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Estrogen in cycling rats alters gene expression in the temporomandibular joint, trigeminal ganglia and trigeminal subnucleus caudalis/upper cervical cord junction

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Abstract

Females report temporomandibular joint (TMJ) pain more than men and studies suggest estrogen modulates this pain response. Our goal in this study was to determine genes that are modulated by physiological levels of 17β -estradiol that could have a role in TMJ pain. To complete this goal, saline or complete Freund's adjuvant was injected in the TMJ when plasma 17β -estradiol was low or when it was at a high proestrus level. TMJ, trigeminal ganglion and trigeminal subnucleus caudalis/upper cervical cord junction (Vc/C₁₋₂) tissues were isolated from the treated rats and expression of 184 genes was quantitated in each tissue using real time PCR. Significant changes in the amount of specific transcripts were observed in the TMJ tissues, trigeminal ganglia and Vc/C₁₋₂ region when comparing rats with high and low estrogen. GABA A receptor subunit $\alpha 6$ (Gabra6) and the glycine receptor $\alpha 2$ (Glr $\alpha 2$) were two genes of interest because of their direct function in neuronal activity and a greater than 29 fold increase in the trigeminal ganglia was observed in proestrus rats with TMJ inflammation. Immunohistochemical studies showed that Gabra6 and Glr $\alpha 2$ neuronal and not glial expression increased when comparing rats with high and low estrogen. Estrogen receptors α and β are present in neurons of the trigeminal ganglia, whereby 17β -estradiol can alter expression of Gabra6 and Glr $\alpha 2$. Also, estrogen receptor α (ER α) but not ER β was observed in satellite glial cells of the trigeminal ganglia. These results demonstrate that genes associated with neurogenic inflammation or neuronal excitability were altered by changes in the concentration of 17β -estradiol.

Keywords

TMJ; Inflammation; 17β -estradiol benzoate; Genes

Introduction

Females report temporomandibular joint (TMJ) disorders more frequently than males (Dworkin et al., 1990; LeResche, 1997). The cycling of hormones during the menstrual cycle is likely a factor in the TMJ nociceptive response. In a study comparing male and female patients, cycling females consistently reported more TMJ pain when the concentration of gonadal hormones were diminishing, such as at menstruation (LeResche et al., 2003). At menopause when estrogen is no longer present there is a reduction in the number of women reporting TMJ pain (LeResche et al., 1997). Notably, the attenuation of symptoms in post-menopausal women is reversed when they are reintroduced to cycling

estrogen or take pharmacological amounts of estrogen (LeResche et al. 1994; LeResche et al., 1997). During pregnancy when gonadal hormone concentrations are much higher than what is observed during the menstrual cycle the reported TMJ pain levels decreased (LeResche et al., 2005) suggesting the concentration of hormone is important.

During the estrous cycle gonadal hormones, such as estrogen, have been shown to modulate TMJ nociceptive responses in animals. When the plasma estrogen concentration reached a maximum (i.e., proestrus, 60–70 pg/ml), Fos staining increased in the TMJ nociceptive pathway as compared to rats having a low (i.e. diestrus) level of estrogen (Bereiter, 2001). In contrast, administration of 17 β -estradiol in a non-cyclical manner, at the same concentration (<70 pg/ml), does not enhance Fos staining, amino acid release nor electrical activity in the TMJ nociceptive pathway suggesting that cyclical administration of 17 β -estradiol is important to the response (Bereiter and Benetti, 2006; Okamoto et al., 2008; Tashiro et al., 2008; Tashiro et al., 2007). Using behavior as an assay for TMJ nociception, researchers showed that increased 17 β -estradiol can be associated with reduced TMJ nociception in cycling rats (Fischer et al., 2009). Consistent with the idea that cycling estrogen is a factor in the TMJ nociceptive response results from our lab demonstrate that a persistent behavioral response, induced by injection of CFA into the rat TMJ, was significantly reduced upon increasing the level of 17 β -estradiol (i.e. proestrus) (Kramer and Bellinger, 2009), whereas during a period low estrogen (i.e., diestrus) the response increased in female rats. The cyclical administration of estrogen can be an important factor but the concentration of estrogen is also a factor in the hormonal response as shown by studies were administration of estrogen in an acute, non-cyclical manner altered the nociceptive response (Favaro-Moreira et al., 2009; Fischer et al., 2008).

We hypothesized that changes in plasma estrogen can modulate TMJ pain, in part, through altering gene expression in tissues required for pain transmission such as the joint, trigeminal ganglion and/or Vc/C₁₋₂ region. To date, little is known about the effect of estrogen has on genes that can modulate TMJ pain. In rats with TMJ inflammation, we identify genes that function in immune or pain responses whose expression was effected by different concentrations of 17 β -estradiol present in a cycling female rat.

Materials and methods

These studies were approved by the Baylor College of Dentistry Institutional Animal Care and Use Committee in accordance of the guidelines of the USDA and National Institutes of Health Guide for Care and Use of Laboratory Animals. A portion of the female Sprague Dawley rats (~250 gm) were ovariectomized by Harlan Industries, Houston TX. Upon arrival, the rats were housed individually and kept on a 14:10 h light/dark cycle with lights on at 0600 h.

Estrogen treatment-artificial estrous cycle

Ovariectomized rats were anesthetized with ketamine (52 mg/kg) and xylazine (5.4 mg/kg). Using sterile technique, the rats were implanted with primed, 28-day Alzet mini-osmotic pumps dispensing the vehicle polyethylene glycol (low E2) or 750 ng/day of 17 β -estradiol benzoate (high E2). Prior to insertion, the alzet pumps are placed into warm physiological saline so the pumps deliver estrogen immediately on implantation. To simulate the proestrus surge the high E2 rats were injected with 2.5 μ g of 17 β -estradiol benzoate in 0.1 ml sesame oil subcutaneously every fifth day (Kramer and Bellinger, 2009). The low E2 rats received a subcutaneous injection of sesame oil with no estrogen.

Vaginal smear protocol

Female rat's vagina was lavaged twice a day at 08:00 h and at 15:00 h using 250 μ l of sterile 0.9% saline, and the solution was then transferred to a glass slide (StatLab, Inc., Lewisville, TX). The slides were completely dried and then fixed and stained using a Hema-Diff Rapid Differential Stain kit from Anapath (StatLab Inc.). After staining, the cell morphologies were observed with a microscope and recorded.

CFA injections

The low E2 and high E2 groups were subdivided such that half the rats were given a bilateral injection (15 μ g) of complete Freund's adjuvant (CFA) and half were given an injection of 0.9% saline (Sal) into the superior joint space of TMJ. The low E2 and high E2 rats were given a TMJ injection in the early phase of the cycle, as determined by charting the cycle by vaginal smears for the previous two weeks. Rats in the low E2 and high E2 groups were given the TMJ injection one hour after the subcutaneous injection of vehicle or 17 β -estradiol benzoate. The four groups included: low E2 + SAL, low E2 + CFA, high E2 + SAL and high E2 + CFA. 24 hours after the last injection the animals were euthanized.

Sample and tissue preparation

The day following TMJ saline or CFA injections, the treated rats were taken to an adjacent room, sacrificed by decapitation, blood was collected for radio immunoassays (RIA) and the heads submerged in an ice bath until dissection. To minimize the release of corticosterone and the subsequent effect of corticosterone on cytokines levels decapitation was necessary (Schultz et al., 1979; Vahl et al., 2005).

The dissected TMJ tissue included the synovial membrane, joint capsule, retrodiscal tissue, articular disc, and a small amount of the lateral pterygoid muscle. The trigeminal ganglia from V1 to 1 mm below V3 and the dorsal portion (i.e., upper lamina layers) of the caudalis/upper cervical region from the obex to the C2 rootlets were dissected from each animal. After dissection, the tissues were placed in liquid nitrogen or fixed in 5% paraformaldehyde and stored in a 25% sucrose solution. Three animals per treatment group were used for each of the procedures.

Quantitation of hormone levels

17 β -estradiol was quantitated in the artificially cycling rats in the blood plasma by first, vigorously mixing 0.5 ml of plasma with 5 ml of ether. Second, the solution was frozen in an ethanol, dry ice bath and the ether phase decanted. The ether phase contains the lipid soluble hormones. The ether was evaporated and the remaining residue suspended in RIA buffer supplied by the manufacturer. The concentration of 17 β -estradiol was quantitated following the manufacturer's directions (ICN Biomedicals, Costa Mesa, CA).

