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### ► To cite this version:

Anne Morales, Chantal Bonnet, Naïck Bourgoïn, Thierry Touvier, Jérémie Nadam, et al.. Unexpected expression of orexin-B in basal conditions and increased levels in the adult rat hippocampus during pilocarpine-induced epileptogenesis. *Brain Research*, 2006, 1109, Issue 1, pp.164-175. 10.1016/j.brainres.2006.06.075 . hal-00090292

**HAL Id: hal-00090292**

**<https://hal.science/hal-00090292>**

Submitted on 29 Aug 2006

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# Unexpected expression of Orexin-B in basal conditions and increased levels in the adult rat hippocampus during pilocarpine-induced epileptogenesis

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31 pages

8 figures

1 table

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Abbreviations used: DG, Dentate Gyrus, ICV, intracerebroventricular, IP, intraperitoneal, LH, lateral hypothalamic area, OX, orexin, OXR, orexin receptor, PeF, perifornical nucleus, Pilo-SE, pilocarpine-induced SE, PVT, paraventricular thalamic nucleus, SD, scopolamine-diazepam, SE, status epilepticus, SmRNA, synthetic external and non-homologous poly(A) Standard RNA.

**Abstract, 1.**

Orexin A (OX-A) and B (OX-B) peptides present in the hippocampus are considered to be exclusively contained in fibers arising from hypothalamus neurons, which were established as the only source of orexins (OXs). Because OX-A is known to exert excitatory actions in the hippocampus, we hypothesized that level of OXs targeted toward the hippocampus may be increased following status epilepticus (SE)-induced epileptogenesis in the rat pilocarpine model of temporal lobe epilepsy. We found that tissue concentration of prepro-OX mRNA, which encodes for both peptides, rapidly decreased in the hypothalamus of rats having experienced pilocarpine-induced SE (Pilo-SE), followed by a reduced density of OX-A and OX-B immunopositive fibers arising from these neurons. By contrast, it was unexpected to detect within the hippocampus the presence of prepro-OX mRNA in basal conditions, and to evidence its up-regulation during the 1-3 day period following Pilo-SE. The number of prepro-OX mRNA copies determined by real-time RT-PCR was ~ 50-fold lower in the hippocampus than that in the hypothalamus, precluding the use of in situ hybridization to localize the cells which synthesize the transcript within the hippocampus. The increase in prepro-OX mRNA level within the hippocampus was accompanied by the detection of OX-B-like immunoreactivity 2-3 days post-SE, not only in pyramidal neurons, granule cells and cell bodies resembling interneurons, but also in some astrocytes scattered throughout the hippocampus. The present data suggest that the gene encoding OXs can be activated in the hippocampus, which may play a role in the pathogenesis of epilepsy.

**Section:** Disease-Related Neuroscience

**Keywords:** Temporal lobe epilepsy, hypothalamus, prepro-orexin, orexin, hypocretin, astrocytes

## **Introduction, 2.**

Orexin-A (OX-A) and orexin-B (OX-B), also known as hypocretin-1 and -2, are two peptides derived by proteolytic cleavage from a 130 amino acid prepro-OX precursor synthesized within neurons located in the hypothalamus (de Lecea et al., 1998; Peyron et al., 1998; Sakurai et al., 1998). By contrast, OX innervation arising from hypothalamus neurons is highly divergent (Peyron et al., 1998; Date et al., 1999) and G-protein-coupled OX receptors 1 and 2 (OX<sub>1</sub>R and OX<sub>2</sub>R, respectively) are widely distributed throughout the central nervous system (Sakurai et al., 1998; Trivedi et al., 1998; Greco and Shiromani, 2001; Hervieu et al., 2001; Marcus et al., 2001; Cluderay et al., 2002). Therefore, OX system has been recognized as a major regulator of various behavioural activities including feeding (Sakurai et al., 1998; Ida et al., 1999; Yamada et al., 2000), sympathetic nerve activity (Shirasaka et al., 1999) and normal sleep-wake cycle (Chemelli et al., 1999; Hagan et al., 1999) and its dysfunction has clearly been associated with pathophysiological disorders such as the sleep-disorder narcolepsy-cataplexy (Chemelli et al., 1999; Lin et al., 1999; Nishino et al., 2000; Peyron et al., 2000). Within the hippocampus, OX<sub>1</sub>R and OX<sub>2</sub>R have been specifically detected in CA1, CA2 and CA3 areas and in Dentate Gyrus (DG) (Trivedi et al., 1998; Hervieu et al., 2001; Marcus et al., 2001; Cluderay et al., 2002), suggesting that OXs may play a crucial role in the functions involving the hippocampus, as evidenced in learning paradigms after intracerebroventricular (ICV) injections of OX-A (Jaeger et al., 2002; Telegdy and Adamik, 2002; Aou et al., 2003). OXs released in the hippocampus by hypothalamus neurons may be directly implicated in the control of hippocampal excitability since: 1) ICV injection of OXs in rats is followed by behavioural seizure activity (Ida et al. 1999), and 2) application of OX-A to hippocampal slices modulates the balance between GABAergic and glutamatergic neurotransmissions (Selbach et al. 2004), the dysregulation of which is a characteristic feature of epilepsy (Bernard et al., 1997; Andre et al., 2001). Thus, OXs may also be involved in epileptogenesis, which is a sequence of biochemical, anatomical and electrophysiological changes that leads to recurrent spontaneous seizures in epilepsy, and that occurs following neurological insults or disorders, such as status epilepticus (SE).

Using calibrated reverse transcriptase (RT) and real time polymerase chain reaction (PCR), we evidenced for the first time within the hippocampus that prepro-OX mRNA was expressed in basal conditions and increased during epileptogenesis provoked by pilocarpine-induced status epilepticus (Pilo-SE). This expression correlated with OX-B synthesis in cell bodies resembling interneurons and

astrocytes. Concomitantly, expression of OXs decreased at both mRNA and protein levels in hypothalamus neurons of rats having experienced Pilo-SE.

### **Results, 3.**

#### **Study 1: Effect of Pilo-SE on body weight, feeding and drinking behaviours**

The main finding of this study is that we confirmed the decrease in body weight and food intake in rats having experienced Pilo-SE. At all time points examined, water intake in rats having experienced Pilo-SE did not differ significantly from that measured in rats treated with scopolamine and diazepam (SD) only, used to stop seizures induced by pilocarpine (not shown). By contrast, food consumption was reduced dramatically to a minimum observed 3 days following Pilo-SE (-82% vs SD rats,  $p < 0.001$ , Fig. 1A) and maximum body weight loss was noticed at day 5 post-SE (-34.8% vs SD rats,  $p < 0.001$ , Fig. 1B). The increase in food intake was initiated at day 5 (+ 123.2% between days 5 and 6,  $p < 0.05$ , Fig. 1A), preceding the gain in body weight noticed at day 7 (+ 14.6% between days 6 and 7,  $p < 0.05$ , Fig. 1B). From day 7 to 14, rats having experienced Pilo-SE exhibited hyperphagia compared to SD-treated rats (Fig. 1A). However, mean body weight values following Pilo-SE were always below those measured in SD-treated rats throughout the period examined (21 days post-SE), even though the difference was no more statistically significant from day 12 (Fig. 1B).

