Tumor necrosis factor-α synthesis inhibitor 3,6′-dithiothalidomide attenuates markers of inflammation, Alzheimer pathology and behavioral deficits in animal models of neuroinflammation and Alzheimer’s disease

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Abstract

Background: Neuroinflammation is associated with virtually all major neurodegenerative disorders, including Alzheimer’s disease (AD). Although it remains unclear whether neuroinflammation is the driving force behind these disorders, compelling evidence implicates its role in exacerbating disease progression, with a key player being the potent proinflammatory cytokine TNF-α. Elevated TNF-α levels are commonly detected in the clinic and animal models of AD.

Methods: The potential benefits of a novel TNF-α-lowering agent, 3,6′-dithiothalidomide, were investigated in cellular and rodent models of neuroinflammation with a specific focus on AD. These included central and systemic inflammation induced by lipopolysaccharide (LPS) and Aβ1–42 challenge, and biochemical and behavioral assessment of 3xTg-AD mice following chronic 3,6′-dithiothalidomide.

Results: 3,6′-Dithiothalidomide lowered TNF-α, nitrite (an indicator of oxidative damage) and secreted amyloid precursor protein (sAPP) levels in LPS-activated macrophage-like cells (RAW 264.7 cells). This translated into reduced central and systemic TNF-α production in acute LPS-challenged rats, and to a reduction of neuroinflammatory markers and restoration of neuronal plasticity following chronic central challenge of LPS. In mice centrally challenged with Aβ1–42 peptide, prior systemic 3,6′-dithiothalidomide suppressed Aβ-induced memory dysfunction, microglial activation and neuronal degeneration. Chronic 3,6′-dithiothalidomide administration to an elderly symptomatic cohort of 3xTg-AD mice reduced multiple hallmark features of AD, including phosphorylated tau protein, APP, Aβ peptide and Aβ-plaque number along with deficits in memory function to levels present in younger adult cognitively unimpaired 3xTg-AD mice. Levels of the synaptic proteins, SNAP25 and synaptophysin, were found to be elevated in older symptomatic drug-treated 3xTg-AD mice compared to vehicle-treated ones, indicative of a preservation of synaptic function during drug treatment.

Conclusions: Our data suggest a strong beneficial effect of 3,6′-dithiothalidomide in the setting of neuroinflammation and AD, supporting a role for neuroinflammation and TNF-α in disease progression and their targeting as a means of clinical management.

Keywords: Neuroinflammation, Neurodegeneration, TNF-α, Neuroprotection, Alzheimer’s disease, Mild cognitive impairment, Amyloid-β peptide, Tau, Thalidomide, Lenalidomide
Background

Neuroinflammation is a critical and invariable component of many neurodegenerative diseases. Indeed, the co-occurrence of inflammatory markers in plasma, cerebral spinal fluid (CSF) and post-mortem brain tissue has become a hallmark feature of Alzheimer’s disease (AD) and Parkinson’s disease (PD) [1-4]. This has given rise to the suggestion that inflammation within the CNS is an important contributing factor in the continuum of neurodegeneration, particularly in AD. The cytokine tumor necrosis factor-alpha (TNF-α) is a major mediator of inflammation that is commonly upregulated in biological samples from patients and animal models of human neurodegenerative disease [5-12]. In brain, TNF-α is primarily synthesized and released by glial cells that can be activated by trauma, infection or the presence of endogenous yet abnormal protein aggregates, such as amyloid-β (Aβ) peptides in AD. Furthermore, cytokines, including TNF-α, have been demonstrated to self-regulate immune/glial cells to increase their expression of amyloid-β precursor protein and Aβ in vitro [13]. Thus, in addition to the synthesis and release of cytotoxic factors, glial cells possess the ability, under appropriate conditions, to be an important non-neuronal, source of abnormal APP in brain, a hallmark feature in AD [14,15]. The existence of an altered brain micro-environment observed in neurodegenerative disease allows for and increases the likelihood of the generation of an unregulated immune response that potentiates any ongoing toxicity toward neurons.

