Chronic melatonin treatment rescues electrophysiological and neuromorphological deficits in a mouse model of Down syndrome

Abstract: The Ts65Dn mouse (TS), the most commonly used model of Down syndrome (DS), exhibits several key phenotypic characteristics of this condition. In particular, these animals present hypocellularity in different areas of their CNS due to impaired neurogenesis and have alterations in synaptic plasticity that compromise their cognitive performance. In addition, increases in oxidative stress during adulthood contribute to the age-related progression of cognitive and neuronal deterioration. We have previously demonstrated that chronic melatonin treatment improves learning and memory and reduces cholinergic neurodegeneration in TS mice. However, the molecular and physiological mechanisms that mediate these beneficial cognitive effects are not yet fully understood. In this study, we analyzed the effects of chronic melatonin treatment on different mechanisms that have been proposed to underlie the cognitive impairments observed in TS mice: reduced neurogenesis, altered synaptic plasticity, enhanced synaptic inhibition and oxidative damage. Chronic melatonin treatment rescued both impaired adult neurogenesis and the decreased density of hippocampal granule cells in trisomic mice. In addition, melatonin administration reduced synaptic inhibition in TS mice by increasing the density and/or activity of glutamatergic synapses in the hippocampus. These effects were accompanied by a full recovery of hippocampal LTP in trisomic animals. Finally, melatonin treatment decreased the levels of lipid peroxidation in the hippocampus of TS mice. These results indicate that the cognitive-enhancing effects of melatonin in adult TS mice could be mediated by the normalization of their electrophysiological and neuromorphological abnormalities and suggest that melatonin represents an effective treatment in retarding the progression of DS neuropathology.

Andrea Corrales¹, Rebeca Vidal^{1,2,3}, Susana García¹, Verónica Vidal¹, Paula Martínez¹, Eva García¹, Jesús Flórez¹, Emilio J. Sanchez-Barceló¹ Carmen Martínez-Cué^{1,*} and Noemí Rueda^{1,*}

¹Department of Physiology and Pharmacology, School of Medicine, University of Cantabria, Santander, Spain; ²Institute of Biomedicine and Biotechnology (IBBITEC, UC-CSIC-IDICAN). Santander, Spa BERSAM Instituto de Salud Carlos III

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Address reprint requests to Carmen Martínez-Cué and Noemí Rueda, Laboratory of Neurobiology of Learning, Department of Physiology and Pharmacology, Faculty of Medicine, University of Cantabria, C/Cardenal Herrera Oria s/n, 39011 Santander, Spain. E-mails: martinec@unican.es and ruedan@unican.es *These authors have contributed equally to this work.

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Introduction

Ts65Dn mice (TS), the most commonly used model of Down syndrome (DS), exhibit numerous phenotypic characteristics of DS, including cognitive deficits due to impairments in hippocampal morphology and function [1]. These cognitive impairments appear in early life stages [2, 3] and become more pronounced during adulthood [4, 5].

One of the neuromorphological substrates of the cognitive deficits in TS mice and in DS individuals is the hypocellularity found in different areas of the central nervous system (CNS), including the hippocampus, presumably due to reduced neurogenesis. Altered pre- and postnatal neurogenesis has been demonstrated in individuals with DS and in TS mice and has been implicated in the cognitive deficits found in both conditions [1, 6]. Restoring neurogenesis by administering fluoxetine, lithium or the α 5-selective negative allosteric modulator of the GABA_A receptor RO4938581 restores cognitive abilities in the TS mouse model of DS [2, 7, 8]. Adult hippocampal neurogenesis has an important role in the establishment of hippocampal long-term potentiation (LTP) [9, 10], which is considered to be the electrophysiological substrate of learning and memory [11]. TS mice show a marked reduction in LTP in the CA1 and DG areas that correlates with their cognitive deficits [12-15].

One of the mechanisms proposed to underlie this altered synaptic plasticity in TS animals is enhanced inhibition due to an imbalance between excitatory and inhibitory neurotransmission. TS mice have increased GABA-mediated inhibition and a concomitant decrease in glutamatergic transmission resulting in impaired hippocampal LTP [12-16]. Different pharmacological manipulations that restore or improve the balance between excitatory and inhibitory transmission rescue neural plasticity and cognitive abilities in the TS mouse model of DS [7, 17-19].

Another mechanism that has been implicated in altering cognitive and neuronal function in DS is increased oxidative stress. In early life stages, DS individuals present enhanced oxidative stress including elevated levels of lipid peroxidation [20-24], which can modify processes such as neurogenesis, differentiation, migration, net connection and neuronal survival [23-25]. In later life stages, oxidative stress can also contribute to the age-related progression of cognitive and neuronal degeneration associated with DS

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[26–28]. Both DS individuals and TS mice overexpress *SOD1*, the gene responsible for the formation of superoxide dismutase, an enzyme that modifies oxygen free radicals into hydrogen peroxide, leading to the overproduction of highly reactive oxygen free radicals. In addition, in DS, the overexpression of the *APP* gene leads to the overproduction of A β peptides and plays a pivotal role in the regulation of oxidative stress [28].