RNA Isolation and Purification

Frozen tissue was homogenized in Trizol reagent (Invitrogen, Carlsbad, CA) which is composed of phenol and guanidine isothiocyanate (2 ml Trizol/30–50 mg tissue). After incubating at room temperature for 5 min, 0.2 ml of chloroform was added per ml of homogenate, the samples mixed and the aqueous phase decanted. RNA was precipitated from the aqueous phase by mixing 0.5 ml of isopropanol per ml of aqueous phase. After a 5 min of incubation at room temperature, the mix was centrifuged at 12,000 g for 15 minutes at 4 $^{\circ}$ C. The supernatant was removed, the RNA pellet washed with 75% ethanol and briefly air-dried. The pellet was dissolved in 0.5 ml of RNase-free water and further purified using

an RNeasy Midi Kit (Qiagen, Valencia, CA). The yield was analyzed using a spectrophotometer (absorbance of 260/280 nm) and the quality of the total RNA was assessed using an Agilent 2100 Bioanalyzer (Agilent Technology, USA). cDNA was synthesized from the total RNA using a RT² First Strand kit following the manufactures directions (Qiagen).

Quantitative gene expression analysis

Gene expression was quantified in the cDNA samples by real time PCR using Superarray RT² ProfilerTM PCR Arrays. Gene expression was analyzed using the cDNA isolated from three replicate animals within each treatment group (low E2 + SAL, low E2 + CFA, high E2 + SAL and high E2 + CFA) from each of the three tissues (TMJ, trigeminal ganglia, Vc/C₁₋₂). The genes that were studied included 91 inflammatory cytokines and receptors (catalog #PARN -011A, Superarray) and 93 additional genes placed on a custom RT² ProfilerTM PCR Array. The genes that were placed on the custom PCR array were determined by a pilot experiment of a whole genome array (>10,000 genes) that determined changes in gene expression from pooled samples of each treatment group listed above (data not shown). PCR reactions were completed on a MX 4000 multiplex quantitative PCR system. For a listing of all the genes analyzed on the PCR arrays see supplemental figures (Fig. 1S through 10S), a summary of the supplemental figures are listed in Table 1.

Immunostaining

Tissues stored in 25% sucrose were frozen, cryosectioned and the 30 μ m sections placed on Histobond slides (VWR international). The tissue was then blocked with a PBS solution containing 5% normal goat serum and 0.3% Triton-X 100 for 2 hours at room temperature. Following three rinses in PBS with 0.3% Tween-20 the slides were incubated in a primary antibody solution overnight at 4°C. Primary antibodies consisted of a rabbit ER α antibody (Santa Cruz, 1:50), a rabbit ER β antibody (ABR, 1:100), a neuronal marker (NeuN) antibody (mouse-Millipore, 1:1000), a satellite glial marker glutamine synthetase antibody (mouse-Millipore, 1:300), a Gabr α 6 antibody (rabbit-Millipore, 1:1000) and a Glr α 2 antibody (rabbit-GeneTex Inc., 1:500). Primary antibodies were diluted with PBS, 5% BSA and 0.3% Triton X-100. After incubation in primary antibody the slides were rinsed three times in PBS and Tween-20 for a total of 60 min and placed for 2 hours in a 1:500 dilution of secondary antibody in PBS and 0.3% Triton X-100. Secondary antibodies included goat anti-rabbit 488 and goat anti-mouse 633 (Invitrogen, 1:500). After rinsing the slides three times in PBS for a total of 60 min, the slides were then mounted with Fluoromount-G mounting medium (Electron Microscopy Sciences, Hatfield, PA). The fluorescent signal was imaged using a Nikon fluorescent microscope and NIS-Elements imaging software and a Photometrics CoolSnap K4 CCD camera (Roper Scientific, Inc, Duluth, GA).

Calculation and statistical analysis

The change in gene expression was calculated first, by determining the Δ Ct value (the change in cycle threshold value) for each gene in each replicate sample using the formula Δ Ct = Ct (GOI) – avg. (Ct (HKG)), where GOI is each gene of interest, and HKG are the housekeeping genes. The Δ Ct value was calculated for each replicate, in each treatment group. The five house-keeping genes included; ribosomal protein P1 (NM_001007604), hypoxanthine guanine phosphoribosyl transferase (NM_012583), β -actin (NM_031144), ribosomal protein L13A (NM_173340) and the lactate dehydrogenase A gene (NM_017025). Second, the relative expression level for each gene was determined by the formula $2^{(-\Delta$ Ct). The relative expression level for the No E2 group was divided by the E2 treatment group to provide the fold increase or decrease in gene expression, values less

than one were converted by the formula $(-1/X)$ to indicate the amount of down regulation in an integer format. Fold changes less than 2 were not reported in the results. To determine how the PCR data was distributed we performed real time PCR on nine replicates for each treatment group, using two of the genes present on the PCR arrays, one cytokine gene (i.e., IL1 β) and one was an Fc receptor gene (i.e., CD16). This data had no appreciable skewness (0.2) or kurtosis (-0.4) (SPSS Software system 17.0) providing validity of parametric assumptions, such as homogeneity of variance and normality. Mean Δ Ct values for each treatment group was analyzed by an unpaired t-test. P-values of < 0.05 were considered to be statistically significant.

Results

Estrogen effects transcript expression

To define the genes and location by which estrogen influences TMJ nociceptive responses (Kramer and Bellinger, 2009), we looked at 17 β -estradiol induced changes in gene expression within the TMJ, trigeminal ganglion and Vc/C₁₋₂. Plasma 17 β -estradiol was low (7–10 pg/ml) in the animals that received vehicle, consistent with previous work (Butcher et al., 1974). This low level of estrogen can be produced from adrenal hormones and fat tissue. A higher concentration of 17 β -estradiol (60–70 pg/ml) was present in the rats that had an Alzet pump releasing estrogen and were given injections of 17 β -estradiol benzoate to produce a natural estrous cycle as determined by vaginal smears (data not shown). A plasma concentration of 60–70 pg/ml 17 β -estradiol is consistent with a proestrus level of estrogen (Butcher et al., 1974).

Animals with low and high (i.e., proestrus) plasma estrogen were given saline or CFA injections into the superior joint space of the TMJ, 24 hours following this injection the animals were sacrificed. Real time PCR was performed with RNA isolated from three different tissues (TMJ, trigeminal ganglia and Vc/C₁₋₂) isolated from the treated rats. The real time PCR results in the treated rats were categorized in two sections; one section was related to the immune response and the second was related to neurons and hormones.

Chemokines, cytokines and immune genes

Retrodiscal tissue Low E2 + SAL versus High E2 + SAL group: Only one chemokine was affected by a proestrus concentration of 17 β -estradiol in the non-inflamed TMJ (Fig. 1S) that was the chemokine Cxcl2, which increased significantly (Fig. 1A).

Retrodiscal tissue Low E2 + CFA versus High E2 + CFA group: When comparing rats with an inflamed TMJ that had low 17 β -estradiol to the rats that had a high proestrus level of 17 β -estradiol (Fig. 2S) the chemokine Cxcl11 was expressed at a 7 fold greater concentration (Fig. 1B) whereas chemokine Ccl25 was significantly less in high E2 (i.e., proestrus) rats with an inflamed TMJ joint (Fig. 1C). Interleukin receptor IL8rb (IL-8R β) was upregulated by 6 fold in the high E2 group (Fig. 1D).

Trigeminal ganglia Low E2 + SAL versus High E2 + SAL group: Several genes with immune function were affected by proestrus levels of 17 β -estradiol (Fig. 3S and 5S). When comparing rats with low 17 β -estradiol to rats having a high proestrus level of 17 β -estradiol Ccr5 and Cfh were expressed at a 2 and 7 fold greater level, respectively (Fig 2A and 2B). In contrast, 17 β -estradiol significantly decreased expression of Ccr2 (Fig. 2C).

