: Facilities of the CAS unit that were used during this project.

(A-D). The subjects were housed in pairs inside cages containing pinewood shavings and an object for the subjects to play with (E-F).

The subjects were given a seven-day habituation period prior to the start of the experimental period where the first dose of LPS and/or PBS was administered. The experimental period

extended for 10 days. Thus, the total housing period of the subjects was 19 days. Trials to assess rodent behavioural patterns were carried out at the University of the Witwatersrand-health sciences campus, CAS laboratory. Histological analyses were conducted at the University of Pretoria (Lynnwood Rd, Hatfield, Pretoria, 0002), Haematology and bone analysis laboratory-Natural Sciences 2 building. The experimental procedures followed during this study are indicated in **figure 9**.



Figure 9: An overview of the procedures that were conducted throughout this study. During the sevenday acclimatization period, subjects were acclimatized to handling and the environment. Next, subjects were administered 0.1M LPS and/or 0.1M PBS via subcutaneous injection for ten days (experimental period). During this 10-day experimental period, Manuka honey was introduced in the intervention groups (PBS+H group and LPS+H group) on day 11 via oral gavage at a volume of 0.5 ml/kg, diluted at 50% v/v with distilled water to enable comfortable consistency. Behavioural analyses were performed between day 17-19. Day 20 was reserved for termination and sample preparation.

The Manuka honey, acquired from Advancis Medical, has no additives and is both filtered and sterilized thus limiting the cofounding effects of possible contaminants. Manuka honey is a medical grade honey sourced from New Zealand that shows potent anti-inflammatory and antioxidant effects as well as enhances immunity by stimulating white blood cells and promotes a good intestinal flora balance. As a result of the filtration process of the medical grade honey there is no variability between samples and thus pooling is not necessary. Day 20 was reserved for termination and sample preparation. Before termination, each animal was weighed, and the core body temperature was measured using a digital thermometer. Isofor[®] was used as an inhalation anaesthetic inside the gas chamber which each animal was placed in for 30 seconds. Next, cardiac puncture was performed to euthanize the animals as well as to enable blood collection into applicable Vacutainer blood collection tubes. Blood samples were used for a separate study that took place in conjunction with this study. Ethical approval was obtained for the use of blood samples- Protocol number: 171/2020. This was followed by perfusion with cold sterile saline solution and decapitation by guillotine. Lastly, the brain was carefully extracted from the skull. Each brain was separated into the left and right hemispheres. The right hemisphere was used for biochemical analyses, while the left hemisphere was used for confocal microscopy. **Figure 10** summarizes the process of sample collection.



Figure 10: A summary showing the sequence of events followed to prepare hippocampal tissue samples for biochemical and histological assays after the subjects were terminated.

3.4 Statistics

This study and all experimental procedural details were discussed with a statistician from the South African Medical Research Council Biostatistics Unit and a letter granting statistical support was been provided (addendum 5). The specific statistical tests and analyses employed per experiment can be found in the method and materials section of their respective following chapters.

Chapter 4: Behavioural analyses

4.1 Chapter Objectives

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This chapter will address the methods and results pertinent to the following objectives:

- The effects of systemically induced neuroinflammation on the short-term spatial working memory of Sprague-Dawley rats using the spatial recognition two-trial Y-maze test.
- The effects of systemically induced neuroinflammation on learning capacity and spontaneous memory of Sprague-Dawley rats using the novel object recognition test.
- The effects of systemically induced neuroinflammation on exploratory behaviour, anxiety, and locomotor activity of Sprague-Dawley rats using the open field test.

4.2 Introduction

Due to the slow-paced progression of dementia and the prolonged pre-clinical period before a provisional diagnosis, researchers are often forced to look at alternative ways to understand the aetiology of the disease while developing new treatment strategies.²¹⁸ Animal models are instrumental resources to better understand the biological mechanisms underlying dementia.²¹⁸ Behavioural assessments that reflect aspects of cognitive, social, and locomotor ability pave the way for advances in novel therapies, such as the possible effects of honey in the alleviation of neuroinflammation.²¹⁸ **Table 2** lists various behavioural tests used to assess specific features of dementia-like pathology in rodent models *in vivo*.

Table 2: A summary of various behavioural tests conducted on dementia-type rodent populations.^{255,256}

TEST	FUNCTION TESTED		
Open field test	Locomotor activity, anxiety, and habituation		
Morris water maze test	Spatial learning and memory		
Fear conditioning	Emotional and contextual memory		
Radial arm water maze test	Spatial learning (episodic-like) and working memory		
Forced alternation Y maze	Working memory and exploratory behaviour		
T maze	Spatial and working memory		
Novel object recognition test	Recognition memory		
Spontaneous alternation Y maze	Spatial working memory, reference working memory, and habituation		

Given that neurological damage is coupled with memory impairment, spatial or working memory is often affected.²²⁰ The Y-maze paradigm provides means to examine cognition and navigational schemes, something that is of significance in identifying brain mechanisms and possible remedial treatments.²²⁰ In addition, studies have shown that the NOR task is a robust and sensitive assay for assessing the cognitive-dysfunction activity of compounds such as LPS.²²¹ As this task allows to assess non-spatial memory that is devoid of emotional and learning factors in rodents.²²² Lastly, the open field test has been recognized as a straightforward assay to evaluate anxiety-like behaviour and locomotor impairments in rodents.²²⁰ It's basic protocol does not require subjects to undergo prior training or pre-conditioning. Reports by Fields, C indicate that a rise in enteric load of LPS amplifies anxiety-like behaviour across both sexes.²²³ Altogether, these assays were chosen as most appropriate to meet the objectives of this chapter.

4.2.1) Spatial recognition two-trial Y-maze test

The spatial recognition two-trial Y-maze test is a behavioural test that examines learning ability, spatial working memory, and reference working memory in rodents.^{218,219} This test is based on the notion that rodents are naturally explorative animals that show a willingness to explore new or strange environments.²¹⁹ Active engagement between multiple areas of the brain such as the hippocampus, prefrontal cortex, and basal forebrain is essential to execute

such activities^{.220} Therefore, it is presumed that rodents without impaired prefrontal functions will recall previously explored arms and show more interest in the unvisited arms by entering more often.²¹⁹ Findings by Deacon, R.M *et al.*²²¹ (2006) showed that animals with hippocampal lesions tend to display a side preference and poorer performance during this task. Y-maze paradigms are also used in disease and transgenic rodent models to assess the effects of novel chemical agents and treatments on cognitive performance and degree of cognitive impairment.²²⁰

4.2.2) Novel Object Recognition (NOR) Test

The NOR test is an exploratory test based on simple visual recall behaviour. It is used to evaluate different stages of learning (i.e., encoding, retention, or retrieval), object recognition, and memory (i.e., short-term memory or long-term memory) in murine models.²²² It has been successfully applied to transgenic rodent models, lesion studies as well as neurodegenerative models.²²³

4.2.3) Open Field Test

The open-field is an exploratory task used to assess anxiety-like behaviour and locomotor activity in murine models.^{224,225} This assay is centered around a parameter called "thigmotaxis", whereby, animals experiencing high levels of anxiety show a tendency to stay within close proximity to the walls of the arena.²²⁰

4.3 Methods and Materials

Behavioural analyses were conducted over a three-day period at the CAS unit, University of the Witswatersrand. The conditions of testing in the laboratory such as temperature, noise level, light intensity, and humidity were monitored and maintained throughout. With each assessment conducted, the apparatus was cleaned with 70% ethanol and allowed to dry completely between each trial run, to remove any olfactory cues.

For each assay conducted, an overhead camera was placed directly above the apparatus to video track the trials for each assessment. ANY-maze Video Tracking System (Version 4.2; Stoelting Co., Wood Dale, II) was used to obtain and collate the results from the behavioural assessments for subsequent analysis.

The weight of each rat was measured and recorded daily throughout the experimental period. The welfare of each rat was also monitored and recorded daily on a welfare monitoring sheet (addendum 4) . Characteristics evaluated include movement, nourishment, breathing, and changes in behaviour during handling and facial expressions. The Rat Grimace scale, illustrated in **figure 11**, uses changes in facial expressions to quantify the degree of pain and discomfort in rats.²²⁴ During the experimental period, pain and discomfort were monitored daily using this grading scale. The four parameters of interest were orbital tightening, nose/cheek flattening, ear changes, and whisker changes. If an "Obviously present" score was observed for any of the parameters, the subject would have been excluded from the study.



Figure 11:The Rat Grimace Scale uses various facial expressions to quantify signs of discomfort in rats during laboratory experiments.²²⁴

Figure 12 gives an overview of the behavioural assays conducted during the experimental period. It was recommended to conduct testing over three days to ensure that the subjects did not experience too much stress or get too tired.



Figure 12: An overview of the study, highlighting the behavioural assays conducted before termination. Open-field analysis was performed on day 17, whilst the two-trial Y-maze test was completed on day 18, and the NOR test was conducted on day 19.

4.3.1) Behavioural analyses

4.3.1.1) Spatial recognition two-trial Y-maze assay

Testing occurred in two trials inside a Y-shaped chamber of three arms distanced at an angle of 120° relative to each other. The subject was placed at the central mid-zone area where all three arms meet, and it is allowed to explore the maze freely. Entry into an arm was characterized by the placement of all four paws within the boundaries of an arm. The time spent in each arm was also noteworthy. The laboratory temperature, noise level, light intensity, and humidity were controlled and maintained for all subjects during all experimental procedures. Light intensity in the testing room was dimmed to stimulate exploratory behaviour.

Apparatus: A Y-shaped maze made from Plexiglas, with three arms equally distributed at an angle of 120°. The arms were classified as follows; long (L), familiar (F), and novel (N). Each arm had a length of 63 cm, 54.9 cm, and 59.4 cm, respectively. All three arms had a width of 10 cm and a height of 15 cm to prevent escape from the maze. In **figure 13 (C)**, an overhead camera was prepared above the maze to video track and record memory and learning behaviour for subsequent analyses.



Figure 13: A graphic representation of the Y-maze apparatus (A) and the test subject inside the arena (B). An overhead camera with a video recording system was used to track behavioural activity (C). Image adapted from Prieur, E *et al.* (2019).²²⁵

Exploration trial: Plexiglas intercepted the novel arm of the maze, leaving only two arms available to explore. Subjects were brought in their housing cage into the laboratory testing room and allowed to acclimatize for one and a half hours in the absence of individuals and the testing apparatus. The rat was then placed in the central mid-zone of the Y-maze and allowed to freely explore the two available arms for two minutes and 30 seconds. The subjects were given a four-hour inter-trial interval before the testing trial of the procedure was conducted.

Testing (recognition) trial: The Plexiglas intercepting the novel arm was removed, leaving all three arms available for exploration. Each rat was brought into the laboratory testing room and allowed to acclimatize for ten minutes. The rat was placed in the mid-zone of the Y-maze and allowed to freely explore all arms of the maze for two minutes and 30 seconds.

Given that this test relies on novelty seeking and the innate tendency of rodents to explore their surroundings, it is anticipated that they will spend more time in the previously inaccessible (novel) arm than the two familiar arms. Therefore, the ability to differentiate the novel arm from familiar ones is used as a marker of spatial recognition memory.²²⁶

For consistency and analysis, the arms of the y-maze were classified as shown in figure 14.



Figure 14: A diagram showing the classification of the arms of the y-maze apparatus.

Once the exploration and testing (recognition) trial were completed, the following parameters were considered for each arm of the maze:

- i. Number of head entries
- ii. Time spent within the arm
- iii. The average speed
- iv. The average number of visits
- v. Time mobile
- vi. Time immobile

4.3.2) Novel Object Recognition Test (NORT)

The assessment involved two trials; the training trial and the test trial.²²⁷

Apparatus: An open arena made from non-porous plastic with the following dimensions: a 65 cm (length) 60 cm (breadth) x 20 cm (height).

Training trial: This was conducted on day 17 of the experimental period. Each subject was brought in its housing cage into the laboratory testing room and allowed to acclimatize for ten minutes. The subject was placed at the centre of the arena and allowed to acclimatize for 30 seconds. Next, it was briefly removed from the arena while two identical objects (X + X), were placed inside the arena at opposite ends of each other (i.e., West and East). The subject was placed in between the two objects with its head facing the interior wall of the arena and

allowed to explore the identical objects for two minutes. Exploratory activity was assessed and used to evaluate memory retention and recall. Exploration was defined as the sweeping or sniffing of an object, with the subjects' nose pointed up towards the object, within a distance of 3 cm or less from the object. Finally, the subject was removed from the arena and returned to its housing cage.

Object selection: The objects were smaller than the subjects. They were made from soft foam and bright contrasting colours. They differed slightly in shape to stimulate exploratory behaviour.

Testing trial: This was performed on day 18 of the experimental period. One of the familiar objects (X) used during the training trial together with a novel object (Y) was placed inside the open arena, at opposite ends of one other. The subject was brought in its holding cage into the laboratory testing room and immediately placed at the centre of the objects (X + Y) inside the arena with its head facing the interior wall of the arena. The rat was allowed to freely explore the environment and the familiar object (X) vs. novel object (Y) for two minutes, as seen in **figure 15**. Exploratory activity was assessed and used to determine memory recall. Exploration was defined as the sweeping or sniffing of the object with the rats' nose pointed up towards the object, within 3 cm or less from the object (X) and the novel object (Y) and will spend more time exploring the novel object using induced preference and memory recall functions.



Figure 15: Testing trial of the novel object recognition test. The familiar object (A) and novel object (B) were placed at opposite ends. The test subject was allowed to explore each object for 2 minutes freely.

4.3.3) Open Field Analysis

Day 19 was reserved for the open-field assessment.

Apparatus: A wall-enclosed Plexiglas chamber with the dimensions: 1 m (length) x 1 m (breadth) and a height of 0.5 m to prevent the rat from escaping the enclosure. Using ANY-maze Video Tracking System, the base was divided into 25 smaller squares of 20 cm x 20 cm, which comprised the outermost area. The outmost area was further divided into an innermost area of 85 cm x 85 cm. **Figure 16 (A)** illustrates the open-field arena during the assessment, whereas **Figure 16 (B)** is a visual representation of the apparatus as displayed on the video tracking software.



Figure 16: An illustration of the open-field arena with the subject inside (A). The surface of the open-field arena as depicted on Any-maze software (B). The outermost area comprises the entire 1 m x 1 m perimeter, and the innermost area includes the shaded region.

On the floor, it was positioned so that an overhead luminous light was the brightest on the innermost region and dim in the outermost area. This was built on the notion that rodents are fearful of large, bright spaces and are likely to seek shelter at the peripheral walls of the open field until they are comfortable and begins to explore, moving towards more brilliant light at the centre of the open field.