In human post-mortem AD brain, a dramatic loss in neurons has been observed in the medial temporal lobe (hippocampus and entorhinal cortex). Over time, AD patients develop classical patterns of brain pathology, as described by Braak and Braak [16], and corresponding cognitive impairments [14]. Early stages of dementia may be associated with neuronal dysfunction causing the observed cognitive impairments rather than overt cell loss. It is possible that neuronal dysfunction is related to the presence of inflammatory factors, as rodent models of chronic neuroinflammation have been shown to induce impaired states of learning and memory in vivo [17,18]. In line with this, the expressions of genes associated with learning and memory have been reported to be altered by LPS administration in mice [19]. Additionally, disturbances in biochemical and functional correlates of learning at the cellular level, such as long-term potentiation (LTP) [20] and neural network activity [21], have been observed in models that mimic AD pathology. These findings support the concept that inflammation exacerbates the existing processes of neurodegeneration observed in AD.

It is plausible that the dysfunction of viable neurons during early stages of AD may not represent a permanent state and may thus be amenable to rescue, potentially providing improved outcomes for long-term nerve cell survival and function. Transgenic animal models are commonly used to investigate pathological processes of neurodegenerative diseases; one such model is the 3xTg-AD mouse model [22]. Data from 3xTg-AD mice suggest a clear involvement of cytokines, particularly TNF-α at pre-symptomatic stages of AD pathology [23]. Additionally, the use of this model has shown that in some circumstances neurons express TNF-α gene products [24], and that reduced TNF-α signaling [25] and microglia activation [26] mitigated disease progression.

Herein, to further define the role of TNF-α in neuroinflammation, neuronal dysfunction and cognitive impairment, we utilized a TNF-α synthesis-lowering agent, 3,6'-dithiothalidomide, developed within our laboratory [27]. This agent has been shown to effectively lower the levels of TNF-α and nitrite, a surrogate of nitric oxide metabolism, in LPS-treated macrophage-like cells in vitro [28], to reverse established hippocampus-dependent cognitive deficits induced by chronic neuroinflammation [29], as well as to reverse learning and memory behavioral deficits in a rodent model of head trauma [30]. We therefore assessed the biochemical and behavioral actions of 3,6'-dithiothalidomide in three models of neuroinflammation and in 3xTg-AD mice to evaluate TNF-α as a neurological drug target in AD.

Methods and materials

Pharmacological interventions

3,6'-Dithiothalidomide was prepared in 100% tissue culture grade dimethyl sulfoxide (DMSO, Sigma-Aldrich, St Louis, MO) for cell culture experiments, or as a suspension in a 1% carboxy methyl cellulose/saline solution (Fluka 21901). A concentration of 56 mg/kg body weight of drug was used for animal work unless stated otherwise. Lipopolysaccharide (LPS) was obtained from Escherichia coli (E coli) serotype 055:B5 (Sigma-Aldrich). Aβ1-42 or Aβ12-1 peptides were from American Peptide, Sunnyvale, CA.

Cell culture

Mouse RAW 264.7 cells were purchased from ATCC (Manassas, VA, USA) and were grown in DMEM media supplemented with 10% FCS, penicillin 100 U/ml and streptomycin 100 μg/ml, and were maintained at 37°C and 5% CO2. Cells were propagated as described by ATCC guidelines. RAW 264.7 cells were cultured as has been previously described [28,31]. Cells were challenged with concentrations of LPS as indicated, and 24 h later, conditioned media was harvested and analyzed for the quantification of secreted TNF-α protein, nitrite and APP levels. Cellular health was assessed by use of the CellTiter 96 AQueous One Solution Cell Proliferation Assay (Promega, Madison, WI).
Acute animal LPS drug study
An in vivo assessment of the effects of 3,6′-dithiothalidomide on the biosynthesis of LPS-induced TNF-α mRNA and protein was undertaken. The levels of hippocampal mRNA, plasma and CNS protein were determined. Male Fisher 344 rats (3 months of age) were challenged with LPS (1 mg/kg body weight, via the i.p. route). A series of blood samples were taken from the rats over a 5-h time period: [−60, 0 (LPS challenge), 30, 60, 90, 120, 180 and 240 min post LPS]. Plasma was generated from blood by conventional means. After 240 min the CNS was harvested, and all samples were immediately frozen to −70°C and stored for analyses.