Trigeminal Ganglia Low E2 + CFA versus High E2 + CFA group: In the trigeminal ganglion (Fig. 4S and 6S), we found that expression of the cytokine Tnfb (TNF- β), Tnfb was significantly greater in the high E2 (i.e., proestrus) rats with an inflamed TMJ (Fig. 2D). In

contrast, *Ccl20*, *Ccl21b*, *Ccl4* and *IL-1ra* were significantly down regulated by 51, 16, 4, and 5 fold, respectively (Fig 2E–H). The IgA Fc receptor and *Mmp9* genes were also significantly down regulated by 23 and 10 fold, respectively (Fig 2I and 2J) and a large 930 fold decrease in *Tlr10* was observed in the trigeminal ganglia of high E2 rats with an inflamed TMJ (Fig. 2K)

Vc/C2 Low E2 + SAL versus High E2 + SAL group: When comparing low E2 + SAL and high E2 + SAL groups (Fig. 7S and 9S) the transcript levels for *Ccl2* and *IL10* were reduced whereas the amount of *Ccr7*, *IL8ra* and *Ltb* transcript were greater at proestrus (Fig. 3A–E).

Vc/C2 Low E2 + CFA versus High E2 + CFA group: No gene decreased in the Vc/C2 region in the high E2 group (Fig. 8S and 10S) but the amount of *Ccl24*, *Ccr3*, *Ccr4*, *IL13ra1* and *Tgfb1* (*Tgf-β1*) transcript increased in the high E2 group (Fig. 3F–J).

Neurotransmitters and hormones—A custom RT² Profiler™ PCR Arrays was not performed for the TMJ tissues after preliminary studies with the whole genome array showed that no change in expression of genes on this array had occurred in response to estrogen (data not shown).

Trigeminal ganglia Low E2 + SAL versus High E2 + SAL group: In a non-inflamed joint, the amount of neurotransmitter receptors *Gabra6* (34 fold) and *Glrα2* (29 fold) increased significantly at in the high E2 proestrus group (Fig 4A and B).

Trigeminal Ganglia Low E2 + CFA versus High E2 + CFA group: In rats with an inflamed joint *Gabra6* expression significantly increased in the high E2 group (Fig. 4C). *Cacna1e* (*Cacna1e*) expression also increased after estrogen treatment (Fig 4D) but olfactory receptor 588 and glycoprotein 49b expression was reduced by 45 and 8 fold, respectively (Fig. 4E and 4F). Growth hormone 1 (*Gh 1*) expression increased nearly 48 fold (Fig. 4G) and a 50 fold increase in the prolactin (*Prl*) gene was observed after administration of 17β-estradiol, consistent with previous results (Diogenes et al., 2006) (Fig. 4H).

Vc/C2 Low E2 + SAL versus High E2 + SAL group: There were no significant findings observed in the subnucleus caudalis tissue samples after treatment with 17β-estradiol (see supplemental Figs. 7S and 9S).

Vc/C2 Low E2 + CFA versus High E2 + CFA group: A small, although significant, increase in *Fgf18* (Fig. 5) was observed in the Vc/C2 after administering estrogen to rats with an inflamed TMJ.

Estrogen increased *Gabra6* and *Glrα2* protein levels in the trigeminal ganglia

Further analysis of *Gabra6* and *Glrα2* expression was of interest because of their direct function in neuronal activity and because the transcript levels of these two genes changed more than 29 fold (Fig. 4). *Gabra6* and *Glrα2* expression increased in proestrus rats (compare Fig. 6A to B, C to D, E to F and G to H), consistent with the increase in transcript observed in the high E2 group (Fig. 4). CFA injection did not appear to effect expression significantly for *Gabra6* (Fig. 6, compare A to C and B to D) or *Glrα2* (Fig. 6, compare E to G and F to H). In cycling rats the expression of *Gabra6* (Fig. 6A–D) and *Glrα2* (Fig. 6E–H) co-localized (yellow) with the neuronal marker NeuN (red). Because NeuN is expressed in the nuclei and the cytoplasm of most neurons the data suggested these proteins are expressed in neurons. Expression was observed in both small (<30 μm) and medium to large neurons (>30 μm) (Fig. 6A–H).

Low amounts of *Gabra6* (Fig. 7A–D) and *Glr2* (Fig. 7E–H) co-localized (yellow) with glutamine synthetase, a marker for satellite glia surrounding the neurons. This result supports the idea that these proteins are expressed primarily in neurons. Proestrus increased the amount of *Gabra6* (compare Fig. 7A to B and C to D) and *Glr2* (compare Fig. 7E to F and G to H). CFA injection had little effect on *Gabra6* (compare Fig. 7B and D) and *Glr2* (compare Fig. 7F and H) expression.

ER α or ER β in trigeminal ganglia neurons and satellite glia

To effect gene expression using a genomic mechanism the cells in the trigeminal ganglia would need to express ER α or ER β . Both small (<30 μ m) and medium to large neurons (>30 μ m) in the trigeminal ganglia express ER α (Fig. 8A, yellow) and ER β (Fig. 8C, yellow), ER α was also expressed in satellite glia of the trigeminal ganglia (Fig. 8B, yellow). Little ER β was observed in the satellite glia (Fig. 8D).

Discussion

Estrogen likely modulates pain sensitivity via altering mechanisms both peripherally and centrally (Duval et al., 1996; Fillingim and Ness, 2000; Smith, 1994). Previous studies have shown that rats in proestrus (i.e., high estrogen) have reduced behavioral nociceptive markers as compared to rats in diestrus (i.e., low estrogen) and that 17 β -estradiol is a contributing factor to this change (Kramer and Bellinger, 2009). To evaluate gene expression in regions that modulate TMJ nociceptive responses, changes in selected genes were analyzed at the transcript and protein level using real time PCR. This analysis was completed in rats having lower and higher amounts of 17 β -estradiol in their plasma. 17 β -estradiol was shown to have an effect on the expression of several genes in the female rat TMJ, trigeminal ganglion and Vc/C2 tissues. These genes were often associated with neurogenic inflammation and neuronal excitability.

Gene transcripts regulated with 17 β -estradiol: Inflammatory cytokines and chemokines

Numerous chemokines and cytokines were affected in the TMJ, trigeminal ganglion and Vc/C2 tissues by changes in 17 β -estradiol. Cytokines are small secreted proteins which mediate and regulate immunity and inflammation (Baggiolini et al., 1997). Chemokines, a subfamily of cytokines with chemotactic properties, have shown to express their receptors throughout the nervous system (Miller et al., 2008). Moreover, release of cytokines and chemokines are known to occur from microglia, satellite glia and astrocytes to effect neurogenic inflammation (Gao et al., 2009; Garrett and Durham, 2008; Gosselin et al., 2008; Gosselin et al., 2010; Gosselin et al., 2005; Vause and Durham, 2010). At the transcript level, we found that *Cxcl2*, *Cxcl11*, *IL8rb* (aka. *Cxcr2*) were upregulated in the TMJ tissue whereas; *Ccl25* was significantly downregulated in the retrodiscal tissue when 17 β -estradiol was high (i.e. proestrus). Consistent with our results, previous studies have shown that *Cxcl11* expression will increase after estrogen treatment (Sentman et al., 2004). Studies have also shown a differential chemokine effect on calcium signaling (Oh et al., 2002; Qin et al., 2005) which is known as an important secondary messenger in primary sensory neurons (Ghosh and Greenberg, 1995). Disrupted Ca²⁺ signaling is a significant feature in peripheral nerve injury induced pain models (Fuchs et al., 2005; Fuchs et al., 2007; Gemes et al., 2010; Hogan et al., 2000). In our study we found that with increased 17 β -estradiol, IL-8R β expression was upregulated and other have demonstrated that upregulation of IL-8R β causes Ca²⁺-regulated opioid release from polymorphonuclear cells and thereby inhibits inflammatory pain (Rittner et al., 2006). In addition, studies have also shown other cytokines such as IL-6 (Torres-Chavez et al., 2010; Yun et al., 2008), IL-1 β and IL-8 (Yun et al., 2008) are modulated in the TMJ tissues after estrogen administration. Our study did not observe any affect on these cytokines. The differences in the action of estrogen on these cytokines could be explained

due to the larger dosage of estrogen used in the previous studies versus the lower dosage used in this report. Also, differences in the models (e.g., inflammatory agent and species of animal) could have influenced the genes that were modulated by estrogen treatment.

