Phosphodiesterase 7 inhibitor TC3.6 ameliorates symptomatology in a model of primary progressive multiple sclerosis.

Running title: PDE7 inhibitors for PPMS

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Abstract

Background and purpose: cyclic Adenosine monophosphate (cAMP) plays an important role in the transduction of signaling pathways involved in neuroprotection and immune regulation. Control of the levels of this nucleotide by inhibition of cAMP specific phosphodiesterases (PDEs) such as PDE7 may affect the pathological processes of neuroinflammatory diseases like multiple sclerosis (MS). In the present study we evaluated the therapeutic potential of the selective PDE7 inhibitor, named TC3.6 in a model of primary progressive multiple sclerosis (PPMS), a rare and severe variant of MS.

Experimental approach: TMEV-induced demyelinated disease (TMEV-IDD) is one of the models used to validate the therapeutic efficacy of new drugs in MS. As recent studies have analyzed the effect of PDE7 inhibitors in the EAE model of MS, here the TMEV-IDD model is used to test the efficacy in a progressive variant of MS. Mice were subjected to two protocols of TC3.6 administration: on the presymptomatic phase and once the disease was established.

Key results: Treatment with TC3.6 ameliorated the disease course and improved motor deficits of infected mice. This was associated with down-regulation of microglial activation and reduced cellular infiltrates. Decreased expression of proinflammatory mediators such as COX-2 and the cytokines, IL-1β, TNFα, IFNγ and IL-6 in the spinal cord of TMEV-infected mice was also observed after TC3.6 administration.

Conclusion: These findings support the interest of PDE7 inhibitors, and specifically TC3.6, as a novel class of agents with therapeutic potential for PPMS raising the possibility to develop regulatory preclinical studies to reach the clinic.

Abbreviations:
PDE, phosphodiesterase; cAMP, cyclic adenosine 3’,5’-monophosphate; PPMS, primary progressive multiple sclerosis; TMEV-IDD, Theiler’s murine encephalomyelitis virus-induced demyelinating disease

**Key words:** Phosphodiesterase-7 Inhibitors; Multiple sclerosis; TMEV model; Neuroinflammation; Neuroprotection
Introduction

Multiple sclerosis (MS) is a chronic inflammatory disease of the central nervous system (CNS) associated with demyelination and axonal damage that impair the normal neurotransmission leading to sensory deficits and deteriorated motor coordination. While the etiology of MS remains elusive, myelin destruction may be a direct consequence of the autoimmune response against myelin epitopes, or may be a neurodegenerative disease where oligodendrocytes get progressively impaired and ultimately death (Ludwin, 2006; Kipp et al., 2012).

Therapeutic strategies for the treatment of MS mainly include immune-based therapies that may have severe drawbacks such as lack of efficacy in long term treatment (Curtin et al., 2014). Moreover, clinical trials of such agents in primary progressive multiple sclerosis (PPMS) have failed and there is limited evidence of their efficacy in secondary progressive disease (Comi, 2013). Today there is an urgent need to discover new neuroprotective treatments able to modify the course of MS, both in the primary and secondary progressive forms, and being the major cause of neurological disability in young adults (Karussis, 2014).

The best animal models used to validate the therapeutic effect of new drugs in MS are the experimental allergic encephalomyelitis (EAE) and the Theiler’s murine encephalomyelitis virus-induced demyelinating disease (TMEV-IDD) (McCarthy et al., 2012). Specifically, IFNβ was one of the first molecules investigated in EAE models which gave satisfactory results in terms of its therapeutic activity in patients. IFNβ short treatment was also effective in TMEV-IDD modulating the expression of the major histocompatibility complex class II (MHC-II) and hence the action of CD8+ T lymphocytes but showed the first problems associated with long term treatments (Njenga et al., 2000). TMEV-IDD infected mice exhibit several clinical deficits such as
progressive impaired motor coordination, incontinence, and paralysis associated with axonal loss and electrophysiological abnormalities (Tsunoda et al., 2010; Sato et al., 2011). This mice model closely resembles the primary and progressive clinical form of MS (PPMS) (Mecha et al., 2013), being an excellent pharmacological tool to discover new treatments with innovative mechanism of action for this rare and severe variant of MS.

Phosphodiesterases (PDEs) are a family of enzymes that hydrolyses the nucleotides cyclic adenosine 3’,5’-monophosphate (cAMP) and cyclic guanosine 3’,5’-monophosphate (cGMP) and are classified into 11 groups on the basis of their sequence homology, cellular distribution, substrate specificity and sensitivity to different PDE inhibitors (Bender et al., 2006). cAMP is a second messenger involved in a variety of cellular responses including inflammation and immunomodulation, and the intracellular levels of this nucleotide should be closely controlled. Specifically, an increase in intracellular levels of cAMP is usually accompanied by an inhibition of certain functions of different cell types involved in the immune response and inflammation (Tasken et al., 2006; Brudvik et al., 2012). Thus, cAMP-specific PDEs inhibitors, such PDE4 and PDE7 inhibitors, have been recently emerged as promising therapeutic agents (Martinez et al., 2014; Maurice et al., 2014). Moreover, and regarding multiple sclerosis, cAMP cascade regulates myelin phagocytosis in microglia and macrophages (Makranz et al., 2006) being also involved in myelin formation (Malone et al., 2013).