#### **Study 2: Effect of Pilo-SE on prepro-OX mRNA expression**

The main finding of this study is that prepro-OX mRNA was detected in the hippocampus and that prepro-OX mRNA levels changed in opposite ways in the hypothalamus and the hippocampus following Pilo-SE. In addition to the hypothalamus, prepro-OX mRNA was detected in the hippocampus following RT-real time PCR. No amplification was obtained when reverse transcriptase was omitted from the reaction mix (not shown), excluding the possibility of genomic DNA amplification. Furthermore, sequencing of the cDNA fragment obtained revealed 100% homology with prepro-OX cDNA sequence (GenBank: [AF041241](#)) of the rat, thus confirming the specificity of the amplification product. The number of copies of prepro-OX cDNA detected in basal conditions was ~50 fold lower in the hippocampus as compared to the hypothalamus ( $15,000 \pm 2,000$  and  $731,000 \pm 125,000$  copies, respectively). In the hypothalamus, prepro-OX mRNA levels decreased in rats having experienced Pilo-SE, the minimum observed being reached 1 day post-SE (-55.6%,  $p < 0.01$ ), levels similar to

controls being recovered at day 3 post-SE (Fig. 2). In the hippocampus, prepro-OX mRNA levels were also affected following Pilo-SE, but in an opposite way to that observed in the hypothalamus. A significant increase was observed as soon as 1 day post-SE (+ 125%,  $p < 0.001$ ), maintained at a plateau until day 3, control values being recovered at day 7 (Fig. 2). In situ hybridization using a DIG-labeled prepro-OX riboprobe allowed to confirm the localisation of prepro-OX mRNA containing neurons organized bilaterally and symmetrically in the lateral (LH) and posterior (PH) hypothalamic areas and in the perifornical nucleus (PeF) (Fig. 3A). Nevertheless, this technique was not sufficiently sensitive to detect the presence of prepro-OX mRNA containing cell bodies in the hippocampus (Fig. 3B), even following Pilo-SE. Indeed, based on quantitative RT-PCR data, the number of prepro-OX mRNA copies found in the hippocampus at day 3 post-SE was still 10 times lower than that determined at day 1 post-SE in the hypothalamus.

### **Study 3: Effect of Pilo-SE on prepro-OX, OX-A and OX-B distribution**

The main finding of this study is that OX expression was dramatically decreased in hypothalamus neurons following Pilo-SE, while OX-B expression was increased in the hippocampus within both neurons and astrocytes.

#### **Specificity of the antibodies used**

The polyclonal antibody used to detect OX-B was obtained from a 19 mer peptide mapping near the C-terminus of human OX-B (78-96 AA; Santa Cruz) and its specificity was tested by using preincubations with OX-B blocking peptide and recombinant mouse prolactin and bombesin. Although the possibility of a non-specific immunostaining cannot be completely eliminated, the antibody used has shown a good specificity. Immunostaining was decreased in a concentration-dependent manner with OX-B blocking peptide whereas immunoreactive staining was not affected after preincubation with 1-10  $\mu\text{M}$  recombinant mouse prolactin and bombesin, as illustrated in the hippocampus at day 3 post-SE (Fig. 4A-D). The specificity of OX-A antibody was not verified because OX-A immunolabelling was strictly identical with the one previously described (de Lecea et al., 1998; Peyron et al., 1998; Sakurai et al., 1998).

#### **OX-A and OX-B immunoreactivity of control rats**

In control rats, OX-A and OX-B immunoreactive (IR) cell bodies were abundantly found within the lateral hypothalamus and the perifornical nucleus (OX-B: Fig. 5A; OX-A: Fig. 6A), while they were

sparsely distributed in the posterior hypothalamic area. Distribution of prepro-OX IR cell bodies correlated with OX-A and OX-B IR cell bodies (not shown). A less intensive staining for OX-B was also observed in cell bodies of numerous regions, including layers 2-3 and 5 of the neocortex, the piriform cortex and the amygdaloid complex (not shown). OX-B IR cell bodies were equally found in the stratum oriens (SO), the stratum pyramidale (SP) and the stratum radiatum (SR) in the CA2-CA3 (Fig. 7A) and CA1 regions of the hippocampus (not shown). The number of OX-B containing cells counted in the hippocampus at IA +6.20 mm (Paxinos and Watson 1998) was  $16 \pm 3$  in SO,  $39 \pm 10$  in SP and  $17 \pm 5$  in SR. OX-A was never detected in cell bodies within these regions as illustrated in the hippocampus (Fig. 7C).

A prominent network of varicose OX-A and OX-B IR fibers was observed within the hypothalamus (OX-B: Fig. 5A; OX-A: Fig. 6A), the paraventricular thalamic nucleus (PVT) (OX-B: Fig. 5D; OX-A: Fig. 6D), and the amygdaloid complex (not shown). In the dorsal hippocampus, few OX-A and OX-B IR fibers were found located in the SO, SP and SR in the CA2-CA3 regions (OX-B: Fig. 7A; OX-A: Fig. 7C) and CA1 region in the Ammon's horn, in the hilus of the DG and in the area fasciola cinerea (not shown). Prepro-OX was rarely observed in both fibers and proximal processes of hypothalamus neurons (not shown).

#### **Alteration of OX-A and OX-B immunoreactivity in the hypothalamus and the paraventricular thalamic nucleus following Pilo-SE:**

In the LH and the PeF, counts of OX-A and OX-B containing neurons revealed no difference at all times examined following Pilo-SE. Cells counts of OX-A containing cells were  $30 \pm 7$  at day 2 post-SE ( $n= 4$ ) vs  $32 \pm 3$  in controls ( $n= 2$ ;  $P>0.05$ ) and  $21 \pm 9$  at day 4 post-SE ( $n= 4$ ) vs.  $25 \pm 6$  in controls ( $n= 2$ ;  $P>0.05$ ) (Figs 6A-C). Cells counts of OX-B containing cells were  $34 \pm 5$  at day 2 post-SE ( $n= 4$ ) vs  $29 \pm 6$  in controls ( $n= 2$ ;  $P>0.05$ ) and  $23 \pm 8$  at day 4 post-SE ( $n= 4$ ) vs  $36 \pm 7$  in controls ( $n= 2$ ;  $P>0.05$ ) (Figs. 5A-C). The density of both OX-A and OX-B IR fibers decreased significantly at day 2 post-SE vs controls (Table 1; OX-B: Figs. 5A, B; OX-A: Figs. 6A, B), where labelling appeared to be disorganised, and culminating at day 4 post-SE (Table 1, OX-B: Fig. 5C; OX-A: Fig. 6C).

In the PVT, a substantial decrease in the density of both OX-A and OX-B IR fibers was observed at day 2 post-SE vs controls (Table 1, OX-B: Figs. 5D, E, OX-A: Figs. 6D, E). The density of OX-B IR fibers tended to recover the pattern observed in controls at day 4 post-SE (Table 1, Fig. 5F) whereas OX-A labelling was lower (Table 1, Fig. 6F).