Testing trial: Habituation was not required for this assay. The subject was brought into the laboratory testing room and allowed to acclimatize for ten minutes without any individuals. The subject was gently placed inside the open-field, at the outermost area of dim light. A five-minute period was given for exploration. Exploratory activity was recorded as the number of lines crossed by the subject in five minutes. A line crossing was defined as all four paws crossing a boundary line that separates adjacent quadrants. It was expected that rats with less anxiety or unimpaired mobility would show more exploratory behaviour, thereby crossing more lines than those who were more anxious or with mobility issues. Behaviour was recorded from the time of subject placement into the field until the end of the trial.

4.4 Results

All statistical tests were completed using GraphPad[™] Prism (version 9.2.0) for Windows (GraphPad[™] Software, San Diego, California USA). The D'Agostino-Pearson normality test was used to establish that the dataset was normally distributed. Data was analysed using one-way analysis of variance (ANOVA). Post-hoc checks with Dunnett's multiple comparisons test were used to compare the three experimental groups to the control group. The results were represented as mean ± standard error of mean (SEM). Error bars indicate SEM. The p-value was set at p <0.05, with a 95% confidence interval.

To characterize the effects of LPS-induced neuroinflammation on spatial working memory, learning capacity, and exploratory behaviour, the control group (PBS only) was compared to the three experimental groups (LPS only vs. LPS + H vs. PBS + H) using the Y-maze test, NORT and open-field test, respectively. Close examination of the mean scores per group reveal non-significant differences between them, although notable differences in performance between the groups were observed and thus discussed.

4.4.1) Spatial recognition Y-maze Analysis

Memory and learning impairments in the long arm of the y-maze were evaluated by analyzing the number of head entries into the arm during the exploration and testing/recognition trial as shown in **table 3** and **figure 17**. The number of head entries were not significantly different across the groups. Data from the testing trials were compared to exploration trial. To determine the performance of subjects, the control group was compared against the experimental groups.

Table 3: A summary of the mean scores and p-values obtained when evaluating the average number of entries into the long arm of the Y-maze chamber during the exploration trial (Trial 1/2), testing trial (Trial 2/2), in comparison to the novel arm (Trial 2/2-Novel arm).

Summary of the descriptive statistics of the number of head entries-long arm						
Group	Trial	Mean	Adjusted P-value	Significant?		
PBS	Trial 1/Z	5.1	12			
PBS vs. LPS	100101000000	4.5	0.6086	No		
PBS vs. LPS+H		4.8	0.9184	No		
PBS vs. PBS+H		4.7	0.8329	No		
PBS	Trial 2/2	4.2				
PBS vs. LPS		2.8	0.2009	No		
PBS vs. LPS+H		3.5	0.7064	No		
PBS vs. PBS+H		3.5	0.7064	No		
PBS	Trial 2/2 (Novel arm)	7.5	-			
PBS vs. LPS		8	08794	No		
PBS vs. LPS+H		7.4	0.9987	No		
PBS vs. PBS+H		7	0.8794	No		



Figure 17 (A-C): Bar graph plot with mean and standard error of mean (SEM) for the number of head entries into the long and novel arm of the Y-maze chamber. Trial 1/2-Long arm shows the average number head entries into the long arm during the exploration trial (A). Trial 2/2-Long arm shows the average number head entries into the long arm during the testing/recognition trial (B). Trial 2/2-Novel arm shows the average head entries into the novel arm during the testing/recognition trial (C).

Next, the time spent in the long arm was analyzed and presented in **table 4** and **figure 18**. Results show that the time spent in the long arm during trial 2/2 was not significantly different than trial 1/2. And the time spent in trial 2/2 in the novel arm was not significantly different than trial 2/2 in the long arm.

Table 4: A summary of the mean scores and p-values obtained when evaluating the average time spent in the long arm of the Y-maze chamber during the exploration trial (Trial 1/2), testing/recognition trial (Trial 2/2), in comparison to the novel arm (Trial 2/2-Novel arm).

Summary of the descriptive statistics of the time spent-long arm						
Group	Trial	Mean	Adjusted P-value	Significant7		
PBS	Trial 1/2	45.23				
PBS vs. LPS		40.81	0.5729	No		
PBS vs. LPS+H		43.12	0.9177	No		
PBS vs. PBS+H		42.56	0.8512	No		
PBS	Trial 2/2	28.17	-			
PBS vs. LPS		24.56	0.7458	No		
PBS vs. LPS+H		31.25	0.8217	No		
PBS vs. PBS+H		21.35	0.2889	No		
PBS	Trial 2/2 (Novel arm)	50.28	-	-		
PBS vs. LPS		50.34	>0.9999	No		
PBS vs. LPS+H		42.31	0.1664	No		
PBS vs. PBS+H		44.69	0.4227	No		



Figure 18 (D-F): Bar graph plot with mean \pm SEM for the average time spent in the long and novel arm of the Y-maze chamber. Trial 1/2-Long arm shows the average time spent in the long arm during the exploration trial (D). Trial 2/2-Long arm shows the average time spent in the long arm during the testing/recognition trial (E). Trial 2/2-Novel arm shows the average time spent in the novel arm during the testing/recognition trial (F).

Then, the average speed scored within the long arm was assessed and presented in **figure 19** and **table 5**. In trial 2/2 of the novel arm, a significant difference (p<0.0347) was noted

when the control group was compared to the PBS + H group . This suggests that the PBS + H treated group may have navigated the novel arm of the maze at higher speeds as a result of honey intake.

Table 5: A summary of the mean scores and p-values obtained when evaluating the average speed of the subject in the long arm of the Y-maze chamber during the exploration trial (Trial 1/2), testing/recognition trial (Trial 2/2), in comparison to the novel arm (Trial 2/2-Novel arm).

Summary of the descriptive statistics of the average speed- long arm						
Group	Trial	Mean	Adjusted P-value	Significant ?		
PBS	Trial 1/2	0.07220	-	1		
PBS vs. LPS	020000000000	0.07110	0.9973	No		
PBS vs. LPS+H		0.07000	0.9797	No		
PBS vs. PBS+H		0.06610	0.7237	No		
PBS	Trial 2/2	0.08440	-			
PBS vs. LPS		0.07520	0.7589	No		
PBS vs. LPS+H		0.06510	0.2314	No		
PBS vs. PBS+H		0.07320	0.6408	No		
PBS	Trial 2/2 (Novel arm)	0.06290	-	5		
PBS vs. LPS		0.07030	0.3504	No		
PBS vs. LPS+H		0.07520	0.0567	No		
PBS vs. PBS+H		0.07630	0.0347	Yes *		



Figure 19 (G-I): Bar graph plot with mean \pm SEM for the average speed of the subject in the long and novel arm of the Y-maze chamber. Trial 1/2-Long arm shows the average speed of the subject in the long arm during the exploration trial (G). Trial 2/2-Long arm shows the average speed of the subject

in the long arm during the testing/recognition trial (H). Trial 2/2-Novel arm shows the average speed of the subject in the novel arm during the testing/recognition trial (I).

Next, the average number of visits into the long arm, as shown in **table 6** and **figure 20**, was recorded and analysis revealed that the number of visits did not significantly differ during the three trials, or between the sample groups.

Table 6: A summary of the mean scores and p-values obtained when evaluating the average number of visits into the long arm of the Y-maze chamber during the exploration trial (Trial 1/2), testing/recognition trial (Trial 2/2), in comparison to the novel arm (Trial 2/2-Novel arm).

Summary of the descriptive statistics of the average number of visita-long arm						
Group	Trial	Mean	Adjusted P- value	Significant?		
PBS	Trial 1/2	9.7	-			
PIBS VS. LPS		10.06	0.9814	No		
PIBS vs. LPS+H		10.30	0.9238	No		
PBS vs. PBS+H		10.37	0.8987	No		
PIBS	Trial 2/2	8.650	-	-		
PIBS VS. LPS		9.730	0.7573	No		
PIBS vs. LPS+H		11.03	0.1967	No		
PBS vs. PBS+H		9.090	0.9759	No		
PBS	Trial 2/2 (Novel arm)	11.81		•		
PIBS vs. LPS		11.35	0.9753	No		
PIBS vs. LPS+H		9.650	0.2893	No		
PBS vs. PBS+H		9.390	0.2095	No		



K



J





L
Figure 20 (J-L): Bar graph plot with mean ± SEM for the average number of visits into the long and novel arm of the Y-maze chamber. Trial 1/2-Long arm shows the average speed of the subject in the long arm during the exploration trial (J). Trial 2/2-Long arm shows the average speed of the subject in the long arm during the testing/recognition trial (K). Trial 2/2-Novel arm shows the average speed of the subject of the subject in the novel arm during the testing/recognition trial (K).

Lastly, the time spent in a state of motion and the time spent immobile in the long and novel arm was evaluated for inter-trial and inter-group comparison. This is shown in **table 7** and **table 8**, and graphically presented in **figure 21** and **figure 22**. **Table 7** shows significant differences (p<0.0440) and (p<0.0237) in the long arm during trial 2/2 was noted when the control group was compared to the LPS group and PBS + H group, respectively.

Table 7: A summary of the mean scores and p-values obtained when evaluating the time lapsed while the subject was mobile in the long arm of the Y-maze chamber during the exploration trial (Trial 1/2), testing/recognition trial (Trial 2/2), in comparison to the novel arm (Trial 2/2-Novel arm).

Summary of the descriptive statistics of the time mobile-long arm						
Group	Trial	Mean	Adjusted P-value	Significant		
PBS	Trial 1/2	19.12	-			
PBS vs. LPS		20.13	0.9403	No		
PBS vs. LPS+H		21.05	0.7144	No		
PBSvs. PBS+H		18.79	0.9976	No		
PBS	Trial 2/2	13.18	-	-		
PBS vs. LPS		8.780	0.0440	Yes *		
PBS vs. LPS+H		10.88	0.4251	No		
PBS vs. PBS+H		8.320	0.0237	Yes *		
PBS	Trial 2/2 (Novel arm)	30.02	-	(c.e.)		
PBS vs. LPS		33.37	0.7187	No		
PBSvs. LPS+H		29.81	>0.9999	No		
PBSvs. PBS+H		31.19	0.9810	No		



Figure 21 (M-O): Bar graph plot with mean ± SEM for time lapsed while the subject was mobile in the long and novel arm of the Y-maze chamber. Trial 1/2-Long arm shows the time mobile in the long arm during the exploration trial (M). Trial 2/2-Long arm shows the time mobile in the long arm during the testing/recognition trial (N). Trial 2/2-Novel arm shows the time mobile in the novel arm during the testing/recognition trial (O).

Table 8 shows that significant differences (p<0.0183) and (p<0.0449) in the novel arm during the testing trial were present when the control group was compared to the LPS + H group and PBS + H group, respectively. This is supported by findings in **table 4**.

Table 8: A summary of the mean scores and p-values obtained when evaluating the time lapsed while the subject was immobile in the long arm of the Y-maze chamber during the exploration trial (Trial 1/2), testing/recognition trial (Trial 2/2), in comparison to the novel arm (Trial 2/2-Novel arm).



Figure 22 (P-R): Bar graph plot with mean ± SEM for time lapsed while the subject was immobile in the long and novel arm of the Y-maze chamber. Trial 1/2-Long arm shows the time mobile in the long arm during the exploration trial (P). Trial 2/2-Long arm shows the time mobile in the long arm during the testing/recognition trial (Q). Trial 2/2-Novel arm shows the time mobile in the novel arm during the testing/recognition trial (R).

Memory and learning deficits in the familiar arm of the y-maze were first evaluated by analyzing the number of head entries into the arm during the exploration and

testing/recognition trial as shown in **table 9** and **figure 23**. The number of head entries were not significantly different between the groups.

Table 9: A summary of the mean scores and p-values obtained when evaluating the average number of entries into the familiar arm of the Y-maze chamber during the exploration trial (Trial 1/2), testing/recognition trial (Trial 2/2), in comparison to the novel arm (Trial 2/2-Novel arm).

Summary of the descriptive statistics of the number of head entries-familiar arm						
Group	Trial	Mean	Adjusted P-value	Significant?		
PBS	Trial 1/2	5	<u> </u>			
PBSvs. LPS		5.1	0.9941	No		
PBSvs. LPS+H		4.3	0.3636	No		
PBSvs. PBS+H		5.1	0.9941	No		
PBS	Trial 2/2	2.8	-	-		
PBSvs. LPS		3.5	0.3227	No		
PBS vs. LPS+H		3.1	0.8585	No		
PBSvs. PBS+H		3.5	0.3227	No		
PBS	Trial 2/2 (Novel arm)	7.5	-	-		
PBS vs. LPS		8	08794	No		
PBS vs. LPS+H		7.4	0.9987	No		
PBS vs. PBS+H		7	0.8794	No		



Figure 23 (A-C): Bar graph plot with mean ± SEM for the number of head entries into the familiar and novel arm of the Y-maze chamber. Trial 1/2-Familiar arm shows the average number head entries into the familiar arm during the exploration trial (A). Trial 2/2-Familiar arm shows the average number head entries into the familiar arm during the testing/recognition trial (B). Trial 2/2-Novel arm shows the average head entries into the novel arm during the testing/recognition trial (C).

Next, the time spent in the familiar arm was analyzed and presented in **table 10** and **figure 24**. A significant difference (p<0.0072) during the trial 2/2 in the familiar arm was noted when the PBS group was compared to the PBS + H group.

Table 10: A summary of the mean scores and p-values obtained when evaluating the average time spent in the familiar arm of the Y-maze chamber during the exploration trial (Trial 1/2), testing/recognition trial (Trial 2/2), in comparison to the novel arm (Trial 2/2-Novel arm).

Summary of the descriptive statistics of the time spent- familiar arm							
Group	Trial	Mean	Adjusted P-value	Significant?			
PBS	Trial 1/2	45.23	÷.	(H)			
PBSvs. LPS		54.11	0.1214	No			
PBSvs. LPS+H		52.39	0.2529	No			
PBS vs. PBS+H		55.80	0.0532	No			
PBS	Trial 2/2	24.54	2	8 3 - 525			
PBSvs. LPS		28.68	0.4728	No			
PBS vs. LPS+H		28.42	0.5232	No			
PBS vs. PBS+H		35.40	0.0072	Yes *			
PBS	Trial 2/2 (Novel arm)	50.28	×				
PIBS vs. LPS		50.34	>0.9999	No			
PBSvs, LPS+H		42.31	0.1664	No			
PBSvs. PBS+H		44.69	0.4227	No			



Figure 24 (D-F): Bar graph plot with mean ± SEM for the average time spent in the familiar and novel arm of the Y-maze chamber. Trial 1/2-Familiar arm shows the average time spent in the familiar arm during the exploration trial (D). Trial 2/2-Familiar arm shows the average time spent in the familiar

arm during the testing/recognition trial (E). Trial 2/2-Novel arm shows the average time spent in the novel arm during the testing/recognition trial (F).