Chronic intracerebroventricularly animal LPS drug study
The rodents used for these experiments where male Fisher 344 rats (3 months of age). Four study groups were utilized: (1) artificial cerebrospinal fluid (aCSF) plus drug vehicle (aCSF-veh; n = 4); (2) aCSF plus 3,6′-dithiothalidomide (aCSF-DT; n = 6); (3) LPS plus drug vehicle (LPS-veh; n = 5) and (4) LPS plus 3,6′-dithiothalidomide (LPS-DT; n = 7). The LPS or aCSF were infused directly into the brain via an intracerebroventricular (i.c.v.) catheter (placement coordinates: 2.5 mm posterior to lambda, on the mid-line and 7.0 mm ventral to the dura) into the lateral ventricle as previously described [17,18,21,29]. Animals received daily i.p. administration of 3,6′-dithiothalidomide (56 mg/kg) or vehicle for 14 days starting the day of the surgery. On day 14 after surgery, each animal was placed in an open field and allowed to explore for 10 min. The open field environment consisted of a circular chamber (130 cm in diameter; 25 cm high) containing four different objects in the center. Total distance moved and time spent in different zones of the chamber were recorded with EthoVision XT software from Noldus (Noldus, Nijmegen Area, The Netherlands). All animals were euthanized by anesthesia in an isoflurane chamber followed by decapitation immediately after exploration. To ensure that transcription induced by euthanasia would not be detectable, the brain was quickly removed (between 120–150 s) and flash frozen in −80°C ice cold isopentane [17,18,21,29]. The fresh frozen brains were stored at −70°C until processing for in situ hybridization and fluorescence immunostaining as previously described [17,18,21,29].

Acute intracerebroventricular Aβ1–42 peptide animal drug study
Adult male C57BL/6 mice (3 months of age) were utilized in this study. Mice received 3,6′-dithiothalidomide (56 mg/kg, i.p.) or vehicle daily for 14 days; after 7 days of treatment with drug animals were challenged with Aβ peptide. Aβ1–42 and the reverse peptide Aβ12–42 were reconstituted in phosphate-buffered saline (pH 7.4) and aggregated by incubation at 37°C for 7 days prior to administration. Aβ1–42 and Aβ12–42 (400 pmol) were then infused i.c.v. into the lateral ventricle as previously described [32].

3xTg-AD animal drug study
Adult and old male 3xTg-AD mice (10 and 17 months of age) were maintained on a 12-h light/dark cycle with free access to water and food. Animals received administration of either 3,6′-dithiothalidomide (42 mg/kg) or the vehicle i.p., daily for 6 weeks. Four to five weeks after the initiation of drug treatment, the animals were tested in the Morris Water Maze to assess acquisition (learning) and retention of spatial memory. After the completion of the Morris Water Maze assessment, the animals were euthanized by decapitation while under isoflurane anesthesia. The brain was immediately removed, and specific regions were excised and instantly frozen to allow later quantification of levels of various cortical proteins of interest: soluble Aβ1–42, total tissue APP, tau and phospho-tau protein, and the presynaptic proteins: SNAP25 and synaptophysin.

All animal studies were undertaken according to protocols approved by the respective Institutional Animal Care and Use Committee's of the Intramural Research Program, National Institute on Aging (331-LNS-2012, 293-LNS-2013 and NICHD # 08–011), and the University of California (AN082537-03A), in compliance with the guidelines for animal experimentation of the National Institutes of Health (Department of Health, Education, and Welfare publication 85–23, revised, 1995).

Quantitative RT-PCR for rat TNF-α mRNA
Total mRNA was isolated from the hippocampus using the RNeasy RNA isolation kit (Qiagen, Valencia, CA). Various concentrations of total mRNA extracted from rat brain were prepared for the generation of absolute and relative standard curves. The RNA samples and serial dilutions for standard curves were reverse-transcribed (Applied Biosystems), and PCR reactions were carried out using primers and probe sets purchased from Applied Biosystems (Foster City, CA): the TNF-α primers and probe set recognize exon 2–3 of TNF-α (assay location 391, GenBank accession NM_012675.2), and the GAPDH primers and probe set recognize exon 3 (assay location 295, GenBank accession NM_017008.3). The signals from the amplified PCR products were detected using the ABI Prism 7700 Sequence Detection System (Applied Biosystems), and obtained Ct values were calculated to the relative amount of RNA from the standard curves for each RNA transcript.

