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Ollscoil na hÉireann, Corcaigh
National University of Ireland, Cork



**Investigating the role of interleukin-6 in neuronal
dysfunction in the *mdx* mouse model of Duchenne
Muscular Dystrophy**

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Declaration

“This is to certify that the work I am submitting is my own and has not been submitted for another degree, either at University College Cork or elsewhere. All external references and sources are clearly acknowledged and identified within the contents. Calcium experiments were conducted by Michael Vaughan and data analysis by Claire Denley. I have read and understood the regulations of University College Cork concerning plagiarism and intellectual property.”

Signed by Kimberley Stephenson, January 2022

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List of Abbreviations

ACSF	Artificial cerebrospinal fluid
ADHD	Attention Deficit Hyperactivity Disorder
AMPA	Alpha-amino-3-hydroxy-5-methyl-4-isoxazolepropionate receptor
ASD	Autism Spectrum Disorder
BMD	Becker Muscular Dystrophy
Ca ²⁺	Calcium
CaCl ₂	Calcium chloride
CaMKII	Ca ²⁺ /calmodulin-dependent protein kinase
CNS	Central nervous system
CS	Cutting solution
DAPC	Dystrophin associated protein complex
DAPI	4',6-diamidino-2-phenylindole
DEPC	Diethyl pyrocarbonate
DG	Dentate gyrus
dH ₂ O	Deionized water
DMD	Duchenne Muscular Dystrophy
DMEM	Dulbecco's modified eagle's medium
EPSP	Excitatory post synaptic potential
FBS	Foetal bovine serum
GABA	γ-aminobutyric acid
H&E	Haematoxylin and Eosin
HBSS	HEPES-buffered saline solution
HPRA	Health Products Regulatory Authority
HPRT	Hypoxanthine-guanine phosphoribosyltransferase
IL	Interleukin
IL-6	Interleukin 6
IL-6Rα	Interleukin-6 Receptor alpha
IL-6st	Interleukin-6 signal transducer

IQ	Intelligence quotient
IVCs	Individually ventilated cage
K ⁺	Potassium
KCl	Potassium chloride
LPS	Lipopolysaccharide
LTP	Long Term Potentiation
mAb	Monoclonal antibody
MAPK	Mitogen-activated protein kinase
MgCl ₂	Magnesium chloride
NaCl	Sodium chloride
NaH ₂ PO ₄	Monosodium phosphate
NaHCO ₃	Sodium bicarbonate
NMDA	N-methyl-D-aspartate
NOR	Novel object recognition
O.C.T	Optimal cutting temperature embedding medium
OCD	Obsessive compulsive disorder
OsMom	Osmolarity
pAb	Polyclonal antibody
PBS	Phosphate buffered saline
PFA	Paraformaldehyde
PI 3-K	Phosphoinositide 3-kinase
P-S	Penicillin-Streptomycin
ROS	Reactive oxygen species
SC	Schaffer Collaterals
SPF	Specific pathogen-free
SR2	Serum Replacement 2
STAT 3	Signal transducer and activator of transcription 3
TNF	Tumour necrosis factor
WT	Wild type

Abstract

Duchenne Muscular Dystrophy (DMD) is a fatal musculoskeletal disorder that results in a loss of dystrophin in muscle fibres, leading to progressive immobility, chronic inflammation and premature death. Inflammation is characterised by increased circulating levels of pro-inflammatory cytokines such as IL-6. However, it is apparent that dystrophin not only functions in skeletal muscle, but in the CNS as well, with an increased incidence of cognitive impairment and deficits in verbal, short-term and working memory in individuals with DMD. The hippocampus is critical for learning and the formation and consolidation of memories and can be modified by neuroimmune cytokines, such as IL-6. The aim of this programme of research was to investigate the possible contribution of the neuroimmune molecule, IL-6 in cognitive impairment associated with loss of the structural protein, dystrophin.

Initially, we used the dystrophic *mdx* mouse to investigate what impact loss of dystrophin had on normal hippocampal function. Significantly, LTP, the molecular correlate of learning and memory was shown to be decreased in *mdx* mice, which has been linked to memory dysfunction. Moreover, changes in the density and structure of *mdx* hippocampal neurons and glia was observed over time.

Subsequently, we examined the possible effects of IL-6 on hippocampal function in *mdx* mice. IL-6 and IL-6R expression were altered in specific regions in the hippocampus, which underpins function relating to learning and memory. Indeed, *in vivo* blockade of IL-6-mediated signalling in the CNS of *mdx* mice restored LTP, although at this age, it did not impact on learning behaviours.

These studies have elucidated previously unknown cellular and molecular changes in the hippocampus of dystrophin-deficient mice. Moreover, we have linked the neuromodulatory cytokine, IL-6, with the observed pathophysiology.

Chapter 1: Introduction

1.1. Duchenne Muscular Dystrophy

Duchenne Muscular Dystrophy (DMD) is a heritable X-linked recessive disorder which affects approximately 1 in 3,500 live male births and subsequently has one of the highest incidences of all inherited disorders (Emery, 1991). DMD results in the loss of the structural protein, dystrophin (Bulfield *et al.*, 1984), due to mutations in the *DMD* gene (Hoffman & Kunkel, 1989). In DMD-affected boys, the loss of dystrophin becomes apparent from an early age, with muscle weakness progressively leading to physical disability and eventual immobility. DMD patients are usually wheelchair-bound by the age of 12 (Wagner, 2008) and suffer a premature death in their early twenties, often through respiratory failure, due to loss of function in the diaphragm muscle, or cardiac failure (Perloff, 1984).

There is currently no cure for DMD with the most effective treatment being corticosteroids that have strong anti-inflammatory effects and which delay the onset and progression of the disease (Simpson *et al.*, 2016). However, there has been an improvement in the efficacy of other interventions (Reinig *et al.*, 2017) including nocturnal ventilation, routine flu vaccinations, early delivery of antibiotics and physiotherapy (Eagle *et al.*, 2002) which, combined with corticosteroids such as deflazacort, has increased the lifespan of individuals with DMD (Biggar *et al.*, 2006). As such, the life expectancy of patients with DMD has increased to approximately 28.1 years (Landfeldt *et al.*, 2020; Broomfield *et al.*, 2021). More recent therapeutic strategies have targeted the genetic origins of the disease, namely, eteplirsen, an oligonucleotide that facilitates skipping of exon 51 mutations in the

dystrophin gene, which slows but does not halt disease progression (Mendell & Rodino-Klapac, 2016).

1.1. Pathology of loss of dystrophin from muscle

In the absence of the protective function served by dystrophin, skeletal muscle fibres in DMD patients are more susceptible to contraction-induced damage. This ultimately results in an increased efflux of calcium (Ca^{2+}) from the sarcolemma into the sarcoplasm, which in turn, triggers an inflammatory response during the repair of the damaged tissue resulting in muscle fibre phagocytosis and necrosis as muscle fibres lose the ability to regenerate and recover (Anderson *et al.*, 1987; Blake *et al.*, 2002; Davies & Nowak, 2006; Zhou & Lu, 2010).

A sustained increase in intracellular Ca^{2+} results in the activation of apoptotic (Tidball *et al.*, 1995) and necrotic cell death pathways in DMD (Morgan *et al.*, 2018). Cytoplasmic Ca^{2+} levels also influence mitochondrial Ca^{2+} uptake which, in turn, leads to altered metabolism and increased production of reactive oxygen species (ROS). The failure to moderate physiological levels of oxygen radicals results in sustained membrane depolarisation leading to cell death (Feno *et al.*, 2019). Following an increase in cytosolic Ca^{2+} , pro-inflammatory immune mediators are secreted from the contraction-induced damaged muscle fibres (De Paepe & De Bleecker, 2013) resulting in chronic inflammation.

Macrophages are the most abundant of the infiltrating immune cells, although T cells, particularly CD4+ lymphocytes, also predominate (McDouall *et al.*, 1990). Elevated concentrations of pro-inflammatory cytokines such as tumour necrosis factor (TNF) (Abdel-Salam *et al.*, 2009), interleukin (IL)-1 (Evans *et al.*, 2009) and IL-6 (Messina *et al.*, 2011) are also detected at early stages of the disease, with the inflammatory response worsening as DMD progresses. Specifically with regard to IL-6, which is considered to be both pro- and anti-inflammatory, plasma levels of this cytokine are consistently raised in DMD patients (Rufo *et al.*, 2011). In fact, chronic stimulation of an immune response and altered immune mediator profiles, are characteristic features of the progressive DMD muscular phenotype. While DMD has been studied extensively primarily with regard to its effects upon skeletal muscle function/physiology, it is important to note that dystrophin is also expressed in cardiac and smooth muscle, endocrine glands and notably, in neurons, in the peripheral and central nervous systems (Knuesel *et al.*, 2000) and such cognitive dysfunction is another aspect of DMD that needs to be considered.

1.2. Cognitive dysfunction in individuals with DMD

In addition to skeletal muscle pathology discussed above, DMD is characterised by significant cognitive and behavioural problems which include; having lower intelligence quotients (IQs) (Felisari *et al.*, 2000; Nardes *et al.*, 2012), significant cognitive impairment (Bresolin *et al.*, 1994; Anderson *et al.*, 2002), communication difficulties and poor affect recognition (Hinton *et al.*, 2006; Hinton *et al.*, 2007), and

specific deficits in verbal, short-term and working memory (Hinton *et al.*, 2000, 2001; Snow *et al.*, 2013) as well as reading deficits similar to those observed in patients with phonological dyslexia (Dorman *et al.*, 1988; Billard *et al.*, 1992; Castles & Coltheart, 1993). Additionally, there is a higher incidence of attention-deficit/hyperactivity disorder (ADHD), obsessive-compulsive disorder (OCD), autism spectrum disorders (ASD), anxiety disorders (Hendriksen & Vles, 2008; Pane *et al.*, 2013a; Banihani *et al.*, 2015) and epilepsy (Pane *et al.*, 2013a) in individuals with DMD.

Although the aetiology of the cognitive impairment and memory deficits seen in patients with DMD is not well understood, neurons in the hippocampus, cerebral cortex, amygdala and cerebellum all express the full-length isoform of dystrophin (Lidov *et al.*, 1990; Bies *et al.*, 1992; Lidov *et al.*, 1993; Knuesel *et al.*, 2000; Sekiguchi *et al.*, 2009). With a loss of dystrophin linked to neuronal loss, neurofibrillary tangles, alterations to dendritic branching, cortical atrophy as well as metabolic abnormalities in the brains of individuals with DMD (Rosman, 1970; Yoshioka *et al.*, 1980; Jagadha & Becker, 1988; Tracey *et al.*, 1995; Lv *et al.*, 2011).

Further, while a hallmark feature of the muscle pathology in DMD is its progressive nature, there are no longitudinal studies that have shown whether the cognitive pathology in humans with DMD is progressive due to the shortened life expectancy of the disease. Interestingly however, while developmental delay was reported in younger boys, more cerebral atrophy was seen in men with DMD, compared to boys, which may indicate a progression in cognitive dysfunction with age (Yoshioka *et al.*, 1980; Pane *et al.*, 2013b; Doorenweerd *et al.*, 2018).

1.3. Dystrophin Gene

The *DMD* gene is large and complex, comprising of at least five promoters. Three independent promoters located on the 5'-end of the *DMD* gene, namely the B(brain) - M(muscle) - and P- (Purkinje cell) promoters, regulate the expression of three full-length isoforms of dystrophin (Dp427) as well as numerous smaller dystrophin proteins, namely, Dp260, Dp140, and Dp116, Dp71, Dp40 (Gorecki *et al.*, 1992; Ahn & Kunkel, 1993), which is illustrated in Figure 1.1 below. Most of these smaller proteins are present in both the central and peripheral nervous systems.

In skeletal muscle, the full-length dystrophin protein has a unique N-terminal amino acid sequence and referred to as Dp427m (muscle) (Howard *et al.*, 1999). In neurons, the start sequence of Dp427m is replaced by a unique N-terminal amino acid sequence, and called Dp427c (cortex) (Nudel *et al.*, 1989). In Purkinje cells of the cerebellum, a unique N-terminal amino acid sequence is expressed, named Dp427p (Purkinje) (Gorecki *et al.*, 1992). All of these full-length dystrophin proteins comprise a central rod domain containing 24 spectrin-like repeats.

Importantly, all isoforms of dystrophin have different locations of expression.

Dp260 is a C-terminal isoform of dystrophin, the formation of which is induced by an alternative promoter located on intron 29 of the *DMD* gene (D'Souza *et al.*, 1995), containing 15 spectrin-like rod repeats in the rod domain, is mainly expressed in retinal tissue. Dp140 is another C-terminal dystrophin isoform formed by the action of another alternative promoter on intron 44 of the *DMD* gene,

contains six spectrin-like rod repeats, and is mainly expressed during foetal brain development and thus is associated with a higher risk of cognitive impairment in DMD and Becker Muscular Dystrophy (BMD) (Chamova *et al.*, 2013). Dp116 is an alternative C-terminal dystrophin short isoform, specific to Schwann cells in peripheral nervous system (Byers *et al.*, 1993). Dp71 is a ubiquitous isoform of dystrophin expressed with an alternative promoter located in intron 62 and is expressed in the brain, namely the hippocampus and is associated with cognitive deficits, specifically learning and memory (Rapaport *et al.*, 1992; Waite *et al.*, 2012). Lastly, Dp40 is the shortest known isoform of dystrophin expressed in hippocampal neurons and is known to interact with various presynaptic proteins (Tozawa *et al.*, 2012; Fujimoto *et al.*, 2014).

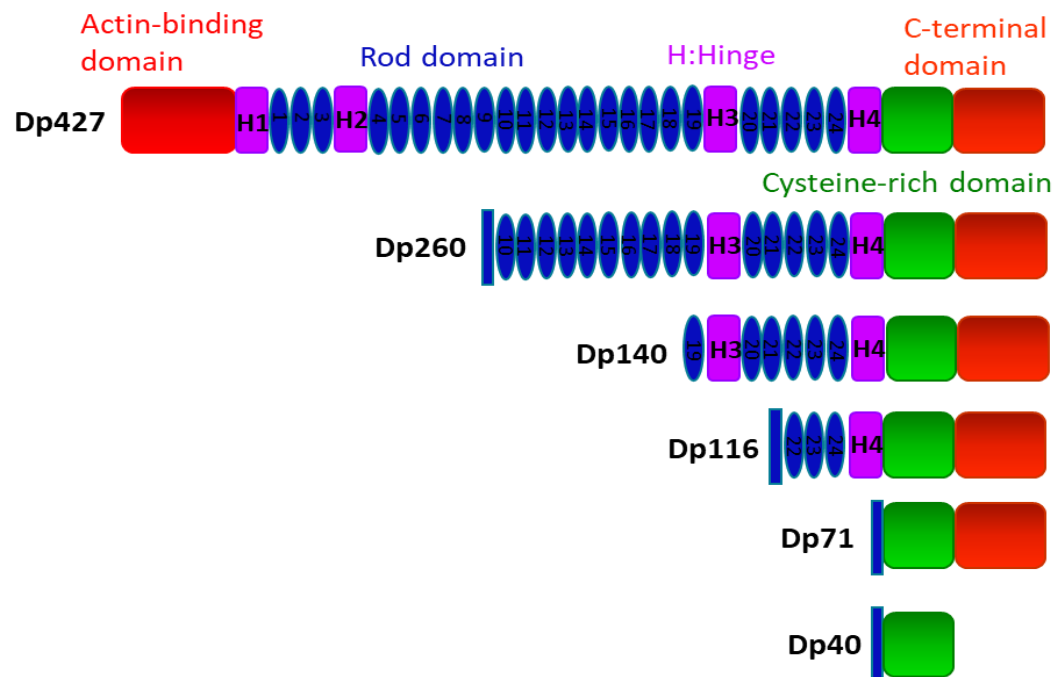


Figure 1.1: Dystrophin protein isoforms present in DMD (adapted from (Nichols et al., 2015). This figure illustrates the different isoforms of the dystrophin protein that are present in different regions of the body. Specifically, Dp427 in muscle and the brain, Dp260 in the retina, Dp116 in Schwann cells in the peripheral nervous system, Dp140, Dp71 and Dp40 in the brain.

1.4. Dystrophin-Associated Protein Complex (DAPC) in myocytes and brain tissue

In muscle, dystrophin forms part of the dystrophin-associated protein complex (DAPC) that functions as an essential link connecting the sarcolemmal cytoskeletal actin to the extracellular matrix (Ervasti & Campbell, 1993; Blake *et al.*, 2002; Davies & Nowak, 2006). This structural function of dystrophin seems to stabilize the sarcolemma and subsequently protect it from any disruption during muscle contraction (Zubrzycka-Gaarn *et al.*, 1988). The DAPC has a dual function in that it supplies mechanical support to the sarcolemma as well as communicating contractile force between muscle fibres, suggesting that the DAPC is also involved in transmembrane signalling (Lapidos *et al.*, 2004). The DAPC maintains physiological handling of Ca^{2+} in muscle fibres as the DAPC acts as an anchor for numerous signalling proteins (Constantin, 2014). The DAPC also provides a strong mechanical link between the sarcolemmal cytoskeleton and the extracellular matrix (thereby mediating interactions between them) in skeletal muscle as it consists of cytoplasmic, transmembrane and extracellular proteins (Rando, 2001). The DAPC is made up of three distinct components: (i) α - and β -dystroglycan, (ii) the sarcoglycan complex (SGC), and (iii) the cytoplasmic subcomplex which is composed of dystrophin, the dystrobrevins and the syntrophin protein family (Blake *et al.*, 2002), as illustrated in Figure 1.2 below.

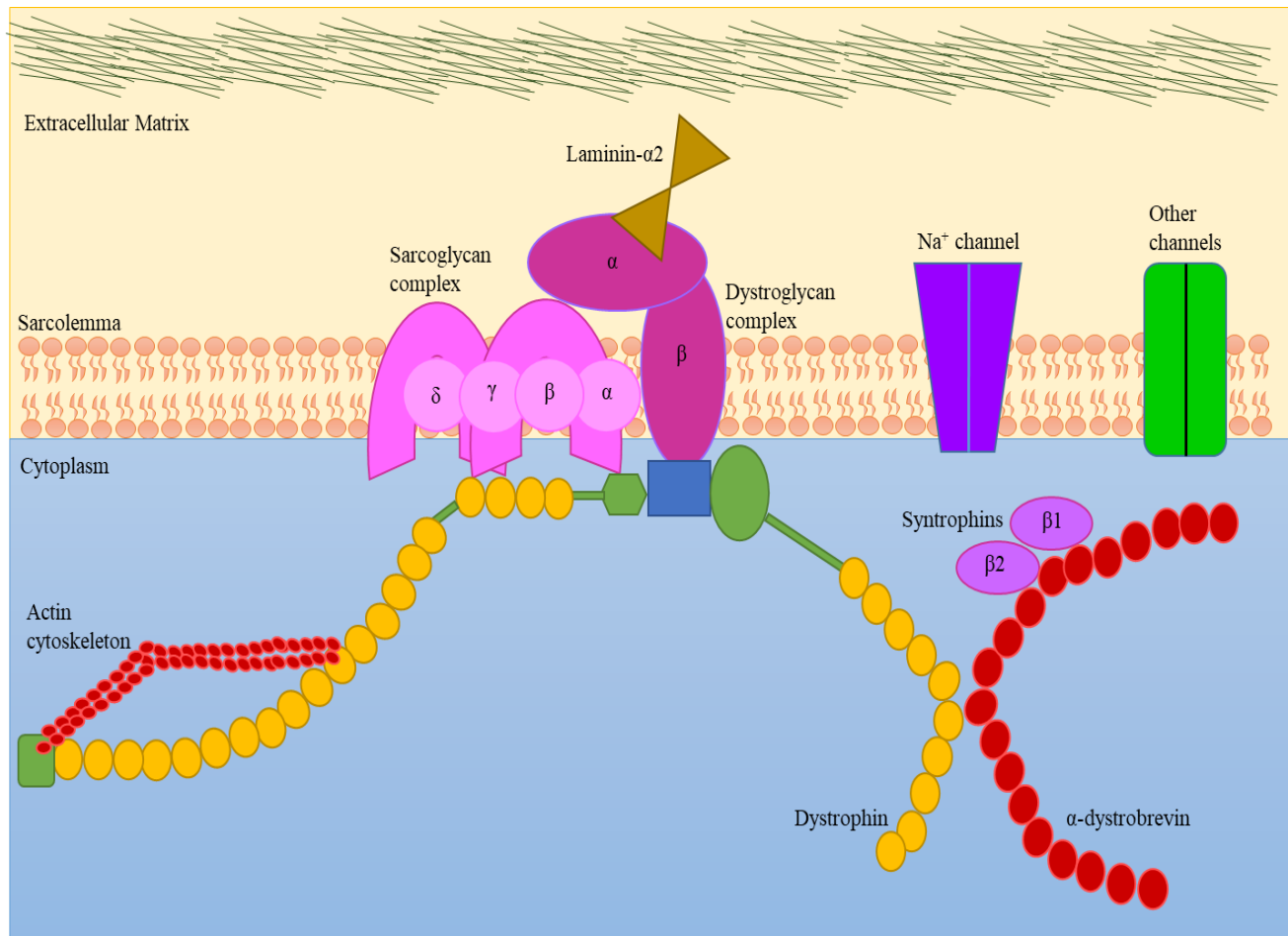


Figure 1.2: Organisation of the Dystrophin-Associated Protein Complex (DAPC) in muscle (adapted from (Fairclough et al., 2013). There are the three distinct components that make up the DAPC: (i) α - and β -dystroglycan, (ii) the sarcoglycan complex (SGC), and (iii) the cytoplasmic subcomplex, which is composed of dystrophin, the dystrobrevins and the syntrophin protein family. Dystrophin provides a crucial structural link between the sarcolemmal cytoskeleton and the extracellular matrix in muscle.

1.5. The role of Dystrophin in the brain

In cells of the central nervous system (CNS), dystrophin and other members of the DAPC partake in macromolecular assemblies that play a role in anchoring receptors to dedicated sites within the membrane (Waite *et al.*, 2009). Furthermore, full length dystrophin, Dp427, co-localizes with γ -aminobutyric acid type A (GABA)_A receptors and is involved in the postsynaptic clustering and stability of inhibitory GABAergic synapses (Knuesel *et al.*, 1999; Anderson *et al.*, 2002) in the postsynaptic membrane of cortical and cerebellar Purkinje neurons (Lidov *et al.*, 1990; Kim *et al.*, 1992; Brunig *et al.*, 2002; Waite *et al.*, 2009). This type of clustering and stabilisation of GABAergic synapses is necessary for effective synaptic signal transduction (Vaillend & Billard, 2002; Kueh *et al.*, 2008).

Importantly, there is evidence that distinct and heterogeneous complexes of the DAPC are present in neurons (Blake *et al.*, 1999; Moukhles & Carbonetto, 2001). Meaningfully, the DAPC in neurons differs from that in muscle in that laminin is replaced by neurexin- α and GABA_A receptors, which are stabilised in the cellular membrane by the DAPC. Further, dystrophin associates with β -dystrobrevin, which plays a role in postsynaptic densities (PSDs) rather than α -dystrobrevin as in skeletal muscle (Blake *et al.*, 1999). These differences in structure of the DAPC in neurons is illustrated in Figure 1.3. Interestingly, defects in the processing and assembly of the DAPC have been associated with brain abnormalities such as mild cognitive impairment and neuronal migration disorders (Blake *et al.*, 1999; Waite *et al.*, 2012), further highlighting the possible impact that a loss of dystrophin, or its

associated proteins, might have on the brain and its normal function. Importantly, in patients with DMD, neuronal loss, neurofibrillary tangles, abnormalities in dendritic development, as well as alterations in the morphology of the presynaptic neuronal cytoskeleton, have been reported (Yoshioka *et al.*, 1980; Jagadha & Becker, 1988; Sogos *et al.*, 1997; Anderson *et al.*, 2002; Lv *et al.*, 2011), which could be the result of altered DAPC function.

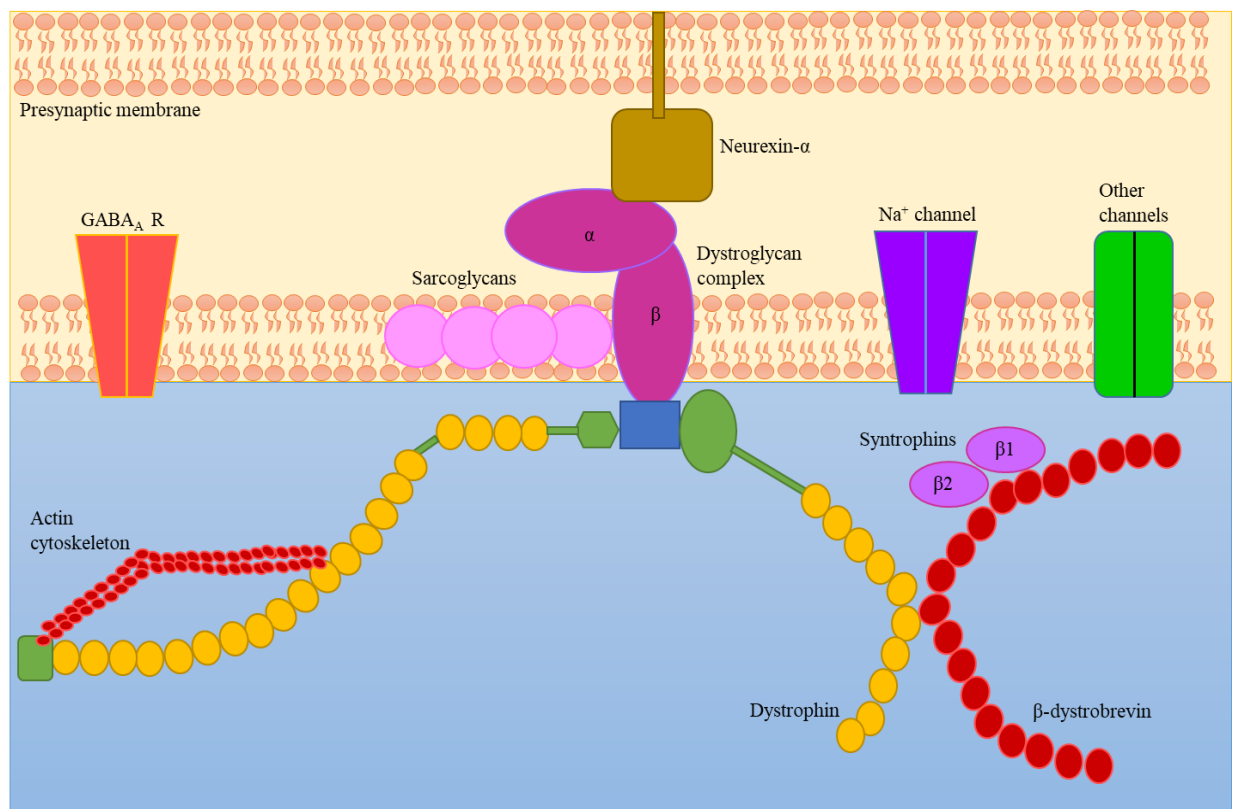


Figure 1.3: The molecular organization of dystrophin and the DAPC in neurons (adapted from (Waite et al., 2009). The DAPC in neurons differs from that in muscle in that laminin is replaced by neurexin- α and GABA_A receptors which are stabilised in the cellular membrane by the DAPC.

1.6. *Mdx* mouse model of DMD

The *mdx* mouse has been used as a pre-clinical model of DMD for more than 30 years, since it arose due to a spontaneous mutation in the premature stop codon of exon 23 in the *DMD* gene (Arechavala-Gomez *et al.*, 2010) of the C57BL/10ScSnJ mouse (Bulfield *et al.*, 1984). The *mdx* mouse is genetically equivalent to DMD in humans and has proven to be advantageous for the investigation of the consequences of the loss of dystrophin on multiple physiological systems. However, despite certain similarities, there are differences between the pathophysiology of DMD in humans compared to that seen in *mdx* mice.

For example, whilst loss of dystrophin in humans and *mdx* mice both result in contraction-induced damage and loss of sarcolemmal integrity in skeletal muscle, the severity in pathology is dependent upon age in *mdx* mice (Moens *et al.*, 1993). This difference in the mouse model is due to the presence of another protein, utrophin, which shares some structural and functional traits with dystrophin such as the fact that it can bind and interact with the same proteins within the cytoskeleton as dystrophin (Matsumura *et al.*, 2011), resulting in a less severe phenotype than in the human disease.

Normally, utrophin is down-regulated in muscle when dystrophin is present (Blake *et al.*, 1996). However, in the absence of dystrophin, utrophin is able to carry out the same function(s) that dystrophin normally would (Keep, 2000). Consequently, in *mdx* mice, the upregulation of utrophin results in the regeneration of sarcolemmal

and plasma membrane integrity, thus saving muscle from deterioration, resulting in a less severe DMD phenotype with regard to skeletal muscle function (Gilbert *et al.*, 1999). In human DMD muscle however, despite there being an upregulation of utrophin relative to unaffected individuals, the levels present in the sarcolemma are not sufficient to prevent the progression of the disease and the subsequent degradation of muscle (Love *et al.*, 1989). Interestingly, the progressive deterioration of the diaphragm of *mdx* mice is similar to that of humans with DMD, with widespread fibrosis present in *mdx* mice from 12 weeks of age until death (Stedman *et al.*, 1991).

While the *mdx* mouse shows differences in skeletal muscle to the DMD profile in humans, there are similarities between humans with DMD and *mdx* mice with regard to dystrophin in the brain. Although no major alterations in gross brain structure have been identified in *mdx* mice relative to controls (Miranda *et al.*, 2009), *mdx* mice do possess fewer neurons, as well as neural shrinkage, in the cerebral cortex and brainstem (Sbriccoli *et al.*, 1995; Carretta *et al.*, 2001). Other architectural changes, as well as reductions in dendritic branching and condensed spinal density in corticospinal neurons of *mdx* mice have also been described (Sbriccoli *et al.*, 1995; Carretta *et al.*, 2003; Carretta *et al.*, 2004). Furthermore, loss of Dp427 appears to be associated with both an increase in the number of inhibitory synapses between axons and dendrites, and alterations in the arrangement of PSDs, in the CA1 region of the hippocampus in *mdx* mice (Miranda *et al.*, 2009). Further, in the *mdx* mouse model of DMD, a study by Bagdatlioglu *et al.*, 2020, showed that hippocampal spatial learning and memory was decreased in

mdx mice, particularly long-term memory, which progressively worsened with age (Bagdatlioglu *et al.*, 2020).

1.7. The role of the hippocampus in memory formation

In the brain, the hippocampus is a paired structure present within the limbic system. The hippocampus forms part of the hippocampal formation, which is comprised of the entorhinal cortex, dentate gyrus (DG) and cornu ammonis (CA) regions (CA1–CA3). The hippocampus facilitates the formation of new memories, mainly declarative memory (Vargha-Khadem *et al.*, 1997; Tulving & Markowitsch, 1998) and has also been shown to play a role in spatial navigation and spatial memory (Snow *et al.*, 2013). The main cell type present in the hippocampus are pyramidal cells, which reside in regions CA1-CA3, whereas within the DG, granule cells are most abundant.

One way in which memory formation and consolidation have been described experimentally is through the phenomenon of synaptic long-term potentiation (LTP) that can be induced in excitatory tri-synaptic circuit that characterises the hippocampal formation (see Figure 1.4). There is evidence that LTP is the molecular mechanism underlying memory formation (Morris *et al.*, 1986; Bliss & Collingridge, 1993; Izquierdo *et al.*, 2007) and, more specifically, that it plays a role in hippocampal spatial learning in rodents (Morris *et al.*, 1986; Martin *et al.*, 2000; Morris *et al.*, 2003). The induction of LTP results in a long-lasting enhancement of

synaptically-evoked potentials (Bliss & Lomo, 1973; Bliss & Collingridge, 1993). The majority of LTP experimentation has been conducted upon circuits within the hippocampus, with the first LTP studies focussed upon the dentate gyrus (Bliss & Lomo, 1973) (although it is now clear that LTP can be elicited within many difference areas of the brain, including the amygdala and cerebellum (Artola *et al.*, 1990; Bliss & Collingridge, 1993)).

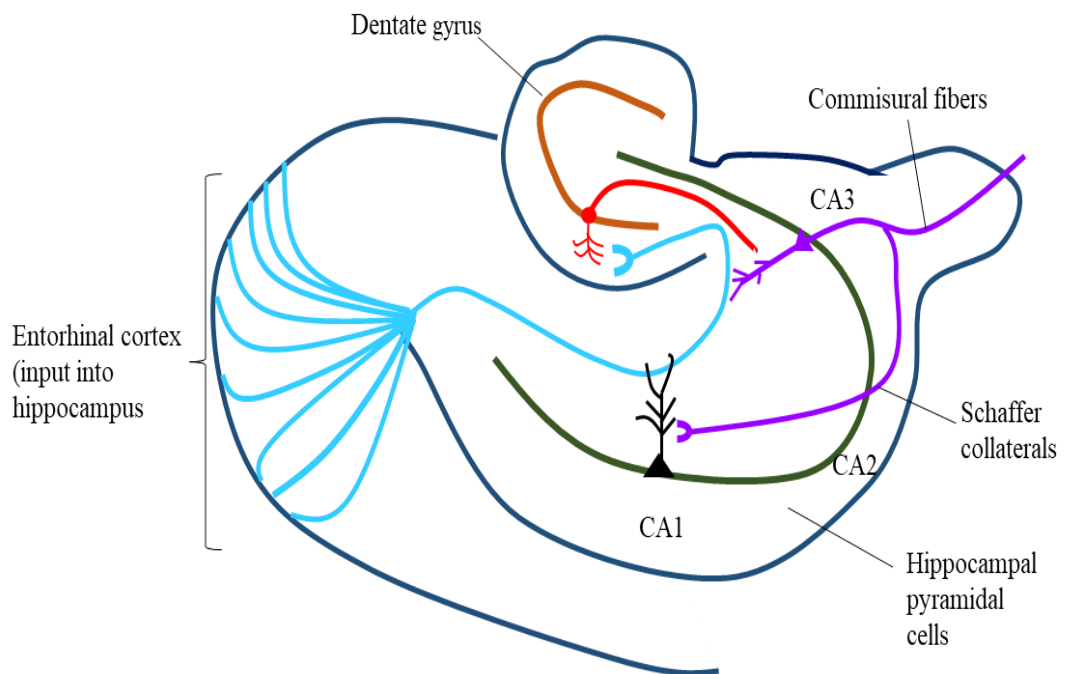


Figure 1.4: Schematic representation of the hippocampal formation, which is comprised of the dentate gyrus, CA1, CA2 and CA3 regions. Most LTP experiments have investigated synaptic plasticity processes that occur between the Schaffer collaterals (purple) and dendrites of CA1 pyramidal cells. Adapted from (Hitti & Siegelbaum, 2014).

In the hippocampus, most experiments investigating LTP have focussed upon the synapses connecting the Schaffer Collaterals (SC) (running from area CA3) to the dendrites of CA1 pyramidal cells, which reside within the *stratum radiatum* above

the hippocampal fissure (Malenka & Nicoll, 1999). When SC axons are electrically stimulated at a basal frequency (*e.g.*, 1 pulse every 20 seconds), this induces the release of the primary excitatory neurotransmitter within the CNS, glutamate, from pre-synaptic terminals of the CA3 axons. The released glutamate then binds to, and opens, a type of ionotropic glutamatergic receptor, known as an AMPA receptor (AMPA), on CA1 dendrites, evoking excitatory post-synaptic potentials (EPSPs) (Collingridge *et al.*, 1983; Muller *et al.*, 1988; Bliss & Collingridge, 1993). Following the induction of LTP at these same synapses using a high frequency stimulus (*e.g.* “theta-burst stimulation” (Larson *et al.*, 1986), several bursts of 4 pulses at 100Hz), when normal basal stimulation is resumed, EPSP amplitude is significantly enhanced for a prolonged period after application of the LTP stimulus (Bliss & Collingridge, 1993). LTP at the SC-CA1 synapses is induced due to the activation of glutamatergic N-methyl-D-aspartate (NMDA) receptors, which are activated by the release of glutamate (Bliss & Collingridge, 1993). The activation of these NMDA receptors are required for LTP to occur (Collingridge *et al.*, 1983) and this is likely due to an increase in the permeability of the NMDA receptors to calcium, after a release of a magnesium ion and the subsequent influx of calcium during LTP (Ascher & Nowak, 1988). After the subsequent elevated Ca^{2+} influx through activated NMDARs, this results in calmodulin-dependent activation of Ca^{2+} /calmodulin-dependent protein kinase (CaMKII) (Lisman *et al.*, 2002), where its function is to phosphorylates the GluR1 subunit in the AMPA receptor which results in increased AMPA conductance and potentiates synaptic transmission (Lisman *et al.*, 2002). As such, LTP results in changes to the post-synaptic synapse and increased synaptic transmission. Importantly, this potentiation is input-specific. That is, the increase in

EPSP amplitude is only observed at the synapses subjected to the LTP stimulus but not at other SC-CA1 synapses (untouched by the LTP stimulus).

The maintenance of LTP beyond 1 to 3 hours in the hippocampus requires de novo protein synthesis (Frey & Morris, 1997). When solely posttranslational modifications are reflected, this is termed early LTP (E-LTP). The duration of E-LTP varies considerably across studies, mainly because conditioning protocols differ in their abilities to produce the relevant posttranslational changes, such as the phosphorylation of synaptic proteins. The stable, protein synthesis-dependent phase is called late LTP (L-LTP); however, this phase can occur relatively early in LTP, newly synthesized proteins to LTP can be detected as early as 20 min after high frequency stimulation with the use of protein synthesis inhibitors (Osten et al., 1996).

Through the ability to experimentally induce LTP in the hippocampus (and several other brain regions), we now have some understanding as to how the hippocampus functions in relation to memory formation and consolidation (Morris *et al.*, 1986; Teyler & DiScenna, 1987; Bliss & Collingridge, 1993; Izquierdo *et al.*, 2007). LTP induction at the post-synaptic neuron can be seen in figure 1.5 below.

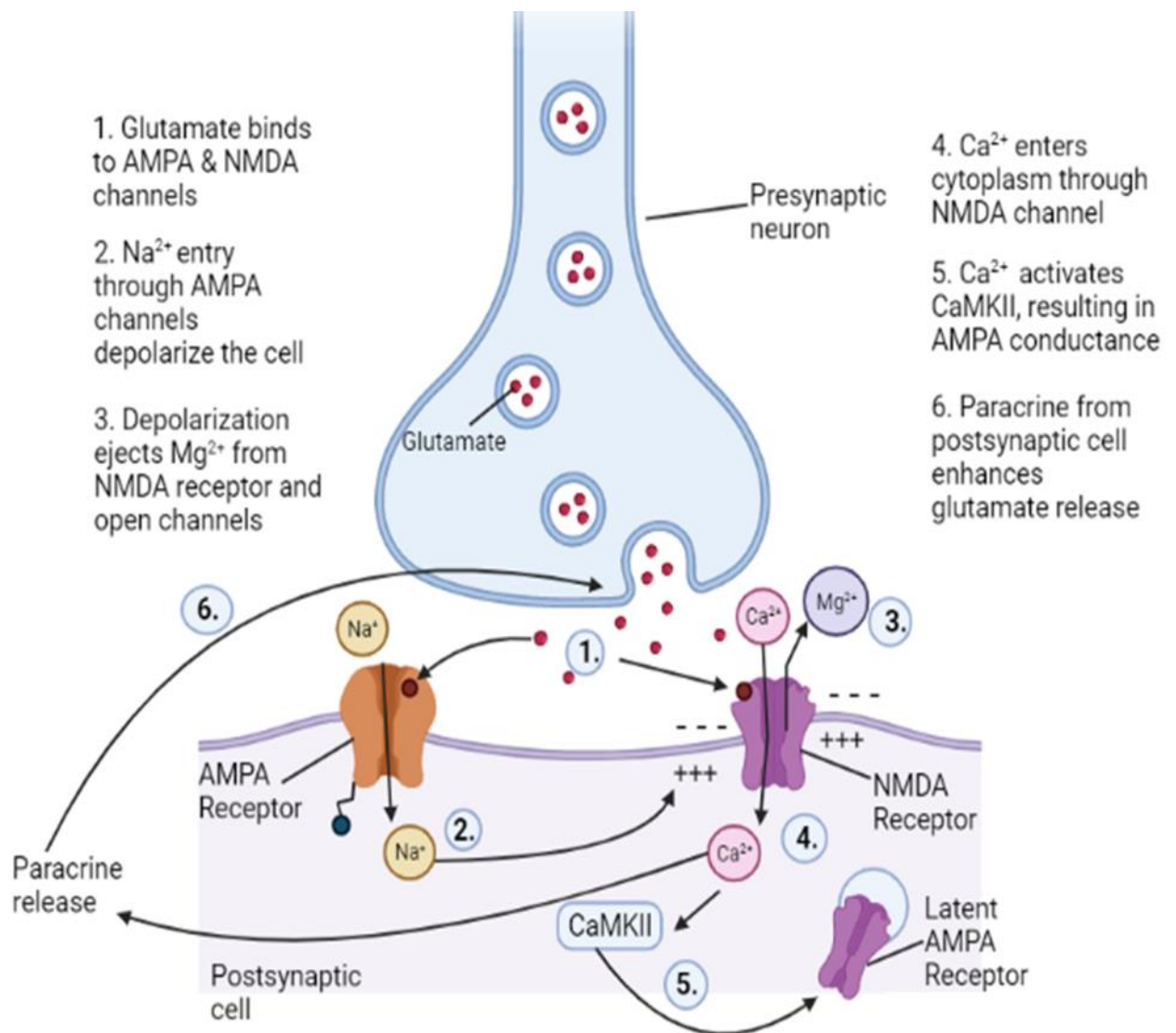


Figure 1.5: A visual representation of the induction of LTP in the post synaptic cell – adapted from (Unglaub Silverthorn, 2013). During normal synaptic transmission, glutamate is released from the presynaptic bouton and acts on both AMPARs and NMDARs. However, Na^+ flows only through the AMPA receptor, but not the NMDA receptor, because Mg^{2+} blocks the channel of the NMDA receptor. Depolarization of the postsynaptic cell relieves the Mg^{2+} block of the NMDA receptor channel, allowing Na^+ and Ca^{2+} to flow into the dendritic spine by means of the NMDA receptor. The resultant rise in Ca^{2+} within the dendritic spine is the critical trigger for LTP.

1.8. Hippocampal dysfunction in the *mdx* mouse

Dystrophic *mdx* mice display a neurological and behavioural profile that is similar to DMD patients (Muntoni *et al.*, 1991; Chausseot *et al.*, 2015) as mentioned in section 1.6. Further, they display a poor capacity to learn new tasks, and deficits in spatial memories (Vaillend *et al.*, 1995; Vaillend *et al.*, 2004; Chausseot *et al.*, 2015) which are consistent with dysfunctions of the hippocampus. They also display deficits in associative learning and in the general processes involved in memory formation and consolidation which are both dependent upon optimal functioning of the hippocampus (Chausseot *et al.*, 2015).

Changes in short- and long-term memory formation and consolidation may be linked with altered synapse morphology and plasticity associated with a loss of Dp427 (Miranda *et al.*, 2009). However, electrophysiological studies conducted on *mdx* mouse brain slices in order to elucidate the role dystrophin might play in synaptic function have generated contradictory findings. For example, initial studies found no obvious deficit in hippocampal NMDA-dependent LTP in *mdx* mice that lacked Dp427 relative to controls (Sesay *et al.*, 1996). Furthermore, no differences in hippocampal-dependent spatial learning tasks were found between *mdx* mice and wild-type controls (Sesay *et al.*, 1996), implying that hippocampal synaptic processes remained intact in *mdx* mice. However, subsequent studies have described deficits in long-term hippocampal spatial memory (Vaillend *et al.*, 2004; Bagdatlioglu *et al.*, 2020), which progressively worsened with age (Bagdatlioglu *et al.*, 2020), as well as deficits in object recognition (Vaillend *et al.*, 2004) in *mdx* mice

relative to controls. Further, hippocampal LTP in *mdx* mice was also abnormally enhanced relative to controls (Vaillend *et al.*, 2004). Although the reason(s) for these contrasting findings remain unclear, one possibility is that the particular training procedures mice were subjected to by the Sesay group may at least partially account for them as the tasks might have helped to alleviate any subtle memory deficits that may have been present before training. Vaillend *et al.* (2004) proposed that the abnormally enhanced hippocampal LTP observed in the *mdx* brain slices could at least, in part, be responsible for the memory deficits seen in the *mdx* mouse. Thus, somewhat counterintuitively, an abnormal enhancement of synaptic plasticity and neuronal excitability might actually inhibit normal cognitive functioning (Vaillend *et al.*, 2004).

Later studies found that the *mdx* brain dysfunction was associated with a loss of Dp427 that resulted in defective synaptic inhibition (Perronnet & Vaillend, 2010) possibly due to Dp427-dependent reduction in the number of GABA_A receptor clusters, which are essential for normal neuronal excitability and synaptic transmission (Dallerac *et al.*, 2011), in the hippocampus (as well as in other brains regions such as the cerebellum and amygdala (Vaillend *et al.*, 2010)). A reduction in GABA_A receptor clusters was also seen in cerebellar Purkinje and hippocampal CA1 neurons of the *mdx* mouse (Knuesel *et al.*, 1999).

It is notable that when Vaillend *et al.* (2010) restored expression of a dystrophin-like protein to 15–25% of normal dystrophin levels in area CA1 of the hippocampus of *mdx* mice (using an AAV vector expressing antisense sequences to modulate

splicing of the dystrophin pre-mRNA), this facilitated a complete recovery of $\alpha 2$ subunit-containing GABA_A-receptor clustering in this brain region of *mdx* mice (Vaillend *et al.*, 2010). This highlights the importance of the role of Dp427 and its association with GABA_A receptors in modulating hippocampal synaptic plasticity. Additionally, using the same treatment, NMDA-dependent LTP was also restored (Vaillend *et al.*, 2010).

1.9. Intracellular Ca²⁺ handling in dystrophic hippocampal neurons

Calcium (Ca²⁺) has a significant role in cell signalling such that many cellular functions are directly or indirectly controlled by the intracellular concentration of Ca²⁺ (Missiaen *et al.*, 2000).

Similarly, in the brain, Ca²⁺ plays an important role in the regulation of numerous neuronal processes such as neuronal plasticity and survival, as well as transcription factor regulation (Brini *et al.*, 2014) and is tightly regulated by complex regulatory mechanisms which balance the influx of Ca²⁺ into the cell and release from intracellular stores (Brini *et al.*, 2014). Intracellular free Ca²⁺ concentration is involved in the control of many physiological functions in all cell types, including neurons. An influx of Ca²⁺ through voltage-dependent and ligand-gated channels in the plasma membrane is a critical signal for the release of neurotransmitters from presynaptic terminals and for responses of the postsynaptic neuron (Yuste *et al.*, 2000; Burnashev & Rozov, 2005; Hartmann & Konnerth, 2005). Ca²⁺ can also play a role in synaptic plasticity which depends on NMDAR activation (Mattson, 2007).

Any disturbance of the normal regulation of intracellular Ca^{2+} (such as an increase in Ca^{2+} from intracellular stores) can result in pathology (Missiaen *et al.*, 2000).

Interestingly, in DMD, there is evidence that a lack of dystrophin is associated with irregular intracellular Ca^{2+} homeostasis. For example, an absence of dystrophin in neurons has been associated with an increase in the open time of Ca^{2+} permeable ion channels in cerebellar neurons in *mdx* mice (Haws & Lansman, 1991), which could ultimately trigger a neuronal cytotoxic cascade resulting in neuronal death.

Most recently, a study by Lopez *et al.* (2018), found that intracellular Ca^{2+} was elevated in cortical and hippocampal pyramidal neurons of adult *mdx* mice compared to WT animals and this increase in Ca^{2+} progressed with age (Lopez *et al.*, 2018). It is notable that the cognitive deficits shown by these *mdx* mice were partially reversed with the injection of GsMTx-4 which blocks stretch-activated channels that are involved in Ca^{2+} regulation (Suchyna *et al.*, 2000; Lopez *et al.*, 2018). Thus, with an injection of GsMTx-4 into *mdx* mice, this resulted in a decrease of intracellular Ca^{2+} in cortical and hippocampal neurons and improvement in spatial learning memory compared to WT mice (Lopez *et al.*, 2018). Interestingly, dystrophin deficiency dysregulated hippocampal intracellular calcium levels to a greater extent in cortical neurons in an age-dependent manner (Lopez *et al.*, 2018). While dystrophin-deficiency undoubtedly impacts upon cognitive function, with regards to intracellular Ca^{2+} (Lopez *et al.*, 2018) and GABA_A-receptor clustering (Vaillend *et al.*, 2010), this may be due entirely to its role as a structural protein. However, another consideration, following with the pathogenesis of DMD, could be to investigate inflammation. DMD is known to result in altered cytokine profiles, thus, we may speculate that this altered cytokine profile may also mediate some of

the negative cognitive effects observed in human DMD patients, for the reasons I will discuss in the next section(s).