Then, the average speed scored within the familiar arm was assessed and presented in figure 25 and table 11. A significant difference (p<0.0347) during trial 2/2 in the novel arm was noted when the PBS group was compared to the PBS + H group.

Table 11: A summary of the mean scores and p-values obtained when evaluating the average speed of the subject in the familiar arm of the Y-maze chamber during the exploration trial (Trial 1/2), testing/recognition trial (Trial 2/2), in comparison to the novel arm (Trial 2/2-Novel arm).



Figure 25 (G-I): Bar graph plot with mean ± SEM for the average speed of the subject in the familiar and novel arm of the Y-maze chamber. Trial 1/2-Familiar arm shows the average speed of the subject

G

Awerage speed (m/s)

in the familiar arm during the exploration trial (G). Trial 2/2-Familiar arm shows the average speed of the subject in the familiar arm during the testing/recognition trial (H). Trial 2/2-Novel arm shows the average speed of the subject in the novel arm during the testing/recognition trial (I).

The average number of visits into the familiar arm, as shown in **table 12** and **figure 26**, was recorded and compared between the exploration and testing trial. The number of visits were not significantly different between the groups.

Table 12: A summary of the mean scores and p-values obtained when evaluating the average number of visits into the familiar arm of the Y-maze chamber during the exploration trial (Trial 1/2), testing/recognition trial (Trial 2/2), in comparison to the novel arm (Trial 2/2-Novel arm).

Summary of the descriptive statistics of the average number of visits-familiar arm						
Group	Trial	Mean	Adjusted P- value	Significant?		
PBS	Trial 1/2	10.21		-		
PBS vs. LPS		10.27	>0.9999	No		
PBSvs. LPS+H		12.08	0.5826	No		
PRSvs. PRS+H		12.21	0.5327	No		
PBS	Trial 2/2	9.220	72 ()	-		
PBS VS. LPS		10.85	0.6720	No		
PBS vs. LPS+H		10.52	0.7993	No		
PBS vs. PBS+H		11.27	0.5129	No		
PBS	Trial 2/2 (Novel arm)	11.81	-	-		
PBS vs. LPS		11.35	0.9753	No		
PBS vs. LPS+H		9.650	0.2893	No		
PBSvs. PBS+H		9.390	0.2095	No		

Thai 1/2- Familiar arm: Average number of visits Trial 2/2- Familiar arm: Average number of visits Trial 2/2- Novel arm: Average number of visits

κ



Figure 26 (J-L): Bar graph plot with mean ± SEM for the average number of visits into the familiar and novel arm of the Y-maze chamber. Trial 1/2-Familiar arm shows the average speed of the subject in the familiar arm during the exploration trial (J). Trial 2/2-Familiar arm shows the average speed of the subject in the familiar arm during the testing/recognition trial (K). Trial 2/2-Novel arm shows the average speed of the subject in the novel arm during the testing/recognition trial (L).

Lastly, the time spent in a state of motion and the time spent immobile was evaluated for inter-trial and inter-group comparison. This is shown in **table 13** and **table 14**, and graphically presented in **figure 27** and **figure 28**. The time mobile in the familiar arm and novel arm were not significantly different between the groups.

Table 13: A summary of the mean scores and p-values obtained when evaluating the time lapsed while the subject was mobile in the familiar arm of the Y-maze chamber during the exploration trial (Trial 1/2), testing/recognition trial (Trial 2/2), in comparison to the novel arm (Trial 2/2-Novel arm).

Summary of the descriptive statistics of the time mobile- familiar arm							
Group	Trial	Trial Mean		Significant			
P85	Trial 1/2	23.26	-	1			
PBS vs. LPS		30.91	0.1215	No			
PBS vs. LPS+H		28.29	0.4083	No			
PBS vs. PBS+H		29.52	0.2426	No			
PBS	Trial 2/2	14.33		-			
PBS vs. LPS		14.84	0.9869	No			
PBS vs. LPS+H		15.21	0.9394	No			
PBS vs. PBS+H		16.87	0.4147	No			
PBS	Trial 2/2 (Novel arm)	30.02	A .				
PBS vs. LPS		33.37	0.7187	No			
PBS vs. LPS+H		29.81	>0.9999	No			
PBS vs. PBS+H		31.19	0.9810	No			



Figure 27 (M-O): Bar graph plot with mean ± SEM for time lapsed while the subject was mobile in the familiar and novel arm of the Y-maze chamber. Trial 1/2-Familiar arm shows the time mobile in the familiar arm during the exploration trial (M). Trial 2/2-Familiar arm shows the time mobile in the familiar arm during the testing/recognition trial (N). Trial 2/2-Novel arm shows the time mobile in the novel arm during the testing/recognition trial (O).

However, when time immobile was analyzed in **table 14** Significant differences in the familiar arm was noted when the PBS group was compared to the PBS + H (p<0.0132) group during trial 2/2. Significant differences were also observed in the novel arm when the PBS group was compared to the LPS + H (p<0.0183) and PBS + H (p<0.0449) groups.

Table 14: A summary of the mean scores and p-values obtained when evaluating the time laps	sed
while the subject was immobile in the familiar arm of the Y-maze chamber during the exploration t	trial
(Trial 1/2), testing/recognition trial (Trial 2/2), in comparison to the novel arm (Trial 2/2-Novel arm	ı).

Summary of the descriptive statistics of the time immobile- familiar arm							
Group	Trial	Mean	Adjusted P- value	Significant?			
PBS	Trial 1/2	21.97	-	(a)			
PBS vs. LPS		23.19	0.9888	No			
PBS vs. LPS+H		24.07	0.9481	No			
PBS vs. PBS+H		26.32	0.6976	No			
PBS	Trial 2/2	10.21					
PBS vs. LPS		13.83	0.4268	No			
PBS vs. LPS+H		13.24	0.5642	No			
PBS vs. PBS+H		18.52	0.0132	Yets *			
PBS	Trial 2/2 (Noval arm)	20.26					
PBS vs. LPS		16.97	0.4828	No			
PBS vs. LPS+H		12.50	0.0183	Yes *			
PBS vs. PBS+H		13.51	0.0449	Yes 🕸			



Figure 28 (P-R): Bar graph plot with mean ± SEM for time lapsed while the subject was immobile in the familiar and novel arm of the Y-maze chamber. Trial 1/2-Familiar arm shows the time mobile in the familiar arm during the exploration trial (P). Trial 2/2-Familiar arm shows the time mobile in the familiar arm during the testing/recognition trial (Q). Trial 2/2-Novel arm shows the time mobile in the novel arm during the testing/recognition trial (R).

4.4.2) Novel Object Recognition Task

After the training and testing trial were complete, the following parameters were considered for each block (object) A and block B:

- i. Number of entries into the perimeter of the block.
- ii. Mean visits.
- iii. Time spent (overall time spent at/close to object).
- iv. Time immobile (time spent while stationary observing/exploring object).
- v. The first block entered.

For the training trial, both Block A and Block B (new block) held identical objects. For the testing trial, Block B (new block) held the novel object, whilst Block A remained the same.

For data analysis, the three experimental groups (LPS only vs. LPS and honey vs. PBS and honey) were compared to the control group (PBS only).

Firstly, the number of head entries into block A and block B during the training phase were analyzed and presented in **table 15 and figure 29.** As shown, The number of head entries was not significantly different across the groups.

Summary of the descriptive statistics of the number of entries during the training phase							
Group	Trial	Mean	Adjusted P- value	Significant			
PBS	Training Phase- Block A	5.5	1	0			
PBSvs. LPS	- 11	6.8	0.5663	No			
P85vs. LPS+H	8	4.7	0.8412	No			
PBSvs. PBS+H		6.1	0.9229	No			
PBS	Training Phase- Block B	5.2	847	19			
PBSvs. LPS	8	6.8	0.5655	No			
PBS vs. LPS+H		5.6	0.9859	No			
PBSvs. PBS+H		5.3	0.9997	No			

Table 15: A summary of the mean scores and p-values obtained when evaluating the number of entries made within the perimeter of Block A and Block B, during the training trial.



JAS TH

Group

A

Number of entries

2

3





Training Phase-Block B (new block): Entries



Figure 29 (A-B): Bar graph plot with mean \pm SEM for the average number of entries made into the perimeter of Block A and Block B during the training trial of the NORT. The number of entries in Block A (A). The number of entries in Block B (B).

Next, the time spent immobile in block A and block B during the training phase was analyzed and presented in **table 16 and figure 30.** The time immobile was not significant between the groups.

Table	16: A	summa	ry of	the mea	an so	cores a	nd p-va	lues	obtained	whe	n eva	luating	time im	mo	bile (ti	me
spent	while	stationa	ry ex	kploring	the	object) within	the	perimeter	of	Block	A and	Block E	З, с	luring	the
trainin	g trial	l.														

Group	Trial	Mean	Adjusted P- value	Significant?	
PBS	Training Phase- Block A	1.590	-	7727	
PBS vs. LPS		1.570	>0.9999	No	
PBS vs. LPS+H	-	1.330	0.9925	No	
PBS vs. PBS+H		2.630	0.7115	No	
PBS	Training Phase- Block B	2.530		-	
PBS vs. LPS		1.380	0.7172	No	
PBS vs. LPS+H		4.450	0.3440	No	
PBS vs. PBS+H		0.9500	0.4972	No	



Figure 30 (C-D): Bar graph plot with mean \pm SEM for the time immobile (time spent while stationary exploring the object) within the perimeter of Block A and Block B, during the training trial of the NORT. Time immobile in Block A (C). Time immobile in Block B (D).

Next, the average number of visits into block A and block B during the training phase was analyzed and presented in **table 17 and figure 31**. The number of visits were not significantly different across the groups.

Table 17: A summary of the mean scores and p-values obtained when evaluating the average number of visits made within the perimeter of Block A and Block B, during the training trial.

Summary of the descriptive statistics of the average number of visits in each block during the training phase						
Group	Trial	Mean	Adjusted P- value	Significant?		
PBS	Training Phase- Block A	0.9700	20			
PBS vs. LPS		1.010	0.9974	No		
PBS vs. LPS+H	-2	1.360	0.3152	No		
PBS vs. PBS+H		1.290	0.4701	No		
PBS	Training Phase- Block B	1.340	-	-		
PBS vs. LPS		0.9000	0.4683	No		
PBS vs. LPS+H		2.080	0.1097	No		
PBS vs. PBS+H		1.430	0.9886	No		



Figure 31 (E-F): Bar graph plot with mean \pm SEM for the average number of visits made into the perimeter of Block A and Block B during the training trial of the NORT. The number of visits in Block A (E). The number of visits in Block B (F).

For the testing phase, a novel object was introduced into block B of the arena. To begin, the number of visits were analyzed during the testing/recognition phase of this assay. Data is presented in **table 18** and **figure 32**. A significant difference (p<0.0044) in the number of entries in block B was noted when the control group was compared to the LPS group during

the testing trial. This suggests that LPS treatment did not affect recognition memory in this trial since the LPS group entered the zone containing the novel object more often than the control group.

Table 18: A summary of the mean scores and p-values obtained when evaluating the number of entries made within the perimeter of Block A and Block B, during the testing trial

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Group	Trial	Mean	Adjusted P- value	Significant?				
PBS	Testing Phase- Block A	4.5	-	3				
PBSvs. LPS		6.9	0.1988	No				
PBSvs. LPS+H		3.8	0.9171	No				
PBSvs. PBS+H		5.8	0.6550	No				
PBS	Testing Phase- Block B	4.0		÷				
PBS vs. LPS		7.5	0.0044	Yes *				
PBS vs. LPS+H		4.3	0.9728	No				
PBSvs. PBS+H		5.8	0.4541	No				

A





Testing Phase-Block B (new block): Entries





Figure 32 (G-H): Bar graph plot with mean \pm SEM for the average number of entries made into the perimeter of Block A and Block B during the testing trial of the NORT. The number of entries in Block A (G). The number of entries in Block B (H).

Next, the time spent in block A and block B were analyzed during the testing phase and presented in **table 19** and **figure 33**. The time spent was not significantly different across the groups.

Table 19: A summary of the mean scores and p-values obtained when evaluating the average time spent within the perimeter of Block A and Block B, during the testing trial.

Group	Trial	Mean	Adjusted P-	Significant
PBS	Testing Phase- Block A	6.970		2
PBSvs. UPS		9.830	0.3162	No
PBSvs. LPS+H		4.330	0.3493	No
PBSvs. PBS+H		8.670	0.7005	No
PB5	Testing Phase- Block B	6.570	*	-
PBS vs. LPS		9.310	0.8046	No
PBSvs. LPS+H		5.030	0.9549	No
PBSvs. PBS+H	1	11.12	0.4819	No





Testing Phase-Block B (new block): Time



Figure 33 (C-D): Bar graph plot with mean \pm SEM for the average time spent within the perimeter of Block A and Block B during the testing trial of the NORT. The number of entries in Block A (C). The number of entries in Block B (D).

Then, the average number of visits block A and block B were analyzed during the testing phase and presented in **table 20** and **figure 34**. The time spent was not significantly different across the groups. The number of visits were not significantly different across the groups.

Table 20: A summary of the mean scores and p-values obtained when evaluating the average number of visits made within the perimeter of Block A and Block B, during the testing trial.

Summary of the descriptive statistics of the average number of visits in each block during the testing phase					
Group	Trial	Mean	Adjusted P- value	Significant?	
PBS	Testing Phase- Block A	1.710	12	31413 	
PBS vs. LPS		1.470	0.7148	No	
PBS vs. LPS+H		1.120	0.0955	No	
PBS vs. PBS+H		1.570	0.9204	No	
PBS	Testing Phase- Block B	1.520		-	
PBS vs. LPS		1.270	0.7811	No	
PBS vs. LPS+H		1.089	0.4332	No	
PBS vs. PBS+H		1.480	0.9986	No	

E



Testing Phase-Block A: Mean Visit







Figure 34 (E-F): Bar graph plot with mean \pm SEM for the average number of visits made into the perimeter of Block A and Block B during the testing trial of the NORT. The number of visits in Block A (E). The number of visits in Block B (F).

Also, the time spent immobile inside block A and block B were analyzed during the testing phase and presented in **table 21** and **figure 35**. The time immobile was not significantly different between the groups.

Table 21: A summary of the mean scores and p-values obtained when evaluating time immobile (time spent while stationary exploring the object) within the perimeter of Block A and Block B, during the testing trial.

