

**ESTROGEN THERAPY IN A VIRAL MURINE MODEL OF
MULTIPLE SCLEROSIS**

A Thesis

by

FRANCISCO PASCUAL GOMEZ

Submitted to the Office of Graduate Studies of
Texas A&M University
in partial fulfillment of the requirements for the degree of

MASTER OF SCIENCE

August 2012

Major Subject: Biomedical Sciences

Estrogen Therapy in a Viral Murine Model of Multiple Sclerosis

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Approved by:

Co-chairs of Committee,	C. Jane R. Welsh Jianrong Li
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ABSTRACT

Estrogen Therapy in a Viral Murine Model of Multiple Sclerosis. (August 2012)

Francisco Pascual Gomez, B.S., Texas A&M University

Co-Chairs of Advisory Committee: Dr. C. Jane R. Welsh
Dr. Jianrong Li

Multiple sclerosis (MS) is an idiopathic neurodegenerative, demyelinating disease of the central nervous system (CNS). MS affects females more than males (3:1) and pregnancy reduces the number of relapses especially during the third trimester when 17- β -estradiol (E_2) and estriol (E_3) are at their highest levels. In order to study the role of estrogens as potential therapeutic agents for MS we investigated their role in Theiler's murine encephalomyelitis virus (TMEV)-induced demyelination (TVID).

SJL female mice were infected intracranially with Theiler's virus or PBS. The mice in the treatment groups were clinically scored and at week 20 they were ovariectomized (OVx) and given a subdermal pellet containing either 1) 0.1mg of E_2 , 2) 5mg of E_3 , or 3) placebo. Four weeks after treatment initiation, the mice were sacrificed and tissue samples were collected and vertebral columns and brains were fixed and placed in paraffin for histological analysis using either hematoxylin and eosin (H&E) stain for general anatomic features or Weil's stain for myelin.

No signs of clinical disease developed in any of the sham-infected mice. Prior to ovariectomy, infected mice had developed significant clinical scores indicative of demyelination. Mice in the placebo and E_3 -treatment groups deteriorated rapidly whereas the E_2 -treated mice improved significantly during the course of the treatment. Uteri were used to assess hormonal effects post-ovariectomy. Hormone treated groups were significantly different from placebo, indicating hormones were present. Hormone treatment showed significant differences among treatment groups for both inflammation and demyelination. E_2 -treatment significantly decreased inflammation compared to placebo and E_3 . E_2 was also effective in reducing demyelination compared to placebo

groups but not E₃. E₃ treatment was effective in reducing inflammation compared to placebo, but no significance was found for demyelination. Both E₃ and E₂ treated mice developed lower antibody levels against TMEV. The improvement in clinical signs, inflammation, demyelination, and the reduction of antibody levels in 17- β -estradiol-treated mice indicate a therapeutic potential for the treatment of MS.

DEDICATION

Para mi mamá, mi papá, y mi hermana. Gracias a su apoyo y su amor estoy donde estoy, son mi todo y este trabajo es prueba de que gracias a ustedes puedo hacer todo lo que me proponga.

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To the members of my committee and my professors for believing in me. Thank you for everything you taught me I take it to heart and I will never forget it.

“In every Triumph there’s a lot of Try”

“Your future depends on many things, but mostly you”

~Frank Tyger

『しゆくふく祝福ときの時は来る、りょうてあ両手上げて』

(“A time for blessings will come, lift up both arms”)

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NOMENCLATURE

MS	Multiple Sclerosis
RRMS	Relapsing-Remitting Multiple Sclerosis
PPMS	Primary Progressive Multiple Sclerosis
SPMS	Secondary Progressive Multiple Sclerosis
CNS	Central Nervous System
BBB	Blood-Brain Barrier
TMEV	Theiler's Murine Encephalomyelitis Virus
TVID	Theiler's Virus Induced Demyelination
L-protein	Leader Protein
EAE	Experimental Autoimmune Encephalomyelitis
MBP	Myelin Basic Protein
PLP	Proteolipid peptide
MOG	Myelin Oligodendrocyte Glycoprotein
pfu	Plaque forming units
OVx	Ovariectomy
E ₂	17- β -estradiol
E ₃	Estriol
ER	Estrogen Receptor
H&E	Hematoxylin and Eosin
RIA	Radio-Immuno Assay
mRNA	Messenger Ribonucleic Acid
Ab	Antibody
ITAbP	Intrathecal antibody production
Ag	Antigen
MHC	Major Histocompatibility Complex
HLA	Human Leukocyte Antigen
ANOVA	Analysis of Variance

DTH	Delayed-Type Hypersensitivity
CD	Cluster of differentiation
T _H	T-helper Cell
T _{CL}	Cytolytic T-cell
T _{Reg}	Regulatory T-cell
nT _{Reg}	Natural Regulatory T-cell
iT _{Reg}	Induced Regulatory T-cell
IL	Interleukin
SCID	Severe Combined Immunodeficient
pg	Picogram (1×10^{-12} grams)
μ g	Microgram (1×10^{-6} grams)
mL	Milliliter (1×10^{-3} liters)
NF- κ B	Nuclear Factor-kappa B
PI3K	Phosphatidylinositol 3-kinase
TNF	Tumor Necrosis Factor
IFN	Interferon
M Φ	Macrophage
ASC	Antibody-secreting Cell
Ig	Immunoglobulin
TGF	Transforming Growth Factor
COX	Cyclooxygenase

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1. INTRODUCTION

Multiple sclerosis, (MS) is a chronic inflammatory, demyelinating, and neurodegenerative disease of the central nervous system (CNS). It can lead to severe disability in young adults, and, according to Gasperini *et al.*, it “is one of the most common causes of neurological disability in this population” (Gasperini *et al.*, 2010). It affects young adults between 15 and 45 years of age, thus making it a high-impact neurological disease because of its effect on the working age group (Owens *et al.*, 2011). It is thought that in MS, different components of the immune system recognize and attack myelin. These include, autoreactive T and B-cells, cytokines, macrophages (MΦ), and antibodies (Ab) directed against myelin. The loss of myelin and axonal damage lead to the characteristic progression of symptoms (Gasperini *et al.*, 2010).

Multiple sclerosis also has more impact on Caucasians and females with a reported female to male ratio of 2:1 and in some places 3:1, and it tends to affect people living in temperate climates (Gold and Voskuhl, 2009). In the United States alone, MS affects more than a quarter million people. MS affects the central nervous system, thus its effects severely affect the individual’s intellect as well as their ability to move and even their sight (Carlson, 2006).

1.1 Etiology

The etiology of MS is thought to be caused by autoimmune damage to myelinated axons followed by degeneration of the axons. As described by Encarta “[It] is chronic, unpredictable, and often progressive” and “[it] attacks and destroys tissues in the brain and spinal cord.”(Carlson, 2006). There are four main types of MS. All forms affect nerve function, resulting in a variety of neurological manifestations. The most common type is the relapsing-remitting MS (RRMS). In this type, the patient has flare-ups and remissions –with periods where the patient does not show symptoms and

This thesis follows the style of the Journal of Neuroimmunology.

periods of pronounced disability. However, these patients do not deteriorate as fast as those with progressive MS. After a period of 10 years or more, 50% of MS patients may exhibit the Secondary-Progressive phase, which is like RRMS, but with increasing deterioration. Another type of MS that is not as common is Primary-Progressive MS (PPMS). In this form the patient just deteriorates steadily from the onset of the disease without any flare-ups and remissions. The rarest form of MS is the Progressive-Relapsing form, which causes the worst symptoms in the shortest period of time. Although patients will have remissions, when the symptoms return they are more severe and the flare-ups experienced become increasingly worse (Carlson, 2006).

There are several schools of thought that address the possible etiology of MS. The most accepted theory is that in MS the immune system has a misdirected response against the myelin sheaths resulting in damage to the axons, which “leads to permanent loss of neurological function” (Gasperini *et al.*, 2010). Myelin acts as an insulator to aid nerve impulses travel at velocities required to accurately relay information. When the immune system disrupts this myelin sheathing, the speed of transmission decreases or is completely interrupted. The neurological effects vary depending on the nerve affected. If the damage is in the pathway supplying the muscles it causes muscular problems, if on a sensory pathway, it may cause the loss of a sense, such as eyesight (Carlson, 2006).

A proposed mechanism for the immunopathogenesis of MS involves activation of autoreactive T-cells by an unknown factor. It has been proposed that molecular mimicry or bystander activation may be one of the factors leading to activation. Once activated, the T-cells can cross the blood-brain barrier (BBB) and possibly lead to its degradation. Once in the CNS, the cells can interact with M ϕ , B-cells, or microglia and lead to the release of pro-inflammatory cytokines. Plasma cells can also migrate into the brain via a disrupted BBB and secrete antibodies against myelin proteins. This causes the degradation of the myelin sheath, recruitment of more inflammatory cells, and the release of free radicals, nitric oxide (NO), and proteases by local microglia. These events eventually culminate in tissue damage and axonal loss leading to the onset of symptoms (Gold and Wolinsky, 2011)

Genetics also plays a role in the susceptibility to MS with an increase in the incidence in patients that have the HLA-DR2 genes of the major histocompatibility complex (MHC) Class II (Gold and Wolinsky, 2011). In addition, the interleukin 2 receptor (*IL-2RA/CD25*) and interleukin 7 receptor (*IL-7RA/CD127*) genes have been identified as possible genetic components for MS (Gold and Wolinsky, 2011; Libbey and Fujinami, 2010; Olsson and Hillert, 2008). These findings give support to the autoimmune hypothesis, as MHC genes are involved in antigen presentation and certain alleles may predispose to reactions against self, or specifically against self-myelin.

Infectious agents, viruses or bacteria, have also been implicated in the pathogenesis of MS. Viruses implicated as “suspect agents” that could trigger MS include: Epstein-Barr virus, *Herpes simplex* virus, *Varicella zooster* virus, Cytomegalovirus, *Herpes hominis* type-6 (Human Herpes Virus 6), measles virus, human T-cell lymphotropic virus, or an endogenous retrovirus known as Human Endogenous Retrovirus (HERV) (Goodin, 2009; Libbey and Fujinami, 2010; Miller *et al.*, 2001; Sato *et al.*, 2011). Other evidence that favors a viral etiology include the following criteria according to Libbey and Fujinami (Libbey and Fujinami, 2010): 1) Characterization of MS lesions demonstrates CD8⁺ T-cells in infiltrates. The CD8⁺ T-cells function in cell-mediated immunity to protect against viral infections; 2) Relapses in MS correlate with increases in viral titers during infection; 3) Antibodies against the various viruses mentioned previously have been detected in MS patients; 4) Peripheral infection can create molecular mimicry that can cause autoimmune reactions. When we consider all these possibilities and the fact that there has not been a clear association between the above viruses and MS patients, it becomes clear that isolating a single causal agent is improbable, especially in the case of mimicry (Libbey and Fujinami, 2010).

Another mechanism, proposed to explain the pathogenesis of MS, hypothesizes that the disease arises as a result of viral infection combined with an overactive immune system which attacks myelin in genetically susceptible individuals. The infectious agent may also mimic host proteins, thus the immune system reacts and attacks both the

pathogen and host tissues (Carlson, 2006). This hypothesis may also explain the relapsing-remitting characteristic progression of the disease as well as other forms of disease progression. Epstein-Barr virus (EBV) infection has been implicated in other autoimmune disorders as well as in MS, including Sjörge's syndrome, rheumatoid arthritis, and systemic lupus erythematosus. In addition, some reports have found viral titers of EBV in MS brains and the diseases mentioned above (Goodin, 2009). Human Herpes Virus type 6 (HHV6) has also been found in MS lesions (Miller *et al.*, 2001). However, there is an important point to be made in that many of the viruses isolated on MS brains have also been found in normal brains or have not been consistently detected in MS brains. Therefore, the etiology of MS may not be due to a single infectious agent but rather a variety of organisms.

A new hypothesis that was proposed by Stadelmann *et al.*, suggests that the symptoms of MS may not only be due to demyelination. The neuro-axonal damage may also be due to a neurodegenerative component. This emerged as a result of evidence from cortical observations of different types of brain lesions in MS patients (Stadelmann *et al.*, 2008). These cortical lesions, according to Calabrese *et al.*, show less inflammation but do have demyelination, axonal loss and neuronal apoptosis, or cell death (Calabrese *et al.*, 2008). Both studies agree that perhaps it is these cortical lesions that cause some of the more severe neurological manifestations of MS (Calabrese *et al.*, 2008; Stadelmann *et al.*, 2008). The Stadelmann study, however, does recommend that these results be retested using more modern imaging techniques in order to obtain better data (Stadelmann *et al.*, 2008).

1.2 Main Basis for This Research Project

Evidence for the autoimmune etiology of MS comes from its preference for the female sex. The most widely recognized autoimmune disorders that shows the strongest female prevalence is Sjörge's syndrome with a 95% incidence rate in females followed closely by systemic lupus erythematosus (SLE) at 90% incidence, and Multiple Sclerosis is in sixth place with an approximately 60% incidence rate (Whitacre, 2001). There is

also a marked difference in how these diseases present in males and females and there is also sexual dimorphism in the immune response. In a recent study by Smith-Bouvier *et al.*, a novel model of mice was used to show if chromosomal gender had an impact in susceptibility to autoimmune disease. The group tested this hypothesis using XX and *Sry*-deficient XY⁻ female mice and XX*Sry* and XY⁻*Sry* male mice that were gonadectomized to examine the effects of chromosome complement without any sex hormones (Smith-Bouvier *et al.*, 2008). They found that XX sex chromosome complement makes SJL mice more susceptible to both experimental autoimmune encephalomyelitis (EAE) and the pristane-induced lupus model (Smith-Bouvier *et al.*, 2008). In addition, the immune response is altered by XX complement, with XY⁻ females producing anti-inflammatory T_H2 cytokines such as IL-13 and IL-5 (Smith-Bouvier *et al.*, 2008). This may help explain why females may be more susceptible to autoimmune diseases, and why they are more often afflicted with MS.

In animal models, females produce more antibodies and their T-cell activation is more robust. However, in human studies the only noticeable difference is that females have an increased absolute number of CD4⁺ lymphocytes but not necessarily increased levels of antibody or cytokines (Whitacre, 2001). Female hormones modulate the immune response and have been shown to dampen the T_H1 response commonly seen in autoimmune disorders during pregnancy (Voskuhl, 2003; Whitacre, 2001). Males seem to have a higher prevalence of primary progressive Multiple Sclerosis (PPMS), especially when they are younger than 30 years of age with the male-to-female ratio of 1:1 (Cottrell *et al.*, 1999). As patients with this rare form of MS age, the ratio begins to switch to the traditional 1:2 seen in relapsing-remitting MS (RRMS) which may indicate a relationship with sex hormones: as female hormones decline with menopause the ratio seems to favor men whose testosterone levels remain into later decades of life (Cottrell *et al.*, 1999). Testosterone has been shown to have a protective role by causing a shift toward a T_H2 (antibody-mediated or humoral immunity) response that reduces the effect of autoimmune disorders, which are primarily T_H1 (cell-mediated immunity) responses, and both testosterone and estrogens have been shown to have neuroprotective roles

(Gold and Voskuhl, 2009). These observations prompted us to further investigate the relationship between sex hormones and the susceptibility of women to MS and the onset and severity of symptoms.

The principal hormones that are addressed by this research project are the estrogens; namely, 17- β -estradiol (E_2) and estriol (E_3). These compounds are similar, with estriol having an extra hydroxyl (-OH) group. However, functionally these hormones perform totally different tasks. E_3 is a hormone that increases to high levels during pregnancy. It is commonly referred to as a hormone of late pregnancy since it is produced by the fetal-placental unit (Sicotte *et al.*, 2002). In contrast, E_2 plays several roles in the body and it can be found in both males and females with females having much higher concentrations. In the Pregnancy in Multiple Sclerosis (PRIMS) European multicenter study, pregnant women with MS were recruited in between January 1993 and July 1995. A total of 254 pregnant women responded to the study and the average relapses were obtained three months before pregnancy, during pregnancy, and three months after. Pregnancy has been shown to decrease the relapse rate especially during the third trimester, coinciding with the highest levels of estrogens and then increases markedly upon delivery as levels of estrogens fall (Abramsky, 1994; Confavreux *et al.*, 1998; Gold and Voskuhl, 2009; Rinta *et al.*, 2010; Stuart and Bergstrom, 2011).