### **Alteration of OX-A and OX-B immunoreactivity in the hippocampus following Pilo-SE**

The quantitative analysis revealed that the density of OX-A and OX-B IR fibers did not change significantly in the SO, SP and SR of the hippocampus throughout epileptogenesis (data not shown), as illustrated at 3 days post-SE vs controls (OX-B: Figs 7A, B; OX-A: Figs 7C, D). Nevertheless, we observed a significant increase in OX-B containing cells in the SO ( $117 \pm 13$  vs  $16 \pm 3$  in controls,  $n=4$ ;  $P<0.001$ ), SP ( $123 \pm 24$  vs  $39 \pm 10$  in controls,  $n=4$ ;  $P<0.001$ ) and SR ( $366 \pm 91$  vs  $17 \pm 5$  in controls,  $n=4$ ;  $P<0.001$ ) of CA3-CA2 (Figs. 7A, B) and CA1 regions of the hippocampus. Prepro-OX (not shown) and OX-A were not detected in cell bodies within these regions following Pilo-SE (Figs. 7C, D). Double-labelling studies using OX-B antibody in combination with NeuN or GFAP antibody showed that OX-B immunoreactivity was present in neurons and astrocytes within the whole hippocampal formation at day 3 post-SE (Figs. 8A, B). All the differences observed following Pilo-SE were specific to the development of SE, and could not be the result of scopolamine and diazepam treatments used to stop SE. Indeed, SD-treated rats exhibited a pattern of immunoreactivity for prepro-OX, OX-A and OX-B similar to control rats (not shown).

### **Discussion, 4.**

Until recently, it was commonly agreed that the hypothalamus was the only source of OXs synthesis in the brain (de Lecea et al., 1998; Peyron et al., 1998; Sakurai et al., 1998). Our results extend the recent observations that OXs could be present in cell bodies outside from the hypothalamus (Ciriello et al., 2003) and highlight the possibility that synthesis of OXs could occur within the hippocampus, by showing, using quantitative RT-PCR: 1) a local presence of prepro-OX mRNA in basal conditions, and 2) an up-regulation of this transcript following Pilo-SE. Hence, we detected OX-B-like immunoreactivity in numerous cell bodies resembling interneurons and astrocytes throughout the hippocampus following Pilo-SE. These results, by evidencing that OXs could be differentially regulated between the hypothalamus and the hippocampus following Pilo-SE, suggest that the decrease of prepro-OX mRNA levels in hypothalamus neurons may support the reduction of food intake, while the increase of prepro-OX mRNA levels in the hippocampus may facilitate increased excitability of hippocampal neurons during epileptogenesis.

Reduction of food intake following Pilo-SE, that we have demonstrated here to be maximum at day 3



post-SE, was associated with a profound alteration of OX synthesis in hypothalamus neurons. Indeed, the density of OX-A and OX-B IR fibers was greatly decreased not only in brain areas receiving projections from orexinergic hypothalamus neurons, as clearly evidenced in the PVT, but also in the hypothalamus. The decreased density of OX-A and OX-B IR fibers may result from the death of some hypothalamus neurons, from an alteration of the axonal targeting of both peptides or from a reduced expression of the prepro-OX gene. Here, the dramatic decrease in prepro-OX mRNA levels to a minimum observed at day 1 post-SE and the unaltered OX cell number in the hypothalamus throughout epileptogenesis argues that only a reduced expression of OXs has caused the depletion of OX-A and OX-B peptides in fibers of hypothalamus neurons. However, it is noteworthy that the altered depletion of OX-A and OX-B peptides in fibers of orexinergic neurons spared those innervating the hippocampus. This result suggests that targeting of OXs in fibers of hypothalamus neurons is precisely controlled and depends of the brain area innervated. The correlation that we evidence here between reduced OX expression in hypothalamus neurons and decreased food consumption is in agreement with the observation that intracerebroventricular injection of either OX-A or OX-B increases food consumption (Sakurai et al., 1998; Haynes et al., 1999). Taken together, these results strongly suggest that regulating OX expression in hypothalamus neurons is a key mechanism in the control of feeding behaviour. These correlations are further reinforced at the mRNA level since 1) prepro-OX mRNA is highly down-regulated in rats exhibiting reduced appetite for food following Pilo-SE, and 2) food deprivation is followed in rats by the up-regulation of prepro-OX mRNA levels (Sakurai et al., 1998; Karteris et al., 2005).

The development of highly sensitive and quantitative methods, such as calibrated RT and real time PCR, to both detect and measure any alteration in specific transcripts, made it possible to locate prepro-OX mRNA in the hippocampus. We determined that the number of prepro-OX mRNA copies in 500 ng of total RNA in the hippocampus was ~50-fold less elevated than in the hypothalamus in basal conditions. This result may explain why prepro-OX mRNA remained undetectable in the hippocampus using colorimetric in situ hybridization, even after its level increased by 2.5-fold through 3 days after Pilo-SE. However, such an increase in prepro-OX mRNA in the hippocampus was sufficient to elevate OX-B synthesis above detection threshold in diverse cell populations, including pyramidal neurons, interneurons and astrocytes. OX-A could not be evidenced in these cells, in spite of the fact that both peptides are derived from the same precursor (prepro-OX) by proteolytic cleavage (Sakurai et al.,

1998). Even if we cannot completely rule out the possibility that the antibody used recognized other targets than OX-B, pre-incubation of the antibody with the peptide sequence used for immunization decreased in a dose-dependant manner the signal obtained in the hippocampus, while pre-incubations with prolactin and bombesin failed to reduce the intensity of the immunolabeling. Previous data have provided evidence that neurons only labeled for OX-B were also identified in others brain regions including the central nucleus of the amygdala and the anterior lateral subnucleus of the bed nucleus of the stria terminalis (Ciriello et al., 2003). As suggested by these authors, lack of OX-A detection in the hippocampus might be explained by the absence of the proteolytic enzyme for the production of OX-A and/or by the selective clearance of OX-A by degrading enzymes.

We initially postulated that OXs originating from hypothalamus neurons may facilitate increased excitability of hippocampal neurons during epileptogenesis, since prior studies reported the implication of OXs in the modulation of neuronal excitability within the hippocampus and in the behavioural seizure activity in rats (Aou et al., 2003; Ida et al., 1999, Selbach et al., 2004). However, instead of an increased presence of OXs originating from the hypothalamus following Pilo-SE, our study shows that the density of OX-A and OX-B immunopositive fibers remained stable in the hippocampus. By contrast, the number of OX-B expressing cells dramatically and transiently increased in the hippocampus following Pilo-SE, as a consequence of the activation of the gene encoding OXs. Whereas OXs synthesized in the hypothalamus are distributed in all compartments (cell bodies, dendrites and axons) of hypothalamus neurons, we noticed that OX-B synthesized in hippocampal cells was restricted to the cell bodies. This observation led us to suggest that OX-B produced in the hippocampus in basal conditions may be released locally to exert a paracrine/autocrine control on neuronal excitability. Such a role may also be played by both interneurons and astrocytes during Pilo-induced epileptogenesis. Such an action could be mediated by OX<sub>1</sub>R and OX<sub>2</sub>R, the specific receptors for OX-A and B, detected within the hippocampus (Trivedi et al., 1998; Hervieu et al., 2001; Marcus et al., 2001; Cluderay et al., 2002). Although OX-A was shown to be from 5 to 100-fold more potent than OX-B in calcium mobilization studies using CHO cells (Sakurai et al., 1998; Smart et al., 1999; Langmead et al., 2004), binding studies resulted in contradictory results in the same transfected cellular model. Indeed, OX-B was shown to have even higher affinity than OX-A in CHO or neuroblastoma cells expressing native OX<sub>1</sub>R (Wieland et al., 2002). Moreover, Rouet-Benzineb et al. (2004) found that OX-A and OX-B suppressed cell growth identically in a cell line expressing OX<sub>1</sub>R