1.10. Interleukin-6 is a pleiotropic cytokine with modulatory actions in the central nervous system (CNS).

Cytokines are traditionally described as being immune molecules that are critical for pro- or anti-inflammatory signalling. Of these, IL-6 has pleiotropic effects, on many cells. In health, plasma levels of IL-6 range from ~1–10 pg/ml (Akira *et al.*, 1993) but this can rise to the lower nanogram/ml range during infection or inflammation (Jones, 2005). IL-6 can be secreted by both immune and non-immune cells (Marz *et al.*, 1998), and has a key role in regulating immune cell function (Hirano *et al.*, 1985), although this 26kDa protein can also modify functions in other cells, including neurons.

IL-6 evokes its biological effects by binding to IL-6 receptors, which can either be soluble or membrane bound. However, membrane-bound IL-6 receptors (IL-6R α) are only expressed in a restricted subset of cells (Rothaug *et al.*, 2016). Whereas soluble IL-6 receptors may form a complex with IL-6 and stimulate distant cells including neurons (Rose-John & Heinrich, 1994; Erta *et al.*, 2012).

1.10.1. Classical signalling mechanism of IL-6 and its receptor

The classical signalling mechanism of IL-6 entails the binding of IL-6 to membrane bound IL-6R α in a complex with glycoprotein (gp)130. This binding results in the homodimerization of the signal transducing protein gp130 and subsequent

phosphorylation of Janus kinase/signal transducer and activator of transcription (JAK/STAT) (Darnell *et al.*, 1994). Not only does the complex of IL-6/ IL-6R α activate STATs, it also activates the mitogen-activated protein (MAPK) and phosphoinositide 3-kinase (PI 3-K) cascades (Eulendorf *et al.*, 2012). IL-6 can, however, also evoke signalling in cells which don't express IL-6R α on their cell membranes.

1.10.2 Trans-signalling mechanism of IL-6 and its receptor

IL-6R α can be proteolytically cleaved from the cell membrane to generate a soluble IL-6R α (sIL-6R α), which is still able to bind to its ligand, IL-6, as seen in classical signalling (Müllberg *et al.*, 1993). And similar to the classical signalling pathway, sIL-6R α /IL-6 complex binds to gp130 expressed on the cell membrane of neurons (Rose-John & Heinrich, 1994; Scheller *et al.*, 2011b) to evoke this trans-signalling mechanism. In humans, sIL-6R α can also be generated by translation of an alternatively spliced mRNA (Lust *et al.*, 1992), although this has not been seen in mice (Scheller *et al.*, 2011a).

Even though a membrane bound receptor for IL-6 is limited to a subset of cells, gp130 is ubiquitously expressed on the cellular membrane of most cell types including neural cells in the CNS (Watanabe *et al.*, 1996). Expression of gp130 is not changed during inflammation but concentrations of sIL-6R α can increase between two and five-fold (Rose-John, 2012; Schaper & Rose-John, 2015), inferring the importance of sIL-6R in the activation of IL-6. Importantly, downstream activation of JAK/STAT or PI 3-kinase occurs in cells that don't express membrane-bound IL-

6R α in a similar manner to classical IL-6R α signalling (Ernst & Jenkins, 2004; Campbell *et al.*, 2014). Thus, as many cells respond to IL-6 during inflammation, which do not express membrane-bound IL-6Rs, the importance of sIL-6R for the activation of IL-6 is evident. See Figure 1.5 below for a visual representation of classical and trans-signalling mechanisms of IL-6.

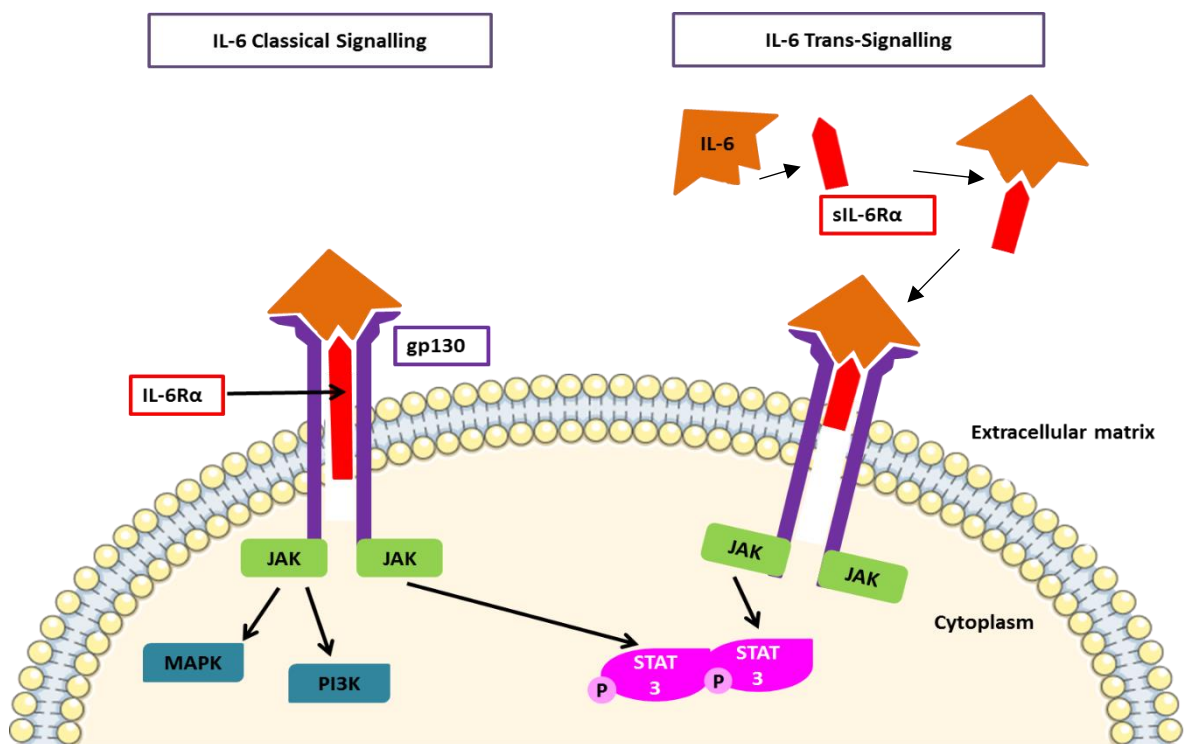


Figure 1.6: A representation of IL-6/ IL-6R α classic and trans-signalling mechanisms – adapted from (Schaper & Rose-John, 2015). Classical signalling involves IL-6 binding to a membrane bound IL-6R α , with this complex subsequently binding to, and activating, the signal transduction of gp130 for downstream activity of JAK/STAT, MAPK and PI3K. Trans-signalling involves IL-6 binding to a soluble IL-6R α in the extracellular matrix and only then binding to gp130 for activation of downstream signalling.

Importantly, differences in the evoked cellular response between classic receptor-mediated IL-6 signalling and trans-signalling can be detected by the cellular outcomes. Classical IL-6 signalling evokes anti-inflammatory signals, in contrast to the pro-inflammatory, and potentially deleterious, actions of IL-6/sIL-6R-evoked trans-signalling (Garbers *et al.*, 2015).

The signalling kinetics of these two distinct mechanisms also differ, leading to fundamentally different cellular responses (Rothaug *et al.*, 2016). In the CNS, neuronal degeneration evoked by IL-6 is mediated by trans-signalling, whereas membrane receptor-mediated IL-6 signalling plays a role in neural tissue regeneration (Campbell *et al.*, 2014) and neuronal protection (Garbers *et al.*, 2015). Understanding the specific signalling mechanisms is particularly relevant when therapeutically targeting this signalling pathway in neurological disorders.

1.11. IL-6 modifies long-term potentiation

Ex vivo hippocampal slices retain the hippocampal architecture and intact neural pathways of the dentate gyrus, CA1 and CA3 areas (Malenka & Nicoll, 1999). As explained in an earlier section, hippocampal LTP is believed to recapitulate the molecular mechanisms underlying the coding of new memories in the brain by modifying the strength of synaptic transmission (Collingridge *et al.*, 1983; Muller *et al.*, 1988; Bliss & Collingridge, 1993).

IL-6 gene expression in the CA1 region of the hippocampus is increased after induction of LTP, although it is important to note that expression of IL-6 mRNA was

specific to nearby astrocytes, not neurons (Jankowsky *et al.*, 2000). Significant increases in the levels of IL-6 mRNA one hour post-stimulation were also noted in *ex vivo* experiments, but only when LTP had been successfully induced (Balschun *et al.*, 2004). Interestingly, blockade of the effects of endogenous IL-6 production using a neutralizing antibody after 90 minutes resulted in a remarkable prolongation of LTP, as well as improved performance in spatial memory tests in mice, further emphasising the pleiotropic actions of this cytokine in the CNS (Balschun *et al.*, 2004). However, it appears that either too little or too much IL-6 may impair memory formation, as IL-6 knock-out mice performed less well in hippocampus-dependent memory tests (Baier *et al.*, 2009) and are more susceptible to hippocampal damage following induced convulsions relative to controls (Penkowa *et al.*, 2001; De Sarro *et al.*, 2004), whereas transgenic mice with cerebral over-expression of IL-6 were characterised by suppressed LTP in the dentate gyrus region (Bellinger *et al.*, 1995).

Although it is recognised that exogenous application of IL-6 may not mimic endogenously released IL-6 during LTP (Balschun *et al.*, 2004), application of IL-6 still inhibited hippocampal LTP using a mechanism that was dependent upon IL-6R activation (Tancredi *et al.*, 2000). Moreover, IL-6-induced LTP inhibition was attenuated by administration of monoclonal antibodies, used to block IL-6 receptors (Li *et al.*, 1997). Thus, IL-6 may fine-tune memory consolidation by acting as a molecular filter, limiting the long-term storage of some types of information.

1.12. IL-6 modifies hippocampal adult neurogenesis and cognitive function

The posterior or dorsal portion of the hippocampus participates in contextual learning, declarative memory and spatial navigation (Kheirbek, 2013, Moser & Moser, 1998). Whereas the ventral portion of the hippocampus is thought to be involved in emotional processing, such as anxiety (Bannerman et al., 2003, (Bannerman et al., 2004, Bannerman et al., 2014). The dentate gyrus is one of the few brain regions where neurogenesis occurs and adult-born neurons are integrated into pre-existing neuronal circuits (Gage, 2000). The process of hippocampal adult neurogenesis is initiated when neuronal progenitor cells from the subgranular zone of the hippocampal dentate gyrus migrate into the granule cell layer and subsequently differentiate into neuronal or glial cells (Gage, 2000). Newly generated neurons then extend their axons into the CA3 region of the hippocampus and eventually form synapses with adjacent cells (Gould *et al.*, 2000; van Praag *et al.*, 2002). Integration of new cells into neuronal circuits is proposed to contribute to hippocampal-dependent learning and memory processes (Wu & Hen, 2014). The dentate gyrus is positioned as the input zone of the hippocampus and has low neuronal activity with upstream cortical areas. Interestingly, IL-6 has been shown to influence neural stem cells in neurogenesis (Islam *et al.*, 2009), with proliferation, survival and differentiation of hippocampal progenitor cells being compromised in the dentate gyrus when IL-6 concentrations are elevated (Vallieres *et al.*, 2002).

In a study by Kong *et al.* (2019), IL-6 concentration-dependently reduced neural stem cell proliferation, decreased neuronal differentiation and ultimately impacted upon hippocampal adult neurogenesis through activation of the JAK2/STAT3 signalling pathway. The importance of JAK2/STAT3 signalling in memory consolidation was further highlighted when cognitive deficits in a mouse model of neuroinflammation (APP/PS1) were rescued by inhibiting this signalling cascade (Kong *et al.*, 2019).

In humans, a deterioration in cognitive function is a hallmark feature of 'sickness behaviour' which is linked to high circulating levels of pro-inflammatory cytokines including IL-6 and IL-1 β (Gibertini, 1998). IL-6 has also been implicated in lipopolysaccharide (LPS) -evoked cognitive dysfunction (Sparkman *et al.*, 2006). Moreover, transgenic mice over-expressing IL-6 exhibit broad memory impairment (Heyser *et al.*, 1997; Wei *et al.*, 2012; Wei *et al.*, 2013) and severe neurological pathologies (Campbell *et al.*, 1993). The explicit effect of this cytokine on learning and memory function is further supported by the finding that transgenic mice deficient in IL-6 possess enhanced spatial learning capabilities in the radial arm maze (Braida *et al.*, 2004) and the Morris water maze (Bialuk & Winnicka, 2018) behavioural task-learning assessments.

In early studies, transgenic mice overexpressing IL-6, developed severe neurological pathologies such as tremor, ataxia, and seizure, concomitant with neurodegeneration, astrocytosis and angiogenesis (Campbell *et al.*, 1993). Further studies using the same transgenic mouse model found a significant reduction in

overall neurogenesis in the dentate gyrus of the hippocampus with neural progenitor cells also reduced in the granule cell layer (Vallieres *et al.*, 2002). These studies suggested that overexpression of IL-6 is detrimental to adult neurogenesis and can have a direct pathogenic role not only in inflammation and infections, but also neurodegenerative CNS disease (Campbell *et al.*, 1993; Vallieres *et al.*, 2002).

1.13. Perspectives

Most research on DMD has focussed on the loss of dystrophin from skeletal muscle. As new therapies have improved life expectancy in patients with DMD, greater attention to other aspects of DMD, such as cognitive impairment, is warranted. DMD patients can have communication difficulties and poor affect recognition (Hinton *et al.*, 2006; Hinton *et al.*, 2007) and specific deficits in verbal, short-term and working memory (Hinton *et al.*, 2000, 2001; Snow *et al.*, 2013). In healthy individuals, dystrophin is expressed in the hippocampus, cerebral cortex, amygdala and cerebellum (Lidov *et al.*, 1990; Bies *et al.*, 1992; Lidov *et al.*, 1993; Knuesel *et al.*, 2000; Sekiguchi *et al.*, 2009) and loss of dystrophin has been linked to neuronal loss, neurofibrillary tangles, alterations to dendritic branching, cortical atrophy as well as metabolic abnormalities (Rosman, 1970; Yoshioka *et al.*, 1980; Jagadha & Becker, 1988; Tracey *et al.*, 1995; Lv *et al.*, 2011).

In parallel, contraction-induced damage to muscle fibres (De Paepe & De Bleecker, 2013) results in chronic inflammation, and elevated circulating levels of IL-6, a pro-inflammatory cytokine which has the capacity to modulate brain function, is raised in DMD patients (Rufo *et al.*, 2011) and *mdx* mice (Pelosi *et al.*, 2015) alike. Further,

IL-6 has also been shown to influence long term potentiation (Li *et al.*, 1997; Tancredi *et al.*, 2000; Stampanoni Bassi *et al.*, 2019), neurogenesis (Islam *et al.*, 2009) and astroglialogenesis (Nakanishi *et al.*, 2007).

Aims

To investigate the physiological mechanisms underpinning the neuromodulatory effects of IL-6 in hippocampal-dependent cognitive deficits in the dystrophin-deficient *mdx* mouse model.

Objectives

1. To investigate the impact of loss of dystrophin on hippocampal function in *mdx* mice.
2. To investigate the importance of IL-6 in hippocampal function in dystrophic hippocampal function.
3. To determine if *in vivo* administration of neutralising monoclonal anti-IL-6 receptor antibodies (xIL-6R) improves cognitive dysfunction in *mdx* mice.

Chapter 2: Methods

2.1. Animals

All animal experiments were approved and performed following guidelines set out by the Health Products Regulatory Authority (HPRA) and following project authorisation (AEI9130/P088), as well as individual authorisation (AEI9130/I303).

Animals were euthanized in accordance with European Directive 2010/63/EU.

Animals aged under P14 (postnatal age) were decapitated with sharp scissors.

Animals older than P21 were anaesthetised with Isoflurane (5% Isoflurane in oxygen, Harvard Apparatus) before decapitation with sharp scissors.

Dystrophin-deficient C57BL/10ScSn-Dmd^{mdx}/J (*mdx*) and wild type C57BL/10ScSnJ (WT) controls were purchased from The Jackson Laboratory (Bar Harbor, Maine, USA). Breeding colonies of both *mdx* and WT mice were established and maintained in the Specific Pathogen Free (SPF) unit of the Biological Services Unit, University College Cork, Ireland. To generate *mdx* mice, homozygous females were bred with hemizygous males. Animals were housed with littermates before being weaned at P21. After sexing, given that boys express the DMD phenotype, only males were housed in individually ventilated cages (IVCs) with no more than 4 male littermates in each cage. IVCs were kept under a 12-hour light/dark cycle between 06:00 and 18:00, with water and standard chow provided *ad libitum*. Further, IVCs contained sawdust made from aspen wood chip, sizzlenest (bedding) and cardboard tubes as play tunnels for environment enrichment.

2.2. Tissue preparation for immunocytochemistry and immunohistochemistry

2.2.1. Hippocampal neuron culture

Primary hippocampal neuronal cultures were prepared using neonatal male *mdx* and WT mice pups (3-5 days old).

In order to minimise the risk of contamination of cultured neurons, the initial hippocampal dissection and subsequent cell culture preparation were carried out in a laminar flow cabinet (OSN Horizontal Laminar Airflow Cabinet, Nicotra), with all equipment sterilized with 70% ethanol (prepared from 99.9% ethanol (Brenntag, Essen, Germany)) solution prior to use. All solutions used in the preparation of cultured hippocampal neurons were syringe filtered (0.2µm pore size; Corning Inc., New York, USA) again, to minimise infection, with strict aseptic technique adhered to throughout for the same purpose.

Primary hippocampal neuron cultures were prepared as described previously in our lab (Kaar *et al.*, 2017). The brain was removed and bisected into two hemispheres. Hippocampi were then isolated and placed into a dissection solution made up with HEPES-buffered saline solution (HBSS) (see Table 2.1 for reagent setup and Table 2.1 for culture medium composition).

Following dissection and isolation of the hippocampi, the tissue was transferred to a papain (Worthington, Ohio, USA) solution (2mg/ml; Table 2.2 for culture medium composition) for 45 minutes at 37°C in order to facilitate the dissociation of individual hippocampal neurons from the hippocampal tissue. Papain has previously

been shown to be the most suitable protease for cell dissociation (Brewer, 1997). Dissociated neurons were then rinsed three times with a trituration solution (see Table 2.2: culture medium composition) and then mechanically dissociated three times with a plastic Pasteur pipette (with a diameter of 2mm, Sarstedt, North Carolina, USA) leaving a suspension of cells in the trituration solution. This cellular supernatant was then transferred to another tube with the plastic Pasteur pipette and centrifuged at 1200rpm for two minutes. After centrifugation, the trituration solution was gently poured out of the tube without dissociating the pelleted hippocampal neurons. Approximately 600-700µl plating solution was then added (Table 2.2: culture medium composition) and the pellet gently detached from the bottom of the tube prior to 'finger' vortexing the cells until they were completely resuspended in the plating solution. Dissociated cells were seeded onto poly-D-lysine hydrobromide (Sigma-Aldrich, Missouri, USA) – coated coverslips (see section 2.2.2 Poly-D-lysine coating of coverslips) in 35mm plastic petri dishes (Corning Inc., New York, USA) and incubated (Galaxy S CO₂ Incubator, New Brunswick Scientific, New Jersey, USA) at 37°C with gas levels maintained at 95% O₂ and 5% CO₂. Culture medium was added after 45 minutes (Table 2.2: culture medium composition). Coverslips were then inverted after a further 2 hours in the incubator. Cultures were routinely maintained in the incubator at 37°C, 95% O₂ and 5% CO₂ for 10-12 days. See Figure 2.1 for an overview of the technique.

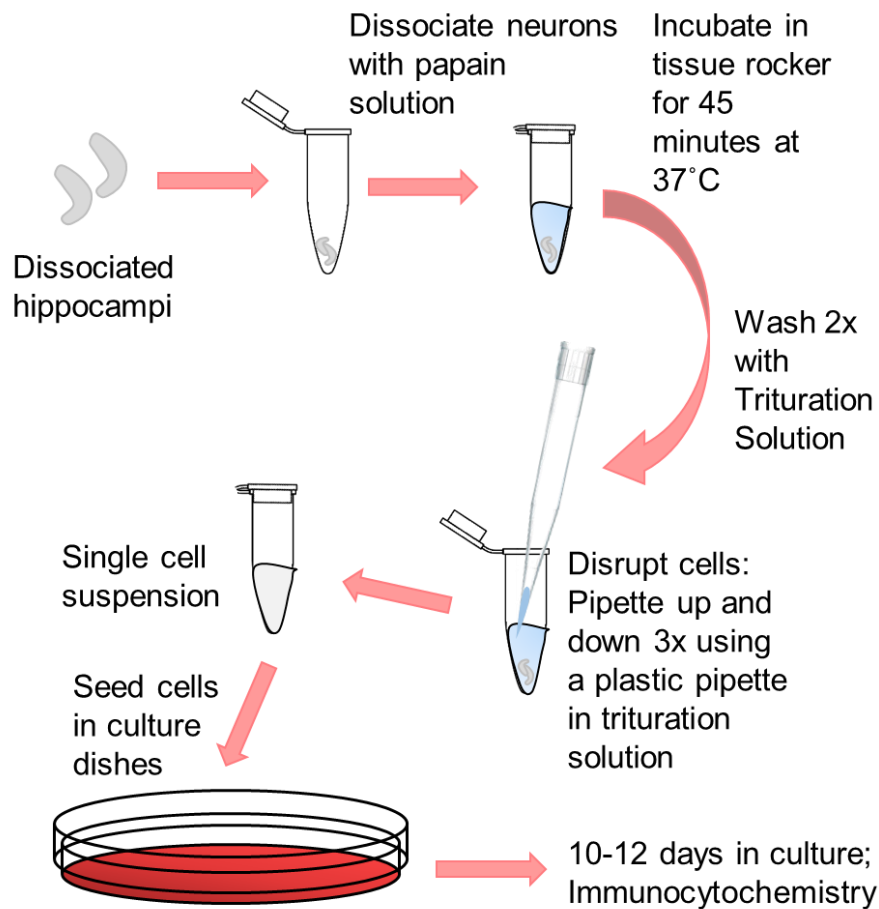


Figure 2.1: Flow diagram displaying the process of primary hippocampal neuron dissociation and culture.

2.2.2. Poly-D-lysine coating of coverslips

In order to minimise the risk of contamination, coverslips were prepared for the culture process in a laminar flow cabinet. Coverslips (13mm - TAAB Laboratories Equipment Ltd., UK) were sterilised in 70% ethanol before being transferred to 35mm dishes (Corning, New York, USA). Poly-D-lysine hydrobromide (35 μ l - 55 μ l, 0.1mg/ml, Sigma-Aldrich, Missouri, USA) was filter-syringed onto each coverslip,

aspirated after 1 hour and the coverslips then washed twice with dH₂O. Coverslips were coated with Poly-D-lysine hydrobromide up to 3 days before a culture.

Table 2.1: Reagent setup for primary hippocampal neuron culture.

Dissection HEPES-buffered saline solution (HBSS), osMom 310, pH 7.35-7.45	Composition (mM)	Source
NaCl	130	Sigma-Aldrich
HEPES	10	Sigma-Aldrich
KCl	5.4	Sigma-Aldrich
MgCl₂	2	Fisher Scientific
D-Glucose	2	Fisher Scientific
CaCl₂	0.5	Sigma-Aldrich

Table 2.2: Primary hippocampal neuron culture medium composition.

Reagents	Components		Source
Dissection solution	HBSS	8ml	See reagent setup
containing	Glutamax™ supplement	0.5mM	ThermoFisher Scientific
	L-Glutamic acid	0.025mM	Sigma-Aldrich
	Penicillin–Streptomycin (P-S)	1%	Sigma-Aldrich
Papain solution	Dissection solution (as	2ml	See reagent set up
containing	above)	2mg/ml	Worthington

	Papain		
Trituration	HBSS	15ml	See reagent set up
solution	Foetal Bovine Serum (FBS)	10%	Sigma-Aldrich
containing	Glutamax™ supplement	0.5mM	ThermoFisher Scientific
	L-Glutamic acid	0.025mM	Sigma-Aldrich
	P-S	1%	Sigma-Aldrich
Plating solution	DMEM	3ml	Sigma-Aldrich
containing	FBS	10%	Sigma-Aldrich
	Glutamax™ supplement	0.5mM	ThermoFisher Scientific
	L-Glutamic acid	0.025mM	Sigma-Aldrich
	P-S	1%	Sigma-Aldrich
Culture medium	DMEM – low glucose		Sigma-Aldrich
containing	Serum Replacement 2 (SR2)	2%	Sigma-Aldrich
	Glutamax™ supplement	0.5mM	ThermoFisher Scientific
	L-Glutamic acid	0.025mM	Sigma-Aldrich
	P-S	1%	Sigma-Aldrich

2.2.3 Preparation of Hippocampal tissue for Immunohisto/cytochemistry

Mice (8–12-week-old WT and *mdx*) were anaesthetised with isoflurane (5% in oxygen) and then immediately euthanized by decapitation using sharp scissors. The whole brain was excised from the skull and submerged in 4% PFA (Sigma-Aldrich,

Missouri, USA) for 72 hours, then cryo-protected by immersing in a 30% sucrose solution (Sigma-Aldrich, Missouri, USA) made up in 0.1mM Phosphate Buffered Saline (PBS, See Table 2.3: reagent setup) for 24 hours. Tissue was then flash frozen in liquid nitrogen and stored at -80°C until cryo-sectioned (CM1850, Leica, Germany) in Optimal Cutting Temperature (OCT) embedding medium (Thermofisher Scientific, Massachusetts, USA) in 20µm sections for subsequent immunohistochemistry and immunofluorescence procedures. Cryo-sectioned tissue was mounted onto positively charged glass slides (SuperFrost® Plus, VWR International, Pennsylvania, USA).

Importantly, in later immunofluorescence experiments, on hippocampal adult neurogenesis specifically, tissue was collected and processed using a different technique. Mice (8–12-week-old WT and *mdx*) were anaesthetised with isoflurane (5% in oxygen) and then were perfused with approximately 15ml intracardiac saline (Sigma-Aldrich, Missouri, USA), followed by approximately 15ml formalin (Sigma-Aldrich, Missouri, USA). Whole brains were excised from the skull and post fixed in formalin (Sigma-Aldrich, Missouri, USA) for 72 hours then cryo-protected by immersing in a 30% sucrose solution (Sigma-Aldrich, Missouri, USA) made up in 0.1mM Phosphate Buffered Saline (PBS, See Table 2.3: reagent setup). Samples were frozen in isopentane and stored at -80°C until cryo-sectioned (CM1850, Leica, Germany) in OCT embedding medium (Thermofisher Scientific, Massachusetts, USA) in 20µm sections for subsequent immunohistochemistry and immunofluorescence procedures. Cryo-sectioned tissue was again mounted onto positively charged glass slides (SuperFrost® Plus, VWR International, Pennsylvania, USA).

Table 2.3: Reagent set up for immunocytochemistry/immunohistochemistry

Phosphate Buffered	Composition (mM)	Source
Saline (PBS)		
NaCl	137	Sigma-Aldrich
KCl	2.7	Sigma-Aldrich
NA₂HPO₄	10	Sigma-Aldrich
KH₂PO₄	1.8	Sigma-Aldrich

2.3. Immunocytochemistry techniques

2.3.1 Immunofluorescent labelling of cultured hippocampal neurons

After 10-12 days in culture, culture medium was removed from the culture dish, and hippocampal cell cultures were washed with PBS and fixed in 4% paraformaldehyde (PFA – Sigma-Aldrich, Missouri, USA) for 90 min at 4°C.

Subsequently, fixed samples were permeabilized with 0.1% Triton X-100 (Sigma-Aldrich, Missouri, USA) at room temperature for 5 min. Samples were then washed three times with PBS and blocked with 1% Donkey Serum (Sigma, Missouri, USA) for 1 hour at room temperature, which is essential for preventing non-specific binding of antibodies to tissue. Primary antibodies were diluted in PBS (see Table 2.4 for antibodies used in this thesis), and preparations were incubated with them for 2 hours at room temperature. After washing three times with PBS (5 mins per wash),

preparations were incubated with the appropriate secondary antibody (see Table 2.4) for 2 hours at room temperature. Finally, samples were washed with PBS three times (5 mins per wash) and the coverslips containing the fixed neurons were mounted onto a microscope glass slide (VWR International, Pennsylvania, USA) using Fluoroshield mounting medium with 4',6-diamidino-2-phenylindole (DAPI - Abcam, Cambridge, UK) which binds to adenine-thymine rich regions in the DNA in order to visualise nuclei of cells.

2.3.2. Immunofluorescent labelling of whole brain sections

After whole brain tissue cryo-sectioning, the tissue was permeabilized with 0.1% Triton X-100 (Sigma-Aldrich, Missouri, USA) at room temperature for 1 hour. Samples were then washed three times in PBS (3x 10 minutes) and blocked with 1% Donkey Serum (Sigma-Aldrich, Missouri, USA) for 2 hours at room temperature. Primary antibodies (Table 2.4) were diluted in PBS, and tissue incubated with them overnight at 4°C. After washing three times with PBS (3x 10 minutes), preparations were incubated with the appropriate secondary antibody (Table 2.4) for 2 hours at room temperature. Finally, samples were washed with PBS (3 x 10 minutes) and a glass coverslip (VWR, Pennsylvania, USA) mounted onto the slide with Fluoroshield mounting medium with DAPI (Abcam, Cambridge, UK). Donkey serum blocking was used for the following primary antibodies: Dystrophin, Synaptophysin, Dystrophin (MANDRA-1), IL-6 and IL-6R α .

Alternatively, after whole brain tissue cryo-sectioning, a Blocking-Permeabilization Buffer, composed of 10% Foetal Bovine Serum (FBS), 1% BSA and 0.1% Triton X-100

(Sigma-Aldrich, Missouri, USA) in PBS, was used. Sectioned tissue was permeabilized and blocked with this Blocking-Permeabilisation Buffer for 1 hour at room temperature. Samples were then washed in PBS (3x 10 minutes). Primary antibodies (Table 2.4) were diluted in PBS, and tissue incubated with them overnight at 4°C. After washing once more with PBS (3x 10 minutes), preparations were incubated with the appropriate secondary antibody (Table 2.4) for 2 hours at room temperature. After a final wash with PBS (3 x 10 minutes), the tissue was mounted onto a glass slide with Fluoroshield mounting medium with DAPI (Abcam, Cambridge, UK) and coverslipped (VWR, Pennsylvania, USA). FBS blocking was used on the following antibodies: doublecortin, GABA_AR, NeuN and GFAP.

Table 2.4: Antibodies used in this thesis

Primary Antibody	Host	Dilution	Source
pAb to Dystrophin	Rabbit	1:250	Abcam
mAb to	Rabbit	1:100	Abcam
Synaptophysin			
mAb to Dystrophin	Mouse	1:100	Abcam
(MANDRA-1)			
mAb to IL-6 (1)	Mouse	1:250	Santa Cruz Biotechnology
mAb to IL-6Rα (D-8)	Mouse	1:250	Santa Cruz Biotechnology
pAb to	Rabbit	1:100	Invitrogen
Doublecortin (DCX)			
mAb to GABA_AR	Mouse	1:100	Abcam

pAb to NeuN	Rabbit	1:100	Invitrogen
mAb to GFAP	Rabbit	1:100	Invitrogen
Secondary Antibody			
Rhodamine (TRITC)	Donkey anti-rabbit	1:250	Jackson Immuno-Research
Fluorescein (FITC)	Goat anti-mouse	1:250	Jackson Immuno-Research
Alexa Flour AP 488	Goat anti-rabbit	1:250	Abcam

2.3.3 Confocal microscopy

Imaging of fluorescently-labelled proteins was carried out using a confocal microscope (Olympus Fluoview, FV10i). Laser and saturation settings were set for each immunofluorescently-labelled protein and kept consistent for all samples within a group.

2.3.4 Image Analysis

When imaging cultured hippocampal neurons, 3 individual fields were imaged at random across the coverslip of cultured hippocampal neurons. Hippocampal neurons from at least three different cultures from each group (*e.g.*, WT and *mdx*) were compared for each experiment.

When imaging the different regions of the hippocampus, namely CA1, CA3 and DG (Figure below 2.2), three individual fields were imaged in each region for an appropriate overview. Hippocampal tissue from at least three different animals from each group (*e.g.*, WT and *mdx*) were compared for each experiment.

Depending on the staining being visualised and assessed, different techniques were applied using Image J. Mean fluorescence intensity (minus background) was assessed in a region of a specified area for the expression of dystrophin, IL-6 and IL-6R. For other antibodies, such as doublecortin (DCX), Neuronal nuclear protein (NeuN), Glial fibrillary acidic protein (GFAP) and γ -aminobutyric acid type A receptors (GABA_AR), the number of cells in the granule cell layer and/or hilus in the DG expressing each protein was counted (see Figure 2.3 below). Further explanation on analysis of each antibody is provided in the results.

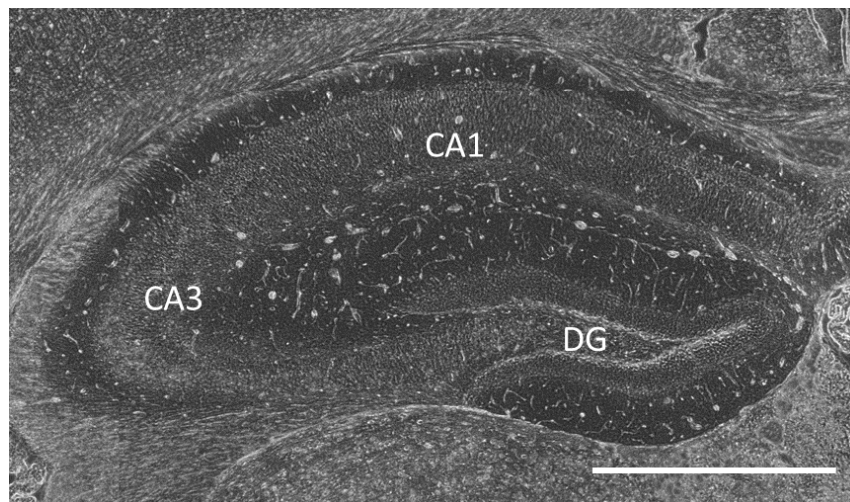


Figure 2.2: Brightfield image of the 3 regions of the hippocampus.

Confocal images were taken from 3 different regions of the hippocampus, namely CA1, CA3 and DG. Scalebars are 1mm.

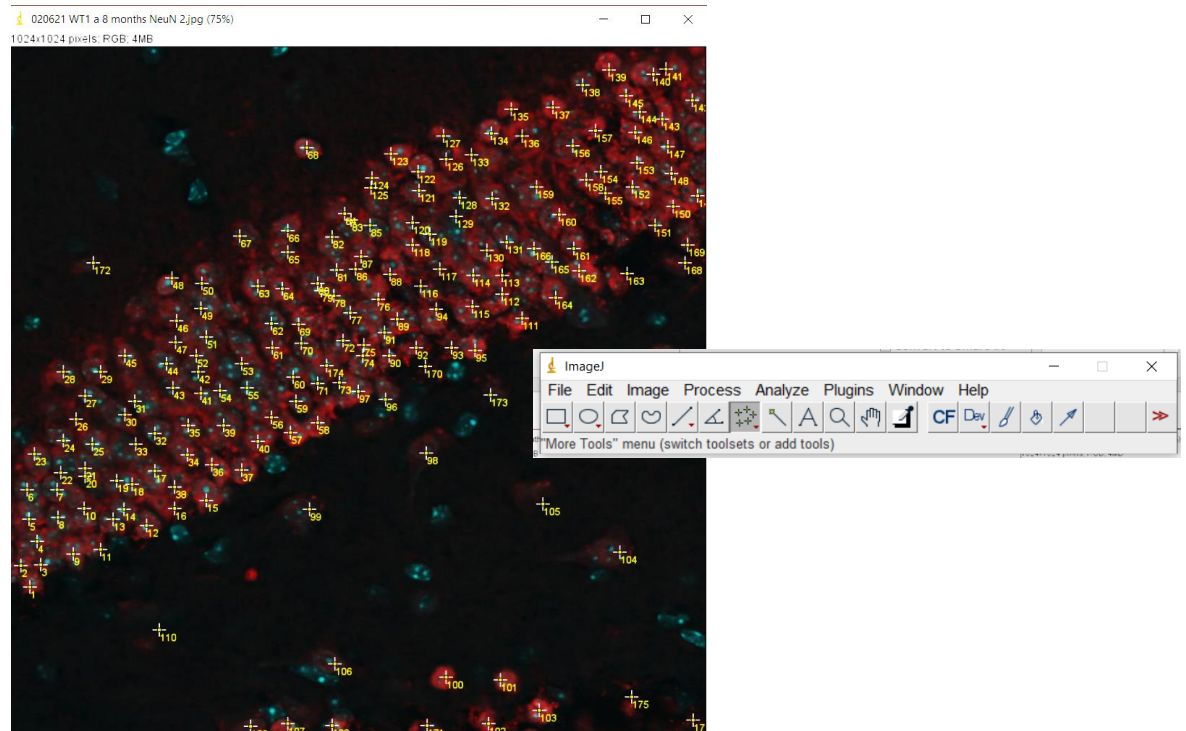


Figure 2.3: Representative image of analysis of whole brain sections for cell counting using Image J.

Image J was used as a tool for marking (white crosses) each neuron (NeuN, red stain) in order to count the number present in both the granule cell layer and hilus in the DG. DAPI (blue stain) showed nuclei of each cell.

2.4. Calcium imaging with cultured hippocampal neurons

2.4.1. Fluorescent calcium dye loading protocol.

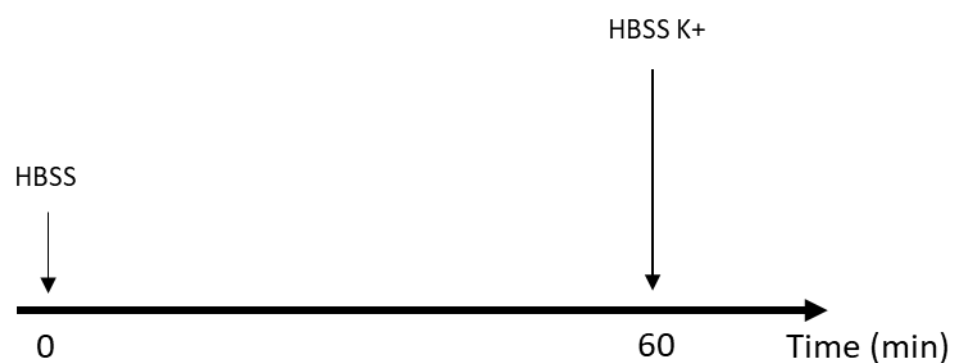
After 3-4 days incubation, intracellular calcium measurements from neuronal somata were conducted as previously described (Rae et al., 2000). Cultured

hippocampal neurons were preloaded with a single wavelength, intensity modulating calcium indicator, Fluo-2 AM (8 μ M dissolved in HBSS, Sigma-Aldrich, Missouri, USA), for 1 hour at room temperature in the dark. After that time, and prior to imaging the dye was washed off the neurons with HBSS.

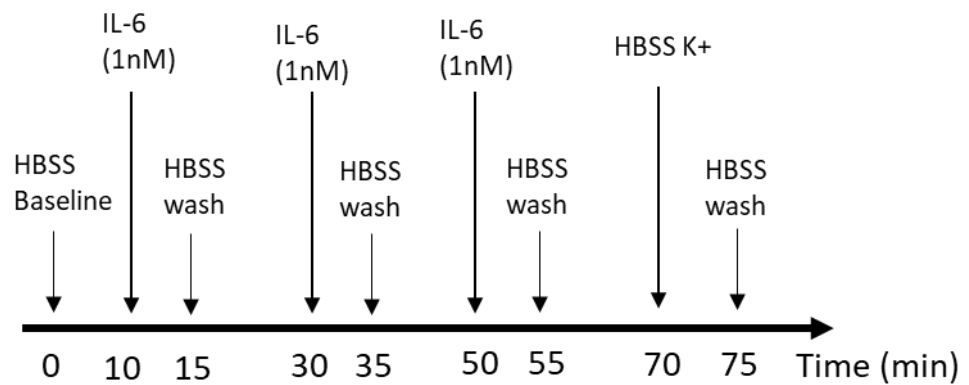
Coverslips upon which the neurons had been plated were then placed into a recording chamber and placed in the light path of an upright microscope (Olympus BX50 WI) in a dark room and superfused with HBSS alone or in the presence of drugs as appropriate (see section 2.4.2 below for experimental design). Initially, normal white light was used to find the appropriate field of view of neurons on the coverslips with a bright light source (Olympus TH4-200). At that point, the neurons were illuminated by a fluorescent light source monochromator (excitation at 488 nm, emission at 550 nm), generated by a monochromator (Cairn Technologies) and visualised with an integrating camera (Hamamatsu ORCA-ER), DAQ device (USB6229) and frame grabber, under both 20x and 40x magnification, as appropriate.

2.4.2 Experimental design

1. HBSS Spontaneous activity



2. IL-6 (1nM)



Images were acquired at one-second intervals using the open-source imaging software, Winfluor (John Dempster, University of Strathclyde, Scotland). Data files with time (seconds) vs fluorescence values (arbitrary units) for each region of interest (ROI) were exported from Winfluor as standard text files. GraphPad Prism was then used to produce XY line graphs to analyse changes in intracellular calcium for each cell indicated by a ROI.

2.4.3 Analysis of calcium imaging data

For each identified neuron (identified by high K⁺ response after experiments), the frequency of peaks (number of peaks per minute) was determined in non-stimulated WT and *mdx* cultures.

In experiments where neurons were exposed to IL-6 (1nM, 5 minutes), the frequency of peaks during the stimulated period were compared to the frequency of peaks during baseline (HBSS only). WT cultures were compared to *mdx* cultured neurons.

2.5. Gene Expression

Hippocampi from *mdx* and WT (8-12 week) whole brains were bisected from each hemisphere and stored in RNALater (ThermoFisher Scientific, Massachusetts, USA) for 24-48 hours. Hippocampi were then flash frozen in liquid nitrogen and stored at -80°C.

2.5.1. RNA Extraction and Preparation

Total RNA was extracted from hippocampi (10 WT and 10 *mdx*) by homogenization using a glass tissue grinder (Dounce tissue grinder, 1ml) in the presence of TriPure Isolation Reagent (Roche Diagnostics Ltd., UK). This was followed by a series of chloroform (Sigma-Aldrich, Missouri, USA) and iso-propanol (Sigma-Aldrich, Missouri, USA) steps, as well as 3 centrifugation steps (Figure 2.4 below). The quantity as well as purity of the RNA was then assessed by spectrophotometry using a Nanodrop 1000 (ThermoFisher Scientific, Massachusetts, USA). The quality of the RNA was assessed using an agarose gel electrophoresis system (E-Gel® X, Invitrogen, California, USA), with the subsequent visualisation of clear 18S and 28S ribosomal RNA bands used to confirm that there was no degradation of the sample during the extraction process (Figure 2.5). For mammalian RNA, a 28S:18S RNA ratio of 2:1 is generally representative of good-quality RNA.

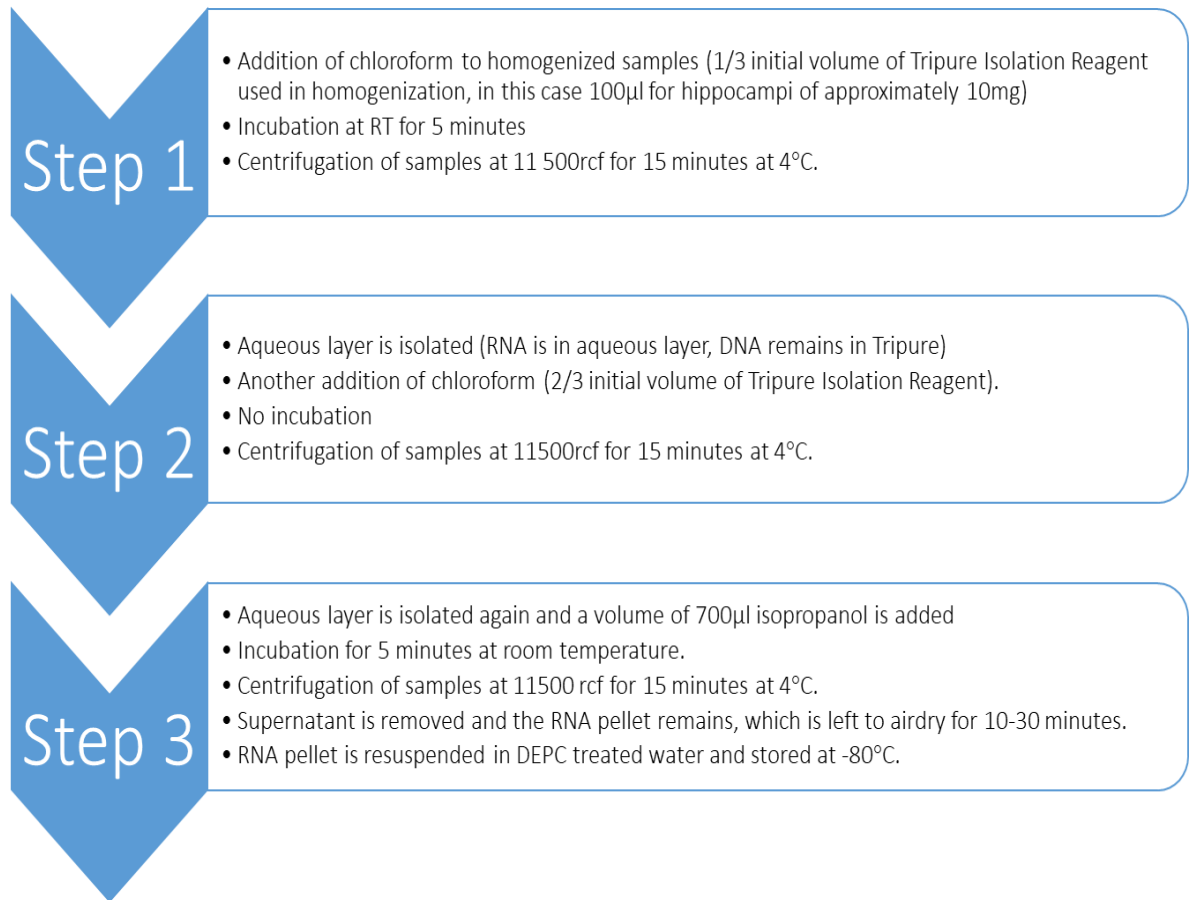


Figure 2.4: Flow diagram displaying the steps involved in RNA extraction from hippocampal tissue.

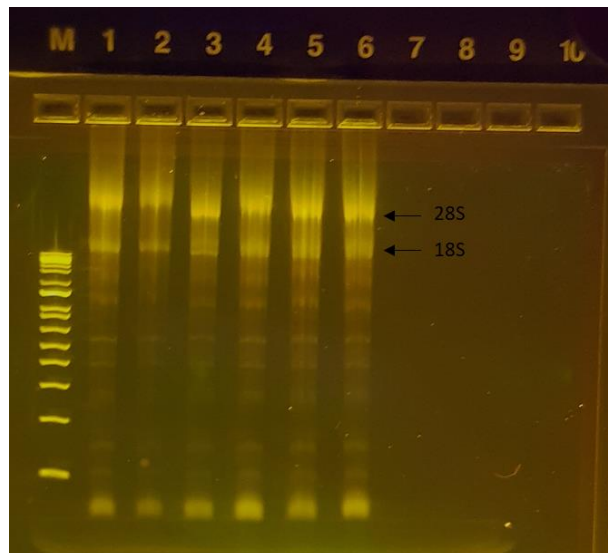


Figure 2.5: RNA analysis using agarose gel electrophoresis.

The bands indicated on the gel show distinct 28S and 18S bands which indicates that there was no degradation of the sample.

2.5.2. Reverse Transcription of RNA

RNA from hippocampal tissue was reverse transcribed using Transcriptor First Strand cDNA Synthesis Kit (Roche Diagnostics Ltd., UK). The following steps were followed:

- After the total mRNA was determined from the Nanodrop 1000 (Thermofisher Scientific, Massachusetts, USA), the sample was diluted in Diethyl pyrocarbonate (DEPC) treated water to a final concentration of 200ng.
- A template-primer mix was made up of 1µg total RNA, anchored-oligo(dT)₁₈ primer (50pmol/µl – with a final concentration of 2.5µM), random hexamer

primer (600 pmol/μl – with a final concentration of 60μM) and DEPC-treated water.

- The remaining components of the reverse transcription (RT) mix are added which contained Transcriptor Reverse Transcriptase Reaction Buffer (8mM MgCl₂), protector RNase inhibitor (20U/μl), a deoxynucleotide mix (10mM – with a final conc. 1mM) and a Transcriptor Reverse Transcriptase (20 U/μl – final concentration of 10 U).
- The RT reaction was then incubated for 10min at 25°C, followed by 30min at 55°C and 5min at 85°C.

2.5.3. *qRT-PCR*

cDNA was amplified using Realtime ready Catalog or Designer Assays (Roche Diagnostics Ltd., UK) (as described above) and Fast Start Essential DNA Probe Master (Roche Diagnostics Ltd., UK) in 20 μl reactions (5 μl cDNA and 15 μl master mix) using the LightCycler 96 (Roche Diagnostics Ltd., UK) on 96-well plates. All reactions were performed in duplicate and reverse transcriptase negatives, RNA negatives, and cDNA negatives (with no template), and plate calibrator controls were used on every plate (an example of this template is shown in figure 2.6 below). Hippocampal gene expression was normalized to a reference gene, HPRT1, to compensate for variations in input amounts of RNA/cDNA and the efficiency of reverse transcription. The relative expression of genes was calculated using the $\Delta\Delta CT$ method, i.e., normalized expression of the gene of interest to that of the

reference gene, with changes in expression displayed as a fold change relative to the control group.