In the trigeminal ganglion, we found that *Ccr2* expression was reduced but that of *Cfh* and *Ccr5* expression was increased in high E2 rats. Consistent with our result, a study by the Pitteri lab showed *Cfh* expression increases after 17 β -estradiol treatment (Pitteri et al., 2009). Expression of chemokines *Ccl20*, *Ccl21b*, *Ccl4* were significantly down regulated in proestrus (high E2) rats with an inflamed TMJ, consistent with previous work showing *Ccl20* expression decreases in the trigeminal ganglia after estrogen treatment (Puri et al., 2006). Many glial cells are present in the trigeminal ganglia and release cytokine and chemokines that contribute to neurogenic inflammation (Cherkas et al., 2004; Chudler et al., 1997; Dublin and Hanani, 2007; Stephenson and Byers, 1995). A reduction in these molecules suggests estrogen can reduce neurogenic inflammation in the trigeminal ganglia.

Another cytokine TNF- β , also known as lymphotoxin A, is an immediate early gene of the inflammatory response. TNF- β was upregulated with high 17 β -estradiol. Expression of TNF- β increases in TMJ osteoarthritis patients (Vernal et al., 2008) suggesting the involvement of TNF- β in inflammation. Further investigations are required to understand effects of 17 β -estradiol on TNF- β in TMJ pain.

Mmp9 and *Tlr10* transcript levels were significantly reduced in the trigeminal ganglia of rats with high 17 β -estradiol. MMP9 protein has been shown to be involved in the early phase of neuropathic pain after dorsal root ganglia injury (Kawasaki et al., 2008) and *Mmp9* gene deletion has shown to reduce neuropathic pain (Chattopadhyay et al., 2007). MMP9 has also shown to mediate the activation of microglia and the development of neuropathic pain in experimental pain models (Kawasaki et al., 2008). A clinical study has shown that an increase in MMP9 occurs during migraine attacks (Leira et al., 2007), consistent with the idea that estrogen can modulate the pain response by effecting MMP9 release from glial cells in the trigeminal ganglia. Toll-like receptors (TLR) are also expressed in microglia of the central nervous system. Increased expression of TLRs following nerve injury suggests its role in neuropathic pain (Guo and Schluesener, 2007). In our study we found that with increased 17 β -estradiol *Tlr10* expression was reduced; suggesting estrogen could affect the pain response by reducing the expression of *Tlr10*.

In the *Vc/C2* region, *Ccl2* expression was reduced in proestrus (high E2) rats without TMJ inflammation. *Ccl2* can modulate pain responses through several mechanisms. One mechanism by which *Ccl2* can increase pain is by stimulating CGRP release from primary nociceptive neurons of dorsal root ganglion thereby producing pain (Qin et al., 2005). A second mechanism involves glutamatergic currents where *Ccl2* functions as an astrocyte-derived pronociceptive factor (Gao et al., 2009). A third mechanism for the pronociceptive action of *Ccl2* may rely on its potent attenuating effect on inhibitory GABAergic currents in spinal neurons (Gosselin et al., 2005). Reduced levels of *Ccl2* in proestrus rats would reduce the pain response consistent with our previous work (Kramer and Bellinger, 2009).

Expression of *Ccr3* and *Ccr4* increased in the spinomedullary tissue of proestrus (high E2) rats. *Ccr3* and *Ccr4* have been implicated in nociception (Oh et al., 2001) but in this example we expected estrogen to decrease *Ccr3* and *Ccr4* causing a decrease in the nociceptive response noted previously (Kramer and Bellinger, 2009). It is unclear the effect that increased *Ccr3* and *Ccr4* has on the nociceptive response but it may be that the effect on these chemokine receptors were counteracted by the effect of estrogen on other cytokine and chemokine genes.

Gene transcripts regulated with 17 β -estradiol: Neurotransmitters and other genes

Gabra6 increased in the trigeminal ganglia of proestrus rats with and without TMJ inflammation. Glra2 was also upregulated in the proestrus, non-inflamed rats consistent with a study showing 17 β -estradiol selectively modulates various glycine receptors (Jiang et al., 2009; Meier et al., 2005). This observation was at the transcript and protein level. Glra2 transcript increased 14 fold in rats with TMJ inflammation but the difference was not significant (60% power, $\alpha=0.05$). GABA and glycine receptors are known for mediating inhibitory neurotransmission in the spinal cord and brain stem (Bohlhalter et al., 1994; Frazao et al., 2007; Maxwell et al., 1995; Ottersen et al., 1988; Popratiloff et al., 1996; Todd et al., 1996). The modulation of glycine receptor content may be the mechanism by which 17 β -estradiol inhibits glycine-activated current in rat hippocampal and spinal dorsal horn neurons (Jiang et al., 2009). Studies have also shown that both the receptors colocalize in spinal cord and their transmitters are released together at nerve endings in the rat (Bohlhalter et al., 1994; Jonas et al., 1998) suggesting estrogen can impact the pain response through changes in receptors specific to inhibitory neurotransmitters. Regions that control of pain transmission express Cacna1e, such as the dorsal root ganglia and the dorsal horn of the spinal cord (Saegusa et al., 2000). Our study showed a significant increase in Cacna1e in proestrus (high E2) rats with an inflamed TMJ. Mice with a deletion of Cacna1e showed increased sensitivity to inflammatory pain as a result of decreased activity in the descending antinociceptive pathways (Saegusa et al., 2000). It is possible the reduced TMJ pain observed at proestrus (Kramer and Bellinger, 2009) was due to the increase in Cacna1e enhancing antinociception.

Studies have shown that estrogen regulates growth hormone (Gh) plasma levels in both rats and humans (Dawson-Hughes et al., 1986; Duursma et al., 1984; Ho and Weissberger, 1990). An impaired growth hormone response leads to increased levels of cytokines and pain severity (Ross et al., 2010). The increase in Gh1 in the trigeminal ganglia of proestrus rats provides us another potential molecule to help unravel the mechanism behind women's sufferings.

Prl expression was increased when the plasma level of 17 β -estradiol was higher, consistent with previous work (Diogenes et al., 2006). PRL has been shown to modulate transient receptor potential cation channel, subfamily V, member 1 (TRPV1) activity in sensory neurons, hence indicating a role PRL in the development of certain pain disorders. The sensitization caused by PRL through TRPV1 contradicts our finding that 17 β -estradiol reduces CFA induced TMJ nociception (Kramer and Bellinger, 2009). One possible explanation for this result is that CFA induces the pain response through a non-TRPV1 dependent mechanism in contrast to a model where the TMJ pain was induced with capsaicin (Diogenes et al., 2006). A few other novel genes such as, Olr588 and Glycoprotein 49b were significantly modulated with estrogen but the role these genes have in the nociceptive response is currently unknown.

ER α and ER β was expressed in neurons of the trigeminal ganglia consistent with previous results (Bereiter et al., 2005; Liverman et al., 2009; Puri et al., 2005). A novel finding was that ER α was also expressed in satellite glia of the trigeminal ganglia but little ER β was observed in the satellite glia. These results are consistent with the idea that estrogen can effect gene expression in both neurons and satellite glia of the trigeminal ganglia through a genomic mechanism. Estrogen's biological effects can be mediated by α and β receptors in the TMJ, trigeminal ganglion or subnucleus caudalis (Bereiter et al., 2005; Puri et al., 2009; Puri et al., 2005; Yamada et al., 2003). In contrast, estrogen effects on pain responses can occur by a non-genomic mechanism. For example, the activation of the nitric oxide-cyclic guanosine monophosphate signaling pathway has been shown to play a role in the antinociceptive action of estradiol (Favaro-Moreira et al., 2009). Future studies would focus

on the mechanism by which estrogen modulated expression of the genes discussed in this study.