Melatonin is an indoleamine that has been consistently demonstrated to have neuroprotective effects [29–32]. Among the mechanisms underlying these effects are its free radical-scavenging and antioxidant properties [33, 34], its beneficial effect on neurogenesis [29, 30] and its ability to promote structural and functional neuroplasticity [30, 35– 38]. Furthermore, it improves cognitive deficits in various mouse models of different neuropathologies [30, 31, 39]. These studies suggest that melatonin could be a useful tool to improve the functional and neuromorphological abnormalities of the DS brain.

In a previous study, we showed that chronic melatonin treatment improves spatial learning and memory and delays the degeneration of cholinergic neurons in adult TS mice [40]. The aim of the present study was to evaluate the mechanisms by which melatonin exerts its cognitiveenhancing effects in TS mice. Therefore, we have studied the effect of chronic treatment with this indoleamine on adult hippocampal neurogenesis, on synaptic plasticity, on excitatory/inhibitory balance and on brain lipid peroxidation in middle-aged TS mice.

Material and methods

Animals and housing

This study was approved by the Cantabria University Institutional Laboratory Animal Care and Use Committee and was carried out in accordance with the Declaration of Helsinki and the European Communities Council Directive (86/609/EEC). Mice were generated by repeated backcrossing of B6EiC3Sn a/A-Ts(17<16>)65Dn (TS) females with C57BL/6Ei x C3H/HeSNJ (B6EiCSn) F1 hybrid males. The parental mouse generation was provided by the Robertsonian Chromosome Resources (The Jackson Laboratory, Bar Harbor, ME, USA), and mating was performed in the University of Cantabria animal facilities. In all experiments, TS mice were compared to euploid littermates (CO). To determine the presence of the trisomy, animals were karyotyped using real-time quantitative PCR (qPCR) as previously described [41]. Because C3H/HeSnJ mice carry a recessive mutation that leads to retinal degeneration (RD), all animals were genotyped by standard PCR to identify and exclude mice carrying this gene. Mice were housed individually and maintained under a 12/12 hr light/dark cycle. Mice were allowed free access to rodent chow and water.

In this study, two cohorts of male mice were used. In the first cohort, eight animals per group (CO-vehicle; CO-Mel; TS-vehicle; TS-Mel) were used to perform the immunohistochemical studies (neurogenesis and inhibitory and excitatory synapse markers). The second cohort (CO-vehicle = 11; CO-Mel = 9; TS-vehicle = 9; TS-Mel = 9) was used to evaluate the effects of melatonin on LTP and brain oxidative stress.

Melatonin treatment

TS and CO male mice were treated with either melatonin (Mel) or its diluents (vehicle) and assigned to one of four experimental groups: TS-Mel, CO-Mel, TS-vehicle, and (100 mg/L; Sigma-Aldrich, CO-vehicle. Melatonin Madrid, Spain) was dissolved in absolute ethanol and then added to drinking water at a final ethanol concentration of 0.06%. Fresh melatonin solution was prepared twice a week in feeding bottles that were protected from light. The estimated daily melatonin intake for each mouse was 0.5 mg, based on an average daily water consumption rate of 5 mL/day. TS and CO mice in the vehicle groups received tap water containing 0.06% ethanol. Daily water consumption was similar for all experimental groups. All mice were 6-6.5 months old at the beginning of the treatment and between 11 and 11.5 months of age at the time of histological procedures. Those animals used for electrophysiology and the ELISA assay continued receiving treatment with melatonin or vehicle until they were 12 months-old.

Hippocampal LTP recording

Mice were decapitated, and the brains were rapidly removed. The hippocampi were dissected, and 400 μ m slices were cut using a tissue chopper. Slices were allowed to recover for at least 1 hr in an interface chamber at room temperature in artificial cerebral spinal fluid (ACSF) containing (in mM) 120 NaCl, 3.5 KCl, 2.5 CaCl₂, 1.3 MgSO₄, 1.25 NaH₂PO₄, 26 NaHCO₃ and 10 D-glucose (saturated with 95% O₂ and 5% CO₂). Field excitatory postsynaptic potentials (fEPSPs) were recorded from the CA1 stratum radiatum with a glass micropipette (1–4 M Ω) containing 2 M NaCl and evoked by stimulation of the Schaffer collaterals using insulated bipolar platinum/iridium electrodes >500 μ m away from the recording electrode. The stimulus strength was adjusted to evoke fEPSPs that were equal to 50% of the relative maximum amplitude, not including any superimposed population spike. After stable baseline recordings (100 μ s pulse-duration, 0.033 Hz), LTP was induced by theta burst stimulation (TBS; 10 trains of 5 pulses at 100 Hz and intervals of 200 ms). The duration of the stimulation pulses was doubled during the tetanus. fEPSPs were amplified, bandpass-filtered (1 Hz-1 kHz) and stored in a computer using the Spike2 program (Spike2, Cambridge Electronic Design, Cambridge, UK). For analysis, fEPSP slopes were expressed as percentages of the baseline values recorded. several slices Results from were expressed as means \pm S.E.M.