only. Therefore, in our model, the neuronal excitability could be mediated at both pre- and post-synaptic sites through both OX<sub>1</sub>R and OX<sub>2</sub>R. Presynaptic orexinergic excitatory actions have been demonstrated on prefrontal thalamocortical synapses (Lambe and Aghajanian, 2003) and in laterodorsal tegmental neurons (Bulet et al., 2002). At a post-synaptic level, excitatory actions of OX-B have been detected in basal forebrain cholinergic neurons (Eggermann et al., 2001), in histaminergic neurons of the tuberomammillary nucleus (Bayer et al., 2001; Eriksson et al., 2001), in sublayer 6b cortical neurons (Bayer et al., 2004) and in serotonin neurons of the dorsal raphe nucleus (Brown et al., 2002). It is thus likely that locally synthesized OX-B supports increased excitability of hippocampal neurons in the early stages of Pilo-induced epileptogenesis. To date, the lack of specific OX<sub>2</sub>R antagonist prevents to verify further the role played by OX-B following Pilo-SE. In addition to a role in excitability modulation, OX-B synthesized in the hippocampus may also play a role in neuronal apoptosis following Pilo-SE (Henshall and Simon, 2005) which is a major event during brain development and maturation and also in neurodegenerative diseases (Lossi and Merighi, 2003). Indeed, Oxs acting at OX<sub>1</sub>R have been shown to induce apoptosis acting at native OX<sub>1</sub>R in colon cancer and neuroblastoma cells (Rouet-Benzineb et al, 2004).

In conclusion, the detection of distinct clusters of neurons and astrocytes containing OX-B in the hippocampus at day 3 following Pilo-SE led us to suggest that this peptide may be involved in neuronal plasticity during the early stages of epileptogenesis and that the hypothalamus is not the only source of OXs throughout the central nervous system at least in pathophysiological conditions. Further studies will be necessary to fully discern whether the ectopic orexinergic expression in the hippocampus induces a pro- or an anti-epileptic effect.

## **Experimental Procedure**

### **Animals**

Male Sprague-Dawley rats (220 ± 19 g) from Harlan (France) were housed for 4 weeks before treatments in groups of five under controlled environmental conditions (lights on from 6.00 to 18.00, 23°C ± 1°C and free access to food and water). All experiments were performed in accordance with the guidelines of the French Ministry of Agriculture (87/848) and the European Community (86/609/EEC). The day of the experimental procedures (induction of epileptogenesis), rats were

transferred to individual cages and kept isolated throughout the experiments. Epileptogenesis was induced in rats following Pilo-SE (Turski et al., 1983). Pilocarpine hydrochloride (Sigma) was injected intraperitoneally (IP) at a dose of 300 mg/kg (Pilo-treated rats). Methylscopolamine nitrate (1 mg/kg) was injected subcutaneously (SC) 30 min prior to pilocarpine in order to minimize peripheral cholinergic effects. Rats entered into SE  $22 \pm 2$  min after the injection of pilocarpine and seizures were stopped 3 hrs after SE onset with an IP injection of 10 mg/kg diazepam (Valium®, Roche). Scopolamine-Diazepam (SD)-treated rats received a SC injection of methylscopolamine nitrate (1 mg/kg), followed 30 min later by an IP injection of 0.9% Sodium Chloride solution and then 3h later, by an IP injection of 10 mg/kg diazepam. Naïve rats (control rats) did not receive any treatment and were used to determine the potential combinatory effects of SD injections on gene expression.

### **Study 1: Effect of Pilo-SE on body weight, feeding and drinking behaviours**

To study the consequences of the Pilo-SE on the general conditions of the rats, body weights and pre-weighted rat food and water were monitored each day at 9:00 am during 21 days following Pilo-SE (day 0) (Pilo rats n= 14 up to day 5, n= 9 at day 6, n= 7 at day 7 and beyond). Values were compared with those obtained from SD-treated rats (n=4).

### **Study 2: Effect of Pilo-SE on prepro-OX mRNA expression**

Reverse Transcriptase real time Polymerase Chain Reaction (RT-real time PCR)

Animals were sacrificed after the onset of SE following a lethal injection of pentobarbital (250 mg/kg) and the hippocampus was rapidly removed and frozen at t=8h (Pilo n=5, SD n=5), t=1 day (Pilo n=6, SD n=4), t=2 days (Pilo n=6, SD n=5), t= 3 days (Pilo n=8, SD n=5) and t=7 days (Pilo n=3, SD n=5). The hypothalamus was collected at t=8h, t=1 day and t=3 days after SE. Both brain structures were collected in naïve rats, sacrificed at t=0 (controls n=5).

Total RNAs were extracted with Tri-reagent LS (Euromedex) and genomic DNA was removed after DNase I digestion (RNase Free DNase Set, QIAGEN). PCR amplifications were performed with reaction in which the reverse transcriptase was omitted to allow for assessment of genomic DNA contamination of the RNA and results revealed the absence of amplified fragment. Messenger RNAs, contained in 500 ng hippocampal or hypothalamic total RNAs were then reverse transcribed with the reverse transcriptase RNase H minus (Promega) using oligod(T)<sub>15</sub>, in the presence of 80 pg of a

synthetic external and non-homologous poly(A) Standard RNA (SmRNA) to normalize the reverse transcription reaction of mRNAs of biological samples (Morales and Bezin, 2004); World Patent number 2004/092414 A12004). cDNAs obtained from the reverse-transcription of prepro-OX mRNA were quantified by real time PCR performed on the LightCycler® System (Roche Diagnostics) using the QuantiTect SYBR®Green PCR Kit (Qiagen). Both reverse (5'-CAGACACCATGAACCTTCCT-3') and forward (5'-TGGTTACCGTTGGCCTGAAG-3') primers (GenBank accession no. [AF041241](#)) generate a 274 bp PCR fragment which was confirmed by sequencing. Results obtained for prepro-OX mRNA were normalized against the SmRNA.

#### In situ hybridization (ISH)

Animals were deeply anesthetized (lethal IP injection of pentobarbital at 250 mg/kg) and were then perfused at 30 mL/min through the ascending aorta with 60 mL of phosphate buffered saline (PBS, 0.1M, pH 7.4) followed by 300 mL of chilled 4% paraformaldehyde (PFA) in PB (0.1 M, pH 7.4). The brains were post-fixed for 2 h in the same fixative at 4°C, transferred to 25% sucrose in PB at 4°C, frozen in isopentane at -40°C and then stored at -80°C until further use. Coronal sections were performed with a cryostat (thickness of 40 µm) at anatomical planes from interaural (IA) 7.40 mm to 4.84 mm according to the rat brain atlas (Paxinos and Watson 1998). A 274 bp segment of rat cDNA encoding Leu28-Ala119 of prepro-OX was generated by conventional PCR and subcloned into a pGEM-T easy plasmid (Promega). Digoxigenin (DIG)-labelled sense and anti-sense riboprobes were generated with T7 and SP6 RNA polymerases, respectively, using the MEGAscript kit (Ambion). ISH protocol was performed on free floating brain tissue sections of rats sacrificed at day 3 post-SE (pilo n=2, control n=2) as already described in details (Tongiorgi et al. 1998), with slight modifications. Briefly, prior to hybridization, slices were post-fixed 2 hours in 4% PFA in PB (0.1 M, pH 7.4) and were permeabilized with 2.3% sodium meta-periodate (Sigma) in H<sub>2</sub>O at RT for 5 min. The sections were incubated in 1% sodium borohydride (Sigma) in 0.1 M Tris-HCl buffer pH 7.5 at RT for 10 min and were digested with 2 µg/mL proteinase K (Roche) at 4°C for 10 min and washed twice in PBST at RT for 5 min. After digestion, the tissue sections were fixed in 4% PFA in PB at RT for 5 min. In situ hybridization was carried overnight at 55°C with the hybridization solution containing the DIG-labeled riboprobes (0.5 ng/µL) and 50% dionized formamide, 10% dextran sulfate, 1x Denhardt's, 20 mM Tris-HCl, 300 mM NaCl, 5 mM EDTA, 0.125 ng/mL denatured salmon sperm DNA. After a series of