The most studied enzymes in relation to its regulatory effect on immune and central nervous system are the cAMP-specific PDE4 and PDE7. In immune cells, inhibition of PDE4 limits the production and release of proinflammatory cytokines from activated peripheral mononuclear cells, including but not limited to, TNFα, IL-2, IL-12 and IFNγ.
(Kaminuma et al., 1998; Muise et al., 2002; Claveau et al., 2004). In addition, the prototypical PDE4 inhibitor, Rolipram, has demonstrated to be a potent anti-inflammatory agent in the EAE model of MS (Sommer et al., 1995; Sanchez et al., 2005; Paintlia et al., 2009; Gonzalez-Garcia et al., 2013) but its secondary emetic effects in humans have hindered its clinical development (Robichaud et al., 2001). An alternative approach is to inhibit other cAMP-specific PDE isoenzyme expressed in immune and neural cells such as PDE7 (Lee et al., 2002). PDE7 participates in the modulation of critical immune processes such as T cell proliferation (Goto et al., 2009) and, PDE7 isoenzymes, PDE7A and PDE7B, have been found in the CNS of rodents (Miro et al., 2001; Morales-Garcia et al., 2011; Johansson et al., 2012). Moreover, recent studies showed that PDE7 inhibitors could play an anti-inflammatory and neuroprotective role in cellular and animal models of Parkinson’s disease (Morales-Garcia et al., 2011), spinal cord injury (Paterniti et al., 2011), Alzheimer’s disease (Perez-Gonzalez et al., 2013) and stroke (Redondo et al., 2012c). In addition, different chemically diverse PDE7 inhibitors have shown efficacy in EAE model of MS (Redondo et al., 2012a; Redondo et al., 2012b) without emetogenic activity (Garcia et al., 2014) pointing to a new era of innovative drugs. In this regard, the therapeutical potential for MS treatment of the thioxoquinazoline derivative TC3.6 synthesized in our laboratory has been recently reported. TC3.6 is able to prevent EAE, reduce IL-17 levels, prevent infiltration in CNS and increase the expression of the T regulator cell marker Foxp3 (Gonzalez-Garcia et al., 2013). Moreover, TC3.6 have recently shown to be an effective drug for promote olygodendrocyte differentiation (Medina-Rodriguez et al., 2013).

With the aim to complete TC3.6 therapeutic profile for MS, here, we present its biological behavior regarding its modulation of cAMP levels and inflammatory
response in microglial cells; its neuroprotective capability in PC12 cells and its efficacy in the primary progressive TMEV-IDD model. In particular, we have assessed the following endpoints: neurological deficits, histological damage, leukocyte infiltration, and pro-inflammatory mediators with the aim to show the therapeutic potential of PDE7 inhibitors, more specifically TC3.6, for this unmet and severe disease.
Material and methods

PDE inhibitors. The PDE inhibitors here used were purchased from Calbiochem (BRL50481) or synthesized according reported procedures (TC3.6) (Redondo et al., 2012c). Purity of the compound (>95%) was assessed by HPLC.

Primary cultures of glial cells. Rat primary microglia was prepared from neonatal (P2) rat cerebral cortex as previously described (Mecha et al., 2011). Briefly, after removal of the meninges the cerebral cortex was dissected and mechanically dissociated by gently pipetting through tips of small diameter. After centrifugation, the pellet was washed with cold HBSS (Gibco) and the cells were plated on 75 cm² flasks and allowed to grow in DMEM medium containing 10% bovine serum (FBS), 10% horse serum and 1% penicillin/streptomycin at 37 °C in a humidified incubator (5% CO₂). After 7 days, cultures were confluent and the flasks were agitated on an orbital shaker for 4 hours at 230 rpm at 37 °C and microglial cells collected from supernatant. After centrifugation at 1050 rpm for 10 minutes, microglial cells were seeded in 96-well dishes. The purity of the cultures was >95%, as determined by immunofluorescence analysis using anti-CD11b antibody.

cAMP assay. To determine the levels of cAMP, microglial cultures were incubated with TC3.6 or BRL50481 (all at 10 µM or 30 µM) for 1h and cAMP quantified as previously described (Morales-Garcia et al., 2011).

Nitrite Determinations. For nitrites quantification TC3.6 (30 µM) was added to the culture medium of microglial cells 1 hour before exposure to LPS (10 µg.mL⁻¹), and cells were harvested 16 h later for evaluation of nitrite levels. To analyze the role of
cAMP, some microglial-containing plates were also preincubated with the cAMP antagonist Rp-cAMP (100 µM, BIOMOL Research Laboratories) or the PKA inhibitor H-89 (20 µM; BIOMOL Research Laboratories) 2 h before the addition of the compounds. The production of NO was monitored by measuring the content of nitrite in media by the standard Griess reaction as previously described (Morales-Garcia et al., 2011).

**PC12 cell cultures.** PC12 cell line was a kindly gift of Dr. Lorenzo Romero (Experimental Neurology Unit, Hospital Nacional de Parapléjicos, Toledo, Spain). The cell line was maintained in DMEM (Invitrogen) supplemented with 10% inactivated fetal bovine serum (FBS) plus 1% v/v penicillin/streptomycin (Life Technologies, Carlsbad, CA, USA) at 37°C in a 5% humidified incubator. Cells were maintained for 6 days until the experiment. Cells were seeded in 96 plate wells (2x10^4 cells/well). First, cells were treated with TC3.6 (10, 30 or 60 µM) to assess the effect of the compound in basal cell survival. In other set of experiments cells were exposed to glutamate (10 mM) for 24 h incubation and 30 min before treated or not with TC3.6 at the doses of 10 or 30 µM.

**Cytotoxicity assay.** PC12 cell death was quantified by measurement of lactate dehydrogenase (LDH) release from damaged cells into the bathing medium by using the LDH kit (Roche, Life Science, Spain). Cells were exposed to glutamate (10 mM) in the absence or presence of TC3.6 (10 and 30 µM) and 24 hours later supernatants were collected and analyzed. TC3.6 was added 30 min before glutamate. Background of LDH levels was determined in paralleled cultures subjected to sham washes and subtracted.
from experimental values. Percentage of cell death in experimental conditions was referred to the value of glutamate alone (set as 100%).