ELISA analysis
TNF-α levels were measured by species-specific ELISAs; Mouse RAW 264.7 cell culture media TNF-α protein were measured with a mouse ELISA, BioLegend, Mouse
TNF-α ELISA MAX™ Deluxe. Rat plasma and CNS protein was measured with a rat ELISA, BD OptEIA Rat TNF ELISA Kit II, BD Biosciences or an Ultrasensitive rat TNF-α, Invitrogen, respectively. Soluble human Aβ1-42 levels were measured by use of a human Aβ42 ELISA kit from Invitrogen. For all ELISA measurements, samples were assayed in duplicate, and the appropriate procedures were followed according to the manufacturer’s instructions.

**Immunohistochemistry**

The brains of animals that received i.c.v. administered LPS or Aβ peptides were processed as previously described for the appropriate procedure [17,18,21,29,32]. For rat i.c.v. LPS studies, tissue was labeled with the following primary antibodies: MHC class II marker OX-6 (Pharmigen, San Diego, CA) for activated microglia and glial fibrillary acidic protein (GFAP; Pharmigen) for activated astrocytes; nuclei were counterstained with Sytox-Green (molecular Probes, Eugene, OR). Arc mRNA detection was performed as previously described [17,18,21]. Activated microglia numbers were quantified in the DG and CA3 area as previously described [18,29]. For mouse i.c.v. Aβ peptide studies activated microglia were identified and labeled with rabbit anti-CD11b (1:200; Chemicon, Temecula, CA). Fluoro-Jade B, a fluorochrome, was used to aid in the quantification of degenerating neurons in the dentate gyrus [33,34]. For mouse 3xTg-AD studies activated microglial cells in the subiculum and CA1 region were visualized using an anti-CD68 (Serotec) primary antibody.

**Morris water maze test**

Spatial learning and memory were assessed using the Morris Water Maze. The target platform was submerged 1 cm below the water surface and placed at the midpoint of one quadrant. Visual cues were placed around the tank to orient the mice. The acquisition training sessions took place over 4 days (for i.c.v. Aβ peptide-injected mice, 3 days after Aβ administration) or over 6 days (for 3xTg-AD mice, 4 to 5 weeks after initiation of drug treatment). The memory retention assessments were performed at 24 h (Aβ peptide-injected mice) or at 4 and 24 h (3xTg-AD mice) after the last training session. The variables of interest were mouse reference memory, the time spent in the platform area and the number of platform crossings. These events were recorded and analyzed to determine age- and drug-related differences between the groups.

**Western blotting**

Secreted APP levels were measured by Western blotting of equal volumes of harvested culture media after separation in a 10% Bis-Tris protein gel and probed with an antibody that recognizes all forms of APP (22 C11; Millipore, Billerica, MA). For 3xTg-AD mouse studies protein from each sample was separated by electrophoresis in Criterion gels (BioRad), then the transferred proteins were probed for APP (clone 6E10, Covance, Emeryville, CA), phospho and non-phospho tau (Thermo Scientific, Rockford, IL); SNAP 25 (Millipore), and synaptophysin (Santa Cruz Biotechnology, Santa Cruz, CA). Protein signals were obtained by chemiluminescence substrate methods, and signals were normalized to β-actin (GE/Amersham) [35,36].

**Statistical analysis**

GraphPad Prism software was used to perform the statistical analysis of the following variables: plasma and CNS TNF-α levels; hippocampus TNF-α mRNA; Morris Water Maze platform escape latency; time spent in quadrants and platform crossings. A one-way ANOVA was performed on each group followed by a Bonferroni’s post test or a Fisher’s post-hoc test, where appropriate. For two-group comparisons, unpaired Student’s t-tests were carried out. One-way ANOVA was performed for variables measured from RAW 264.7 cells, TNF-α, nitrite and sAPP, and were followed by Bonferroni’s post hoc tests, where appropriate. For the immunostaining analysis, the control and experimental groups were the independent variable, and the percentages of neurons expressing Arc or activated microglia from various categories were the dependent variables. When an ANOVA was significant (p < 0.001), individual between-group comparisons were performed with Bonferroni’s post hoc tests to correct for multiple comparisons. Statistical analyses are provided in each figure legend and, where appropriate, involved one- or two-tailed t-tests for specific comparisons.