Pregnancy seems to have an effect on the immune system shifting responses from T_{H1} to T_{H2} and in the case of diseases like MS, this setup could explain the marked decrease in the relapse rate (Confavreux *et al.*, 1998). These observations have also been made when comparing pre-menopausal and post-menopausal patients noting that as endogenous estrogen levels wane in post-menopausal patients, the relapse rate increases (Smith and Studd, 1992). Clinical trials reported by Sicotte *et al.* and by Soldan *et al.* have shown estriol treatment in human patients as a promising treatment option for MS and suggests that hormones may indeed be playing a role in disease modulation in patients with MS (Sicotte *et al.*, 2002; Soldan *et al.*, 2003). In addition, hormonal therapy with oral contraceptives (OC) has shown a decrease in the number of relapses (Stuart and Bergstrom, 2011), the severity of symptoms, and the incidence of MS in OC

users was reported as 40% lower if the drugs used contained ethinyl estradiol and a progestagen when compared to non-users (Alonso *et al.*, 2005; Nicot, 2009). Progesterone treatments in nonovariectomized experimental autoimmune encephalomyelitis (EAE) C57Bl/6 mouse models have shown that treatment with 100mg pellets give levels of progesterone comparable to pregnancy (~75ng/mL) and lead to an attenuation of symptoms and increased levels of protective cytokines, such as interleukin-10 (IL-10) (Garay *et al.*, 2007; Yates *et al.*, 2010). However, we will focus solely on the effects of estrogens as a means to observe hormonal effects on a viral model of MS.

1.3 Theiler's Murine Encephalomyelitis Virus (TMEV) Model

Researchers have used two murine – mouse – models in the majority of research related to MS, and each uses different approaches to establish the disease. The first model, which will not be addressed by this project, is the Experimental Autoimmune Encephalomyelitis (EAE) model. This model uses injections of myelin proteins with an adjuvant to cause an autoimmune attack on myelin. This is a purely non-infectious model that focuses exclusively on the autoimmune aspect of MS. To induce EAE, mice are inoculated with a myelin peptide, either Myelin Basic Protein (MBP) or proteolipid peptide (PLP), in complete Freund's Adjuvant (CFA) containing *Mycobacterium tuberculosis*, and induced via injection with pertussis toxin (Bebo *et al.*, 2001).

This project utilizes the Theiler's murine encephalomyelitis virus (TMEV), which causes Theiler's virus-induced demyelination (TVID) in susceptible mice strains. TMEV is a non-enveloped, positive sense, single stranded RNA virus belonging to the genus *Cardiovirus*, in the family *Picornaviridae* (Sato *et al.*, 2011). TMEV has been divided into two subgroups: GDVII and Theiler's original (TO). The Theiler's original strains cause a persistent infection of the CNS in susceptible strains of mice, followed by demyelination whereas the GDVII strains cause lethal encephalitis in all strains of mice. This project will use SJL/J female mice since this strain is highly susceptible to developing a persistent demyelinating infection of the CNS.

The choice of model relates back to the proposed post-viral autoimmune etiology hypothesis. The TMEV model offers several similarities to human MS: First, it is a chronic inflammatory disease of the white matter with gray matter involvement; second, symptoms are caused by myelin breakdown, which is in turn mediated by a T_H1 response (mainly CD4⁺ cells); and third, mice develop autoimmune responses. In addition, disease can be spontaneously induced in mouse colonies because Theiler's is a natural pathogen of mice (Miller *et al.*, 2001). The most virulent groups of Theiler's virus are the FA and GD-VII groups and they cause a fatal disease in mice known as acute polioencephalomyelitis (Oleszak *et al.*, 2004; Roos, 2010). The onset of disease post-infection with either of the less virulent members of the Theiler's Original (TO) group, the Daniels (DA) or the "tissue culture-adapted BeAn strain" is usually characterized by infection of the gray matter, especially motor neurons, leading to a poliomyelitis that may or may not be fatal (Miller *et al.*, 2001; Sato *et al.*, 2011). The mice that survive this period, develop a "chronic, progressive, inflammatory demyelinating disease" that persists primarily in the spinal cord (Miller *et al.*, 2001; Roos, 2010). Like MS, the exact defined mechanism for demyelination induced by TMEV is unclear. Demyelination may arise as a result of either direct lysis of infected oligodendrocytes by the virus or by the immune system, or lysis of uninfected oligodendrocytes by an over-reactive immune system or via bystander responses (Libbey and Fujinami, 2010). Comparable to progressive MS there is no clear gender bias with the Theiler's virus model. However, in other strains such as the C57L/J, the opposite gender bias exists with males being severely affected and females showing more resistance (Fuller *et al.*, 2005; Kappel *et al.*, 1990). All these characteristics make the Theiler's murine encephalomyelitis virus-induced demyelination (TMEV-ID) model, ideal to study the effect of hormones on MS. In this model, H&E stains characteristically show lymphocytic infiltrates in the parenchyma and around blood vessels (perivascular cuffing), meningitis, and status spongiosus. The lack of uptake of the Weil's stain for myelin emphasizes areas of demyelination. The demyelinating lesions seen in TVID are very similar to those seen in human MS. In addition, there was a recent discovery of a

cardiovirus related to TMEV that can infect humans, the Saffold virus (Chastain *et al.*, 2011; Roos, 2010). Thus the TMEV model may prove to be important to determine the mechanisms of MS pathogenesis.

The main goal of this study was to determine the effectiveness of E_2 and E_3 as therapeutic agents for the treatment of (TMEV)-induced demyelination (TVID). We hypothesize that the immune modulatory properties of these two hormones will cause a decrease in clinical signs of disease and also decrease inflammation and consequently demyelination and neurodegeneration in the spinal cords of TMEV infected mice.

2. MATERIALS AND METHODS

2.1 Project Design

The project was divided into two main parts: Development of demyelinating disease as a result of infection and then hormone treatment. Figure 2.1 is a project timeline showing the events that took place from infection to termination. Table 2.1 shows the project design and distribution of mice into groups. Originally eight (8) mice were assigned to each group. During the course of the experiment, two mice were lost in the TMEV-infected placebo treatment group. One mouse from each of the hormone treated uninfected groups died.

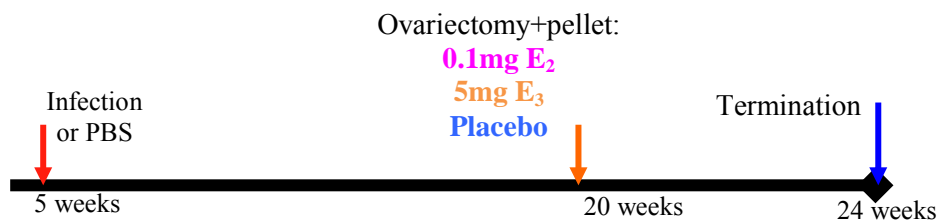


Figure 2.1 Project timeline from infection to termination.

Table 2.1 Project design.

Group distributions for each treatment option

17- β -estradiol (E ₂) (0.1 mg) treatment	Estriol (E ₃) (5 mg) treatment	Placebo treatment (Control)
n=7 - Non-infected (PBS)	n=7 - Non-infected (PBS)	n=8 - Non-infected (PBS)
n=8 - TMEV Infected	n=8 - TMEV Infected	n=6 - TMEV Infected

2.2 Subjects

Groups of four female SJL mice (Jackson Labs) were assigned to twelve different cages ($N_{\text{Total}}=48$). The mice had access to food and water *ad libitum*. For this experiment six different experimental groups were created: three groups were infected with virus at 5 weeks of age and three groups were used as age-matched sham-infected controls. At 20 weeks of age (15 weeks post infection), all mice were ovariectomized (OVx) and then given either: a 0.1mg 17- β -estradiol (E_2) pellet, a 5mg estriol (E_3) pellet, or a placebo pellet placed subdermally (Bebo *et al.*, 2001; Palaszynski *et al.*, 2004; Wang *et al.*, 2009). All animals were housed and handled in accordance with the Texas A&M University Institutional Animal Care and Use Committee (SACC) (AUP# 2007-218, 2007-2010).

2.3 Virus and Infection

For this project, we utilized the BeAn strain of Theiler's virus (obtained from Dr. H.L. Lipton, Department of Neurology, Northwestern University, Chicago, IL). The virus was previously propagated and amplified in BHK-21 cells and the culture supernatant was stored at -70°C (Welsh *et al.*, 1987).

At 5 weeks of age, 24 mice ($n=24$) were anesthetized with isoflurane and infected intracranially into the right cerebral cortex with 5×10^4 plaque forming units/mL (pfu/mL) of the BeAn strain of TMEV. Controls ($n=24$) were given a similar injection with Phosphate Buffered Saline (PBS) intracranial injection into the right cerebral cortex

2.4 Behavioral Measurements – Clinical Symptoms

All mice were evaluated for clinical signs of disease weekly. This project focused on behavioral evaluations during the chronic phase of TMEV infection, paying close attention to trends in improvement after ovariectomy and treatment. We utilized a scoring system based on Borrow *et al.*, 1998 and Sieve *et al.* (2004). The scoring system provides a range of values from 1 to 6. The scores were given as follows

- 1=some weakness in the hind limbs,
- 2=slight wobble,
- 3=wobbly gait,
- 4=pronounced wobbly gait, loss of righting reflex, mice hunch or arch their backs,
- 5=incontinence in addition to symptoms in 4, and
- 6=moribund mice (Sieve *et al.*, 2004).

The data was obtained, averaged and plotted as a score vs time chart using Microsoft Excel.

2.5 Tissue Samples and Preparation

All mice were euthanized at 19 weeks p.i., which represents the 24th week of the experiment. The mice were perfused via the left ventricle with PBS followed by 10% formalin in phosphate buffer pH 7.2. The uteri, spleens, brains, and vertebral columns were collected. Uteri were weighed to assess exogenous hormonal effects on the mice and discarded after their weight was recorded. The spleens were flash frozen in liquid nitrogen and stored at -80°C. The brains and spinal cords were dehydrated and embedded in paraffin blocks. They were stored at 4°C until they were sectioned and stained (Campbell *et al.*, 2001).

The present study focused on lesions in the spinal cord since previous research has shown that the spinal cord is regularly affected during the chronic phase of the TMEV infection (Blakemore *et al.*, 1988; Young *et al.*, 2010). Vertebral columns containing intact spinal cord were removed and sectioned transversely into 12 pieces. Spinal cords were serial sectioned on a microtome at either 5µm for Hematoxylin and Eosin (H&E) staining or at 10µm for Weil's staining. They were mounted on individual slides containing 12 sections per animal and covering cervical, thoracic, lumbar, and sacral/*conus medullaris* divisions of the spinal cord.

2.6 Light Microscopy and Image Captures

An Olympus VANOX AHBS3 microscope with an adapted digital camera was used for all image captures. Hematoxylin and Eosin (H&E) slides were captured at 40x magnification using a SPOT Insight QE digital camera (Diagnostic Instruments, Inc). Due to a software update and the purchase of a new camera, the Weil's-stained slides were captured at 40x magnification using an SPOT RT Slider digital camera (Diagnostic Instruments, Inc). Both cameras had a 1.0x magnification. The software used for image capturing was SPOT Software version 4.6 (Imaging Solutions division of Diagnostic Instruments, Inc.)

2.7 Hematoxylin and Eosin (H&E) Staining and Lesion Evaluation

Sections of the paraffin-embedded vertebral column were cut at 5 μ m and then stained with H&E by Lin Bustamante at the Veterinary Integrative Biosciences (VIBS) Histotechnology Lab according to their established Hematoxylin and Eosin (H&E) staining protocol presented here:

The sections were deparaffinized with three series of 15-20 dips each in Pro-Par Clearant and rehydrated in two 10-15 dips in 100% ethanol (EtOH). The sections were then dipped for two series of 10-15 times each in 95% EtOH and then in one series of 10-15 dips in 70% EtOH. The sections were washed three times with distilled water. The sections were stained in Hematoxylin 560 (Surgipath cat # 01571/Leica Microsystems cat # 3801571) for 4 minutes. Afterwards they were washed in tap water until clear. The slides then underwent a series of 2-3 quick dips in a solution made up of 1% hydrochloric acid (HCl) and 70% EtOH (0.5% acid alcohol) to differentiate them. The sections were rinsed with tap water three times and were dipped 2-3 times in a 0.5% lithium carbonate (Li₂CO₃) solution (1.54g Li₂CO₃ in 100mL distilled H₂O) to give them a blue tint. Once the sections are stained blue, they are washed in running tap water four times and rinsed with 10-15dips in 80% EtOH. The sections can then be counterstained in 0.5% Alcoholic Eosin Y (StatLab, cat# SL98-1) in a series of 1-10 dips. The

sections are dehydrated using two series of 3-5 dips in 95% EtOH and three series of 10-20 dips in absolute EtOH. The stain was finalized by clearing using 10-15 dips in xylene and then four series of 10-15 dips in Pro-Par Clearant. The slides were then mounted and cover-slipped.

The TMEV model is characterized by inflammatory lesions of the spinal cord, and previous studies have shown restraint stress alters the immune systems response to TMEV in these areas using a categorical system of measurement based on lesion expression (Campbell *et al.*, 2001; Sieve *et al.*, 2006; Sieve *et al.*, 2004). This project utilized area measurements to determine percent inflammation in the different spinal cord segments evaluated making the results more quantitative and facilitating interpretation (Campbell *et al.*, 2001; Meagher *et al.*, 2007; Sieve *et al.*, 2004).

Independent raters blind to the subjects' treatment and condition were assigned different groups of mice and were asked to measure perivascular cuffing (perivascular accumulation of lymphocytes and macrophages) using ImageJ Software (National Institutes of Health). *Status spongiosus* (where possible demyelination could occur) were noted on the sections. All divisions of the spinal cord were measured on the slides – with the exception of the *conus medullaris* – at 40x magnification.

The three measurements obtained from each section were: Total cord area, gray matter area, and area of inflammation (sum of areas, including cuffing and parenchymal) if applicable. White matter area was obtained using the equation: [Total cord area (pix²)]-[Gray matter area (pix²)]. Percent white matter inflammation was obtained by the equation: [Area of inflammation (pix²)]/[Total white matter area (pix²)]x100. The author supervised the raters and their data was checked for accuracy. Data was recorded on Microsoft Excel software. Analyses of the inflammatory lesions of the spinal cord were conducted for comparison of the percent parenchymal inflammation among groups.

2.8 Weil's Staining and Lesion Evaluation

Demyelination was evaluated following staining of serial sections with the Weil's method for myelin. The Weil's stain uses iron alum and hematoxylin to visualize areas

with diminished myelin, or lipid-rich contents (Young *et al.*, 2010). Sectioning (10 μ m) and staining were done by Lin Bustamante at the VIBS Histotechnology lab according to a modified protocol based on Arthur Weil's original method (Weil, 1928):

A stock solution of Weil's solution is prepared by mixing 45mL of 4% ferric alum ($\text{FeNH}_4(\text{SO}_4)_2 \cdot 12\text{H}_2\text{O}$) with 5mL of 1% stock hematoxylin solution. A solution of Weigert's Borax ferricyanide solution is prepared by mixing 2g Borax (Sodium borate [$\text{Na}_2\text{B}_4\text{O}_7 \cdot 10\text{H}_2\text{O}$]), 2.5g potassium ferricyanate ($\text{K}_3\text{Fe}(\text{CN})_6$) in 200mL of distilled H_2O . A 0.5% solution of Lithium carbonate (Li_2CO_3) is prepared by mixing 0.5g Li_2CO_3 with 100mL distilled H_2O .

The sections were deparaffinized in absolute EtOH and placed in 0.5% celloidin for 10 minutes to prevent detachment from the slide, and were left to air dry for 15 minutes. The slides were then placed in 70% EtOH for 5 minutes and then washed with distilled H_2O . The slides were then placed in prepared Weil's solution for 10-45 minutes at 50-60°C. After that period of time, the slides are washed with tap water. Using the microscope, the slides are placed back in the Weil's solution and checked for differentiation of white matter or areas of degeneration. Once the desired level is reached, the slides are washed with distilled H_2O (several changes). To complete the differentiation, the slides are placed in the Weigert's borax ferricyanide solution and checked under the microscope. Once the white matter/lesion areas are differentiated, the slides are again washed in several changes of distilled H_2O , and then in tap water. The slides are placed in the Li_2CO_3 solution for one minute and then rinsed in sequence with tap and distilled H_2O respectively. The slides are then dehydrated in EtOH, cleared in xylene and mounted and cover-slipped.

Independent raters blind to the subjects' treatment and condition used ImageJ Software (National Institutes of Health) to separately identify and outline areas of demyelination for each group assigned. All divisions of the spinal cord were measured on the slides – with the exception of the *conus medullaris* – at 40x magnification.

Once again, the three measurements obtained from each section were: Total cord area, gray matter area, and area of demyelination (sum of areas) if applicable. Each measurement was repeated twice and the values averaged. Average White matter area was obtained using the equation: [Average Total cord area (pix²)]-[Average Gray matter area (pix²)]. Percent white matter demyelination was obtained by the equation: [Average Area of demyelination (pix²)]/[Average Total white matter area (pix²)]x100. The average values were used for all analyses of the demyelinating lesions of the spinal cord. Statistical analyses were conducted for the percent parenchymal demyelination. Data was recorded on Microsoft Excel software.