washes in 0.1X saline sodium citrate (SSC), 0.1% Tween 20 at 60°C for 15 min, sections were incubated for 30 min with 20 µg/mL RNase A (Qiagen) at 37°C. Slices were washed three times in 2x SSC, twice in PBS 0.1 M for 15 min and then were processed for immunodetection with an anti-DIG antibody F(ab) fragment (Roche) diluted 1:1000 in 1X blocking reagent (Roche) at RT for 1h. Sections were incubated in developing buffer containing 1 mM Levamisol at RT for 15 min and finally incubated overnight at 30°C in a chromogen solution composed by the developing buffer containing nitro blue-terazolium (NBT) (Roche) and 5-bromo-4-chloro-3-indolyl-phosphate (BCIP) (Roche). Sections were washed three times in PBS 0.1 M, mounted on a glass slide (SuperFrost Plus), dehydrated and defatted in graded ethanols and xylene and then mounted in DPX (Fluka).

### **Study 3: Effect of Pilo-SE on prepro-OX, OX-A and OX-B distribution**

Immunohistochemistry was performed for prepro-OX, OX-A and OX-B using immunohistochemistry on adjacent free-floating fixed brain tissue sections of rats sacrificed at 8h (Pilo n=4, SD n=2), 1 day (Pilo n=4, SD n=2), 2 days (Pilo n=4, SD n=2), 3 days (Pilo n=4, SD n=2) and 4 days (Pilo n=4, SD n=2) after Pilo-SE. After rinses in PBS and permeabilization in PBS containing 0.3% triton X100 (PBS-T), sections were preincubated for 2 h at room temperature (RT) in 2% Normal Donkey Serum in PBS-T (PBS-T-D). Incubations with primary antibodies were then performed at 4°C for 48 h, using either a polyclonal goat anti-OX-A antibody (1:1,000; sc-8070; Santa-Cruz Biotechnology), a polyclonal goat anti-OX-B antibody (1:1,000; sc-8071; Santa-Cruz Biotechnology) or a polyclonal rabbit anti-prepro-OX antibody (1:250; AB3096; Chemicon), diluted in PBS-T-D. For chromogenic reaction, sections were washed in PBS-T and were incubated for 24 h at 4°C, either with a biotinylated donkey anti-goat IgG (1:10,000; AP180B, Chemicon) or a biotinylated donkey anti-rabbit IgG (1:5,000; 711-066-152, Jackson ImmunoResearch) diluted in PBS-T-D. Sections were then incubated for 5 min in 1% H<sub>2</sub>O<sub>2</sub>, rinsed in PBS and then incubated for 1h30 min at RT with avidin biotin peroxylase (1:500; Vectastain Elite ABC kit, Vector) diluted in PBS. The sections were reacted with 0.4 mM 3', 3-diaminobenzidine (DAB, Sigma Fast) for 5 min, then mounted, dehydrated and coverslipped. The staining results were observed with a video camera 3CCD (DXC-930P; Sony) connected to a microscope (Leitz Diaplan).

In order to characterize cell types, fluorescent double immunostaining was performed using the polyclonal goat anti-OX-B antibody described above (1:1,000) with either a mouse anti-NeuN (1:1,000, MAB 377, Chemicon) or a rabbit anti-GFAP (Glial Fibrillary Acid Protein, 1:3000; AB 5804; Chemicon)

antibody to detect neurons and astrocytes, respectively. Sections were washed in PBS-T before being exposed to donkey anti-mouse IgG (1:500; Alexa-488, A-21202; Molecular probes), donkey anti-rabbit IgG (1:500; Alexa-488; A-21206; Molecular Probes) or donkey anti-goat IgG (1:500; Alexa-633; A-21082; Molecular Probes) antibodies. Specificity controls of the OX-B antiserum used were done by preincubating the primary antiserum with 1-10  $\mu\text{M}$  of either OX-B blocking peptide (78-96 AA; sc-8071 P, Santa-Cruz Biotechnology), or recombinant mouse prolactin (1445-PL RnDSystems) or bombesin (H-2155, Bachem, Switzerland). Sections were mounted, dehydrated and coverglassed with Prolong Gold Antifade reagent (P 36930; Molecular Probes). Images captured using a TCS SP2 confocal microscopy system (Leica) were imported into Adobe Photoshop 8.0.1 (Adobe Systems, San Jose, USA) for further editing.

### **Neuronal counts and measurements of density of fibers**

Sections immunostained for OX-A and OX-B were observed under a light microscope (Diaplan; Leitz), and images were captured with a video camera 3CCD (DXC-930P; Sony) coupled to an image analysis system (Visilog® 6.0; Noesis). The system allows to scan at magnification 10X adjacent fields throughout the brain structures (LH, PeF, dorsal hippocampus and PVT) and to reconstruct a single image composed of a mosaic of the digitized adjacent fields. These structures were selected at IA +6.20 mm (Paxinos and Watson 1998). Neurons expressing OX-A and OX-B were counted within the surface area of the LH and the PeF occupied by OX-A and OX-B immunopositive cell bodies at both sides and were expressed as mean cells  $\pm$  SEM. Cells expressing OX-B were measured within the SO (measurement window: 50,000  $\mu\text{m}^2$ ) SP (measurement window: 50,000  $\mu\text{m}^2$ ) and SR (measurement window: 170,000  $\mu\text{m}^2$ ) of the Ammon's Horn of the hippocampus at both sides and were expressed as mean cells  $\pm$  SEM. The density of fibers expressing OX-A and OX-B was determined in measurement windows, the surface area of which is indicated for each structure, and expressed as the surface area occupied by the specific immunolabeling within the measurement window. The structures analyzed were the PVT (measurement window: 190,000  $\mu\text{m}^2$ ) and the LH (measurement window: 280,000  $\mu\text{m}^2$ ). In controls, mean densities of OX-A and OX-B fibers were expressed as  $\mu\text{m}^2$  of fibers/ $\mu\text{m}^2$  of measurement windows. Mean densities of OX-A and OX-B fibers following Pilo-SE were expressed as the percentage of variations vs controls.

### **Data analysis**

All results were presented as mean  $\pm$  S.E.M. Group comparisons were performed following two-way analysis of variance (ANOVA II) (factor 1: treatment, factor 2: time), using a post-hoc Fisher's LSD test.