*Animal and Theiler's virus inoculation.* All experimental procedures followed the European Communities Council Directive 2010/63/EU and the Spanish regulations (BOE67/8509-12; BOE1201/2005) on the use and care of laboratory animals, and approved by the local Animal Care and Ethics Committee of the CSIC. We used female SJL/J mice, susceptible to TMEV-IDD development, from Harlan Interfauna Ibérica (Barcelona, Spain). The mice were housed in the Animal Resource Facility at the Cajal Institute, CSIC (Madrid) and maintained on food and water *ad libitum* in a 12 hours dark-light cycle. Four-to six week-old mice were inoculated intracerebral in the right cerebral hemisphere with $10^6$ plaque forming units (PFU) of Daniel’s (DA) TMEV strain as previously described (Arevalo-Martin *et al.*, 2003). Another set of mice (four to six week-old) were injected intracerebral with 30 µl of Dulbecco’s modified Eagle’s medium supplemented with 10% of fetal calf serum as Sham animals.

*Experimental in vivo procedure.* Two protocols of TC3.6 administration were used. i) During the pre-symptomatic phase (30 days post-infection) TMEV infected mice received TC3.6 (10 mg·kg$^{-1}$ i.p.; n=6) or appropriate vehicle (0.2% DMSO, Tocrisolve 5% in phosphate-buffered saline; n=6) for 12 consecutive days and then, animals were maintained and sacrificed at 70 days after TMEV infection; ii) At 60 days after TMEV infection and once established disease the mice were treated daily for 14 days with TC3.6 (5 mg·kg$^{-1}$, i.p.; n=10) or appropriate vehicle (0.2% DMSO, Tocrisolve 5% in phosphate-buffered saline; n=10) and sacrificed at day 75 post-infection.
Evaluation of symptomatology and motor function. General health conditions (weight and clinical score) and motor function of animals were periodically evaluated every 5 days from day 30 until day 50 and every 10 days from day 50 to 70 (presymptomatic treatment) or 75 (established disease) after TMEV infection. Clinical scores were assigned based on the classical scale of 0-5: score 1, mice show waddling gait; score 2, mice show more severe waddling gait; score 3, mice had a loss of righting ability associated with spastic hind limbs; score 4, mice had paralysis of hind limbs; and score 5, mice were moribund.

The screening for locomotor activity was performed using an activity cage (Activity Monitor System Omnitech Electronics, Inc., Colombus, OH, USA) coupled to a Digiscan Analyser. The number of times that the animals broke the horizontal or vertical sensor beams was measured in two 5 min sessions for horizontal and vertical activity.

Tissue processing and immunohistochemistry. Spinal cords were removed from sham and infected mice at the indicated days after the infection (n=4 of each group). Animal tissue was processed as previously described (Mestre et al., 2009). Free-floating transversal thoracic spinal cord sections (30 µm thick) were processed as detailed to visualize the proinflammatory cytokines with specific anti-COX-2 antibody (Santa Cruz Biotechnology, Inc. CA, USA, goat polyclonal IgG, 1:500 in PBS, v/v), anti-TNF-α ligand antibody (Santa Cruz Biotechnology, Inc. CA, USA, goat polyclonal IgG 1:500 in PBS, v/v), anti-IL-1β ligand antibody (Santa Cruz Biotechnology, Inc. CA, USA, rabbit polyclonal IgG 1:500 in PBS, v/v). Microglial cells were stained with a rabbit polyclonal anti-Iba-1 antibody (1:1000; Wako Chemical Pure Industry, GmbH); CD4 and CD8 positive T cells were stained with rat anti-mouse polyclonal antibodies
Axonal density was visualized with pan neurofilament primary antibody (rabbit polyclonal antiserum cocktail to neurofilaments, NA1297; 1:2000 Biomol International) and axonal neurofilaments in longitudinal sections of spinal cord were stained with a rabbit polyclonal antibody anti-Neurofilament H (1:1000; Merck Millipore, Spain). Longitudinal sections were also stained with RIP a monoclonal antibody isotype IgG1 (1:1000; DSHB University of Iowa Ames, IA, USA).

In all cases specificity of staining was confirmed by omitting the primary antibody. Six spinal cord sections per animal (n=4 of each group) were analyzed and two microphotographs per section were taken. Quantification of staining was performed using the Image J software designed by National Institutes of Health.

Eriochrome cyanine staining. Myelin staining was performed as previously described (Arevalo-Martín et al., 2010). Six spinal cord sections per animal (n=4 of each group) were analyzed and two microphotographs per section were taken. Briefly, the slides were dried and warmed at 37 °C, immersed in acetone for 5 min, and stained in eriochrome cyanine solution for 30 min at room temperature (rt). Slides were rinsed in water, differentiated in 5% iron alum for 10 min at rt, rinsed again in water, and fully differentiated in borax-ferricyanide solution for 10 min at rt. Slides were then dehydrated through graded ethanol solutions, immersed in xylene, and coverslipped using DePeX mounting medium.

Mouse Neurofilament ELISA assay Levels of mouse neurofilament were determined in tissue homogenates of spinal cord (n=6 per group) by a commercial Kit (My Biosource, San Diego, CA, USA) following the manufacturer’s recommendations. The detection
range was between 20ng/ml to 0.312 ng/ml. The sensitivity of the assay was 0.06ng/ml and the intra and inter assay coefficients of variations were 8% ± 1.2 and 12 % ± 2.3, respectively. Results were expressed as ng/mg protein.

**RNA extraction.** After saline perfusion, tissue samples were frozen in dry ice and stored at −70 °C until required (n=6 of each group). Total RNA was extracted using the RNeasy Lipid RNA extraction kit (Qiagen, UK), given the high content of lipids found in the spinal cord. Contaminating genomic DNA was degraded by a treatment with DNaseI (Qiagen, UK). The yield of RNA was determined using a Nanodrop® spectrophotometer (Nanodrop Technologies).