To conclude, plasma levels of estrogen in female rats modulate gene expression in the joint, trigeminal ganglion and subnucleus caudalis region. These studies implicate that neurogenic inflammation and changes in neurotransmitter receptors in both the trigeminal ganglia and upper lamina layers of the Vc/C2 region are potential mechanisms by which estrogen influences TMJ pain. The rat estrous cycle, like the human ovarian or menstrual cycle has a maximal level of estrogen during proestrus but we do recognize that after proestrus there is no increase in plasma estrogen in a rat (i.e., metestrus, estrus) in contrast to a gradual increase in estrogen observed during the luteal phase of the menstrual cycle (Butcher, Collins et al. 1974, Evans and Long 1922). These differences in hormonal cycles could impact gene expression and lead to differences between a rat model and humans thus justifying future studies with human samples.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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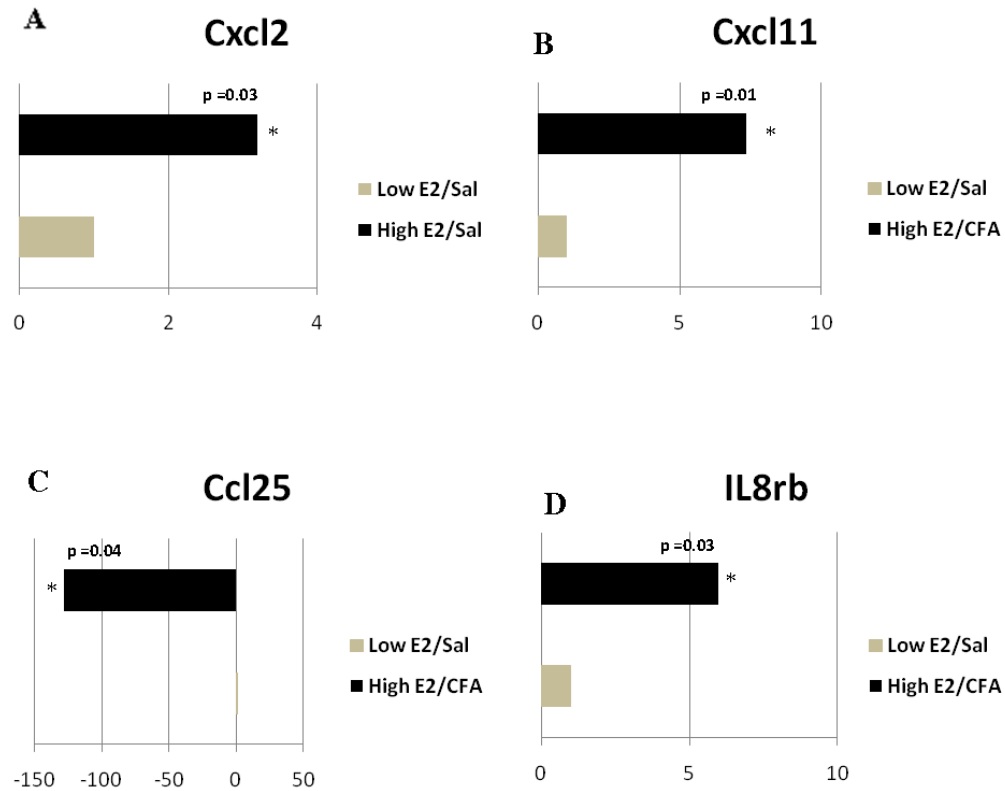
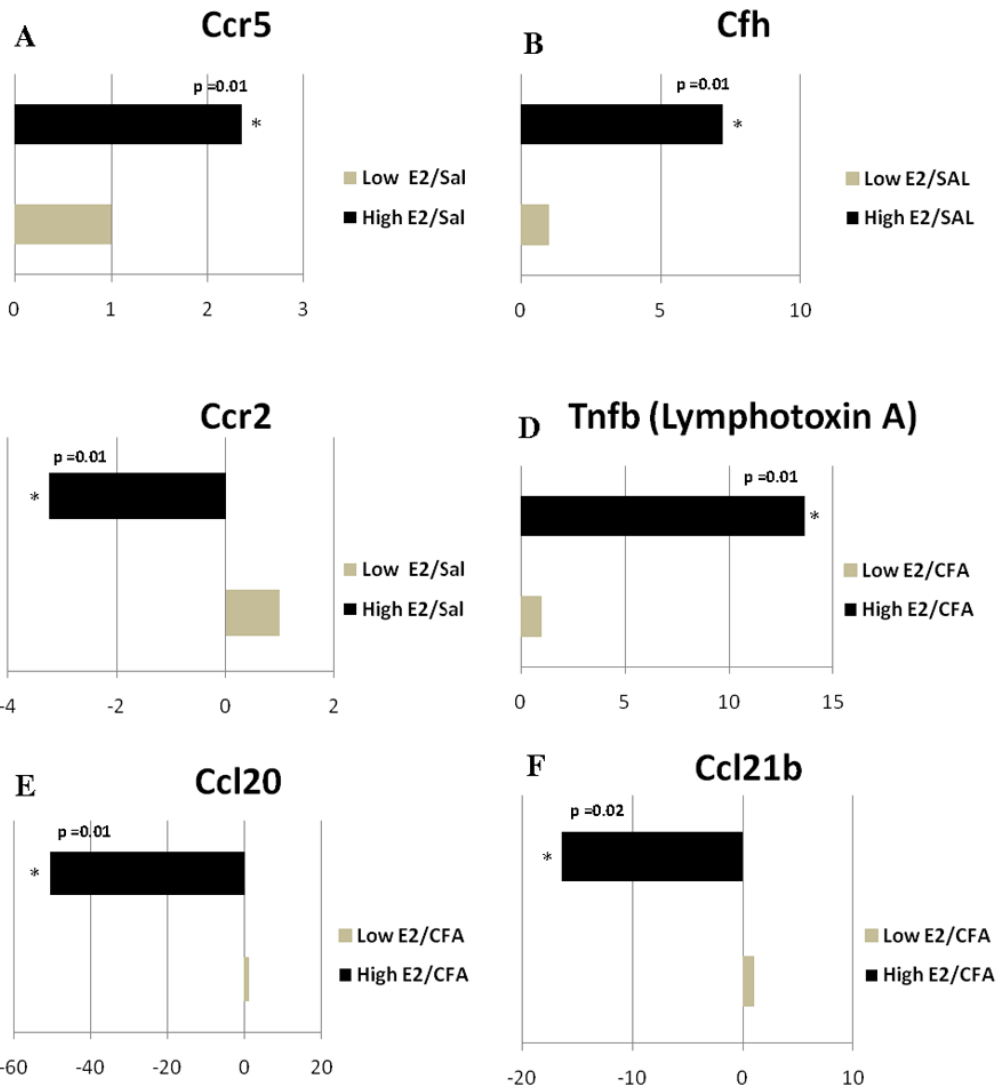


Figure 1. Real time PCR: Histogram showing expression of genes with immune function in the TMJ tissue

The animals were ovariectomized and given vehicle (Low E2) or 17 β -estradiol (High E2) to produce an estrous cycle. The upper joint space of the TMJ was injected with 0.9% Sal or CFA in both the Low E2 and High E2 rats. 24 hrs after TMJ injection the tissue was collected for analysis of gene expression. Expression of A) chemokine (C-X-C) motif ligand 2 (Cxcl2) changed when comparing the Low E2 + SAL and High E2 + SAL groups. B) Cxcl11, C) Chemokine (C-C) motif ligand 25 (Ccl25), and D) interleukin receptor 8 β (IL8rb) changed when comparing the Low E2 + CFA and High E2 + CFA groups. The transcript levels in rat tissue from three independent experiments were determined for each treatment group and statistics were conducted on the change in cycle threshold (Δ Ct) values using a Student's *t*-test. Results were considered statistically significantly when $p < 0.05$. Values are the mean \pm SEM.