LightCycler® 96



Experiment: kim stephenson il6st 060421.lc96p

Plate Id: <None>

*	1	2	3	4	5	6	7	8	9	10	11	12
A	U mdx 1 gp130	U mdx 2 gp130	U mdx 3 gp130	U mdx 4 gp130	U mdx 5 gp130	U mdx 7 gp130	U mdx 10 gp130	U mdx 11 gp130	U mdx 12 gp130	U mdx 13 gp130	- NTC None	- NTC None
B	U mdx 1 gp130	U mdx 2 gp130	U mdx 3 gp130	U mdx 4 gp130	U mdx 5 gp130	U mdx 7 gp130	U mdx 10 gp130	U mdx 11 gp130	U mdx 12 gp130	U mdx 13 gp130	- RNA NEG None	- RNA NEG None
C	U WT 1 gp130	U WT 3 gp130	U WT 5 gp130	U WT 7 gp130	U WT 9 gp130	U WT 12 gp130	U WT 13 gp130	U WT 16 gp130	U WT 15 gp130	U WT 14 gp130	- RT NEG HPRT	- RT NEG HPRT
D	U WT 1 gp130	U WT 3 gp130	U WT 5 gp130	U WT 7 gp130	U WT 9 gp130	U WT 12 gp130	U WT 13 gp130	U WT 16 gp130	U WT 15 gp130	U WT 14 gp130	+ Cal HPRT	+ Cal HPRT
E	U mdx 1 HPRT	U mdx 2 HPRT	U mdx 3 HPRT	U mdx 4 HPRT	U mdx 5 HPRT	U mdx 7 HPRT	U mdx 10 HPRT	U mdx 11 HPRT	U mdx 12 HPRT	U mdx 13 HPRT	- RT NEG gp130	- RT NEG gp130
F	U mdx 1 HPRT	U mdx 2 HPRT	U mdx 3 HPRT	U mdx 4 HPRT	U mdx 5 HPRT	U mdx 7 HPRT	U mdx 10 HPRT	U mdx 11 HPRT	U mdx 12 HPRT	U mdx 13 HPRT		
G	U WT 1 HPRT	U WT 3 HPRT	U WT 5 HPRT	U WT 7 HPRT	U WT 9 HPRT	U WT 12 HPRT	U WT 13 HPRT	U WT 16 HPRT	U WT 15 HPRT	U WT 14 HPRT		
H	U WT 1 HPRT	U WT 3 HPRT	U WT 5 HPRT	U WT 7 HPRT	U WT 9 HPRT	U WT 12 HPRT	U WT 13 HPRT	U WT 16 HPRT	U WT 15 HPRT	U WT 14 HPRT		

Figure 2.6: Template used for all qRT-PCR reactions. Half the plate (Rows A to D, column 1 to 10) are dedicated to samples and the probe of interest (in this case, IL-6st) and the other half is dedicated to the reference housekeeping gene (which was HPRT, and was included in all probe runs). Column 11 and 12 are dedicated to reverse transcriptase negatives for both the reference gene and target gene, RNA negatives, and cDNA negatives (with no template), and plate calibrator controls.

2.6. Immunoassay for protein quantification

2.6.1. *Animals*

WT and *mdx* mice were euthanised by cervical dislocation and decapitation at 8-12 weeks, with whole brains removed from the skull. Hippocampi were then excised from each hemisphere and immediately flash frozen in liquid nitrogen and stored at -80°C.

2.6.2. *Protein Extraction and Quantification*

Protein was extracted from one hippocampi from each sample (10 WT and 10 *mdx*) by homogenisation using a glass tissue grinder (Dounce, 1ml tissue grinder) in 500 µl lysis buffer (Table 2.5 below) made up with Tris-HCL (10mM, pH7.0), 0.5% Triton X-100 (Sigma-Aldrich, Missouri, USA) as well as cOmplete™, Mini, EDTA-free Protease Inhibitor Cocktail (Roche Diagnostics Ltd., UK). Samples were then centrifuged at 10,000 rpm for 10 minutes at 4°C. Aliquots of the supernatant were then stored at -80°C to limit freeze thaw cycles.

2.6.3. *Bradford Protein Assay*

Bradford reagent (Sigma-Aldrich, Missouri, USA) was used to quantify the concentration of protein in each sample. Bradford protein quantification is a colorimetric assay which measures the shift in absorbance of Commasie Brilliant Blue G-250 (Bradford 1976). Using a standard curve of known protein concentration of bovine serum albumin (BSA, Sigma-Aldrich, Missouri, USA), with concentration

standards ranging from 0mg/ml to 1.5mg/ml, the concentration of the unknown samples was determined. Samples were diluted with Tris-HCL (10mM, pH7.0, Sigma-Aldrich, Missouri, USA) in a 1:4 dilution. Absorbance was read at 595nm on a spectrophotometer (Molecular Devices, SpectraMax M3, USA) and the concentration of protein was determined using SoftMax Pro 6 software package.

Table 2.5: Lysis Buffer Recipe.

Reagent	Composition	Source
Tris- HCL (10 mM, pH7.0)	1M Tris-HCL	Sigma
Triton X-100	0.5%	Sigma
cOmplete™, Mini, EDTA-free Protease Inhibitor Cocktail	1 tablet per 10ml	Roche Diagnostics Ltd., UK

2.6.4. Enzyme Linked Immunosorbent Assay (ELISA)

An ELISA was performed using the LEGEND MAX™ Mouse IL-6 ELISA Kit (BioLegend, San Diego, USA) (see Figure 2.7 below), on WT and *mdx* hippocampi samples of known total protein concentration after completion of a Bradford protein assay (as described above). A standard curve was prepared using serial dilutions of standard IL-6 protein supplied with the kit (0 pg/ml, assay diluent as blank, to 500pg/ml).

The ELISA plate was read at 570nm and 450nm using a multi well spectrophotometer (Molecular Devices, SpectraMax M3, USA). The standard curve

was generated using GraphPad Prism software and each sample concentration was calculated from it.

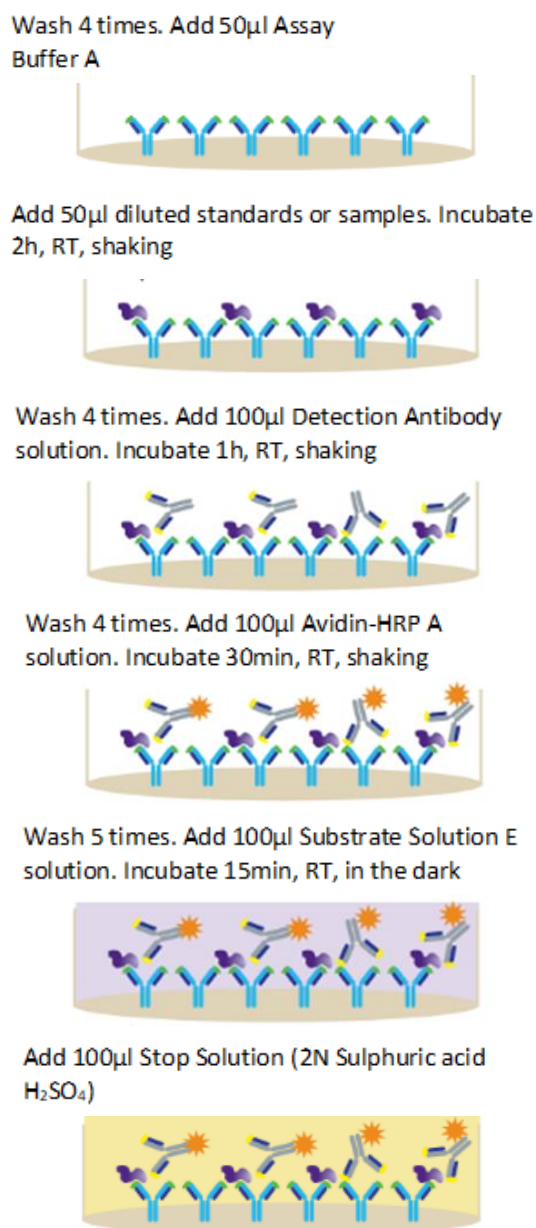


Figure 2.7: ELISA protocol followed using the LEGEND MAX™ Mouse IL-6 ELISA Kit.

2.7. Behavioural tests

2.7.1. *Animals*

Behavioural experiments were carried out on 8–12-week-old WT and *mdx* mice.

All behavioural experiments were carried out between 8:30-14:00 and room temperature and humidity were maintained at $22\pm 1^{\circ}\text{C}$ and $57\pm 2\%$, respectively, throughout.

Mice were handled for 4 days prior to the start of trials in order to habituate them to any handling that may have occurred during the behavioural test phase. This type of handling involved picking up the mice 10 times on each day of handling and letting them walk along the experimenter's sleeve each time. Mice were also moved from the holding room to the testing room an hour before the start of each trial in order to habituate them to the room.

2.7.2. *Novel Object Recognition (NOR) Test*

NOR tests are used to test a rodent's ability to recognise a previously presented object or stimulus (Ennaceur & Delacour, 1988) and as such, are used to assess recognition memory. NOR tests were carried out in an open field arena (30cm x 30cm x 30cm) under a light intensity of approximately 15 Lux. Firstly, mice were allowed to habituate to the open field arena without any objects for 10 minutes. The arena was then wiped down with 70% ethanol after each mouse to remove any olfactory cues. Trial 1 (training phase) took place 24 hours after the open field habituation, where mice were exposed to two identical objects (200ml purple plastic bottles filled with

water; figure 2.8 below) for 10 minutes. Trial 2 took place 24 hours later, where one object was exchanged for a novel object (a 200ml flat bottomed rectangular flask partially filled with blue dye; figure 2.7 below). Exploration of an object included directing the snout at a distance less than 1cm to the object, sniffing or touching it with the snout, plus leaning on the object while sniffing or touching it with the snout. In contrast, turning or running around the object, or climbing or sitting on the object was not considered as exploration. Mouse activity was recorded on a camera (Akaso EK7000, 4K camera) placed 50cm directly above the open field area and behaviour was analysed at a later time point by two researchers, blinded to the groups. The objects were unable to be displaced by the mouse and were wiped down thoroughly with 70% ethanol after each trial to ensure that no olfactory cues remained.

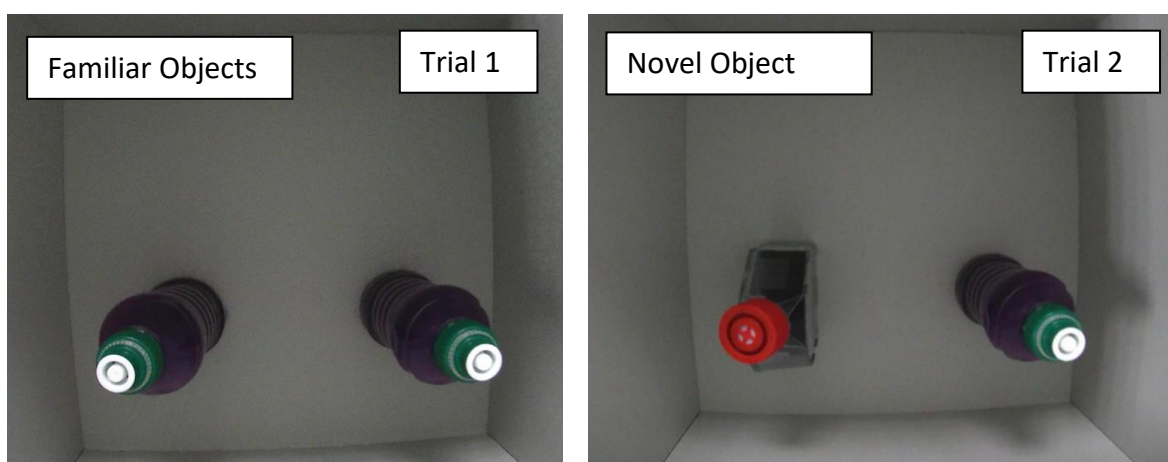


Figure 2.8: Novel Object Recognition test setup. The familiar objects used were two purple plastic bottles, filled with water (left image). The novel object, used in trial 2, was a 200ml flat bottomed rectangular flask partially filled with blue dye (right image)

The percent of time spent exploring the novel object relative to the total time spent exploring both objects can be a measure of novel object recognition (Benice *et al.*, 2006; Sarkisyan & Hedlund, 2009; Broadbent *et al.*, 2010; Oliveira *et al.*, 2010).

Thus, *Recognition Index* (RI), i.e., the time spent investigating the novel object relative to the novel object and the familiar object investigation [$RI = T_N / (T_N + T_F)$] was used (Mumby *et al.*, 2002; Piterkin *et al.*, 2008; Botton *et al.*, 2010; Gaskin *et al.*, 2010; Schindler *et al.*, 2010). Overall, an exploration ratio of greater than 50% indicated that the mouse investigated the novel object more than the familiar object, which suggests a memory recall for the for the familiar object (Gareau *et al.*, 2011).

Exclusion criteria (Lueptow, 2017):

- Mice were excluded from the study if the total exploration time (sum of both objects) was less than 20 seconds either on trial 1 or trial 2.
- During trial 1 (training phase), if mice had a less than 40%, or greater than 60%, exploration ratio (suggesting a bias towards one object) they were excluded from further analyses.

For these experiments, 5 *mdx* mice and 3 WT mice were excluded.

	Left Object	Right Object
	Time Exploring (seconds)	Time Exploring (seconds)
<i>mdx</i> 19/034	10,02	6,5
<i>mdx</i> 19/039	5,17	9,42
<i>mdx</i> 19/040	11,16	17,63
<i>mdx</i> 19/004	3,89	6,16
<i>mdx</i> 19/022	5,14	1,11
BL10 19/011	3,25	2,14
BL10 19/021	5,47	12,84
BL10 19/013	1,84	0,72

For the intervention experiments, 1 WT mouse was excluded.

BL10	8,7	8,4
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2.7.3. Open field behavioural test

Mice were allowed to habituate to the open field arena for 10 minutes prior to NOR testing, with the mouse activity recorded (Akaso EK7000, 4K camera) placed 50cm directly above the open field area and behaviour was analysed at a later time point by two researchers, blinded to the groups. Between mice, the arena was wiped down with 70% ethanol after each mouse to remove any olfactory cues.

Importantly, this open field behavioural test was not set out as a separate experimental protocol, rather as an additional analysis to the data acquired in habituation in the NOR test.

The behaviours assessed were the following and can be visualised in figure 2.9 below:

- a) Faecal output (number of boli)
- b) Grooming (number of times the mouse scratched its face and fur with its forepaws)
- c) Central area frequency (number of times the mouse had 4 paws in central area)
- d) Freezing (number of times the mouse did not move for more than 2 seconds).
- e) Rearing (number of times the mouse reared its front paws, against the wall or mid-air)
- f) Jumping (number of times the mouse jumped against the wall, with all 4 paws in the air).

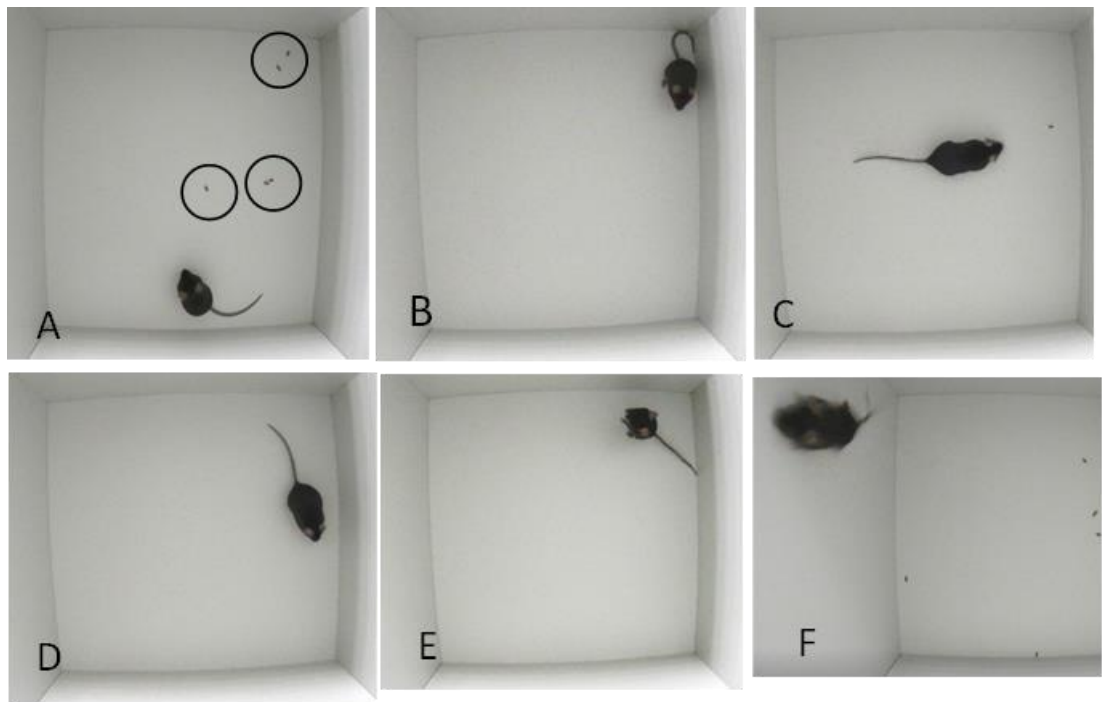


Figure 2.9: Representative images showing behaviours in the open field arena. Number of boli, shown by black circles (A), grooming (B), central area frequency (C), freezing (D), rearing (E) and jumping (F).

2.8. Slice Electrophysiology

2.8.1. Preparing hippocampal slices

Male WT and *mdx* mice (8–12-week-old) were used for *ex vivo* extracellular electrophysiological studies.

Following decapitation, the brain was quickly removed from the skull and placed in a slush of frozen, carbogenated (95% O₂, 5% CO₂) cutting solution (CS – Table 2.6). Transverse slices of 300 µm thickness were sectioned using a vibrotome (5100mz, Campden Instruments Ltd., UK) in the same CS slush. Slices were then transferred

to a solution made up of 50:50 CS and artificial cerebrospinal fluid (ACSF – Table 2.7). After 30 minutes in this solution, slices were then transferred to 100% aCSF for at least 1 hour before recording, in order for the slices to stabilise in the artificial media. Throughout the slicing and recovery periods, the slices were constantly carbogenated (95% O₂, 5% CO₂) and maintained at room temperature (23-25°C). Importantly, the hippocampus was sliced in an orientation to display the dorsal portion of the hippocampus.

Table 2.6: Electrophysiology reagents – cutting solution recipe.

Cutting Solution	Composition (mM)	Source
Sucrose	100	Sigma-Aldrich
HEPES	20	Sigma-Aldrich
NaCl	50	Sigma-Aldrich
KCl	3	Sigma-Aldrich
NaH₂PO₄	1.25	Sigma-Aldrich
NaHCO₃	28	Sigma-Aldrich
CaCl₂	0.5	Sigma-Aldrich
MgCl₂	7	Fisher Scientific
D- Glucose	5	Fisher Scientific

Table 2.7: Electrophysiology reagents – aCSF recipe.

aCSF	Composition (mM)	Source
HEPES	20	Sigma-Aldrich
NaCl	125	Sigma-Aldrich

KCl	2.5	Sigma-Aldrich
NaH₂PO₄	1.25	Sigma-Aldrich
NaHCO₃	25	Sigma-Aldrich
CaCl₂	2	Sigma-Aldrich
MgCl₂	1	Fisher Scientific
D-Glucose	25	Fisher Scientific

2.8.2. Extracellular electrophysiological recording

After at least a 1-hour recovery period following slicing, hippocampal slices were transferred to a recording chamber, and secured in placed under a slice holder consisting of a grid of nylon threads within a circular metal frame (ALA Scientific Instruments, United States, see figure 2.10 below). The slices were continually superfused with oxygenated aCSF (2ml/min) through a feedback-controlled temperature regulator (ThermoClamp-1, Digitimer, UK) and kept at a constant 28°C for the duration of all experiments. Keeping the temperature stable at 28°C resulted in stability of recordings as well as the temperature in which we saw the largest EPSPs, whereas higher or lower temperatures resulted in fluctuation.

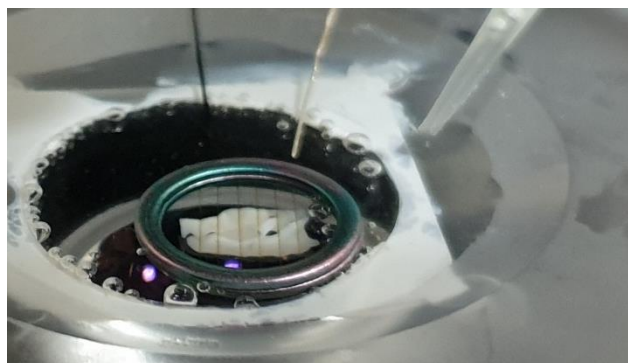


Figure 2.10: Recording chamber set up. Brain slices were secured in the recording bath by a slice holder.

2.8.3. Induction of LTP

An advantage of using the hippocampus for experimental procedures, is that it is anatomically distinctive in that the cell layers in individual regions are conserved throughout. Thus, signalling pathways between the regions can be investigated in a systematic manner between different slices. A bipolar stimulating electrode was used to stimulate axons of the Schaffer-collateral pathway (located between the CA3 region and CA1 region of the hippocampus; see figure 2.11) in order to evoke field excitatory postsynaptic potentials (fEPSPs) in the stratum radiatum of the CA1 region, which were recorded using an aCSF-filled, low resistance (2–5 M Ω), monopolar borosilicate glass electrode (premium thin wall borosilicate capillary glass with filament, Model G150TF-4, Multichannel systems, Germany). Once the evoked fEPSP amplitude had stabilised, a process that usually took approximately 1 hour, an input/output curve was generated by evoking fEPSPs at a frequency of 0.05Hz, whilst steadily increasing the stimulation current from a minimum of 0.11mA up to a maximum of 2mA. Three fEPSPs were recorded at each current, with the average slope value (mV/ms) of the three recordings used for analysis. Current stimulation intensity was then adjusted to evoke 50% of the maximal fEPSP response (as determined by the input/output curve) and a control 20-minute recording period initiated, with fEPSPs again evoked at 0.05Hz.

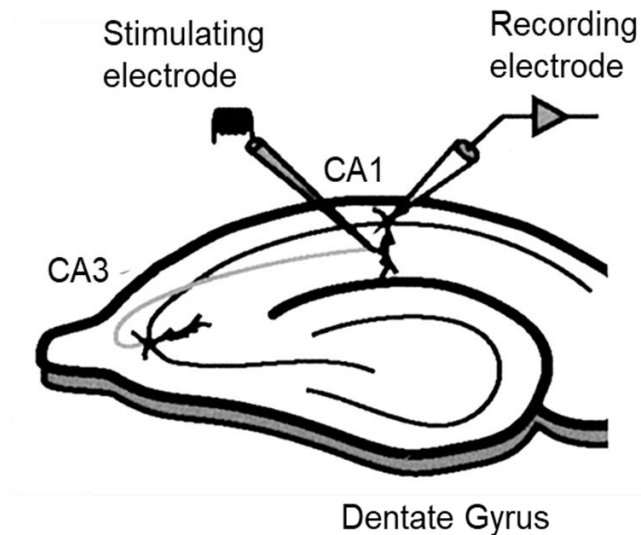


Figure 2.11: A schematic of the placement of the stimulation-recording path (Schaffer collateral-CA1 synapses) in the mouse hippocampus. A bipolar stimulation electrode (stimulating electrode) is placed in the Schaffer-collateral pathway, with responses recorded from the stratum radiatum of the CA1 region, using an aCSF-filled, low resistance (2–5 M Ω), monopolar borosilicate glass electrode (recording electrode).

After this control period, stimulation intensity was adjusted to evoke 30% of the maximal fEPSP response, and then four high frequency trains of stimuli (100Hz for 1 second, 60 second intervals) were applied to the slice. Stimulation intensity was then again returned to that required to evoke 50% of the maximal fEPSP response and evoked fEPSPs (0.05Hz) were recorded for at least a further hour. Each experiment was completed with the generation of a final input/output curve in an identical manner to that generated at the beginning of the protocol. Experiments were excluded from further analysis if the maximum fEPSP slope at the end of the experiment deviated by more than 0.5mV/ms of the control value, that experiment was excluded.

2.8.4. Analysis of electrophysiological recordings.

For analysis, fEPSP slopes were normalised to 100% with respect to the mean slope of the fEPSPs recorded during the 20-minute control period preceding induction of LTP. Post-high frequency stimulation, fEPSPs were normalised to the same baseline. (Figure 2.12: shows the manner in which fEPSP slopes were analysed).

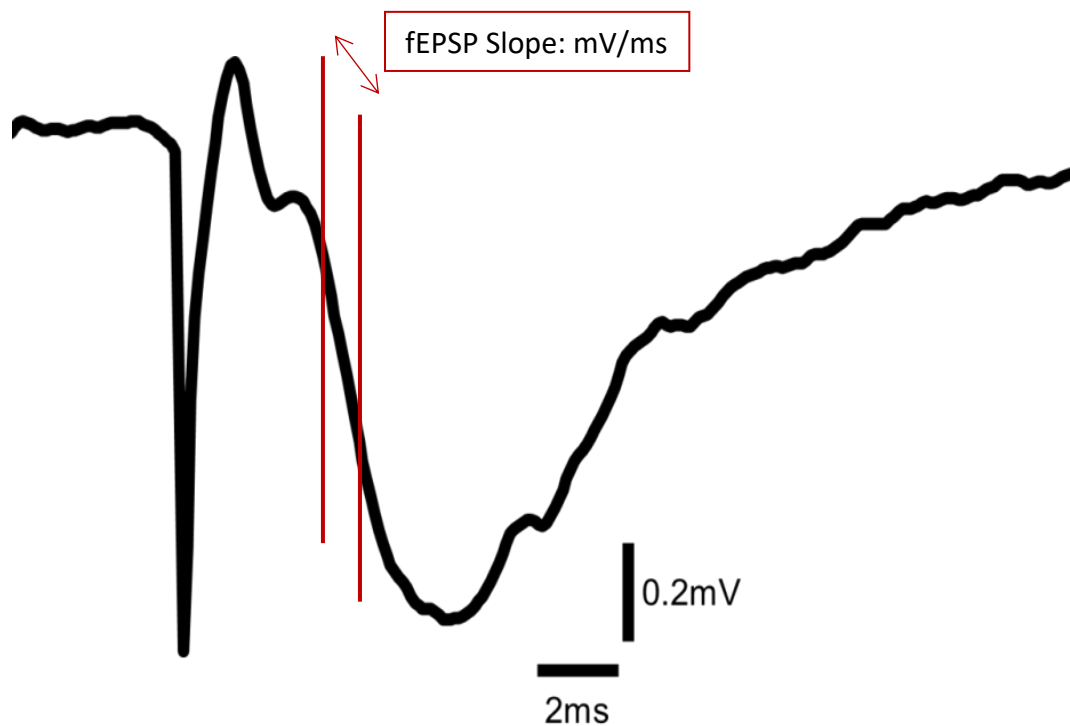


Figure 2.12: A representative trace of how the fEPSP slope (mV/ms) is analysed on each sweep and normalised to 100%.

2.9. Intervention Study

The intervention study included an intrathecal injection of xIL-6R or saline, NOR as well as LTP. The NOR and LTP protocols used were the same as those outlined in the control experiments described above.

2.9.1. Intrathecal Injection

Following the protocol described in Hylden *et al.* (1980), intrathecal injections were administered to male WT and *mdx* mice (8 – 12 weeks old) using disposable 30-gauge sterile BD Microlance™ Stainless Steel Needles (Becton Dickinson, New Jersey, USA) attached to a Hamilton™ 700 Microliter™ Syringe: Luer tip (Hamilton, Reno, USA).

Mice were anaesthetised with isoflurane (5% Isoflurane in oxygen, Harvard Apparatus), and the hair from the lower lumbar region removed. Each mouse was placed over a 50ml tube in order to expose the space between vertebrae. 70% ethanol and surgical Povidone-iodine were used to prepare the exposed area for injection. The mouse was then injected intrathecally into the subarachnoid space between L5 and L6 (Figure 2.13 below). The success of accurate spatial insertion of the needle into the subarachnoid space was confirmed by the observation of a tail flick from the anaesthetised animal caused by the insertion of the needle into the correct area. A volume of 8µl of either xIL-6R antibody (made up in saline to 0.2mg/kg) or saline was injected. After injection, mice were placed into an animal holding chamber, where warm air was circulated (Thermacage, Datasand limited,

United Kingdom) at an even temperature (30°C) until they woke up from anaesthesia as providing thermal support to rodents following anaesthesia or a surgical procedure is an important intervention and is emphasised in the Guide for the Care and Use of Laboratory Animals (*National Research Council Committee for the Update of the Guide for the Care and Use of Laboratory, 2011*). Mice were observed in order to confirm normal ambulation as incorrect injection into the spine can cause paralysis (which was an immediate humane endpoint following injection) and then placed back into their home cage for 24 hours before further experimentation (NOR and LTP).

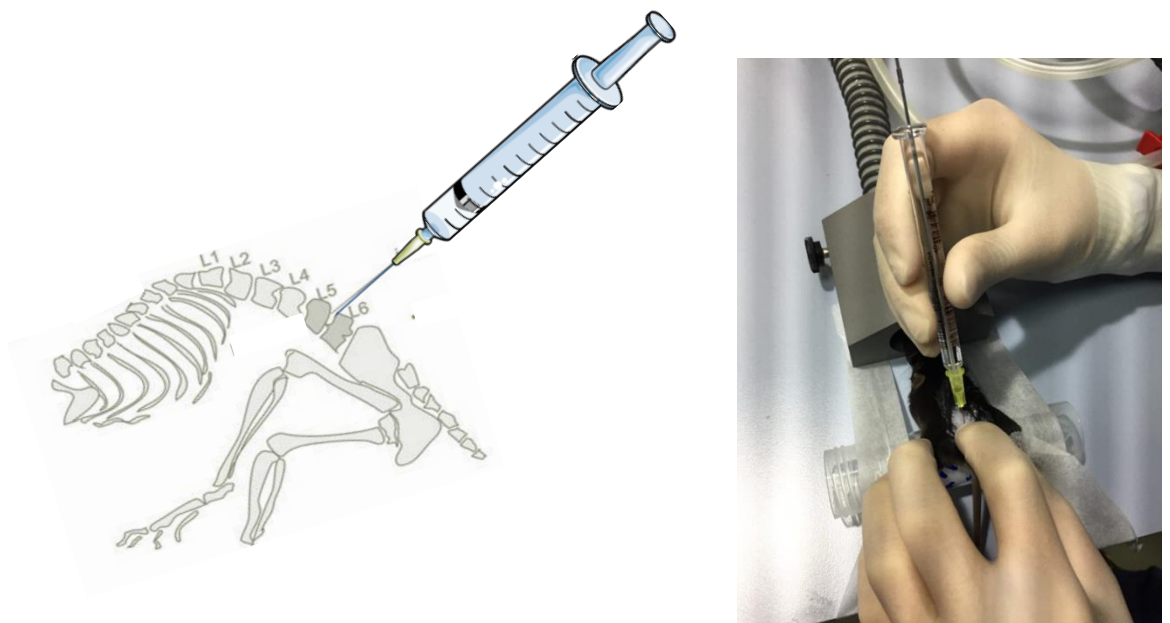


Figure 2.13: The schematic and photograph illustrate the procedure whereby an intrathecal injection was administered between the L5 and L6 vertebrae.

2.10. Statistical Analysis

Unless stated otherwise, GraphPad Prism Version 8.4.2 (GraphPad Software Inc., USA) was used for all statistical analyses. In addition, to test for normality and variance (Shapiro-Wilk test), unpaired or paired Student's t- tests were performed for comparisons between two groups with Welch's Correction or Mann-Whitney corrections, with one-way ANOVA or two-way ANOVA being performed for comparisons between multiple groups, unless stated otherwise. Appropriate *post-hoc* tests (*e.g.*, Bonferroni's test) were used for multiple comparisons. Statistical significance was accepted with p values < 0.05. Statistical significance is illustrated as *p<0.05, **p<0.01, ***p<0.001, on graphs throughout. Data are expressed as mean $\pm$ SD.

2.11. Materials

Anti-IL-6 receptor antibody (MR16-1) used throughout this study was a kind gift from Chugai Pharmaceuticals Ltd., Tokyo, Japan. This monoclonal anti-IL-6 receptor antibody is referred to as xIL-6R through this thesis. All other reagents were purchased commercially from Sigma Aldrich, Fischer Scientific or as stated in the text.

**Chapter 3: Dystrophin-deficient
mdx mice exhibit deficits in
molecular correlates of learning
and memory.**

3.1. Introduction

The *mdx* mouse is an animal model of DMD that mimics some aspects of the neurological and behavioural profile exhibited by individuals with the condition (Muntoni *et al.*, 1991; Chaussenot *et al.*, 2015), that are consistent with dysfunctions of the hippocampus and amygdala, such as a poor capacity to learn new tasks, and deficits in storing spatial memories as well as the processes generally involved in memory formation and consolidation (Vaillend *et al.*, 1995; Vaillend *et al.*, 2004; Chaussenot *et al.*, 2015). The hippocampus plays a central role in learning and memory (Anand & Dhikav, 2012) and is one of the regions where long term potentiation (LTP), proposed to be the molecular correlate of learning and memory (Bliss & Collingridge, 1993), occurs.

Interestingly, the literature has reported contradictory findings relating to LTP in *mdx* mice. For example, initial studies found no obvious deficits in hippocampal NMDA-dependent LTP or hippocampal-dependent spatial learning tasks in *mdx* mice that lacked Dp427 relative to controls (Sesay *et al.*, 1996), implying that hippocampal synaptic processes were intact within the *mdx* mouse. However, a later study by Vaillend *et al.* (2004) found that hippocampal-dependent LTP in *mdx* mice was in fact abnormally enhanced. This study also revealed deficits in long-term spatial memory and object recognition in *mdx* mice (Vaillend *et al.*, 2004). It has been suggested that one possible reason for the contrasting findings of Sesay *et al.* (1996) and Vaillend *et al.* (2004) may have been the particular training procedures utilised by Sesay *et al.* (1996) as they may have helped to alleviate any subtle memory deficits that were possibly present before training. Furthermore, Vaillend

et al. (2004) proposed that the abnormally enhanced hippocampal LTP observed in *mdx* mouse brain slices could, at least in part, be responsible for the memory deficits seen in these mice. Thus, somewhat counterintuitively, abnormally-enhanced synaptic potentiation and neuronal excitability might actually inhibit normal cognitive functioning (Vaillend *et al.*, 2004). Interestingly, a recent study by Bagdatlioglu *et al.*, 2020, showed that hippocampal-dependent spatial learning and memory was decreased in *mdx* mice, particularly long-term memory, which progressively worsened with age (Bagdatlioglu *et al.*, 2020).

Neurogenesis, where progenitor cells differentiate into neurons, occurs in the dentate gyrus, with adult-born neurons being integrated into pre-existing neuronal circuits (Gage, 2000). Such integration of new cells into neuronal circuits is proposed to contribute to hippocampal-dependent learning and memory processes (Wu & Hen, 2014). The dentate gyrus is positioned as the input zone of the hippocampus and normally has low neuronal activity compared to upstream cortical areas (Treves & Rolls, 1994). One study has shown that a soluble mediator released from the muscle of *mdx* mice (nitrous oxide) can have an effect on hippocampal neurogenesis (Deng *et al.*, 2009). Also, this study found that the cognitive defects in DMD could result from the pathologically high number of immature neurons in the circuitry of the DG, which could disrupt the established circuitry (Deng *et al.*, 2009). Given the contradictory reports in the literature about the effect of loss of dystrophin on behaviour and the processes of learning and memory consolidation, it is apparent that further studies need to be conducted to establish the role of dystrophin in the dysfunction of hippocampal neural circuitry.

3.1.1. Study objectives

- To determine if cellular pathways involved in cognitive function are altered in the absence of dystrophin.
- To characterise behaviours in the *mdx* mouse model of DMD.
- To assess hippocampal adult neurogenesis in *mdx* mice aged 8 weeks, 4 months and 8 months compared to age-matched WT mice.

3.1.2. Animals

- *Mdx* mice aged 8-12 weeks were used in this study for electrophysiological and behavioural experiments with age-matched WT comparators.
- *Mdx* mice aged 8-weeks-old, 4 months and 8-months-old were used for immunofluorescence with age-matched WT comparators.

3.1.3. Techniques used

- Extracellular electrophysiology (more detail in methods section 2.8, pg. 74).
- Immunofluorescence on whole brain sections (see methods section 2.3.2, pg. 54).
- Novel Object Recognition test (see methods section 2.7.2, pg. 70)
- Open field behaviour (see methods section 2.7.3, pg. 73).
- Calcium imaging (see methods section 2.4. pg. 59).

3.2. Dystrophin is absent in all regions of the hippocampus in the *mdx* mouse.

Immunofluorescence labelling of the hippocampus of 8–12-week WT and *mdx* mice was carried out as described (see Methods section 2.3.2, pg. 39). A dystrophin polyclonal antibody was used for immunofluorescent labelling of this structural protein. Dystrophin is present in all 3 regions of the hippocampus investigated, namely CA1, CA3 and the dentate gyrus (DG) in WT mice with dystrophin present in the pyramidal cell layer of the CA1 and CA3 regions, the granule cell layer of the DG and in interneurons (indicated by white arrows in Figure 3.1). In contrast however, there was a complete absence of dystrophin labelling in any of the same regions in *mdx* mice (Figure 3.2). The appropriate positive controls and negative controls were also conducted to ensure that no random binding of either the primary or secondary antibody was occurring.

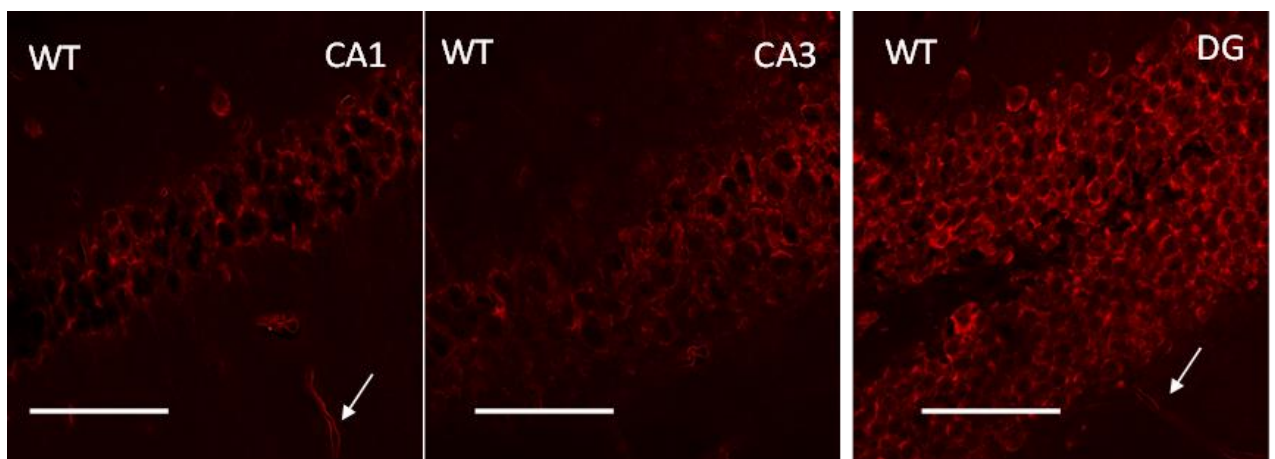


Figure 3.1: Immunofluorescent labelling of dystrophin in CA1, CA3 and DG regions of the hippocampus of WT mice. White arrows indicate interneurons. Scalebars = 70 μ m.

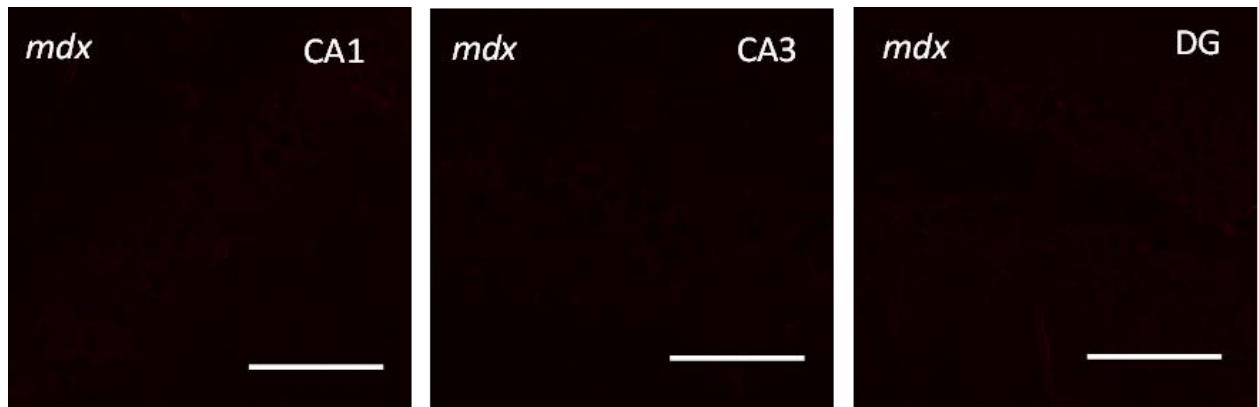


Figure 3.2: Immunofluorescent labelling of dystrophin in CA1, CA3 and DG regions of the hippocampus of *mdx* mice. Scalebars = 70 μ m

3.3. Hippocampal long-term potentiation (LTP) is reduced in dystrophin-deficient *mdx* mice.

Given the contradictory findings regarding the induction of LTP, the proposed molecular correlate of learning and memory formation, in *mdx* mice (Sesay *et al.*, 1996; Vaillend *et al.*, 2004), we investigated LTP induction in *mdx* mice and compared it to that induced in WT slices.

Field excitatory post-synaptic potentials (fEPSPs) generated by stimulating the Schaffer Collaterals (SC) running from area CA3 to synapses which terminate on the dendrites of CA1 neurons, which reside within the *stratum radiatum* above the hippocampal fissure (Malenka & Nicoll, 1999), were recorded from the CA1 region of the hippocampus.

Monopolar stimulation of SC fibres was performed by placing a stimulating electrode on the border between areas CA1 and CA3. First, input/output curves

were constructed from a range of stimulus intensities before the induction of LTP, to assess synaptic transmission and the relationship between excitatory input and firing efficiency (Figure 3.3). No significant difference was detected between the two genotypes (Two-way ANOVA, fEPSP slope: $F(1, 8) = 0.2327$, $p = 0.6425$) under these basal conditions.

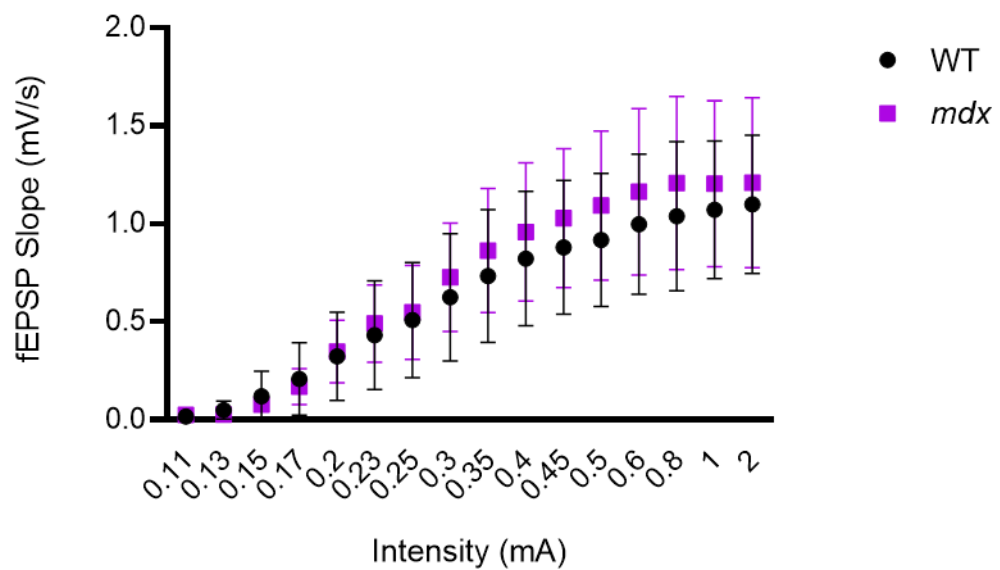


Figure 3.3: Input/output curves constructed during baseline recordings. There is no difference between pooled baseline recordings in WT ($n=6$) and mdx mice ($n=6$). Mixed-effect analysis ($p>0.05$)

Following stabilisation of the electrical recordings, which could take up to an hour, and generation of the baseline input/output curve, fEPSP baseline recordings were made for 20 minutes, stimulating every 30 seconds, using a current that induced fEPSPs that were 50% of the maximal size that could be generated. This recording period was followed by the application of four trains of high-frequency stimuli (HFS – 4 x 100 pulses, 100Hz for 1s/60s), with a current that induced fEPSPs that were 30% of the maximal size that could be generated. Immediately following the HFS

protocol, stimulation frequency and intensity was returned to baseline parameters, where we saw a dramatic increase in the slope (mV/ms) of the fEPSP for 5 minutes after HFS application. This increase in slope was observed in both WT (n=9) and *mdx* (n=6) mice. However, the mean size of the fEPSP slope was decreased in *mdx* mice compared to WT (WT $231 \pm 34.63\%$ vs *mdx* $167.8 \pm 37.61\%$; $p = 0.0052$, figure 3.4b). Furthermore, for the entire duration of recording, 80 minutes, following HFS, LTP was significantly decreased in dystrophin-deficient *mdx* hippocampal slices (n=6) compared to that in slices from WT mice (n=9) (Genotype factor, $F(1, 13) = 8.517$, $p = 0.0120$, Interaction of time and genotype factors, $F(179, 2327) = 1.370$, $p = 0.0012$, two-way repeated-measures ANOVA starting at 1 min after stimulation).

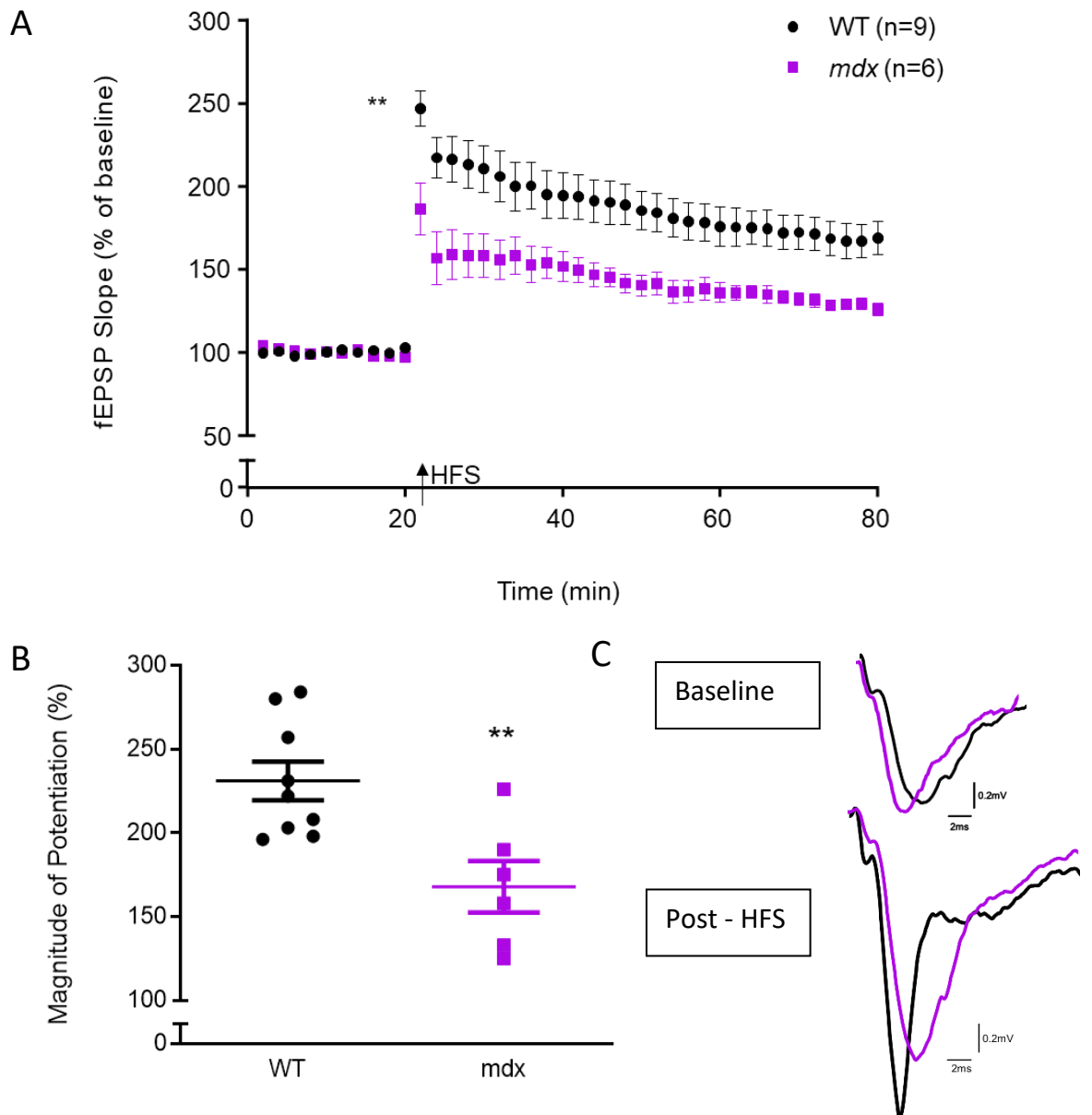


Figure 3.4: Long-term potentiation in the SC-CA1 pathway of the hippocampus is significantly decreased in mdx mice relative to WT mice.