Summary of the descriptive statistics of the time immobile in each block during the testing phase					
Group	Trial	Mean	Adjusted P-value	Significant?	
PBS	Testing Phase- Block A	3.150	8	-	
PBS vs. LPS		4.360	0.7691	No	
PRSss. 1PS+H		1.570	0.6062	No	
PBSvs. PBS+H		5.030	0.4756	No	
PES	Testing Phase- Block B	2.370	-		
PBSvs. LPS		3.190	0.9929	No	
PES vs. LPS+H		2.580	>0.9999	No	
PBSvs. PBS+H		7.180	0.4506	No	



Figure 35 (G-H): Bar graph plot with mean \pm SEM for the time immobile (time spent while stationary exploring the object) within the perimeter of Block A and Block B, during the testing trial of the NORT. Time immobile in Block A (G). Time immobile in Block B (H).

Lastly, the first zone entered (block explored first) during the training and testing trial was also noted and illustrated in **figure 36**. It was predicted that the novel (Block B) object would be visited first, before the familiar (Block A) object during the testing trial. Results of the training phase indicate that all three experimental groups explored block A first, while the control group explored block B first. However, this was contrasted during the testing phase. The LPS-treated group and control group

explored block B, containing the novel object, first. The LPS + H group showed no preference in the object first explored. And, the PBS + H group explored block A, containing the familiar object, first.



Figure 36 (A-B): A vertically stacked bar graph plot showing the proportion of the first zone entered (object explored first) upon commencement of the NORT trial. First zone entered during the training trial (A). First zone entered during the testing trial (B). In the training trial, all the experimental groups explored Block A (grey) first, at the start of the trial. However, during the testing trial, a majority of subjects in the LPS group and PBS explored Block B (blue), which contained the novel object, first. Surprisingly, the LPS + H group showed equal preference for both the familiar and novel object. The PBS + H group showed bias towards to the familiar object.

Altogther, the results of this assay suggest that low systemic LPS does not cause impairments in learning and recognition memory.

4.4.3) Open-Field Analysis

Habituation was not required for this assay. To begin, the number of head within the outer zone during the training phase were analyzed and presented in **table 22 and figure 37.** As shown, the number of head entries was not significantly different across the groups.

Table 22: A summary of the mean scores and p-values obtained when evaluating the number of entries completed by the subject in the outer zone of the open-field arena during the testing trial.

Summary of the descriptive statistics of the number of entries within the outer zone during the testing phase					
Group	Trial	Mean	Adjusted P-value	Significant?	
PIES	Testing Phase	6.30	· ·	•	
PIBS vs. UPS		8.30	0.1271	No	
PBSvs. LPS+H		5.10	0.4891	No	
PBS vs. PBS+H		5.60	0.8223	No	

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Figure 37: A bar graph plot with mean ± SEM showing the number of entries completed in the outer zone of the open-field arena during the testing trial.

Next, the average speed within the outer zone during the training phase were analyzed and presented in **table 23 and figure 38.** The average speed was not significantly different between the groups.

Table 23: A summary of the mean scores and p-values obtained when evaluating the average speed undertaken by the subject in the outer zone of the open-field arena during the testing trial.

Summary of the descriptive statistics of the average speed within the outer zone during the testing phase						
Group	Trial	Mean Adjusted P-value :	Mean Adjusted P-value S	Mean Adjusted	Adjusted P-value Signific	Significant?
PBS	lesting Phase	0.08160	· ·	-		
PBS vs. LPS		0.08790	0.7665	No		
PBS vs. LPS+H		0.08810	0.7503	No		
PBS vs. PBS+H		0.09050	0.5450	No		

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Figure 38: A bar graph plot with mean ± SEM showing the average speed undertaken by the subject in the outer zone of the open-field arena during the testing trial.

Then, the average number of visits in the outer zone during the training phase were analyzed and presented in **table 24 and figure 39.** The average number of visits was not significantly different between the groups **Table 24:** A summary of the mean scores and p-values obtained when evaluating the average number of visits in the outer zone of the open-field arena during the testing trial.

Summary of the descriptive statistics of the average number of visits within the outer zone during the testing phase					
Group	Trial	Mean	Adjusted P-value	Significant?	
PBS	Testing Phase	30.21	· ·	-	
PBS vs. LPS		23.68	0.5961	No	
PBSvs. LPS+H		37.31	0.5343	No	
PBSvs. PBS+H		36.62	0.6092	No	

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Figure 39: A bar graph plot with mean ± SEM showing the average number of visits in the outer zone of the open-field arena during the testing trial.

Then, the average speed within the outer zone during the training phase were analyzed and presented in **table 25** and **figure 40**. Again, the average was not significantly different between the groups.

Table 25: A summary of the mean scores and p-values obtained when evaluating the time spent in the outer zone of the open-field arena during the testing trial.

Summary of the descriptive statistics of the time spent in the outer zone during the testing phase					
Group	Trial	Mean	Adjusted P-value	Significant?	
PB5	Testing Phase	172.2	· ·	-	
PBS vs. LPS		171_3	0.8984	No	
PBS vs. LPS+H		173.7	0.6914	No	
PBSvs. PBS+H		172.5	0.9952	No	





Figure 40: A bar graph plot with mean ± SEM showing the time spent completed by subject in the outer zone of the open-field arena during the testing trial.

Further, the number of lines crossed within the outer zone during the training phase was analyzed and presented in **table 26** and **figure 41**. Again, the average was not significantly different between the groups.

Table 26: A summary of the mean scores and p-values obtained when evaluating the number of line crossings in the outer zone of the open-field arena during the testing trial.

Summary of the descriptive statistics of the number of line crossings in the outer zone during the testing phase					
Group	Trial	Mean	Adjusted P-value Signi	Significant?	
PBS	Testing Phase	69.7	· ·	-	
PBS vs. LPS		73.1	0.9049	No	
PIBS vs. LPS+H		73.9	0.8395	No	
PBS vs. PBS+H		76.8	0.5333	No	

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Figure 41: A bar graph plot with mean ± SEM showing the number of line crossings completed by subject in the outer zone of the open-field arena during the testing trial.

For the middle zone, the number of head entries made within was analyzed and presented in **table 27 and figure 42.** Results show that the average was not significantly different between the groups. **Table 27:** A summary of the mean scores and p-values obtained when evaluating the number of entries in the middle zone of the open-field arena during the testing trial.

Summary of the descriptive statistics of the number of entries in the middle zone during the testing phase					
Group	Trial	Mean	Adjusted P-value	Significant?	
PBS	Testing Phase	6.6	· ·	•	
PBS vs. LPS		8.6	0.1907	No	
PBSvs. LPS+H		4.9	0.3040	No	
PBSvs. PBS+H		5.7	0.7567	No	





Figure 42: A bar graph plot with mean ± SEM showing the number of entries in the middle zone of the open-field arena during the testing trial.

Next, the time spent in the middle zone during the training phase was analyzed and presented in **table 28 and figure 43.** Data suggests that the time spent was not significantly different between the groups.

Table 28: A summary of the mean scores and p-values obtained when evaluating the time spent in the middle zone of the open-field arena during the testing trial.

Summary of the descriptive statistics of the time spent in the middle zone during the testing phase					
Group	Trial	Mean	Adjusted P-value	Significant?	
PBS	Testing Phase	6.980	-	-	
PBS vs. LPS		8.130	0.7921	No	
PBS vs. LPS+H		5.550	0.6711	No	
PBS vs. PBS+H		7.0	>0.9999	No	



Figure 43: A bar graph plot with mean ± SEM showing the time spent in the middle zone of the open-field arena during the testing trial.

Next, the average speed detected in the middle zone during the training phase was analyzed and presented in **table 29 and figure 44.** Data suggests that the average speed was not significantly different between the groups. **Table 29:** A summary of the mean scores and p-values obtained when evaluating the average speed undertaken in the middle zone of the open-field arena during the testing trial.

Summary of the descriptive statistics of the average speed in the middle zone during the testing phase					
Group	Trial	Mean	Adjusted P-value	Significant?	
PBS	Testing Phase	0.2680	· · ·	-	
PBS vs. LPS		0.2530	0.9132	No	
PBS vs. LPS+H		0.2713	0.9988	No	
PBSvs. PBS+H		0.2526	0.9070	No	

SBAAL TODA: Arendys Baard



Figure 44: A bar graph plot with mean ± SEM showing the average speed in the middle zone of the open-field arena during the testing trial.

Finally, the number of lines crossed within the middle zone during the training phase was analyzed and presented in **table 30** and **figure 45**. Data suggests that the average speed was not significantly different between the groups.

Table 30: A summary of the mean scores and p-values obtained when evaluating the number of line crossings completed by the subject in the middle zone of the open-field arena during the testing trial.

Summary of the descriptive statistics of the number of lines crossed in the middle zone during the testing phase					
Group	Trial	Mean	Adjusted P-value	Significant?	
PBS	lesting Phase	11.90	· ·	-	
PBS vs. LPS		13.70	0.7071	No	
PBS vs. LPS+H		9.4	0.4752	No	
PBS vs. PBS+H		10.70	0.8842	No	

Middle Zone: Basches of Lize Cosseinge



Figure 45: A bar graph plot with mean ± SEM showing the number of line crossings completed by subject in the middle zone of the open-field arena during the testing trial.

For the central zone, the number of head entries made was analyzed and presented in **table 31** and **figure 46**. Results show that the average was not significantly different between the groups. **Table 31:** A summary of the mean scores and p-values obtained when evaluating the number of entries completed by the subject in the central zone of the open-field arena during the testing trial.

Summary of the descriptive statistics of the number of entries in the central zone during the testing phase				
Group	Trial	Mean	Adjusted P-value	Significant?
PB5	lesting Phase	1.2	· · ·	-
PBS vs. LPS		1.3	0.9916	No
PBSvs. LPS+H		0.8	0.6854	No
PBSvs. PBS+H		0.9	0.8320	No

Cesides: Number of Endless



Figure 46: A bar graph plot with mean ± SEM showing the number of entries completed by subject in the central zone of the open-field arena during the testing trial.

Then, the time spent in the central zone was analyzed and presented in **table 32 and figure 47.** Results show that the time spent was not significantly different between the groups.

Table 32: A summary of the mean scores and p-values obtained when evaluating the time spent in the central zone of the open-field arena during the testing trial.

Summary of the descriptive statistics of the time spent in the central zone during the testing phase				
Group	Trial	Mean	Adjusted P-value	Significant?
P85	lesting Phase	0.82	-	-
PBS vs. LPS		0.64	0.9479	No
PBSvs. LPS+H		0.71	0.9870	No
PBSvs. PBS+H		0.47	0.7334	No

Semler: They Speak



Figure 47: A bar graph plot with mean ± SEM showing the time spent in the central zone of the open-field arena during the testing trial.

Furthermore, the average speed in the central zone was analyzed and presented in **table 33 and figure 48**. Results show that the average speed was not significantly different between the groups.

Table 33: A summary of the mean scores and p-values obtained when evaluating the average speed undertaken in the central zone of the open-field arena during the testing trial.

Summary of the descriptive statistics of the average speed in the central zone during the testing phase				
Group	Trial	Mean	Adjusted P-value	Significant?
PBS	Testing Phase	0.2063	-	-
PBS vs. LPS		0.2994	0.3101	No
PBS vs. LPS+H		0.2983	0.4517	No
PBS vs. PBS+H		0.2689	0.6156	No



Figure 48: A bar graph plot with mean ± SEM showing the average speed undertaken in the central zone of the open-field arena during the testing trial.

Next, the average number of visits in the central zone was analyzed and presented in **table 34 and figure 49.** Results show that the average number of visits was not significantly different between the groups. **Table 34:** A summary of the mean scores and p-values obtained when evaluating the average number of visits in the central zone of the open-field arena during the testing trial.

Summary of the descriptive statistics of the average number of visits in the central zone during the testing phase				
Group	Trial	Mean	Adjusted P-value	Significant?
PB5	Testing Phase	0.7750	· ·	-
PBS vs. LPS		0.4429	0.2784	No
PBSvs. LPS+H		0.6750	0.9580	No
PBSvs. PBS+H		0.5286	0.5105	No





Figure 49: A bar graph plot with mean \pm SEM showing the average number of visits in the central zone of the open-field arena during the testing trial.

To conclude, the number of lines crossed in the central zone was analyzed and presented in **table 35** and **figure 50**. Results show that the average number of visits was not significantly different between the groups.

Table 35: A summary of the mean scores and p-values obtained when evaluating the number of line crossings in the central zone of the open-field arena during the testing trial.

Summary of the descriptive statistics of the number of line crossings in the central zone during the testing phase				
Group	Trial	Mean	Adjusted P-value	Significant?
P85	Testing Phase	1.1	· ·	•
PBS vs. LPS		1.3	0.9382	No
PBS vs. LPS+H		0.8	0.8278	No
PBSvs. PBS+H		0.9	0.9382	No

Conter: Number of Line Consumers



Figure 50: A bar graph plot with mean \pm SEM showing the number of line crossings in the central zone of the open-field arena during the testing trial.

Altogether, the open-field analysis indicates that ten-day systemic exposure to 0.1 M LPS was not potent enough to induce anxiety-like behaviour and locomotor impairments.

4.5 Discussion

Neurodegenerative diseases are characterized by a crippling intellectual decline. The development of murine models has been exceptionally important because they have been adopted to assess neurodegeneration with qualities like those in the human brain.²²⁸ Models

that restate different features of AD, including deficits in cognitive areas disturbed in patients provide fundamental tools for extensive research into the diseases and provide beneficial means for effective screening *in vivo*.²²⁸ As yet, the exact aetiology of neurodegeneration remains unclear.²²⁹

Presently, there are no therapies available to cure AD.²³⁰ Medications that are available on the market are disease-modifying, in that they target the inherent cause of the disease.²³⁰ This suggests an earnest requirement for effective pre-clinical, and new remedial approaches for the disease.²³¹

Changes to features of cognitive behaviour can provide meaningful markers of neurodegeneration, and they can reveal signs relating to the disease prognosis, treatment, and complications. To identify signs of cognitive decline and/or group differences in behaviour between healthy subjects and those exposed to systemic LPS. This chapter investigated three behavioural tests commonly used in the laboratory to assess such impairments, and its association to neuroinflammation in a homogenous group of Sprague Dawley rats. The results of the control group (PBS only) were compared to the three experimental groups (LPS vs. LPS + H vs. PBS + H). Only significant results will be discussed.

4.6.1) Y-maze Test

Collectively, the findings reveal no statistical significance in the differences between the control and experimental groups, and thus no conclusive association between acute LPS-induced systemic neuroinflammation and spatial recognition. However, when comparing just the means of each group, per parameter, notable differences are seen.