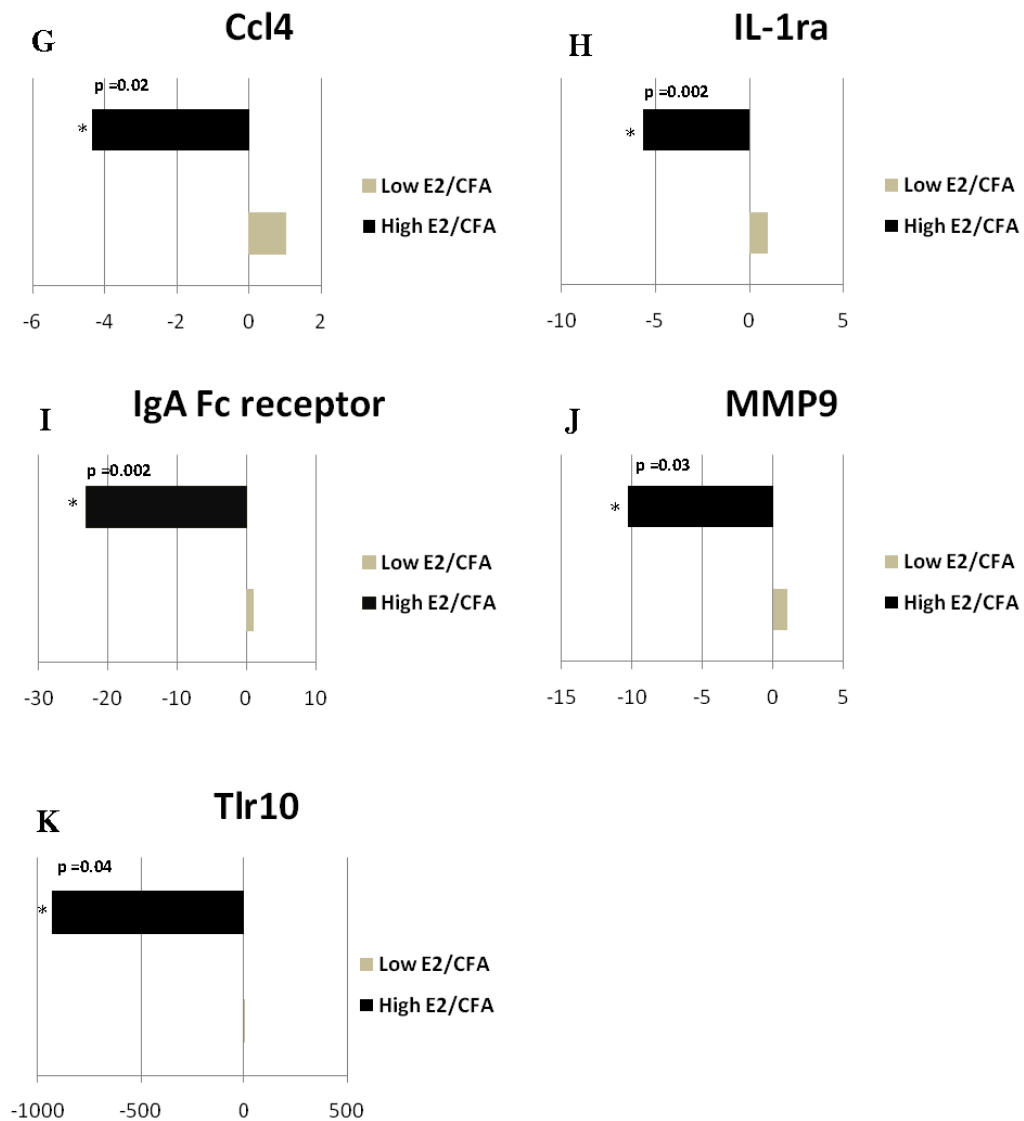
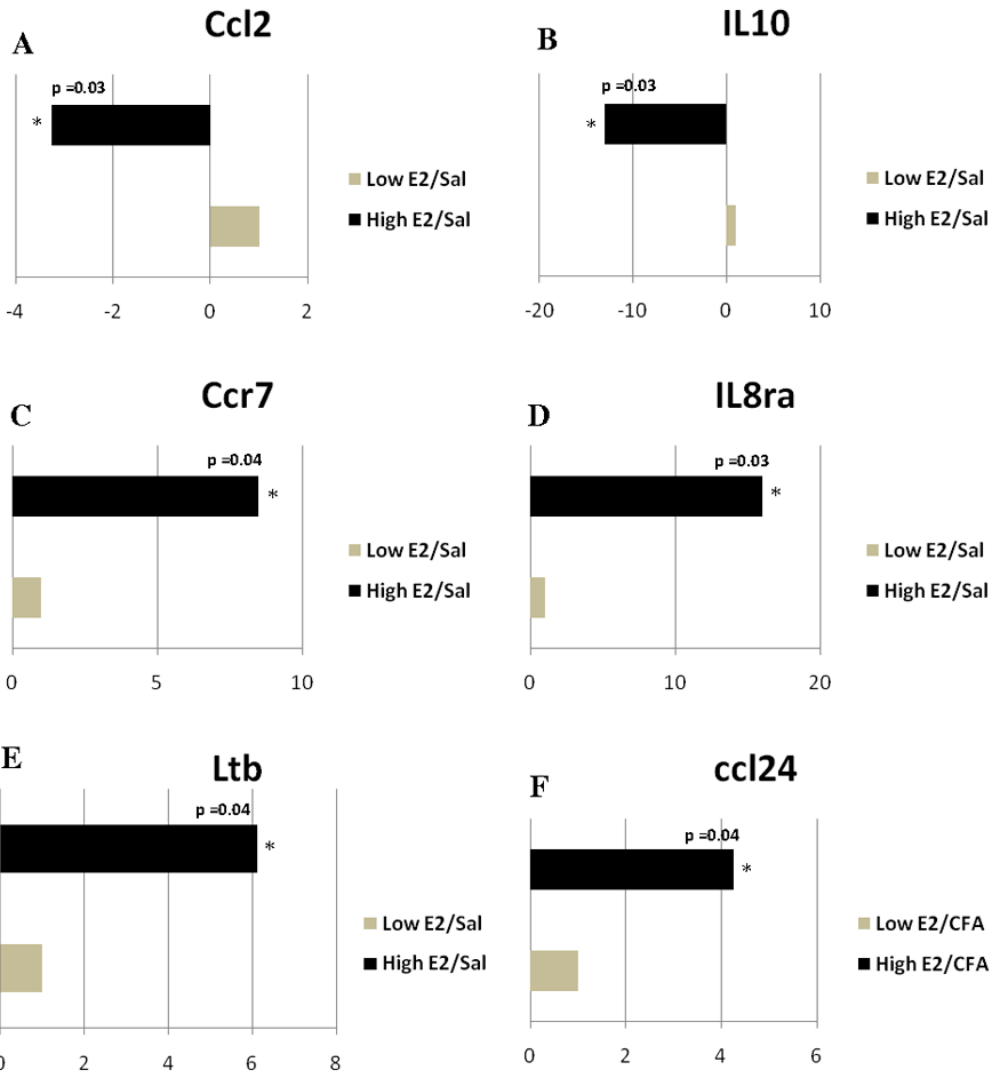


Figure 2. Real time PCR: Histogram showing expression of genes with immune function in the trigeminal ganglia

Expression of A) Ccr5, B) complement factor H (Cfh), C) Ccr2, comparing the Low E2 + SAL and High E2 + SAL groups; D) tumor necrosis factor β (Tnfb), E) Ccl20, F) Ccl21b, G) Ccl4, H) Interleukin-1 receptor antagonist (IL-1ra), I) IgA Fc receptor, J) matrix metalloproteinase 9 (Mmp9), and K) toll-like receptor 10 (Tlr10) changed when comparing the Low E2 + CFA and High E2 + CFA groups. The transcript levels from three independent experiments were collected and values were compared using the Student's *t*-test. The results were considered statistically significantly with $p < 0.05$. Values are the mean \pm SEM.