**Reverse transcription (RT) and real-time polymerase chain reaction (PCR).** Total RNA (1 μg in 20 μL) was reverse transcribed into cDNA using the Promega reverse transcription kit (Promega, Spain) with poly-dT primers and amplified with the primers summarized in Table 1. All gene expression was quantified using SYBR Green. Cycling conditions were as follows: 50 °C for 2 min, 95 °C for 10 min, followed by 40 cycles of amplification (95 °C for 15 s, 60 °C for 1 min). Samples were assayed on the Applied Biosystems PRISM 7500 Sequence detection system. Each sample was assayed in duplicate and a 6-point standard curve was run in parallel. To ensure the absence of genomic DNA contamination, a control sample of non-reverse-transcribed RNA was run for each set of RNA extractions. Relative quantification was obtained by calculating the ratio between the values obtained for each gene of interest and the house-keeping gene GAPDH.
Table 1. Primers used in SYBR PCR

<table>
<thead>
<tr>
<th>Gene</th>
<th>Sense (5'→3')</th>
<th>Antisense (5'→3')</th>
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<tr>
<td>IL1β</td>
<td>TGGTGTGTGACGTTYCCCATT</td>
<td>TCCATTGAGGTGGAGAGCTTTC</td>
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<tr>
<td>TNFα</td>
<td>AGAGGCACCTCCCCAAAAGA</td>
<td>CGATCACCCCCGAAGTTGTCAGT</td>
</tr>
<tr>
<td>IL6</td>
<td>TCCAGAAACCGCTATGAAGTC</td>
<td>CACCAGCATCAGTCCCCAAGA</td>
</tr>
<tr>
<td>IFNγ</td>
<td>GGCCATCAGCAACACAATAAGCGT</td>
<td>TGGGTTGTTGACCTCAAACTTGCC</td>
</tr>
<tr>
<td>GAPDH</td>
<td>TGATGCTGGTGCTGAGTATGTCG</td>
<td>TCTCCTGGGTCACACCACCCTCAAC</td>
</tr>
</tbody>
</table>

Statistical analysis. For microglia in vitro experiments, values represent the means ± SD of six replications in three different experiments and were analyzed by one way analysis of variance (ANOVA) and the Tukey’s test for multiple comparisons. For experiments with PC12 cells values represent the mean ± SEM of three independent experiments in triplicate. For in vivo experiments, values represent the mean ± SD and ANOVA for repeated measures followed by appropriate post-hoc test were used to determine statistical significance (95%; p<0.05). The statistical analysis in the first in vivo experiment (treatment in presymptomatic phase) assessed differences in the day of disease onset between TMEV+vehicle and TMEV+TC3.6 groups, as well as the clinical score and motor activity deficit progression of each group. The statistical analysis of second in vivo experiment (treatment in established disease) showed the differences in the clinical score and in the motor performance in activity cage at day 75 after TMEV infection.
Results

TC3.6 increases the levels of cAMP on microglia cultures promoting an anti-inflammatory response. TC3.6 is a synthetic heterocyclic small molecule with an IC$_{50}$ of 0.55 μM on PDE7 inhibition (Fig.1A). It is rather selective regarding other isoenzymes such as PDE3, PDE4B, PDE4D and PDE10, with IC$_{50}$ values of 70.7, 57.9, 23.9 and 50.1 μM respectively. Here, we have tested if TC3.6 has the expected cellular behaviour being able to modulate cAMP levels. Briefly, cells were seeded at 3x10$^4$/well in 96-well dishes and incubated overnight before the assay. After 15 min incubation with 10 μM or 30 μM of TC3.6, intracellular levels of cAMP were significantly increased (Fig. 1B). In line with this result, the treatment with a commercial PDE7 inhibitor (BRL50481) used as standard reference showed the same increase of the intracellular cAMP at doses of 10 μM and 30μM. The antinflammatory potential of TC3.6 was also determined on microglial cells exposed to lipopolysaccharide (LPS) by measuring nitrites accumulation. To analyze the role of cAMP, in nitrite changes some microglial-containing plates were also preincubated with the cAMP antagonist Rp-cAMP (100 μM) or the PKA inhibitor, H-89 (20 μM) for 24 h before the addition of the different compounds. In these conditions the antinflammatory properties of TC3.6 (30 μM) were completely abolished (Fig. 1C).
Fig. 1 TC3.6 increases the levels of cAMP on microglia culture modulating the anti-inflammatory response. A) Chemical compounds used. B) Intracellular levels of cAMP measured in rat primary microglial cultures in the presence of TC3.6 and BRL50481 (10 or 30 µM). Values represent the means ± SD from three different experiments. *p<0.05, **p<0.01 vs. control. C) Nitrite production in rat primary microglial cultures treated with LPS (10 µg/ml) in the absence or presence of TC3.6 and BRL50481 (30 µM). Some of the cultures were pretreated with the protein kinase A inhibitor H-89 or the cAMP antagonist Rp-cAMP. Values represent the means ± SD of six replications in three different experiments. ***p<0.001 vs LPS-treated cells.

TC3.6 mediates neuroprotective effects. Since TC3.6 present an anti-inflammatory profile, we decided to explore its potential neuroprotective properties by using PC12 cell cultures subjected to glutamate-induced cell death. PC12 cells were treated with
glutamate (10mM) and lactate dehydrogenase (LDH) release was quantified as a cell death read-out. First, we assessed whether TC3.6 per se could affect the viability of PC12 cells. As shown in Fig 2A, the higher dose of TC3.6 (60µM) reduced PC12 cell survival; thus, we discarded the use of 60µM for glutamate experiments. Exposure to glutamate induced the LDH release to a level that we set as the highest cell death reached (100%). When the cells were pre-treated with 10µM or 30µM TC3.6, glutamate-induced cell death was diminished by 40 and 60% respectively, thus TC3.6 exerted a dose-response neuroprotective activity in PC12 cells (Fig 2B).