Statistical analysis showed a significant difference in the average speed between the control group and PBS + H group (mean=0.07630, p<0.0347) in novel arm during the testing trial (**table 5**). Although speculative, these results suggest that honey may facilitate increased short-term spatial working memory by alleviating symptoms of memory decline in rodents, hence the PBS + H group showed preference to the novel arm and moved about at a higher speed than the other groups. This is supported by a study by Akanmu *et al.*, (2011), who examined the neurological impacts of Nigerian honey using the Y-maze paradigm, and concluded that honey improved spatial working memory in mice.²³² Examination of the time spent in motion in the long arm (**table 7**) during the testing trial indicated that in general, all four groups spent much more time in motion inside the novel arm than they did in the long arm (although not significant). The amount of time that the subjects remained in motion in the

novel arm, was suggestive of their short-term spatial working memory ability. Surprisingly, upon further scrutiny of the mean scores during the testing trial in the long arm, significant differences were noted when the control group was compared to LPS (mean=8.780, p<0.0440) and PBS + H (mean=8.320, p<0.0237) group. The motion-time profile of these two groups when evaluated against the training trial in the long arm and testing trial in the novel arm could be suggestive of weakened spatial memory consolidation processes in the LPS populations. This is consistent with previous findings that LPS administration impairs cognitive ability and exploratory behaviour.²²⁹ Furthermore, table 8 reveals significant differences in the time that the LPS + H (mean=12.50, p<0.0183) and PBS + H (mean=13.51, p<0.0449) groups remained immobile in the novel arm. A possible explanation to why the honey-fed experimental groups showed increased mobility (less time immobile) in the novel arm is that honey improves spatial memory, thus they explored the novel arm with greater enthusiasm and mobility. Examination of the time spent in the familiar arm (table 10) during the testing trial showed a significant difference between the control group and PBS + H group (mean=35.40, p<0.0072), where the PBS + H group spent more time in the familiar arm than they did in the novel arm. However, non-conclusive associations can be drawn from these results. On the other hand, data shown in table 5 indicates a significant difference in the average speed undertaken by the PBS + H (mean=0.07630, p<0.0347) group in the novel arm during the testing trial. The PBS + H group explored by novel arm at a greater pace than the other three groups, which may suggest that honey consumption induces hyperactivity in rodents. This is consistent with findings of Marwitz and colleagues²³³ who concluded that rats exposed to a Western-style Diet, comprising high sugar and saturated fats, exhibit behaviours characteristic to hyperactive-impulsive type ADHD. Lastly, observations of the time lapsed while the subjects were immobile show that the PBS + H group (mean=18.52, p<0.0132) spent significantly more time immobile in the familiar arm during the testing trial, than the control group (table 14). Plausible justifications for this observation can't be made. When time immobile in the novel arm was examined, it was noted that both the LPS + H (mean=12.50, p<0.0183) and PBS + H (mean=13.51, p<0.0449) groups spent significantly less time immobile in the novel arm than the control group. Again, this emphasizes previous findings that a high-sugar diet (honey) contributes to hyperactivity and impulsivity in Sprague Dawley rats.233

4.6.2) Novel Object Recognition Test

Collectively, the findings reveal no statistical significance in the differences between the control and experimental groups. These observations are consistent with findings by De La Torre²³⁴ that indicated that male C57BL/6 mice exhibit decreased speed (cm/s) and distance (cm) after LPS exposure. However, when the number of entries in Block B (which contained the novel object) during the testing trial were analyzed, a significant difference between the LPS group (mean=7.5, p<0.0044) and the control group (mean=4.0) was observed (table 18). This indicates that the LPS group exhibited more curiosity in the novel object by exploring it more frequently than the control group. Similarly, analysis of the first zone entered (figure 35) during the testing trial shows that LPS group and PBS group explored Block B first, which contained the novel object. Surprisingly, the LPS + H group showed equal preference for the familiar and novel object. Meanwhile the PBS + H group showed bias towards to the familiar object. Conclusions by Czerniawski J et al., (2015) indicate that even though the subjects were able to differentiate a familiar object from a novel one, LPS-induced neuroinflammation does not hinder memory recall in every hippocampal-dependant task, but instead it may distinctly interfere with tasks that demand context discrimination.²³⁵ For the most part, the results obtained in this assessment may be attributed to the age (ten-weeks old) of the subjects which limits them from displaying significant cognitive dysfunction, and recognition memory. However, studies have reported that significant brain development in rats occurs until 9 weeks of age, and CNS myelination of limbic structures occurs until 6 weeks of age.^{236,237} Alternatively, the two-minute trial period may have not been sufficient for memory consolidation and recall, which consequently resulted in the task not being sensitive enough to identify subtle differences between the groups.

4.6.3) Open-field Test

Taken together, the outcomes reveal that there were no significant differences between the control group and experimental groups. This implies that LPS-induced neuroinflammation does not induce anxiety or motor deficits of rodents. This is supported by Bassi and colleagues²³⁸ who concluded that LPS administration up to 200 µg/kg does not cause motor dysfunction. Given that this study sought to simulate a period of sustained inflammation that may arise naturally in healthy individuals, a low concentration of LPS treatment administered regularly was sufficient. In summary, a study by Tanaka, *et al.*, (2012) identified four factors that may explain differences underlying rodent behaviour in the open-maze paradigm.²³⁹

These include activity, sequential organization, diversive exploration and inspective exploration.²³⁹ The authors suggest that this four-factor model be used in drug efficacy and psychiatric studies, to ensure systematic and consistent characterization of rodent behaviour.

4.6 Conclusion

Research has validated the importance of LPS-induced murine models of neuroinflammation in understanding the pathological mechanisms underlying neurodegeneration.²²⁹ Compounding reports reveal that the administration of LPS promotes neuroinflammation, coupled with damage to the BBB and memory dysfunction.⁹⁶ The administration of LPS has been shown to reduce cognitive ability in animals and cause a complex range of behaviours including weight loss, depression, reduced mobility, and intensified anxiety and sleepiness.^{228,240,241}

Sufficient evidence has been collected in this chapter to deduce that 0.1 M LPS systemically administered at a 0.1 mg/ml concentration and a volume of 0.1 ml/kg for a period of 10 days does not cause significant cognitive dysfunction in Sprague Dawley rats. This may be attributed to a considerable number of elements that affect the results observed in behavioural testing, such as group selection, task complexity, individuality, data analysis, animal-investigator interactions and repetition.^{242,} At the same time, there are several factors that can impact animal response to LPS administration like the dosage of LPS given, type of LPS exposure, route of exposure, and period of exposure.²⁴³ For example, if it is administered acutely, prior to training, it impairs cue-fear conditioning, whereas chronic administration of LPS, has been found to impair spatial memory and promote memory and learning deficits. ²⁴⁴ A single intraperitoneal injection of LPS in a dose of 100 µg/kg in adult male Wistar rats, impaired memory object recognition.²⁴⁵ Shaw *et al.*²⁴⁶ reported that a single intraperitoneal injection of LPS in a dose of 250 µg/kg impaired hippocampal dependent spatial learning in the Morris water maze behavioral test. In another study, in adult male Wistar rats, a single injection of LPS in a dose of 1mg/kg, impaired cognitive performance in the Barnes Maze test and in the inhibitory avoidance test.²⁴⁷ Therefore, to mimic a period of sustained inflammation, repeated exposure to low dose LPS (concentration of 0.1mg/ml) was determined using reference articles.

Group selection and randomization can influence the cognitive and sensory-motor abilities of each group.²⁴² In this study, the sample groups were categorized by weight range. This may have impacted mobility performance in the behavioural tests in such a way that the heavier
cohort had fewer entries into the arms/blocks, showed longer periods of immobility (which may have been incorrectly interpreted as increased exploratory behaviour) and completed the task at slower speeds when compared to lighter group. Task complexity may challenge subject performance if the task is too easy or too demanding.²⁴² Although speculative, results from the NORT and open-field test, which show no significant differences between the control and experimental groups, may suggest a negative impact possibly caused by the notion that the tasks were too complex for the low dosage of LPS and short exposure time. Furthermore, it is important to note that the subjects are individuals and will consequently behave differently despite the treatment method selected.²⁴² Animal-investigator interactions, which includes the experience level of the investigator and their degree of comfort working with rodents, may have an adverse effect on the results. In this study, the principal investigator was a novice at animal studies, and therefore, may have unintentionally induced fear and anxiety in the subjects causing their performance to deviate from the predicted outcomes. Altogether, it is conceivable that the non-significant differences observed in this study may be attributed to the factors listed above, in addition to the suggestion that acute neuroinflammation plays a protective role in the body.^{248,249,250}

To confirm whether or not the LPS treatment induced histological changes similar to what is observed in AD-dementia, amyloid formation and deposition into the hippocampus was examined by enzyme-linked immunosorbent assay (ELISA) specific for the A β_{42} isoform, which is considered to be the most toxic.



5.1 Chapter Objectives

This chapter will address the methods and results pertinent to the following objective;

 Assessing the progression of amyloid β (Aβ) formation by measuring the amount (concentration) of soluble Aβ₄₂ levels in the hippocampal tissue of aged Sprague-Dawley rats using a sandwich-ELISA.

5.2 Introduction

The aggregation of soluble amyloid oligomers and deposition into fibrils and plaques is a pathogenic phenomenon central to AD dementia. The most prevalent amyloid β peptides comprise 40 (A β_{40}) and 42 (A β_{42}) amino acids.¹⁹ Despite the fact that the A β_{40} peptides are produced in larger quantities than A β_{42} , A β_{42} isoforms account for a larger portion of amyloid aggregates in the brain because of its tendency to misfold-which renders it more toxic.²⁵¹ Nonetheless, extracellular A β is transported from the blood to the brain and vice versa. Translocation into the brain is facilitated by carrier-receptor transport across the BBB whilst degradation occurs by proteins such as astrocytes, plasmin and apolipoprotein E (ApoE).^{252,253,254,255} The concentration of soluble A β peptides in the CNS plays a significant role in the production of toxic oligomers, and this concentration is profoundly regulated by A β transport across the BBB.²⁵⁶ Moreover, recent studies show that soluble A β aggregates trigger neuronal dysfunction and activate microglia.²⁵⁷ And neurotoxic A β oligomers activate gliosis, which stimulates the release of pro-inflammatory mediators by astrocytes and microglia *in vitro*.²⁵⁸ Essentially, enzyme-linked immunosorbent assay (ELISA) is a simple technique used to detect the concentration of A β in plasma, the brain and CSF.²⁵⁹

ELISA is a plate-based analytical procedure used in biochemistry to quantitatively determine the concentration of soluble substances in solution.²⁶⁰ The assays is conducted in 96-well polystyrene plates which bind antibodies. Multiple wells per plate allow for several samples to be run at once. The bottom surface of each well of the plate is pre-coated with the antigen of interest. Plates are purchased with the wells already coated and the concentration of the fixed antigen differs for each ELISA kit therefore the detection range / sensitivity of the antigen-antibody complex is always indicated. During the analysis an intense colour change indicates a higher concentration of antibody (i.e. protein) present. For the analysis, the generated data is evaluated in comparison to a standard curve (optical density vs. log concentration) and the antibody of interest is quantified.

5.3 Methods and Materials

All the samples were prepared at the Laboratory of Microscopy and Microanalysis unit, University of Pretoria, Prinshof campus. In this study, the concentration of soluble $A\beta_{42}$ present in the hippocampal tissue was determined. Commercially available ELISA kit from Elabscience[®] catalogue number E-EL-R1402 was used.

5.3.1) Sample preparation

After termination, the right hemisphere was used for ELISA. On an ice-cold metal plate, the right hemisphere was separated into the cortex, cerebellum, and hippocampus. The hippocampus was extracted and placed inside an empty round bottom Eppendorf tube. The tube was immersed in liquid nitrogen for snap freezing and the tissue was maintained in dry ice until final storage in a -80°C freezer until the assay was conducted. The hippocampal tissue was brought to room temperature before it was homogenized in EzLys[™] tissue protein extraction reagent (purchased from Biocom Africa) containing DNAse 1 protease inhibitor cocktail (purchased from Sigma[®], South Africa).

5.3.2) Soluble Aβ₄₂ Assay

<u>Brain homogenate</u>: The tissue sample was weighed by recording the weight of the empty eppendorf tube and subtracting that from the weight of the tube containing the right hemisphere. For each gram of tissue to be homogenized, 10-15 ml of cold tissue protein extraction reagent and 100-150 U of DNAse I protease inhibitor was added. The tissue was homogenized with 15 strokes using Dounce disposable homogenizer. The sample was then centrifuged at 10,000 ϑ at 4°C for 10 minutes. The supernatant was collected, aliquotted and used to quantify the amount of soluble A β_{42} protein present in the hippocampal tissue.

<u>Reagent preparation</u>: All reagents were brought to room temperature before use. To prepare the wash buffer, 30 mL wash buffer concentrate was diluted in 720 mL of distilled water. To prepare standard working buffer; the standard was centrifuged at 10,000 ϑ for 1 minute

followed by the addition of 1.0 mL of reference standard and sample diluent. This was gently mixed and allowed to stand for 10 minutes before serial dilutions of the gradient 1000, 500, 250, 125, 62.5, 31.25, 15.63, 0 pg/mL were made. To prepare biotinylated detection antibody solution, 100x concentrated biotinlyated detection antibody was diluted to 1x working solution with biotinylated detection antibody diluent. To prepare HRP solution, 100x concentrated https://www.concentrated.concentrated HRP conjugate was diluted to 1x working solution with concentrated HRP conjugate diluent.

Assay procedure: Standard working solution was added to all the wells of the first two columns (100 µL each well). Next, 100 µL of sample was added to the other appropriate wells and the plate was covered and incubated for 90 minutes at 37°C. The solution was drained and 100 µL of biotinylated detection antibody working solution was added to each well. The plate was covered, gently mixed, and incubated for one hour at 37°C. The solution was removed and 350 µL of wash buffer was added to each well, allowed to soak for two minutes before it was decanted and pat dry using clean absorbent paper. This step was repeated three times. Then, 100 µL of HRP conjugate working solution was added to each well; the plate was covered and incubated for 30 minutes at 37°C. The solution was removed, and the wells washed five times with wash buffer. To facilitate antibody detection, 90 µL of substrate reagent was added to each well and the plate was covered and incubated for 15 minutes at 37°C in the dark. Finally, 50 µL stop solution was added to each well and gently mixed. The optical density (OD value) of each well was determined, using a BioTek[®] Epoch microplate reader set to 450 nm absorbance. The generated data was plotted on an optical density vs. log concentration curve and then compared to standards to quantify soluble A β_{42} present in the hippocampal tissue.