### **Acknowledgments**

We thank B. Smatti and Y. Tourneur from the "Centre Commun de Quantimétrie" of Université Claude Bernard Lyon 1 for their helpful assistance in confocal microscopy analysis. This study was supported by the Centre National de la Recherche Scientifique (CNRS) and the University Claude Bernard Lyon 1, France.



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## Figure legends

**Fig.1** Food intake (A) and body weight (B) monitoring following Pilo-SE reveals a rapid and drastic reduction in food consumption until day 5, the increase in food intake observed between days 5-6 preceding the significant gain in body weight observed at day 7. Note that rats having experienced Pilo-SE exhibited hyperphagia from day 7 to day 14 post-SE, and that the difference in body weight between these rats and Scopolamine-Diazepam (SD) controls was not significant by day 12. ▲:Difference in pilocarpine-treated rats at one time point as compared to the prior one. ▲  $p<0.05$ ; ▲▲▲  $p<0.001$ . \*: Difference at a given time point between Pilo-treated and SD-treated rats. \* $p<0.05$ ; \*\* $p<0.01$ ; \*\*\* $p<0.001$ . Pilo rats:  $n=14$  up to day 5,  $n=9$  at day 6,  $n=7$  at day 7 and beyond. SD rats:  $n=4$ .

**Fig.2** Levels of expression of prepro-OX mRNA in the hippocampus and in the hypothalamus following Pilo-SE by RT-real time PCR. Messenger RNAs were reverse transcribed in the presence of a synthetic external and non-homologous poly(A) Standard RNA (SmRNA) to normalize the reverse transcription reaction. In the hypothalamus, prepro-OX mRNA levels decreased in rats having experienced Pilo-SE, the minimum observed being reached at day 1 post-SE (-55.6%,  $p<0.01$ ), levels similar to controls being recovered at day 3 post-SE. In the hippocampus, the induction of prepro-OX mRNA levels contrasts with the inhibition observed in hypothalamus neurons following Pilo-SE. Values are expressed as percentage of control rats. \*\* $p<0.01$ ; \*\*\* $p<0.001$ .

**Fig. 3** Bright-field photomicrographs in the perifornical nucleus of the hypothalamus and the CA3-CA2 regions of the hippocampus at day 3 post-SE in rat brain sections hybridized with digoxigenin-labeled riboprobes for prepro-OX mRNA. In situ hybridization technique allowed to confirm the localisation of prepro-OX mRNA containing neurons in the PeF (A) but was not sufficiently sensitive to detect the presence of prepro-OX mRNA containing cell bodies in the CA3-CA2 regions of the hippocampus (B) even following Pilo-SE. Scale bar: 100  $\mu\text{m}$  (A-B).

**Fig. 4** Confocal images of the CA3 region of the hippocampus at day 3 post-SE labelled with OX-B antiserum in the absence (A) and in the presence of 10  $\mu\text{M}$  of either OX-B blocking peptide (the 19 mer fragment used to obtain the antibody) (B) or recombinant mouse prolactin (C) and bombesin

(D). OX-B immunoreactivity was strongly decreased when 10  $\mu$ M OX-B blocking peptide was incubated with OX-B antibody (B) while immunolabeling was not affected when 10  $\mu$ M of either recombinant mouse prolactin (C) or bombesin (D) was added. Scale bar: 15  $\mu$ m (A-D).

**Fig. 5** Immunohistochemical detection of OX-B in the lateral hypothalamus and the paraventricular thalamic nucleus in control rats and in rats having experienced Pilo-SE. In the LH, counts of OX-B containing neurons revealed no difference at day 2 (B) and at day 4 post-SE (C) vs controls (A). The density of OX-B IR fibers decreased significantly at day 2 post-SE (B) where labelling appeared to be disorganised and culminating at day 4 post-SE (C). In the PVT, a substantial decrease in the density of OX-B IR fibers was observed at day 2 post-SE (E) vs controls (D) which tended to recover the pattern observed in controls at day 4 post-SE (F). Scale bar: 250  $\mu$ m (A-F).

**Fig. 6** Immunohistochemical detection of OX-A in the lateral hypothalamus and the paraventricular thalamic nucleus in control rats and in rats having experienced Pilo-SE. In the LH, counts of OX-A containing neurons revealed no difference at day 2 (B) and day 4 post-SE (C) vs controls (A). The density of OX-A IR fibers diminished significantly at day 2 post-SE (B), culminating at day 4 post-SE (C). In the PVT, the density of OX-A IR fibers decreased at day 2 post-SE (E) vs controls (D) and was lower at day 4 post-SE (F). Scale bar: 250  $\mu$ m (A-F).

**Fig. 7** Immunohistochemical detection of OX-B and OX-A in the hippocampus. OX-B was detected in control rats (A) and at day 3 post-SE (B). A significant increase in OX-B containing cells was observed in the stratum oriens (SO), the stratum pyramidale (SP) and the stratum radiatum (SR) of the CA3-CA2 region of the hippocampus (B). Nevertheless, OX-A labeling was not detected in these regions both in control rats (C) and at day 3 post-SE (D). Scale bar: A,B,C,D 100  $\mu$ m

**Fig. 8** Double-immunostainings were performed using either a mouse anti-NeuN or a rabbit anti-GFAP antibody (to characterize neurons and astrocytes, respectively) in combination with the anti-OX-B antibody in the stratum radiatum of CA3 region of the hippocampus at day 3 post-SE. OX-B appears as red fluorescence and both NeuN and GFAP appear as green fluorescence. Confocal microscopy images of the rat hippocampus double labelled with OX-B antiserum and NeuN showed that OX-B

immunoreactivity was present in neurons localised in the stratum radiatum of the CA3 region of the hippocampal formation (A). Straight arrow indicates representative neuron with coincident NeuN (green fluorescence) and OX-B labeling (red fluorescence). OX-B immunoreactivity was equally present in astrocytes (B) within this region. Crossed arrow indicates representative astrocyte with coincident GFAP (green fluorescence) and OX-B labeling (red fluorescence). Scale bar: 10  $\mu$ m (A-B).

Table 1: Altered density of OX-A and OX-B fibers in the lateral hypothalamus and the PVT following Pilo-SE.

	OX-A		OX-B	
	Day 2	Day 4	Day 2	Day 4
Lateral HPT	- 38 ± 6 <sup>*</sup> (n=4)	- 68 ± 8 <sup>**</sup> (n=4)	- 41 ± 4 <sup>*</sup> (n=4)	- 60 ± 7 <sup>**</sup> (n=4)
PVT	- 39 ± 7 <sup>*</sup> (n=4)	- 51 ± 6 <sup>**</sup> (n=4)	- 56 ± 5 <sup>**</sup> (n=4)	- 8 ± 3 (n=4)

In controls, mean density of OX-A IR fibers expressed as  $\mu\text{m}^2 / \mu\text{m}^2$  of measurement area was:  $0.17 \pm 0.02$  and  $0.12 \pm 0.03$  in the LH and the PVT respectively. In controls, mean density of OX-B IR fibers expressed as  $\mu\text{m}^2 / \mu\text{m}^2$  of measurement area was:  $0.14 \pm 0.03$  and  $0.09 \pm 0.02$  in the LH and the PVT respectively. At day 2 and day 4 post-SE, mean densities of OX-A and OX-B IR fibers are expressed as the percentage of variations vs controls. \*  $P < 0.05$ , \*\* $P < 0.01$  vs controls.