2.9 Theiler's Virus Antibody (Ab) Levels

Radio-Immuno Assays (RIAs) using Theiler's virus (GD-VII) as antigen (Ag) were performed using the sera collected from each mouse at termination according to the protocol described by Sieve *et al.* (2004). Flexible U-shaped, 96-well polyvinyl chloride plates (Costar, Cambridge, MA) were rinsed five times with Tween 20 (0.05%v/v) in Reverse Osmosis water (RO-H₂O) and then rinsed five times with RO-H₂O. The wells were then blot dried and 1µg/well of the Theiler's virus antigen (GD-VII, 1.0x10⁻⁷ pfu/100µL) in carbonate buffer (pH=9.6) was added to coat the bottom of the wells. To block the plate, 100µL of a buffer made of 495mL of a solution made from 0.08M Trizma HCl, 0.03M Trizma base, and 0.15M NaCl (final pH of 7.2), and 5mL of a solution made from 1.0% nonfat dry milk (NFDM) and 0.5% Tween-20 in RO-H₂O. The plates were incubated for 24h at 4°C. The plates were washed with Tween 20 (0.05%v/v) in Reverse Osmosis water (RO-H₂O) and rinsed with RO-H₂O once again and blocked with 3.0% NFDM in phosphate buffered saline (PBS, pH 9.0) using 200µL/well. The plates were incubated at 37°C for 1h. After washing and blot drying, 10µL of mouse sera were added to the plates and diluted in 190µL of PBS containing 0.1% Bovine Serum Albumin (BSA) and NaN₃ to create a 1/20 dilution. Serial dilutions were then made by taking 100µL of the 1/20 dilution, adding it to 100µL of PBS and mixing. The dilutions made were 1/40, 1/80, 1/160, 1/320, and 1/640. At the end, 100µL of the 1/640 dilution

were discarded to keep a 100 μ L volume per well. The plates were then incubated for 1h at 37°C and washed as described above. Following the wash, 100 μ L of rabbit anti-mouse IgG diluted 1/500 from stock (Accurate Chemical and Scientific, New York) were added to each well. The incubation and washing was repeated, and then 100 μ L of Protein-A-I¹²⁵ (1x10⁵ counts per minute [CPM]) were added to the wells and incubated at room temperature for 1h. After rinsing, the wells were cut and placed in tubes and radioactive counts read using a micromedic 4/200 plus automatic gamma counter (Sieve *et al.*, 2004).

The results were transposed to Microsoft Excel and CPM to dilution charts were made for uninfected and infected mice and the respective treatment groups.

2.10 Statistical Analysis

The data is presented as Mean \pm SEM. Data for behavioral/clinical scores was analyzed with Prism Software (GraphPad Software, Inc.) using the non-parametric Kruskal-Wallis test with Dunn's Multiple comparison *post hoc* test. Data for uterine weights, spleen weights, and radio-immuno assays (RIAs) was analyzed with SPSS Statistics 20 (IBM) using a Two-way Analysis of Variance (2-way ANOVA) with a Least Squares Difference (LSD) *post hoc* test. Data for percent inflammation and percent demyelination, and comparison of the infected treatment groups in the RIAs was analyzed using a One-Way Analysis of Variance (1-way ANOVA) with a Least Squares Difference (LSD) *post hoc* test. For all data analyzed, statistical significance was achieved if $p < 0.05$.

3. RESULTS

3.1 Estradiol (E_2)-treatment Ameliorates Clinical Signs of Disease

Mice were evaluated once weekly for signs of disease and data is shown for scores beginning at week 17 post-infection (pi) in order to evaluate the chronic phase of TMEV infection. Scores for the weeks post treatment (Weeks 21-24) were averaged and evaluated for an effect of treatment on clinical symptoms of disease. Table 3.1 shows the mean weekly scores per treatment group and the overall mean score for the four-week treatment period. The overall mean clinical score for the placebo treated mice was 3.8616 ± 0.1057 ; whereas E_2 treated mice scored 3.3750 ± 0.0846 and for E_3 treated mice 3.6719 ± 0.0771 . Figure 3.1 is a graphical representation of the mean scores. Non-parametric Kruskal-Wallis analysis showed a significant difference among the all three groups for the mean weekly scores ($H_{(2)}: 6.865, p < 0.05$). The scores began to diverge and become significantly different among the three groups at weeks 23 ($H_{(2)}: 7.337, p < 0.05$) and 24 ($H_{(2)}: 10.39, p < 0.05$). The Dunn's Multiple comparison *post hoc* test showed that E_2 treatment significantly decreased clinical scores compared to placebo treatments at ameliorating scores post ovariectomy ($p < 0.05$) but scores were not different from E_3 -treatment groups ($p > 0.05$). There was no difference between placebo-treated groups and E_3 -treated groups ($p > 0.05$). Regression lines were fitted to each plot revealed that both placebo and E_3 -treated groups tend to increase in score severity as indicated by their positive slopes (m), while the E_2 -treated group seems to ameliorate based on its negative slope (m); however, the R^2 -value ($R^2=0.082$) for the E_2 regression line is not high enough to claim this effect.

Table 3.1 Mean weekly clinical scores.

The mean clinical scores per experimental group are presented in this table in addition to the overall averages for the four-week treatment period

	Week	Placebo Tx n=6	E₂ Tx n=8	E₃ Tx n=8
Avg	21	3.7500	3.3125	3.5938
SEM		0.0945	0.1619	0.2001
Avg	22	3.6429	3.6250	3.5000
SEM		0.1429	0.1250	0.1637
Avg	23	3.9286	3.2500	3.7500
SEM		0.1304	0.1336	0.2113
Avg	24 (term.)	4.1250	3.3125	3.8438
SEM		0.1548	0.1315	0.1151
	Overall Average	3.8616	3.3750	3.6719
	SEM	0.1057	0.0846	0.0771

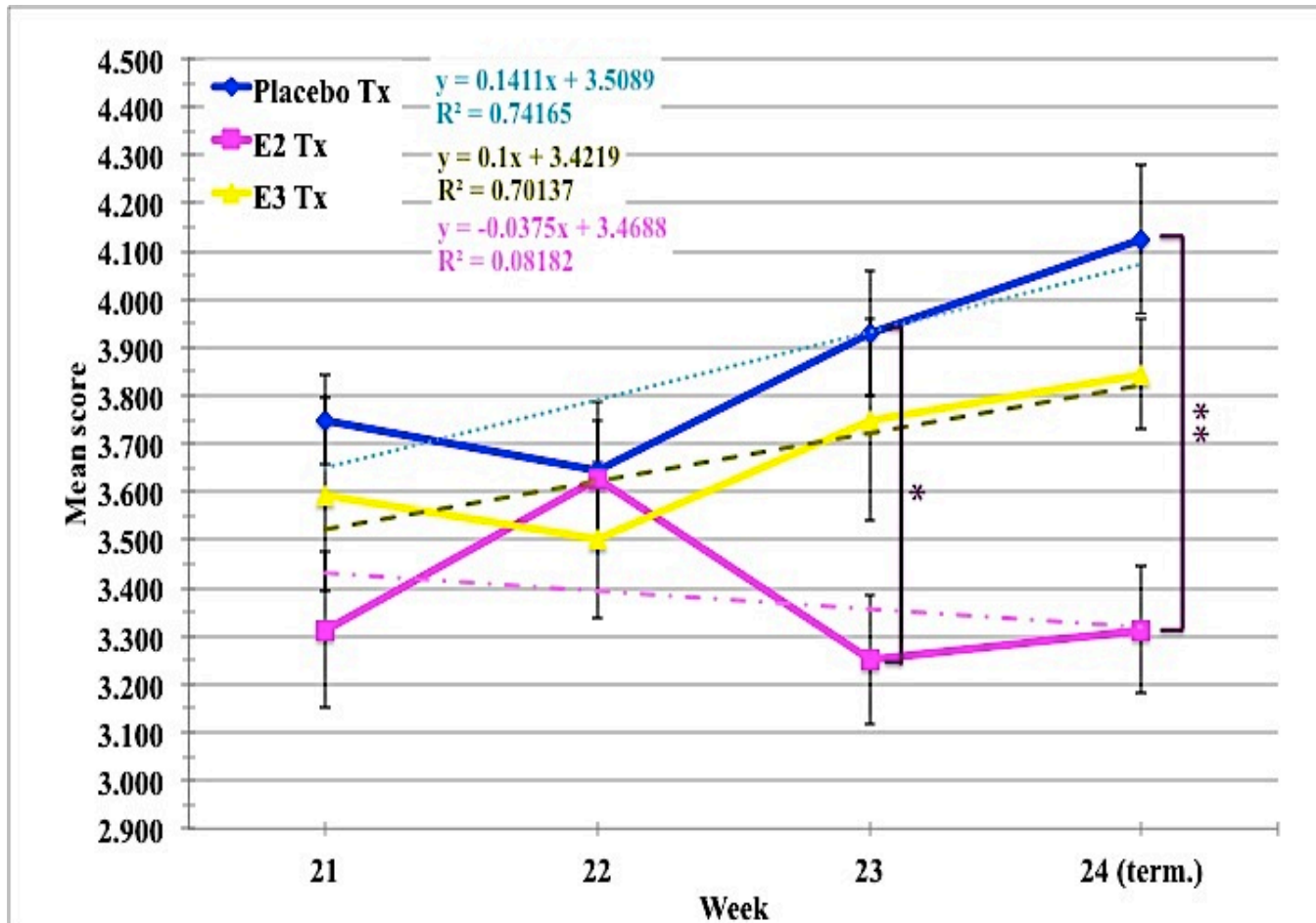


Figure 3.1 E_2 -treatment ameliorates the mean clinical scores when compared to placebo treatment. Scores following ovariectomy were significantly different among treatment groups ($p < 0.05$). Hormone treatment with E_2 began showing a marked effect compared to placebo.

3.1.1 Before-and-After Clinical Score Comparison Showed Both E₂ and E₃ Treatment Groups Exhibit Similar Rates of Disease Progression, but E₂-treatment Maintained Lower Clinical Scores

Table 3.2 shows the mean before and after scores per group comparing week 17, when we began measuring scores, to week 24 (termination). Figure 3.2 depicts this data graphically for the different treatment groups between week 17 and week 24 post-infection. Regression lines fitted to the plots showed that all scores had a tendency to increase. Treatment with E₂ had the same increase in score as E₃ as indicated by the slope (m) of their respective regression lines ($m=0.5938$) and placebo-treatment showed the steepest increase in score ($m=1.1875$). Non-parametric Kruskal-Wallis analysis was used to compare the three groups at week 17 and week 24, respectively. No significant difference was found among groups during week 17 ($H_{(2)}: 3.572; p>0.05$); however, the group scores were significantly different at termination ($H_{(2)}: 10.39; p<0.05$). Though E₂-treatment increased at the same rate as E₃, the range of scores remained lower than E₃. The final score of E₂ treatment was significantly different from placebo ($p<0.05$) but not from E₃-treated groups. There was no significant difference between E₂ and E₃ treatment groups ($p>0.05$).

Table 3.2 Mean clinical scores from week 17 and week 24.

Mean clinical scores for both endpoints per group with the standard error are presented in this table. The average score per group is also shown

	Week	Placebo n=6	E₂-Tx n=8	E₃-Tx n=8
Avg	17	2.93750	2.71875	3.25000
SEM		0.14752	0.26490	0.16366
Avg	24 (term.)	4.12500	3.31250	3.84375
SEM		0.15478	0.13153	0.11512
Average		3.53125	3.01563	3.54688
SEM		0.59375	0.29688	0.29688

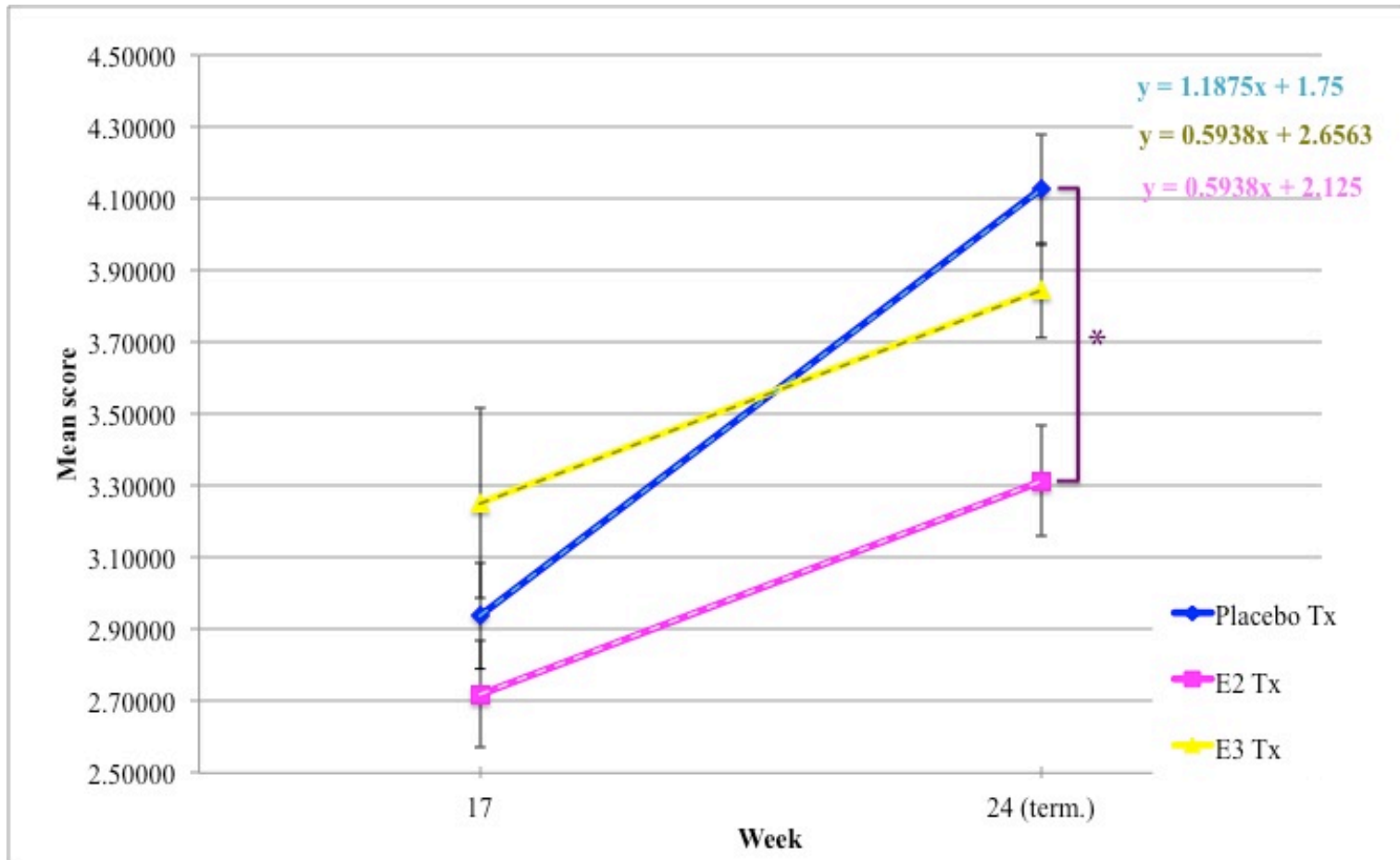


Figure 3.2 Mean clinical scores from weeks 17 and 24 show an increase in severity over the observation period. No significant difference was found among groups during week 17 ($p > 0.05$). Group scores were significantly different at termination (week 24) ($p < 0.05$). Though E₂-treatment and E₃-treatments both had the same slope, E₂-treatment group scores remained lower than both E₃ and Placebo-treated group scores.

3.2 Hormone Replacement Treatment Post-ovariectomy Causes Uteri to Increase in Size and Weight

The mean uterine weights for both the uninfected and infected groups and hormone treatment subgroups are presented in Table 3.3. The uteri were collected and weighed after sacrificing the mice on week 24 of the experiment. The uteri were used to assess the effect of the hormones on the mice post-ovariectomy. Figure 3.3 presents the mean weight of the uteri graphically. A Two-way Analysis of Variance (ANOVA) was done to evaluate the difference between both uninfected and infected mice and their respective hormone treatment subgroups. There was a very significant difference in uterine weights among groups ($F_{(5,43)} : 68.098, p < 0.05$) that was influenced by hormone treatment ($F_{(2,43)} : 166.696, p < 0.05$) but not infection with virus ($F_{(1,43)} : 0.026, p > 0.05$). There was no interaction between virus and hormone ($F_{(2,43)} : 1.186, p > 0.05$). Further analysis, using Least Squares Difference (LSD) *post hoc* test showed no significant difference in uterine weights between E₂ and E₃-treated mice in either group ($p > 0.05$), but a significant difference between these groups and placebo-treated mice ($p < 0.05$) in both infected and uninfected mice.

Table 3.3 Mean uterine weights in grams.