- A) Graph showing averaged time course of Schaffer collateral-evoked synaptic responses from WT (black circles) and mdx (purple squares) in the hippocampal region using extracellular field potential recordings in CA1 stratum radiatum (WT n=9, mdx n=6). Two-way ANOVA, $p < 0.01^{**}$
- B) After high-frequency stimulation (HFS – 4 x 100 pulses, 100Hz for 1s/60s), the fEPSP slope in the mdx slices was significantly reduced compared to WT ($p = 0.0012$). Unpaired t test, $p < 0.01^{**}$
- C) Representative traces show fEPSPs before and after high frequency stimulation to induce LTP in WT and mdx mice. At baseline, WT (black) and mdx (purple) fEPSP slopes (mV/ms) are similar, whereas, post-HFS, fEPSP slopes are significantly decreased in treated mdx mice.

3.4. Recognition memory is not deficient in *mdx* mice.

To determine if the suppression of hippocampal LTP resulted in changes in memory formation in the *mdx* mice, we assessed WT and *mdx* mice using Novel Object Recognition (NOR) testing. NOR tests are used to assess recognition memory by testing a rodent's ability to recognize a previously presented object or stimulus (Ennaceur & Delacour, 1988). NOR uses two similar objects on a first testing day, and the mouse is left to explore for 10 minutes. 24 hours later, one of the similar objects is replaced with a novel object and once again the mouse is left to explore the area for 10 minutes. A percent of time spent exploring the novel object relative to the total time spent exploring both objects can be a measure of novel object recognition (Benice *et al.*, 2006; Sarkisyan & Hedlund, 2009; Broadbent *et al.*, 2010; Oliveira *et al.*, 2010). This concept can be represented by *Recognition Index* (RI), i.e., the time spent investigating the novel object relative to the total object investigation [$RI = T_N / (T_N + T_F)$], and it is the main index of retention (Mumby *et al.*, 2002; Piterkin *et al.*, 2008; Botton *et al.*, 2010; Gaskin *et al.*, 2010; Schindler *et al.*, 2010). In our study however the RI was not significantly altered between WT (0.75 ± 0.02 ; n=14) and *mdx* (0.76 ± 0.01 , n=14; p=0.7628) mice (figure 3.5).

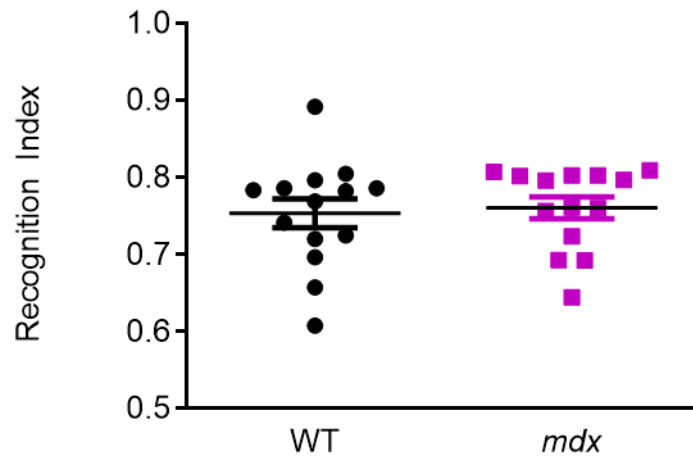


Figure 3.5: Recognition memory is not significantly different between WT and mdx mice. The ratio of time spent exploring a novel object versus a familiar object (recognition index) was comparable in mdx mice and WT mice. Unpaired t test, $p > 0.05$

3.5. Behavioural characteristics are altered in *mdx* mice relative to WT.

Even though recognition memory was not altered in *mdx* mice compared to controls, other behavioural and cognitive issues are also prevalent in individuals with DMD. For example, boys with DMD are commonly diagnosed with behavioural problems such as Attention-Deficit/Hyperactivity Disorder (ADHD), anxiety, depression, or other mood disorders (Hendriksen & Vles, 2008; Pane *et al.*, 2013a; Banihani *et al.*, 2015). We examined the *mdx* mouse model to determine if any behaviours related to these disorders were presented as part of the phenotype. Using a behavioural assay in an open field arena, we assessed *mdx* mice for central area frequency, freezing frequency, rearing frequency, jumping frequency, grooming frequency and faecal output (methods section 2.7.3, pg. 43).

Central area frequency is often used as a measure of anxiety in mice, such that the more time spent on the periphery of an open field, close to the walls, the more anxious the mouse (Holmes *et al.*, 2003). The central area frequency (number of times the mouse had 4 paws in central area) of *mdx* mice was comparable to WT (figure 3.6a, n=19, p=0.8186, WT 23.94 ± 6.847 vs *mdx* 23.37 ± 8.214).

Increased faecal output is also linked to increased anxiety (Hall, 1934), however, the faecal output (number of boli) in this study was comparable in *mdx* and WT mice (figure 3.6b, n=19, p=0.3431, WT 3.300 ± 2.536 vs *mdx* 2.632 ± 1.707).

Another measure of anxiety is freezing, which has been linked to fear conditioning (Blanchard & Blanchard, 1988). Interestingly, *mdx* mice displayed a significant increase in freezing frequency (number of times the mouse did not move for more than 2 seconds) in comparison to WT (figure 3.6c, WT 10.95 ± 4.839 vs *mdx* 13.89 ± 5.098 ; n=18, p=0.0383).

Rearing frequency (number of times the mouse reared its front paws, against the wall or mid-air) and jumping frequency (number of times the mouse jumped against the wall, with all 4 paws in the air) are related to exploratory and escape behaviours and was shown to be comparable between *mdx* and WT mice, - rearing frequency (figure 3.6d, n=17, p=0.2607, WT 88.56 ± 18.63 vs *mdx* 96.65 ± 23.08) and jumping frequency (figure 3.6e, n=18, p=0.1687, WT 0.8235 ± 1.334 vs *mdx* 1.722 ± 2.321).

Lastly, grooming frequency has been linked to autism spectrum disorder (ASD) as it is regarded as a repetitive behaviour (Silverman *et al.*, 2010), in *mdx* mice, the grooming frequency was comparable to WT mice (figure 3.6f, n=19, p=0.9108, WT 2.947 ± 1.224 vs *mdx* 2.889 ± 1.844).

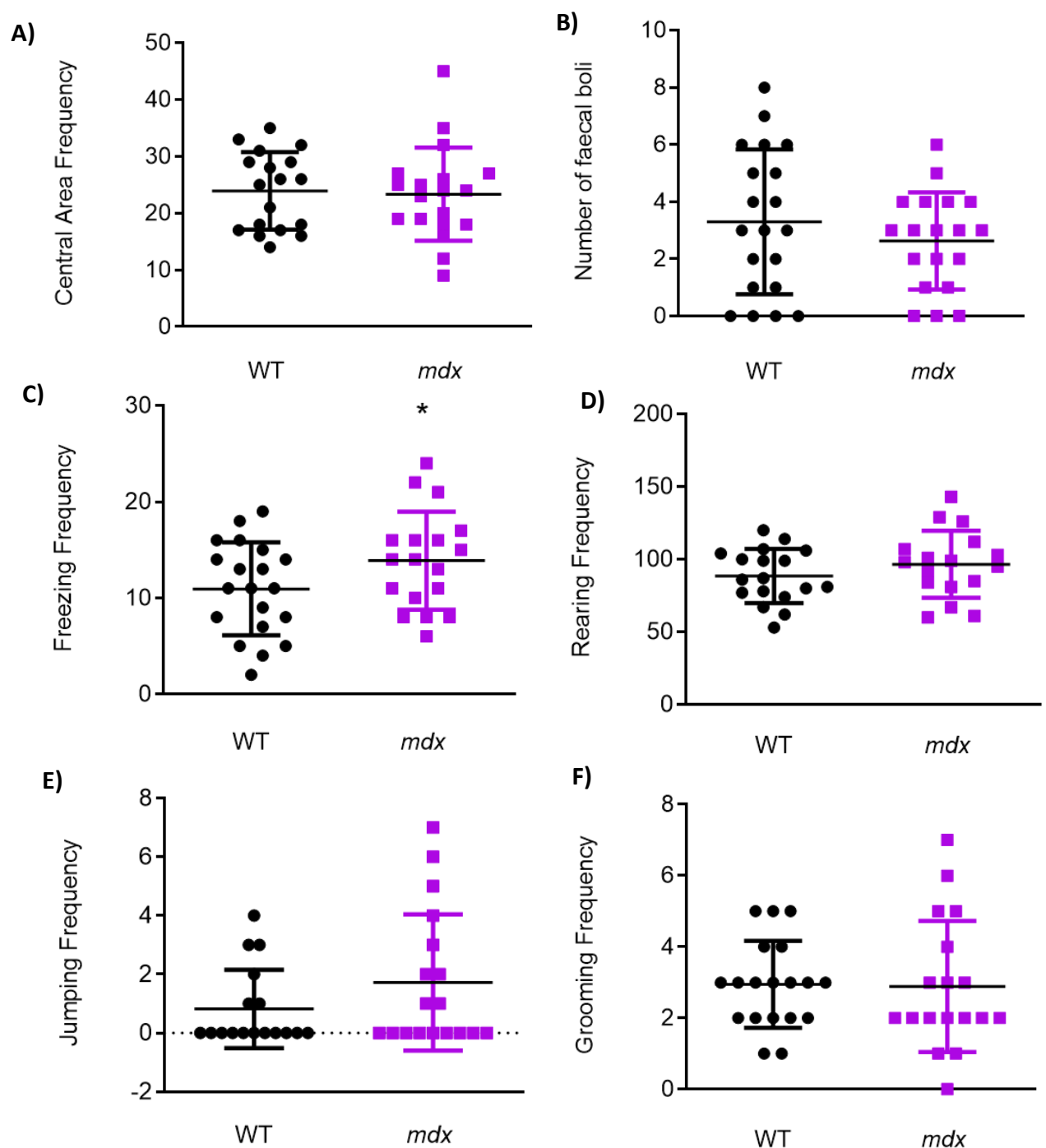


Figure 3.6: Behaviours of mdx and WT mice in an open field.

The data plot illustrates that A) Central area frequency, where all four paws were in the central area of the open field arena, in mdx mice was similar to WT. B) Faecal output (number of boli) in mdx mice was similar to WT. C) Freezing frequency was significantly higher in mdx mice in comparison to WT. D) Rearing frequency in mdx mice was similar to WT. E) Jumping frequency was similar in mdx mice to WT. F) Frequency of grooming in mdx mice was similar to WT. Unpaired t test, $p < 0.05$ *

3.6. GABA_A receptor expression in neurons in the dentate gyrus of *mdx* mice is similar to that in WT mice.

GABA has long been associated with mood disorders such as anxiety spectrum disorders (Nutt & Malizia, 2001; Lydiard, 2003; Hasler *et al.*, 2008) and depression (Gold *et al.*, 1980; Krystal *et al.*, 2002; Chang *et al.*, 2003).

Previous research has indicated that GABA_A receptor clustering is aberrant in the absence of dystrophin (Vaillend *et al.*, 2010). The dentate gyrus in the hippocampus is one of two regions in the brain where neurogenesis occurs (Gage, 2000) and GABA is known to have a depolarizing effect on neural progenitor cells and immature neurons, both during developmental and adult neurogenesis (Ge *et al.*, 2006). As such, the dentate gyrus is a region of interest for any possible aberrant changes that could impact cognition. Using immunofluorescence, we examined GABA_A receptor expression in whole brain sections, looking specifically at the dentate gyrus in the hippocampus of WT and *mdx* mice at 4 and 8 months of age (we chose these two different time points in order to determine if there are any progressive changes with age).

The number of GABA_A receptor-positive neurons was counted in three different regions of the dentate gyrus in WT and *mdx* mice (n=3-4 animals), specifically within two random regions of the granule cell layer and hilus, which were then pooled for a total number of GABA_A receptor-positive neurons in the dentate gyrus as a whole. The numbers of GABA_A receptor positive neurons did not differ significantly between WT and *mdx* dentate gyrus in either of the two age groups (figure 3.7, Two-way ANOVA, n=3-5 animals per group, 2-3 whole brain sections per animal, 3

regions in the dentate gyrus, of the granule cell layer, $F(1, 10) = 0.08065$, $p=0.7822$), although this does not discount the fact that these neurons may be dysfunctional.

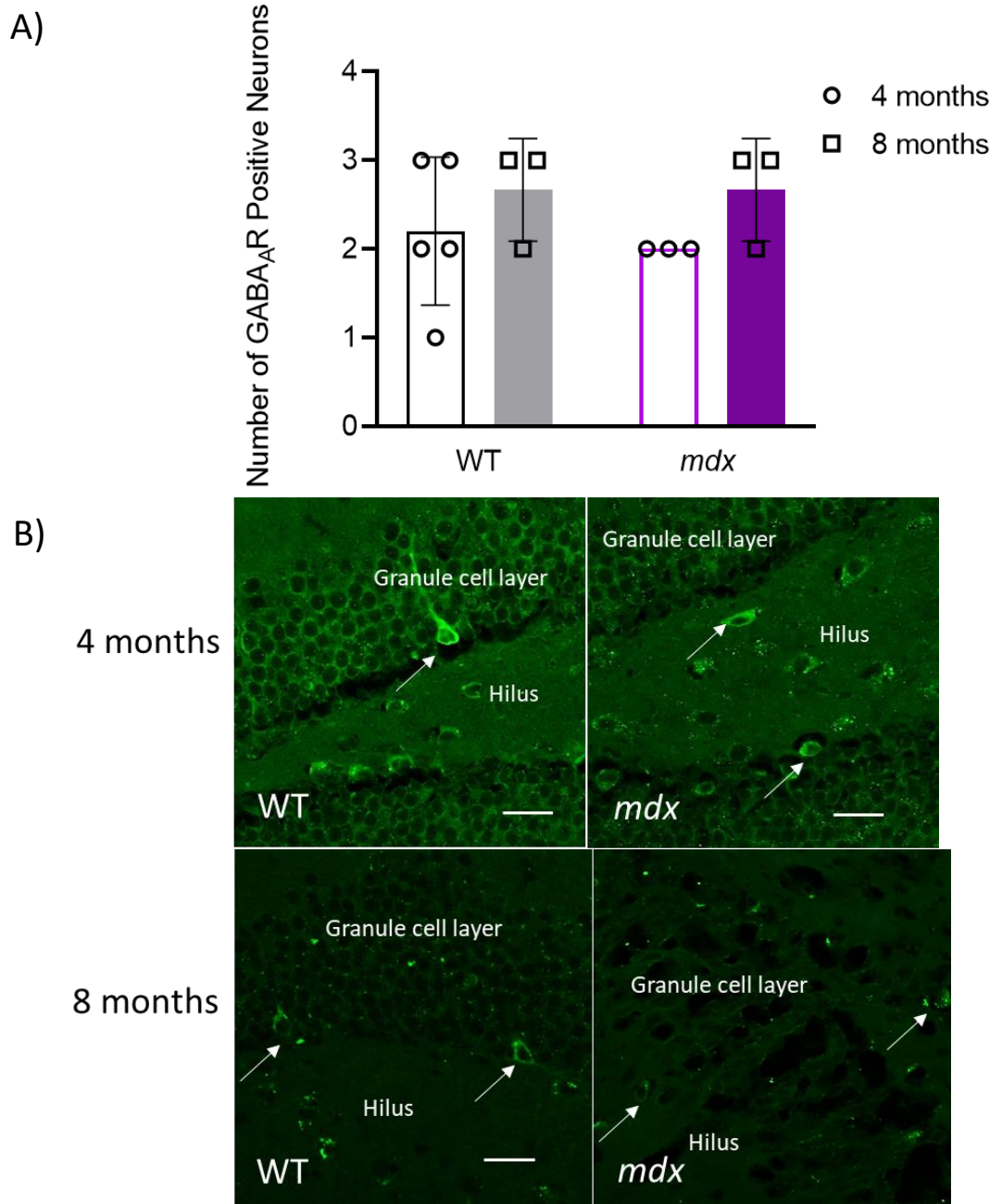


Figure 3.7: The number of GABA_AR positive neurons in the granule cell layer and hilus of the dentate gyrus of WT and mdx mice at two different ages (4 months and 8 months) is similar.

The number of GABA_AR-positive neurons did not differ significantly in the dentate gyrus of WT and mdx mice between the two age groups.

Representative immunofluorescence images of the number of GABA_AR positive neurons in WT and mdx mice at 4 and 8 months. GABA_AR-positive immunostaining is similar in WT and mdx mice at 4 and 8 months. Tissue was stained with a mouse GABA_AR primary antibody and a FITC (goat anti-mouse) secondary antibody. White arrows indicate GABA_AR-positive neurons in the granule cell layer and hilus in both WT and mdx. Scale bars are 30μm. Two-way ANOVA, $p>0.05$

3.7. Loss of dystrophin increases neuronal excitability in cultured hippocampal neurons

** Experiments completed by Kimberley Stephenson and Michael Vaughan and data analysed by Claire Denley.*

Although we did not observe changes in the absolute number of GABA-expressing hippocampal interneurons in dystrophic tissue, aberrant GABA_A receptor clustering (Vaillend *et al.*, 2010) could nonetheless have an impact on the neuroregulatory actions of GABAergic neurons (McCormick & Contreras, 2001; Saxena & Caroni, 2011). Using hippocampal neurons cultured (4-5 days) from neonatal mice (aged 3-5 days old), we next sought to examine basal neuronal activity by monitoring intracellular neuronal calcium levels. No apparent differences were observed in cell survival or network formation of hippocampal neurons cultured from WT and *mdx* mice (Figure 3.8). However, when we examined baseline cellular calcium levels, unstimulated *mdx* hippocampal neurons exhibited a significantly greater frequency of calcium peaks than observed in WT controls (Figure 3.9c, unpaired t test, WT 0.3234 ± 0.1635 ; n=16 neurons vs *mdx* 0.7443 ± 0.1797 ; n=19 neurons, $p < 0.001$) over a period of 60 minutes (4000 seconds). This can be observed from the representative traces in figure 3.9, intracellular calcium activity in WT neurons (figure 3.9a) is almost non-existent in comparison to that observed in *mdx* neurons (figure 3.9b) indicating neuronal hyperexcitability in *mdx* neuronal cells.

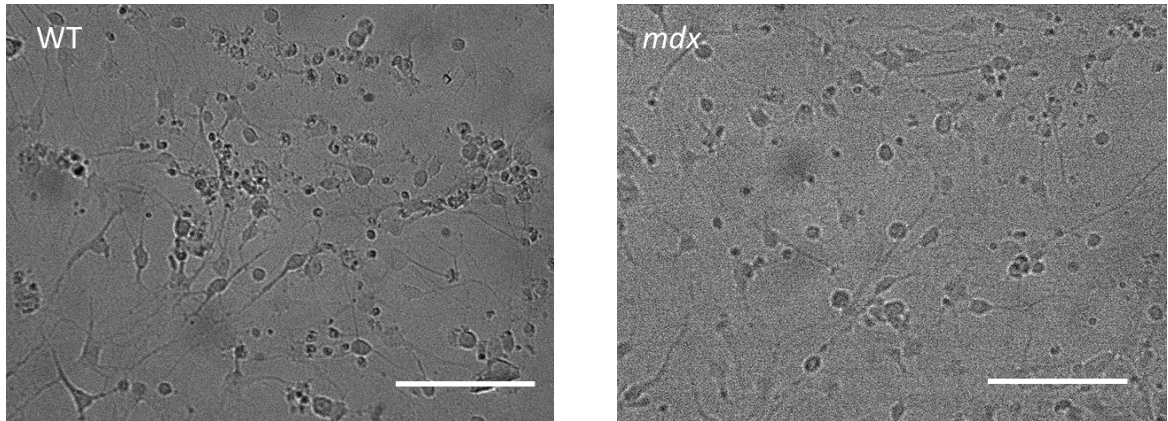


Figure 3.8: Brightfield images of mdx and WT neurons, 4-5 days after primary culture. The morphology of WT and mdx neurons is similar. Scalebars are 35 μ m

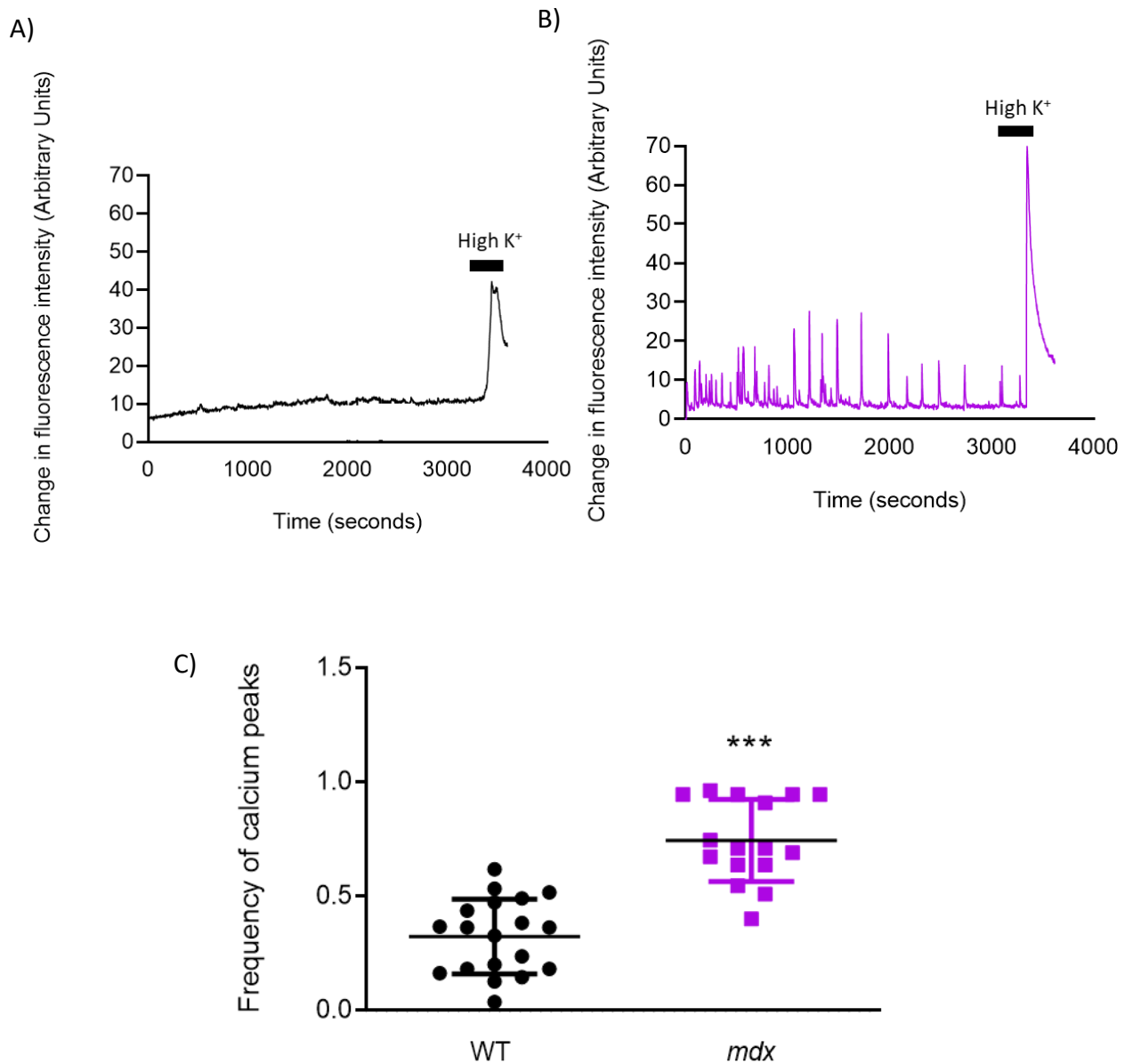


Figure 3.9: Baseline activity in cultured mdx hippocampal neurons is increased.

Representative traces of cellular activity and frequency of peaks under basal conditions (HBSS) in A) WT mice are different to the cellular activity in B) mdx mice (purple). C) This difference is significantly different. Unpaired t test, $p < 0.001^{***}$

3.8. Neuronal density in the dentate gyrus decreases with age in *mdx* mice.

Using a pan-neuronal biomarker, NeuN (neuronal nuclear protein) to identify postmitotic neurons, we counted the number of neurons in the granule cell layer of the dentate gyrus. In dystrophin-expressing WT mice, neither neuronal density nor neuronal number in the granule cell layer of the dentate gyrus was changed between 4 months (adult mouse) and 8 months (aged mouse). However, we found that there was a significant decrease in neuronal number in the 8-month-old *mdx* mice relative to age-matched WT controls (Two-way ANOVA, $n=3-4$ animals per group, 2-3 whole brain sections per animal, 3 regions in the dentate gyrus, of the granule cell layer). The interaction between the WT and *mdx* (strain vs age) at 8 months was significantly different (figure 3.10a, $F(1, 9) = 136.8$, $p < 0.0001$) and the age effect of 4 months vs 8 months in the *mdx* mice was also significantly different (figure 3.10, $F(1, 9) = 121.6$, $p < 0.0001$).

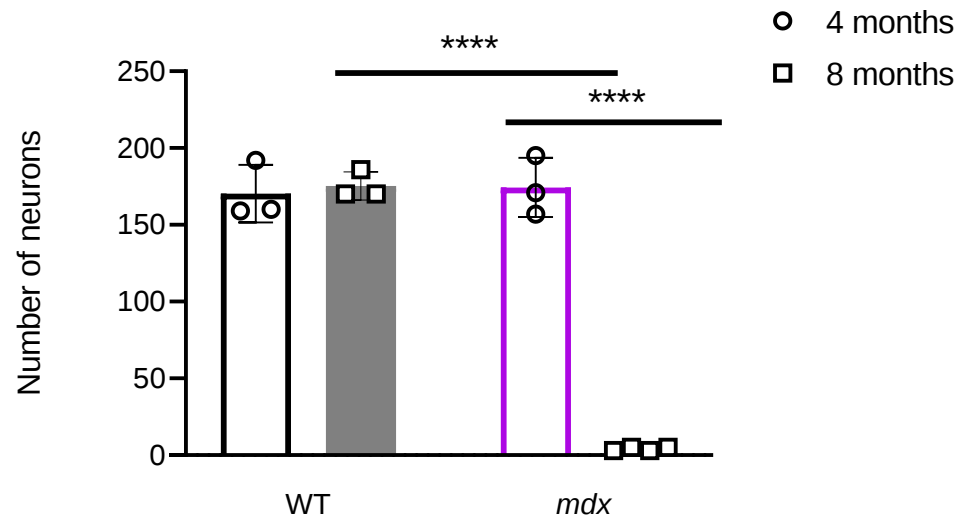
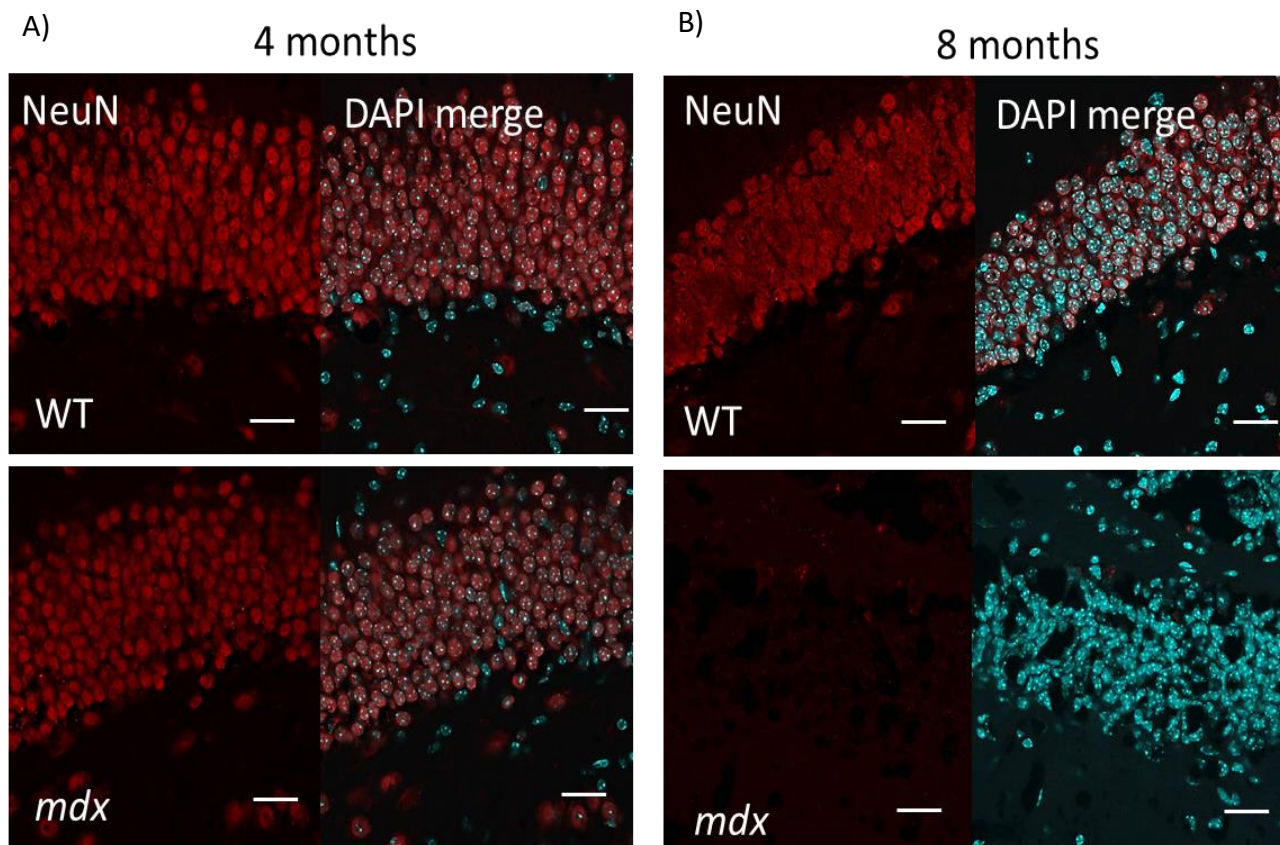


Figure 3.10: Neuronal density in the dentate gyrus decreases with age in mdx mice (4 and 8 months).

*Mean neuronal number in WT slices was the same in the two age groups studied whereas there was a significant decrease in neuronal number in the 8-month-old mdx slices compared to 4-month-old mdx and WT. Two-way ANOVA, $p < 0.001$ *****

Representative immunofluorescence images (figure 3.11a) show that there was no difference in the number of neurons in WT and *mdx* mice at 4 months old. DAPI, a nuclear marker, was used as a control stain to determine the number of cells per region. However, in the representative images (figure 3.11b), the significant decrease in the number of neurons in the granule cell layer of the dentate gyrus in 8-month-old *mdx* can be seen as an absence of neurons. However, there are other cell types present as evidenced by the DAPI staining. It is also clear that there is a change in gross cell morphology within the 8-month-old *mdx* mice in that the cells within the granule cell layer are disorganised and less densely packed relative to either the WT or 4-month-old *mdx* mice.



*Figure 3.11: Immunofluorescence images looking at NeuN staining for neurons in the dentate gyrus. A) There is no difference between the number of neurons in WT and *mdx* mice at 4 months old. However, B) there is a significant decrease in the number of neurons in *mdx* mice at 8 months relative to age-matched WT comparators and 4-month-old *mdx* mice. DAPI is used as a nuclei marker for all cells. Scalebars are 30µm.*

That said, there is a difference in the gross cell morphology in 8-month-old WT mice in that the neurons are more densely packed, which could be an age-related change. Further, in 8-month-old *mdx* mice, there appear to be large fluid filled structures disrupting the cell layer in the granule cell layer (figure 3.12b and c), that are not present in the 4-month-old WT and *mdx* mice (figure 3.12a). This large fluid filled structure may have displaced the neurons, resulting in a significant decrease in 8-month-old *mdx* mice.

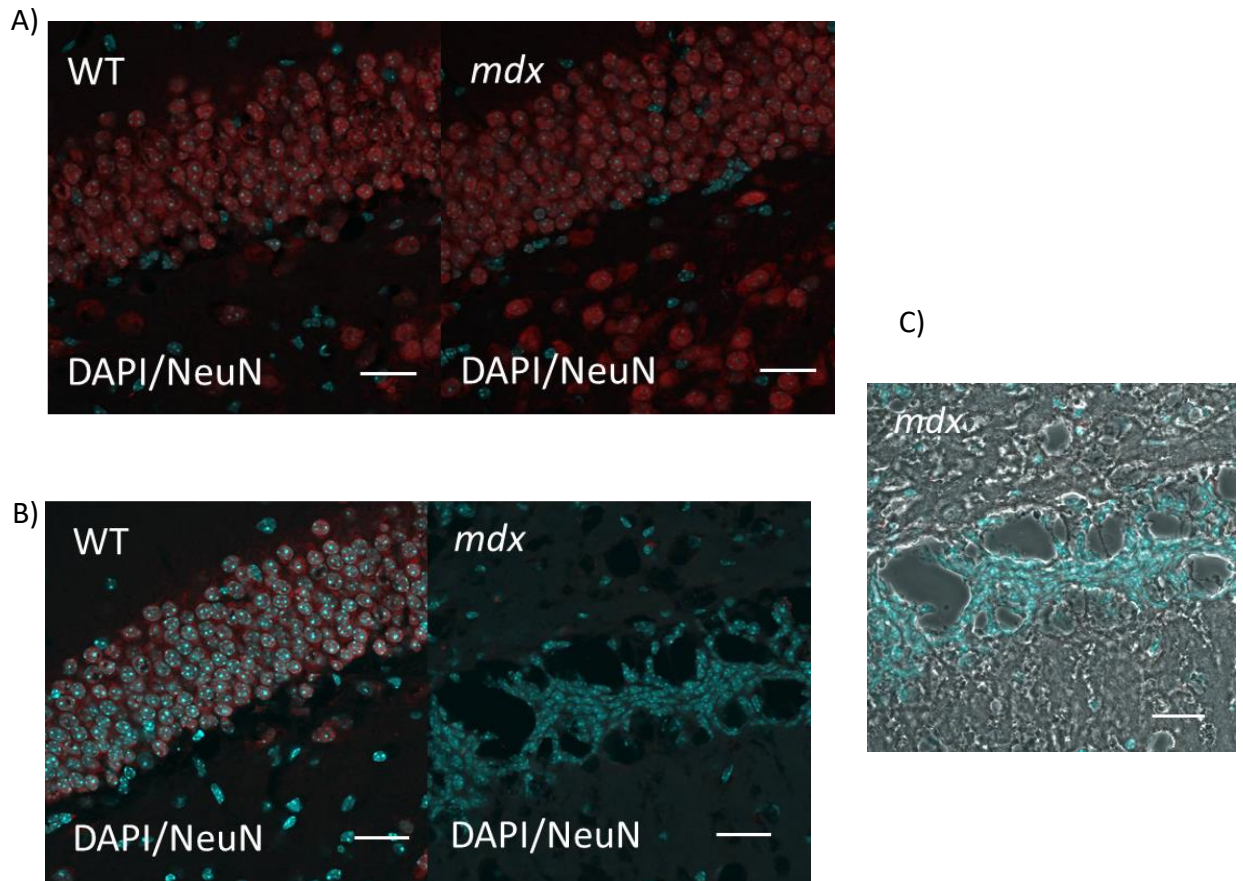


Figure 3.12: Immunofluorescence and brightfield images looking at NeuN staining for neurons in the dentate gyrus. Brightfield and DAPI/NeuN stained images of A) 4-month-old and B) 8-month-old WT and mdx mice. C) Brightfield image of the fluid filled structures that were present in the granule cell layer of 8-month-old mdx mice. Scalebars are 30 μ m.

3.9. The number of DCX-positive cells in the dentate gyrus of *mdx* mice decreases with age.

Doublecortin (DCX) is a microtubule-associated protein that is required for normal neocortical and hippocampal development and which has also been associated with hippocampal adult neurogenesis. As such, it is a suitable marker for neurogenesis (Couillard-Despres *et al.*, 2005).

For this part of the investigation, the number of DCX-positive cells in 8-week-old, 4-month-old and 8-month WT and *mdx* mice was counted. Three different age groups were chosen for study as hippocampal adult neurogenesis and synaptic plasticity are both greatly affected by the aging process (Rao *et al.*, 2005). For our analysis, three regions of the granule cell layer in the dentate gyrus were investigated. These regions were determined as an area where there was the greatest accumulation of cells in a 3cm x 3cm area (indicated by a white box on representative images).

In WT mice, the number of DCX-positive cells decreased between 8-week-old and 4-month-old mice, but this decrease was not statistically significant (unpaired t test, $n = 5$ animals, 8-week-old 5 ± 1 vs 4-month-old 3.8 ± 0.8367 , $p = 0.0736$). Further, there was no change in the number of positive DCX cells between 4-month-old and 8-month-old WT mice ($n = 4-5$ animals, unpaired t test, 4-month-old 3.8 ± 0.8367 vs 8-month-old 3.750 ± 0.9574 , $p > 0.05$). On the other hand, the number of DCX-positive cells was significantly decreased in *mdx* mice in comparison to WT mice as such an age-effect was noted in *mdx* mice only.

When comparing the number of DCX-positive cells at each of the three ages between WT and *mdx* mice, there are significant differences in the number of cells in each group. Fewer cells were positively stained for DCX in 2-month-old *mdx* mice as compared to WT mice (figure 3.13, unpaired t test, n=5 animals, WT 5 ± 1 vs *mdx* 2.2 ± 1.095 , $p=0.0029$). At 4-months-old, the number of DCX-positive cells was also significantly decreased in *mdx* mice in comparison to WT (figure 3.14, unpaired t test, n=3-5 animals, WT 3.8 ± 0.8367 vs *mdx* 2.667 ± 0.5774 , $p=0.0436$). At 8-months of age, the number of DCX-positive cells was also significantly decreased in *mdx* mice in comparison to WT (figure 3.15, unpaired t test with Welch's correction, n=4 animals, WT 3.75 ± 0.9574 vs *mdx* 1.75 ± 0.5 , $p=0.0167$).

Representative immunofluorescence images for each age group showed a decrease of DCX-positive cells (red stain) in the granule cell layer of *mdx* mice in comparison to WT. In all analysis, DAPI, a marker of nuclei, was used as a cellular marker.

Decreased neurogenesis in aging (Mathews *et al.*, 2017), was exacerbated in *mdx* mice with a clear difference between strains ($F(1, 20) = 31.6$, $p < 0.0001$). No interaction between strain and age was noted.

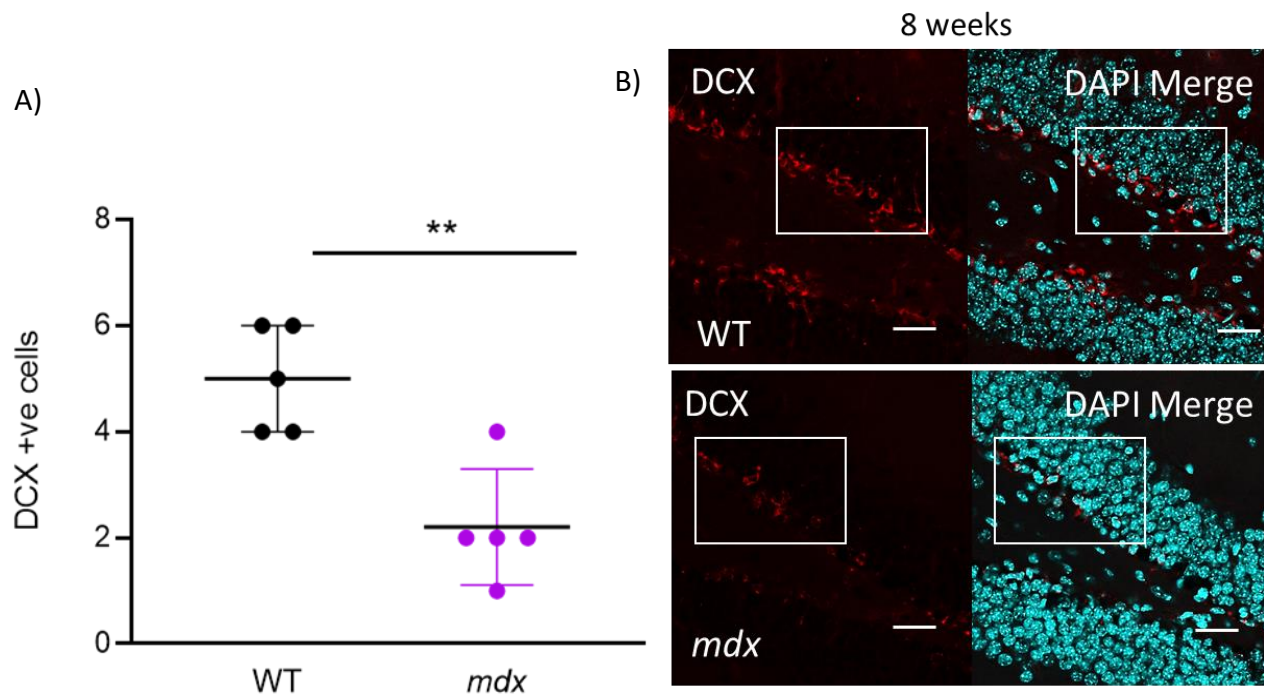


Figure 3.13: The number of DCX-positive cells in the granule cell layer of the dentate gyrus is decreased 8-week-old mdx mice.

The number of positive DCX cells is significantly reduced in mdx mice in comparison to WT.

*Immunofluorescence images of DCX-positive cells (red) counted in the white box is decreased in mdx compared to WT. Scalebars are 30µm. Unpaired t test, $p < 0.01^{**}$*

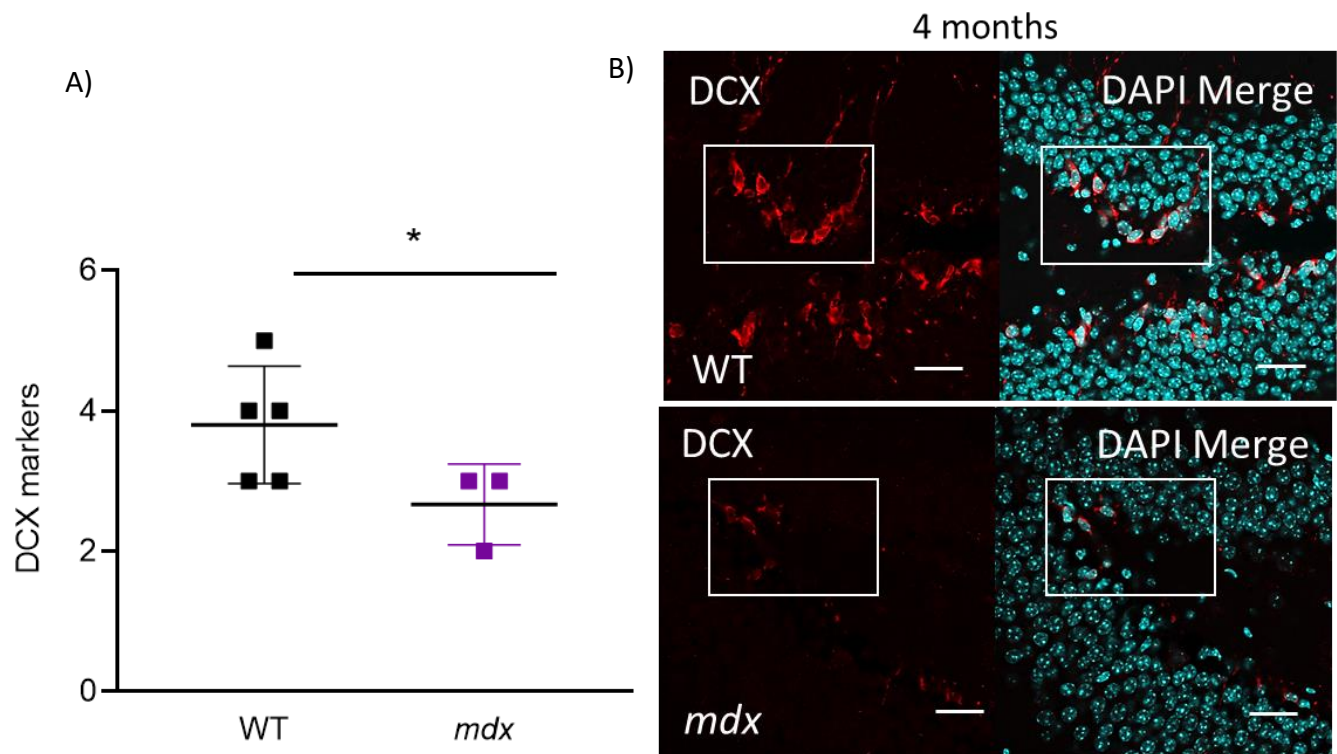


Figure 3.14: The number of DCX-positive cells in the granule cell layer of the dentate gyrus is decreased in 4-month-old mdx mice.

The number of positive DCX cells is significantly reduced in mdx mice in comparison to WT.

Immunofluorescence images of DCX-positive cells (red) counted in the white box is decreased in mdx compared to WT. Scalebars are 30µm. Unpaired t test, $p < 0.05^$*

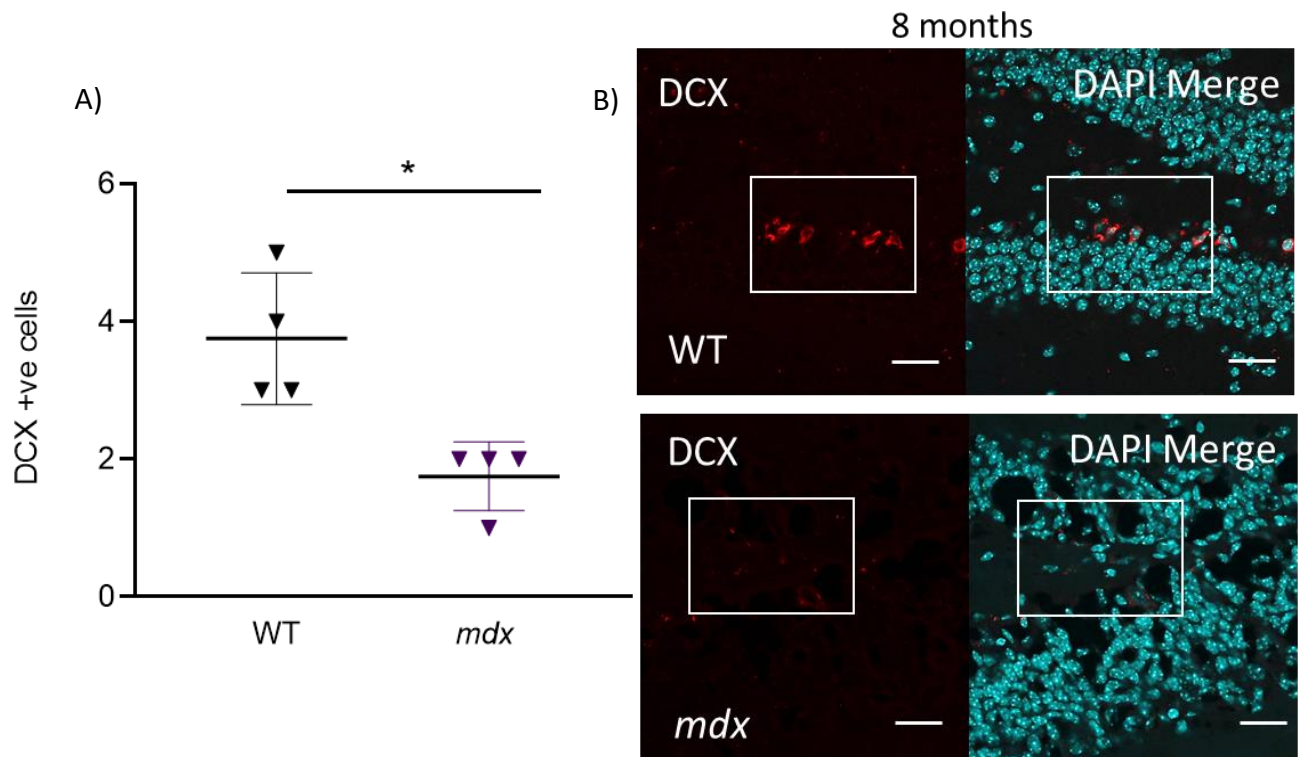


Figure 3.15: The number of DCX positive cells in the granule cell layer of the dentate gyrus is decreased in in 8-month-old mdx mice.