5.4 Results

All statistical analyses were completed using GraphPad[™] Prism (version 9.2.0) for Windows (GraphPad[™] Software, San Diego, California USA). The data was analysed using one-way ANOVA. The p-value was set at p <0.05, with a 95% confidence interval.

To characterize the effects of LPS-induced neuroinflammation on total A β_{42} quantity in the hippocampus, the control group (PBS only) was compared to the three experimental groups (LPS only vs. LPS + H vs. PBS + H). Biochemical analysis indicated that there are no significant differences between the four groups. **Table 36** and **figure 51** are a representative summary and graph of the quantity of soluble A β_{42} present.

Total Aβ42 (pg/mL)			
Group	Mean ± SD	p-value	Significant? (yes/no)
PBS	0.1528±0.1959		
PB\$+H	0.1695 ± 0.2306	0.8585	no
LP\$	0.1142 ± 0.1592		
LPS + H	0.1164 ± 0.1583		

Table 36: A summary of the soluble $A\beta_{42}$ mean scores ± standard deviation (SD) per sample group.

The p-value (**table 36**) indicates that the quantity of A β_{42} detected in the hippocampi of the experimental group did not significantly differ from the amount detected in the control group after LPS-induced neuroinflammation.



Figure 51: A dot plot (mean ± SEM) graph showing quantity of A β_{42} peptide found in the hippocampal area using A β_{42} sandwich ELISA. The p-value indicates that the of soluble A β present in the hippocampus did not signifantly differ between the groups.

Results reveal no statistical differences in A β_{42} levels among the groups. However, when comparing just the means of each group, notable differences were seen.

5.5 Discussion

The deposition and aggregation of soluble A β_{42} fibrils into plaques in the brain are known to contribute to the pathogenesis of AD.²⁶¹ Various neurodegenerative diseases exhibit an inflammatory element.²⁶² It is possible that chronic or acute systemic inflammation provides means by which A β_{42} plaques steadily increase in the brain, consequently resulting to it's association with cognitive dysfunction.²⁶³ However, the key components linking acute inflammatory activities, synthesis of soluble A β_{42} peptides, A β_{42} plaque deposition and cognitive decline remain unclear.²⁶³

The objective of this chapter was to quantitively determine the level of A β_{42} present in the hippocampal tissue of each group, following LPS-induced systemic neuroinflammation. According to findings in **table 36**, no significant differences were detected in AB₄₂ level among the four groups. This suggests that the SC administration of 0.1 M LPS systemically administered at a 0.1 mg/ml concentration and a volume of 0.1 ml/kg for a period of 10 days does not significantly elevate $A\beta_{42}$ levels in the hippocampal region of Sprague Dawley rats. This is consistent with reports by Mechnikov²⁶⁴ that pre-conditioning with low-dose LPS causes physiological inflammation –which is characterized by the removal of cellular debris, tissue repair, and thus return to CNS homeostasis and protection- as opposed to a pathological immune response which contributes to neurodegenerative diseases.²⁶⁵ In the AD model, LPS pre-conditioning was shown to improve cognitive impairment by hindering amyloid formation and aggregation.^{266,267,268} This is due to low-dose LPS pre-conditioning inhibits the expression of pro-inflammatory agents, such as IL- β and TNF- α .²⁶⁸ In addition, it leads to the preferential activation of anti-inflammatory microglia instead of pro-inflammatory microglia.²⁶⁹ Although speculative, the concept of "physiological inflammation" may provide insight to the observations of these results.

5.6 Conclusion

The sandwich ELISA used to quantify $A\beta_{42}$ levels showed that LPS did not produce increased $A\beta_{42}$ levels. Instead, it appears that the administered LPS may have triggered an antiinflammatory response. However, assays which detect markers of pro- and anti-inflammatory mechanisms would provide a useful tool to confirm the results from this study. Also, it is imperative to note that the cytokine profile in the brain and blood are influenced by different doses of LPS.²⁶⁵ Pre-conditioning with LPS is said to induce neuroprotection by promoting anti-inflammatory and anti-oxidative processes, and autography.²⁶⁵ Being that LPS is amply present in traditional remedies and foods ingested by humans^{270,271,272}, it is clear that humans consume LPS regularly. This suggests that a higher dose of LPS may have steered the innate immune response to LPS, away from physiological to a pathogical response that eventually results in amyloid accumulation. Therefore, it is valuable for future studies to examine the pro- and anti-inflammatory effects at increasing doses of LPS, in order to further elucidate the shift from regulated inflammation to pathological inflammation.

Chapter 6: Immunofluorescence Assay

6.1 Chapter Objectives

6

This chapter will address the methods and results pertinent to the following objectives:

- Determine the number of astrocytes present in the hippocampal tissue of Sprague-Dawley rats by immunofluorescence staining with glial fibrillary acidic protein (GFAP) antibody using confocal microscopy and ImageJ.
- Identify astrocyte activity (astrogliosis) in the hippocampal tissue of Sprague-Dawley rats by measuring fluorescent intensity using a GFAP antibody using confocal microscopy and ImageJ.
- Identify microglial activity (microgliosis) in the hippocampal tissue of Sprague-Dawley
 rats by co-labeling tissue sections with Iba1 and CD68 antibodies and counting the
 number of Iba1 positive cells using confocal microscopy and ImageJ and measuring
 the fluorescent intensity of co-localized Iba1 and CD68 positive cells using ImageJ.
- Identify the protective effects of honey on the astrocyte and microglial expression/quantity in the hippocampal tissue of Sprague-Dawley rats using ImageJ.
- Identify the presence and fluorescent intensity of fibril amyloid proteins in the hippocampal tissue of Sprague-Dawley rats by measuring fluorescent intensity using Thioflavin-T stain using confocal microscopy and ImageJ.

6.2 Introduction

Immunohistochemistry (IHC) forms an important component of diagnostic pathology, drug development, and research methodology, especially in the fields of neuropathology, hematopathology and oncopathology.²⁷³ It is a commonly used approach for examining protein expression and distribution in health and pathohistological studies.²⁷³ The use of IHC

techniques to examine tissue samples provides details that otherwise can't be attained using standard hematoxylin and eosin (H & E) stains, on pathways that contribute to pathological states.²⁷³ Tissue samples are either embedded in paffarin wax or frozen for microtome or cryostat sectioning, respectively, followed by subsequent staining and visualization by light or confocal microscopy. The technique uses antigen-antibody reactions to identify antigens of interest in biological samples, that are subsequently visualized by a chromogenic-substrate marker.²⁷⁴ Additionally, the technique maintains the composition, cellular and structural integrity of the tissue.²⁷⁴ Therefore, it is a valuable assay with a diverse series of applications that include the; identification of infectious agents in tissue samples, classification and diagnosis of neuropathologies, and diagnosis of muscular dystrophies. Also, IHC can be used to determine the functions of gene products, and serves as a predictive marker for cancer progression.²⁷³

Immunofluorescence (IF) assay is a specialized subcategory of IHC, that is split into two types; direct and indirect IF. This technique uses a fluorescent dye to capture antibody binding, intracellular processes and cellular structures under fluorescent microscopy.²⁷⁴ The antigen of interest binds to a fluorochrome-tagged antibody that emits light when a reaction is present.²⁷⁵ This light is captured by fluorescent microscopes attached with filters specific for the wavelength of light emited.²⁷⁵ Some advantages of IF (compared to IHC) include: higher resolution imaging and stability of fluorescent-labelled tissue as well as the ability to label multiple antigens at a time.²⁷⁶ Lastly, the enzymatic nature of IHC techniques prevents the quantitative analysis of results, whilst the opposite is true for the IF approach. In fact, modern high-output techniques are facilitated by fluorescent detection, for rapid and quantitative microscopy.²⁷⁷

Previous research has shown that GFAP is mostly expressed by astrocytes.²⁷⁸ Additionally, ionized calcium-binding adapter molecule 1 (Iba-1) is distinctly expressed in microglia and macrophages.²⁷⁹ It is induced by cytokines and IFNs and contributes to inflammation.²⁸⁰ Therefore, GFAP and Iba-1 are recognized as biomarkers of astrocytes and microglial cells.²⁸¹ An investigation by Kang, J ²⁸¹ aimed at elucidating the role of LPS in regulating activations of neuroglia in the cerebral cortex used anti-GFAP and anti-Iba-1 fluorescent antibodies to observe Iba-1 and GFAP positive microglia and astocytes. To assess the expression value of these antibodies, they quantified the expression as a ratio of the intensity of the control group, and found that the LPS-treated group expressed greater levels of both antibodies. Moreover, Belfiore and colleagues co-labeled hippoacampal sections with CD68

and Iba-1 antibodies to identify microglial activation.²⁸² Activation was defined as the colocalization of CD68 and Iba-1. Also, they found that the quantity of microglia showing colocalized binding was greater in the LPS-treated group.

In this study, whole brain sections of the left hemisphere were prepared for IF staining with antibodies specific for microglia (anti-CD68 and anti-Iba1) and astrocytes (anti-GFAP). Microscope slides were visualized by fluorescent confocal microscopy. Quantitative data was generated and interpreted using ImageJ software (ImageJ 1.52a, National Institutes of Health, USA) and the results of the experimental groups were compared to the control group. The efficacy of honey as a protective/mopping agent was determined by comparing the quantitative results of the intervention (honey) group to the experimental groups. In line with the foregoing literature in chapter 2, it was anticipated that honey will suppress microglial and astrocyte reactivity.

6.3 Methods and Materials

All tissue samples were prepared at the Laboratory of Microscopy and Microanalysis unit, University of Pretoria, Prinshof campus, following completion of the this first trial of the study, as indicated in **figure 9**. A detailed list of the reagents used in this chapter can be found in **addendum 6**.

6.3.1) Antibodies used for the immunoflorescence

Purchased from Abcam: Anti-GFAP antibody (ab33922,1:1000 dilution); anti-Iba1 antibody (ab5076, 1:1000 dilution); anti-CD68 antibody (ab31630, 1:800 dilution); Thioflavin T stain (T3516, 20µM).

6.3.2) Tissue collection and preservation

Subjects were terminated by inhalation of Isfor[®] followed by perfusion with saline and 4% formaldehyde, which was then followed by decapitation. The brain was carefully removed from the skull and bisected into the right and left hemisphere. The left hemisphere was dropped and fixed in 4% formaldehyde solution for 48 hours at room temperature.

6.3.3) Tissue processing

The tissue was rinsed with 0.1 M phosphate buffered saline (PBS) three times for 10 minutes each to remove formaldehyde precipitates. It was then placed in 15% sucrose solution until it sank to the bottom. The sucrose solution was replaced with a 30% sucrose solution containing 0.05% sodium azide. The tissue was stored at 4°C until sectioning. Sagittal sections of 30 μ m were sectioned using a cryostat set to -20°C.

6.3.4) Free-floating sections

<u>Cryo-protection</u>: All four sides of a 96-well plate were labelled for each brain and cryoprotectant medium was prepared using 125 ml glycerine, 150 ml ethylene glycol and 250 ml 0.1 M PO₄. With the use of a multi-channel pipette, the wells were filled with cryo-protectant and stored at 4°C.

<u>Sectioning</u>: Tissue sectioning was performed using a cryostat (Leica CM 1850,Leica Biosystems) set at -20°C. Sections of 30 µm sections were prepared and floated in wells containing cryo-protectant.

<u>ThT staining</u>: 10 mM stock solution was prepared by dissolving 31.8 mg ThT powder in 0.1M PBS.

<u>Permeabilization</u>: The slides were incubated in 0.1 M TBS containing 0.3% Triton X-100, three times for 15 minutes. Tris buffered saline (TBS) was prepared from 26.44 g trizma hydrochloride, 3.88 g trizma base and 18 g sodium chloride + 2 L ddH₂O. The buffer was removed and the sections were blocked.

Preparation of TBS+: Add 1.25 ml Triton X-100 to 500 ml 0.1 M TBS.

<u>Blocking</u>: Sections were incubated in a blocking buffer prepared using 38 ml TBS+, 0.3 M glycine and 2 ml bovine serum albumin (BSA) for one hour with gentle agitation. The blocking solution was removed and the sections were rinsed in 0.1 M TBS, three times for 15 minutes each.

<u>Immunostaining</u>: The primary antibodies (GFAP, Iba-1, and CD68) were diluted according to manufacturer suggestions. Antibodies were diluted in TBS+. Each well was filled with the solution and incubated in the fridge at 4°C for 72 hours. This was aspirated and sections were washed in 0.1 M TBS three times for 15 minutes each.

<u>Mounting</u>: The sections were mounted on a 1 mm thick clear glass slide (Labocare[®] microscope slides) and allowed to air dry for less than one minute. Fluoromount[™] aqueous mounting medium (Sigma-Aldrich) was used to mount the sections. These were placed flat overnight and stored in a dark slide box, at 4°C, until visualization with a Zeiss LSM 880 confocal laser scanning microscope (Carl Zeiss Microscopy GmbH, Jena, Germany).

6.3.5) Quantitative analysis using ImageJ

Sample analysis involved capturing a series of micrographs that represented features of interest to the study. A minimum of ten representative micrographs per sample was taken to show the general overview of the sample at low magnification (40x objective with a 1.5x digital zoom). Features of interest were identified and further series of micrographs were taken that showed these features at progressively higher magnifications. The following properties were noted:

1. Fluorescent intensity of GFAP stained sections for each subject in all three sample groups using ImageJ.

- 2. The number of GFAP stained cells in each sample using ImageJ.
- 3. The number of Iba1 positive cells in each sample using ImageJ
- 4. Fluorescent intensity of CD68 stained sections for each subject in all three sample groups using ImageJ.
- 5. Co-localization of Iba1 positive cells with CD68 stained lysosomal cells using ImageJ.
- 6. Fluorescent intensity of ThT stained cells using ImageJ.

To quantify co-localization in the confocal micrographs, ten images per rat were quantified and averaged. To count astrocytes and microglia, GFAP and Iba1- stained sections were visualized using a 10x and 40x objective with a 1.5x digital zoom. Micrographs were transferred to ImageJ for cell counting.

6.3.5.1) Cell counting with ImageJ

<u>Astrocyte activity</u>: Automated counting was employed. The image to be analysed was opened, and the cells to be counted were highlighted. The background was subtracted accordingly. Customization (i.e. size or circularity of particles to be counted) of the analysis tool was made accordingly, the image was processed and the results were obtained. The boxes next to the information of interest were checked and all the relevant data was recorded. The results log was saved on a excel spreadsheet.