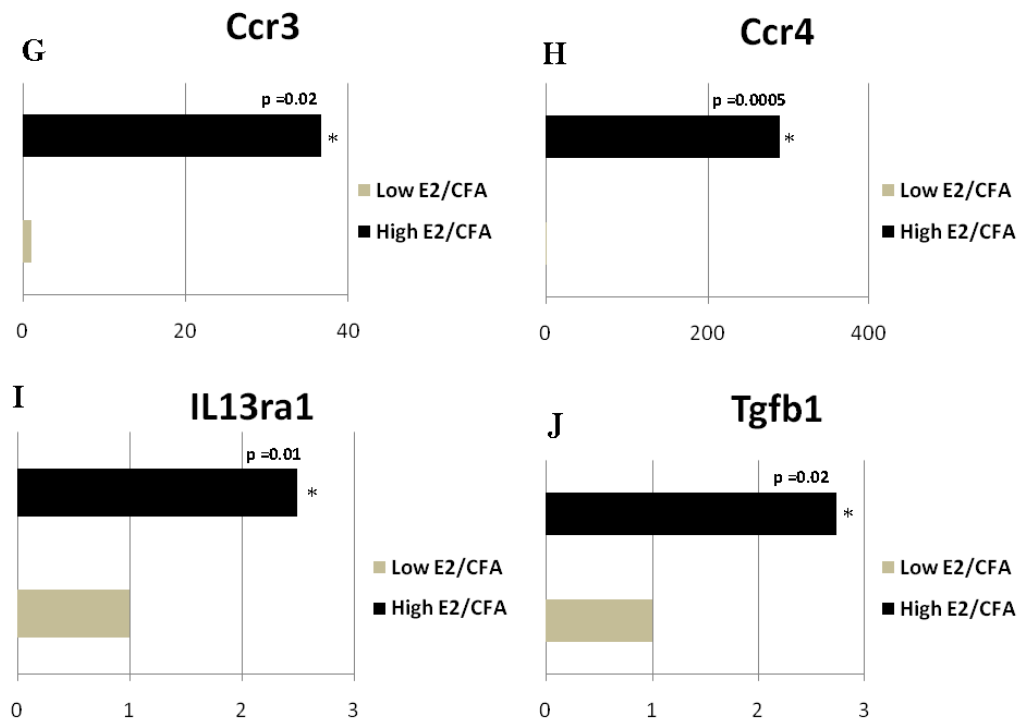


Figure 3. Real time PCR: Histogram showing expression of genes with immune function in the caudalis/upper cervical cord junction
 The expression of A) Ccl2, B) IL10, C) Ccr7, D) IL8ra and E) lymphotoxin b (Ltb) changed when comparing the Low E2 + SAL and High E2 + SAL groups. Expression of F) Ccl24, G) Ccr3, H) Ccr4, I) IL13ra1 and J) transforming growth factor β 1 (Tgfb1) changed when comparing the Low E2 + CFA and High E2 + CFA groups. The transcript levels from three independent experiments were collected and values were compared using the Student's *t*-test. The results were considered statistically significantly with $p < 0.05$. Values are the mean \pm SEM.

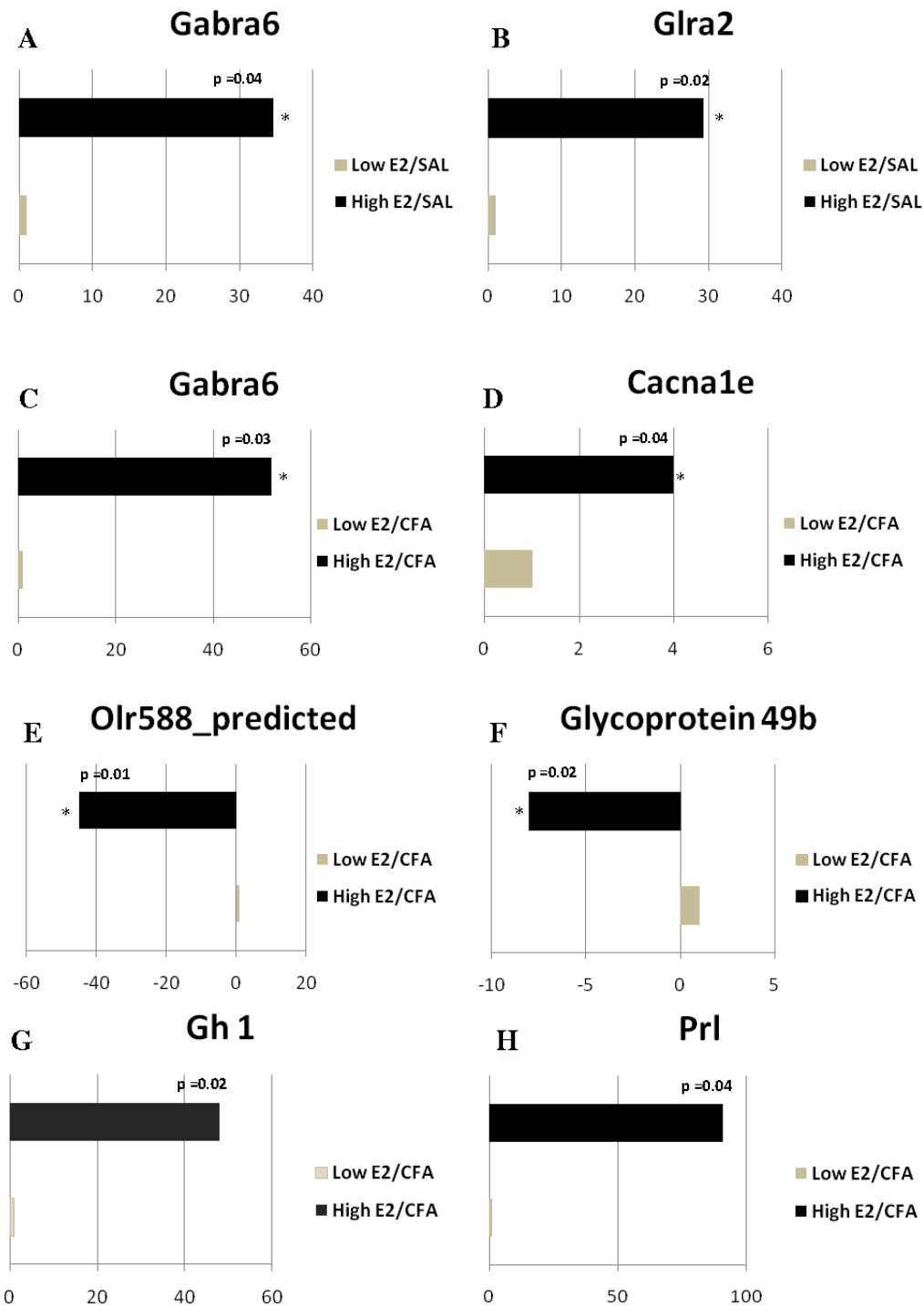


Figure 4. Real time PCR: Neurotransmitters and hormones genes changed in the trigeminal ganglion

Quantitative analysis using real-time PCR showed A) GABA A receptor subunit $\alpha 6$ (Gabra6), B) glycine receptor $\alpha 2$ (Glr2) comparing Low E2 + SAL and High E2 + SAL groups; and C) Gabra6, D) calcium channel voltage-dependent L type subunit $\alpha 1E$ (Cacna1e), E) Olr588, F) Glycoprotein 49b, G) growth hormone (Gh1), and H) prolactin (Prl) changed when comparing Low the E2 + CFA and E2 + CFA groups. The transcript

levels from three independent experiments were collected and values were compared using the Student's *t*-test. The results were considered statistically significantly with $p < 0.05$. Values are the mean \pm SEM.

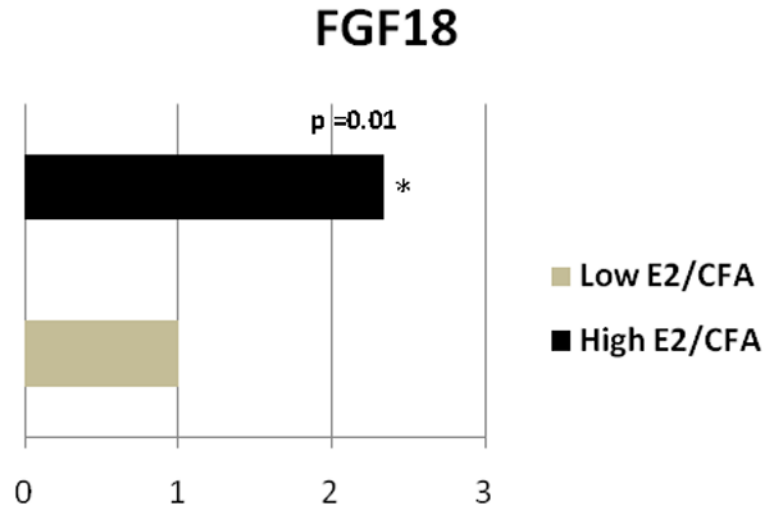


Figure 5. Real time PCR: Fibroblast growth factor 18 (Fgf18) expression changed in the subnucleus caudalis/upper cervical cord junction when comparing the Low E2 + CFA and High E2 + CFA groups

The transcript levels from three independent experiments were collected and values were compared using the Student's *t*-test. The results were considered statistically significantly with $p < 0.05$. Values are the mean \pm SEM.