Immunohistochemistry

The animals used for histology and cell counting were deeply anesthetized with pentobarbital and transcardially perfused with saline followed by 4% paraformaldehyde. After being removed and postfixed in 4% paraformaldehyde overnight at 4°C and transferred into 30% sucrose, the brains were frozen in dry ice and sliced coronally in a cryostat (50 μm thick sections). Series of brain slices were randomly made up of one section out of every nine for the immunohistochemistry protocol. A randomly chosen series was used to perform Nissl staining to calculate the subgranular zone (SGZ) total area of each mouse. The total SGZ extension was measured by the standard Cavalieri method as described previously by Llorens-Martin et al.
[42] us semiautomatic system (ImageJ v.1.33, NIH, USA, http://rsb.info.nih.gov/ij/).

Mature granule cell count

Mature granule cells in the hippocampal granule cell layer (GCL) were counted in series of one-in-nine sections stained with 4'6-diamidino-2-phenylindole (DAPI, Calbiochem, 1:1000) for 10 min in 0.1 M phosphate buffer (PB). Cell counts were performed using a previously described physical dissector system coupled with confocal microscopy [42]. Random numbers were generated to select the points at which to locate the dissectors. Six dissectors in each section were measured. A selected points, the 4 confocal microscope (Leica SPE) was directed toward a position previously established randomly inside the GCL. Next, at each point, a series of confocal images was serially recorded, keeping to the general rules of the physical dissector and the unbiased stereology. The confocal images were then analyzed on a computer with the aid of the ImageJ software (ImageJ, v. 1.33, NIH, Bethesda, MD, USA, http://rsb.info.nih.gov/ij). Every successive pair of images was used, with one considered as the reference image and the other the sample image. Next, the sample image became the reference image for the next pair of images, and so on. The cells were counted with the NIH ImageJ Cell Counter, labeling each cell on the screen the first time it appears in the series of confocal images. The software generated the total number of cells when the dissector brick was completed. For counting the mature granule neurons in the GCL, the dissector frame was a square situated randomly inside the GCL. The number of cells was then divided by the reference volume of the dissector (this parameter is the volume of a cube formed by the area of the frame multiplied by the height of the dissector) to obtain the number of cells per volume unit (cell density).

Cell proliferation in the SGZ (Ki67 immunofluorescence)

Slices were initially pre-incubated in PB with Triton X-100 0.5% and bovine serum albumin (BSA) 0.1%, and then immunohistochemistry was performed as described previously [7]. Briefly, free-floating slices were incubated with ry antibodies (rabbit anti-Ki67, 1:750; Neo Markers, diluted in PB with Triton X-100 0.5% and BSA 0.1% (PBTBSA) for 2 days at 4°C. Then, slices were incubated overnight at 4°C with secondary antibody (donkey anti-rabbit-Alexa Fluor 488, 1:1000, Molecular Probes, Eugene, OR, USA). The sections were counterstained with DAPI and mounted in gelatin-covered slides to be analyzed and photographed. The total number of Ki67-positive cells was counted in the selected sections with the help

of an optical fluor ce microscope (Zeiss Axioskop 2 plus, $40 \times$ objective) using the optical dissector method **\boxed{3}** previously described [43].

Neurodifferentiation (doublecortin/calretinin immunofluores-cence)

One-in-nine series of 50 μ m sections of mouse brains were used for the determination of cells expressing immature markers: doublecortin (DCX) and/or calretinin (CLR). Slices were initially preincubated in PBTBSA, and then dual immunohistochemistry was performed as described previously [42]. The primary antibod sed were goat anti-doublecortin (1:250; San ___ruz, VUSA) and rabbit 7 anti-calretinin (1:3000; Swant, switzerland). The primary 8 antibodies were recognized with an Alexa Fluor 594-conjugated donkey anti-goat and an Alexa Fluor 488-conjugated donkey anti-rabbit antibody (Alexa-conjugated antibodies from Molecular Probes, 1:1000). The sections were incubated with both primary antibodies at the same time and then with both secondary antibodies at the same time. The sections were then analyzed *hotographed* under a confocal microscope (Leica SP37. Quantification 9 of DCX and CLR expression was performed according to stereological procedures of physical dissector previously described by Llorens-Martin et al. [42]. At each point of the section to be registered, we counted all the immature cells in the reference area using ImageJ. Next, the number was divided by the reference area of the side of the prism aligned with the SGZ to obtain the number of cells per area unit (the reference area for the side of the prism is the length of the side of the frame multiplied by the height of the dissector). The cell density was then multiplied by the SGZ area to obtain the total number of cells for each of the different populations of immature neurons. The number of immature neurons is presented as either DCX+/ CLR- or total CLR+ (i.e. DCX+/CLR+ plus DCX-/ CLR+).