Mean uterine weights for each group are presented along with the standard error. The top row represents uninfected groups and the bottom row represents infected groups

	E ₂ +PBS n=7	E ₃ +PBS n=7	Placebo+PBS n=8
Avg	0.14614	0.14943	0.04288
SEM	0.00896	0.00803	0.00505
	E ₂ +Virus n=8	E ₃ +Virus n=8	Placebo+Virus n=6
Avg	0.15863	0.14250	0.03943
SEM	0.00821	0.00389	0.00290

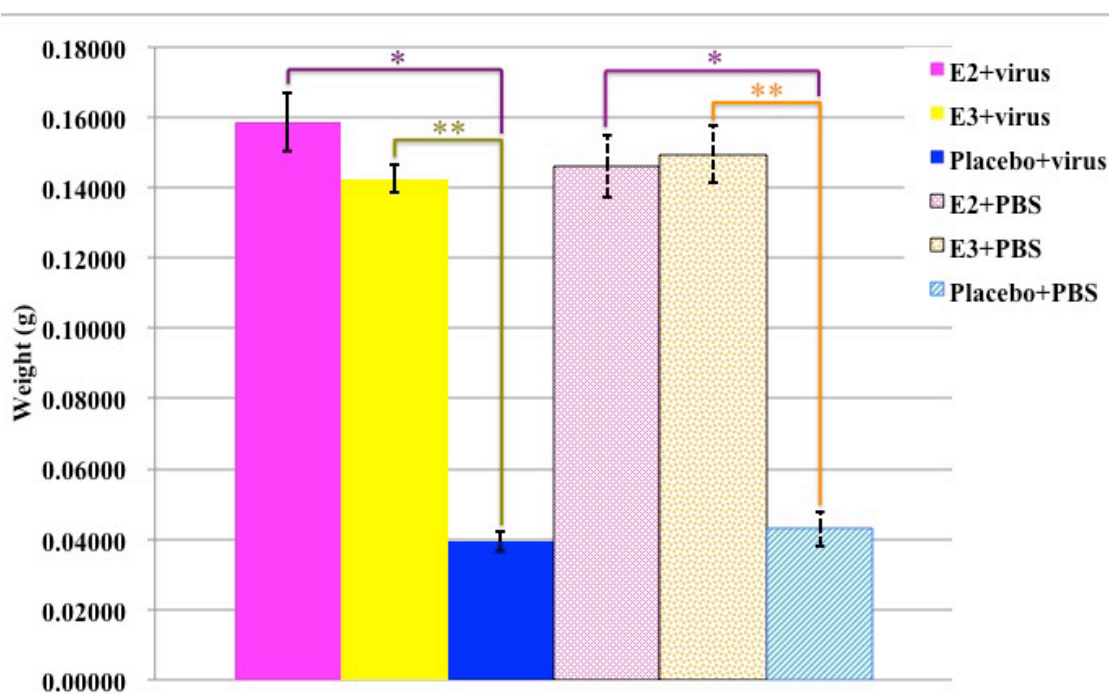


Figure 3.3 Graphical representation of the mean uterine weights obtained at termination. There was a significant difference among groups ($p < 0.05$) with a main effect of hormone treatment ($p < 0.05$). There was a significant difference between E₂ and placebo (*) as well as between E₃ and placebo groups in mean uterine weight (**) in both infected and uninfected groups. No significant difference ($p > 0.05$) was found between E₂ and E₃-treated groups in either infected or uninfected groups. Data shown as Mean \pm SEM. * $p < 0.05$, ** $p < 0.05$

3.3 Infection with Theiler's Virus Has No Effect on Spleen Weights

The spleens were collected and weighed after sacrificing the mice on week 24. Their weights were used to assess immune activation. Table 3.4 presents the mean weights and standard errors for the uninfected and infected groups and their hormone treatment subgroups, respectively. The three infected groups seemed to have smaller spleens than the uninfected groups, however, this did not reach significance. Figure 3.4 presents the mean weight of the spleens graphically. A Two-way Analysis of Variance (ANOVA) was done to evaluate the difference between both uninfected and infected mice and their respective hormone treatment subgroups. There was no significant difference in spleen weights among groups ($F_{(5,43)}: 1.599, p > 0.05$). Infection with virus had an effect on spleen weights ($F_{(1,43)}: 7.912, p < 0.05$); however, hormone treatment

had no effect ($F_{(2,43)} : 0.004, p > 0.05$). There was no interaction between virus and hormone ($F_{(2,43)} : 0.012, p > 0.05$).

Table 3.4 Mean splenic weights in grams.

Mean splenic weights for each group are presented along with the standard error. The top row represents uninfected groups and the bottom row represents infected groups

	E ₂ +PBS n=7	E ₃ +PBS n=7	Placebo+PBS n=8
Avg	0.13271	0.13200	0.13175
SEM	0.00984	0.00581	0.00467
	E ₂ +Virus n=8	E ₃ +Virus n=8	Placebo+Virus n=6
Avg	0.11613	0.11463	0.11500
SEM	0.00873	0.00729	0.00572

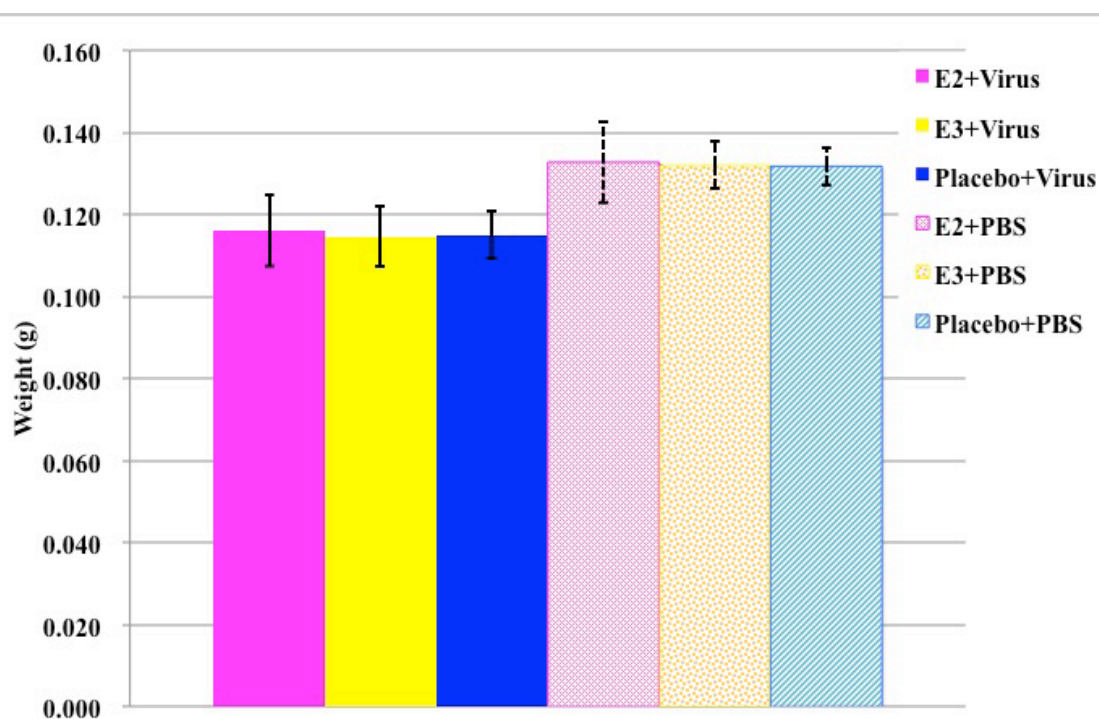


Figure 3.4 Graphical representation of mean splenic weights obtained at termination.

There was no significant difference among groups ($p > 0.05$) despite having a main effect of infection with virus ($p < 0.05$). Data shown as Mean \pm SEM.

3.4 Treatment with E₂ and E₃ Diminishes the Percent Inflammation in the Spinal Cord Sections Compared to Placebo Treatment

As previously mentioned, inflammatory lesions of the spinal cord characterize the Theiler's Murine Encephalomyelitis Virus (TMEV) model. Area measurements were used to determine percent inflammation in the different spinal cord segments. We measured perivascular cuffing (accumulation of leukocytes around blood vessels) at 40x magnification. In addition, a note was made of areas with *status spongiosus*, which represent areas where demyelination may be inferred using H&E.

The TMEV model's inflammatory lesions are commonly found in cervical and thoracic spinal cord sections. Figure 3.5 A-D shows a representative sampling from each group using cervical spinal cord sections. The sections selected as representative for each treatment group had the mean scores of inflammation of that particular experimental group. All sections were taken at 40x magnification and were edited using SketchbookExpress (Autodesk) to remove excess structures (bone, muscle, etc). Panel A depicts an uninfected mouse, panel B depicts a placebo-treated infected mouse, panel C depicts an E₂-treated infected mouse, and Panel D shows an E₃-treated infected mouse. Within each panel, the red circles indicate perivascular cuffing and the red arrows represent *status spongiosus*, or the beginnings of demyelination. Similarly, Figure 3.6 A-D shows a representative sampling in a similar manner but using thoracic spinal cord sections.

Total cord area, gray matter area, and area of inflammation (sum of areas, including cuffing and parenchymal) were obtained using ImageJ (NIH). The white matter area was calculated and these data were used to obtain a percent white matter inflammation. The data was collected per mouse and later compiled leaving only white matter area and area of inflammation. These numbers were used to obtain a global percent inflammation per mouse and the averages and standard errors are presented in Table 3.5. This was the data used for statistical analysis. One-Way Analysis of Variance (ANOVA) with a Least Squares Difference (LSD) *post hoc* test was used to analyze the differences between groups. There was a significant difference among groups in

inflammation ($F_{(2,21)} : 8.770, p < 0.05$). The LSD *post hoc* test revealed there was also significant difference between E₂ and placebo ($p < 0.05$), E₃ and placebo ($p < 0.05$), and between E₂ and E₃-treated groups ($p < 0.05$) indicating an overall effect of treatment with E₂ having the greatest effect. Figure 3.7 presents the compiled data graphically as Mean \pm SEM.

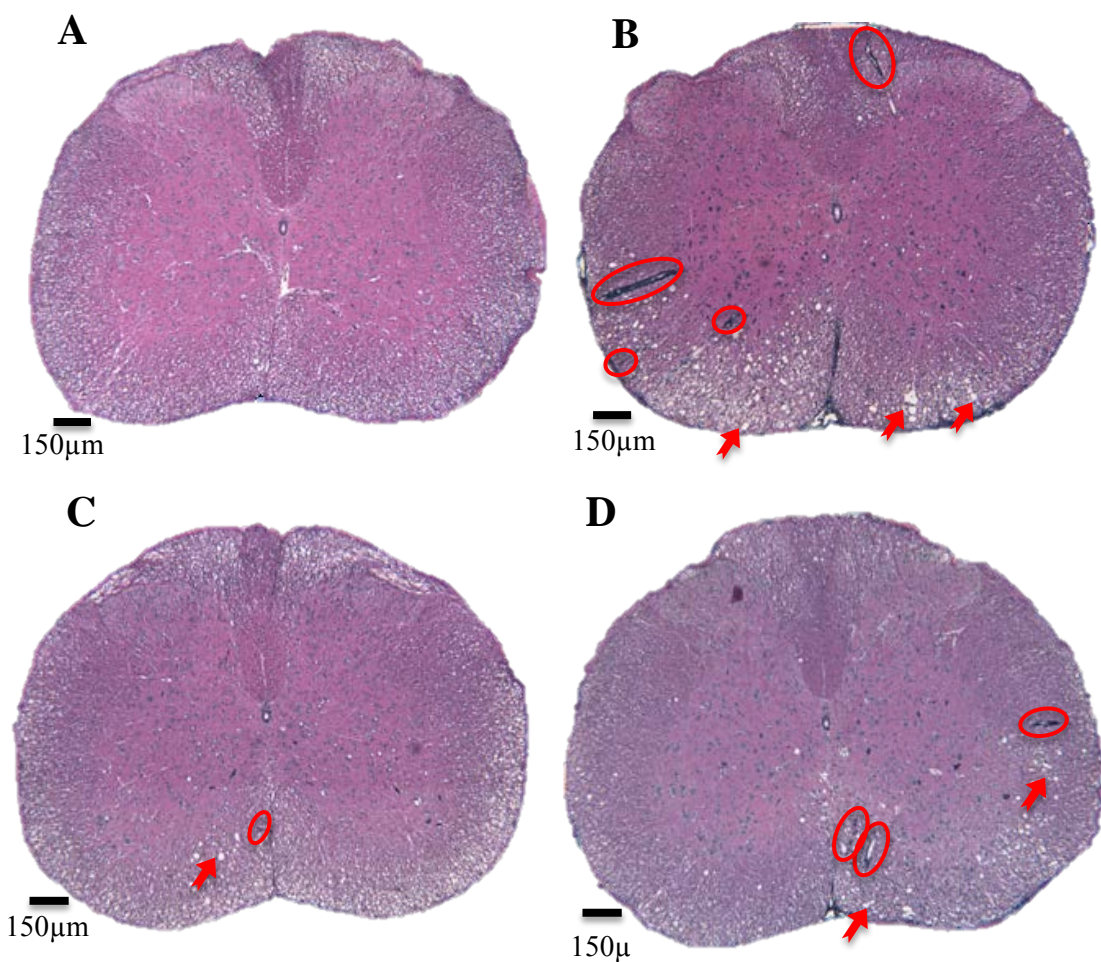


Figure 3.5 H&E-stained cervical spinal cord sections from control and three different treatment groups.

Panel A depicts a placebo-treated uninfected mouse. Panel B represents a placebo-treated infected mouse. Panel C represents an E₂-treated infected mouse. Panel D represents an E₃-treated infected mouse. The red circles depict areas of perivascular infiltration and the red arrows represent areas of parenchymal inflammatory infiltration and *status spongiosus*. All images captured at 40x magnification

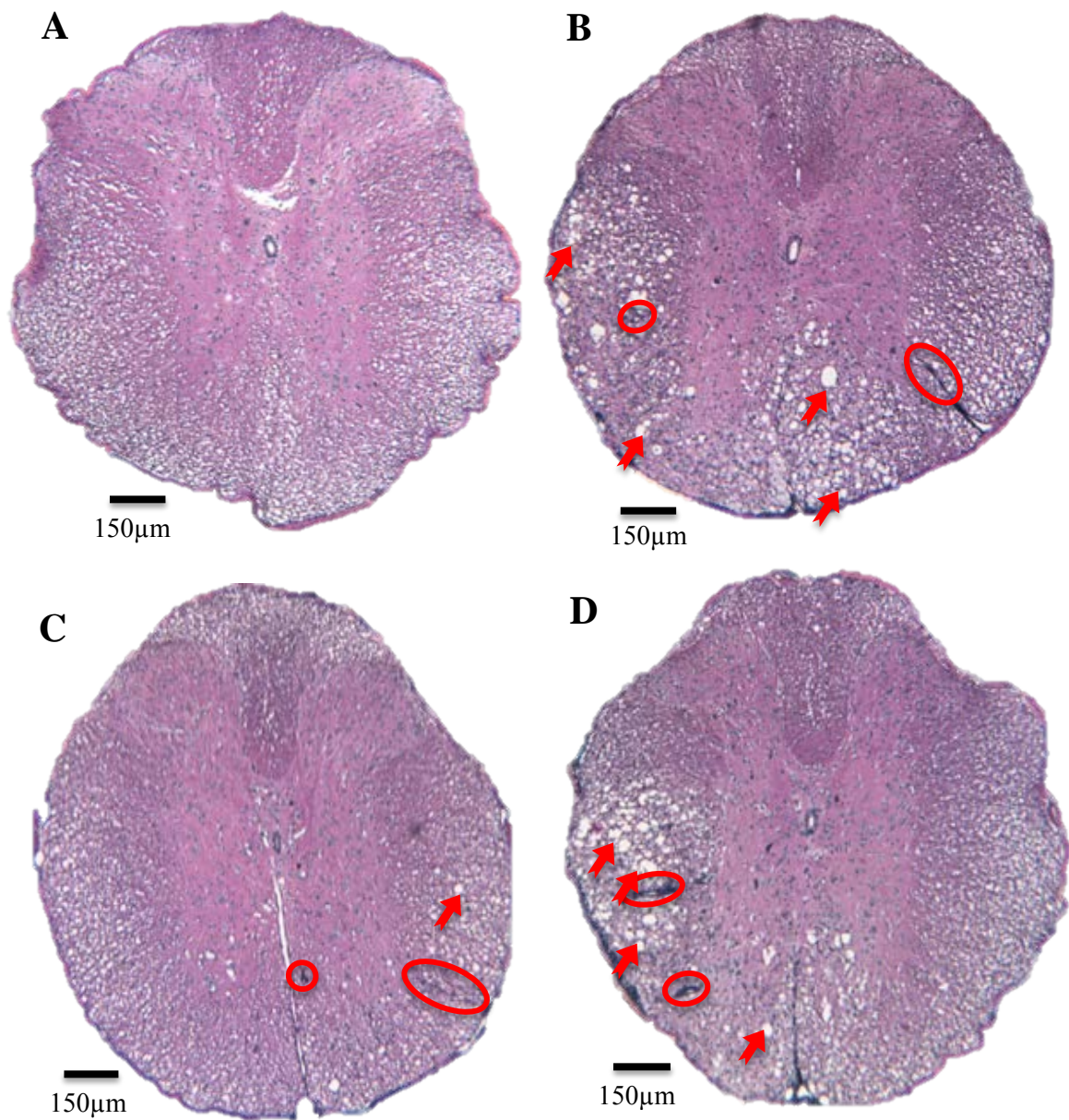


Figure 3.6 H&E-stained thoracic spinal cord sections from control and three different treatment groups.