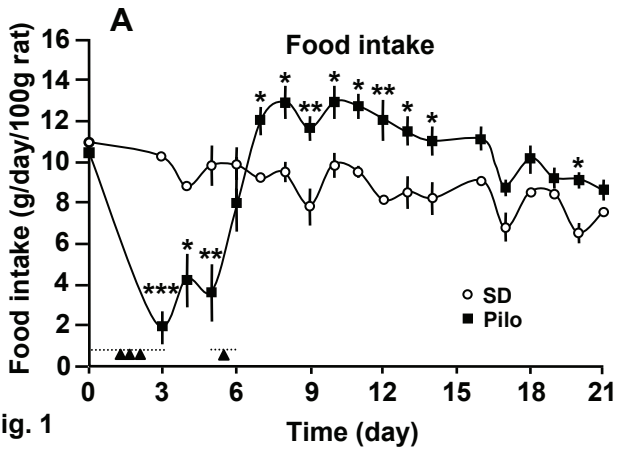
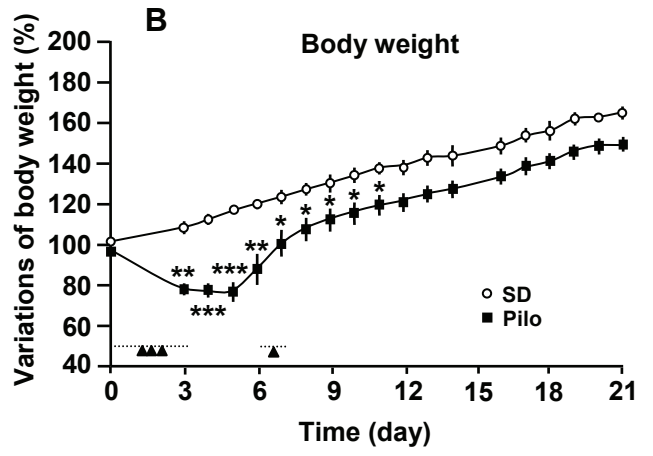


Fig. 1



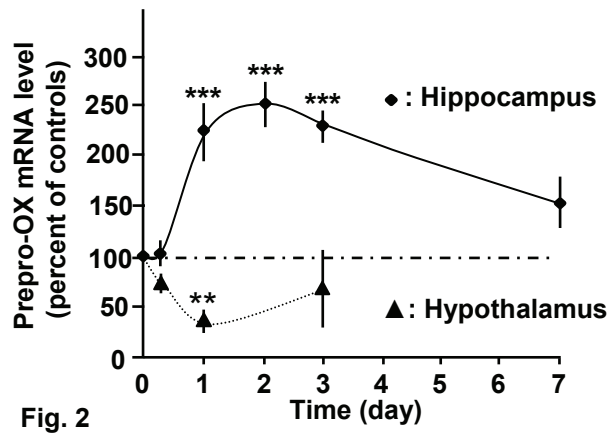
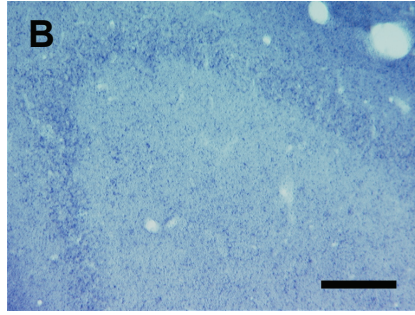
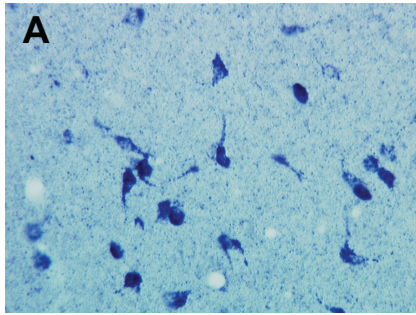
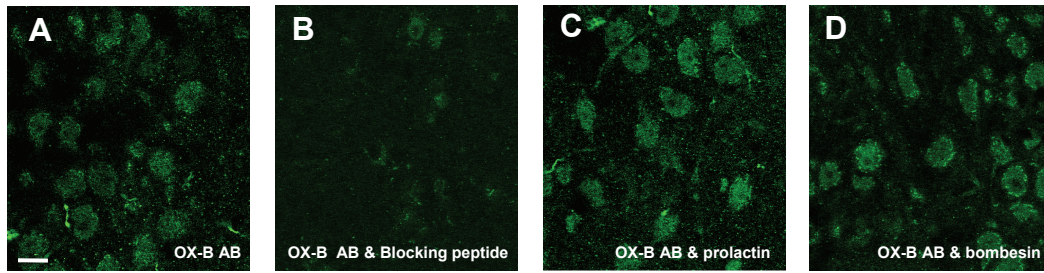