Density of glutamatergic and GABAergic synapse markers (VGLUT1 and GAD65 immunofluorescence)

One-in-nine series of 50 μ m sections of mouse brains were used for the determination of GABAergic and glutamatergic synapses. Slices were initially preincubated in PBTBSA, and then dual immunohistochemistry was performed as described previously [44]. Glutamatergic and GABAergic boutons in the molecular layer (ML) of the hippocampus were identified with anti-vesicular glutamate transporter 1 (VGLUT1, 1:2500; Chemicon, Temecula, CA, USA), a glutamatergic synapse marker, followed by Alexa Fluor 488-conjugated goat anti-guinea pig Ig (1:1000; Invitrogen, Carlsbad, CA, USA) and with antiglutamic acid decarboxylase (GAD65, 1:250; Chemicon), a GABAergic synapse marker, followed by Alexa Fluor 594conjugated donkey anti-mouse Ig (1:1000; Invitrogen).

Measurements were performed in images obtained with a confocal microscope (Leica SP5), using a 63×1.4 NA objective and a $9 \times$ zoom. For each marker, four sections per animal were used comprising the entire hippocampus, and one random area in the hippocampus per section was measured. Image analysis was performed with the aid of the NIH ImageJ software. Briefly, boutons with positive immunofluorescence (either VGLUT1 or GAD65 because these markers never colocalize) were measured separately applying the same threshold to all pictures. Images were first converted to grey scale to improve the contrast between signal and noise. Areas were measured inside a reference circle with a standard size of $325 \ \mu\text{m}^2$. A reference space was located in the inner ML of the hippocampal DG, lining the most external layer of granule neurons in the GCL. The percentage of reference area occupied by VGLUT1- and GAD65- positive boutons was calculated.

Quantitation of lipid peroxidation in brain tissue

To quantify lipid peroxidation levels, 4-hydroxynonenal (HNE) Adduct ELISA Kit (OxiSelect[™] HNE Adduct ELISA Kit, STA-338; Cell Biolabs, San Diego, CA, USA) was used to detect and measure hippocampal and cortex levels of HNE protein adducts in seven animals of each group. Briefly, tissue samples were homogenized in 1X PBS containing proteinase inhibitors (1:100, Protease Inhibitor Cocktail Set III, Merck, Darmstadt, Germany) and centrifuged at 12,000 g for 10 min. The supernatant fraction was collected and stored at -80° C. The protein concentration was determined with a Lowry protein assay, and samples were diluted to 10 μ g/mL. To quantify HNE levels, supernatant fractions were analyzed using an HNE Adduct ELISA Kit following the manufacturer's instructions. Analyses were always performed in duplicate. OD_{450} values were detected on a microplate reader (Multiskan EX; Thermo Electron Corporation, Vantaa, Finland). HNE levels were calculated according to the standard curve.

Statistical analysis

LTP data were analyzed by repeated-measures (RM)-ANOVAs ('time' × 'treatment' × 'genotype'). The remaining neuromorphological and ELISA data were analyzed using two-way ('genotype' \times 'treatment') ANOVAs. The means of each experimental group were compared posthoc by Student's *t*-test if two groups were compared or by Bonferroni tests if more than two groups were compared. All he analyses were performed using SPSS (version 21.0% for Windows. The *F* values of RM-ANOVAs, two-**10** way ANOVAs, and post hoc analysis of each independent neuromorphological and electrophysiological variable tested are shown in Table 1.

Results

Figure 1(A) shows the immunocytochemical detection of mature granule neurons (DAPI+ cells) in the GCL of the DG of TS and CO mice treated with either melatonin or vehicle. The quantitative analysis of this cell population showed that vehicle-treated TS mice had a significantly lower density of DAPI+ cells than CO animals (P = 0.03). Melatonin increased the density of DAPI+ cells in both genotypes (P = 0.003; Table 1). After chronic treatment with melatonin, the density of granule neurons was significantly increased in the GCL of TS mice to levels similar to those observed in vehicle-treated CO mice (TS-melatonin versus CO-vehicle: P = 0.25; Fig. 1B). In CO mice, melatonin slightly increased the density of DAPI+ cells, although this effect did not reach statistical significance (CO-melatonin versus CO-vehicle: P = 0.07; Fig. 1B).