Panel A depicts a placebo-treated uninfected mouse. Panel B represents a placebo-treated infected mouse. Panel C represents an E₂-treated infected mouse. Panel D represents an E₃-treated infected mouse. The red circles depict areas of perivascular infiltration and the red arrows represent areas of parenchymal inflammatory infiltration and *status spongiosus*. All images captured at 40x magnification

Table 3.5 Mean percent white matter inflammation.

The mean percent inflammation and standard errors are presented for each treatment group

	Placebo Tx n=6	E ₃ Tx n=8	E ₂ Tx n=8
Mean	0.76068	0.49727	0.25470
SEM	0.10324	0.09385	0.05026

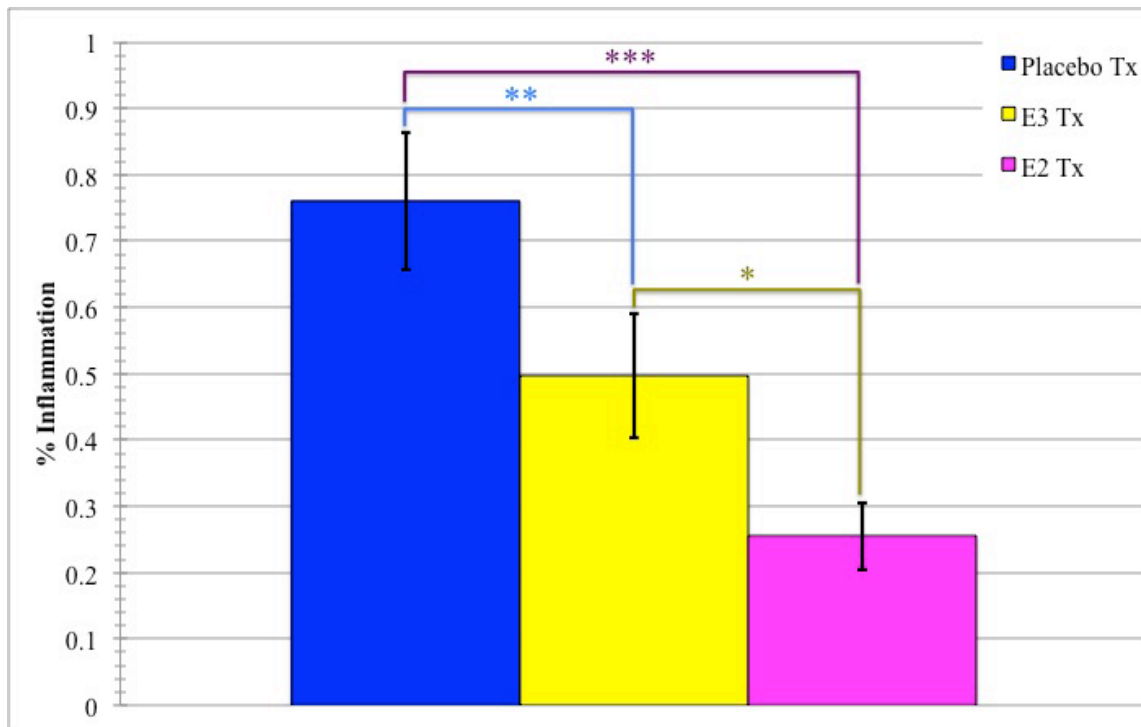


Figure 3.7 Graphical representation for the mean white matter inflammation per treatment group.

There is a significant difference among all three groups ($p < 0.05$). *Post hoc* tests revealed a significant difference between E₂ and E₃-treated groups (*), E₃ and placebo groups (**), and between E₂ and placebo groups (***). Data presented as Mean \pm SEM. * $p < 0.05$, ** $p < 0.05$, *** $p < 0.05$

3.5 Treatment with E₂ Reduces Demyelination in the Spinal Cord When Compared to Placebo

The Weil's stain for myelin was used to detect demyelination. We identified and outlined areas of demyelination at 40x magnification.

Similar to the location of inflammatory lesions, demyelinating lesions in TMEV are also found on primarily in cervical and thoracic spinal cord sections. Figure 3.8 A-D show a representative sampling from each group using cervical spinal cord sections. The sections selected as representative for each treatment group had the mean scores of demyelination of that particular experimental group. All sections were taken at 40x magnification and were edited using SketchbookExpress (Autodesk) to remove excess structures (bone, muscle, etc.). Panel A in Figure 3.8 depicts an uninfected mouse, panel B depicts a placebo-treated infected mouse, panel C depicts a E₂-treated infected mouse, and Panel D shows an E₃-treated infected mouse. Within each panel, the red arrows indicate areas of demyelination. Similarly, Figure 3.9 A-D shows a representative sampling using thoracic spinal cord sections.

Once again, three measurements were obtained from each section: Total cord area, gray matter area, and area of demyelination (sum of areas) if applicable. The white matter area affected was obtained from calculations and used to obtain a percent white matter demyelination. The white matter area was calculated and these data were used to obtain a percent white matter demyelination. The data was collected per mouse and later compiled leaving only white matter area and area of demyelination. These numbers were used to obtain a global percent demyelination per mouse and the averages and standard errors are presented in Table 3.6. This was the data used for statistical analysis. One-Way Analysis of Variance (ANOVA) with a Least Squares Difference (LSD) *post hoc* test was used to analyze the differences between groups. There was a significant difference among groups in inflammation ($F_{(2,21)}: 4.732, p<0.05$). The LSD *post hoc* test revealed there was also significant difference between E₂ and placebo ($p<0.05$). No significant difference was found between E₃ and placebo ($p>0.05$), or between E₂ and

E_3 -treated groups ($p>0.05$). Figure 3.10 presents the compiled data graphically as Mean \pm SEM.

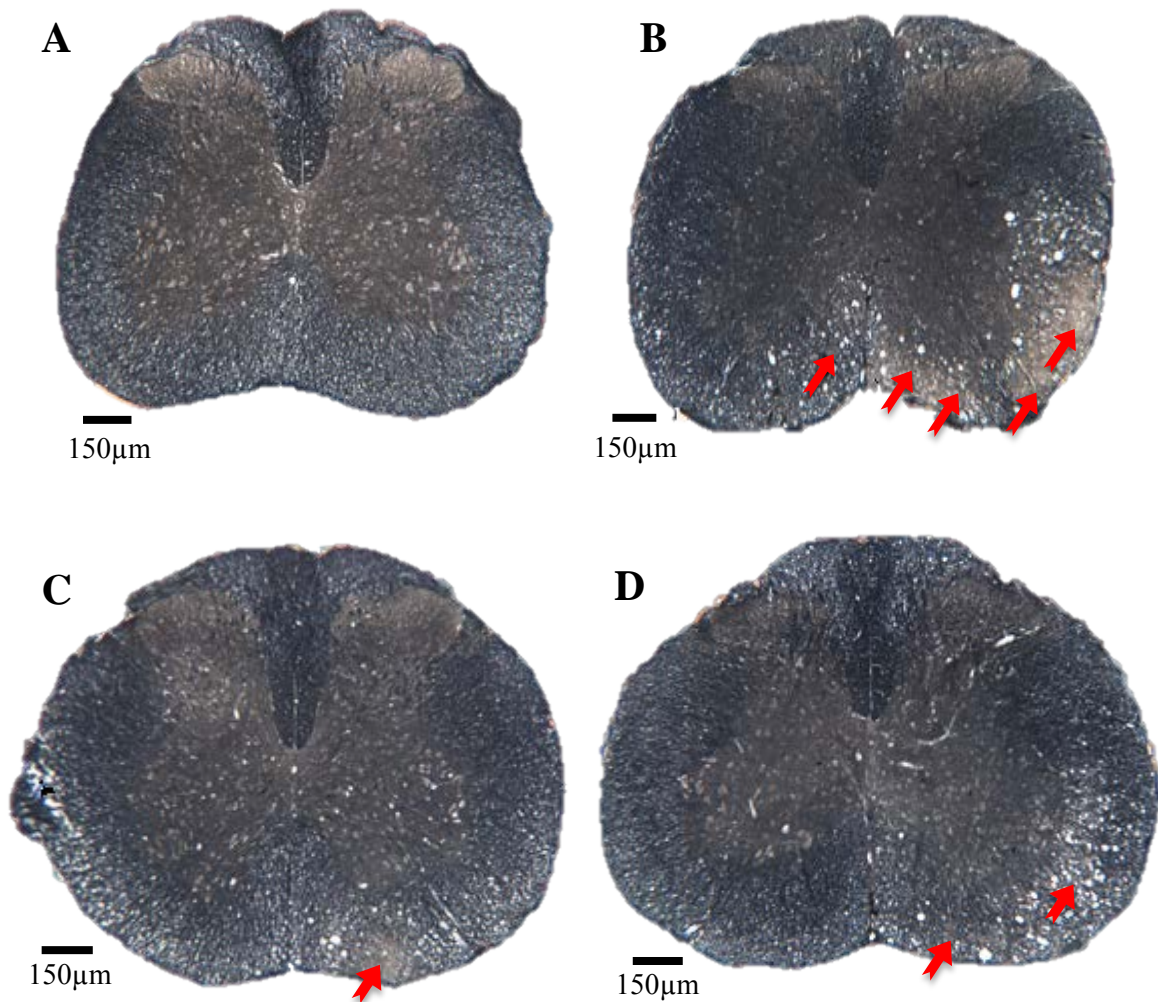


Figure 3.8 Weil's-stained cervical spinal cord sections from control and three different treatment groups.

Panel A depicts a placebo-treated uninfected mouse. Panel B represents a placebo-treated infected mouse. Panel C represents an E_2 -treated infected mouse. Panel D represents an E_3 -treated infected mouse. The red arrows show areas of demyelination. Panel D is at the early stages of demyelination and the arrows point to primarily *status spongiosus* with the left arrow pointing to a small area of demyelination. All images captured at 40x magnification

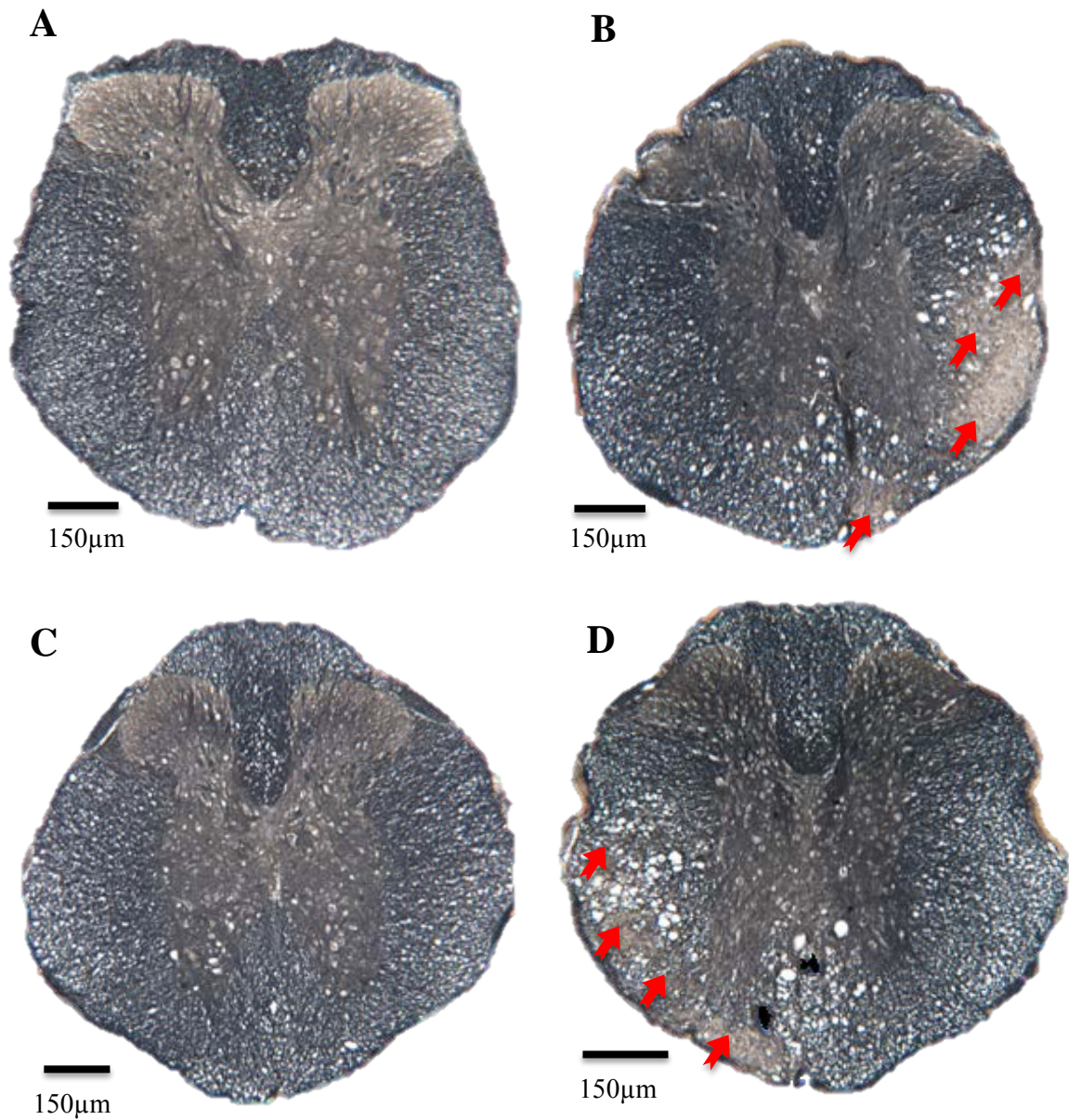


Figure 3.9 Weil's-stained thoracic spinal cord sections from control and three different treatment groups.

Panel A depicts a placebo-treated uninfected mouse. Panel B represents a placebo-treated infected mouse. Panel C represents an E₂-treated infected mouse. Panel D represents an E₃-treated infected mouse. The red arrows show areas of demyelination. Panel C shows no apparent signs of demyelination. All images captured at 40x magnification

Table 3.6 Mean percent white matter demyelination.

The mean percent white matter demyelination and standard errors are presented for each treatment group

	Placebo Tx n=6	E ₃ Tx n=8	E ₂ Tx n=8
Mean	13.35095	8.07803	4.54065
SEM	2.02817	2.27615	1.47107

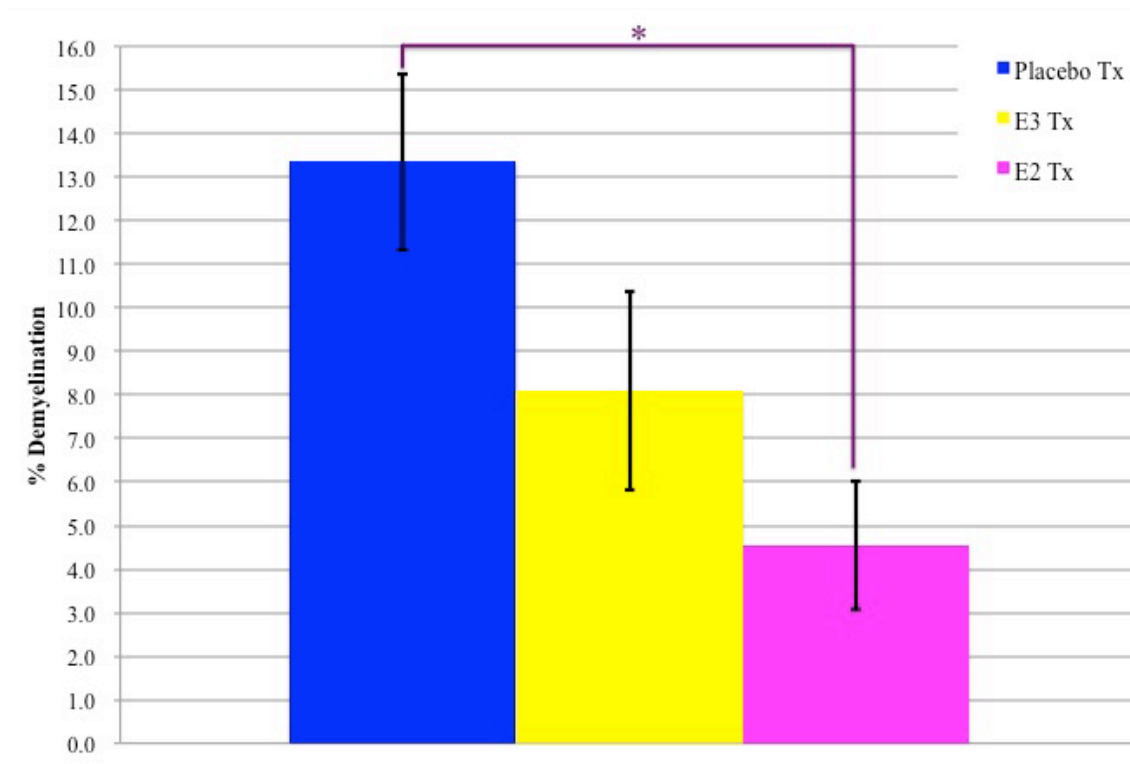


Figure 3.10 Graphical representation of the mean percent white matter demyelination per group.