The number of positive DCX cells is significantly reduced in mdx mice in comparison to WT

Immunofluorescence images of DCX positive cells (red) counted in the white box is decreased in mdx compared to WT. Scalebars are 30µm. Unpaired t test, $p < 0.05^$*

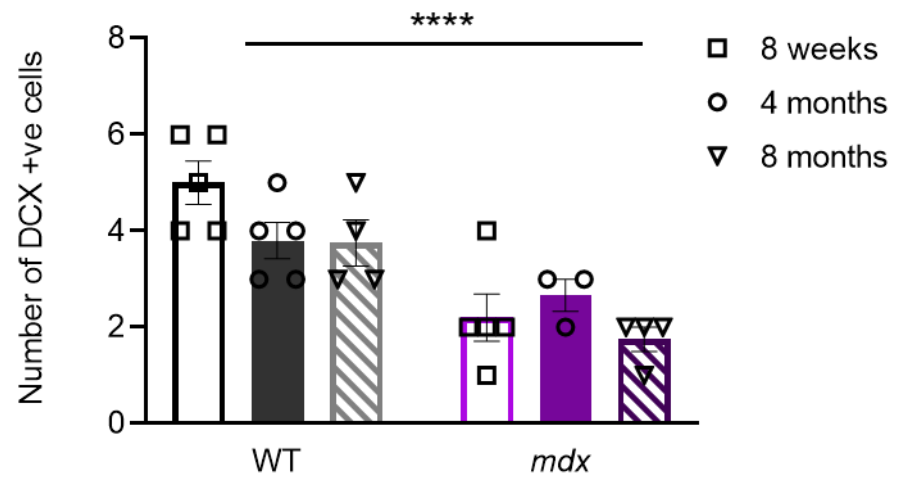


Figure 3.16: The number of DCX-positive cells in the cell layer of the dentate gyrus is decreased in mdx mice across 3 ages.

The number of positive DCX cells decreases with age in WT mice and is significantly decreased in mdx mice with no change across age groups. Two-way ANOVA, $p < 0.001$ ****

3.10. Glial filamentary acidic protein is significantly decreased in the *mdx* mouse.

Glial filamentary acidic protein (GFAP) is an intermediate filament protein which is expressed primarily by astrocytes in the CNS (Eng *et al.*, 2000). Its main function is to maintain astrocyte structural integrity and aid in cell movement and shape change (Eng *et al.*, 2000). As such, GFAP is widely accepted as the marker for astrocytes in neurological studies (Yang & Wang, 2015). Astrocytes serve an important supportive role for neurons, providing a favourable ionic environment, modulating extracellular levels of neurotransmitters, aiding the organization of neurons (Hatten & Mason, 1990) and being actively involved in the modulation of neuronal signalling (Araque *et al.*, 2014). As astrocytes are likely to modulate neuronal signalling, they have shown to subsequently regulate glutamate transmission between neurons (Angulo *et al.*, 2004; Amiri *et al.*, 2012; Amiri *et al.*, 2013) and such are involved in long term potentiation (Henneberger *et al.*, 2010). For instance, astrocytic glutamate release has been shown to activate presynaptic NMDA receptors and promote an increase in excitatory communication between neurons (Jourdain *et al.*, 2007). As we have already demonstrated that neuronal calcium activity as well as long term potentiation is altered in *mdx* mice in comparison to WT, we wanted to determine if GFAP and thus, astrocytes, may play a role in mediating these effects.

Using a GFAP primary antibody (labelled with a AP488 secondary antibody) in the hilus and granule cell layer of the dentate gyrus, we found that the number of astrocytes significantly increased with age in WT mice, from 8-weeks to 8-months

(figure 3.17, n=3-5 animals, two-way ANOVA, $F(2, 17) = 6.844$, $p=0.0066$). We also found that although there was a significant decrease in the number of astrocytes in *mdx* mice in comparison to WT (n=3-5 animals, two-way ANOVA, $F(1, 17) = 44.35$, $p<0.0001$), there was no significant differences in the numbers of astrocytes between the 2-, 4- and 8-month-old *mdx* mice. There was an overall interaction between strain and age in the two-way ANOVA with n=3-5 animals, $F(2, 17) = 9.858$, $p=0.0014$.

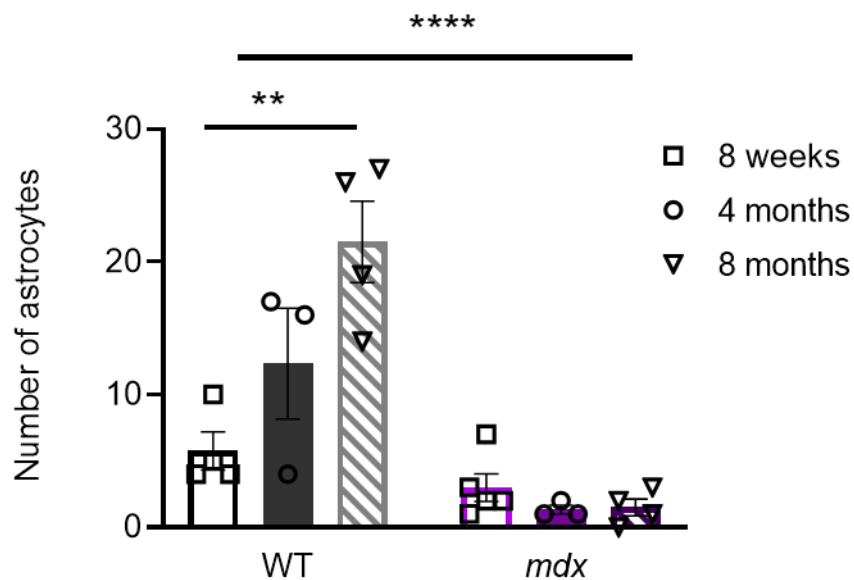


Figure 3.17: The number of astrocytes was significantly decreased in mdx mice and significantly increased in WT mice in 8-week-old, 4-month-old and 8-month-old mice.

*The number of astrocytes was significantly decreased in mdx mice relative to WT in 8-week-old, 4-month-old and 8-month-old mice. Two-way ANOVA, $p<0.01$ ** and $p<0.001$ *****

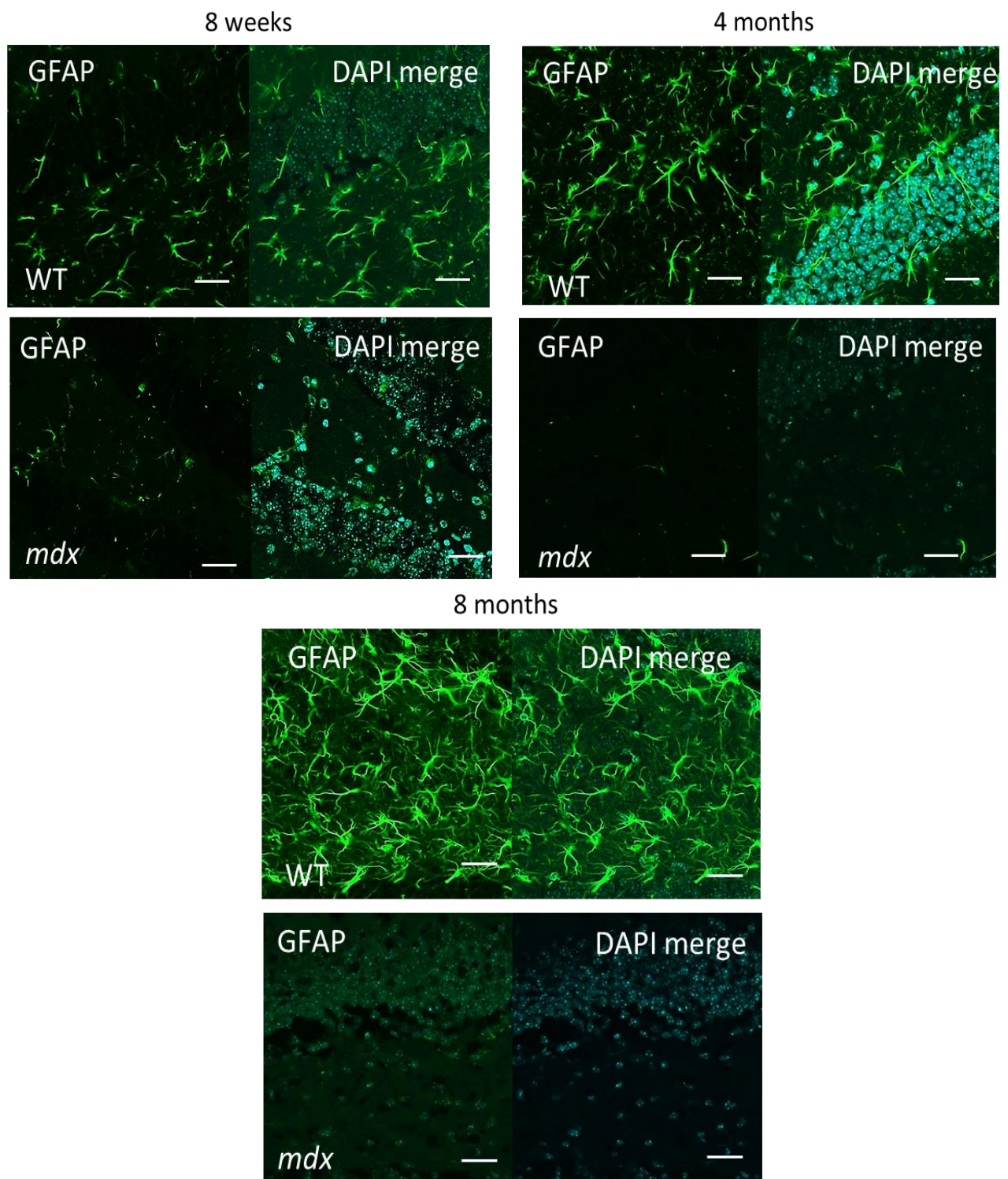


Figure 3.18: Immunofluorescence images looking at GFAP staining of astrocytes in the dentate gyrus

The number of astrocytes increases with age in WT mice compared to a significant reduction in astrocytes at all ages in mdx mice. DAPI was used as a nuclei marker for all cells. Scalebars are 30µm.

3.11. Discussion

Duchenne Muscular Dystrophy is an X-linked recessive disorder resulting in the loss of the structural protein dystrophin. Boys with DMD usually present with muscle weakness from an early age, which progresses to physical disability and eventual immobility. While most research on DMD has focused on the effect the condition has on muscle, as treatment has increased life expectancy in those affected, greater attention is now being paid to another aspect of DMD pathology: cognitive impairment. In this respect, boys with DMD have a higher incidence of mild-to-moderate cognitive impairment, with deficits in verbal, short-term and working memory (Snow *et al.*, 2013).

In our studies, LTP, proposed to be the molecular correlate of learning and memory, was decreased in *mdx* mice in comparison to WT mice, which is a novel finding as results of previous studies investigating LTP in *mdx* mice differed. In a study by Sesay and colleagues, no obvious deficit in hippocampal NMDA-dependent LTP was found in *mdx* mice (Sesay *et al.*, 1996). One explanation for the difference in our results from the study by Sesay *et al.* (1996) could be that the study was carried out on animals that were 10-12 weeks old, which is similar to our study (we used 8-12-week-old mice), but the LTP experimentation took place *in vivo* on anaesthetised animals held in a semi-stereotaxic apparatus. This methodology differs significantly to that used by us and may be one explanation for our differing results. In contrast, Vaillend *et al.* (2004), found that *ex vivo* LTP was enhanced. Although they did justify that enhanced LTP could underpin cognitive dysfunction, it is counterintuitive as increased LTP is normally associated with improved memory

consolidation. Our findings of suppressed LTP are more consistent with the cognitive deficits in learning and memory processes in *mdx* mice (Vaillend *et al.*, 2004).

Moreover, a significant decrease in LTP could impact memory in *mdx* mice leading to deficits in LTP which were associated with memory dysfunction in the spatial navigation task, the Morris water maze (Goussakov *et al.*, 2019).

Recognition memory was also investigated using the novel object recognition test, but this failed to reveal any significant differences in this study between *mdx* and WT mice. However, a recent study using NOR has shown that cognitive deficits were only observed in 4 month and older *mdx* mice, and that these cognitive deficits progressed with age (Bagdatlioglu *et al.*, 2020). As such, when we examined the *mdx* hippocampus over time, we found striking changes in the cellular structure.

We found that the total number of neurons in the hippocampal dentate gyrus, where neural progenitor cells migrate into the granule cell layer and differentiate into neuronal or glial cells (Ming & Song, 2011), were substantially reduced in *mdx* mice across all ages in comparison to WT. However, we did not see this change in adult *mdx* mice. In parallel with the observed decrease in neuronal density in aged *mdx* mice, fluid-filled structures were evident in the cell layer. These fluid-filled structures may be displacing neurons present in the cell layer and such, are likely to impact on processes that are intricately dependent on the neuronal circuitry of the hippocampus.

Another marker that has been associated with hippocampal adult neurogenesis is DCX (Couillard-Despres *et al.*, 2005). The expression of DCX is decreased with

ageing in accordance with an overall decrease in hippocampal adult neurogenesis (Kuhn *et al.*, 1996; Brown *et al.*, 2003), and as expected from previous studies in this area, we found that the number of DCX-positive cells decreased in WT mice as they aged, whereas there was a significant decrease in the number of DCX-positive cells in *mdx* mice relative to WT, which persisted up to 8 months. Decreased DCX in *mdx* mice from an early age and a reduction in hippocampal adult neurogenesis could result in a reduction in neuronal migration and proliferation (Pramparo *et al.*, 2010) resulting in disruption to the hippocampal cellular circuitry in the hippocampus. Such a disruption could result in aberrant synaptic plasticity and thus the mechanisms surrounding learning and memory would be affected, ultimately contributing to the deficits observed in individuals with DMD.

For many years the role of astrocytes within the CNS was thought to only support neurons, it is now clear that they have several diverse functions such as being actively involved in the control of neuronal functions, communication pathways, and synaptic plasticity, as they are able to receive signals from neurons and release neuroactive substances (Kirchhoff *et al.*, 2001). Astrocytes in the CNS release and regulate several neuroactive molecules that can affect neuronal activity and modulate plasticity and LTP (Theodosis *et al.*, 2008) and, as such, are involved in synaptogenesis and in regulating communication between already formed connections (Bains & Oliet, 2007). Additionally, adult neural stem cells in the subgranular zone of the dentate gyrus are both positively and negatively regulated by contact with astrocytes (Cassé *et al.*, 2018). Any change to astrocytic function or number could alter neuronal and/or overall cognitive function. In our study, we report that the number of astrocytes expressing the GFAP marker is significantly

reduced in *mdx* mice across three age groups; 2,4 and 8 months old whereas in WT mice, there is a significant increase across the same three age groups. A decrease in astrocytes expressing GFAP, in *mdx* mice, could also be a contributory factor in the age-related changes in hippocampal structure in these mice. Thus, a loss of astrocytes and altered neurogenesis on a whole could contribute to the loss of cells in the aged hippocampus which would have a deleterious effect on brain function in these animals as normal neuronal activity and plasticity are likely to be affected by this loss.

While we were primarily focussed on hippocampally-mediated learning and memory consolidation, we also took a look at other centrally-regulated behaviours that might be affected by loss of dystrophin. When we investigated the behaviour of *mdx* mice, we found that they displayed increased freezing time in an open field compared to WT mice. Freezing behaviour is a known fear response in rodents, with studies linking this behaviour to projections from the amygdala to the brainstem (LeDoux *et al.*, 1988). In fact, an increase in freezing has previously been seen in *mdx* mice with a reduction of GABA receptors in the amygdala (Sekiguchi *et al.*, 2009). While we did not find any difference in the number of GABA_A-receptor expressing interneurons in the hippocampal dentate gyrus in either of the two ages investigated, in *mdx* and WT mice, this does not necessarily imply that receptor expression or function in these cells is normal. In *mdx* mice, dystrophin has been linked to clustering of GABA_A receptors (Knuesel *et al.*, 1999; Miranda *et al.*, 2009; Vaillend *et al.*, 2010). Disruption in GABA_A receptors has also been identified in the brain of patients of DMD (Suzuki *et al.*, 2017), and interestingly, the decline in

GABA_A receptors was more evident in older adults, suggesting that there is an age-related progression of DMD on the brain.

Our studies also revealed that under basal conditions there was an increase in intracellular calcium activity in cultured hippocampal neurons of *mdx* mice resulting in neuronal hyperexcitability. Such calcium dysregulation has also been noted in other diseases characterised by cognitive impairment. For example, in Alzheimer's disease (Toescu & Verkhratsky, 2007). Dysregulated calcium can be cytotoxic and thereby contribute to premature brain aging, neuronal dysfunction and cell death (Orrenius & Nicotera, 1994) as well as neurodegeneration (Khachaturian, 1987; Landfield, 1987). At a cellular level, calcium disruption may be key to hippocampal dysfunction in *mdx* mice however, further studies are needed to elucidate this.

3.12. Conclusions

Although in early adulthood (8-12 weeks) we did not detect behavioural deficits in memory in *mdx* mice, a striking suppression of LTP, the molecular correlate of learning and memory, was apparent. Moreover, at a cellular level, calcium dyshomeostasis was evident, resulting in spontaneous neuronal hyperexcitability, which if left unregulated can lead to cellular cytotoxicity. This could be due to defective inhibitory regulation by GABAergic neurons. Although no change in the absolute number of GABAergic interneurons was detected in *mdx* hippocampal tissue, GABA_A receptor post-synaptic clustering is dependent upon the presence of neuronal dystrophin (Vaillend *et al.*, 2010) thus function in these neurons could be impacted. In longitudinal studies, we also noted a loss of neurons and astrocytes

and the pathophysiological changes in the cellular structure of the hippocampus in aged *mdx* mice. Impaired hippocampal adult neurogenesis is likely to impact acutely on the formation of new memories, but may also contribute to the long-term loss of neuronal cells. The density of astrocytes, which are critical to normal neuronal function, was suppressed to an even greater extent across all age-points in *mdx* mice. Several of the cellular changes are evident even in early adulthood when the behavioural expression of these changes is not yet evident. However, compensatory mechanisms may exist to preserve these critical cognitive functions.

This research has revealed an array of structural and functional changes in the hippocampi of dystrophin-deficient *mdx* mice, which begins to elucidate the role of dystrophin in non-contractile tissue, such as hippocampal neurons.

Chapter 4

**Interleukin-6: its potential as a
neuromodulatory factor
underpinning aberrant
hippocampal activity in
dystrophin-deficient *mdx* mice.**

4.1. Introduction

IL-6 is a pleiotropic cytokine which plays a key role in the immune system and also in CNS physiology and pathophysiology.

IL-6 signalling occurs through two mechanisms, classical and trans-signalling.

Classical IL-6 signalling occurs through the binding of IL-6 to its membrane bound receptor, IL-6R α . Whereas trans-signalling, which is often occurs when IL-6 binds to the soluble form of the receptor. This results in homodimerization of a glycoprotein, gp130, resulting in phosphorylation of JAK/STAT (Darnell *et al.*, 1994) and further downstream activation of MAPK and PI 3-K cascades (Eulenfeld *et al.*, 2012).

In the CNS, neuronal degeneration induced by IL-6 is mediated by trans-signalling. IL-6 can also be produced locally within the CNS, mainly by astrocytes (Dong & Benveniste, 2001; Farina *et al.*, 2007) and microglia (Ye & Johnson, 1999) but may also cross the blood brain barrier *via* saturable transport systems (Banks *et al.*, 1994). Thus, acute increases in circulating levels of this pro-inflammatory cytokine as are seen in DMD patients (Rufo *et al.*, 2011) and *mdx* mice (Pelosi *et al.*, 2015), could also raise levels within the CNS.

The effects of IL-6 on the brain have been studied with regards to cognitive function. In humans, deterioration of cognitive function, a hallmark of 'sickness behaviour', is linked to high circulating levels of pro-inflammatory cytokines such as IL-1 β (Gibertini, 1998) and IL-6, which has also been implicated in lipopolysaccharide (LPS) -evoked cognitive dysfunction (Sparkman *et al.*, 2006).

Moreover, transgenic mice over-expressing IL-6 exhibit broad memory impairments (Heyser *et al.*, 1997; Wei *et al.*, 2012; Wei *et al.*, 2013) and severe neurological pathologies such as tremor, ataxia, and seizure, with neurodegeneration, astrogliosis and angiogenesis also being reported (Campbell *et al.*, 1993). The explicit effect of this cytokine on learning and memory function is further supported by transgenic mice deficient in IL-6, which possess enhanced spatial learning capabilities in the radial arm maze (Braidá *et al.*, 2004) and the Morris water maze (Bialuk & Winnicka, 2018) behavioural task-learning assessments.

IL-6 may also be involved directly in synaptic plasticity and the hippocampus. Application of IL-6 was shown to inhibit hippocampal LTP using a mechanism that was dependent upon IL-6R activation (Tancredi *et al.*, 2000). Moreover, IL-6-induced LTP inhibition was attenuated by administration of monoclonal antibodies, used to block IL-6 receptors (Li *et al.*, 1997).

Anatomical and biochemical abnormalities, such as a loss of neurons, the appearance of neurofibrillary tangles, alterations in dendritic branching, cortical atrophy and metabolic defects (Rosman, 1970; Jagadha & Becker, 1988; Tracey *et al.*, 1995; Lv *et al.*, 2011) are observed in both DMD patients and *mdx* mice (Miranda *et al.*, 2009). This is paired with an increased incidence of mild to moderate cognitive impairment (Bresolin *et al.*, 1994; Anderson *et al.*, 2002; Cotton *et al.*, 2005) and deficits in verbal, short-term and working memory (Snow *et al.*, 2013) in DMD patients. Cognitive deficits are also present in the *mdx* mouse. In the *mdx* mouse, there is evidence of cognitive dysfunction such as a poor capacity to learn new tasks as well as deficits in storing spatial memories, relative to controls

(Vaillend *et al.*, 1995; Vaillend *et al.*, 2004; Chaussonot *et al.*, 2015). They also display deficits in associative learning and in the general processes involved in memory formation and consolidation which are both dependent upon optimal functioning of the hippocampus and amygdala (Chaussonot *et al.*, 2015).

Given that loss of dystrophin from skeletal muscle is associated with chronic inflammation, which is characterised by elevated circulating levels of *IL-6 in humans* (Rufo *et al.*, 2011) and the *mdx* mouse (Pelosi *et al.*, 2015), IL-6 may act as an immune mediator leading to cognitive dysfunction in the absence of dystrophin.

4.1.1. Study objectives

- To determine if IL-6, which is elevated in DMD and *mdx* plasma, is increased in dystrophic hippocampal tissue.
- To investigate if IL-6 contributes to aberrant hippocampal activity in *mdx* mice.

4.1.2. Techniques used

- Neuronal cell culture (see methods section 2.2.1, pg. 33).
- Immunocytochemistry (see methods section 2.3.1, pg. 39)
- Gene expression - qRT-PCR (see methods section 2.5, pg. 47).
- Immunoassay - ELISA (see methods section 2.6, pg.52).
- Calcium imaging (see methods section 2.4, pg. 44).
- Immunofluorescence of whole hippocampal sections (see methods section 2.3.2, pg. 40).
- Extracellular electrophysiology (see methods section 2.8, pg.59).

4.2. Dystrophin isoform is co-localised at synaptophysin-labelled hippocampal synapses.

As dystrophin expression has been reported in the postsynaptic density (PSD) of cortical and cerebellar neurons (Lidov *et al.*, 1990), we examined immunofluorescent expression of dystrophin in hippocampal neurons (10 to 12 days in culture) cultured from dystrophin-expressing C57BL/6 mice (n= 15 fields of view from 3 different cultures). Immunofluorescent staining to the monoclonal mouse MANDRA-1 antibody which identifies full length dystrophin, Dp427 as well as dystrophin variants including Dp71, the most prevalent form of dystrophin in the CNS (Hugnot *et al.*, 1992; Lederfein *et al.*, 1992; Jung *et al.*, 1993) was used. Our results demonstrate that both forms were present in both cultured neurons and non-neuronal cells. In the neuronal cell body, diffuse dystrophin staining was evident in the cytosol, whereas immunolabelling in the processes was clustered into distinct puncta. Co-labelling with a rabbit monoclonal antibody against synaptophysin, an integral membrane protein localised to synaptic terminals, further indicated localisation of dystrophin at hippocampal synapses. This pattern of staining was consistent across several cultures and all neurons imaged (Figure 4.1).

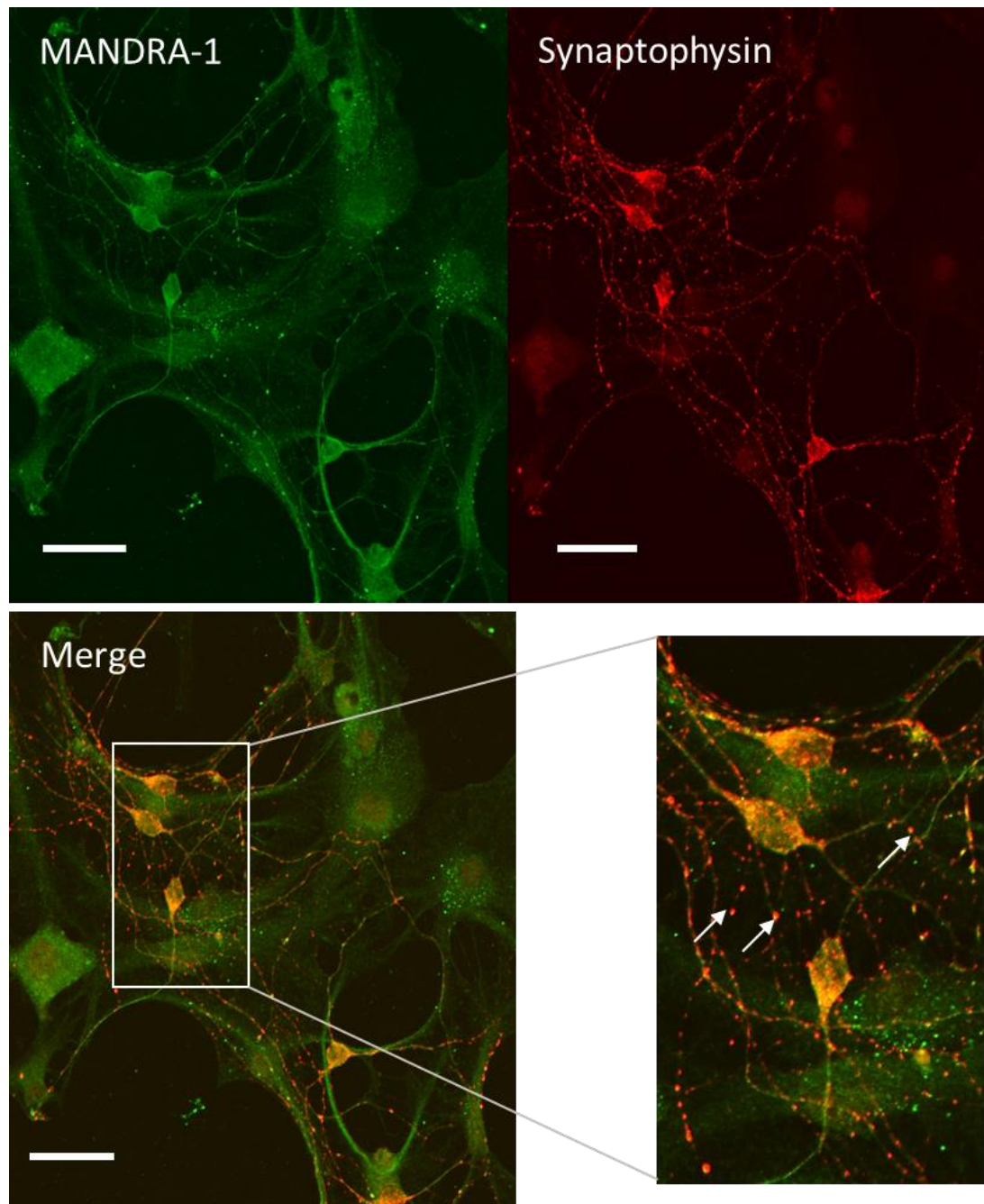


Figure 4.1: Dystrophin is expressed at hippocampal synapses.

Representative images of cultured hippocampal neurons co-stained with the MANDRA-1 antibody for several isoforms of dystrophin (green stain), and synaptophysin (red stain), a specific marker of synapses, indicate that dystrophin is expressed on synapses on neuronal cell bodies and neuronal processes (indicated by white arrows). Scalebar: 35 μ m.

4.3. IL-6Rs are clustered at synaptophysin-labelled hippocampal synapses.

Hippocampal neurons cultured from dystrophin-expressing C57BL/6 mice, labelled with a monoclonal mouse antibody against IL-6R α (n= 15 fields of view from 3 different cultures), showed a similar pattern of staining to that of dystrophin. Specifically, cytosolic IL-6R was diffusely expressed in neurons, with punctate patterns of expression in dendritic processes (figure 4.2), and IL-6R expression was shown to cluster and co-localise with synaptophysin.

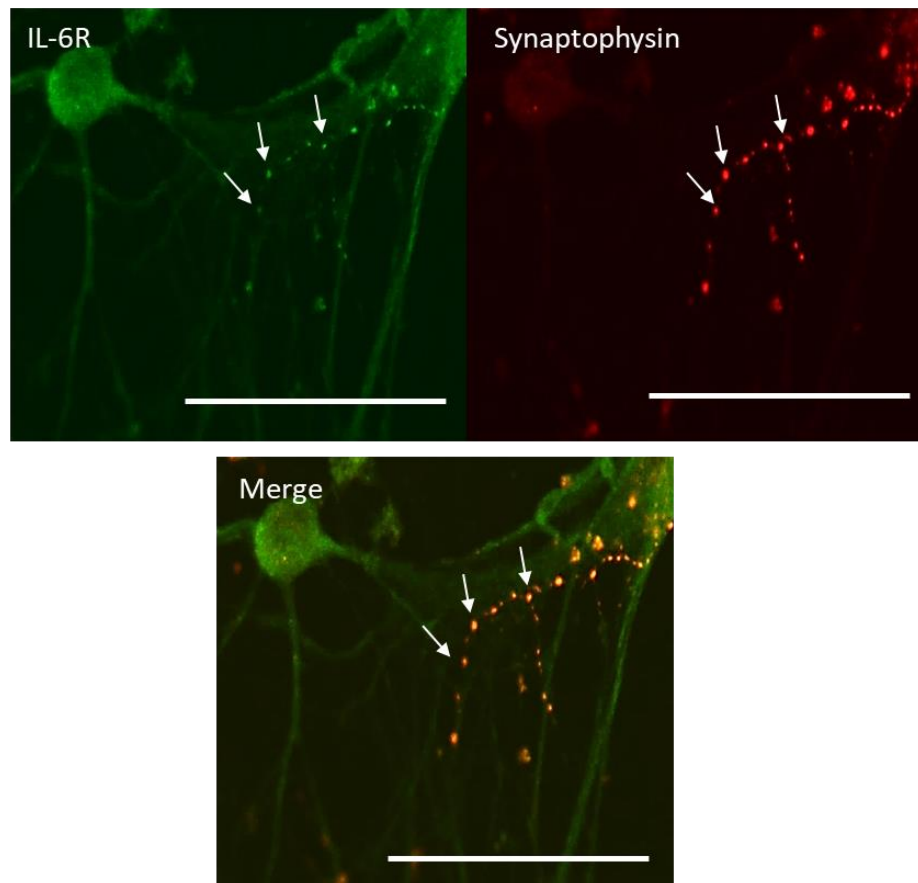


Figure 4.2: IL-6 receptors are expressed at hippocampal synapses.

Representative images of cultured hippocampal neurons illustrate that IL-6R (green staining) co-localises (yellow staining in merged image) with the pre-synaptic marker, synaptophysin (red staining, illustrated with white arrows) in cultured hippocampal neurons. Scalebar: 35 μ m.

4.4. IL-6 exposure increases IL-6R clustering in dystrophin containing hippocampal neuronal processes.

Given that neuronally-expressed dystrophin isoforms and IL-6Rs are both located in hippocampal synapses, we next sought to investigate possible if IL-6 stimulated synaptogenesis in control dystrophin-expressing hippocampal neurons. Neurons grown in the presence of recombinant IL-6 (1nM) were compared to control neurons without exogenous IL-6. The IL-6R-labelled immunofluorescent images generated from these neurons indicated that exposure to IL-6 induced an increase in the density of IL-6R clustering in the neuronal processes compared to untreated neurons (n=19 neurons/3 different cultures, 20mm of process from cell body, $p=0.0137$, figure 4.3).

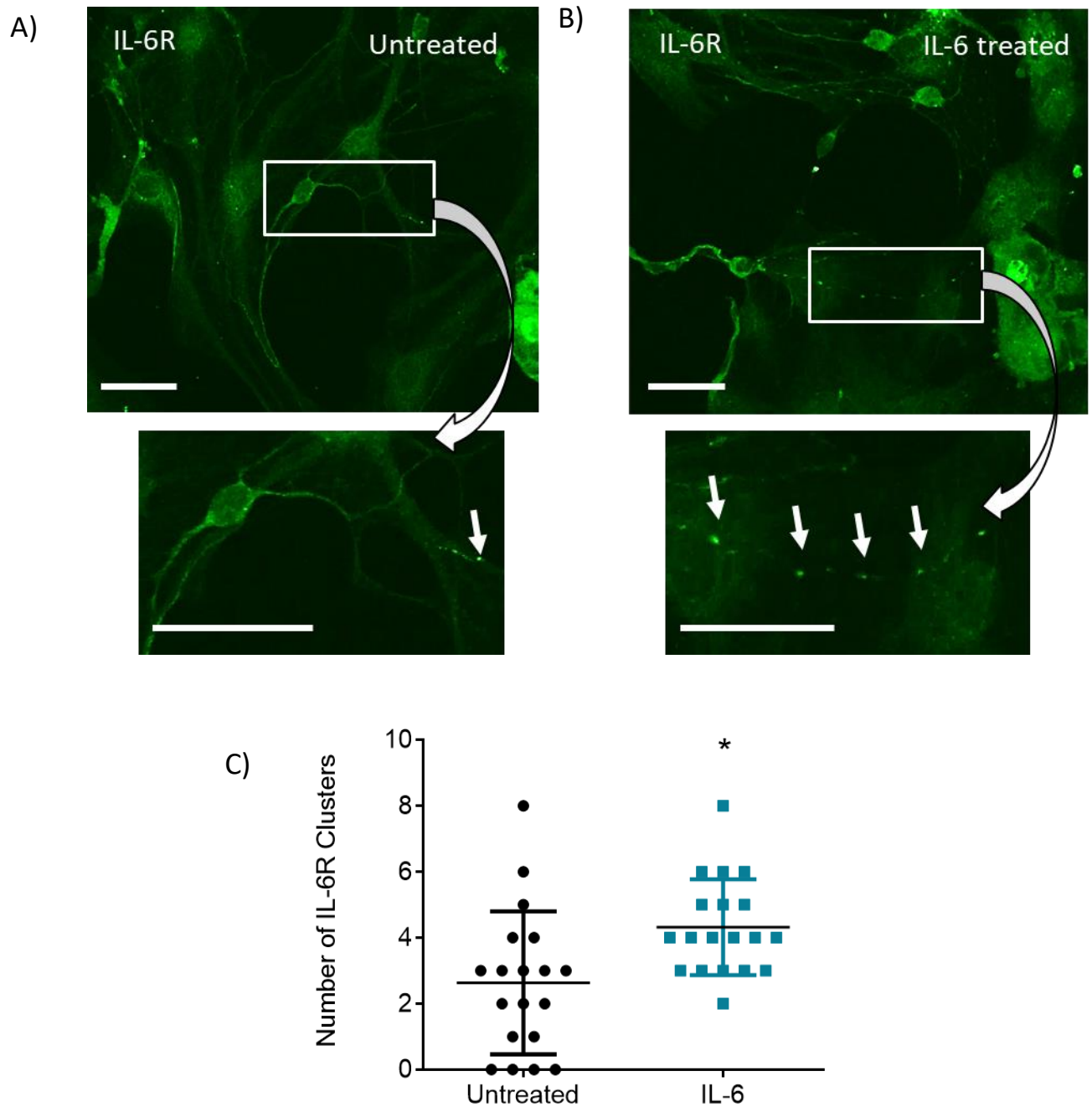


Figure 4.3: Exposure to IL-6 stimulates increased IL-6-positive puncta on hippocampal processes.

C) The dot plot illustrates the number of IL-6R clusters per 20mm of process in dystrophin-containing C57BL/6 mice in untreated neuronal cell cultures and in neurons treated with 1nM recombinant IL-6 in culture media over 10-12 days, $n=19$ neurons/3 cultures. This can be seen in the representative images A) and B). The arrows indicate IL-6R clusters along the process. Scalebar: 35 μ m. Unpaired t test, $p<0.05^*$

4.4. Hippocampal interleukin-6 (IL-6) expression is comparable between WT and *mdx* mice.

Given the elevations in circulating IL-6 in *mdx* mice (Pelosi *et al.*, 2015) we sought to determine if IL-6 was elevated in the hippocampus of *mdx* mice. qRT-PCR (where quantification of amplified DNA is achieved using fluorescence measurements) was conducted to determine the gene levels of IL-6 in the whole hippocampus of 8–12-week-old WT (n=10) and *mdx* mice (n=10). The hippocampal gene expression of IL-6 was not significantly different between WT (n=10, 2 ± 2.662) and *mdx* (n=10, 2.367 ± 3.156) mice ($p > 0.9999$, figure 4.3). However, as gene expression does not always correlate with protein expression (Liu *et al.*, 2016), we also compared hippocampal IL-6 protein levels between WT and *mdx* mice. Using an ELISA assay, we found that IL-6 protein in hippocampal homogenates was similar between WT (n=10, 5.059 ± 0.8282) and *mdx* (n=10, 4.977 ± 0.6414) mice ($p = 0.8070$, figure 4.4).

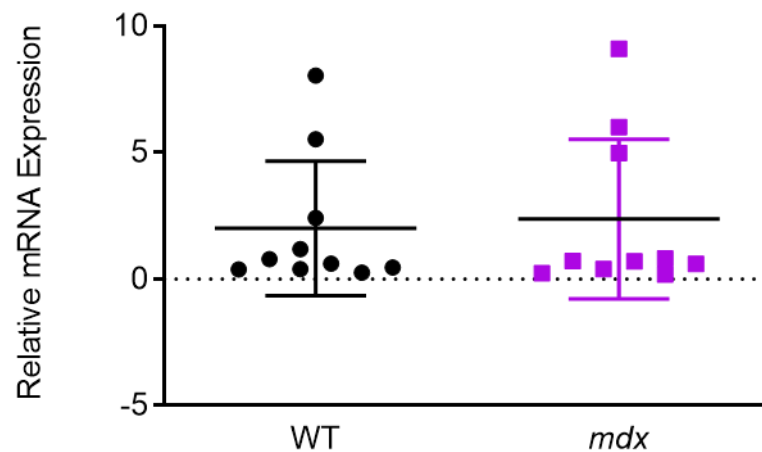


Figure 4.3: Hippocampal IL-6 gene expression is not different in *mdx* mice.

The data plot indicates that no difference in the relative mRNA expression of IL-6 was detected between *mdx* and WT mice (n=10, $p > 0.9999$). Unpaired t test, $p > 0.05$

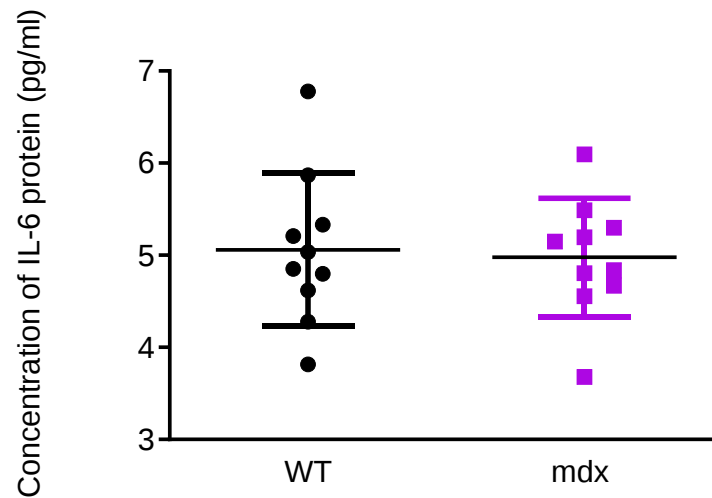


Figure 4.4: Hippocampal protein expression of IL-6 did not differ significantly between WT and mdx mice.

The dot plot illustrates the concentration of IL-6 protein (pg/ml) in WT and mdx hippocampal homogenates (n=10, p=0.8070). Unpaired t test, p>0.05

4.5. Hippocampal IL-6 receptor expression and intracellular signalling is not altered in the absence of dystrophin.

Although IL-6 at the gene and protein level was not significantly altered in *mdx* hippocampal tissue compared to WT controls, the sensitivity to this cytokine could be altered by changes in receptor expression or modification of downstream signalling pathways. We therefore examined gene expression of IL-6 receptors (IL-6Rs). Using RT-qPCR we found that IL-6R gene expression was similar in WT (n=10, 1.078 ± 0.139614) and *mdx* (n=10, 1.077 ± 0.1386) mice ($p=0.9990$, figure 4.5).

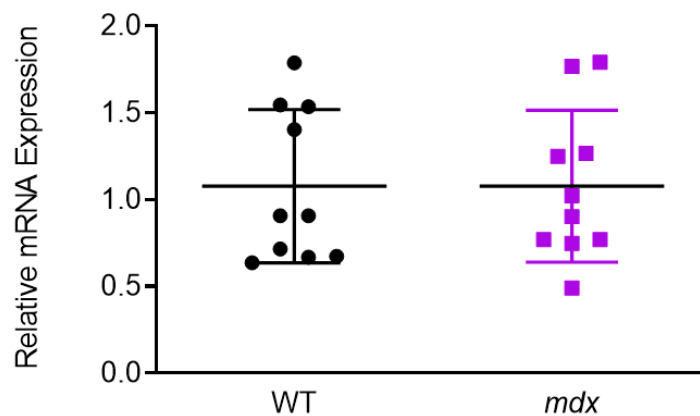


Figure 4.5: Hippocampal IL-6R gene expression is not different in mdx mice.

The data plot indicates that no difference in the relative mRNA expression of IL-6R was detected between mdx and WT mice ($n=10$, $p=0.9990$). Unpaired t test, $p>0.05$

As the IL-6/IL-6R complex subsequently binds to gp130, a membrane-bound signal transducer for IL-6, which leads to the downstream activation of the JAK/STAT pathway, and downstream cellular effects, we also measured gp130 expression in WT and *mdx* mice but, once again, we did not detect any significant differences between WT ($n=10$, 1.155 ± 0.6439) and *mdx* ($n=10$, 1.036 ± 0.3121) mice ($p=0.6048$, figure 4.6).

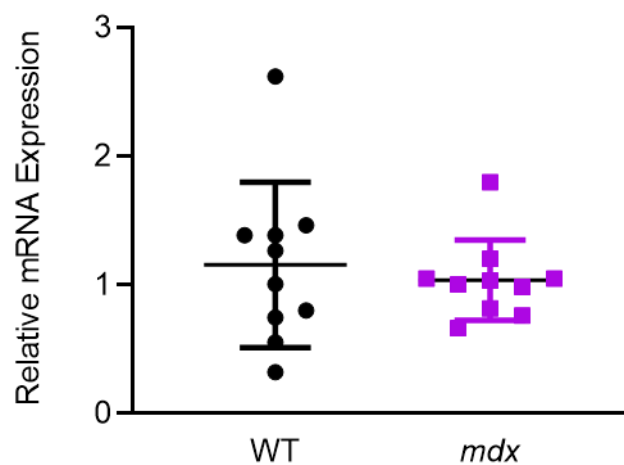


Figure 4.6: Hippocampal gp130 gene expression does not differ significantly in mdx mice.

The box plot illustrates that there is no difference in the relative mRNA expression of gp130 between mdx and WT mice ($n=10$, $p=0.6048$). Unpaired t test, $p>0.05$

Next, we examined the expression of STAT3, which is the main downstream target of IL-6 receptor activation. Its downstream actions include gene activation, cell growth and division, cell movement and apoptosis. At the gene level, in unstimulated tissue, no differences were detected in STAT3 gene expression between WT ($n=10$, 1.271 ± 0.2976) and *mdx* ($n=10$, 1.115 ± 0.1623 , $p=0.6502$, figure 4.7) either.

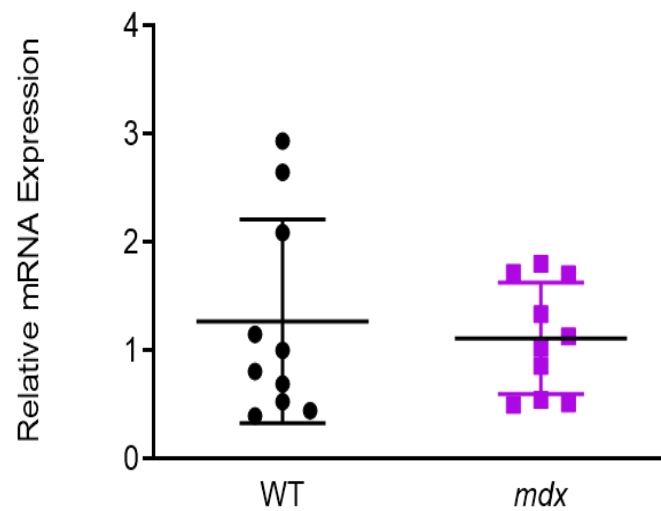


Figure 4.7: Hippocampal STAT3 gene expression is not different in mdx mice.

There is no significant difference in the relative mRNA expression of IL-6R between mdx and WT mice ($n=10$, $p=0.6502$). Unpaired t test, $p>0.05$

4.6. Regional differences in hippocampal IL-6 expression were detected in *mdx* mice.

Although useful for quantifying IL-6 expression, both qRT-PCR and ELISA use homogenised hippocampal tissue, therefore detection of region-specific changes in IL-6 or IL-6R was not possible with these techniques. However, immunofluorescence is a useful technique for identifying specific regions of protein expression. Using hippocampal slices, where the specific cell body layer remains intact, IL-6 expression was visualised in CA1, CA3 and the dentate gyrus (DG) regions using a monoclonal mouse antibody against IL-6. IL-6 expression was determined by analysing the mean fluorescent intensity in the pyramidal cell layer of the CA1 and CA3 regions and the granule cell layer in the DG. Confocal images were taken under identical conditions for laser and saturation between WT and *mdx* for IL-6 for accurate later comparison in each of the hippocampal regions.

The expression of IL-6 was increased in the CA1 region of *mdx* mice (% increase) compared to WT (*n*=20 confocal images in this region from 5 WT and *mdx* animals, WT 1.610 ± 0.9609 vs *mdx* 2.601 ± 1.741 , *p* = 0.04). Further, there was also a significant increase in IL-6 expression observed in the DG region of *mdx* mice (36% increase) in comparison to WT (*n* = 17 confocal images in this region from 5 WT and *mdx* animals, WT 0.8728 ± 0.3478 vs *mdx* 2.268 ± 1.559 , *p* = 0.0058). However, there was no difference detected between IL-6 expression in the CA3 region in *mdx* and WT mice (*n*=20 confocal images in this region from 5 WT and *mdx* animals, WT 1.346 ± 0.8657 vs *mdx* 1.214 ± 1.146 , *p*=0.3506). This data can be seen in the data plot in figure 4.8 below.

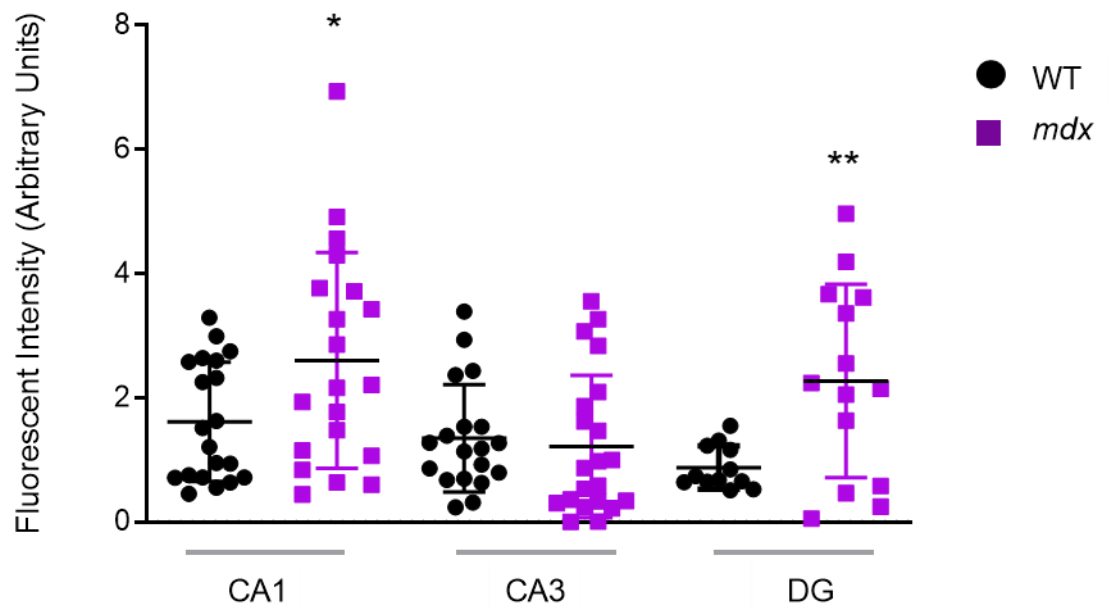


Figure 4.8: Regional hippocampal immunofluorescent expression of IL-6 is increased in dystrophin-deficient mdx mice.