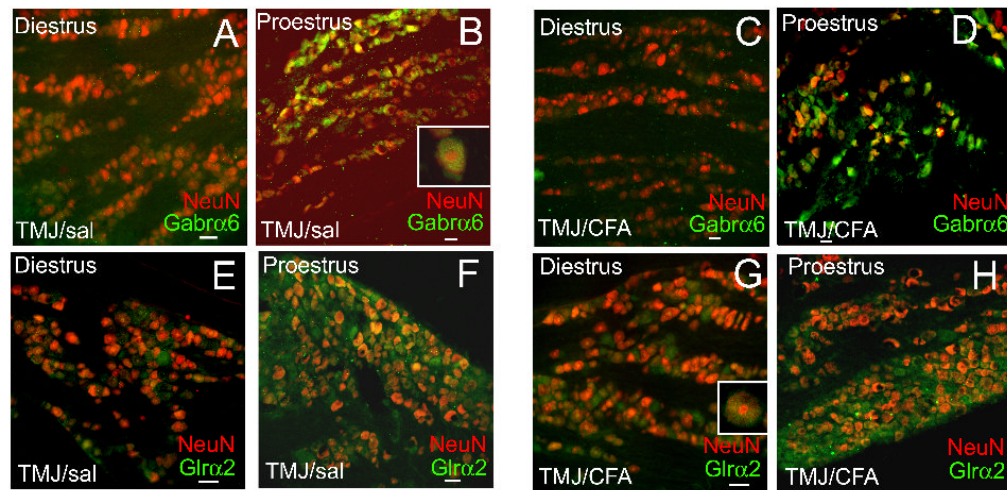


Figure 6. Fluorescent staining of Gabra6 or Glra2 in trigeminal ganglia tissue of intact female rats

Rats were given a bilateral injection (15 μ g) of CFA or were given an injection of 0.9% Sal into the superior TMJ space in the phase immediately preceding diestrus or proestrus. The phase of the estrous cycle was determined using vaginal smears. 24 hours after TMJ injection the rats were sacrificed in either the diestrus or proestrus phase, the trigeminal ganglia was isolated, fixed, sectioned and double stained with antibodies to Gabra6 (rabbit-Millipore, 1:1000) or Glra2 (rabbit-GeneTex, Inc., 1:500) and the neuronal marker NeuN (mouse-Millipore, 1:1000). Goat anti-rabbit 488 (green) or goat anti-mouse 633 (red) were the secondary antibodies used. Insert is a magnified image of a cell within that panel. A no primary antibody control showed no signal (data not shown). Size bar equals 50 μ m.

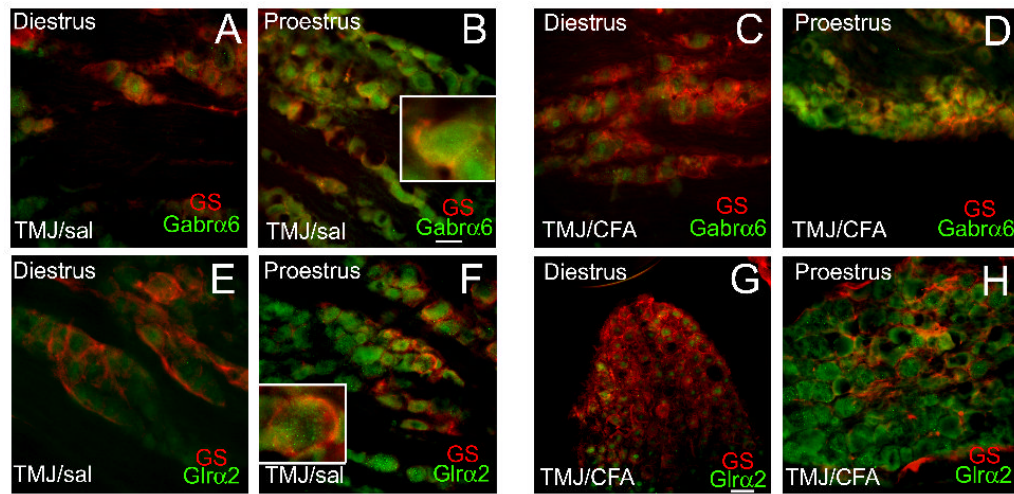


Figure 7. Fluorescent staining of trigeminal ganglia tissue from intact cycling rats-satellite glial gene expression

Rats were treated and stained as in Figure 6, a glutamine synthetase antibody (GS, mouse-Millipore, 1:300) was used as a marker for satellite glial cells. Goat anti-mouse 633 (red) was then used (1:500) as a secondary. Insert is a magnified image of a cell within that panel. A no primary antibody control showed no signal (data not shown). Size bar equals 50 μ m.

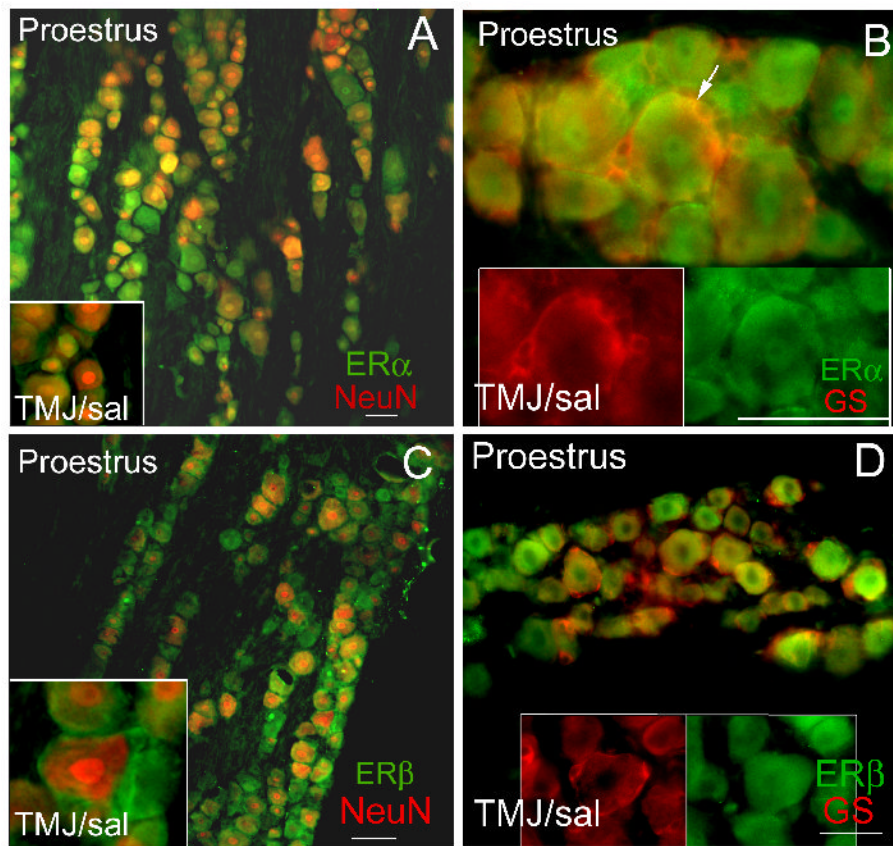


Figure 8. Estrogen receptor content in the trigeminal ganglia of a proestrus rat after saline TMJ injection

Trigeminal ganglia tissue was collected from an intact proestrus rat 24 hours after injecting the TMJ with saline. Frozen sections were double stained with a rabbit ER α antibody (panel A) (Santa Cruz, 1:50, green) or rabbit ER β antibody (panel C) (ABR, 1:100, green), with an antibody against neuronal marker NeuN (mouse-Millipore, 1:1000, red). Sections were also stained with an ER α (panel B, green) or ER β antibody (panel D, green) and an antibody against satellite glial marker glutamine synthetase (GS, red) (mouse-Millipore). The secondary antibodies were goat anti-rabbit 488 (green) and the goat anti-mouse 633 (red). A no primary antibody control showed no signal (data not shown). Arrow points to yellow double labeling. Size bar equals 50 μ m.