There is a significant difference in percent white matter demyelination among all three groups ($p < 0.05$). *Post hoc* tests revealed no significant difference between E₂ and E₃-treated groups ($p > 0.05$), or E₃ and placebo groups ($p > 0.05$). However, there was a significant difference between E₂ and placebo groups (*). Data presented as Mean \pm SEM. * $p < 0.05$

3.5.1 Serial Section Comparison between H&E and Weil's-stained Spinal Cord Sections Allow Visual Co-localization of Inflammatory and Demyelinating Lesions

H&E sections from representative mice were chosen as described previously and matched to the Weil's stain serial sections to visually correlate inflammatory lesion location to demyelinating lesion location. All sections were captured at 40x magnification and were edited using SketchbookExpress (Autodesk) to remove excess structures (bone, muscle, etc.). Figure 3.11 A-D depict serial cervical sections. Row A represents cervical sections from control-uninfected mice from H&E and Weil's stains, respectively. Row B represents sections from placebo-treated mice, row C represents E₂-treated infected mice, and row D represents E₃-treated infected mice in the same manner as described above. Figure 3.12 A-D depict serial thoracic sections. Row A represents thoracic sections from control uninfected mice from the infection and demyelination studies, respectively. Row B represents sections from placebo-treated mice, row C represents E₂-treated infected mice, and row D represents E₃-treated infected mice in the same manner as described above.

All the serial sections that show inflammation, also have signs of demyelination that can be co-localized to areas near perivascular cuffing and areas that show *status spongiosus* in H&E-stained sections.

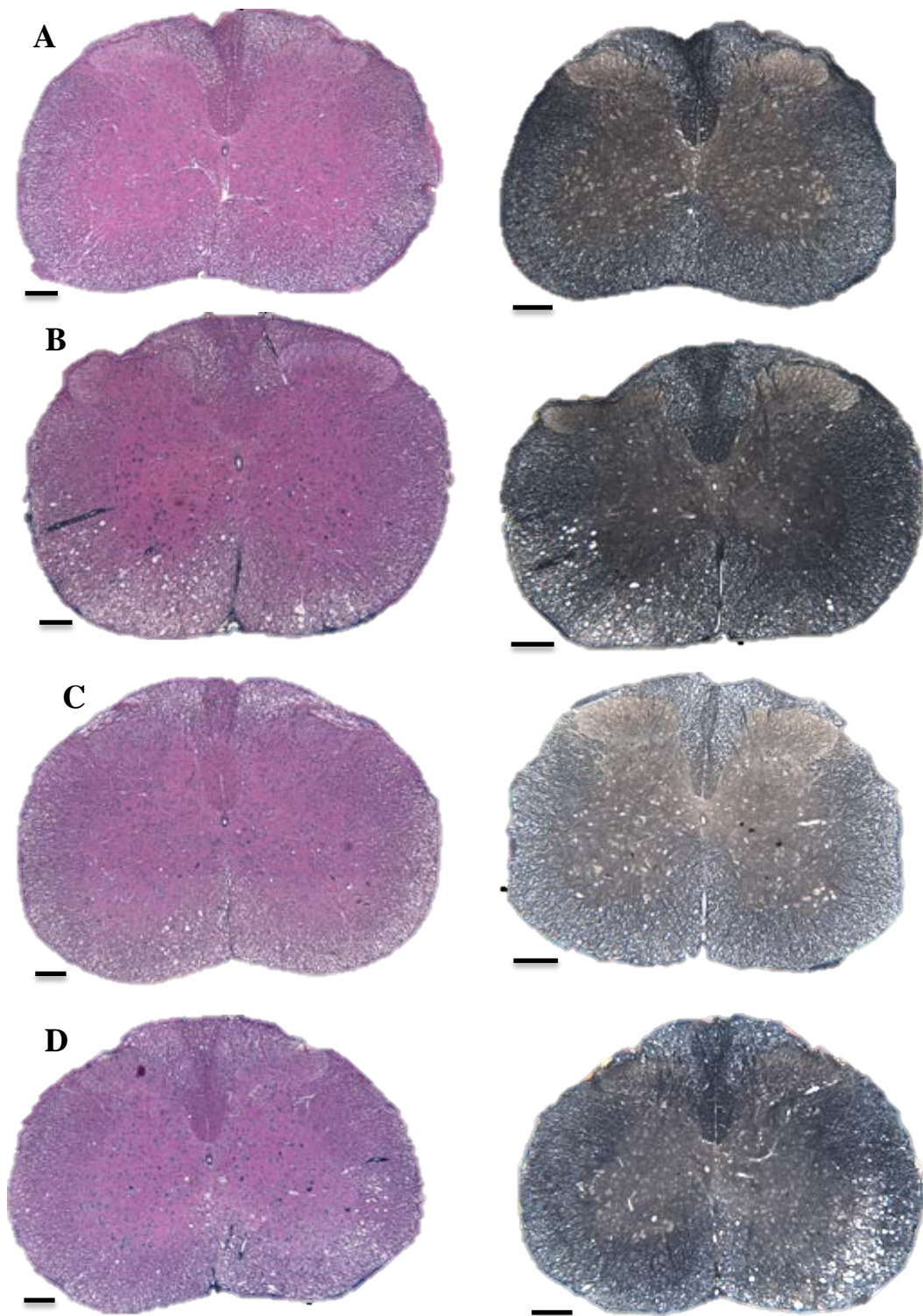


Figure 3.11 Cervical spinal cord serial section comparisons between H&E (inflammation) and Weil's (demyelination) stains.

Row A represents control-uninfected mice, B represents placebo-treated infected mice, C represents E₂-treated infected mice and D represents E₃-treated infected mice. Bar = 150µm. 40x magnification

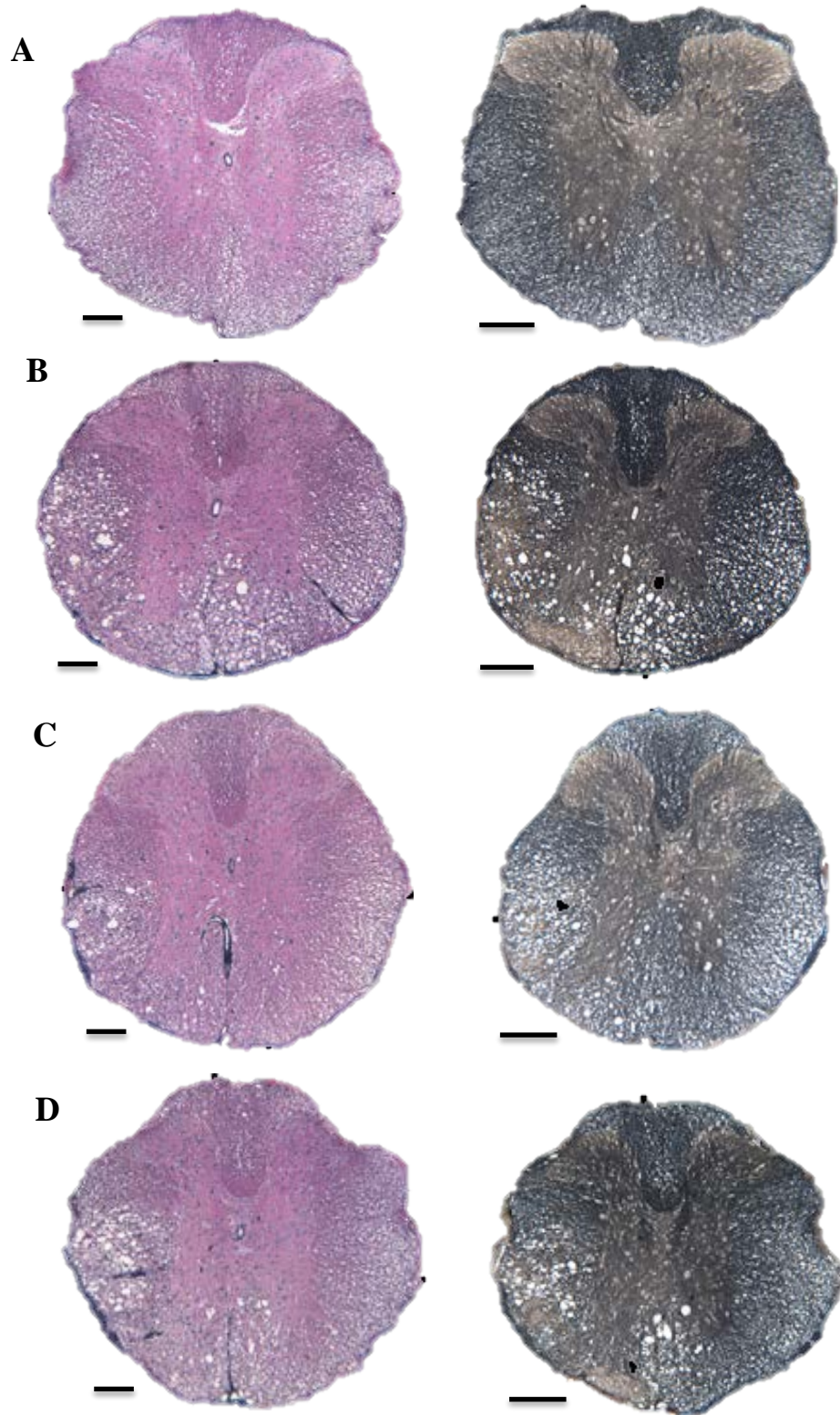


Figure 3.12 Thoracic spinal cord serial sections between H&E (inflammation) and Weil's (demyelination) stains.

Row A represents control-uninfected mice, B represents placebo-treated infected mice, C represents E₂-treated infected mice and D represents E₃-treated infected mice. Bar = 150 μ m. 40x magnification

3.6 Radio-immuno Assay Measurements of Theiler's Virus Antibody (Ab) Levels Show Differences between Infected and Uninfected Groups and That Both E₂ and E₃-Treatments Differ from Placebo in the Infected Groups

Radio-Immuno (RIAs) for Theiler's virus antibody (GD-VII) were conducted on the sera collected from the mice at termination. The data was compiled and averaged by group to create Tables 3.7 and 3.8 and graphed as counts per minute (CPM) versus dilution as shown in Figure 3.13. Figure 3.14 shows the graphed CPM vs. dilution values for the infected groups only. By the 1/160 dilution, there are no detectable differences among control or infected groups.

A Two-way Analysis of Variance (ANOVA) was done to evaluate the difference between both uninfected and infected mice and their respective hormone treatment subgroups at the 1/40 dilution, as this dilution is a good representative choice for the range of dilutions up to 1/160. There was a significant difference in Theiler's Virus antibody (Ab) levels among groups ($F_{(5,43)}: 6.447, p<0.05$). Infection with virus had a significant effect on Ab levels ($F_{(1,43)}: 15.124, p<0.05$) as did hormone treatment ($F_{(2,43)}: 7.841, p<0.05$). There was no interaction between virus and hormone ($F_{(2,43)}: 2.780, p>0.05$) at the 1/40 dilution. *Post hoc* Least Squares Difference (LSD) analysis revealed that there was a significant difference between placebo and E₃ ($p<0.05$) and between placebo and E₂ ($p<0.05$) in the 1/40 dilution. No significant difference was found between E₃ and E₂ ($p>0.05$).

A comparison of only the infected groups using 1-way ANOVA revealed similar significant differences among the three groups in the 1/40 dilution ($F_{(2,21)}: 5.579, p<0.05$). *Post hoc* Least Squares Difference (LSD) analysis revealed that there was a significant difference between placebo and E₃ ($p<0.05$) and between placebo and E₂ ($p<0.05$) at the 1/40 dilution. No significant difference was found between E₃ and E₂ ($p>0.05$).

Table 3.7 Average RIA counts per minute for treated, uninfected groups.

The mean counts per minute for serial dilutions from 1/20 to 1/160 are presented in this table for each uninfected and treated mouse group. The 1/40 dilution was used for statistical analysis

Control (n=8)				
Dilution	1/20	1/40	1/80	1/160
Average	610.500	390.875	266.250	202.125
SEM	200.219	114.143	63.818	59.379
E ₃ Control (n=7)				
Dilution	1/20	1/40	1/80	1/160
Average	307.857	194.857	147.429	115.143
SEM	77.346	40.130	17.389	4.758
E ₂ Control (n=7)				
Dilution	1/20	1/40	1/80	1/160
Average	401.143	182.857	164.857	142.857
SEM	178.105	15.251	14.287	13.869

Table 3.8 Average RIA counts-per-minute for treated, infected groups.

The mean counts per minute for serial dilutions from 1/20 to 1/160 are presented in this table for each infected and treated mouse group. The 1/40 dilution was used for statistical analysis

Placebo Tx (n=6)				
Dilution	1/20	1/40	1/80	1/160
Average	1,797.667	1,142.000	761.833	477.500
SEM	446.942	328.822	210.086	112.335
E ₃ -Tx (n=8)				
Dilution	1/20	1/40	1/80	1/160
Average	475.500	315.500	262.875	222.875
SEM	86.689	45.293	33.625	16.576
E ₂ -Tx (n=8)				
Dilution	1/20	1/40	1/80	1/160
Average	976.250	577.375	399.750	299.250
SEM	217.872	112.100	67.546	60.535

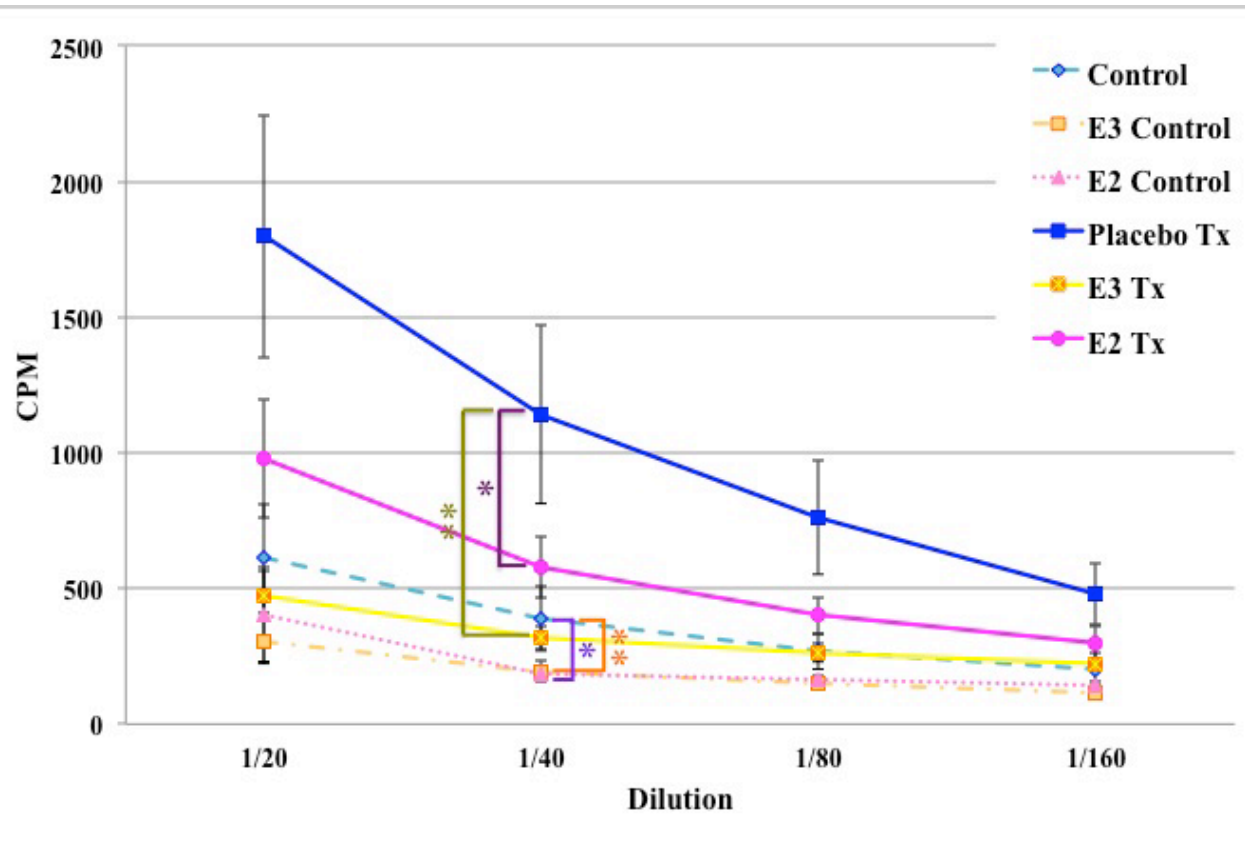


Figure 3.13 Radio-immuno assay analysis for TMEV antibody levels in the sera of uninfected and infected mice.