The data plot illustrates a significant increase in IL-6 fluorescence in the CA1 and DG regions in mdx mice in comparison to WT mice and DG: WT 0.8728 ± 0.3478 vs mdx 2.268 ± 1.559 , $p=0.0058$). There is no significant difference in IL-6 fluorescence between CA3 regions in mdx and WT mice (CA3: WT 1.346 ± 0.8657 vs mdx 1.214 ± 1.146 , $p=0.3506$). Scale bars are $70\mu\text{m}$. Unpaired t test, $p<0.05^*$ and $p<0.01^{**}$

In the pyramidal cell layer of CA1 and CA3 and granule cell layer of DG, IL-6 expression is diffuse in the cytosol and localised to the cell membrane of both WT and mdx mice. Although, in CA1 and DG regions in mdx mice, IL-6 expression in the cell membrane of these cell layers is increased in comparison to WT which can be seen in the representative images in figure 4.9.

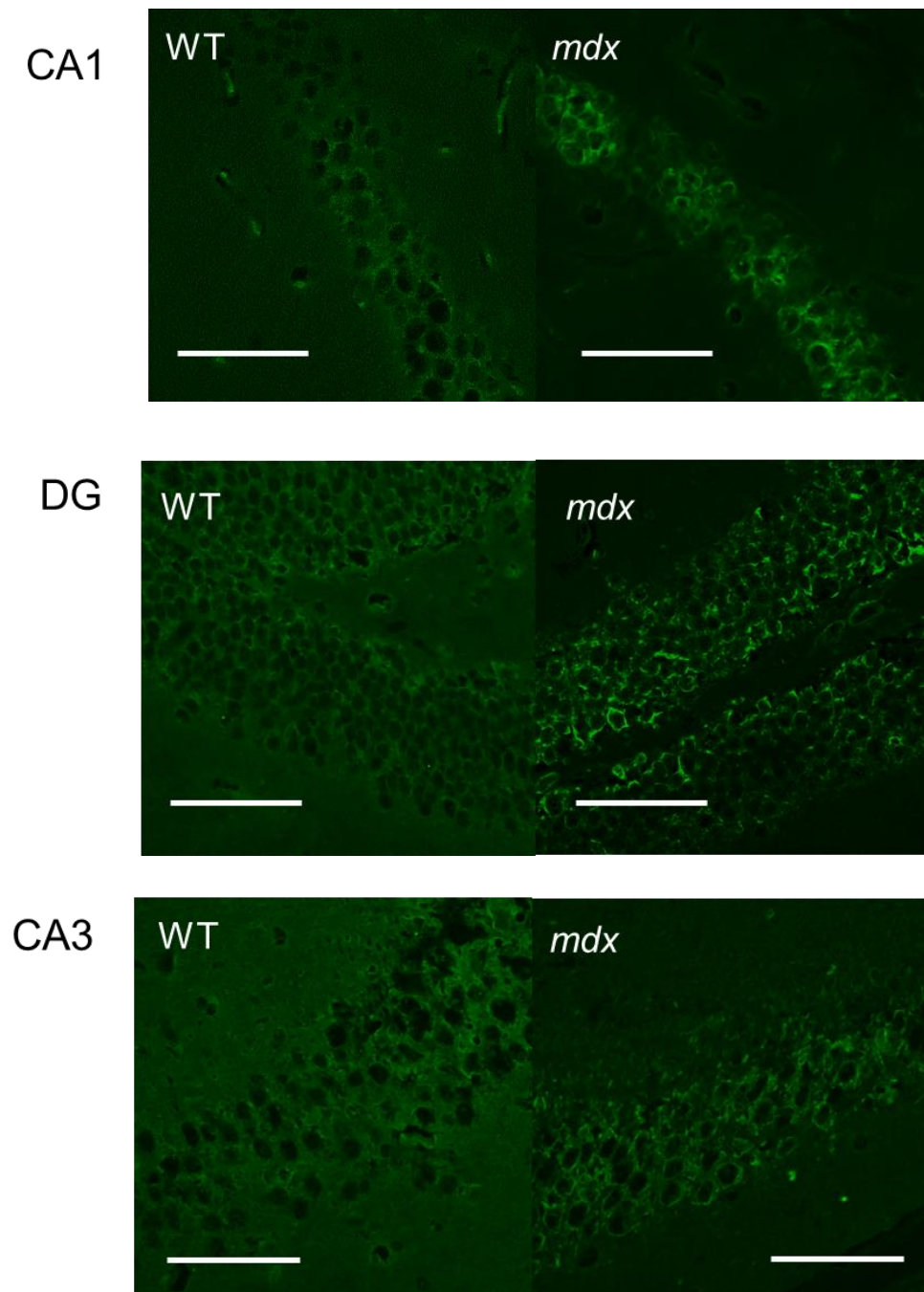


Figure 4.9: Representative images of regional hippocampal immunofluorescent illustrate that the expression of IL-6 is increased in dystrophin-deficient mdx mice.

In the pyramidal cell layer of CA1 and CA3 and granule cell layer of DG of both WT and mdx mice, IL-6 expression is diffuse in the cytosol, with IL-6 localised in the cell membrane. In mdx mice, this expression of IL-6 (fluorescence) in the cell membrane is increased in CA1 and DG in comparison to WT.

4.7. Regional differences in hippocampal IL-6 receptor expression were detected in *mdx* mice.

As IL-6 expression was shown to be increased in CA1 and DG, in the previous section, we next used a monoclonal mouse antibody against IL-6R. Similarly, IL-6R was determined by analysing the mean fluorescent intensity in the pyramidal cell layer of the CA1 and CA3 regions and the granule cell layer in the DG. With confocal images were taken under identical conditions for laser and saturation between WT and *mdx* for IL-6R for accurate later comparison in each of the hippocampal regions.

When immunofluorescence intensity was quantified, a significant increase in relative IL-6R protein expression was detected in both CA1 (63% increase, $n=17$ confocal images in this region from 5 WT and *mdx* animals, WT 7.072 ± 3.764 vs *mdx* 11.61 ± 4.543 , $p=0.0126$) and DG (52% increase, $n=12$ confocal images in this region from 5 WT and *mdx* animals, WT 2 ± 0.9930 vs *mdx* 3.8 ± 2.347 , $p=0.035$) regions in *mdx* mice compared to WT (Figure 4.10). Lastly, and differing from IL-6, IL-6R expression in the CA3 region was significantly decreased in *mdx* and WT mice ($n=12$ confocal images in this region from 5 WT and *mdx* animals (56% decrease, WT 6.996 ± 3.355 vs *mdx* 3.963 ± 2.206 , $p=0.0242$). This data can be seen in the data plot in figure 4.10 below.

Similarly, to IL-6, in the pyramidal cell layer of CA1 and CA3 and granule cell layer of DG, IL-6R expression is diffuse in the cytosol and localised to the cell membrane of both WT and *mdx* mice. Although, in CA1 and DG regions in *mdx* mice, IL-6R expression in the cell membrane is increased and for IL-6R, CA3 expression in the

cell membrane is decreased in these cell layers in comparison to WT which can be seen in the representative images in figure 4.11.

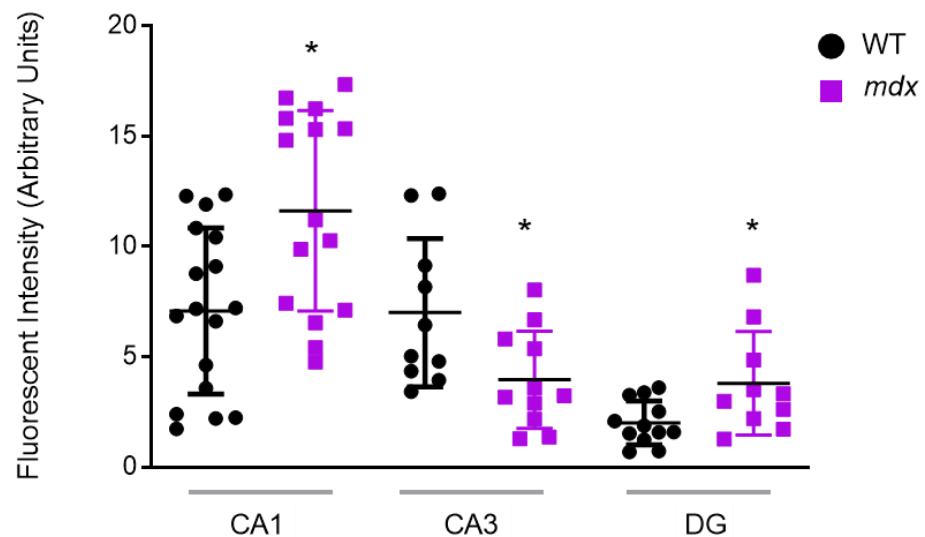


Figure 4.10: Regional hippocampal immunofluorescent expression of IL-6R is increased in dystrophin-deficient mdx mice.

The data plot illustrates a significant increase in IL-6R fluorescence in the CA1 and DG regions in mdx mice in comparison to WT mice (CA1: WT 7.072 ± 3.764 vs mdx 11.61 ± 4.543 , $p = 0.0126$ and DG: WT 2 ± 0.9930 vs mdx 3.8 ± 2.347 , $p = 0.035$). There is a significant decrease in IL-6R fluorescence between CA3 regions in mdx and WT mice (CA3: WT 6.996 ± 3.355 vs mdx 3.963 ± 2.206 , $p = 0.0242$). Scale bars are $70 \mu\text{m}$. Unpaired t test, $p < 0.05^$.*

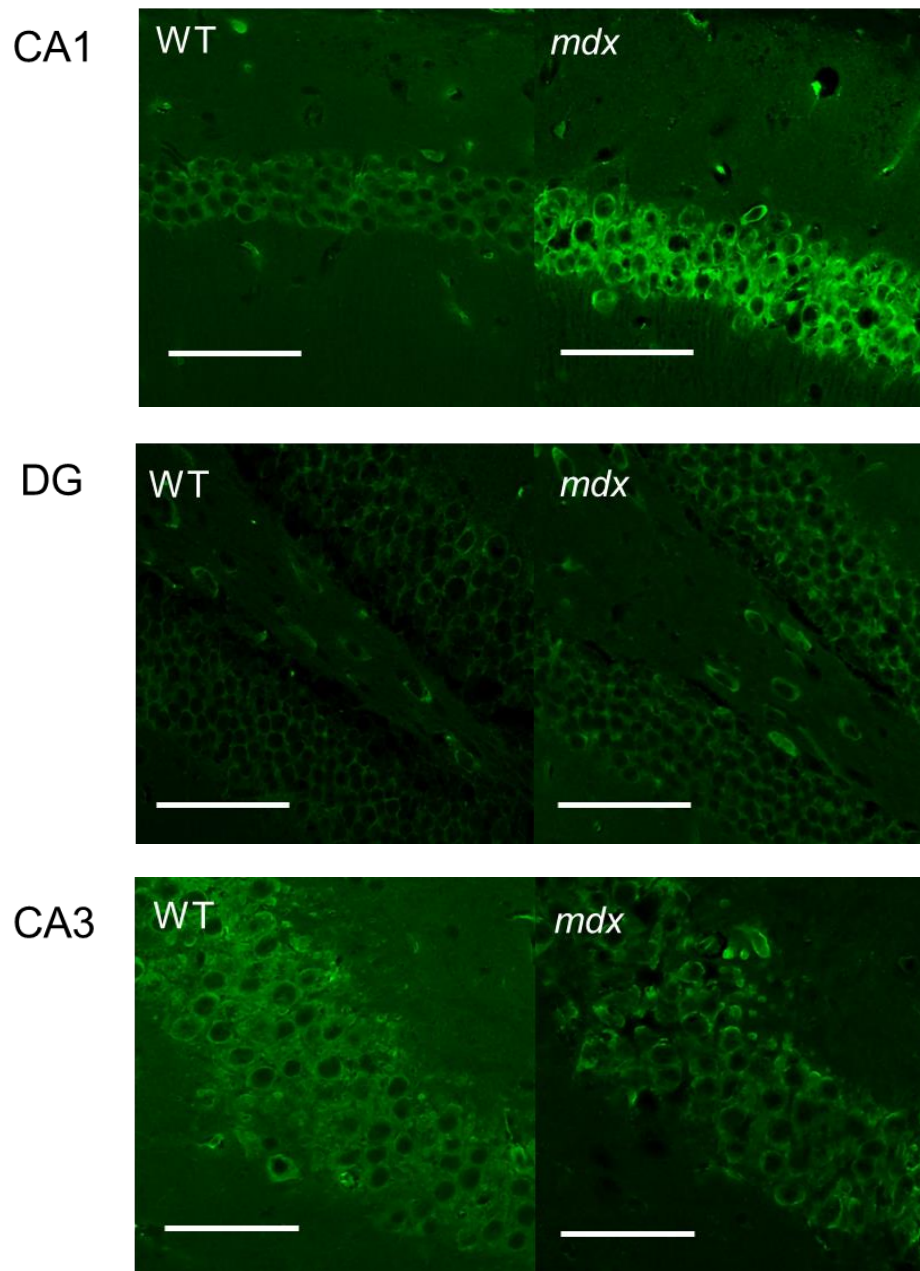


Figure 4.11: Representative images of regional hippocampal immunofluorescence illustrate that the expression of IL-6R is increased in dystrophin-deficient mdx mice.

In the pyramidal cell layer of CA1 and CA3 and granule cell layer of DG of both WT and mdx mice, IL-6R expression is diffuse in the cytosol, with IL-6R localised in the cell membrane. In mdx mice, this expression of IL-6R (fluorescence) in the cell membrane is increased in CA1 and DG in comparison to WT.

4.8. IL-6 suppresses spontaneous calcium activity in dystrophin-deficient hippocampal neurons.

** Experiments completed by Kimberley Stephenson and Michael Vaughan and data analysed by Claire Denley.*

As presented in chapter 3.4, under basal conditions spontaneous intracellular activity was increased in cultured *mdx* neurons in comparison to WT mice. To determine if the neuromodulatory cytokine, IL-6, modified this activity, the cultured neurons were exposed to exogenous recombinant IL-6 (1nM).

The pooled data and representative traces show that the elevated frequency of calcium peaks was unaffected by IL-6 in dystrophic *mdx* neurons (n= 29 neurons from 3 cultures and IL-6 did not modify activity in WT neurons (n=31 neurons from 3 cultures) either ($F(1,1027) = 0.25$, $p = 0.62$, two-way ANOVA). However, the strain difference was highly significant ($F(1, 1027) = 86.5$, $p < 0.0001$) and an interaction between treatment and strain was detected ($F(1, 1027) = 6.7$, $p = 0.01$).

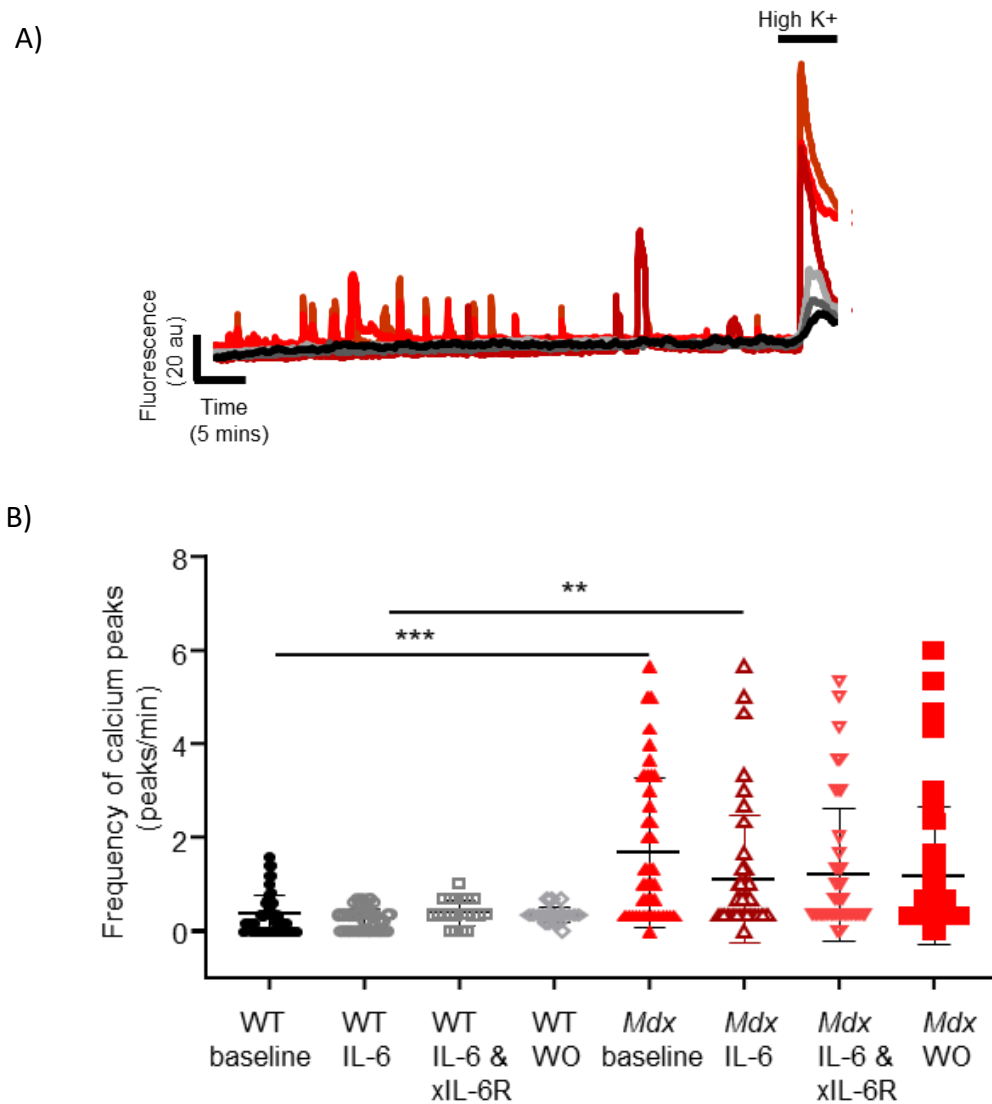


Figure 4.12: Spontaneous calcium activity in mdx hippocampal neurons is not changed by exogenous IL-6.

*The representative traces in A) show the effect of IL-6 WT neurons (grey) and mdx neurons (red) and B) The data plot illustrates the frequency of calcium peaks in WT and mdx mice following addition of IL-6 (1nM, 5 mins). Two-way ANOVA, $p < 0.01$ ****

4.9. Exposure to IL-6 modifies LTP in WT but not *mdx* hippocampal slices.

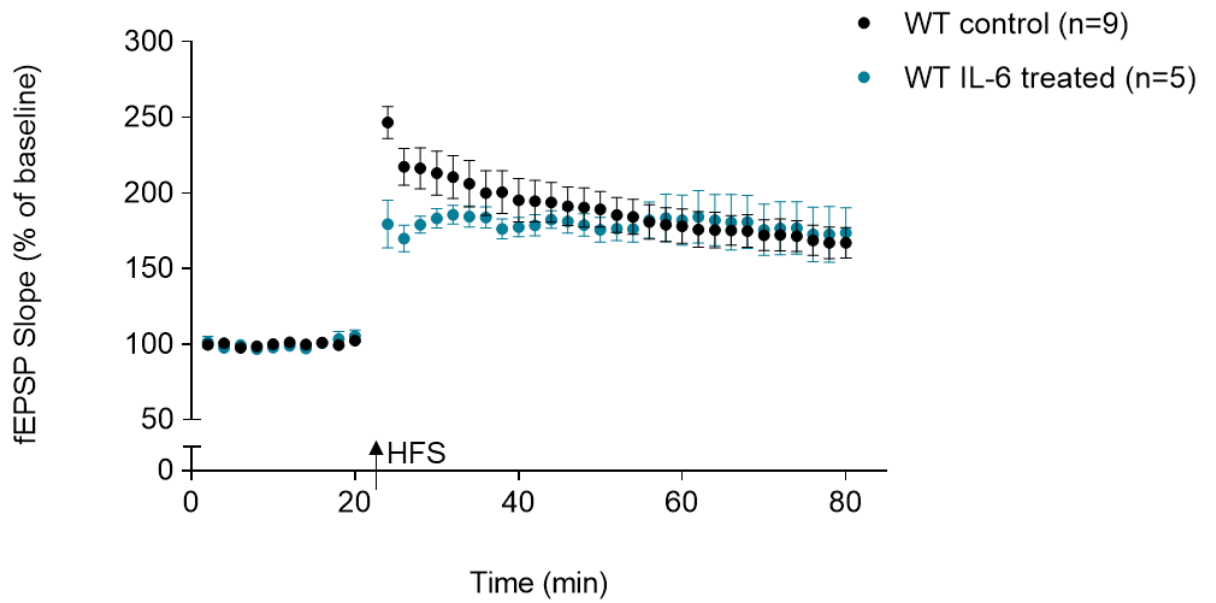
While the previous experiments demonstrated that IL-6 has an impact on neuronal excitability in individual neurons, we next used extracellular electrophysiology to determine if IL-6 modified signalling pathways underpinning the formation and consolidation of new memories. Additionally, as detailed in the previous chapter (section 3.6), LTP could be generated in the Schaffer collaterals which span the CA3 to CA1 regions by applying high frequency stimulation (HFS – 4 x 100 pulses, 100Hz for 1s/60s) resulting in a long-lasting increase in the fEPSP slope, we found in control experiments, hippocampal LTP in *mdx* mice was reduced in comparison to LTP in WT mice (Chapter 3, section 3.3, pg. 77) And as IL-6 has shown to disrupt LTP with addition of exogenous IL-6 (Tancredi *et al.*, 2000), we wanted to investigate whether IL-6 has the same effect in *mdx* mice.

Hippocampal slices were perfused with aCSF containing recombinant IL-6 (1nM) for 1-hour prior to recording, as well as throughout the LTP protocol. LTP was induced as described in the previous chapter (as well as methods section 2.8, pg.58).

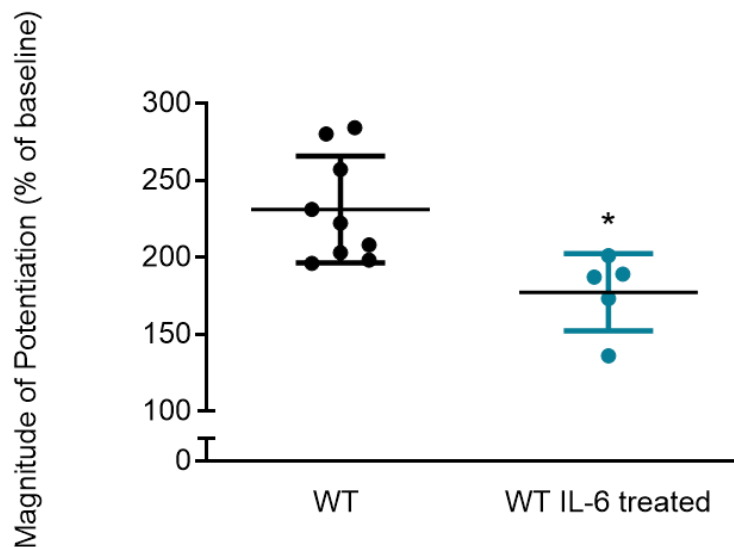
As in previous experiments, fEPSP baseline recordings were conducted for 20 minutes, with stimulation occurring every 30 seconds, using a current that induced fEPSPs that were 50% of the maximal size that could be generated. This recording period was followed by the application of four trains of high-frequency stimuli (HFS – 4 x 100 pulses, 100Hz for 1s/60s), with a current that induced fEPSPs that were 30% of the maximal size that could be generated.

In these experiments, immediately following the HFS, we saw the magnitude of potentiation of the fEPSP slope, was significantly decreased in WT slices perfused with exogenous IL-6 in comparison to WT controls (WT $231 \pm 35\%$ vs WT + IL-6 $177 \pm 25\%$; $p = 0.01$, unpaired t test, figure 4.13b). Importantly, this was only different in the early phase of LTP. Early LTP depends primarily on short-term kinase activity and is important in increasing synapse sensitivity but not *de novo* protein synthesis. Further, IL-6 seems to have stabilised fEPSP slopes over time in WT slices perfused with exogenous IL-6 in comparison to WT controls, which have a natural decay (figure 4.13a).

A



B



C

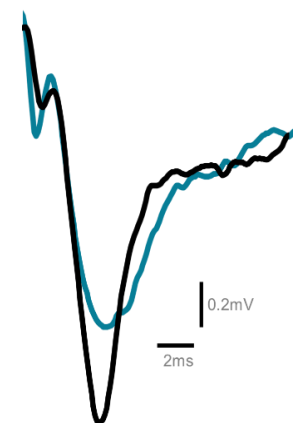


Figure 4.13: Early LTP in WT hippocampal Schaeffer collaterals-CA1 synapses was suppressed by IL-6.

- A) Graph showing averaged time course of Schaffer collateral-evoked synaptic responses from WT controls (black circles) and WT slices perfused with IL-6 (blue circles) in the hippocampal region using extracellular field potential recordings in CA1 stratum radiatum (WT n=9; WT+IL-6 n=5). LTP was induced by high frequency stimulation (HFS) at the point indicated by the arrow.
- B) Immediately after LTP induction by HFS (5 minutes after LTP) the magnitude of the potentiation in in WT slices treated with IL-6 is significantly reduced in comparison to control WT slices. Unpaired t test, $p < 0.05^*$
- C) Representative traces show fEPSP traces from WT controls (black line) and WT IL-6 treated (teal line).

Following HFS in *mdx* slices perfused with exogenous IL-6, we did not see the same effect as we did in WT. Interestingly, there was no difference between *mdx* slices perfused with exogenous IL-6 and *mdx* control slices (*mdx* 168±38% vs *mdx* + IL-6 181±20%; $p=0.5$, unpaired t test, figure 4.14b). Also, the fEPSP slope over time in *mdx* slices perfused with exogenous IL-6 look nearly identical to *mdx* controls (figure 4.14b).

In fact, the fEPSP slope in WT slices perfused with exogenous IL-6 mirrored the fEPSP slope over time of control *mdx* mice (WT+IL-6, 177±25% vs *mdx* 168±38%, $p=0.65$, unpaired t test, figure 4.15b). This similarity was seen in the fEPSP slope for 50 minutes (figure 4.15a), after which WT slices perfused with exogenous IL-6 remain stable, whereas there is the decay in *mdx* controls that we also saw in WT controls. As expected, given that exposure to IL-6 reduced early LTP in WT mice but did not modify this LTP in *mdx* mice, the fEPSP slopes were also similar in WT and *mdx* hippocampal slices exposed to IL-6 (WT+IL-6, 177±25% vs *mdx* + IL-6 181±20%; $p=0.81$, unpaired t test, figure 4.16b). In the fEPSP slope over time (figure 4.16a), early 'stabilisation' of both WT and *mdx* by exogenous IL-6 was apparent.

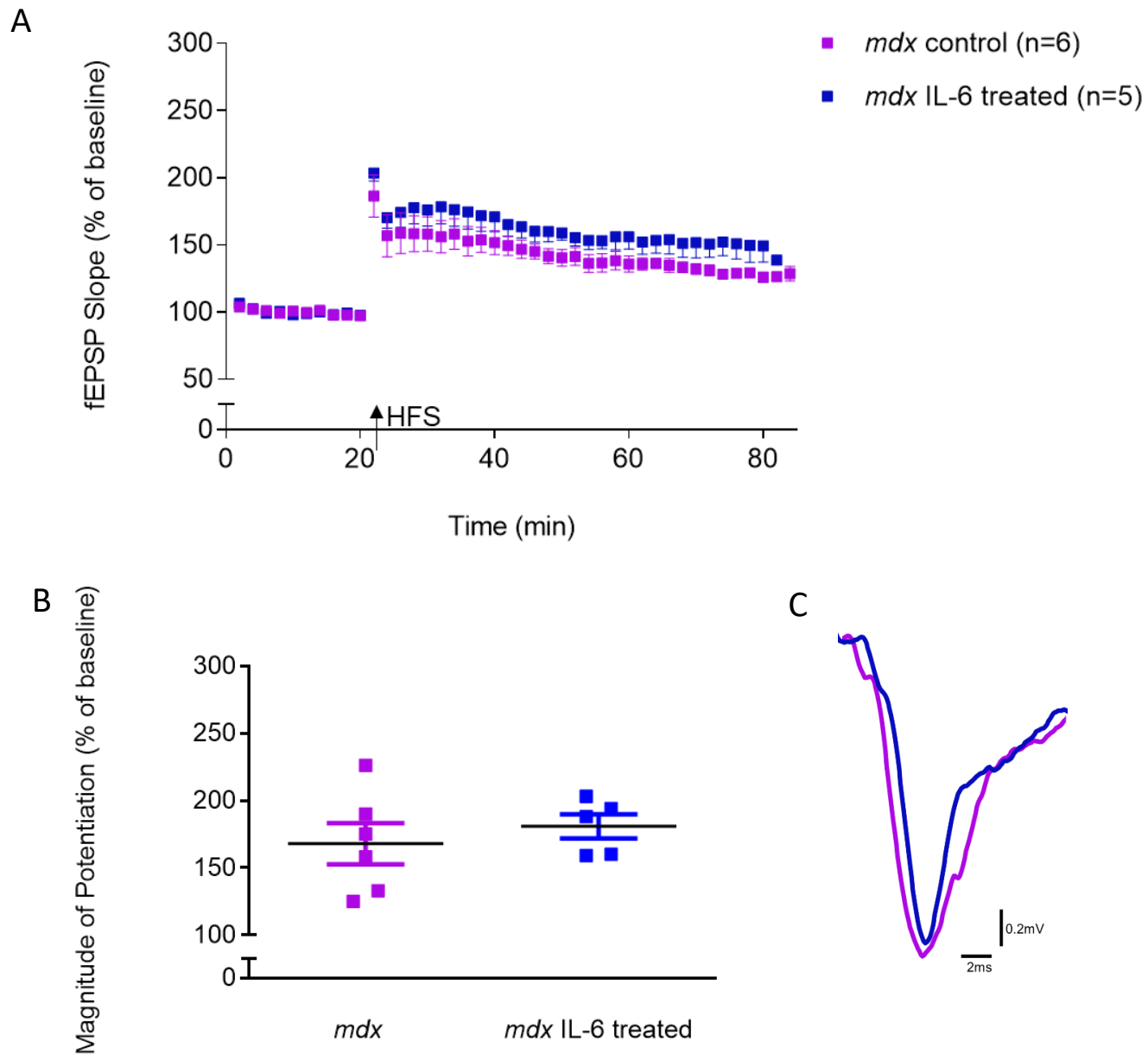
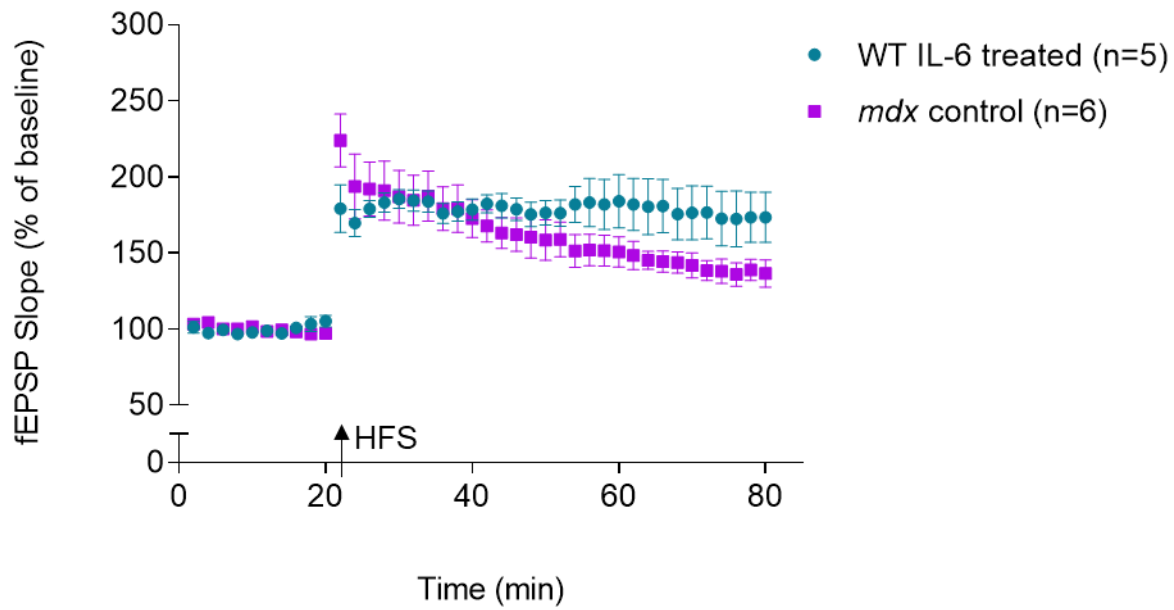


Figure 4.14: LTP in mdx hippocampal Schaeffer collaterals-CA1 synapses was not changed by IL-6.

- A) Graph showing averaged time course of Schaffer collateral-evoked synaptic responses from mdx controls (purple squares) and mdx slices perfused with IL-6 (blue squares) in the hippocampal region using extracellular field potential recordings in CA1 stratum radiatum (mdx n=6; mdx+IL-6 n=5). Exposure to IL-6 (1nM) for 60 minutes prior to recording and for the duration of experiments.
- B) Immediately after LTP induction by HFS (5 minutes after LTP) the magnitude of the potentiation in mdx slices treated with IL-6 is similar to control mdx slices. Unpaired t test, $p > 0.05$
- C) Representative fEPSP traces from mdx control (purple line) and mdx IL-6 treated (dark blue line) immediately after HFS.

A



B

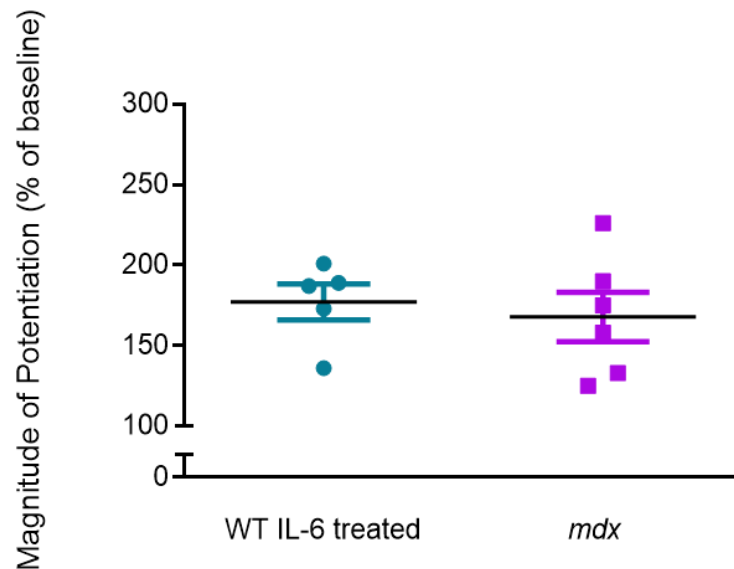


Figure 4.15: LTP generated in WT hippocampal Schaeffer collateral-CA1 synapses exposed to IL-6 is similar to untreated mdx LTP

- A) Graph showing averaged time course of Schaeffer collateral-evoked synaptic responses from mdx controls (purple squares) and WT slices perfused with IL-6 (blue circles) in the hippocampal region using extracellular field potential recordings in CA1 stratum radiatum (mdx n=5; mdx+IL-6 n=5). Exposure to IL-6 (1nM) for 60 minutes prior to recording and for the duration of experiments.
- B) Immediately after LTP induction by HFS (5 minutes after LTP), the magnitude of the potentiation slope in WT slices treated with IL-6 is similar to control mdx slices. Unpaired *t* test, $p > 0.05$

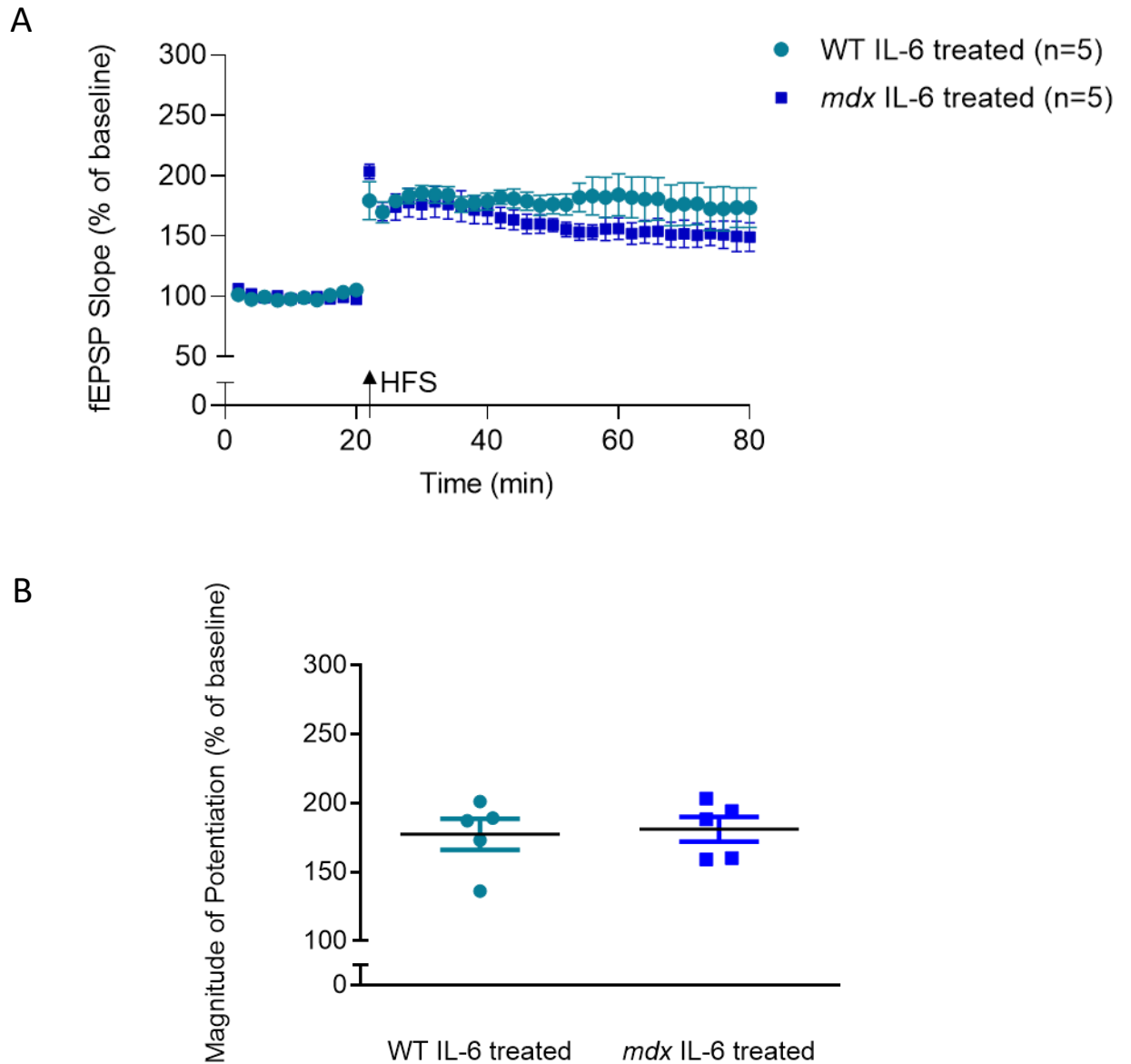


Figure 4.16: LTP generated in WT hippocampal Schaeffer collateral-CA1 synapses exposed to IL-6 is similar to LTP generated in mdx exposed to IL-6.

- A) Graph showing averaged time course of Schaeffer collateral-evoked synaptic responses from mdx perfused with IL-6 (blue squares) and WT slices perfused with IL-6 (blue circles) in the hippocampal region using extracellular field potential recordings in CA1 stratum radiatum (WT+IL-6 n=5; mdx+IL-6 n=5). Exposure to IL-6 (1nM) for 60 minutes prior to recording and for the duration of experiments.*
- B) Immediately after LTP induction by HFS (5 minutes after LTP), the magnitude of the potentiation in WT slices treated with IL-6 is similar to mdx slices treated with IL-6. Unpaired t test, $p>0.05$*

4.10. Discussion

As brain dysfunction associated with loss of full-length dystrophin, Dp427, is reported to be due to dysfunctional synaptic inhibition (Perronnet & Vaillend, 2010), and neuronal dystrophin is normally expressed at PSDs, it is likely that dystrophin plays a role in synapse structure and function (Blake & Kroger, 2000). Indeed, we found that dystrophin was expressed in clusters along neuronal processes that co-localised with synaptophysin, a biomarker of synapses. We also found that IL-6R is expressed in similar way to dystrophin in that it also clusters along neuronal processes at synapses, as such the effects of IL-6R activation on synapses may be modified if dystrophin is absent from this region.

Given the elevated levels of peripheral IL-6 that is characteristic of DMD, we investigated whether IL-6 in the CNS was also elevated in the absence of dystrophin. Gene and protein expression of IL-6 in whole homogenised hippocampal tissue was not different to WT control tissue. Similarly, gene expression of IL-6R, gp130, the transmembrane protein which initiates the downstream signalling cascade, and STAT3, a signalling molecule activated by IL-6 were unchanged. However, the caveat of this experimental design is that regional differences, which are critical to the function of the hippocampus, cannot be elucidated. Further investigation, using immunofluorescent localisation demonstrated region-specific changes in IL-6 and IL-6R expression in dystrophin-deficient *mdx* tissue. There was an increase in IL-6 and IL-6R expression in the CA1 and DG region of the hippocampus of *mdx* mice in comparison to WT with a surprising decrease in IL-6R in the CA3 region in *mdx* mice compared to WT. The

CA1 and CA3 region of the hippocampus is important in LTP, the proposed correlate of learning and memory, as LTP can be induced along the Schaeffer-collateral (emanating in CA3)– CA1 pathway. Further, the DG, in the hippocampus, is of importance as it is one of the few brain regions where neurogenesis occurs and adult-born neurons are integrated into pre-existing neuronal circuits (Gage, 2000). In fact, IL-6 has been shown to influence neural stem cells in hippocampal adult neurogenesis (Islam *et al.*, 2009), with proliferation, survival and differentiation of hippocampal progenitor cells being compromised in the dentate gyrus when IL-6 concentrations are elevated (Vallieres *et al.*, 2002). Thus, an increase in IL-6 in *mdx* mice in these regions, could in part, explain deficits in neurogenesis and LTP. As well as a possible impact of IL-6 on neurogenesis and LTP, LTP itself can impact on neurogenesis (Bruehl-Jungerman, 2006). A study by Bruehl-Jungerman (2006) found that electrically induced LTP in the dentate gyrus *in vivo* provides a cellular/molecular environment that favours both proliferation and survival of adult-generated neurons, thus there likely may be a sort of coupling between synaptic plasticity and neurogenesis. Further, LTP can be induced more easily in newly born cells such as granule cells in the dentate gyrus (Schmidt-Hieber *et al.*, 2004), thus an increase of neurogenesis in *mdx* DG that has previously been reported (Deng *et al.*, 2009) may underlie changes in LTP and associated cognitive deficits seen in DMD.

Our results showed that exposure to IL-6 reduced early-phase LTP induction in WT slices and this finding supports previous work which has also shown that IL-6 can modify LTP (Li *et al.*, 1997; Tancredi *et al.*, 2000; Stampanoni Bassi *et al.*, 2019). However, this difference was only seen in early LTP. At a molecular level, early LTP

includes activation of AMPA receptors, calcium influx through activated NMDA receptors which subsequently activates CaMKII, which eventually potentiates synaptic transmission. Later (> 1 hour following the high frequency stimulation) stimulation of gene transcription leads to *de novo* protein synthesis. However, if early LTP is disrupted or modified, that might impact on later cellular events. Importantly, IL-6 can serve as a negative regulator of both early and late LTP in the hippocampus (Bellinger et al., 1995; Li et al., 1997; Tancredi et al., 2000; Balschun et al., 2004) through multiple signalling pathways including p42/44 mitogen-activated protein kinase (MAPK) (Engberink et al., 2017). Interestingly though, after endogenous addition of IL-6 to *mdx* slices, there was no effect on LTP induction. Although, the exact mechanism through which IL-6 could modify LTP is not yet known, this could be due to decreased IL-6R expression in the CA3 region of *mdx* mice, where the Schaffer collaterals are stimulated. Thus, chronic exposure to elevated IL-6 could have resulted in receptor internalisation or it is possible that reduced LTP observed in *mdx* mice could be due to a complex interplay between local IL-6 levels paired with abnormal neuron functioning due to a loss of synaptic dystrophin. As was noted with aberrant GABA_AR clustering in *mdx* hippocampal neurons (Knuesel et al., 1999; Knuesel et al., 2001), it is possible that hippocampal IL-6Rs may not be appropriately expressed on the neuronal cell membrane, or, perhaps down-stream signalling is dysfunctional resulting in insensitivity to IL-6. While further research is needed to reveal the precise signalling mechanisms, these studies suggest that while overall IL-6 levels or IL-6R levels in the brain are not changed in dystrophic mice, there may be local changes in expression that could modify hippocampal LTP. Increases in IL-6 expression could be due to increases in

circulatory IL-6 that has been transported from the periphery across the BBB (Banks *et al.*, 1994) or IL-6 could be released locally in the brain from astrocytes and microglia during pathophysiological events such as inflammation (Gruol & Nelson, 1997), both of which could occur in DMD due to increased peripheral IL-6 (Rufo *et al.*, 2011; Pelosi *et al.*, 2015) and inflammation caused by a loss of dystrophin.

We also investigated the effect of exogenous IL-6 on calcium homeostasis in cultured hippocampal neurons. Though, exogenous IL-6 had no effect on neuronal activity in the *mdx* mouse. This aligned with our findings that the addition of IL-6 does not alter hippocampal LTP in *mdx* mice. Importantly, the experiments on neuronal activity were completed on cultured hippocampal neurons whereas, LTP experiments were done on intact hippocampal slices which could imply that there are other complex mechanisms at play on a physiological level, not just at a cellular level.

4.11. Conclusions

Although we did not detect changes in mRNA expression of components of the IL-6 pathway, namely IL-6, IL-6R, gp130 and STAT3, in whole hippocampi of *mdx* mice, we did see regions specific increases in both IL-6 and IL-6R in the CA1 and DG region of the hippocampus. This increase in IL-6 could possibly be due to the elevation of IL-6 in the periphery of *mdx* mice (Pelosi *et al.*, 2015) as IL-6 has been shown to cross the BBB *via* saturable transport mechanisms (Banks *et al.*, 1994) or local production of IL-6 from astrocytes and microglia, which are activated during inflammation.

Consistent with the findings of others (Li *et al.*, 1997; Tancredi *et al.*, 2000; Stampanoni Bassi *et al.*, 2019) and our own experiments, addition of exogenous IL-6 suppressed early-phase LTP in dystrophin-expressing WT mice. Following tetanic stimulation, changes in post-synaptic expression and activation of AMPA and NMDA receptors lead to depolarisation of the post-synaptic neuron and influx of calcium. This in turn results in activation of intracellular CaMKII and other protein kinases and are the key first steps in creating new memories. Elevated IL-6 suppresses this activity under normal circumstances, however, *mdx* mice were resistant to this effect. This could be due to altered expression of IL-6Rs in *mdx* hippocampal tissue. While increased IL-6R expression was observed in the CA1 and DG regions, IL-6R expression was specifically decreased in the CA3 region, the site where the tetanic stimulus is applied to generate LTP. While further work is needed to elucidate why these changes in IL-6R expression have occurred in specific dystrophic hippocampal regions, it does appear that these changes have had functional consequences in the cellular and molecular processes that underpin learning and the consolidation of memories

Chapter 5

**Intervention study: Neutralisation
of IL-6 receptor-mediated
signalling improves cellular
correlates of learning and memory.**

5.1. Introduction

As demonstrated in the previous chapter, IL-6 could modulate the cellular processes underpinning learning and memory (figure 4.14, pg. 76). Research has shown that circulating levels of IL-6 are higher in severe neurological pathologies such as Alzheimer's and dementia (Campbell *et al.*, 1993) and similarly, transgenic mice over-expressing IL-6 exhibit broad memory impairment (Heyser *et al.*, 1997; Wei *et al.*, 2012; Wei *et al.*, 2013). Other studies have shown that transgenic mice deficient in IL-6 possess enhanced spatial learning capabilities in the radial arm maze (Braida *et al.*, 2004) and the Morris water maze (Bialuk & Winnicka, 2018) behavioural task-learning assessments, further emphasising the pleiotropic actions of this cytokine in the CNS. Consistent with these findings, IL-6 has also been shown to inhibit LTP (Li *et al.*, 1997; Tancredi *et al.*, 2000; Stampanoni Bassi *et al.*, 2019) as well as a blockade of endogenous IL-6 with a neutralizing IL-6 antibody, resulted in a remarkable prolongation of LTP, as well as improved performance in spatial memory tests (Balschun *et al.*, 2004).