**Fig.2 Neuroprotective activity of TC3.6.** A) PC12 cells were incubated for 24 h with TC3.6 (10, 30 and 60µM) and cell viability was quantified; results are shown as mean ± SEM from 3 independent experiments in triplicate. Statistics: ***p<0.001 vs control. B) PC12 cells were cultured and incubated with 10mM glutamate for 24 h to induce cell death in the absence or presence of TC3.6 (10 and 30µM) added 30 min before glutamate; results are shown as mean ± SEM from 3 independent experiments in triplicate. Statistics: **p<0.01, *p<0.05 vs control and # p<0.05 vs glutamate +TC3.6 (10µM).

*Inhibition of PDE7 improves motor function during established neurological symptomatology.* TMEV-IDD is a model of chronic progressive MS in which following virus inoculation there is a latency period until the symptoms and motor deficits appear.
In our first studies, TC3.6 administration during the pre-symptomatic phase (30 days post-infection) for 12 consecutive days (10 mg.kg⁻¹, i.p.) was capable to significantly delay the TMEV-IDD onset at 60 days post infection instead of 45 days in case of TMEV vehicle mice (Fig. 3A). Moreover, at 50, 60 and 70 days post-infection the treatment with TC3.6 reduced the clinical scores in comparison to those observed in TMEV vehicle mice at the same day indicating a slow course evolution of the disease.

Data from motor performance in the activity cage revealed an improvement in motor function by TC3.6 treatment (Fig. 3B and 3C). Deambulatory activity displayed by TMEV-infected mice receiving the inhibitor of PDE7 was significantly higher at 45, 60 and 70 days post-infection when compared to the horizontal activity data obtained from TMEV-vehicle mice at the corresponding days (Fig. 3B). This improved behaviour was also noticed when we assessed the vertical activity parameter (Fig. 3C).
Fig. 3 The treatment with TC3.6 delays TMEV-IDD progression. TMEV-infected mice were treated with TC3.6 (10 mg.kg\(^{-1}\)) during pre-symptomatic phase (30 days post-infection) for 12 consecutive days, then were maintained and sacrificed at 70 days after TMEV infection. A) Clinical score of animals were recorded every 5 days from day 30 until day 70 after TMEV infection. Statistics: ++p<0.001 vs. TMEV+vehicle (30 days); ^^p<0.01 vs. TMEV+TC3.6 (30 days); # p<0.05 vs. TMEV+vehicle (at the same day). The motor function was analyzed (days 30, 45, 60 and 70 post-infection) by activity cage including the horizontal activity B) and the vertical activity C) for 10 minutes. Statistics: ++p<0.001 vs. TMEV+vehicle (30 days); ^^p<0.01 vs. TMEV+TC3.6 (30 days), ANOVA for repeated measures ###p<0.01 vs. TMEV+vehicle (at the same day).

Therefore, in an attempt to study the efficacy of TC3.6 during established disease for the therapeutic interest of its potential utility, we designed a set of experiments in which the inhibitor of PDE7 was administered to diseased mice showing significant neurological deficits. Clinical symptoms and motor function of TMEV-infected mice were recorded every 5 or 10 days starting from day 30 post infection. The treatment of
TMEV-infected mice with TC3.6 (5 mg.kg\(^{-1}\)), at day 60 post-infection for 14 consecutive days slowed the clinical score progression. Thus, the clinical score reached by TMEV-vehicle mice was significantly higher (Score=3) that TMEV-TC3.6 treated mice (Score<2) (Fig. 4A). In support of this, the evaluation of spontaneous motor activity, at day 75 postinfection, showed that the administration of TC3.6 significantly improved the performance of the activity cage test as horizontal and vertical activity parameters were better in mice that received the inhibitor of PDE7 (Fig. 4B and 4C, respectively).

**Fig. 4** TC3.6 improves motor function during established neurological symptomatology. Once established disease (60 days postinfection) TMEV-infected mice were treated with
TC3.6 (5 mg.kg$^{-1}$) daily for 14 days or appropriated vehicle. A) Clinical score of animals were recorded every 5 days from day 30 until day 50 and every 10 days until day 75 after TMEV infection. Statistics: **p<0.01 vs. *p<0.05 vs Sham; ++p<0.001 vs. TMEV+vehicle (day 60), ANOVA for repeated measures. ##p<0.01 vs. TMEV+vehicle (day 75), ANOVA. The motor function was assessed by activity cage including the horizontal activity B) and the vertical activity C) for 10 minutes. Statistics: **p<0.01 vs. Sham; ANOVA for repeated measures. #p<0.05 and ##p<0.01 vs. TMEV+vehicle (day 75). Each mice was represented by a circle (O) and the mean of each group was represented by a line (—).