Data for the 1/40 dilution was chosen as a representative sample for the assay. Dashed lines represent control-uninfected groups and the solid lines represent treated-infected groups. There was a significant difference among groups in Ab levels to TMEV ($p < 0.05$) with a main effect of virus infection ($p < 0.05$) and hormone treatment ($p < 0.05$). There was no interaction between virus and hormone ($p > 0.05$). There was a significant difference between E_2 and placebo (*) as well as between E_3 and placebo groups for TMEV Ab levels (**) in both infected and uninfected groups. No significant difference ($p > 0.05$) was found between E_2 and E_3 -treated groups in either infected or uninfected groups. Data shown as Mean \pm SEM. * $p < 0.05$, ** $p < 0.05$

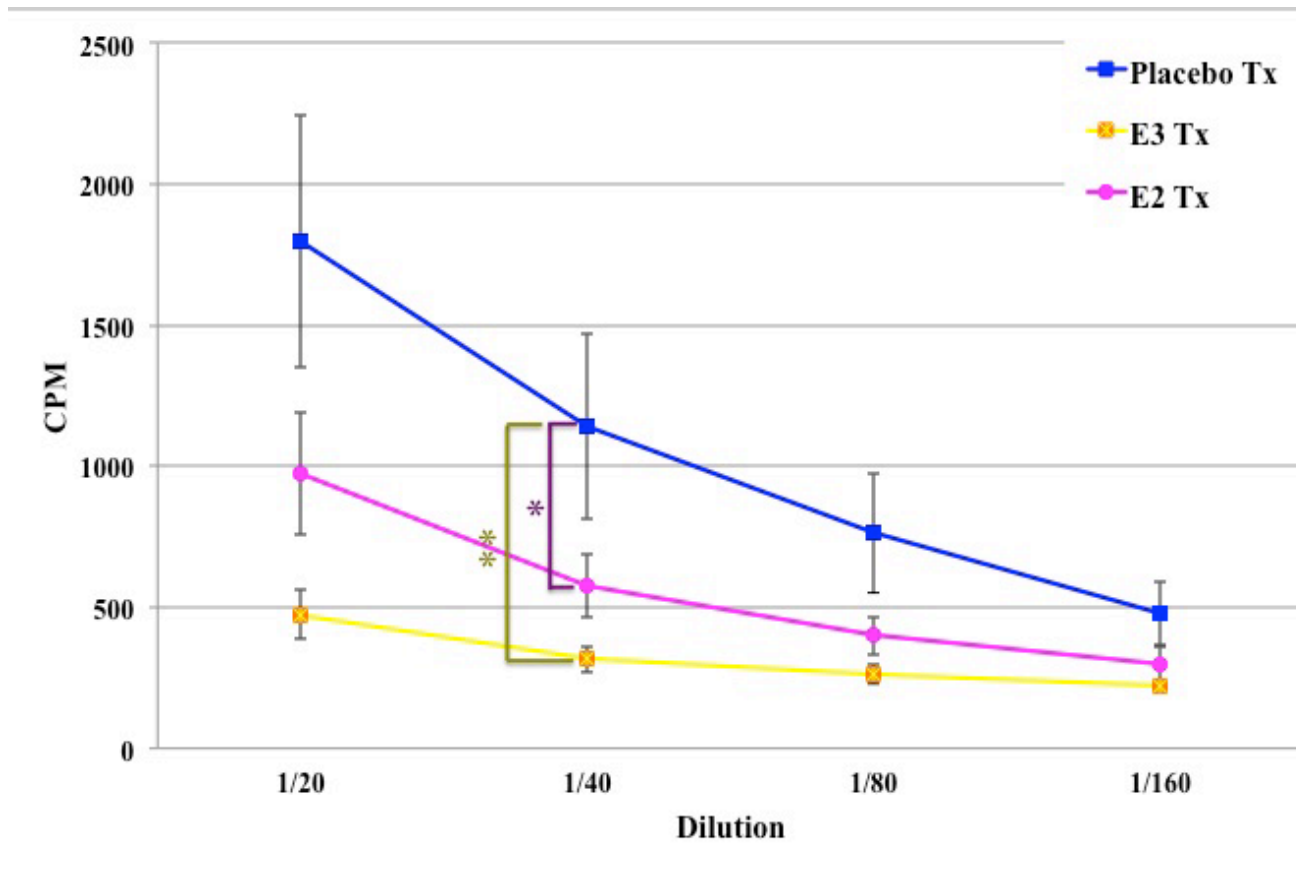


Figure 3.14 Radio-immuno assay analysis for TMEV antibody levels in the sera of infected mice

This graph presents the data for the 1/40 dilution of TMEV infected groups only. There is a significant difference ($p < 0.05$) among all three groups in Ab levels. No significant difference was found between E_3 and E_2 ($p > 0.05$). Placebo treated mice differ significantly from both E_2 -treated (*) and E_3 -treated mice (**). Data is presented as Mean \pm SEM. * $p < 0.05$, ** $p < 0.05$

4. DISCUSSION

This project assessed the effect of hormone-replacement therapy on the chronic phase of Theiler's Murine Encephalomyelitis Virus (TMEV) infection. The data analysis focused on observations from week 15 post-infection (pi), which corresponds to week 17 of the experiment. Chronic demyelination in the Theiler's infection model is considered to begin 2-5 months pi and remain for the life of the mouse causing progressive disease similar to progressive human multiple sclerosis (Miller *et al.*, 1995). For our treatments, we chose a 0.1mg pellet based on previous experiments performed by Bebo *et al.*, which indicated this dose, equivalent to diestrus levels of 17- β -estradiol (E_2) in mice (25-50pg/mL) was effective in significantly reducing EAE clinical scores compared to placebo (Bebo *et al.*, 2001). Because estriol (E_3) is a hormone of pregnancy produced by the fetal-placental unit (Sicotte *et al.*, 2002), we used information from Bebo *et al.* and from Palaszynski *et al.* to select a 5.0mg E_3 pellet, which would give pregnancy hormone level equivalents (2000-3000pg/mL), this level of E_3 in circulating blood reduced incidence of EAE compared to placebo (Bebo *et al.*, 2001; Palaszynski *et al.*, 2004). The levels of E_2 and E_3 chosen led to increases in uterine size. For both infected and uninfected groups, the weight of the uteri was significantly different than placebo-treated mice. Because the mice were ovariectomized, this measurement allowed us to determine the effectiveness of the hormone replacement. Our results indicate that the doses of E_2 and of E_3 we selected have a positive effect on inflammation. The spleen is an important secondary lymphoid organ, our results showed that the spleens in the infected groups on average weighed less than those of the uninfected groups; however, our results showed no significance in weight. This is consistent with findings by Mendez-Fernandez *et al.*, who showed the spleen was not necessary for TMEV-specific $CD8^+$ cell-activation and recruitment (Mendez-Fernandez *et al.*, 2005).

The current study showed that E_2 has therapeutic effects on the Theiler's Virus-Induced Demyelination (TVID) model of multiple sclerosis. Our experiments demonstrated that treating young ovariectomized female mice with estradiol reduced clinical signs of disease as well as histological signs of inflammation and demyelination

in the spinal cord. These findings were consistent with our hypothesis. Treatment with E₃ also was effective in reducing signs of inflammation; however, it showed no difference from placebo-treated mice in improvement of clinical scores or demyelination. In the experimental autoimmune encephalitis (EAE) model, females that were ovariectomized but not treated did not show altered disease severity when compared to placebo-treated non-ovariectomized mice. This indicates that physiological levels of estrogens may not be enough to cause any effect on disease progression in this model once the disease has begun (Voskuhl and Palaszynski, 2001). Conflicting results were shown if treatment with physiologic levels of E₂ and E₃ were administered prior to the onset of EAE (Bebo *et al.*, 2001). Ovariectomy may have been a factor for the lessened effect of E₃ in our experiment; however, EAE model is a pure autoimmune model that does not account for the effects of the virus present in TVID; rather it directs a CD4⁺ response to myelin components (Miller *et al.*, 1995). Nevertheless, E₃ has been shown to be protective in studies with human MS patients and in the non-ovariectomized EAE model of MS (Palaszynski *et al.*, 2004; Sicotte *et al.*, 2002; Spence and Voskuhl, 2012). It is difficult to assess the reason behind the inability of E₃ treatment to diminish demyelination. Perhaps the length of treatment was not sufficient to allow the hormone to have an effect.

The discrepancy in our results could be explained by differences in the binding of estriol *versus* estradiol to estrogen receptors (ERs). Both estriol and estradiol can bind to estrogen receptors α and β (ER α and ER β , respectively). However, estradiol has a higher affinity for ER α while estriol has a higher affinity for ER β (Spence and Voskuhl, 2012). ER α has been implicated in EAE as having a potent anti-inflammatory response. Binding of E₂ to ER α inhibits the transport of the pro-inflammatory transcription factor NF- κ B into the nucleus via interaction with PI3K, most notably in microglia, the resident macrophages of the CNS (Spence *et al.*, 2011; Vegeto *et al.*, 2008). In ischemia, ER α -mediated suppression of these cells and other inflammatory cells seems to be responsible for neuroprotection in young female rats post-stroke (Selvamani and Sohrabji, 2010). *In vitro* experiments have shown that stimulation of ER α by E₂ or another agonist

suppresses transcription of NF- κ B in astrocytes and TNF- α -mediated production of pro-inflammatory CCL2 (Giraud *et al.*, 2010) and to modulate infiltration of peripheral lymphocytes into the CNS, whereas ER β did not show any effect (Spence and Voskuhl, 2012). In addition, treatment with ER α ligand reduced spinal cord inflammation, whereas ER β ligand treatment only showed protective effects in the long term, but no reduction in inflammation (Tiwari-Woodruff and Voskuhl, 2009). These observations help explain the relative abundance of inflammatory cells in the parenchyma of the spinal cord sections for E₃-treated mice *versus* E₂-treated mice in this project, as well as the decreased demyelination observed on the Weil's stained sections. One final observation is that E₂ increases the ability of astrocytes to resorb excess glutamate, thus preventing neuronal loss due to excitotoxicity and explaining the relative effectiveness of E₂ *versus* E₃ therapeutic effects on treated mice (Spence and Voskuhl, 2012).

The role of E₂ and astrocytes is more complex. Astrocytes are essential components of the blood-brain barrier (BBB), and they are important in cell-to-cell communication, not only in the CNS, but also with inflammatory cells (Kipp and Beyer, 2009). E₂ has been shown to prevent astrocytes from expressing MHC-II and to enhance the induction of T-cell apoptosis, via astrocytes by reduction of H³-thymidine incorporation by the T-cells (Kipp and Beyer, 2009). Expression of ER α is necessary in astrocytes to prevent axonal loss, gliosis, and diminish or prevent lymphocyte and monocyte infiltration into the CNS (Spence *et al.*, 2011). Estradiol is known to have positive effects in brain circulation and in young animal models, and reduce stroke severity (Cipolla *et al.*, 2009; Selvamani and Sohrabji, 2010). The reduction in stroke severity is due to E₂'s influence on the BBB, both endothelial cells and astrocytes. During stroke, edema causes damage to the parenchyma. Cipolla *et al.* found that ovariectomy in rats increases BBB permeability, but that treatment with E₂/E₃ combination therapy restores the integrity of the barrier almost to control level (Cipolla *et al.*, 2009) The levels of E₂ found in this study are almost at the levels found at estrous in the rat ovarian cycle (Shaikh, 1971), making them a comparable measure to the dose of E₂ given in this study. A similar mechanism may also be taking place in our study:

with the maintenance of BBB integrity, less lymphocytes and macrophages would be able to migrate into the parenchyma, thus reducing inflammation. The Cipolla study did not clarify whether E₂ or E₃ had a more significant effect on BBB permeability.

The Theiler's virus model is effective in showing pathological signs consistent with human MS. This model is unique in its biphasic disease onset. Strains such as the Daniel's (DA) and BeAn strain set up an acute gray matter inflammation. Studies using Severe Combined Immunodeficient (SCID) mice, showed that the virus could infect neurons and glia early on (6h to 7d pi) in both SCID and immunocompetent mice establishing infection in the corpus callosum and in hippocampal pyramidal neurons (Njenga *et al.*, 1997) leading to the development of pathological polioencephalomyelitis, though most mice remain asymptomatic (Sato *et al.*, 2011). In addition, SCID mice had virus in neuronal cell bodies in the spinal cord after 7d pi compared to immunocompetent mice. This indicated that virus was cleared before 7d pi in immunocompetent mice. This study revealed that both cellular and humoral immune responses are clearing the virus from gray matter, with the humoral branch being less efficient and also that both CD4⁺ and CD8⁺ cells have similar functions in gray matter viral clearance (Njenga *et al.*, 1997).

It is not entirely clear why the virus cannot be fully cleared from white matter. It has been shown that it can persist in, macrophages, oligodendrocytes, axons, or glia, and that the attempt to clear it may be the main cause of the demyelinating chronic pathology that follows acute infection (Njenga *et al.*, 1997; Sato *et al.*, 2011). Infection in the TMEV model seems to be modulated by T_H1 mechanisms and based on our results, inflammation in the spinal cord parenchyma corresponds to areas of demyelination. This confirms the previous observation. A possible mechanism for the observed pathogenesis of TMEV can be elucidated from observations done by previous groups and observations from this work. The virus is capable of causing damage to axons, and by consequence to myelin and oligodendrocytes or directly infect and kill oligodendrocytes (Sato *et al.*, 2011).

It was determined that the L-protein in Theiler's virus can inhibit gene

transcription, interfere with translation by preventing the export of mRNA from the nucleus, and regulate apoptotic signals. Thus the L-protein can cause cell death without inflammation (Ghadge *et al.*, 2011). This may explain why the polioencephalomyelitis symptoms are limited. E₂ treatment has also been shown to inhibit Theiler's virus replication (Kipp and Beyer, 2009), which may help explain the reduction in pathological findings in this treatment group.

Theiler's virus is unique among non-enveloped viruses in that it can exit a cell without lysis whereas other non-enveloped viruses need to lyse the cell once their replication is complete (Roussarie *et al.*, 2007). This ability may help explain how TMEV is able to move from neuron to oligodendrocytes without causing neuronal cell death in the majority of cases and setting up a persistent infection in these white matter cells. Roussarie *et al.* propose that TMEV uses conventional cell machinery to migrate from axon to myelin and the group mentioned a hypothesis proposed by Gatzinski *et al.* which states axons may use myelin to remove unwanted materials, in this case virions (Roussarie *et al.*, 2007). Because of the number of internodes and axons one oligodendrocyte can myelinate, its death can cause extensive demyelination. However, in the event of axonal damage, demyelination can also occur. Damage in the EAE model is accepted to come from "bystander" CD4⁺-mediated injury to CNS tissue when these cells respond to myelin antigen, principally myelin proteolipid protein (PLP) (Sato *et al.*, 2011) and myelin basic protein (MBP) is also involved in disease pathogenesis in EAE (Dal Canto *et al.*, 2000). The PLP epitope used to induce EAE (PLP 139-152) seems to be involved in TMEV autoimmunity and as the disease progresses this epitope "spreads" to include other regions of the protein and other similar proteins such as Myelin Oligodendrocyte Glycoprotein (MOG) (Dal Canto *et al.*, 2000).

Organotypic spinal cord cultures from SJL/J mice were prepared and treated with lymphocytes from TMEV-infected mice. They were then stimulated with MBP or PLP. Under MBP stimulation, the axons appeared normal and their numbers did not differ from naïve or sham-infected mice cultures; however, under PLP stimulation, the majority of the axons were heavily demyelinated (Dal Canto *et al.*, 2000). These results

indicate that TMEV mechanisms of action may be different from the EAE model, where anti-MBP lymphocytes can cause culture demyelination and transfer disease to naïve mice (Dal Canto *et al.*, 2000). EAE could be protected by anti-PLP Ab when whole myelin was used to prime the mice while anti-MBP Ab had no effect (Dal Canto *et al.*, 2000). The TMEV model exhibited similar properties due to the liberation of myelin fragments early in the disease (Dal Canto *et al.*, 2000). In addition CD8⁺ T-cells expressing MHC-I infiltrate the CNS to clear the virus, but in doing so cause damage to myelin by recognizing self-epitopes (Sato *et al.*, 2011; Tsunoda *et al.*, 2005). Experimentally, self-recognizing CD8⁺ T-cells from MS patients can elicit an immune response against CNS tissue in naïve mice, demonstrating the ability of this cell subtype to cause CNS pathology (Tsunoda *et al.*, 2005). CD8⁺ T-cells are the most likely candidates for causing axonal damage that leads to demyelination. Roos *et al.* report that both CD4⁺ and CD8⁺ T-cells can induce demyelination since depletion of either population leads to myelin breakdown, and that the DA strain of TMEV can damage myelin in both MHC-I and MHC-II-deficient mice (Roos, 2010). However, mice that lack either CD8⁺ cells or MHC-I have less severe disease outcomes, thus the CD8⁺ population may cause more damage to axons, impede repair mechanisms, and lead to demyelination (Roos, 2010).