In the *mdx* mouse model of DMD, inhibition of IL-6 signalling with the monoclonal anti-IL-6 receptor antibody (xIL-6R) improves the muscular physiology of DMD such as increasing diaphragm muscle force (Manning *et al.*, 2017) and overall respiratory function (Burns *et al.*, 2018) in *mdx* mice. Skeletal muscle regeneration is also improved by xIL-6R treatment (Wada *et al.*, 2017). It is also notable that another recombinant (humanized) monoclonal IL-6R antagonist (tocilizumab), which has been approved as an anti-inflammatory drug for inflammatory diseases such as rheumatoid arthritis, Castleman's disease, and systemic juvenile idiopathic arthritis,

has also shown promise in alleviating the muscular symptoms of DMD (Wada *et al.*, 2017). Tocilizumab blocks IL-6-mediated signalling via an inhibition of the binding of IL-6 to both soluble and transmembrane IL-6Rs in the same way as xIL-6R.

DMD patients have an increased incidence of mild-to-moderate cognitive impairment (Bresolin *et al.*, 1994; Anderson *et al.*, 2002) and exhibit impaired verbal, short-term and working memory (Snow *et al.*, 2013). Similarly, *mdx* mice display behaviours that are consistent with dysfunctions of the hippocampus and amygdala, such as a poor capacity to learn new tasks and deficits in spatial learning tasks, relative to controls (Vaillend *et al.*, 1995; Vaillend *et al.*, 2004; Chausseot *et al.*, 2015). They also display deficits in associative learning and in the general processes involved in memory formation and consolidation which are both dependent upon optimal functioning of the hippocampus and amygdala (Chausseot *et al.*, 2015). Along with cognitive impairment, DMD is also associated with a number of behavioural issues, such as anxiety, depression and Attention-Deficit / Hyperactivity Disorder (ADHD) (Rubinshtein *et al.*, 1998; Hendriksen & Vles, 2008). Behavioural abnormalities have also been observed in the *mdx* mouse model, with increases in both depressive behaviour and anxiety and fearfulness observed relative to controls (Manning *et al.*, 2014), sometimes independently of any skeletal muscle impairment (Sekiguchi *et al.*, 2009). Interestingly, increased levels of IL-6 in the blood have been associated with major depressive disorder (Dowlati *et al.*, 2010; Haapakoski *et al.*, 2015). Similarly, IL-6 knockout mice (IL-6 KO) show alterations in anxiety-like and depression-like behaviours (Armario *et al.*, 1998; Butterweck *et al.*, 2003; Chourbaji *et al.*, 2006; Aniszewska *et al.*, 2015).

Given that inhibition of IL-6-mediated signalling in the brain appears to enhance learning and memory, we hypothesised that central administration of neutralising αIL-6R antibodies to block IL-6 signalling, would have beneficial effects on hippocampal function in dystrophic animals.

5.1.1. Study objectives

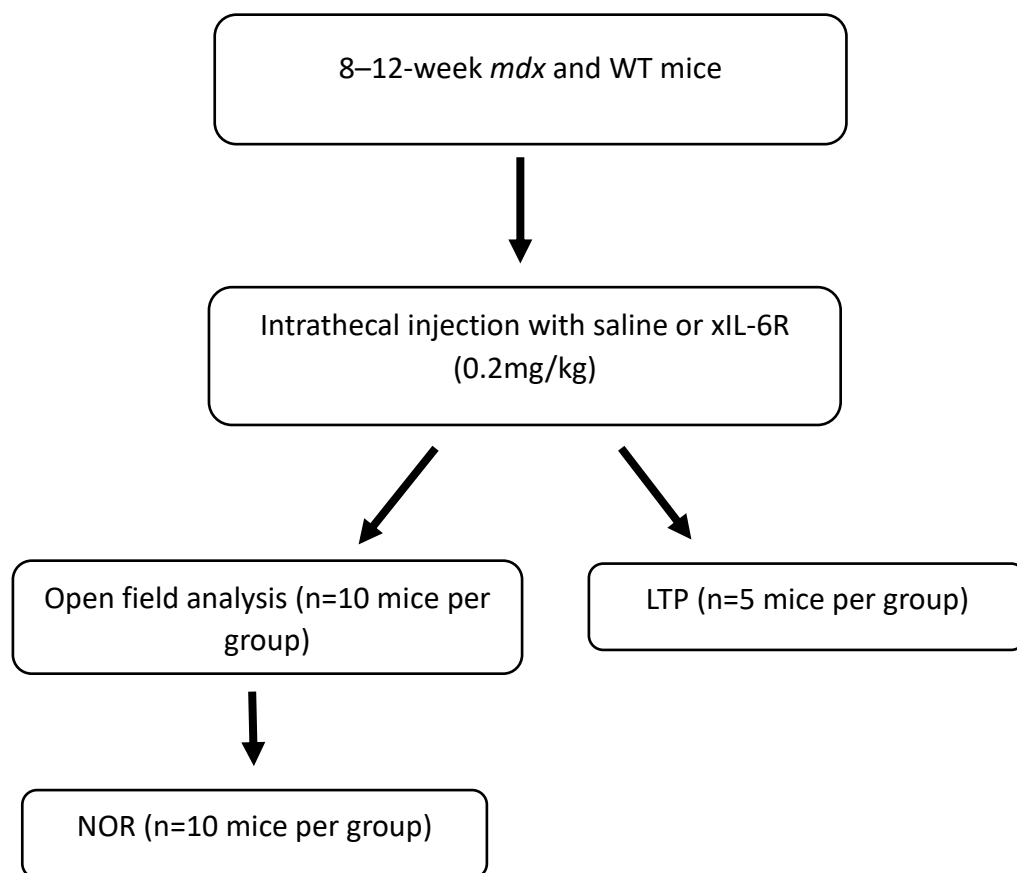
- To investigate if intrathecal delivery of monoclonal anti-IL-6 receptor antibodies modifies hippocampal long-term potentiation in dystrophin-deficient *mdx* mice.
- To determine if behavioural patterns are modified by the administration of monoclonal anti-IL-6 receptor antibodies.

5.1.2. Techniques used

- Intrathecal injection (see methods section 2.9, pg. 65).
- Extracellular electrophysiology (see methods section 2.8, pg. 59).
- Open field behavioural test (see methods section 2.7.3, pg. 58).

5.2. Study design for intervention

WT and *mdx* mice aged 8-12 weeks were intrathecally injected with a monoclonal anti-IL-6 receptor (αIL-6R) antibody between the L5 and L6 region of the spine (see methods section 2.9, pg. 64). Mice were allowed to recover for 24 hours and then underwent behavioural studies and electrophysiological studies.



5.3. Intrathecal administration of xIL-6R antibodies normalises hippocampal LTP in *mdx* mice.

In chapter 3, we reported that LTP was significantly reduced in *mdx* mice in comparison to WT (Figure 3.4, pg. 76) thus, we sought to determine if *in vivo* neutralisation of central IL-6 signalling modified this.

Following all LTP experiments in this thesis, monopolar stimulation of the Schaffer collaterals was performed through an electrode located on the border between areas CA1 and CA3. Baseline stimulations were recorded for 20 minutes following input/output curves at the intensity (mA) required to obtain 50% of the maximum output. High-frequency stimulation (HFS – 4 x 100 pulses, 100Hz for 1s/60s) followed the baseline, with stimulation intensity at 30% required for maximum output.

In *mdx* mice administered with intrathecal xIL-6R, immediately after HFS, the magnitude of LTP was significantly increased compared to saline treated *mdx* animals (saline treated $199.5 \pm 32\%$ vs xIL-6R $282.3 \pm 6.11\%$, $p=0.0074$, $n=3$ animals, unpaired students t test, figure 5.1b, c). Furthermore, this potentiation was observed for the remaining 80 minutes of each experiment following HFS, such that LTP was significantly increased in *mdx* mice treated with xIL-6R ($n=3$) hippocampal slices as compared to saline treated *mdx* animals ($n=3$, Genotype factor, $F(1, 3) = 10.51$, $p=0.0478$, two-way repeated-measures ANOVA starting at 1 min after stimulation, figure 5.1a).

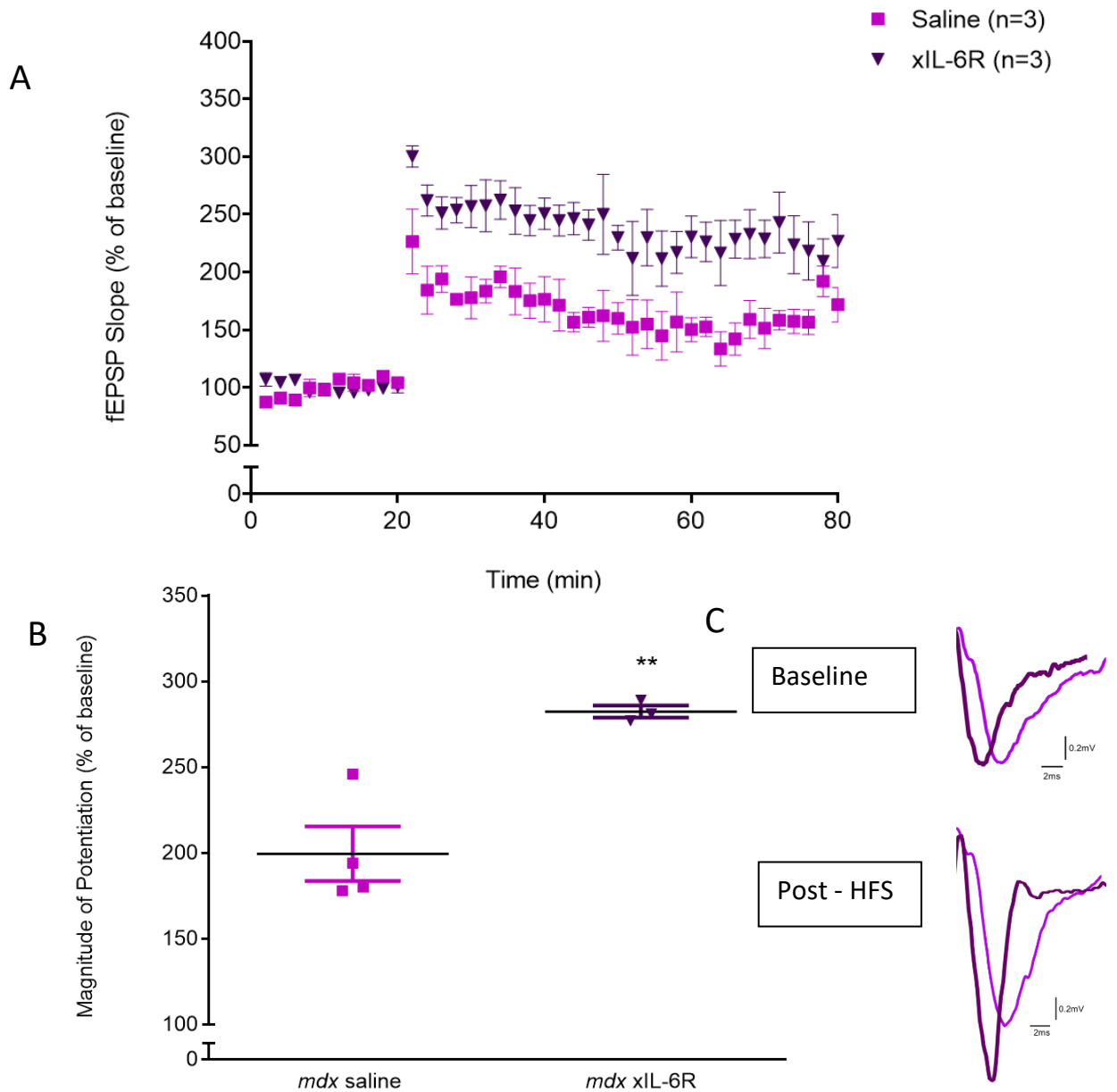


Figure 5.1: Long-term potentiation in the SC-CA1 pathway of the hippocampus in mdx mice intrathecally-injected with xIL-6R is enhanced compared to saline treated mdx controls.

- A)** Graph showing averaged time course of Schaffer collateral-evoked synaptic responses from saline treated mdx (purple squares) and xIL-6R-treated mdx (dark purple triangles) in the hippocampal region using extracellular field potential recordings in CA1 stratum radiatum (mdx n=3; xIL-6R-treated mdx n=3). Two-way ANOVA, $p < 0.001^{***}$
- B)** Immediately after LTP induction by HFS, the magnitude of potentiation in mdx mice intrathecally injected with xIL-6R is enhanced in comparison to saline treated controls (n=3, $p = 0.0074^{**}$). Unpaired t test, $p < 0.01^{**}$
- C)** Representative traces show fEPSPs before and after high frequency stimulation to induce LTP in saline treated and xIL-6R treated mdx mice. At baseline, untreated (light purple) and treated (dark purple) fEPSP slopes (mV/ms) are similar, whereas, post-HFS, fEPSP slopes are significantly increased in treated mdx mice.

Unlike its restorative effects on LTP in *mdx* mice, *in vivo* treatment of WT mice with xIL-6R did not have the same effect. Immediately following HFS, there was no significant difference between the magnitude of potentiation of the fEPSP for the first 5 minutes in WT mice treated with xIL-6R and saline treated WT controls (saline $192 \pm 24\%$ vs xIL-6R $209 \pm 41.9\%$, $p=0.3835$, $n=3-4$ animals, unpaired students t test; Figure 5.2b, c). Furthermore, this potentiation was observed for the remaining 80 minutes of each experiment following HFS, with no change in LTP magnitude between slices from WT mice treated with xIL-6R ($n=4$) hippocampal slices compared to saline treated WT controls ($n=3$, Genotype factor, $F(1, 4) = 0.1953$, $p=0.6813$, two-way repeated-measures ANOVA starting 1 min after stimulation, figure 5.2a).

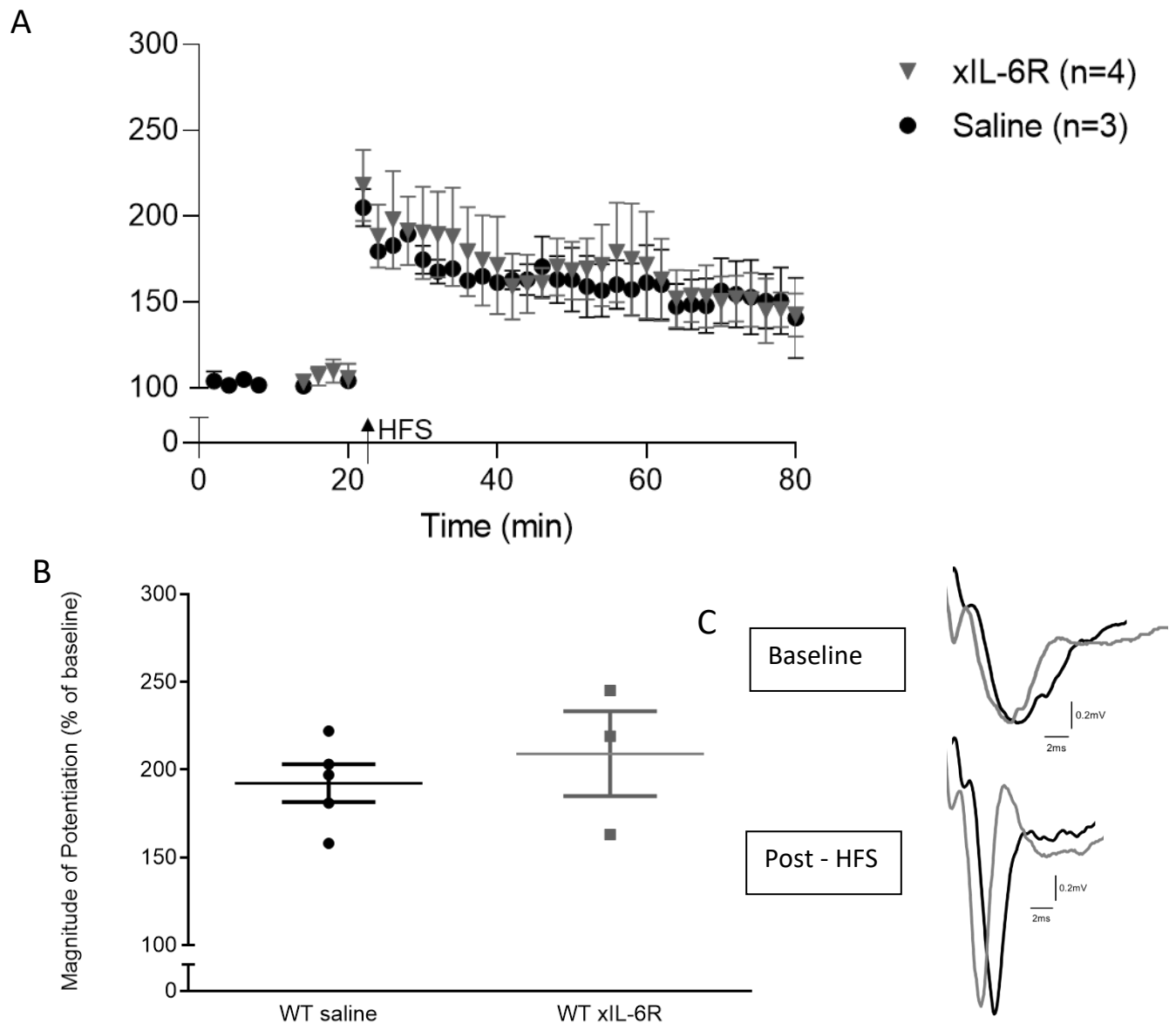


Figure 5.2: Long-term potentiation in the SC-CA1 pathway of the hippocampus in WT mice intrathecally-injected with xIL-6R is similar compared to untreated WT controls.

- A) Graph showing averaged time course of Schaffer collateral-evoked synaptic responses from saline treated WT (black circles) and xIL-6R-treated WT (grey triangles) in the hippocampal region using extracellular field potential recordings in CA1 stratum radiatum (saline treated WT n=3; xIL-6R-treated WT n=3). Two-way ANOVA, $p > 0.05$
- B) Immediately after LTP induction by HFS, the magnitude of potentiation in WT mice intrathecally injected with xIL-6R is similar to saline treated controls (n=3-4, $p = 0.4885$). Unpaired t test, $p > 0.05$
- C) Representative traces show fEPSPs before and after high frequency stimulation to induce LTP in saline treated and xIL-6R treated WT mice. At baseline and post-HFS, untreated (black) and treated (grey) fEPSP slopes (mV/ms) are similar.

As a control for the injection procedure itself, the effects of intrathecal administration of saline on LTP induction and maintenance were examined and compared to those from the original, untreated WT animals and untreated *mdx* animals.

In WT mice, LTP magnitude was not significantly different between saline-injected and untreated WT mice, suggesting that the injection procedure itself had no effect on either LTP induction and/or maintenance. The initial magnitude of potentiation following HFS did not differ significantly between untreated and saline-injected WT animals (untreated WT, 216.4 ± 22 vs saline 200.8 ± 16.9 , $p = 0.1308$; unpaired student t test, $n=4-9$ animals, figure 5.3b). Furthermore, this potentiation was observed for the remaining 80 minutes of each experiment following HFS, with no change in LTP between WT mice injected with saline ($n=4$) as compared to untreated controls ($n=9$, factor genotype, Genotype factor, $F(1, 11) = 0.8061$, $p = 0.3885$, two-way repeated-measures ANOVA starting at 1 min after stimulation, figure 5.3a).

The injection procedure had a similar lack of significant effect on LTP induction and/or maintenance in slices from *mdx* mice. The initial magnitude of potentiation following HFS did not differ significantly between untreated and saline-injected *mdx* animals (untreated *mdx* 174.8 ± 37.5 vs saline, vs 199.5 ± 31.8 , $p = 0.33$; unpaired student t test, $n = 4-5$ animals, figure 5.4b). Furthermore, this potentiation was observed for the remaining 80 minutes of each experiment following HFS, with no change in LTP between *mdx* mice injected with saline ($n=4$) as compared to untreated controls ($n=5$, factor genotype, Genotype factor, $F(1, 7) = 0.1513$,

$p=0.71$, two-way repeated-measures ANOVA starting at 1 min after stimulation, figure 5.4a). This provided us with confidence that the restoration of LTP in *mdx* mice was due to central neutralisation of IL-6Rs.

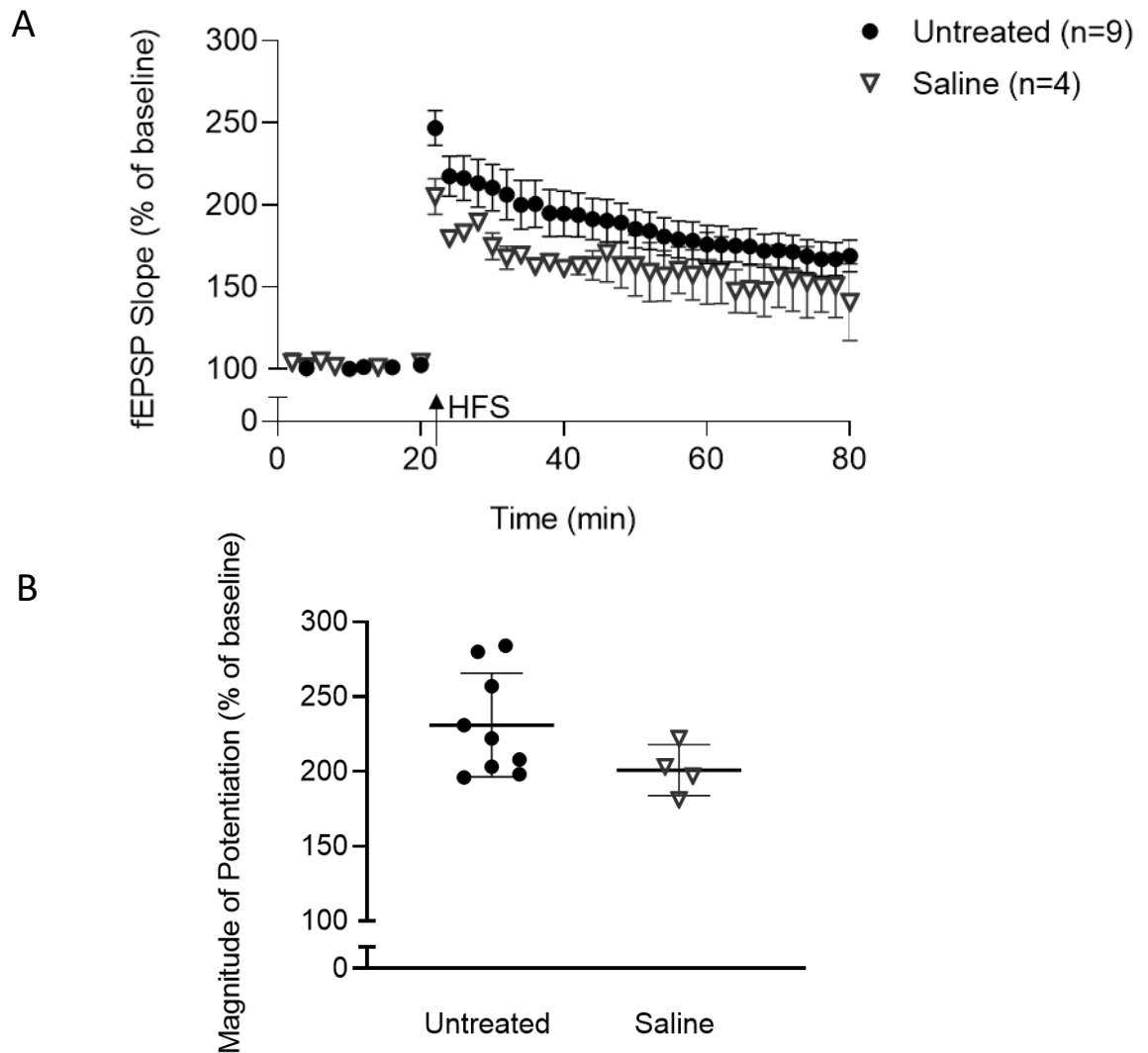


Figure 5.3: Intrathecal injection of saline had no impact on LTP induction and/or maintenance in WT mice.

- A) Graph showing averaged time course of Schaffer collateral-evoked synaptic responses from untreated WT (black circles) and saline-treated WT (grey triangles) in the hippocampal region using extracellular field potential recordings in CA1 stratum radiatum (WT n=9; saline treated WT n=4). Two-way ANOVA, $p>0.05$
- B) Immediately after LTP induction by HFS, the magnitude of the potentiation in WT mice intrathecally injected with saline is similar to untreated WT mice (n=4-9, $p=0.252$). Unpaired t test, $p>0.05$

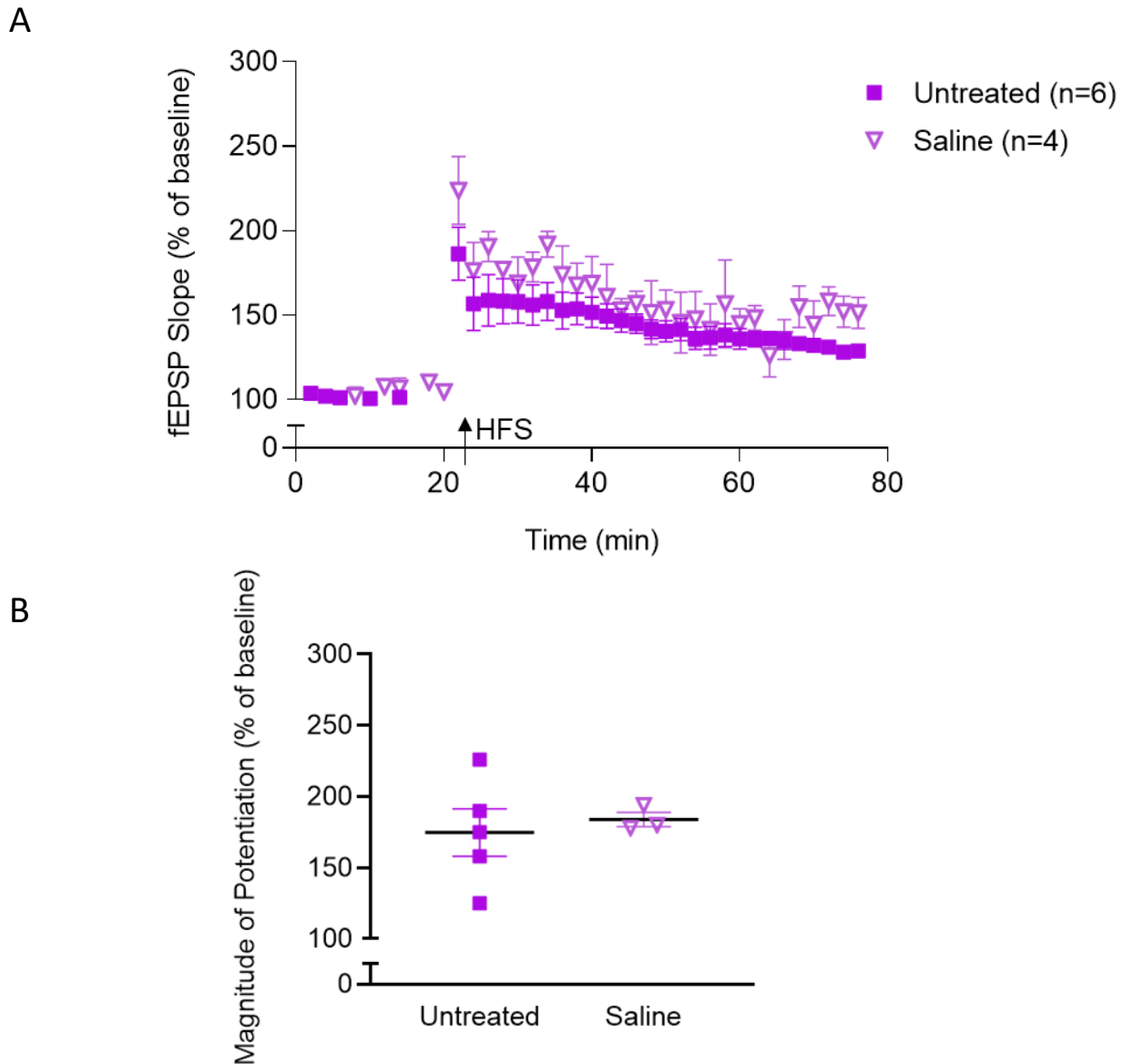


Figure 5.4: Intrathecal injection of saline had no impact on LTP induction and/or maintenance in mdx mice.

- A) Graph showing averaged time course of Schaffer collateral-evoked synaptic responses from untreated mdx (purple squares) and saline-treated mdx (open purple triangles) in the hippocampal region using extracellular field potential recordings in CA1 stratum radiatum (mdx n=6; saline treated mdx n=4). Two-way ANOVA, $p > 0.05$
- B) Immediately after LTP induction by HFS, the magnitude of potentiation in mdx mice intrathecally injected with saline is similar to untreated mdx mice (n=4-6, $p = 0.33$). Unpaired t test, $p > 0.05$

5.4. xIL-6R administration did not modify NOR recognition memory in WT or *mdx* mice.

Having observed such a dramatic restoration of the cellular correlate of learning and memory in *mdx* mouse hippocampal slices intrathecally injected with xIL-6R, we next investigated whether or not this potentiation of LTP in *mdx* mice translated into functional changes in behaviour, by repeating the NOR test in these mice.

As described in chapter 3, NOR uses two similar objects on a first testing day, and the mouse is left to explore for 10 minutes. 24 hours later, one of the similar objects is replaced with a novel object and once again the mouse is left to explore the area for 10 minutes. A percent of time spent exploring the novel object relative to the total time spent exploring both objects can be a measure of novel object recognition (Benice *et al.*, 2006; Sarkisyan & Hedlund, 2009; Broadbent *et al.*, 2010; Oliveira *et al.*, 2010). This concept can be represented by *Recognition Index* (RI), i.e., the time spent investigating the novel object relative to the novel object and the familiar object investigation [$RI = T_N / (T_N + T_F)$], and it is the main index of retention (Mumby *et al.*, 2002; Piterkin *et al.*, 2008; Botton *et al.*, 2010; Gaskin *et al.*, 2010; Schindler *et al.*, 2010).

Despite its clear effects on LTP, intrathecal injection of xIL-6R did not appear to have any effect on recognition memory. No significant differences were seen in the time spent investigating the novel object relative to the novel object and the familiar object investigation (RI) between WT and *mdx* mice treated with either saline or xIL-6R (WT saline 0.76 ± 0.04 vs xIL-6R 0.77 ± 0.027 , $p=0.7088$ and *mdx* saline 0.79 ± 0.036 vs *mdx* xIL-6R 0.8 ± 0.02 , $p=0.7601$, unpaired t test, Figure 5.6).

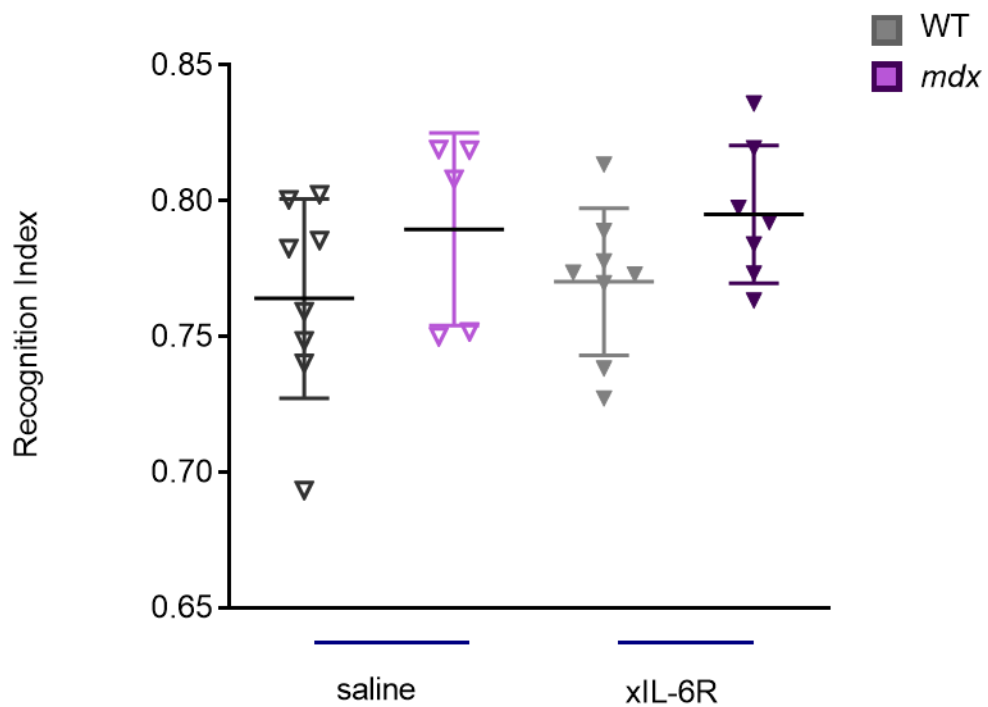


Figure 5.5: The NOR recognition index values for *mdx* and WT mice injected intrathecally injected with either saline or xIL-6R did not differ significantly from each other. Unpaired *t* test, $p > 0.05$

5.5. Anxiety-like behaviours in *mdx* mice.

We showed in chapter 3 that freezing behaviours, which are associated with a fear response in mice (Sekiguchi *et al.*, 2009), are significantly increased in untreated *mdx* mice in comparison to untreated WT mice. Therefore, we wanted to next investigate the effects of intrathecal administration of xIL-6R on freezing and anxiety-like behaviours (methods section 2.7.3, pg. 58).

These experiments showed that *mdx* mice administered with saline alone ($n=7$) were significantly more reluctant to enter the centre of the open field arena in comparison to saline-treated WT mice ($n=10$, two-way ANOVA, Strain, $F(1, 29) =$

9.063, $p=0.005$, Figure 5.6a) which is a behaviour consistent with increased anxiety (Hölter *et al.*, 2015) in this strain. Intrathecal injection of xIL-6R had no impact on this anxiety-like phenotype in the *mdx* mice ($n=7$, Intervention, $F(1, 29) = 0.14$, $p=0.71$).

Increased freezing, a recognised fear response in mice (Sekiguchi *et al.*, 2009) did not significantly differ between WT ($n=9$) and *mdx* mice ($n=7$) injected with either saline (two-way ANOVA, Strain, $F(1, 29) = 2.0$, $p=0.17$, figure 5.6b) or xIL-6R (two-way ANOVA, intervention, $F(1, 29) = 0.1$, $p=0.75$).

Grooming frequency also did not differ significantly between WT ($n=9$) and *mdx* mice ($n=7$) treated with either saline (two-way ANOVA, Strain, $F(1, 27) = 0.006$, $p=0.9$, figure 5.6c) or xIL-6R (two-way ANOVA, Intervention, $F(1, 27) = 0.1538$, $p=0.7$).

Escape behaviours were also assessed and showed that rearing frequency increased significantly in saline-treated *mdx* mice ($n=6$) in comparison to saline-treated WT mice ($n=10$, two-way ANOVA, Strain, $F(1, 28) = 6.054$, $p=0.02$, figure 5.6d).

However, xIL-6R administration had no significant effect on the rearing frequency observed in the WT ($n=9$) and *mdx* mice ($n=7$, two-way ANOVA, Intervention, $F(1, 28) = 0.006$, $p=0.94$).

When we next counted the number of faecal boli excreted over 10 minutes as a measure of anxiety, we found that the number excreted by saline-treated *mdx* mice ($n=7$) did not differ significantly to that excreted by the saline-treated WT mice ($n=9$, two-way ANOVA, Strain, $F(1, 27) = 1.76$, $p=0.2$, figure 5.6e). Further, administration of xIL-6R had no significant effect on the number of faecal boli

excreted by WT (n=8) or *mdx* mice relative to the saline-treated animals (n=7, two-way ANOVA, Intervention, $F(1, 27) = 0.65$, $p = 0.4$).

Finally, we found no significant difference in the frequency of jumping in the saline-treated *mdx* mice (n=7) relative to the saline-treated WT mice (n=10, two-way ANOVA, Strain, $F(1, 24) = 0.77$, $p = 0.4$, figure 5.6f). Further, administration of xIL-6R also had no significant effect on the frequency of jumping observed in either the WT (n=9) or *mdx* mice relative to the saline-treated animals (n=7, two-way ANOVA, Intervention, $F(1, 24) = 0.03$, $p = 0.9$).

Overall, the intrathecal administration of xIL-6R had no effect on these behavioural outputs.

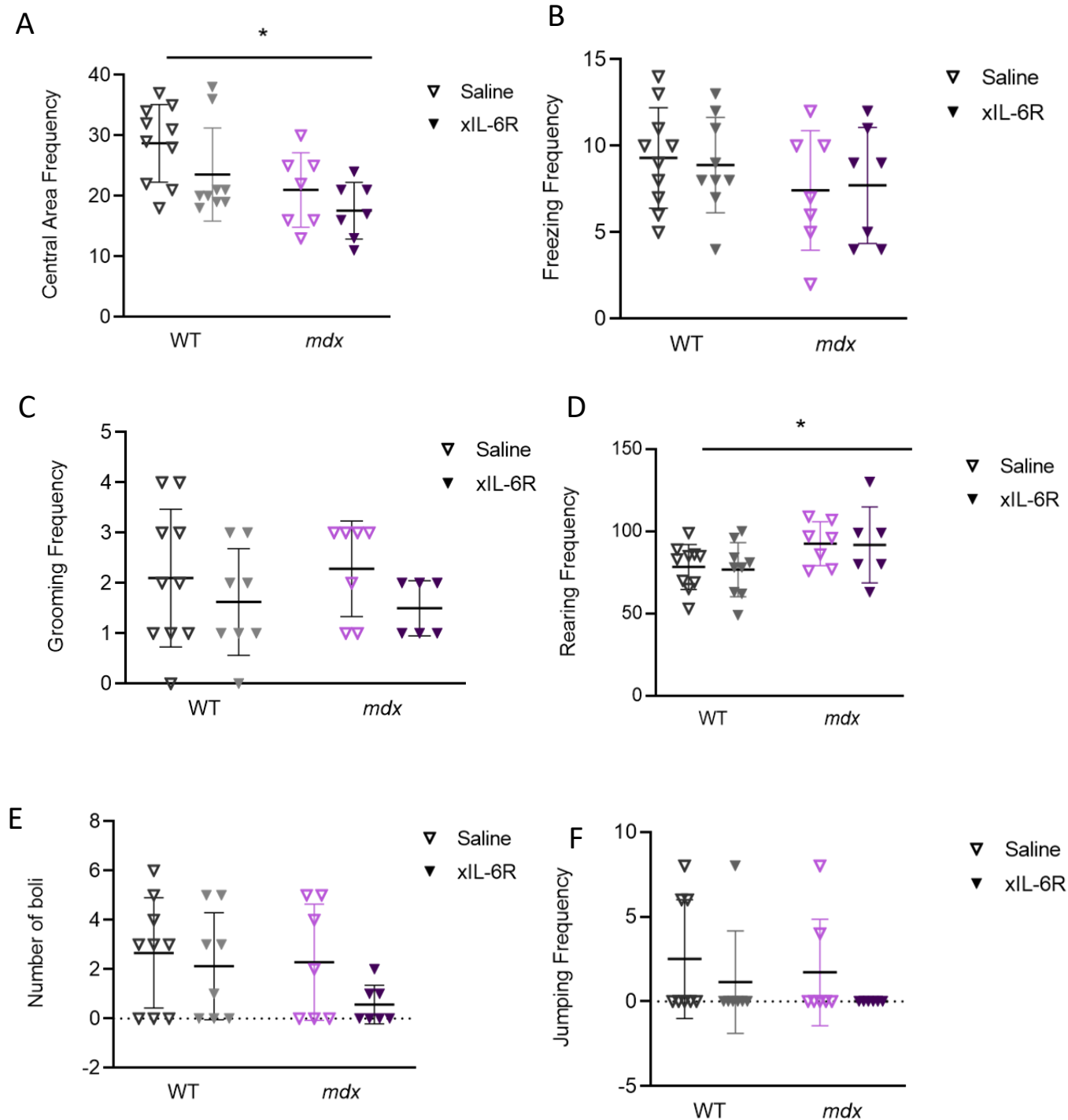


Figure 5.6: Behaviours of saline-treated and xIL-6R treated mdx and WT mice in an open field.

The data plots illustrate that the central area frequency A) and rearing frequency D) is increased in saline-treated mdx mice but with no significant change observed in xIL-6R-treated mice. Freezing frequency B), grooming frequency C), faecal output (number of boli) E) and jumping frequency F) did not differ significantly between saline-treated mdx and WT mice, with no additional changes observed in either WT or mdx xIL-6R-treated mice. Open triangles indicate saline treated mice, closed triangles indicate xIL-6R treated mice. Two-way ANOVA, $p < 0.05$ *

5.6. Discussion

Given the deleterious effects of IL-6 on cognitive function (Campbell *et al.*, 1993; Heyser *et al.*, 1997; Wada-Isoe *et al.*, 2004; Wei *et al.*, 2012; Wei *et al.*, 2013; Lyra *et al.*, 2021), we set out to determine if intrathecal injection of xIL-6R monoclonal antibodies, which bind to IL-6 receptors and block IL-6-mediated signalling, modified hippocampal function and associated behaviours.

Compared to wildtype mice, *mdx* mice exhibit reduced hippocampal LTP as shown in Chapter 3. However, this could be reversed by intrathecal injection of xIL-6R in *mdx* mice. Not only was the initial magnitude of potentiation enhanced immediately after high frequency stimulation, this potentiation was maintained for at least 80 minutes. In fact, the enhanced LTP in *mdx* mice treated with xIL-6R surpassed the magnitude of the LTP evoked in untreated WT controls. Importantly however, neutralisation of central IL-6 receptors with xIL-6R did not modify LTP in WT mice, suggesting that the importance of IL-6 signalling in modulating LTP only becomes evident and significant in the context of dystrophin loss.

As we have already described region-specific changes in the expression of IL-6 and IL-6 receptors in *mdx* hippocampi (chapter 3), in cells present in the pyramidal cell layer of the CA1 region and granule cell layer of the DG. Our findings suggest that normal LTP induction may be disrupted by aberrant signalling caused by this neuroimmune molecule in dystrophic *mdx* mice.

Despite restoring LTP to control levels *ex vivo*, at the behavioural level, xIL-6R had no effect on recognition memory in *mdx* mice. Had time permitted, we would also

have liked to have investigated if spatial memory was altered in *mdx* mice relative to WT, by using behavioural tests such as the Morris water maze or Barnes maze. Although the use of the Morris water maze to measure hippocampal-dependent spatial navigation and reference memory is supported by numerous studies (Vorhees & Williams, 2006), the Barnes maze may be more valuable in this context as it does not involve forcing animals to swim, which is associated with increased levels of stress in mice (Harrison *et al.*, 2009). Furthermore, swimming may also reduce animals' core body temperature which could affect performance (Iivonen *et al.*, 2003).

In addition to the Novel Object Recognition test which was used to investigate recognition memory, we also looked at several behaviours in an open field. Behavioural issues linked to Autism, depression, anxiety and ADHD have been reported in DMD patients (Rubinsztein *et al.*, 1998; Hendriksen & Vles, 2008; Latimer *et al.*, 2017). Importantly, behavioural issues have also been observed in the *mdx* mouse model (Perronnet *et al.*, 2012) such as increased depressive behaviour, and increased anxiety and fearfulness (Sekiguchi *et al.*, 2009; Manning *et al.*, 2014). In mice, behaviours such as a reluctance to enter the centre of an open field arena, and escape behaviours such as rearing and jumping (Kraeuter *et al.*, 2019), are postulated to be a measure of animals' anxiety (Hölter *et al.*, 2015). Interestingly, we observed a significant increase in reluctance of the *mdx* mice to enter the centre of the open field compared to WT mice, suggesting a higher level of anxiety in these mice. Faecal output is yet another proposed measure of anxiety (Deacon & Rawlins, 2005). However, no alterations were noted in faecal output over the 10-minute study period in either control or xIL-6R-treated WT or *mdx*

mice. For jumping and freezing behaviours, we did not find any significant differences between *mdx* and WT mice in either saline- or xIL-6R-treated groups. Thus, the intrathecal injection itself, of xIL-6R did not affect these behaviours.

Grooming frequency and repetitive jumping in mice has been associated with other centrally-regulated behaviours such as autism-spectrum disorder and OCD (Kalueff et al., 2016). Our results showed that there were no significant differences in these behaviours between WT and *mdx* mice.

However, the lack of effect of the dystrophin-deficient phenotype on these behaviours may be due to the relatively young age of the *mdx* mice used in this study. Thus, memory and cognitive deficits we were investigating may not have fully developed in the 8–12-week-old *mdx* mice we studied here. This suggestion is supported by a recent study showing that cognitive deficits in *mdx* mice were progressive and developed from 4 months of age (7), a finding which also parallels with the fact that the severity of loss of dystrophin in skeletal muscle is also age-dependent in *mdx* mice (Moens et al., 1993). Although *mdx* mice do show progressive muscle weakness and deterioration with age, this deterioration is most evident after 52 weeks of age (Pastoret & Seville, 1995). This slowing of deterioration is primarily due to the compensatory upregulation of utrophin, which is able to perform the same function(s) as dystrophin does (Keep, 2000), resulting in the regeneration of sarcolemmal and plasma membrane integrity, thus saving muscle from deterioration, resulting in a less severe DMD phenotype with regard to skeletal muscle function (Gilbert et al., 1999). Under normal circumstances, utrophin is down-regulated in muscle in the presence of dystrophin (Blake et al.,

1996). It has been suggested that similar compensatory mechanisms may also occur in the dystrophic *mdx* brain. However, upregulation of brain utrophin fails to rescue behavioural alterations such as exploration, emotional reactivity and spatial and fear memories in dystrophin-deficient *mdx* mice (Perronnet *et al.*, 2012) suggesting that either utrophin-dependent or -independent mechanisms have no significant impact on cognitive and behavioural functions (Perronnet *et al.*, 2012). It remains possible however that other, currently unidentified, dystrophin-compensating mechanisms/proteins may exist within the brain to slow down the progression of cognitive deficits in *mdx* mice or other pre-clinical models of DMD. Whether or not the cognitive deficits in people with DMD are progressive in nature has not, to the best of our knowledge, been studied, probably due to the fact that patients with DMD have a relatively short life expectancy of 28.1 years (Landfeldt *et al.*, 2020; Broomfield *et al.*, 2021).

5.7. Conclusions

Having observed that hippocampal LTP was suppressed in untreated *mdx* mice, which would be consistent with dysfunctional learning and memory consolidation processes, we investigated whether or not an intervention procedure could improve these cognitive deficits. As elevated IL-6 is associated with memory impairment (Heyser *et al.*, 1997; Wei *et al.*, 2012; Wei *et al.*, 2013), and we and others (Li *et al.*, 1997; Tancredi *et al.*, 2000) have shown not only that exogenous IL-6 negatively impacts upon LTP in WT mice (Chapter 4), but also that suppression of IL-6 signalling restores normal LTP in *mdx* mice without impacting on LTP in

dystrophin-expressing WT mice, we examined whether centrally-administered, neutralising xIL-6R could improve cognitive outcomes in *mdx* mice. Our results showed that whilst the beneficial effect of xIL-6R on the molecular processes underlying hippocampal learning and memory was convincing at a cellular level, it did not translate into whole animal behavioural changes, with no deficits in recognition memory being observed. However, we hypothesise that this may be due, at least in part, to the relatively young age of the mice used for our study. Some anxiety-like behaviours were displayed by our *mdx* mice; however, these are largely likely to be regulated by other brain regions and were not modified by administration of xIL-6R.

Restoration of hippocampal LTP in *mdx* mice is an important finding and treatment with monoclonal neutralising IL-6R antibodies may be important in preventing future cognitive decline caused by loss of dystrophin.

Chapter 6:

Summary and Conclusions

6.1. Overview of research findings

In this research program, the possible contribution of the neuroimmune molecule, IL-6 in cognitive impairment associated with loss of the structural protein, dystrophin, was investigated. The main cognitive abnormalities seen in DMD patients involve specific deficits in verbal, short-term and working memory (Hinton et al., 2000, 2001; Snow et al., 2013). Using the *mdx* mice model of DMD, which lacks dystrophin, we focused on the hippocampus, as this brain region is critical for learning and the acquisition of new memories.

Little is known about the functional role of dystrophin in non-contracting cells such as neurons. Thus, we started by investigating the localisation of dystrophin in dystrophin-expressing mice. Full-length dystrophin (Dp427) was detected in the cell body and processes of cultured hippocampal neurons. Puncta at the synapses labelled strongly for dystrophin, implicating dystrophin in synapse structure and function. Indeed, others have reported a role for dystrophin in regulating GABA_AR clustering at synapses (Knuesel *et al.*, 1999; Knuesel *et al.*, 2001), thus, a loss dystrophin could be associated with synaptic dysfunction.

Long-term potentiation (LTP), the proposed molecular correlate underpinning the formation and consolidation of new memories, results in structural and molecular changes in hippocampal synapses. This includes activation of NMDA receptors, increased expression and activation of AMPA receptors, calcium-induced activation of downstream signalling molecules and structural changes, such as the formation of new synapses and synaptic remodelling. Given that dystrophin is a structural

protein, it could be involved in both processes. Consistent with this role, *ex vivo* hippocampal slices from dystrophic *mdx* mice displayed reduced LTP in comparison to WT controls. This finding, although different to previous reports (Sesay *et al.*, 1996; Vaillend *et al.*, 2004) is consistent with the cognitive deficits reported in individuals with DMD (Hinton *et al.*, 2000, 2001; Snow *et al.*, 2013). That said, we did not observe differences between *mdx* and WT mice in terms of their recognition memory. However, this may have been due to the age of the mice used, as deficits have been observed in older *mdx* mice (Bagdatlioglu *et al.*, 2020). It is possible that compensatory mechanisms preserve learning behaviours despite pathophysiological changes at the cellular level. Other behavioural issues such as autism, depression, anxiety and ADHD, have been reported in DMD patients and in *mdx* mice (Hendriksen & Vles, 2008; Sekiguchi *et al.*, 2009; Perronnet *et al.*, 2012; Manning *et al.*, 2014). Although behavioural outputs for these were not different in *mdx* mice, we did observe increased freezing behaviours, that are associated with a fear response (Hendriksen & Vles, 2008; Pane *et al.*, 2013a; Banihani *et al.*, 2015). In humans, emotional disturbances have also been directly associated with this loss of dystrophin (Ricotti *et al.*, 2016).