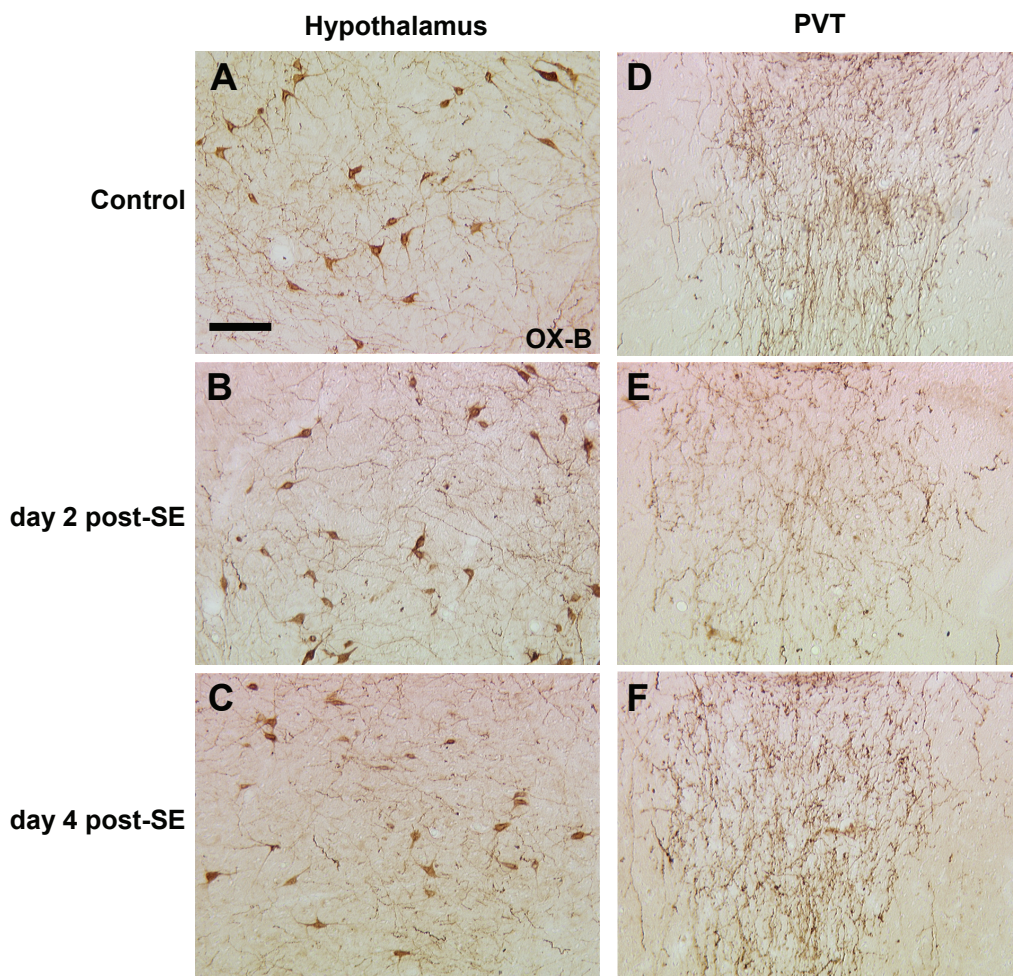
Fig. 2



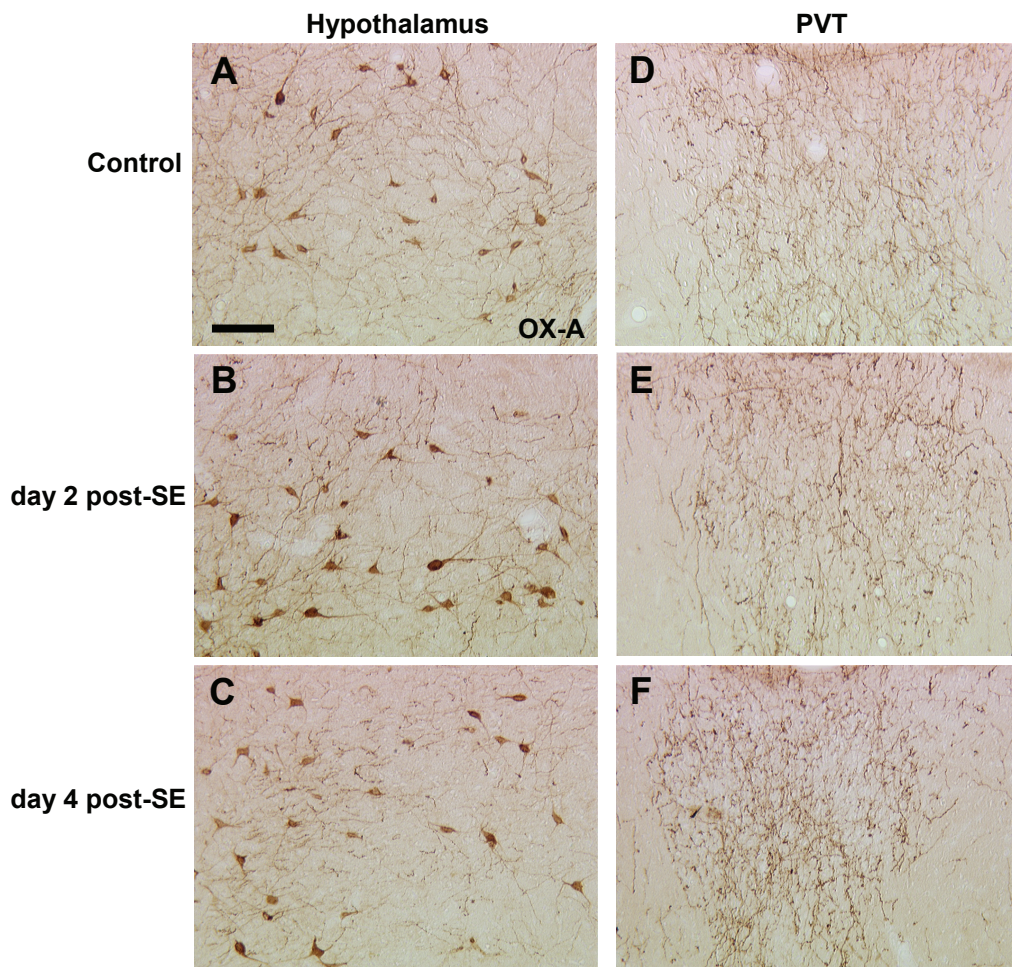
**Fig.3**



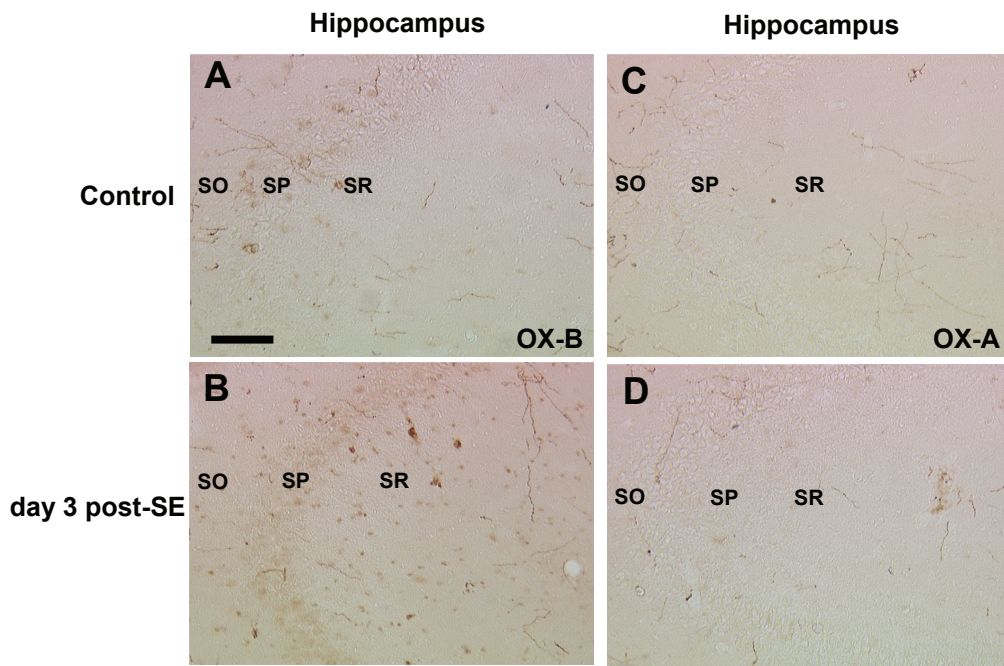
**Fig.4**



**Fig. 5**



**Fig. 6**



**Fig. 7**

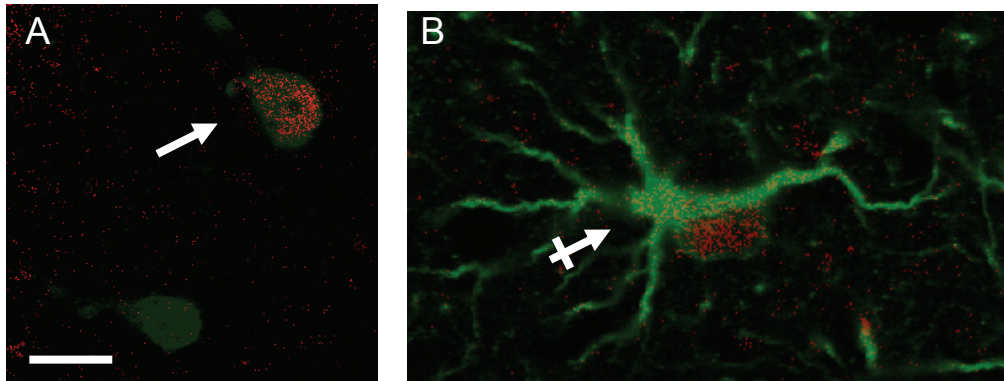


Fig. 8