Cells undergoing different stages of adult hippocampal neurogenesis were identified and quantified using specific markers for proliferation and differentiation. To estimate the density of the actively dividing cells, we used Ki67 immunohistochemistry (Fig. 2B, up and second row). The density of Ki67+ cells in the SGZ was reduced in vehicletreated TS mice compared to the vehicle-treated CO group (P < 0.001; Fig. 2C). Statistical analysis showed that melatonin treatment significantly increased the density of Ki67+ cells in TS mice compared to those of vehicle-treated TS mice (TS-vehicle versus TS-melatonin: P = 0.01; Table 1; Fig. 2C). By contrast, melatonin treatment did not modify the density of this cell population in CO mice.

Table 1. F values of RM-MANOVA, MANOVA (genotype x treatment) and post hoc analysis of each variable tested

	Genotype	Treatment	Genotype × Treatment
Inmunohistochemistry			
DAPI+ cells	$F_{1,29} = 4.99; P < 0.05$	$F_{1,29} = 11.00; P < 0.01$	$F_{1,29} = 0.05; P = 0.82$
Ki67+ cells	$F_{1,29} = 27.73; P < 0.001$	$F_{1,29} = 1.52; P = 0.22$	$F_{1,29} = 1.16; P = 0.29$
DCX+ cells	$F_{1,29} = 3.89; P = 0.059$	$F_{1,29} = 7.61; P < 0.01$	$F_{1,29} = 1.52; P = 0.22$
CLR+ cells	$F_{1,29} = 6.30; P < 0.05$	$F_{1,29} = 1.51; P = 0.22$	$F_{1,29} = 5.29; P < 0.05$
GABAergic boutons (GAD65)	$F_{1,29} = 137.95; P < 0.001$	$F_{1,29} = 0.04; P = 0.99$	$F_{1,29} = 1.02; P = 0.32$
Glutamatergic boutons (VGLUT)	$F_{1,29} = 28.49; P < 0.001$	$F_{1,29} = 13.57; P < 0.001$	$F_{1,29} = 4.46; P < 0.05$
Levels of HNE			
Cortex	$F_{1.25} = 9.64; P < 0.01$	$F_{1.25} = 6.91; P < 0.05$	$F_{1.25} = 2.20; P = 0.15$
Hippocampus	$F_{1,25} = 2.50; P = 0.12$	$F_{1,25} = 0.29; P = 0.59$	$F_{1,25} = 2.82; P = 0.10$
LTP results			
Basal	$F_{1,35} = 0.19; P = 0.65$	$F_{1,35} = 0.01; P = 0.91$	$F_{1.35} = 0.19; P = 0.65$
After theta burst stimulation	$F_{1,35} = 7.85; P < 0.01$	$F_{1,35} = 4.86; P < 0.05$	$F_{1,35} = 1.80; P = 0.18$
Post hoc comparisons between each pair	of potentiation curves	-,	
CO-Veh versus TS-Veh	$F_{1.17} = 11.09; P < 0.01$		
TS-Veh versus TS-mel		$F_{1.15} = 15.10; P < 0.001$	
CO-Veh versus CO-mel		$F_{1,17} = 0.25; P = 0.62$	
CO-Veh versus TS-mel		-, ,	$F_{1,17} = 0.19; P = 0.66$



Fig. 1. (A) Representative images of DAPI in the DG region of hippocampus of vehicle- and melatonin-treated TS and CO mice. (B) Means \pm S.E.M. of the density of mature granule neurons in the GCL of TS and of CO mice treated with melatonin or vehicle. **P* < 0.05 TS versus CO; #*P* < 0.05 melatonin versus vehicle; Bonferroni post-hoc tests after significant MANOVAs.

We then performed dual immunohistochemistry using specific markers for the determination of proteins expressed by different subpopulations of hippocampal neurons in consecutive developmental stages in TS and CO mice treated with either melatonin or vehicle (Fig. 2A). For the determination of immature neurons, we performed immunohistochemistry for the early neuronal marker DCX in the different groups of mice (Fig. 2B, third row). Vehicle-treated TS animals showed a reduced density of DCX+ cells compared to CO animals (P = 0.003), and chronic melatonin treatment rescued this alteration in TS mice (P = 0.01; Fig 2C). Indeed, after chronic melatonin treatment, TS mice did not differ in the number of DCX+/CLR- cells from vehicle-treated CO mice (CO-vehicle versus TS-Mel: P = 0.58; Table 1; Fig. 2C).

Cells undergoing later differentiation stages were evaluated using the neuronal marker CLR (Fig. 2B, fourth row). The density of CLR+ cells (Fig. 2C; including both DCX+/CLR+ and DCX-/CLR+ phenotypes) was reduced in TS mice compared to vehicle-treated-CO mice (P = 0.018; CO-vehicle versus TS-vehicle: P = 0.006). However, the density of CLR+ cells was normalized after chronic melatonin treatment in TS mice (CO-vehicle versus TS-Mel: P = 0.48; Fig. 2C).