*TC3.6 administered on established disease reduces inflammation in the spinal cord of TMEV-infected mice.* Immunohistochemical analysis of the spinal cord of TMEV-infected mice revealed that the treatment with TC3.6 induced a significant attenuation in the level of neuroinflammation. H&E staining revealed that infection with TMEV provoked the infiltration of immune cells into the spinal cord (Fig. 5A). Treatment with TC3.6 partially diminished this infiltration (Fig. 5A) and specifically decreased the accumulation of CD4 and CD8 lymphocytes into the parenchyma (Fig. 5B). In Fig 5C is shown a reduction in microglial reactivity (p<0.05) in the spinal cord of TC3.6-treated mice in comparison to microglial images obtained from spinal cord sections of TMEV-infected mice that received vehicle. Quantification analysis revealed significant reduction in cell infiltrates (Fig 5D; p<0.01), CD4$^+$ (Fig 5E; p<0.05) and CD8$^+$ T lymphocytes (Fig 5F; p<0.05) in TMEV-infected mice treated with TC3.6 vs TMEV-infected mice treated with the vehicle. In addition, it was observed a reduction in the percentage of Iba-1 staining (Fig 5G; p<0.05) in TMEV mice treated with TC3.6 vs TMEV+vehicle mice. These findings support the data from our in vitro studies with primary microglial cultures.
Fig. 5 TC3.6 administered on established disease reduces inflammation in the spinal cord of TMEV-infected mice. Representative sections of ventral columns of thoracic spinal cord of TMEV-infected mice subjected to vehicle or TC3.6 treatment (dose-schedule: 5 mg.kg⁻¹ daily for 14 consecutive days). Sections were stained A) by hematoxiline/eosine to visualize leukocyte infiltrates; B) by an antibody against CD4 T cells and C) by Iba-1 for microglia. Quantification analysis of cell infiltrates (D), CD4⁺ T cells (E), CD8⁺ T cells (F) and percentage of area stained by Iba-1 (confocal images with constant laser beam). Statistics: **p<0.01; +p<0.05 vs TMEV+vehicle. Scale bars, 50 or 100µM

PDE7 inhibition diminishes the expression of proinflammatory mediators in the spinal cord of TMEV-infected mice. Given the reduced microglia reactivity by TC3.6 treatment, we then assessed the expression of proinflammatory markers like COX-2 and proinflammatory cytokines in the TMEV-IDD model. As shown in Fig. 6A immunohistochemical staining of IL-1β, TNFα and COX-2 was drastically reduced in
the spinal cord of TMEV-infected mice treated with TC3.6. Analysis of mRNA expression, by real time PCR, of the proinflammatory cytokines IL1β, TNFα, IL6 and IFNγ, showed a significant induction in the spinal cord of TMEV-infected mice. Accordingly with immunohistochemical analysis, transcripts for IL-1β and TNFα were significantly reduced in the spinal cord of TMEV-infected mice subjected to TC3.6 administration (Fig. 6B and 6C). The expression level of IL-6 mRNA was also significantly decreased (Fig. 6D), but levels of IFN-γ, although lower than those observed in TMEV-vehicle mice, did not reach statistical significance (Fig. 6E).
The treatment with TC3.6 diminishes the expression of proinflammatory mediators in spinal cord of TMEV-infected mice. A) Representative sections of ventral columns of thoracic spinal cord (at day 75 postinfection) of TMEV-infected mice

**Fig. 6**
treated with TC3.6 or appropriated vehicle for 14 consecutive days were stained for IL1β, TNFα and COX-2. Scale bar 100 µm. The effect of TC3.6 on IL1β (B), TNFα (C), IL6 (D) and IFNγ (E) mRNA levels in thoracic sections of spinal cord were analysed by semi-quantitative PCR. Each mice was represented by a circle (O) and the mean of each group was represented by a line (─) (n=6). *p<0.05 and **p<0.01 vs. Sham; #p<0.05 and ##p<0.01 vs. TMEV+vehicle.

**PDE7 inhibition reduces demyelinated lesions and improves spinal cord white matter disorganization increasing neurofilament in the spinal cord of TMEV-infected mice.** The data of reduced infiltrates and microglia reactivity by the treatment with TC3.6 were supported by histological examinations of spinal cord sections from experimental animals. Staining was performed using eriochrome of cyanine to visualize demyelination. Sections from spinal cord of TMEV-infected mice given vehicle show demyelination and vacuolization (Figure 7A) that was much lesser in spinal cord sections from TC3.6-treated TMEV mice. Previous studies from our group (Loria et al., 2010; Mecha et al., 2013) have shown that at the chronic symptomatic phases there is a reduction in the axonal density by using pan neurofilament staining of transversal spinal cord sections. The same was observed in the present study (Fig 7B). However, the treatment with TC3.6 improved noticeably the white matter disorganization. The quantification of intensity of staining by pan neurofilament revealed a significant effect (p<0.05) of TC3.6 treatment on the preservation of the axonal package (Fig 7D). Similarly, the axonal swelling evident when longitudinal spinal cord sections of TMEV-infected animals were stained with Neurofilament-H and CNPase for myeling labeling (Fig. 7C) was less frequent and milder following TC3.6 administration. Although the quantification of axonal loss by using electronic microscopy is considered to be the best technique, biochemical methods by measuring neurofilament levels in the spinal cord tissue homogenates were also employed to estimate neurodegeneration (Petzold et al., 2003). Here, using ELISA assays we showed that the treatment with TC3.6 increased
the levels of neurofilament (p<0.05) in spinal cord in comparison to the levels observed in TMEV-infected mice given vehicle (Fig 7E)

**Fig. 7** Effects of inhibition of PDE7 by TC3.6 in demyelination and neurodegeneration in the TMEV-IDD model. A) Representative transversal sections of ventral columns of thoracic spinal cord of TMEV-infected mice treated with TC3.6 or appropriated vehicle stained by eriochrome cyanine to visualize myelin; scale bar 50 μm. B) Representative transversal sections of ventral columns of thoracic spinal cord of TMEV-infected mice treated with TC3.6 or vehicle stained with pan-neurofilament; scale bars: 100 and 25 μm; confocal images with constant laser beam were analysed for quantification. Scale bars; C) Representative longitudinal sections of thoracic spinal cord of TMEV-infected mice treated with TC3.6 or vehicle stained with NFH and RIP; scale bar 50 and 10 μm. D) Intensity of pan-neurofilament staining in white matter was quantified: results are the mean ± SEM of six sections per spinal cord (4 mice) *p<0.05 vs TMEV+vehicle. E) Levels of neurofilament in spinal cord tissue homogenates were increased by treatment with TC3.6 in TMEV infected mice. Results are the mean± SEM from 5-6 mice (+p<0.05 vs TMEV+vehicle).
Discussion