<u>Microglial activity</u>: Manual counting was employed. After installing the Cell Counter plug-ins, the image was opened. The plug-in tool was selected and two new windows were opened, a counter window with the image on top of a row of buttons, and another with a results window where cells tallied. To begin counting, "type 1" or "type 2" at the bottom of the counter window was selected followed by a direct click on the cell/object to be counted. After counting, the results button was selected and a total for each cell type plus a grand total of all clicks at the bottom of the results window was generated. The results were saved on a excel spreadsheet.

6.3.5.2) Intensity quantification with ImageJ

To get the intensity of a defined area within the image; ImageJ program was opened. Using the toolbar, the "analyze" tool was selected followed by "set measurements". The boxes next to the information of interest were checked (i.e. area, diameter and intensity). To create a plot of intensity values across features in the image, the "analyze" and "plot profile" tools were used. The background was subtracted accordingly. Customization of the analysis tool was made accordingly, the image was processed and the results were obtained. The results log were saved on a excel spreadsheet.

6.4 Results

The data is presented as microphotographs of hippocampal sections, which show the fluorescent profiles observed. Microphotographs of each sample group were captured and used to compare the experimental groups to the control group. The samples were visualized using a 40x and 10x objective, during which the fluorescent trends and cell counts were noted. ImageJ was used to determine the fluorescent intensity observed in addition to performing cell counts. To calculate fluorescent intensity, the corrected total cell fluorescence (CTCF) was used to reliably compare the groups. Statistical tests were completed using GraphPadTM Prism (version 9.2.0) for Windows (GraphPadTM Software, San Diego, California USA). The Shapiro-Wilk normality test was used to establish that the dataset was normally distributed. The dataset was analyzed using one-way ANOVA. Post-hoc checks with Dunnett's multiple comparisons test were used to compare the three experimental groups to the control group. The p-value was set at p <0.05, with a 95% confidence interval. The results were represented as mean ± standard error of mean (SEM). Error bars indicate SEM.

To characterize the effects of LPS-induced neuroinflammation on hippocampal astrocytes, microglia, and amyloid presence, the control group was compared to the three experimental

groups by immunostaining with GFAP, Iba1 antibodies (and CD68 for colabelling) and ThT stain respectively. Analysis of the mean scores, per group revealed non-significant differences between them, although notable differences in fluorescent intensity and quantity of cells was observed between the groups and thus discussed.

6.4.1) Astrocytes

Sections stained with the GFAP antibody emitted a blue fluorescencethat is observable on the microphotographs of **figure 51 (A-D)**.



Figure 52 (A-D): Hippocampal microphotographs from the four groups stained with anti-GFAP. Scale bar was set at 10 μ m. (A) Microphotograph from the control group. (Label 1) An astrocyte cell emitting slight fluorescence. (B) Microphotograph from the PBS + H group. (Label 2) An astrocyte cell emitting more fluorescence than the control group. (C) Microphotograph from the LPS group. (Label 3)

Astrocytes showing a greater degree of fluorescence and cell quantity than the control group. (D) Microphotograph from the LPS + H group. (Label 4) Astrocytes emitting more fluorescence and cell quantity than the control group.

A summary of the descriptive statistics of the fluorescent intensity observed when the hippocampal tissue was stained with anti-GFAP antibody is shown in **table 37**. Analysis of the CTCF values show that fluorescent intensity was not significantly different between the groups.

Table 37: A summary of the mean CTCF scores and p-values obtained when evaluating the fluorescent intensity of the four groups after immunostaining with anti-GFAP.

Astrocyte Immunostaining			
Group	Mean CTCF	Adjusted p-value	Significant?
PDS	35186	-	E
PBS vs. PBS + H	43382	0.5701	No
PBS vs. LPS	49168	0.2665	No
PBS vs. LPS + H	48426	0.3053	No

Astrocytes: Fluorescent Intensity



Figure 53: Bar graph plot with mean and standard error of mean (SEM) for the fluorescent intensity of astrocytes (expressed as CTCF score) across all four groups.

6.4.2) Microglia

Sections stained with anti-Iba-1 antibody emitted a red fluorescencethat is observable on the micrographs of **figure 54 (A-D)**. Here, activation was defined as an increase in fluorescent intensity, indicated by the CTCF value, and density of microglia, indicated by cell counts **(table 41).** Data shows that the fluorescent intensity of the LPS group was significantly higher than the control group. This indicates that low, systemic LPS exposure induced the activation of microglia.



Figure 54 (A-D): Hippocampal microphotographs of the four groups stained with anti-Iba1. Scale bar was set at 10 µm. (A) Micrograph from the control group. (Label 1) Microglia showing slight activation (fluorescence). (B) Microphotograph from the PBS + H group. (Label 2) Microglia showing more activation than the control group. (C) Microphotograph from the LPS group. (Label 3) Microglia showing a significanty higher degree of fluorescence and cell quantity than the control group. (D)

Microphotograph from the LPS + H group. (Label 4) Microglia showing a fair degree of fluorescence and cell quantity than the control group.

Table 38: A summary of the mean CTCF scores and p-values obtained when sections were immunolabeled with anti-Iba-1 evaluating the fluorescent intensity of the four groups.

Microglia Immunostaining			
Group	Mean CTCF	Adjusted p-value	Significant?
PBS	84360	-	<u>.</u>
PBS vs. PBS + H	42923	0,3029	No
PBS vs. LPS	244103	<0.9999	Yes
PBS vs. LPS + H	96160	0,7683	No

Microglia: Fluorescent Intensity



Figure 55: Bar graph plot with mean and standard error of mean (SEM) for the fluorescent intensity of microglia (expressed as CTCF score) across all four groups.

6.4.3) Colocalization - Microglial Activation

Sections co-labelled with Iba1 and CD68 antibody emitted a combination fluorescencethat is observable as a orange on the micrographs of **Figure 56 (A-D)**. Here, microglial activation was defined as the co-localization of Iba-1-positive cells and CD68-positive cells. Data showed that no significant differences in microglial activation was present between the experimental and control groups.



Figure 56 (A-D): Hippocampal micrographs from the four groups co-stained with anti-Iba1 and CD68. Scale bar was set at 10 μ m. (I) Micrograph from the control group. (Label 1) Microglia showing minimal activation. (J) Micrograph from the PBS + H group. (Label 2) Microglia in close-proximity to CD68-labelled lysosomes. (K) Micrograph from the LPS group. (Label 3) CD68 positive microglia cells indicating a degree of microglial activation and cell quantity than the control group. (L) Micrograph from the LPS + H group. (Label 4) A CD68 positive microglia cell indicating a degree of microglial activation than the control group.

A comparison of the mean CTCF values, presented in **table 39**, show that the fluorescent intensity of Iba-1 and CD68-positive cells between the groups were not significantly different.

Table 39: A summary of the mean CTCF scores and p-values obtained when evaluating the fluorescent intensity of the four groups after co-staining with anti-Iba1 and CD68.

Co-localization immunostatining			
Group	Mean CTCF	Adjusted p-value	Significant?
PDS	8832	-	-
PBS vs. PBS + H	10991	0,575	No
PBS vs. LPS	19861	0,696	No
PBS vs. LPS + H	12983	0.610	No

Colocalization: Fluorescent Intensity



Figure 57: Bar graph plot with mean and standard error of mean (SEM) for the fluorescent intensity of activated microglia (expressed as CTCF score) across all four groups.

6.4.4) Thioflavin T

Sections stained with ThT stain emitted a green fluorescenceobservable on the representative Micrographs of **Figure 58 (A-D)**. Here, amyloid progression was determined by the comparing the CTCF values of the experimental groups to the control group.



Figure 58 (A-D): Hippocampal Micrographs of the four groups stained with ThT. Scale bar was set at 10 μ m. (M) Micrograph from the control group. (Label 1) A β fibrils staining positive with ThT. (N) Microphotograph from the PBS + H group. (Label 2) An amyloid fibril showing more fluorscent intensity than the control group. (O) Micrograph from the LPS group. (Label 3) A β fibrils appear to be increased than the control group. (P) Micrograph from the LPS + H group. (Label 4) A β fibrils showing less fluorescence and cell quantity than the LPS group.

A comparison of the mean CTCF values, presented in table 40, show that the fluorescent

intensity of ThT-positive cells between the groups was not significantly different.

Table 40: A summary of the mean CTCF scores and p-values obtained when evaluating the fluorescent intensity of the four groups after immunostaining with ThT.

Thioflavin T Immunostaining			
Group	Mean CTCF	Adjusted p-value	Significant?
PDS	1093	8 4 0	£5
PBS vs. PBS + H	2059	0,408	No
PBS vs. UPS	4253	0,301	No
PBS vs. LPS + H	3384	0,386	No

ThT: Fluorescent Intensity



Figure 59: Bar graph plot with mean and standard error of mean (SEM) for the fluorescent intensity of amyloid fibrils (expressed as CTCF score) across all four groups. No significant differences were detected for the fluorescence intensities between the groups.

6.4.5) Cell counts

The dataset in **table 41** was analyzed using one-way ANOVA. Post-hoc checks with Dunnett's multiple comparisons test were used to compare the three experimental groups to the control group. The p-value was set at p <0.05, with a 95% confidence interval.

Table 41: A summary of the number of cells observed in the micrographs for each IF analysis performed. Significant differences in the number of cells observed between the PBS control group and LPS group apply both astrocytes and microglia.

Antibody/Dye	p-value.	Significant? (yes/no)
GFAP	0.0207	yes
lba-1	0.0041	yes
lba-1 and CD68 (co-localization)	0.6038	no
ТһТ	0.1342	no

6.5 Discussion

There is a rising interest in the study of neuroinflammation and the role in neuropathology, therefore, the use for reliable biomarkers of inflammation are important for diagnosis. Moreover, inflammatory biomarkers were found in the rat brain even two years after exposure to AD-inducing pathogens.²⁸³ In this study, popular biomarkers of inflammation were used to identify the impact of systemic LPS on the profile of neuroglial cells implicated in AD pathology.

6.5.1) Astrocytes

Collectively, the findings reveal no statistical significance in the differences between the control and experimental groups, and thus no conclusive association between LPS-induced systemic neuroinflammation and astrocyte activity. However, when comparing just the mean fluorescence (CTCF score) of each group, notable differences were seen.

Taken as a whole, results in **table 37** show that there were no differences in fluorescent intensity, thus astrocyte activation, between the groups. Although speculative, it suggests that LPS exposure did not have a major effect on the activity of astrocytes. Furthermore, the groups treated with honey did not show decreased astrocyte activation, as anticipated.

These findings are contradictory to previous research which suggest that honey is an effective neuroprotective agent against the effects of LPS-induced neuroinflammation. It is known that honey contains phenol and hydrogen peroxide elements which inhibit the spread of infectious agents^{217,284}, like LPS, that can cause a leaky gut, spread into circulation and the CNS where it triggers neuroinflammation.²⁸⁵ Secondly, Ali and colleagus examined the *in vitro* effects of honey on the viability of astrocytes exposed to oxidative stress and found that honey at 1% (v/v) concentration inhibits apoptosis.²⁸⁶

On the hand, our findings are supported Mohr, K whose investigation into the discrepancy of GFAP as a glial marker showed that there is a discrepancy between species and experimental models that limit the use of GFAP as a marker for neuroglial cell activation.²⁸⁷ This provides insight to the non-significant differences observed in fluorescence (activation) expressed by the groups.

6.5.2) Microglia

Findings from **table 38** reveal a statistical significance in the differences between the control group and LPS group (p<0.0001). The LPS group (mean= 244103, p<0.0001) displayed a considerable degee of fluorescent intensity which suggests that LPS-induced systemic neuroinflammation facilitates microglial activation.²⁸⁸ The activation of microglia facilitates restoration and homeostasis in the neuron microenvironment.²⁸⁹ Since microglia are known to play a vital function of degrading apoptotic cells and aggregated proteins ²⁹⁰, in addition to multiplying in number and reactivity within close proximity to Aβ plaques²⁹¹, the outcomes observed were expected. Microglia are one of the first cells to be activated during neuroinflmammation²⁸⁹, which further validates the signifant fluorescent differences observed between the control and LPS group. Activated microglia show varied interactions with Aß proteins and neuronal pathways, which result in wide-ranging effects on the progression of AD, based on the stage of disease and individual vulnerability.²⁹² The mechanisms linking systemic inflammation and microglial activation remain unclear however, age is a key factor regulating the degree of microglial activation following systemic inflammation.²⁹³ These findings are consistent with reports by Norden and colleagues who observed an increase in Iba-1 immunoreactivity 24 hours after LPS exposure.²⁹⁴ Interestingly, they also found that repeated LPS exposure was linked to immune and behavioural tolerance, and a minimized inflammatory microglial profile opposed to observations of acute LPS exposure.

Furthermore, the fluorescent intensity observed in LPS + H group (mean=98160, p=0.7683) was not significantly different from the intensity observed in both the control and LPS group. This is contradicted with findings by Candiracci, M *et al.* (2012) who report that 0.5 μ g/mL and 1 μ g/mL honey flavonoid extract sourced from unprocessed multifloral honey strongly hinders the secretion of proinflammatory cytokines and reduces the quantity of degenerated neuronal cells in the hippocampus.²⁹⁵ Our findings differ from previous research possibly due to the origin of honey used and the short exposure time, which might have been insufficient to influence microglial activation.

6.5.3) Colocalization - Microglial Actvitation

It is known that CD68 levels are higher in reactive microglia and extensively reduced in resting microglia²⁹⁶. The results indicate that there were no significant differences between the control group and experimental groups. Although **figure 55 (C)** illustrates an increased number of Iba1-positive microglial cells as well as activated (Iba1 and CD68 positive)

microglia when compared to the control group **figure 55** (**A**), in addition to the statistical differences between the PBS group (mean=8832) and LPS group (mean=19661, p=0.698), conclusive associations cannot be derived. The Iba and CD68 antibodies are commonly used as microglial markers, however, due to heterogeneity in gene regulation, they may recognize different activation stages of microglia.²⁹⁷ This may possibly explain the low amount of activated microglia (Iba1 and CD68 positive) seen in **figure 55** (**C**) in comparison the amount of microglia observed in **figure 54** (**C**).

6.5.4) Thioflavin T

The aggregation of soluble A β proteins influences AD progression, which may result in synaptic aberration and subsequent neurodegeneration.²⁹⁰ On the other hand, astrocytes may, in principle, contribute to A β production since they increase β -secretase and APP in pathology states, however, this has not been validated.²⁹⁸ To quantitatively examine fibril A β deposition in the hippocampal region, ThT stain was used as an A β biomarker for which fluorescent intensity was determined and used to compare between the four groups. Changes in intensity levels were used as a marker of fibril synthesis or degradation, where a decline was attributed to the reduction of fibril synthesis caused by the neuroprotective effects of honey.