The CNS is considered to be an immunoprivileged site. In infection, however, this is not the case. The cytokine environment created by inflammatory cells determines whether or not the brain will have inflammation and demyelination. CD4⁺ cells respond to TMEV infection early on. Of the two subtypes of T-helper (T_H) cells, the response mounted against TMEV seems to be T_H1-mediated, or pro-inflammatory. T_H1 cells are polarized by interleukin-12 (IL-12), which is a pro-inflammatory cytokine. CD4⁺-T_H1 cells will secrete cytokines such as interleukin-2 (IL-2), interferon gamma (IFN γ) and tumor necrosis factor beta (TNF β). These cytokines are involved in cell-mediated immunity and work to activate macrophages (M Φ) (via IFN γ), recruit lymphocytes (via TNF β +IFN γ), cause CD8⁺ cells to differentiate into cytolytic T-cells (T_{CL}) (via IL-2+IFN γ), and switch IgG to IgG_{2a}, which binds to F_{C γ} and binds complement to help in

opsonization (Abbas *et al.*, 1996). In MS and EAE, a new subset of T_H cells has also been implicated in inflammation and demyelination. T_H17 cells secrete IL-17 and are induced by IL-6 and transforming growth factor beta (TGF β), and maintained by IL-23, a cytokine related to IL-12 by its p40 subunit (Aranami and Yamamura, 2008; Fletcher *et al.*, 2010); thus, for a long time, the presence of T_H17 cells was unnoticed until 2003, when Cua *et al.* discovered this subgroup of cells (Aranami and Yamamura, 2008). IL-17 attracts neutrophils into the brain, and causes the release of more pro-inflammatory cytokines (Fletcher *et al.*, 2010). T_H17 CD4⁺ cells can induce EAE in naïve mice, thus they are now associated with the pathogenesis of chronic demyelinating disease of the CNS. It is not surprising that this cytokine environment causes damage to CNS tissue as observed in the placebo-treated infected mice.

Estrogens have been shown to induce a switch to a T_H2, or anti-inflammatory environment. E₂ was shown to inhibit T_H1 and T_H17 cell differentiation in the EAE model of MS via interaction with ER α (Lélu *et al.*, 2011), which is of significance to this project. Estrogens are known to induce a shift toward T_H2-mediated immunity by influencing T-cells directly via interaction with ER α (Giraud *et al.*, 2010), especially during pregnancy (Nicot, 2009).

Theiler's virus has been shown to increase expression of cyclooxygenase-2 (COX-2) in oligodendrocytes and astrocytes. The enzyme cyclooxygenase-2 (COX-2) converts arachidonic acid to prostanoids, which are known to mediate responses during inflammation. Of notable interest is the pro-inflammatory molecule Prostaglandin E2 (PGE2), which is known to induce excitotoxic neuronal death by binding to EP1 receptors. Similar mechanisms are thought to be at play for oligodendrocyte death (Carlson *et al.*, 2010). Treatment with E₂ has been shown to protect oligodendrocytes from glutamate excitotoxic cell death by reducing expression of COX-2, which renders them vulnerable to the toxic effects of glutamate (Carlson *et al.*, 2010; Molina-Holgado *et al.*, 2002), and increasing astrocyte reuptake of glutamate (Spence and Voskuhl, 2012).

The T_H2 cytokine environment is anti-inflammatory and it causes a shift toward humoral immunity. T_H2 cells produce cytokines that help reduce inflammation such as IL-3, 4, 5, 10, and 13. Some of these cells produce $TGF\beta$, but it is produced in other tissues (Abbas *et al.*, 1996) and depending on the cytokine milieu, it can induce a shift toward T_H17 or induced regulatory T-cells (iT_{Reg}) (Aranami and Yamamura, 2008). IL-4 causes a shift in IgG subclass to neutralizing (non-complement binding) IgG_1 . In B-cells it switches to IgE expression, which helps in mast cell-mediated reactions (response to allergens). IL-3, 4, and 5 cause activation of eosinophils and increase the production of eotaxin to guide them when dealing with helminthes. IL-4 also inhibits $IFN\gamma$ and prevents $M\Phi$ activation, IL-10 inhibits $M\Phi$ responses, and $TGF\beta$ inhibits activation of other leukocytes. Thus, T_H2 -mediated immunity prevents acute and chronic inflammation – including the delayed-type hypersensitivity (DTH) reactions brought about by T_H1 $CD4^+$ cells and limit their responses (Abbas *et al.*, 1996).

During classical infection, it is common to initiate T_H1 -mediated immunity early and then switch to T_H2 for control and memory functions (Abbas *et al.*, 1996). However, during autoimmune chronic inflammation or infection, having T_H1 responses may not be ideal. The switch from T_H1 to T_H2 -mediated immunity explains the observations made in the inflammation and demyelination sections of this project.

In addition T_H2 $CD4^+$ cells can induce the shift of uncommitted T-cells into iT_{RegS} such as T-regulatory-1 (T_{R1}), T_{R3} , and $CD8^+T_{Reg}$ via secretion of anti-inflammatory cytokines $TGF\beta$ and IL-10, and retinoic acid (RA) (Fletcher *et al.*, 2010). Natural T_{RegS} (nT_{RegS}) are found in the thymus and they express CD25 and Forkhead box P3 (FoxP3) transcription factor on their cell surface ($CD25^+FoxP3^+$) to elicit regulatory functions (Fletcher *et al.*, 2010). It is known that nT_{RegS} regulate tolerance to self and are important in autoimmunity. In MS patients, increased T_{Reg} activity correlates with symptom remission as the cells try to combat increased inflammation from T_H1/T_H17 cells, and one can speculate that perhaps the relapsing-remitting onset of MS may be related to success of these cells in controlling inflammation, this however is beyond the scope of this project. In the TMEV model, specifically in SJL/J mice, T_{RegS} are

preferentially activated, and this prevents the virus from being properly cleared from the CNS, which brings us closer to understanding the reason behind the chronic inflammation caused by the virus, and also to understanding MS pathogenesis (Getts *et al.*, 2010).

In a recent study by Valor *et al.*, MS pregnant and non-pregnant women were recruited to measure the effects of E₂ on T_{Reg}S (Valor *et al.*, 2011). The results of this study showed that E₂ enhances T_{Reg} generation, in accordance to other studies that showed the same (Yates *et al.*, 2010), and increases perforin expression (Valor *et al.*, 2011). Perforin has been postulated as a mechanism of T_{Reg} *modus operandi*, complete with degranulation as in the granzyme-perforin pathway found in T_{CLS}. It also found that at higher doses, or levels of E₂, such as in pregnancy, T_{Reg} perforin expression is diminished in accordance to the recruitment of T_{Reg}S to the fetal-maternal interface (Valor *et al.*, 2011). This novel finding helps explain the reduced levels of inflammation and demyelination in the E₂-treated mice with more accuracy, and aids in the proposal of a mechanism of action for estrogens in the TMEV model of MS.

B-cells and plasma cells in human MS and in TVID have recently received considerable attention. CD138⁺ cells are commonly attributed the title of mature plasma cells. In the CNS these antibody-secreting cells (ASCs) are part of a secretion system known as intrathecal antibody production (ITAbP) (Pachner *et al.*, 2007). ITAbP ASCs are non-existent in normal CNS tissue, but upon infection with TMEV their numbers increase within months of infection and they begin secreting IgG_{2a} class anti-TMEV antibodies to try to eliminate the virus (Pachner *et al.*, 2007). Mice infected with the BeAn strain of TMEV showed that the levels of antibody were high in cerebrospinal fluid (CSF), and that the ITAb-producing cells were found around or in close proximity to blood vessels (Pachner *et al.*, 2011). However, repeated exposure may result in cross-reaction with myelin components, namely galactocerebroside (Sato *et al.*, 2011). In this pro-inflammatory milieu mediated by T_H1 responses, the principal cytokine interferon gamma (IFN- γ) helps promote IgG_{2a} secretion by plasma cells (Dal Canto *et al.*, 2000; Nicot, 2009; Peterson *et al.*, 1992; Tsunoda *et al.*, 2005). Antibody titers measured by

enzyme-linked immunosorbent assay (ELISA) in SJL/J mice infected with TMEV show that IgG_{2a} is preferentially expressed between days 29 and 108 pi. The CNS in these mice is under T_H1-mediated immunity, thus IgG_{2a} production is regulated by IFN γ . In TMEV-resistant C57BL/6 mice, the T_H2 environment regulates IgG₁ production via IL-4 (Miller *et al.*, 1995; Peterson *et al.*, 1992). In addition, these cells were found to be in locations that paralleled demyelinating lesions, predominantly in ventral white matter areas of the spinal cord (Pachner *et al.*, 2011). As mentioned before, E₂ and E₃ are known to switch the T_H1-mediated immunity to T_H2-mediated (Giraud *et al.*, 2010; Nicot, 2009). We used a Radio-Immuno Assay (RIA) using protein-A-I¹²⁵-labeling to measure the amount of TMEV (GD-VII) Antibody in mouse sera. We expected to see a difference between the estrogen-treated groups and placebo-treated group. We did not find significant differences between E₂ and E₃-treated mice at any dilution; however, both groups were significantly different from placebo-treated mice. These results agree with the observations made by Peterson *et al.* as our experiments show that E₂ and E₃ both cause a shift to T_H2-mediated immunity and IgG₁, while placebo-treated mice stay on the T_H1-mediated milieu and secrete IgG_{2a} (Peterson *et al.*, 1992).

5. CONCLUSION AND FUTURE DIRECTIONS

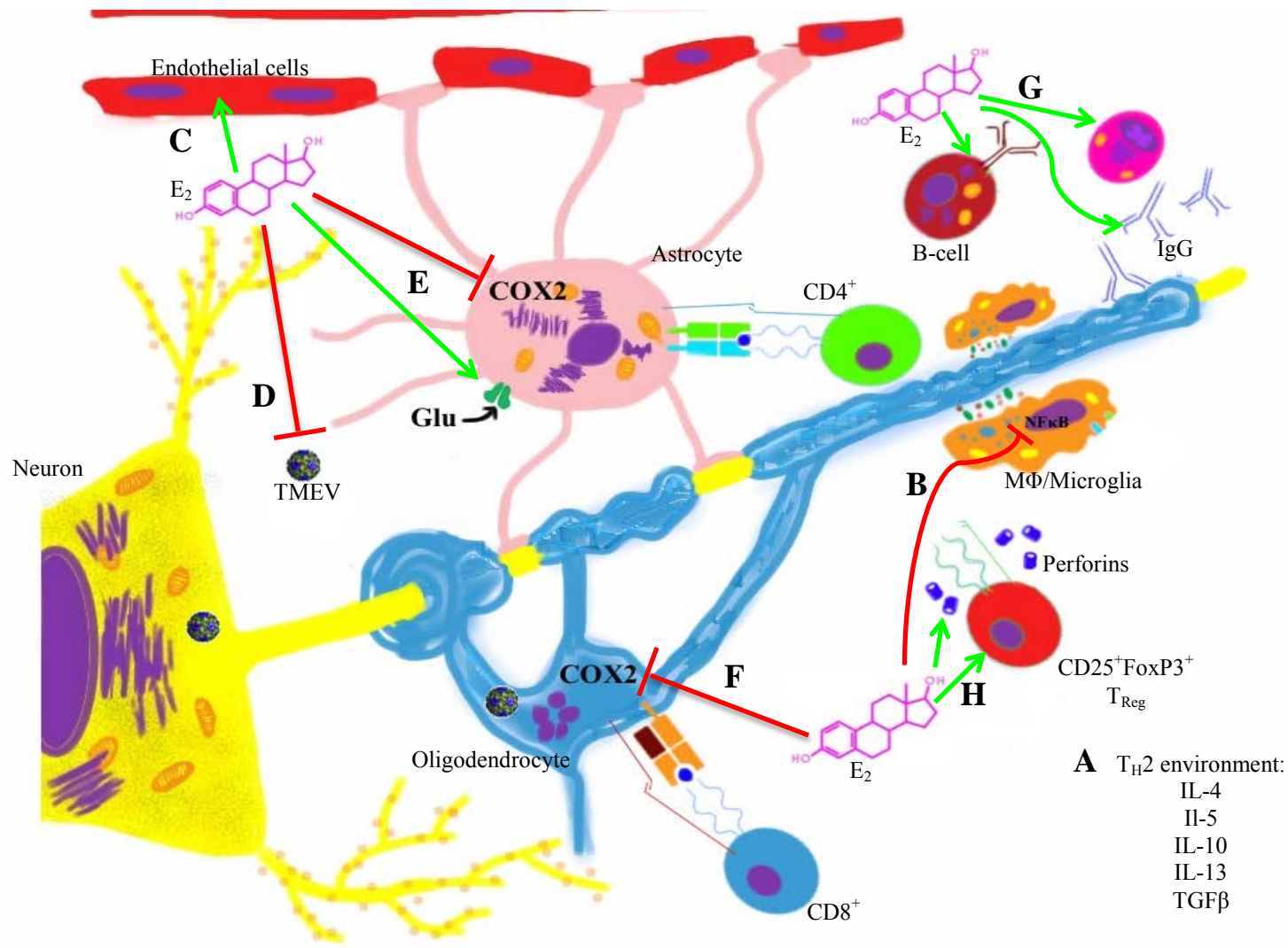
Our study concludes that the use of estrogens as potential therapeutic agents in a murine model of multiple sclerosis is plausible in early disease or as a preventive treatment (Vegeto *et al.*, 2008). The use of estrogens, in particular 17- β -estradiol ameliorated clinical symptoms, reduced signs of inflammation and demyelination, and reduced TMEV antibody counts. This effect is supported by observations made by other research groups. Treatment with estriol is not helpful in ameliorating symptoms. Though it seems to reduce inflammation mildly and diminish TMEV antibody counts, it does not seem to help with demyelination. Though these results are in disagreement with other experimental models of MS, there may be differences in the pathogenic mechanisms of TMEV that can account for our findings. Figure 5.1 presents a proposed mechanism for the beneficial effects of estradiol on TMEV.

Future studies will need to explore the populations of T_H-cells present in the lesions, explore the cytokine environments present in the different treatment groups and confirm the role of T_{RegS} in this model. Studies of the brain in TMEV have not been conducted, thus it would be important to look at the brains in the different treatment groups and assess inflammation and demyelination to create a neuroanatomical model for the pathogenesis of Theiler's Murine Encephalomyelitis Virus. In addition further analysis of cytokine profiles in the experimental groups may provide potential mechanisms of action of estrogens in modulating the immune response and protecting inflammatory demyelination. Studies using hormones such as progesterone for the treatment of TVID may also be in order. Finally, to assess whether hormone treatments may be effective in males, studies with male mice should be carried out.

Figure 5.1 Proposed mechanism of action of estradiol in TVID.

We propose that estrogens, in particular 17- β -estradiol (E_2), reduce inflammation and demyelination via interaction with estrogen receptors (ERs) in cells of the CNS. The result is a decrease in clinical scores and severity of disease.

- A) E_2 induces a switch from a T_H1 (pro-inflammatory) to a T_H2 (anti-inflammatory) cytokine environment. Interleukin (IL)-4, IL-5, IL-10, IL-13, and TGF β are cytokines produced in this environment and they inhibit macrophage activity and modulate T-cell responses
- B) E_2 prevents the pro-inflammatory p65 transcription factor of the (NF- κ B family) from entering the nucleus to promote transcription of inflammatory genes.
- C) E_2 restores the impermeability of the blood-brain barrier (BBB) following ovariectomy and prevents leukocyte infiltration into the CNS.
- D) E_2 can inhibit viral replication of TMEV which prevents direct lysis or infection of other cells.
- E) E_2 interacts with ERs in astrocytes inducing the reuptake of glutamate (Glu) reducing its excitotoxic effects. It also reduces the expression of COX2 in astrocytes reducing the production of prostanoids
- F) COX2 expression in oligodendrocyte has been associated with cell death. E_2 inhibits COX2 production in Oligodendrocytes, preventing death and demyelination.
- G) E_2 induces a switch in immunoglobulin (Ig) class in both B-cells and plasma cells. IgG $_1$ has been associated with a T_H2 response and less myelin damage due to antibodies.
- H) E_2 induces T_{Reg} generation and perforin production, which may be the mechanism by which these cells regulate autoimmune responses



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