Recent studies point out PDE7 inhibition as a new therapeutic approach for neuroimmune diseases (Giembycz et al., 2006; Paterniti et al., 2011; Redondo et al., 2012a; Redondo et al., 2012b; Gonzalez-Garcia et al., 2013). Our data show that inhibition of PDE7 by the compound TC3.6 administered systemically (i.p.) ameliorates clinical signs and improves motor function in the TMEV-IDD progressive model of MS. A sub-chronic schedule of TC3.6 administration was efficacious in two windows of treatment, during the pre-symptomatic phase and during the established disease. Our results also show that inhibition of PDE7 results in a reduction of cellular infiltrates, microglial immunoreactivity as well as in the expression of COX-2 and proinflammatory cytokines in the spinal cord of TMEV-infected mice. This was accompanied by reduced demyelination, preservation of axons package and increased neurofilament levels. These observations suggest that inhibition of PDE7 may have potential for alleviating MS-like pathology, and especially in the progressive state of the disease which is severe and without any treatment up to date.

It is well known the importance of controlling cAMP levels in the regulation of critical aspects of immune responses (Giembycz et al., 1992; Van Wauwe et al., 1995). Here, we show that TC3.6 elevates cAMP levels and significantly attenuates nitrite release in activated microglial cells. The reversion of the anti-inflammatory effect of TC3.6 by inhibitors of PKA suggests that the activity of PDE7 inhibitors is regulated by microglial cAMP concentration, and that its anti-inflammatory activity involves cAMP-dependent PKA activation. The neuroprotective profile of TC3.6 in glutamate induced cytotoxicity in PC12 cells is also supported by previous findings about the family of
PDEs as PDE4 and more recently PDE7 have been shown to play a role in neuroprotection and neuroinflammation (Paterniti et al., 2011). Previous studies have shown that inhibition of PDE4 with the selective inhibitor Rolipram in EAE models ameliorated the clinical course of the disease (Robichaud et al., 2001; Paintlia et al., 2009). Unfortunately, clinical trials with PDE4 inhibitors revealed the major adverse effects of these drugs, namely nausea and vomiting (Robichaud et al., 2001). PDE7 inhibitors are considered a good strategy to overcome these adverse effects with a subtle modulation on cAMP levels (Giembycz, 2005) without emetogenic activity (Garcia et al., 2014). We showed in the present study that TC3.6, a selective PDE7 inhibitor able to penetrate into the CNS (unpublished data), is efficacious in diminishing symptomatology and neuroinflammation in the Theiler’s model of MS. Several studies described changes in the expression of several PDE7 isoforms in the brain of rodents and humans in neurodegenerative diseases (Miro et al., 2001; Perez-Torres et al., 2003; Reyes-Irisarri et al., 2005). Therefore, PDE7 might be a pharmacological target in the context of neurological diseases such as MS.

PDE7 inhibitors have been shown to alleviate EAE symptomatology but the underlying histopathological changes were not described (Redondo et al., 2012a; Gonzalez-Garcia et al., 2013). A mechanism by which the compound TC3.6 ameliorated TMEV symptomatology may be related to its anti-inflammatory and neuroprotective profile. Recent data seem to support this notion. Thus, in a comparative study about Rolipram and TC3.6 effects in EAE found that both compounds prevented the clinical signs of EAE after MBP inoculation by reducing IL-17 and increasing the T regulator cell marker Foxp-3 as (Gonzalez-García et al., 2013). Here, in a viral model of MS, we found that transcript levels of TNFα, IL-1β and IL-6 were significantly decreased in the TC3.6-treated groups and immunohistological analysis also supported this
antinflammatory activity. In agreement to the above study (Gonzalez-Garcia et al., 2013) we found reduced cellular infiltrates in TMEV-infected mice treated with TC3.6. In addition, COX-2 expression was reduced in the spinal cord of TMEV-infected mice subjected to PDE7 inhibition. COX-2 immunoreactivity has been found in MS and in animal models (Mestre et al., 2006; Yiangou et al., 2006). In the Theiler’s virus model, COX-2 immunoreactivity has been mainly associated to the demyelinating process (Carlson et al., 2006). Selective PDE7 inhibitors promote oligodendrocyte precursor survival and differentiation towards myelin-forming phenotypes in both murine and human OPCs (Medina-Rodriguez et al., 2013). In the present study by immunohistochemistry techniques, confocal microscopy and biochemical assays we described an improvement in the organization of axons package and a partial recovery in neurofilament levels together a diminished demyelination in TMEV-IDD after the treatment with a selective inhibitor of PDE7. Further studies using electronic microscopy would add support to these first findings.

In summary, recent data in EAE underline PDE7 inhibition as a new therapeutic target for inflammatory diseases. Here we observed that the quinazoline derivative PDE7 inhibitor TC3.6 exhibits anti-inflammatory and neuroprotective activity, reduces neuroinflammation, decreases axonal damage and improves symptomatology and motor function in an autoimmune viral model of progressive MS. Collectively, our data confer an added value to the consideration of TC3.6 as a promising new candidate for MS management, including the primary progressive one, with both immunomodulatory and neuroprotective properties. Development of this small molecule to clinical trials will be the only way to finally translate its therapeutic potential to the patients.
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Conflict of Interest
The authors declare no conflict of interest
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Legends

**Fig. 1** TC3.6 increases the levels of cAMP on microglia culture modulating the anti-inflammatory response. A) Chemical compounds used. B) Intracellular levels of cAMP measured in rat primary microglial cultures in the presence of TC3.6 and BRL50481 (10 or 30 µM). Values represent the means ± SD from three different experiments. *p<0.05, **p<0.01 vs. control. C) Nitrite production in rat primary microgial cultures treated with LPS (10 µg/ml) in the absence or presence of TC3.6 and BRL50481 (30 µM). Some of the cultures were pretreated with the protein kinase A inhibitor H-89 or the cAMP antagonist Rp-cAMP. Values represent the means ± SD of six replications in three different experiments. ***p<0.001 vs LPS-treated cells.