Figure 3 shows the levels of HNE in the cortices and hippocampi of TS and CO mice. Vehicle-treated TS mice had increased levels of lipid peroxidation in the cortex (P = 0.005) and in the hippocampus (P = 0.015; Table 1). Melatonin administration significantly reduced the levels of HNE in the cortex (P = 0.015) but not in the hippocampus (P = 0.59) of TS or CO animals. However, post hoc analysis revealed that the increase in HNE levels in the hippocampus of TS-vehicle mice compared to COvehicle mice was no longer significant after melatonin treatment (P = 0.42; Fig. 3).

To assess the effect of chronic melatonin treatment on hippocampal synaptic plasticity in TS and CO mice, we used a theta burst stimulus (TBS) to induce LTP in the Schaffer collaterals-CA1 region (SC-CA1) of the hippocampus. No significant differences were found in fEPSP amplitudes in the baselines of the four groups of mice (P = 0.93, Table 1; Fig. 4). Analysis of fEPSPs in vehicle-treated TS mice revealed deficits in TBS-induced LTP in CA1 with respect to vehicle-treated CO mice (P = 0.004; Fig. 4). Melatonin produced an enhancement of LTP in TS mice (P = 0.001). These findings indicate that chronic melatonin treatment completely rescued LTP in TS mice, as the mean fEPSP slopes did not differ between melatonin-treated TS mice and vehicle-treated CO mice (P = 0.66; Fig. 4). However, chronic melatonin administration did not modify LTP in CO animals (P = 0.62).

The analysis of the number of GABAergic (GAD65+) and glutamatergic (VGLUT1+) synaptic boutons in the hippocampal ML revealed that vehicle-treated TS mice showed reduced VGLUT1 immunoreactivity (Fig. 5A, upper row) and an increased area occupied by GAD65+ boutons compared to vehicle-treated CO mice (Fig. 5A, lower row; VGLUT1; P < 0.001; GAD65; P < 0.001).

Remarkably, melatonin treatment significantly increased the percentage of the area occupied by VGLUT1 boutons in TS mice, but no changes in the GABAergic synapse marker GAD65 were observed in this group of animals (VGLUT1: P = 0.001; GAD65: P = 0.99; Table 1). In CO animals, melatonin treatment did not modify the percentage of the area pied by GABAergic or glutamatergic boutons (Fig. 5b

Discussion

In this study, we investigated the putative mechanisms that mediate the cognitive-enhancing effects of chronic melatonin treatment in adult TS mice. In particular, we studied the effects of melatonin on the main electrophysiological and neuromorphological alterations that have been proposed to underlie learning and memory deficits in the TS mouse model of DS, i.e. reduced hippocampal cellularity, adult neurogenesis, altered synaptic plasticity due to enhanced inhibition and increased oxidative stress [1, 28, 45].

Reductions in neuronal number in different areas of the CNS appear to compromise cognitive function in DS and in the TS mouse. This hypocellularity is particularly important in the hippocampus due to its major impact on spatial learning and memory. As previously shown by

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Fig. 2. (A) Time courses for the expression of specific molecular markers during different stages of adult DG neurogenesis. Modified from **IS** Contestabile et al. [8]. (B) Representative images of Ki67 (up row), coimmunostaining of Ki67 and DAPI (second row) and of immature neurons expressing DCX (third row) and CLR (fourth row) in the DG of TS and CO mice treated with vehicle or melatonin. (C) Means \pm S.E.M. of the density of Ki67+ cells and of immature neurons DCX+/CLR- and CLR+ neurons in the SGZ of the hippocampus of vehicle- or melatonin- treated TS and CO mice. ***P* < 0.01; ****P* < 0.001 TS versus CO; #*P* < 0.05, ##*P* < 0.01 melatonin versus vehicle; Bonferroni post-hoc tests after significant MANOVAs.

numerous studies [7, 43, 46], the density of mature granule neurons is reduced in the GCL of TS mice. This reduction may be due to increased cell death and/or to reduced proliferation in the SGZ. Interestingly, we found that melatonin treatment rescued the density of granule cells in TS mice.

Reduced neurogenesis is likely to be one of the mechanisms implicated in the impaired learning and memory



Fig. 3. Means \pm S.E.M. of the HNE levels in the cortex and hippocampus of vehicle-or melatonin-treated TS and CO mice (n = 7 animals of each group). **P* < 0.05; ***P* < 0.01 TS versus CO; #*P* < 0.05 vehicle-treated versus melatonin-treated mice. Bonferroni post-hoc tests after significant MANOVAs.

found in TS mice. Numerous studies have demonstrated that adult hippocampal neurogenesis is important for normal spatial learning and memory. When newborn cells are removed genetically, animals display impaired capacity for learning and memory [10, 47]. Similarly, most of the treatments that impair adult hippocampal neurogenesis also disrupt spatial cognition [47, 48]. Indeed, a significant negative correlation has been observed between performance in the Morris water maze (MWM) and the number of newly generated cells derived from adult neurogenesis in the DG of TS mice and euploid littermates [1, 49].