Results showed that there was noticeable green fluorescencein the hippocampal area of all four groups **figure 57 (A-D)**. The synthesis and deposition of A β peptides occurs in healthy brains without causing cognitive decline.^{299,300} This may provide insight for the presence of A β proteins observed in the control group (**figure 57 A**).

Emergence to AD then, is regulated by the synthesis and degradation of A β proteins and how well they are eliminted from the brain.³⁰⁰ The results in **table 40**, though not significant, reveal that 10 days after s.c injection of LPS, A β deposition was the highest in the LPS group (mean=4253, p=0.301). This observation suggests that the period of exposure to LPS or the concentration of LPS administered was enough to impact amyloid formation or deposition in the hippocampus.

The results in **table 40** suggest that honey treatment did not have a mopping effect on amyloid formation. This is inconsistent with findings by Wan and colleagues which report that honey significantly reduces LPS-induced neuronal loss and presents anti-inflammatory potential against oxidative stress and A β protein deposition.³⁰¹

6.6 Conclusion

Research has validated the importance of glial cells in neuroinflammation and neurodegeneration.³⁰¹ Compounding data present strong evidence that astrocytes actively facilitate the pathogenesis of AD.³⁰² Additionlly, microglia have become the focal point amongst researchers as the cells have been implicated in neurodegeneration. This is attributed to the fact that microglia have the ability to upregulate neuroinflammation, causing subsequent neuronal death.³⁰³ Lastly, foregoing literature highlights the role, significance and pathogenesis of amyloid proteins in AD. Taken as a whole, it is imperative to examine the activity of the glial cells in neuroinflammation models *in vivo*.

Sufficient evidence has been collected in this chapter to deduce that 0.1 M LPS systemically administered at a 0.1 mg/ml concentration and a volume of 0.1 ml/kg for a period of 10 days does not cause significant astrogliosis, microgliosis and amyloid deposition in Sprague Dawley rats.

Literature may be used to explain the lack of significant differences in these observations. Firstly, there exists inter- and intra-regional differences between astrocyte populations, which translates into heterogenous functional features.³⁰² Secondly, though GFAP has been regarded a reliable biomarker for astrocytes, not all astrocytes positively bind to GFAP.³⁰² Finally, astrocytes can provide neuroprotective effects at various stages of AD.³⁰² Both activated astrocytes and microglia, when exposed to A β proteins, release transforming growth factor (TGF- β) which functions to protect neuronal cells from amyloid toxicity and stimulate the removal of A β .³⁰² Altogether, this implies that the activated astrocytes came to a peak level and undertook an adaptive state.

Numerous lines of evidence suggest that microglia are instrumental in AD progression.³⁰⁴ The findings presented in this study reveal that LPS-induced neuroinflammation significantly stimulates microglial activation in LPS rodent populations. However, depending on the stage of disease, activated microglia operate in both beneficial and deleterious ways.³⁰⁴ Therefore, the degree of activation and, thus, their influence on pathogenesis may be contingent on the type and duration of injury, in addition to the CNS area under examination.^{305,306,307} Also, while Iba1 antibody is commonly used to identify microglia, it does not specify microglial polarization since both pro- and anti-inflammatory microglia express Iba1.³⁰⁴

The ThT stain was used to assess fibril amyloid deposition, however, ThT also has several intrinsic disadvantages including poor BBB penetration and short emission wavelength.³⁰⁸ Notably, it has been shown that distinct fibril structural configurations, derived from the same protein, can exhibit varying ThT fluorescent intensities.^{309,310,311} Furthermore, the signal intensity can be regulated by a other agents present in solution, through fluorescence quenching or molecular interactions.^{312,313} Additionally, the binding of ThT is not restricted to Aβ fibrils but it extends to certain resident proteins.³¹⁴ Lastly, fibrils are capable of binding to ThT through various binding modes with varying affinities.³¹⁵

Chapter 7: Integrated discussion and conclusion

Substantial research is currently underway to find a cure for AD globally. Many remarkable advances have been successful in this effort, and it is probable that available treatments will soon shift from symptomatic to curative, thus decreasing the rate of progression towards AD and improving the quality of life in elderly populations. What seems to be lacking in this global undertaking is the comprehensive understanding of the biological pathways and molecular mediators contributing to the aetiology of AD.

The aim of this study was to investigate a concept that has been gaining recognition in neurodegenerative conditions, namely the influence of LPS-induced neuroinflammation on AD-like symptoms in the hippocampal region. These are symptoms resulting in damaged cortical structures, functions, and subsequent cognitive decline. Consequently, it is important to elucidate the role of bacterial agents natural to the GIT, specifically LPS, in inflammation because the chronic release of such agents has been implicated in BBB damage and AD development. In this study, it was noted that systemically distributed 0.1 M LPS does not induce pathological neuroinflammation in the hippocampus. Observations from the assays (behavioural and biochemical) conducted in this study suggest that neuroinflammation did occur in subjects that received LPS, however, examination of the findings imply that this reaction was neuro-protective rather than disease-causing. As discussed in foregoing literature, acute inflammation is beneficial to the immune system since it initiates mechanisms involved in neuro-immune conditioning and cellular repair.⁷⁷

The results obtained from the various behavioural assessments (y-maze, NOR, and openfield test) show that cognitive abilities were not severely affected by LPS. Although analysis of results from the y-maze experiment indicate that the PBS + H group maintained high speeds while exploring the novel arm, it appears that honey may have a role in spatial and working memory. Albeit not conclusive, it is supported by Gasparrini's investigations into the protective impact of Manuka honey on LPS-treated macrophages that show that honey inhibits LPS-induced inflammatory molecules and TLR₄ /NF-k β signaling.³¹⁶ In line with literature, this means that an innate immune response may have been supressed by the Manuka honey since it is known that honey has anti-tumor, anti-oxidant and anti-inflammatory properties.^{316,317,318} It is important to note, however, that the composition and physiochemical properties of honey are contingent on its floral source and origin.³¹⁹ Factors such as the

phenolic and flavonoid content, peroxidase activity, type of sugars present and pH differs among honey subtypes.³²⁰ This may have posed a limitation on the remedial effects observed in this study. Nonetheless, this presents an opportunity for future studies to evaluate the effects of Tualang honey, Malaysian honey and Nigerian honey on LPS-induced systemic inflammation using both a low and high dose of LPS. Then, assess the anti-inflammatory effects of Manuka honey.

It is evident that a diverse array of variables influence behavioural and physiological performance.³²¹ These limiting factors can be broadly categorized into empirical, animal and experimental factors.³²¹ Some empirical factors that may have influenced observations of this study include: the sequence which the assays were performed, measures of health at the beginning of the study, environmental factors that influence innate behaviour, method for measuring performance. It is recommended that the sequence of the assays proceed from the least stressful to the most stressful. In this study, the Y-maze test was performed first, followed by the NOR task and open-field test, respectively. Since the open-field test is the most facile assay of all three, it should have been conducted first, followed by the NOR task and finally, the Y-maze analysis.³²¹ Also, starting with measures of health eliminates the possibility of health confounds effecting physical performance. Moreover, animal factors such as sex, age and vendor source play a role in behavioural performance.³²¹ The population of subjects used in this study comprised 10-week old, male Sprague-Dawley rats sourced from one vendor. Considering that AD is prevalent among older human populations, rodents of an older age may have presented results similar to what is seen in humans. Also, investigating the effects of LPS-induced neuroinflammation in both sexes may provide insight to the role of gender in neuroinflammation. Lastly, since there are variations in the methods for measuring and evulating performance, data from this study may be subject to differences in interpretation of previous studies.

To assess the progression of amyloid deposition, the quantity of soluble $A\beta_{42}$ was measured by sandwich ELISA. The negative results observed from this assay may be linked to the lack of specificity of the type of amyloid peptide detected. Since A β peptides are naturally present and harmless in the brain, and the progression to plaque deposition and AD rises from the aggregation of peptides into fibrils then plaques; the assay used was unable to distinguish the structure of A β_{42} , thus toxicity, detected. The results show that in totality, the quantity of A β_{42} in the hippocampus was not affected by LPS-induced neuroinflammation. However, this

does not reveal whether the ratio of A β peptides, fibrils and plaques differed between the three experimental groups. This is an important variable because the presence of A β plaques has profound effects on pathology than the presence of A β peptides. Techniques such as western blot, in conjunction with ELISA, may be used to qualitatively determine A β deposition and verify results from ELISA. Alternatively, more sophisticated techniques such as magnetic resonance spectroscopy can be used to detect A β plaques, however, this was not feasible in this study. ³²² This is because the quantification of A β using ELISA is mostly used in transgenic rodent models, so not much data is available to support results obtained from wild-type rodents.²⁵⁹ Therefore, it is imperative to examine the effects of LPS-induced neuroinflammation on glial cells that are directly implicated in AD pathology namely astrocytes and microglia.

According to the conventional hypothesis, astrocytes and microglia are major contributors to neuropathologies. These macrophage-like cells are involved in brain homeostatic activities involving neurogenesis, BBB intrigrity, neurotransmitters and synapse growth.³²³ Apart from homeostasis, microglia and astrocytes contribute to inflammation by assuming the role of CNS-resident immune cells.³²⁴ Pathogens like LPS trigger the activation of these cells, and the phenotype (repair or intensify damage) adopted by the cells thereof is regulated by the nature and severity of damage, presence of anatagonistic mediators and activation status of other macrophages.³²⁴

Results from this present study revealed a significant difference in the quantity of activated microglia in the LPS group than the control group, who showed normal levels of microglia. It has been alluded that activated microglia recruit astrocytes, which then exacerbate the brain's inflammatory response to amyloid proteins resulting in cytokine-mediated inflammation.³²⁴ Consistent with this present study, the concentration of microglia increased significantly compared to astrocytes, which are the most prevalent in the brain.¹¹⁸ This suggests that microglia respond to LPS injury within a shorter period of time than astrocytes. Also, the considerable increase in microglial cells compared to astrocyte show a tendency to co-localize with A β plaques.¹¹⁸ In view of this, we can further deduce that in this study, the integrity of BBB was not severely compromised by systemic LPS, as indicated by a non-significant difference in A β 42 concentration observed in the ELISA. The significant increase of microglia in the LPS groups was expected. This is because a mere single injection

of LPS is known to increase microglial density in Sprague Dawley rats.²⁶³ Also, seeing that only microglia greatly increased in quantity, it is plausible that the inflammatory response triggered, was a physiological reaction, aimed at brain repair and homeostasis.²⁶³ However, the categorization of the immune response observed requires the analysis of pro- and antiinflammatory mediators. To overcome this limitation, it is advisable to combine the assays of this study with assays like ELISA to confirm the presence and quantity of inflammatory mediators produced by the activated microglial cells like IL1- α , TNF α and complement component 1q (C1q), all of which are capable of inducing reactive astrogliosis themself.³²⁵

In conclusion, it is suggested that a low dose of systemic LPS may induce neuroprotective molecular interactions that increase the density of microglia in the hippocampus, without significantly influencing astrocytes or the formation of amyloid proteins. Furthermore, honey may be an effective mopping agent that suppresses these molecular interactions, preventing cognitive deterioration and weakened physical abilities.



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Appendice

Addendum 1: Animal ethics committee approval-181/2020



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Addendum 2: Animal Research Ethics Committee-2019/07/44/C



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ANREAL REPRESENCE ETHICS (CORNETTED MARSIC)

GLEARANCE CERTIFICATE NO. 2012/07/44/2

APPLICANT Prof W Daniels

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Addendum 3: MSc committee approval



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Addendum 4: Letter of statistical support



BIOSTATISTICS UNIT

LETTER OF STATISTICAL SUPPORT

18 July 2019

This letter is to confirm that student Nonkululeko Dhlamini Faculty of Health Sciences; University of Pretoria discussed the Project with the title "The effects of neuroinflammation induced by systemic lipopolysaccharide on the hippocampi of aged Sprague-Dawley rats". We therefore confirm that we are aware of the project and undertake to assist with the statistical analysis of the data generated from the project.

Data Analysis

The aim of this study is to investigate if systemic induced lipopolysaccharide (LPS) from Escherichia coli 055:B5 at a volume of 0.1 ml per kg of rat for 10 days induces neuroinflammation in aged Sprague-Dawley rats. Descriptive statistics like mean, median, standard deviation and range will be given for continuous data. Independent samples t-test will be undertaken to assess the effect of systemic induced LPS on the spatial memory between the control, vehicle and experimental aged Sprague-Dawley rats. Data will be captured into Excel and all statistics will be evaluated at 5% level and will be undertaken using STATA 15.

Sample size

To achieve the objectives of this study a sample of 18 aged Sprague-Dawley rats will be used.

Name: Tshifhiwa M Nkwenika Biostatistics Unit tshifhiwa.nkwenika@mrc.ac.za 012 339 8519

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Biostatistics Unit
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Pretoria
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Addendum 5: Study log-Welfare monitoring form

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Addendum 6: List of materials and reagents

Material / Reagent	Vendor	
1 mm glass slide	Labocare®	
24-well plates	Sigma-Aldrich [®]	
Coverslips	Lasec [®]	
Amyloid-β ₄₂ staining kit (E-EL-R1402)	Elabscience®	
Anti-GFAP antibody (ab33922)	Abcam [®]	
Anti-Iba-1 antibody (ab5076)	Abcam [®]	
CD68 antibody (ab31630)	Abcam [®]	
Thioflavin T dye (T3516)	Sigma-Aldrich [®]	
Fluoromount [™] mounting medium	Sigma-Aldrich [®]	
Formaldehyde	Merck	
Manuka honey	Ample Resources South Africa	
Sodium phosphate monobasic (S0751)	Sigma-Aldrich [®]	
Sodium phosphate dibasic (S0876)	Sigma-Aldrich [®]	
Sucrose (S-8501)	Sigma-Aldrich [®]	
Glycerine (G31-500)	ThermoFisher Scientific™	
Ethylene Glycol (BP230-1)	ThermoFisher Scientific™	
Bovine serum albumin	Sigma-Aldrich [®]	
Triton X-100	Sigma-Aldrich [®]	
Sodium chloride (S-9888)	Sigma-Aldrich [®]	
Trizma base (T-1503)	Sigma-Aldrich [®]	
Trizma hydrochloride (T-3253)	Sigma-Aldrich [®]	
Sodium azide (S2002)	Sigma-Aldrich [®]	
Isfor®	Safeline Pharmaceuticals	
Lipopolysaccharide 055:B5 (L2880)	Sigma-Aldrich [®]	
EzLys™ tissue protein extraction reagent	Biocom Africa	
DNAse 1 protease inhibitor cocktail	Sigma-Aldrich [®]	