**Fig.2 Neuroprotective activity of TC3.6.** A) PC12 cells were incubated for 24 h with TC3.6 (10, 30 and 60µM) and cell viability was quantified; results are shown as mean ± SEM from 3 independent experiments in triplicate. Statistics: ***p<0.001 vs control. B) PC12 cells were cultured and incubated with 10mM glutamate to induce cell death in the absence or presence of TC3.6 (10 and 30µM) added 30 min before glutamate; results are shown as mean ± SEM from 3 independent experiments in triplicate. Statistics: **p<0.01, *p>0.05 vs control and # p<0.05 vs glutamate +TC3.6 (10µM).

**Fig. 3** The treatment with TC3.6 delays TMEV-IDD progression. TMEV-infected mice were treated with TC3.6 (10 mg.kg⁻¹; n=6) or appropriate vehicle (0.2% DMSO, Tocrisolve 5% in phosphate-buffered saline; n=6) during presymptomatic phase (30 days postinfection) for 12 consecutive days, then were maintained and sacrificed at 70 days after TMEV infection. A) Clinical score of animals were recorded every 5 days
from day 30 until day 50 and every 10 days from day 50 to 70 after TMEV infection. The motor function was analyzed (days 30, 45, 60 and 70 postinfection) by activity cage including the horizontal activity B) and the vertical activity C) for 10 minutes. ++p<0.001 vs. TMEV+vehicle (30 days); ^^p<0.01 vs. TMEV+TC3.6 (30 days), ANOVA for repeated measures # p<0.05 and ##p<0.01 vs. TMEV+vehicle (at the same day).

**Fig. 4** TC3.6 improves motor function during established neurological symptomatology. Once established disease (60 days postinfection) TMEV-infected mice were treated with TC3.6 (5 mg.kg$^{-1}$; n=10) daily for 14 days or appropriated vehicle (0.2% DMSO, Tocrisolve 5% in phosphate-buffered saline; n=10). A) Clinical score of animals were recorded every 5 days from day 30 until day 50 and every 10 days from day 50 to 75 after TMEV infection. The motor function at day 75 postinfection was assessed by activity cage including the horizontal activity B) and the vertical activity C) for 10 minutes. **p<0.01 vs. Sham; ++p<0.001 vs. TMEV+vehicle (day 60), ANOVA for repeated measures. #p<0.05 and ##p<0.01 vs. TMEV+vehicle (day 75), ANOVA. Each mice was represented by a circle (O) and the mean of each group was represented by a line (─)

**Fig. 5** TC3.6 administered on established disease reduces inflammation in the spinal cord of TMEV-infected mice. Representative sections of ventral columns of thoracic spinal cord (at day 75 postinfection) of TMEV-infected mice subjected to vehicle or TC3.6 treatment (dose-schedule: 5 mg.kg$^{-1}$ daily for 14 consecutive days). Sections were stained A) to visualize leukocyte infiltrates by hematoxiline/eosine; B) CD4$^+$ T cells and C) for microglia by Iba-1 staining. Quantification analysis of cell infiltrates
(D), CD4⁺ T cells (E), CD8⁺ T cells (F) and percentage of area stained by Iba-1 (confocal images with constant laser beam). Statistics: **p<0.01; +p<0.05 vs TMEV+vehicle.. Scale bars, 50 or 100µM

**Fig. 6** The treatment with TC3.6 diminishes the expression of proinflammatory mediators in spinal cord of TMEV-infected mice. A) Representative sections of ventral columns of thoracic spinal cord (at day 75 postinfection) of TMEV-infected mice treated with TC3.6 or appropriated vehicle for 14 consecutive days were stained for IL1β, TNFα and COX-2. Scale bar 100 µm. The effect of TC3.6 on IL1β B), TNFα C), IL6 D) and IFNγ E) mRNA levels in thoracic sections of spinal cord were analysed by semi-quantitative PCR. Each mice was represented by a circle (O) and the mean of each group was represented by a line (—) (n=6). *p<0.05 and **p<0.01 vs. Sham; #p<0.05 and ##p<0.01 vs. TMEV+vehicle.

**Fig. 7** Effects of inhibition of PDE7 by TC3.6 in demyelination and neurodegeneration in the TMEV-IDD model. A) Representative transversal sections of ventral columns of thoracic spinal cord of TMEV-infected mice treated with TC3.6 or appropriated vehicle (at day 75 postinfection) stained by eriochrome cyanine to visualize myelin; scale bar 50µm. B) Representative transversal sections of ventral columns of thoracic spinal cord of TMEV-infected mice treated with TC3.6 or vehicle (at day 75 postinfection) stained with pan-neurofilament; scale bars: 100 and 25 µm; confocal images with constant laser beam were analysed for quantification. Scale bars; C) Representative longitudinal sections of thoracic spinal cord of TMEV-infected mice treated with TC3.6 or vehicle (at day 75 postinfection) stained with NFH and RIP (CNPase); scale bar 50 and 10 µm.
D) Intensity of pan-neurofilament staining in white matter was quantified. Statistics: *p<0.05 vs TMEV+vehicle. E) Levels of neurofilament (at day 75 postinfection) in spinal cord tissue homogenates were increased by treatment with TC3.6 in TMEV infected mice. Statistics: *p<0.05 vs TMEV+vehicle.