To evaluate the effects of chronic melatonin treatment on adult hippocampal neurogenesis in TS mice, we analyzed different subpopulations of newborn cells (Ki67positive cells) and of new neurons under different stages of differentiation (identified by DCX- and CLR-labeling). In accordance with the results of a number of studies [2, 43, 49–51], adult TS mice had a lower density of proliferating cells and decreases in both populations of immature neurons in the SGZ of the DG. This reduction in the density of proliferating and differentiating cells is likely to be responsible for the hypocellularity found in TS hippocampi.

Previous studies showed that exogenous melatonin administration rescued or attenuated the reduction of neurogenesis found during aging in normal mice [29, 37, 52] and in several models of different neuropathologies [31, 53–55]. Similarly, our data demonstrate that in adult TS mice, melatonin enhanced adult hippocampal neurogenesis by increasing the density of proliferating cells. However, melatonin did not completely restore the density of progenitors under proliferation (Ki67+ cells) in TS mice. In DS, $A\beta PP$ contributes to impaired neurogenesis by influencing neural precursor cell proliferation [56]. In our previous work [40], melatonin did not reduce the increased levels of $A\beta PP$ expression in the hippocampus of TS mice, which may contribute to the incomplete recovery of cell proliferation in these animals. In addition, cell cycle disruptions and decreased cell proliferation beginning at early developmental stages in the TS mouse model of DS [2, 46, 57]. Thus, it is also possible that because the melatonin treatment regimen in the present study started at 6 months of age, the hypocellularity established in early developmental stages was not affected by this indoleamine treatment. Further studies will be necessary to clarify whether early melatonin administration may exert larger beneficial effects on cell proliferation in TS mice.

Melatonin completely rescued the density of immature neurons undergoing different stages of differentiation by normalizing the density of cells in intermediate DCX- and CLR-expressing stages. In agreement with this result, in vitro and in vivo studies have demonstrated that melatonin predominantly facilitates the transformation of precursor cells into mature neurons in the SVZ and SGZ of the adult mice [29, 58].

Fig. 4. (A) Representative fEPSPs traces **IS** recorded before issuing the stimulus (black) and at 60 min after stimulation (red) for each experimental group. Scale bars are 1 mV and 10 ms. (B) Means \pm S.E.M. of fEPSPs (expressed as % versus basal) values from vehicle-treated CO (n = 11 slices from 11 different mice), vehicle-treated TS (n = 9 slices from 9 different mice), melatonin-treated CO (n = 9 slices from 9 different mice), respectively.





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Increase cell death is another putative mechanism implicated in the hippocampus hypocellularity and the altered cognitive abilities found in TS mice. Indeed, melatonin has been shown to induce neuroprotection due to its antiapoptotic action through the prevention of the activation of the mitochondrial pathway [59].However, previous studies showed that apoptotic cell death during development or in adulthood is not involved in the reduced number of neurons found in the TS brain [60, 61]. Therefore, it is more likely that melatonin rescues the density of mature granule neurons in the hippocampus by increasing adult neurogenesis and survival.

Both DS individuals and TS mice present enhanced oxidative stress leading to premature neuronal death and cognitive dysfunction [22, 23, 62]. Consistent with previous results [22, 62], the levels of HNE, a specific marker of lipid peroxidation, were significantly increased in the hippocampus of vehicle-treated TS mice compared to CO mice. By contrast, TS mice did not show higher levels of HNE in cortex, suggesting that this region may exhibit more tolerance against oxidative stress than the hippocampus.

Melatonin rescued the levels of HNE in the hippocampus of TS mice; thus, its cognitive-enhancing effects in trisomic animals could also be partially due to its wellknown antioxidant properties [33, 39, 63]. Although the antioxidant SGS-111, an analogue of the nootropic piracetam, administered from conception failed to improve learning and memory in the TS mouse model of DS [64], other antioxidants, such as vitamin E, delay impairment in cognitive performance, preserve markers of cholinergic cell survival, sustain hippocampal morphology and normalize oxidative stress in adult TS mice [62]. The free radical scavenging ability of melatonin is twice that of vitamin E [65], and melatonin is particularly effective in crossing the blood-barrier [66]. Thus, the reduction in lipid peroxidation in the hippocampus of TS mice after melatonin treatment might be implicated in the increase in granule cell density and the decrease in age-associated cognitive decline observed in these animals.

MANOVAs.

16 Fig. 5. (A)

animals

microscope

immunopositive for

Representative

of

VGlut1

images

glutamate synapses; up row) and GAD65 (red, GABAergic synapses; down row) in the inner ML of the hippocampal DG,

lining the most external layer of granule

neuron in the GCL of the four groups of

Means \pm S E M

percentage of total immunopositive

surface occupied by VGlut1 (B) and

GAD65 (C) boutons of vehicle- and

melatonin-treated TS and CO mice.