A body of work has examined the progressive nature of DMD in skeletal muscle, where compensatory mechanisms allow near-normal development early in life. However, eventually, when these mechanisms fail, loss of mechanical function is inevitable. Adverse effects of dystrophin loss may progress similarly in cells of the CNS. Indeed, cellular changes were apparent in hippocampi from dystrophic *mdx* mice. We saw a decrease in the density of astrocytes in *mdx* mice in the DG region. This is relevant to hippocampal function as astrocytes release and regulate several

neuroactive molecules that can affect neuronal activity and modulate plasticity and LTP (Bains & Oliet, 2007; Theodosis *et al.*, 2008). Additionally, adult neural stem cells in the subgranular zone of the dentate gyrus are both positively and negatively regulated by contact with astrocytes (Cassé *et al.*, 2018), which is consistent with our observed changes in hippocampal adult neurogenesis, as indicated by decreased doublecortin immunolabelling in *mdx* hippocampi. Neurogenesis plays a critical role in neural plasticity, brain homeostasis and is crucial in preserving the cognitive function and repair of damaged brain cells. Consistent with this function, we also observed an overall decrease in the density of mature neurons in the DG of aged *mdx* mice (8 months old). Obvious deterioration in hippocampal structure was also evident in these aged mice. Fluid-filled structures were present in the granule cell layer of aged *mdx* mice but not in age-matched WT animals. These fluid-filled structures disrupted the cellular morphology of the granule cell layer which, is likely to impact on the neurocircuitry that underpins learning and memory formation.

Although others have linked GABA_A receptor clustering with dystrophin (Knuesel *et al.*, 1999; Knuesel *et al.*, 2001) we did not observe any notable differences in the number of neurons expressing GABA_A receptors in the dentate gyrus of the hippocampus in *mdx* mice relative to controls. Normally, GABA_A R inhibits activity in the postsynaptic cell by functioning as ligand-gated chloride channels, where binding of GABA to the receptor results in an influx of Cl⁻ ions, hyperpolarising the cell. However, that is not to say that GABA_A receptor function is intact in *mdx* mice. Indeed, increases in spontaneous cellular excitability, manifested as increased intracellular calcium activity in neuronal cultures of *mdx* mice, could be due to dysfunction in the regulatory actions of inhibitory GABAergic interneurons. Indeed,

others in the lab have demonstrated that the GABA_A receptor agonist, muscimol, can suppress neuronal hyperexcitability in *mdx* hippocampal cultures.

Figure 6.1. below summarises our observations, combined with other reports (Bagdatlioglu *et al.*, 2020) of the consequences of loss of dystrophin from hippocampal neurons in young, adult and aged *mdx* mice.

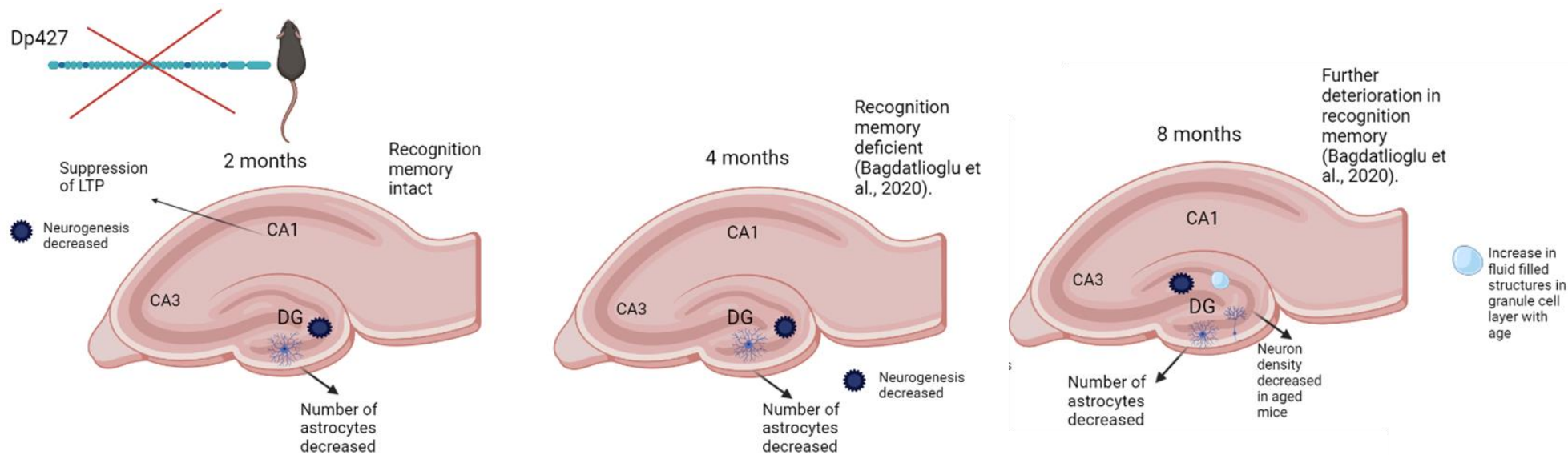


Figure 6.1. Loss of hippocampal dystrophin results in progressive pathophysiological changes.

The schematic illustrates the progressive consequences of the absence of dystrophin in the hippocampus, a brain region key to learning through the formation and consolidation of memories.

- Young mdx mice (2 months) exhibit intact recognition memory with suppressed LTP and reduced astrocytes.*
- Mature adult mdx mice (4 months) exhibit with deficits in recognition memory, decreased astrocytes as well as decreased neurogenesis.*
- Aged mdx mice (8 months) exhibit further deterioration in memory (Bagdatlioglu et al., 2020), decreased neurogenesis, decreased astrocytes and alterations to the cellular layer in the DG due to the presence of fluid-filled structures that result in decrease in neuronal density.*

Having established the morphological and cellular changes associated with the loss of dystrophin in *mdx* hippocampi, we went on to investigate the potential role of the pleiotropic cytokine, IL-6, in the pathophysiological changes occurring in dystrophic hippocampal tissue.

Our interest in IL-6 was piqued by the observed increase in circulating IL-6 both in DMD patients (Rufo *et al.*, 2011) and *mdx* mice (Pelosi *et al.*, 2015), likely reflecting chronic inflammation associated with this disease. This cytokine has been shown to influence LTP (Li *et al.*, 1997; Tancredi *et al.*, 2000; Stampanoni Bassi *et al.*, 2019), neurogenesis (Islam *et al.*, 2009) and astrogliogenesis (Nakanishi *et al.*, 2007), however, IL-6 in the specific context of the brain of *mdx* mice and patients with DMD has not been studied until now.

Somewhat surprisingly, we did not detect any changes in hippocampal IL-6 or IL-6Rs at the gene or protein level. Similarly, expression of the transmembrane protein gp130, which facilitates transcellular signalling by soluble IL-6 receptors, and STAT3, the downstream signalling molecule activated by IL-6Rs was unchanged in *mdx* mice. However, IL-6 can be transported from the periphery across the BBB (Banks *et al.*, 1994) and IL-6 can be released locally in the brain from astrocytes and microglia during pathophysiological events such as inflammation (Gruol & Nelson, 1997) and during physiological processes such as LTP (Balschun *et al.*, 2004) so a lack of global hippocampal differences between WT and *mdx* mice does not necessarily reflect possible local effects. It is also important to note that these results reflect expression levels in the whole hippocampus and are not region-

specific. As described above, the inter-regional neural circuitry is critical to hippocampal function and when we explored regional expression of IL-6 and IL-6Rs using immunofluorescence, regional differences were apparent as there was an increase in IL-6 and IL-6R expression in both the CA1 and DG region of the hippocampus of *mdx* mice, relative to WT. In the CA3 regions, IL-6 expression was similar to WT, whereas IL-6R expression was decreased in the CA3 regions of *mdx* mice. Additionally, we noted in dystrophin-expressing mice, that IL-6Rs were clustered at synapses, similarly to dystrophin on hippocampal processes. Interestingly, exposure to elevated extracellular IL-6 resulted in an increased density of IL-6R-expressing clusters on hippocampal processes. It is not clear if this effect is beneficial or otherwise in terms of cognitive function, or if it is dependent upon the expression of dystrophin, however, it may be important in the observed effects of IL-6 on the generation of LTP in WT and *mdx* mice.

As circulating IL-6 is elevated in the absence of dystrophin and we explored whether elevated levels of IL-6 could affect LTP in dystrophin-expressing hippocampi. Exposure to exogenous IL-6 reduced early LTP in WT animals, although this reduction was not seen in LTP maintenance. This suggested that IL-6 mechanisms could underlie the suppression of the initial kinase activity in LTP. Indeed, this is consistent with broad memory impairment characteristic of transgenic mice overexpressing IL-6 (Heyser *et al.*, 1997; Wei *et al.*, 2012; Wei *et al.*, 2013). In the same transgenic mouse model, a significant reduction in overall neurogenesis in the dentate gyrus of the hippocampus was also found with neural progenitor cells also reduced in the granule cell layer (Vallieres *et al.*, 2002). This is

consistent with our observations that hippocampal adult neurogenesis and the density of mature neurons is decreased the dentate gyrus of *mdx* hippocampi.

However, *mdx* hippocampal slices appeared to be insensitive to the suppressive effect of IL-6 observed in WT mice. This could be due to decreased IL-6R expression in the CA3 region, where the Schaffer collaterals are stimulated. Chronic exposure to elevated IL-6 could have resulted in receptor internalisation or it is possible that reduced LTP observed in *mdx* mice could be due to a complex interplay between local IL-6 levels paired with abnormal neuron functioning due to a loss of synaptic dystrophin. As was noted with aberrant GABA_AR clustering in *mdx* neurons (Knuesel *et al.*, 1999; Knuesel *et al.*, 2001) it is possible that hippocampal IL-6Rs may not be appropriately expressed on the neuronal cell membrane, or, perhaps down-stream signalling is dysfunctional resulting in insensitivity to IL-6. While further research is needed to reveal the precise signalling mechanisms, these studies suggest that while overall IL-6 or receptor levels in the brain are not changed in dystrophic mice, there may be local changes in expression that could modify hippocampal LTP. Increases in IL-6 expression could be due to increases in circulatory IL-6 that has been transported from the periphery across the BBB (Banks *et al.*, 1994) or IL-6 could be released locally in the brain from astrocytes and microglia during pathophysiological events such as inflammation (Gruol & Nelson, 1997), both of which could occur in DMD due to increased peripheral IL-6 (Rufo *et al.*, 2011; Pelosi *et al.*, 2015) and inflammation caused by a loss of dystrophin.

Further support for this suggestion was generated by administering a monoclonal antibody against the IL-6 receptor (xIL-6R) *in vivo* to block IL-6R-evoked signalling in the CNS. Intrathecal administration of xIL-6R resulted in restoration of the suppressed LTP recorded from Shaffer collaterals in treated *mdx* mice. Our results showed that whilst the beneficial effect of xIL-6R on the molecular processes underlying hippocampal learning and memory was convincing at a cellular level, it did not translate into whole animal behavioural changes, with no deficits in recognition memory being observed. However, as discussed above, cognitive deficits were not apparent in this age category (8-12 weeks). It would however, be very interesting to study the effects of xIL-6R in aged mice, where behavioural deficits are apparent (Bagdatlioglu *et al.*, 2020). That said, the progressive loss of hippocampal neurons and glia may mean that the beneficial effects of xIL-6R on LTP in young animals is not effective at later disease stages. Indeed, as is the case with other neurodegenerative diseases when the behavioural deficits become apparent, neuronal deficits are already advanced and resistant to treatment. One small virtue of a DMD diagnosis early in life is that preventative treatments could be employed before cognitive deficits become advanced. Some anxiety-like behaviours were displayed by young *mdx* mice; however, these are largely likely to be regulated by brain regions other than the hippocampus and were not modified by administration of xIL-6R.

We proposed that dysfunctional GABAergic regulation of hippocampal networks could underpin the observed hyperexcitability in hippocampal neurons. It was therefore interesting to observe that acute exposure of *mdx* hippocampal neurons, which had been cultured in IL-6-free media for >7 days, to IL-6 resulted in

suppression, although not abolition, of the spontaneous calcium activity. Given that unregulated calcium can result in neuronal cytotoxicity and cell death (Orrenius & Nicotera, 1994) in this scenario, brief exposure to IL-6 appears to have a beneficial effect.

Figure 6.2 below summarises the findings from this study as well as others work on IL-6 in the brain.

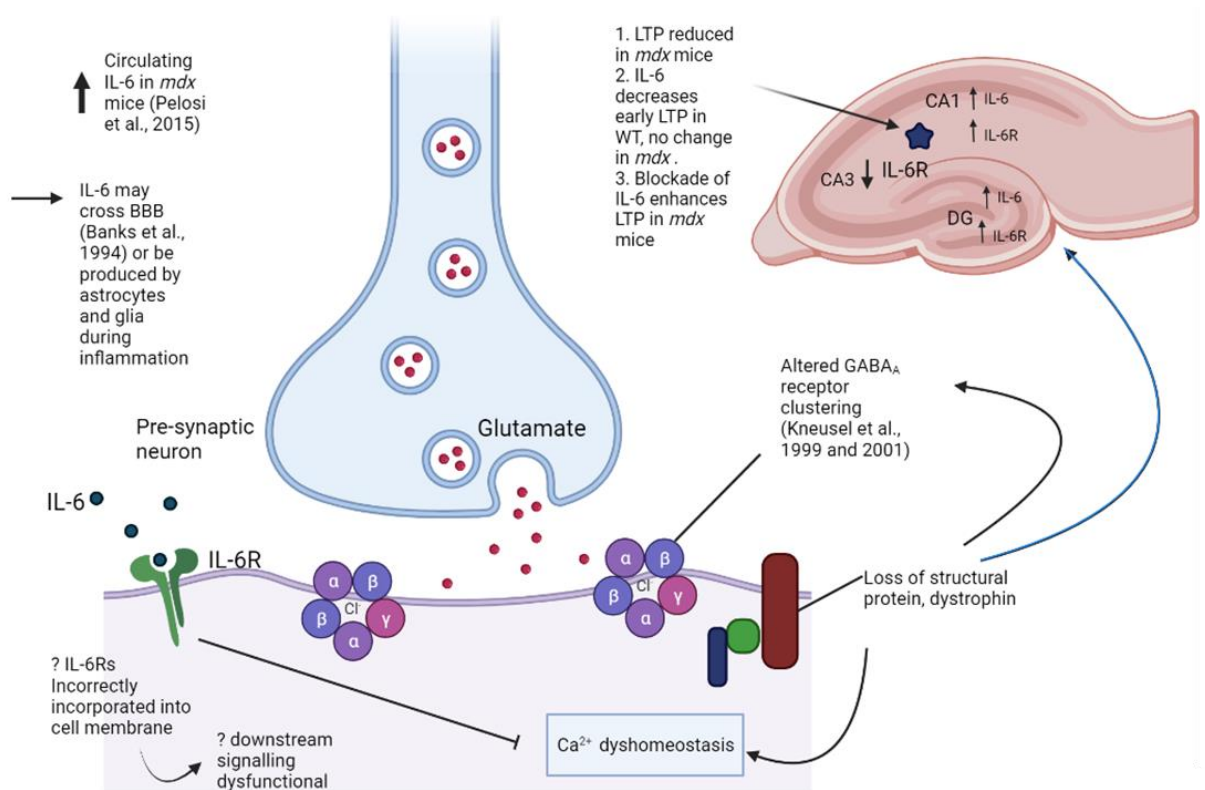


Figure 6.2. The figure illustrates the potential mechanisms of how IL-6Rs could be involved in hippocampal dysfunction associated with a loss of dystrophin.

6.2. Conclusion

This program of work has investigated the importance of dystrophin in hippocampal function and its interaction with the neuroimmune molecule IL-6. Suppression of LTP in *mdx* mice indicates the importance of dystrophin to the molecular correlate of learning and memory. However, in young adult mice this does not yet result in behavioural deficits in learning and memory. We hypothesised that chronically elevated levels of the pro-inflammatory cytokine, IL-6 contributed to dysfunctional LTP, and, indeed, blocking central IL-6 signalling resulted in recovery of LTP, even though baseline levels of IL-6 were not different in dystrophic hippocampal tissue. Impaired hippocampal adult neurogenesis and altered hippocampal cellularity were also observed in *mdx* mice. Others have shown that IL-6 can have similar detrimental effects, however, further studies are needed to determine if IL-6 is fundamental to these changes in the context of dystrophic tissue. Finally, and somewhat counterintuitively, IL-6 exhibited neuroprotective effects when it was applied acutely to cultured hippocampal neurons, where it suppressed neuronal hyperexcitability. However, in the context of physiological homeostasis, this may not be unusual. Acute stimuli often evoke a physiological response to restore baseline function. Generally, detrimental effects are observed when a stimulus is not inhibited through negative feedback, as would be the case with elevated IL-6 caused by chronic inflammation associated with the loss of dystrophin.

In conclusion, loss of dystrophin underlies early cellular and molecular deficits in the hippocampus leading to progressive deterioration in the structure and function of this brain region. However, altered hippocampal sensitivity to chronically

elevated IL-6 also appears to be a contributory factor in the pathophysiological consequences of dystrophin loss, as noted by the restoration of LTP when central IL-6 receptor mediated signalling was inhibited. It is likely that any therapeutic strategies to target IL-6 as a signalling intermediary would need to be applied prior to the presentation of cognitive decline, when neurocircuits may be irreversibly damaged. However, in contrast to other neurodegenerative conditions, DMD is diagnosed early in life. With significant advances in treatment of this disorder and prolongation of the duration and quality of life of individuals with DMD, increased attention should be focussed on understanding the cognitive aspects of the disease.

6.3. Limitations

In this thesis, there were a few limitations that need to be mentioned. Some data-sets were missing in hippocampal adult neurogenesis due to a lack of tissue availability. Further, deficits were not observed in learning at the age we studied (8-12 weeks), and differences could possibly only be seen at an older age, which would be interesting to look at. With regards to LTP, additional experiments on more animals as well as older animals could be beneficial as well. Importantly, where IL-6 was present in LTP experiments, the effect was only seen in the early phase of LTP. After 1 hour, LTP was similar between groups. This could possibly display that IL-6 is only effecting early changes and processes in LTP and further studies would be needed to confirm and explain this.

6.4. Future directions

Given our findings that dystrophin impacts on hippocampal morphology and function, which appears to involve the direct effects of dystrophin loss from neurons, in addition to the indirect actions of the pro-inflammatory cytokine, IL-6, which is elevated in *mdx* mice, there is a possibility that different cell types are having an effect on the hippocampus and the deficits we have seen in the *mdx* mouse with regards to IL-6 and LTP as well as hippocampal adult neurogenesis. One cell type of interest is microglia. Microglia are of particular interest as they are involved in the release of IL-6 as well as involved in the modulation of hippocampal plasticity processes. Thus, there is a necessity to extend LTP and behavioural test analysis to later time points such as 4-8 months and possibly older than a year. More analysis regarding hippocampal adult neurogenesis at older ages would be beneficial as well. With regards to hippocampal adult neurogenesis specifically, as neurogenesis occurs in the dentate gyrus, it could be beneficial to record neuronal plasticity in this region and to perform behavioural tests related with this particular hippocampal region in order to determine if any deficits seen are adding to the cognitive deficits in DMD. Further, GABAergic function in the CA1 could be analysed. Although we did not observe changes in the absolute number of GABA-expressing hippocampal interneurons in dystrophic tissue in the dentate gyrus, aberrant GABA_A receptor clustering could nonetheless have an impact on the neuroregulatory actions of GABAergic neurons. Interestingly, we found that a loss of dystrophin increases neuronal excitability in cultured hippocampal neurons and thus, other regions of the hippocampus, such as CA1 could have altered GABAergic

function. There are a few future directions that this study could take and thus, this work is continuing in the lab and will all be pivotal to establishing a thorough understanding of the functional role of dystrophin in non-contracting neurons and how that feeds through to altered cognitive function in DMD.

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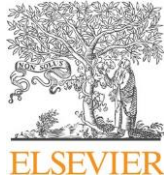
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Appendix



Contents lists available at ScienceDirect

Cytokine

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Review article

Interleukin-6: A neuro-active cytokine contributing to cognitive impairment in Duchenne muscular dystrophy?

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Memory

ABSTRACT

Interleukin-6 (IL-6) is a pro-inflammatory cytokine, classically associated with orchestrating an immune response to pathogens. However, IL-6 can also directly or indirectly modify central nervous system function, and thereby alter cognitive functions, such as learning and the consolidation of memories. IL-6 is chronically elevated in Duchenne Muscular Dystrophy (DMD), a neuromuscular disorder arising when a mutation causes the loss of the structural protein, dystrophin. Dystrophin leads to progressive immobility, chronic inflammation and premature death. However, the role of dystrophin as a mechano-transducing signalling molecule is unnecessary in non-contracting cells such as neurons, where it may be involved in synapse formation. Specific brain regions, including the hippocampus, which is the site of memory acquisition, are affected in DMD. Therefore, loss of this protein may underlie variable deficits in cognitive function that are common in DMD. This review will evaluate the potential role of IL-6 in cognitive dysfunction in dystrophin-deficient DMD.

1. Introduction

Dystrophin is expressed in skeletal, cardiac and smooth muscle and in endocrine and neuronal tissue. Thus, loss of this protein, which underlies the progressive symptoms of Duchenne Muscular Dystrophy (DMD), results in deficiencies in multiple physiological systems. In the central nervous system (CNS), high levels of dystrophin have been detected in neurons of the cerebellum, cerebral cortex and also, the hippocampus [1], which is a region important for learning, and in memory formation. In contrast to its function as a sarcolemma-spanning structural protein, which provides stability to contracting peripheral myocytes, neuronal dystrophin is thought to play a role in the formation of new synaptic connections [2]. Individuals with DMD often exhibit nonprogressive cognitive deficiencies, which vary in severity [3], in addition to difficulties in communication and poor facial affect recognition [4,5]. Deficits in verbal, short-term and working memory, functions that are likely to be facilitated by the hippocampus, have also been observed in individuals with DMD [6–8]. Elevated levels of proinflammatory cytokines,

including TNF α , IL-1 β and IL-6 [9–13] have been detected in DMD muscle biopsies, which is likely to be due to degeneration and regeneration of myofibres. In addition to their well characterized role in the immune response, cytokines can also modify neuronal function, both directly by binding to receptors expressed on neurons [14,15] and indirectly, by stimulating secretion of neuromodulatory molecules from glia or endothelial cells [16,17]. Cross-talk between immune and neuronal cells are key to maintaining normal homeostasis of the nervous system [18]. Thus, chronic inflammation, which is prevalent in individuals with dystrophin-deficient DMD, in addition to structural changes at synapses, could contribute to dysfunctional synaptic transmission (Fig. 1) leading to deficits in cognitive function.

2. Interleukin-6, a pleiotropic cytokine with modulatory actions in the central nervous system (CNS)

Cytokines are molecules of the immune system that are critical for

Abbreviations: AMPA, α -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid; CNS, central nervous system; DAPC, dystrophin-associated protein complex; DMD, Duchenne Muscular Dystrophy; ERK, extracellular-signal-regulated kinase; EPSPs, excitatory post-synaptic potentials; GABA, gamma-aminobutyric acid; IP $_3$ R, inositol – 1,4,5-trisphosphate receptor; IQs, intelligence quotients; IL-6, interleukin-6; IL-6R, IL-6 receptor; JAK/STAT, Janus kinase/signal transducer and activator of transcription; LTP, long-term potentiation; nAChRs, nicotinic acetylcholine receptors; NMDA, N-methyl-D-aspartate; PI 3-kinase, phosphoinositide 3-kinase; (RyR), ryanodine receptor

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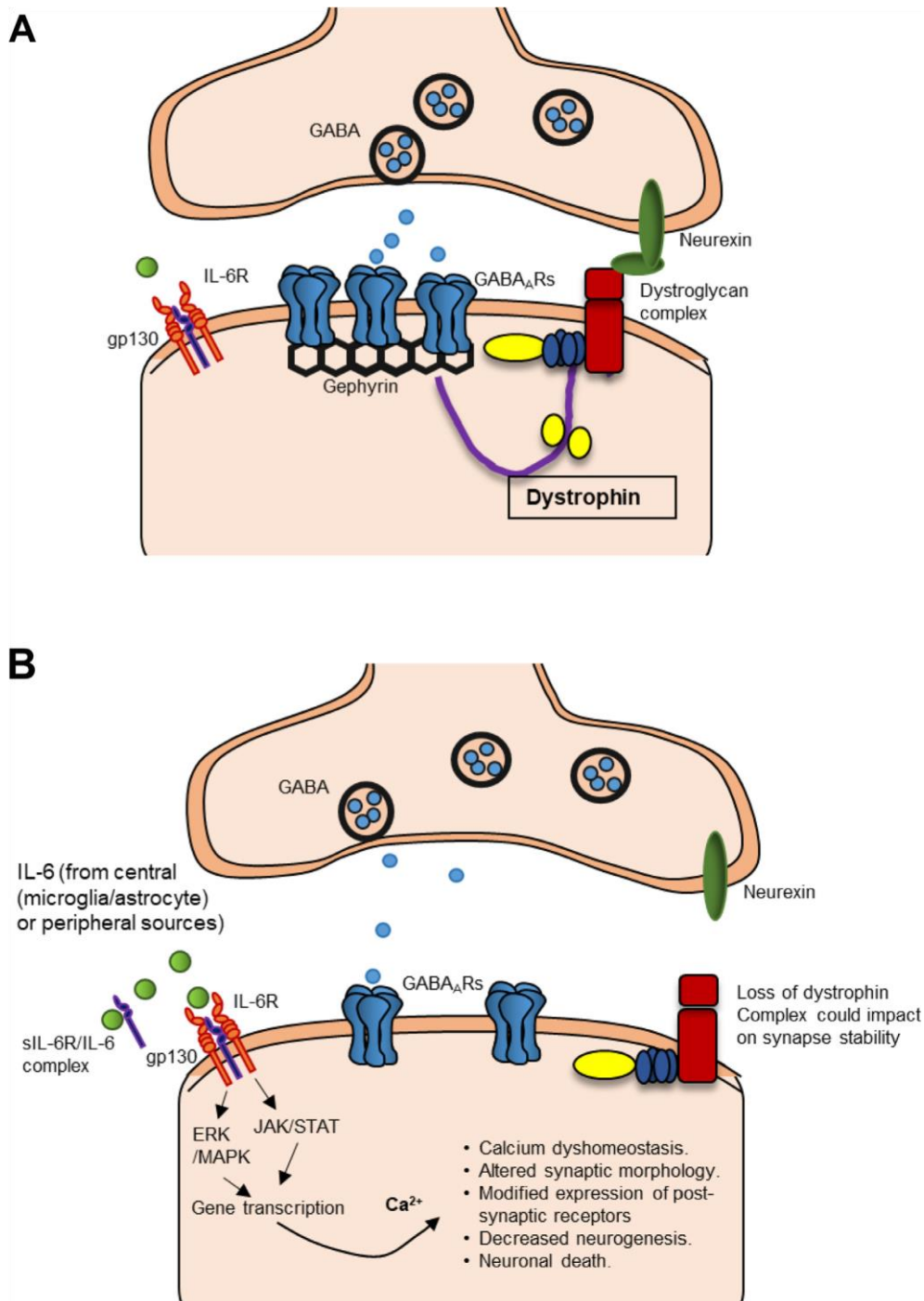


Fig. 1. Mechanisms by which IL-6 could contribute to dysfunction hippocampal neurotransmission in the absence of dystrophin. The schematic compares inhibitory synapses in hippocampal neurons from A: healthy individuals and B: individuals with dystrophin-deficient Duchenne Muscular Dystrophy. Extracellular signal-regulated kinase/Mitogen-activated protein kinase: ERK/MAPK; gamma-aminobutyric acid receptors: GABA_ARs; interleukin-6: IL-6; IL-6 receptor: IL-6R; Janus kinase/signal transducer and activator of transcription: JAK/STAT; sIL-6R soluble IL-6R.

pro- or anti-inflammatory signalling. One such pleiotropic cytokine, interleukin-6 (IL-6) is secreted by both immune and non-immune cells [19], and has diverse effects on many different cell types. In health, plasma levels of IL-6 vary between ~ 1–10 pg/ml [20]. However, this can rise to the lower nanogram/ml range during infection or inflammation [21]. IL-6 has a key role in regulating immune cell function [22], although this 26 kDa protein can also modify function in other cells, including neurons [23]. This cytokine evokes its biological effects by binding to IL-6 receptors, which can either be membrane bound or soluble. Membrane-bound IL-6 receptors (IL-6R α) are only expressed in a restricted subset of cells [24], but this does include many neurons in the brain [25]. Although IL-6R α lacks intrinsic signal transduction capacity, binding of IL-6 to membrane-bound IL-6R α and glycoprotein 130 kDa (gp130) complex results in a plethora of downstream cellular responses. Dimerization of gp130 and subsequent phosphorylation of Janus kinase/signal transducer and activator of transcription (JAK/ STAT), extracellular-signal-regulated kinase (ERK) and Phosphoinositide 3-kinase (PI 3-kinase) signal transduction molecules represents the classical signalling pathways for this cytokine. IL-6 can, however, also trigger signalling in cells which do not express IL-6R α via soluble IL-6 receptors (sIL-6R).

sIL-6R α uses a trans-signalling mechanism where sIL-6R α forms a complex with IL-6 which then binds to gp130 [26,27]. Gp130 is expressed in most cell types including neural cells in the CNS [28]. Although expression of gp130 is not changed by inflammation, the concentrations of sIL-6R can increase between two and five-fold [29,30], thereby increasing trans-membrane signalling. In fact, many cells that respond to IL-6 during inflammation do not express membrane-bound IL-6Rs, but are activated by IL-6/sIL-6R complexes. Furthermore, downstream activation of JAK/STAT, ERK or PI 3-kinase proceeds in a similar manner in cells that don't express membrane-bound IL-6R α as cells that do express IL-6R α [31,32].

Differences in the evoked cellular response between classic receptormediated IL-6 signalling and trans-signalling can however, be detected. IL-6 evokes anti-inflammatory signals using classical signalling, which contrasts with the pro-inflammatory, and potentially deleterious actions of IL-6/sIL-6R-evoked trans-signalling [33]. Furthermore, the kinetics of how these two IL-6 receptor systems signal, also differ, which can lead to the generation of fundamentally different cellular responses [24]. For example, within the CNS, neuronal degeneration evoked by IL-6 is mediated by trans-signalling, whereas membrane receptormediated IL-6 signalling facilitates neural tissue regeneration [34] and neuronal protection [35]. Thus, understanding the specific signalling mechanisms of this and other cytokines is particularly pertinent when therapeutically targeting this signalling pathway for the treatment of neurological disorders.

3. Neuromodulatory actions of IL-6 in the hippocampus.

The hippocampus, a structure within the medial temporal lobe upon which the acquisition of new declarative/spatial memories is contingent, has abundant expression of IL-6R α [15] and indeed, IL-6 induces multiple effects in this brain region. An initial clue to the role that IL-6 may play in modulating cognitive function came from investigating 'sickness behaviour', which describes behavioural symptoms during infection and inflammation. High levels of pro-inflammatory cytokines, including IL-6 and IL-1 β [36], are associated with a deterioration of cognitive function during infection. IL-6 has also been implicated in lipopolysaccharide-evoked cognitive dysfunction [37]. Moreover, transgenic mice over-expressing IL-6 also exhibit broad memory impairment [38–40] and severe neurological pathologies [34]. Further support for the suggestion that IL-6 negatively impacts learning and memory function comes from the finding that transgenic IL-6-deficient mice possess enhanced spatial learning capabilities in the radial arm [41] and Morris water mazes [42] as well as behavioural task-learning assessments. However, it is notable that various forms of learning were also deficient in IL-6 knock-out mice [43], suggesting that basal levels of IL-6 are essential for normal cognitive function.

4. IL-6 modifies synaptic transmission

Ex vivo hippocampal slices retain the hippocampal architecture and intact neural circuitry between the dentate gyrus, CA1 and CA3 areas [44] and as such, can be utilised to study the proposed molecular correlate of memory formation, long-term potentiation (LTP). LTP induction in the hippocampus recapitulates the molecular mechanisms underlying the coding of new memories by modifying the strength of synaptic transmission [45–47]. Using this experimental procedure, a high frequency electrical stimulus induces a long-lasting potentiation of glutamatergic excitatory post-synaptic potentials (EPSPs).

Interestingly, IL-6 gene expression in the CA1 region of the hippocampus is increased after induction of N-methyl-D-aspartate (NMDA) receptor-dependent LTP via stimulation of the Schaffer collateral-CA1 pathway, although it is important to note that expression of IL-6 mRNA was specific to nearby astrocytes, not neurons [48]. Significant increases in the levels of IL-6 mRNA one hour post stimulation were also noted in ex vivo experiments, but only when LTP had been successfully induced [49]. Intriguingly, blockade of endogenous IL-6 using a neutralizing antibody resulted in a remarkable prolongation of LTP, as well as improved performance in spatial memory tests, further emphasising the pleiotropic actions of this cytokine in the CNS [49]. However, it appears that a delicate balance between too little or too much IL-6 is required to maintain optimal memory function, as IL-6 knock-out mice performed less well in hippocampus-dependent memory tests [43] and were more susceptible to hippocampal damage following induced convulsions [50,51], whereas transgenic mice with cerebral overexpression of IL-6 were characterised by suppressed LTP in the perforant pathway dentate gyrus synapses [52].

Although it is recognised that exogenous application of IL-6 may not mimic endogenously released IL-6 during LTP [49], ex vivo application of IL-6 inhibited hippocampal LTP using a mechanism that was dependent upon IL-6R activation and stimulation of the JAK/STAT pathway [53,54]. Moreover, inhibition of LTP by IL-6 was attenuated by administration of monoclonal antibodies, used to block IL-6 receptors [55]. Thus, IL-6 may fine-tune memory consolidation by acting as a molecular filter that

limits the long-term storage of some types of information. Although the mechanistic connections between IL-6-mediated JAK/STAT pathway activation and LTP inhibition remain unclear, work conducted in rat hippocampal neurons suggests that at least one component may be due to an alteration in intracellular neuronal homeostasis of calcium, which is a critical mediator of synaptic plasticity, by increasing its influx through NMDA receptors [54]. This mirrors, to a certain extent, earlier work conducted in rat cerebellar and Purkinje cells which also showed that IL-6 not only raised basal cytosolic calcium concentrations, but also increased flux through NMDA receptors and voltage-gated calcium channels in cerebellar granule cells [56,57], and also enhanced the magnitude of group 1 metabotropic glutamate receptor-mediated responses in Purkinje neurons [58,59].

Neuroinflammatory molecules, including IL-6, also have the capacity to modify synaptic transmission [60] and hippocampal synaptic network activity [61] by modulating γ -aminobutyric acid (GABA)-mediated signalling. GABA is the key inhibitory neurotransmitter in the adult CNS. It can be synthesised and secreted by both neurons and astrocytes and binds to GABA receptors on nearby neurons. Altered distribution of GABA receptors, abnormal subunit composition and aberrant GABA_A receptor-mediated signalling have been detected in disorders such as Huntington's disease, which are characterised by neuroinflammation [62].

5. Cytokines modify neurogenesis

The posterior or dorsal portion of the hippocampus participates in contextual learning, declarative memory and spatial navigation [63]. Within this region, the "input zone" of the classical, tri-synaptic hippocampal circuit, the dentate gyrus region of the hippocampus, which may serve to facilitate translation of cortical signals into coding patterns suitable for memory encoding [64], is one of only two brain regions where neurogenesis occurs and adult-born neurons are integrated into pre-existing neuronal circuits [65]. Indeed, it is this integration of new cells into existing neuronal circuits that is proposed to contribute to hippocampal-dependent learning and memory processes [66]. However,

IL-6 can directly influence neural stem cells during neurogenesis [67], with proliferation, survival and differentiation of hippocampal progenitor cells being compromised in the dentate gyrus when IL-6 concentrations are elevated [68]. Indeed, IL-6 dose-dependently reduces proliferation of neural stem cells, decreases neuronal differentiation and ultimately impacts upon neurogenesis through activation of the JAK2/STAT3 signalling pathway. The importance of JAK2/STAT3 signalling in memory consolidation is highlighted by the fact that cognitive deficits can be rescued in a mouse model of neuroinflammation (APP/PS1) by inhibiting this signalling cascade [69].

Microglia are macrophage-like cells which reside in the CNS and coordinate the brain's inflammatory response. Interestingly, microglia form a particularly dense populations in the subventricular zone [70], which, in addition to the dentate gyrus, is the other brain region where neurogenesis occurs. Activated microglia enhance neurogenesis and oligodendrogenesis in the subventricular zone through their release of cytokines, which include IL-6, IL-1 β , TNF- α and IFN- γ [71]. Others have specifically described the potential role of astrocyte-derived IL-6 in hippocampal neurogenesis [72]. As the number of hippocampal neural stem cells correlates inversely with microglial density, it was proposed that they inhibited adult hippocampal neurogenesis [73]. Indeed, although the contribution of microglia to neurogenesis remains complex, more recent studies have demonstrated that homeostasis of the adult hippocampal neurogenic cascade is dependent upon microglial phagocytosis (Diaz-Aparicio et al., preprint 2019). The prevailing conditions appear to modify both cytokine profile and quantity secreted by microglia, with different cytokines having varying effects on neural stem cells at different concentrations [74].

It is understood that neuro-immune interactions are critical to the maintenance of homeostasis in the nervous system [18], and therefore it is interesting to note that elevated levels of pro-inflammatory cytokines, including IL-6 have been detected in cognitive diseases characterised by neuroinflammation, including Alzheimer's disease [75], Multiple Sclerosis [76], Parkinson's disease [77] and Huntington's disease [78]. The remainder of this review will consider the evidence for IL-6 as a contributory factor in cognitive dysfunction in individuals with DMD.

6. DMD, a multi-system disorder.

DMD is a heritable X chromosome-linked recessive disorder which affects approximately 1 in 3500 live male births, thus ranking it as one of the most prevalent inherited disorders worldwide [79]. Symptoms of DMD are evident in many physiological systems, however, progressive tissue inflammation and loss of function in cardiac and skeletal muscles leading to immobility and eventual cardio-respiratory failure [80] and premature death, have justifiably taken priority in terms of research. The manifestation of the aforementioned symptoms is due to mutations in the DMD gene leading to loss of the structural protein dystrophin [81]. In muscle, dystrophin forms part of the dystrophin-associated protein complex (DAPC) that functions as an essential link in connecting the sarcolemmal cytoskeletal actin to the extracellular matrix [82–84]. As dystrophin stabilises the sarcolemma, myofibres lacking this protein are more susceptible to contraction-induced damage [85]. Repair mechanisms are eventually exhausted and skeletal myofibres are replaced by connective and adipose tissue, which underlies loss of ambulation and respiratory complications [86]. Absence of dystrophin also contributes to cardiac muscle dysfunction, resulting in cardiac myopathies [87] and arrhythmias [88]. Furthermore, expression of dystrophin in smooth muscle underscores symptoms of bowel dysfunction [89] and urologic diagnoses [90] in DMD patients.

Due to improvement in the efficacy of therapeutic interventions [91] including nocturnal ventilation, routine influenza vaccinations, early delivery of antibiotics and physiotherapy [92], life-expectancy for individuals with DMD has increased considerably [93]. Moreover, new gene-editing technologies have made progress in restoring dystrophin to dystrophin-deficient mdx mice [94–97] and human induced pluripotent stem cells [98]. In individuals with DMD, eteplirsen, an oligonucleotide that

facilitates skipping of exon 51 mutations in the dystrophin gene, slows but does not halt disease progression [99]. Given this progress in treating the peripheral symptoms of DMD, it is becoming increasingly important to consider less researched aspects of the disorder, including DMD-associated cognitive impairment [100,101].

7. Cognitive dysfunction in individuals with DMD

Varying degrees of non-progressive intellectual impairment, including lower intelligence quotients (IQs) [3,102], cognitive impairment [103,104], communication difficulties, poor affect recognition [4,5] and specific deficits in verbal, short-term and working memory [6–8], are exhibited by individuals with DMD. Changes in neuronal function in the hippocampus, cerebral cortex, amygdala and cerebellum, where the full length isoform of dystrophin is normally expressed [1,105–108], are likely to contribute to the aetiology of the cognitive impairment and memory deficits seen in patients with DMD. Neuronal loss, neurofibrillary tangles, alterations in dendritic branching, cortical atrophy and metabolic abnormalities have all been reported in the brains of DMD individuals [109–113]. However, not all individuals with DMD exhibit brain abnormalities and such pathological changes do not correlate well with the degree of IQ impairment [103].

8. Cognitive dysfunction in a rodent model of DMD

Understanding the functions of specific proteins in the CNS has been facilitated enormously by the generation of transgenic rodent models where the expression of individual proteins has been selectively deleted or suppressed. Thus, the fortuitous discovery of the dystrophin-deficient mdx mouse [81], which has a nonsense mutation within exon 23 of the dystrophin gene [114], similar to the mutation found in human DMD, has provided an opportunity to understand the mechanistic underpinnings of this disease. As such, mdx mice do indeed recapitulate striated muscle dysfunction and inflammation [115], although the severity of fibrosis and limb muscle dysfunction is substantially less than the human disease [116–118]. Nonetheless, similar to studies of humans with DMD [104,119–121], studies in mdx mice [81,122,123] did not detect gross abnormalities in brain structures. However, ultrastructural architectural changes, as well as reduced dendritic branching and condensed spinal density in corticospinal neurons have been reported in mdx mice [124–126]. The density of hippocampal neurons is decreased in mdx mice [127] and lateral ventricles are enlarged, possibly due to grey matter atrophy [128]. Similar to DMD patients, muscle levels of IL-6, TNF α and IL-1 β are elevated in mdx mice [129–131]. However, changes in circulating or CNS levels of inflammatory mediators in this mouse model have not yet been determined.

Behaviours consistent with hippocampal dysfunction, such as deficits in associative learning and storage of spatial memories and an inability to learn new tasks, have been observed in mdx mice [132–134]. Interestingly, no variance in spatial learning, or the magnitude of LTP stimulated in the dentate gyrus and CA1 regions of the hippocampus, was detected by Sesay and colleagues [135]. In contrast, Vaillend et al. (2004) reported an enhanced LTP magnitude in mdx mice compared to that in wild type mice [132]. Given the known structural function of dystrophin in muscle cells, loss of this protein could lead to disruptions in normal neuronal physiology which could, in turn, alter cellular processes underlying learning and memory. In this vein, it is noteworthy that mdx cortical and hippocampal neurons display significant intracellular calcium dyshomeostasis relative to controls due to both excessive plasmalemmal cation influx and intracellular release through the endoplasmic reticulum-located calcium-release channels, inositol – 1,4,5 – trisphosphate receptors (IP₃Rs) and ryanodine receptors (RyRs) [136]. Significantly, a spatial learning deficit in the mdx mice could be significantly improved using a compound which inhibited the extracellular calcium influx [136]. Changes in long- and short-term memory formation linked to a loss of dystrophin have been associated with altered synapse morphology and plasticity [122,132,133].

9. Altered synaptic receptor clustering in dystrophin-deficient neurons.

The density and expression of several types of receptor is changed in hippocampal neurons from mdx mice. The full-length form of dystrophin is exclusively expressed in dense puncta on the post-synaptic cell membranes of inhibitory neurons [106,137–141], where it is believed to anchor GABA [for review see 142] and other neurotransmitter receptors [143]. Significantly, this clustering is absent in dystrophic membranes, which has implications for the regulation of synaptic excitability. It is notable, however, that normal clustering of GABA_A receptors can be restored by administering specific antisense oligonucleotides, designed to restore functional dystrophin [144]. Moreover, NMDA-dependent LTP can be normalised using the same treatment, thus providing strong support for the suggestion that dystrophin-dependent clustering of GABAergic neurotransmitter receptors at inhibitory synapses is essential for the regulation of normal neuronal excitability and synaptic transmission [145].

A muted response to nicotine in passive avoidance memory tasks, indicated that mdx mice have either altered expression or responsiveness of nicotinic acetylcholine receptors (nAChRs) [146]. Indeed, there are lower levels of hippocampal α 3 nAChR mRNA [147] and α 7 nAChR binding sites in mdx brains [148]. However, such changes, which should result in decreased nicotinic cholinergic synaptic signalling, appear to be mitigated by an upregulation of nAChR-mediated cholinergic transmission in dystrophic animals [149]. Intriguingly, the fact that there is no change in expression of hippocampal muscarinic receptors in dystrophic mice [123] indicates that dystrophin does not exert a generalised effect on the expression of all cholinergic receptor subtypes. However, dystrophic hippocampal neurons did express lower densities of kainic acid / α -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA) receptors, although this could reflect a compensatory response to increased glutamatergic input [122,150]. Combined, these studies demonstrate that a loss of dystrophin from post-synaptic neurons is linked to modified expression of post-synaptic receptors [122,139,150–152], which will inevitably impact upon the neuronal response to afferent stimuli [153] and will also inevitably affect their efferent output. The exact and synchronised neural network activity critical for normal hippocampal function will thereby be disrupted [for review see 154].

10. Pro-inflammatory cytokines could contribute to cognitive dysfunction linked to loss of dystrophin.

The pro-inflammatory characteristics of DMD is due, in part, to the secretion of cytokines from damaged dystrophin-deficient muscle fibres [155,156]. Given that plasma levels of IL-6 are consistently raised in both individuals with DMD [13], and mdx mice [157], and peripheral IL-6 can cross the blood brain barrier using a saturable transport system [158], inflammatory cytokines originating in the periphery could potentially impact upon brain function. As cytokines can modify channels and receptors at both pre- and post-synaptic membranes [159], this could facilitate deficits in cognition linked to a loss of dystrophin. For example, over-expression of IL-6 not only promotes abnormal dendritic spine formation at both inhibitory and excitatory synapses, which mimics the abnormal spines formed by neurons lacking dystrophin, but it is also, as previously discussed, associated with impaired cognitive abilities and deficits in learning [38,40,160–162]. Prenatal exposure to IL-6 in rats leads to elevated hippocampal levels of IL-6, altered expression of NMDA and GABA_A receptors and deficits in spatial learning [163]. Furthermore, as previously mentioned, IL-6 can modify synaptic transmission [60] and hippocampal synaptic network activity [61] by modulating GABA-mediated signalling. Thus, IL-6 is a multifaceted neuroinflammatory molecule which can modify neural function by modulating synaptic transmission and synaptic plasticity, promoting neurite outgrowth, neurogenesis and/or neuronal survival [18,25,164,165].

11. Conclusions

The emergence of treatments, which have extended both the life expectancy and the life quality of individuals with DMD, have highlighted the unmet need to investigate and understand the deficits in cognition, that are common in this dystrophin-deficient neuromuscular disorder. In healthy individuals, dystrophin expression is restricted to specific brain regions, including the hippocampus, a region which is critical for learning and acquisition of declarative memory. Loss of dystrophin from this region is thereby implicated in the presentation of deficiencies in verbal, short-term and working memory, which vary in severity in boys with DMD. Altered density and clustering of postsynaptic receptors, in addition to malformed and malfunctioning hippocampal synapses, have been linked to the loss of dystrophin. These pathological changes are likely to impair the precise and coordinated activation of neural networks underlying learning. As DMD patients and the mdx dystrophin-deficient animal model of DMD both exhibit elevated circulating levels of the pro-inflammatory cytokine, IL-6, we evaluated the potential role that IL-6 plays in the deficits in hippocampal function in dystrophin-deficient individuals. IL-6 has neuromodulatory effects in the CNS, and elevated levels of IL-6 have been linked to cognitive dysfunction and memory impairment. IL-6 has the capacity to modify expression of post-synaptic receptors, a feature which has been described in dystrophin-deficient neurons. Moreover, intracellular calcium homeostasis, a critical mediator of synaptic plasticity, is also modulated by IL-6 and could underlie neuronal dysfunction in the absence of dystrophin. Neurogenesis, where adult-born neurons are incorporated into pre-existing neuronal circuits, is impaired when levels of IL-6 are elevated. In parallel, the density of hippocampal neurons is decreased in mdx mice, which lack dystrophin and have elevated levels of circulating IL-6 [127]. This review has focussed on the possible role of IL-6 in cognitive dysfunction (Fig. 1) associated with the loss of dystrophin. The evidence supports a possible role for this neuromodulatory immune molecule in the pathophysiological deficits in cognition exhibited by individuals with DMD. However, further studies are needed to pin down the precise role IL-6 plays in synaptic transmission in hippocampal neurons that lack dystrophin.

Author contributions

K.A.S. drafted manuscript; M.G.R. and D.O.M. edited and revised the manuscript and approved final version of manuscript.

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Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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