P < 0.01; *P < 0.001 TS versus CO;

###P < 0.001 melatonin vs. vehicle; Bonferroni post-hoc tests after significant

Numerous studies have demonstrated that TS mice present altered hippocampal synaptic plasticity that correlates with their cognitive alterations. In agreement with previous evidence [13–15], the present study showed that LTP was reduced in TS mice, and melatonin treatment for 5–6 months completely rescued LTP in the CA1 area in trisomic animals. Consistent with these findings, chronic melatonin treatment has been shown to rescue deficits in hippocampal LTP and spatial memory after intra-hippocampal injection of A β [30], although the acute application of melatonin directly to mouse and rat hippocampal slices inhibits LTP in the CA1 region [67–69].

Increasing evidence indicates that impaired LTP and the associated cognitive dysfunction in TS animals are mediated by a dysregulation between excitatory and inhibitory neurotransmission [15, 16, 19]. Consistent with previous results [7, 51], TS mice presented a reduced density of glutamatergic (VGLUT1-positive boutons) and an increased density of GABAergic (GAD65-positive boutons) synapse markers in the hippocampal ML indicating enhanced inhibition. Chronic melatonin administration to adult TS mice increased the density of glutamatergic synapse markers, thereby reducing inhibition. By contrast, this treatment did not have any effect on the area occupied by GAD65-positive boutons. Reducing synaptic inhibition by administering different antagonists and negative allosteric modulators of the GABA_A receptor has been proven to rescue synaptic plasticity, neuromorphological and cognitive alterations in the TS mouse model of DS [7, 17, 18]. In addition, LTP in the SC-CA1 pathway is known to be dependent upon the activation of NMDA receptors. Recently, a high-affinity selective GABA_B antagonist improved LTP in TS mice by increasing the NMDA receptor-mediated component [19]. Melatonin also protects and

increases NMDA receptor subunits 2A and 2B concentrations in rat hippocampus [70, 71]. Thus, at the postsynaptic level, the enhanced activation of NMDA receptors represents a possible mechanism by which melatonin could increase LTP in TS mice. Melatonin administration to TS mice might also restore LTP by increasing the density and/or activity of glutamatergic synapses. In addition, the restoration of the density of granule cells could improve afferent synaptic input from the DG and enhance the efficiency of excitatory signaling through the trisynaptic circuit, thereby enhancing LTP.

In young TS mice, several trmacotherapies, includ-ing several selective GABAL reptor antagonists, the selective GABAA a5 negative allosteric modulator RO4938581, lithium, fluoxetine and the long-acting $\beta 2$ adrenergic agonist formoterol, have been shown to improve or normalize their cognitive performance, concomitantly to their neuromorphological and electrophysiological deficits [7, 8, 19, 72, 73]. In aged TS animals, like the ones used in this study, several drugs including the uncompetitive NMDA receptor antagonist memantine, the antioxidant vitamin E and the anti-inflammatory drug minocycline have also been reported to reduce the cognitive deterioration of these mice, and some agerelated neuropathological hallmarks of DS [51, 62, 74]. However, none of these treatments have been proven to completely rescue the neuromorphorphological and electrophysiological deficits proposed to underlie DS-cognitive alterations as demonstrated in this work after chronic melatonin treatment. In addition, melatonin is normally well tolerated in adults, does not induce important side-effects and has been approved for human use; therefore, this indoleamine is a promising preventive therapy to slow the age-related progression of DS cognitive alterations.

In summary, melatonin administration to adult TS mice significantly increased the density of proliferating cells, the density of differentiating neuroblasts, and the density of mature granule cells while reducing the levels of lipid peroxidation. Furthermore, it reduced synaptic inhibition in the hippocampus of TS animals, leading to a complete recovery of LTP. Therefore, the normalization of the function and/or morphology of the hippocampus is likely to account for the cognitive-enhancing effects of chronic melatonin treatment previously demonstrated in TS mice. Our results provide evidence supporting the use of melatonin in clinical trials to alleviate cognitive deficits in the DS population prior to advanced memory decline. Regarding its efficacy as a preventive therapy in adults with DS, the initial dose, derived from this study to be administered to humans could be calculated according to the body surface area (BSA) normalization method previously described by Reagan-Shaw et al. [75].

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Author contributions

A.C. performed pharmacological treatments and histological studies. A.C. and R.V performed electrophysiological studies. P. M., S. G. and V.V. karyotyped the animals. E.G. performed ELISA studies. J. F. and E.S.-B. designed and discussed experiments. C.M.-C. and N.R. conceived the project, designed, supervised, conducted and interpreted experiments, analyzed the data and wrote the manuscript.

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