**Results**

3,6'-Dithiothalidomide attenuates the synthesis of inflammatory mediators in vitro and in vivo

The lipophilic, small molecular weight (mw: 290), TNF-α synthesis-lowering agent 3,6'-dithiothalidomide [27] was utilized in a series of studies aimed at assessing the role of TNF-α on neuroinflammation directed at an AD-like phenotype. RAW 264.7 cells treated with LPS recapitulate aspects of microglial cells observed in neurodegenerative diseases exemplified by AD. A challenge of RAW 264.7 cells with increasing concentrations of LPS induced a dose-dependent generation of TNF-α protein (Figure 1A), nitrite an intermediate of nitric oxide metabolism (Figure 1A) and the secretion of amyloid precursor protein (sAPP) into culture media, suggestive of an upregulated processing of APP (Figure 1B). These changes occurred without loss of cell viability (Figure 1B). Pretreatment of cells with 3,6’-dithiothalidomide prior to LPS (10 ng/ml) administration reduced the synthesis of each factor in a dose-dependent
manner (Figure 1C and D), likewise without loss of cell viability (Figure 1D). The action of 3,6′-dithiothalidomide was additionally assessed in an acute in vivo rodent LPS model. Rats challenged with systemic LPS (i.p.) displayed a marked time-dependent increase in plasma TNF-α protein that was significantly attenuated by pretreatment with the agent (Figure 1E). Notably within the CNS, hippocampal TNF-α mRNA levels (Figure 1F) were elevated by LPS and markedly suppressed by drug treatment, as were TNF-α protein levels (Figure 1G).

3,6′-Dithiothalidomide treatment reduces LPS-induced chronic neuroinflammation and restores LPS-mediated abnormal hippocampal neuronal plasticity

The potential benefits of lowering TNF-α levels on neuroinflammation-induced altered neuroplasticity were assessed. To recreate the hostile brain microenvironment associated with chronic neuroinflammation, we utilized the properties associated with direct chronic CNS infusion of a small quantity of LPS in rats. This infusion, with and without systemic 3,6′-dithiothalidomide treatment, was well tolerated. When animals were allowed to explore an open field arena for 10 min, no differences were observed in either speed or exploration pattern (Figure 2A), indicating that such treatments had no impact on levels of anxiety or motility of animals. However, chronic i.c.v. LPS did induce a significant increase in the number of granule cells expressing the plasticity-related immediate early gene (IEG) Arc within the suprapyramidal blade of the dentate gyrus (Figures 2B and 3) compared to aCSF animals. No differences were observed within the infrapyramidal blade (not shown). The IEG Arc is required for synaptic plasticity and...
is essential for memory processing [37,38]; behaviorally induced Arc expression within the hippocampus is significantly modified by chronic inflammation [39]. Notably, the centrally mediated LPS-induced disturbances in Arc expression were abolished by systemically administered 3,6′-dithiothalidomide. In parallel with Arc mRNA levels, the number of OX-6 positive (MHC class II) microglial cells was markedly elevated in LPS-treated animals, whereas drug treatment prevented this rise (Figures 2C and 4). Furthermore, GFAP immunofluorescence staining indicated a marked upregulation in both the number and activated microglial cells.
morphology of astrocytes in response to LPS treatment; these effects likewise were significantly attenuated by 3,6′-dithiothalidomide (Figure 5).

3,6′-Dithiothalidomide treatment attenuates the effects of central administration of toxic Aβ₁₋₄₂ peptide on behavior, cell viability and microglia activation

The direct administration of aged Aβ peptide, allowing its oligomerization, into the CNS of wild-type (wt) mice, was undertaken to emulate the inflammatory microenvironment of the AD brain. In our study, a single i.c.v. administration of Aβ₁₋₄₂ was undertaken 7 days after the initiation of a daily schedule of systemic (i.p.) 3,6′-dithiothalidomide, utilizing a dose determined to be effective to ameliorate LPS-induced CNS elevations in TNF-α in the prior studies. Control animals were administered reverse peptide (with or without similar 3,6′-dithiothalidomide treatment). As illustrated in Figures 6A and B, Aβ₁₋₄₂ alone induced a marked deficit in the learning ability of mice in the Morris Water Maze paradigm (for both acquisition and retention). This was accompanied by neuronal degeneration (assessed by Fluoro-Jade B staining) and an increased presence of CD11b-positive staining microglial cells within the dentate gyrus (DG) of the same animals (Figure 6C, D and E). Treatment of mice with 3,6′-dithiothalidomide, prior to Aβ₁₋₄₂, markedly mitigated the actions of this toxic peptide. Specifically, drug-treated animals performed at a level similar to control mice in the Morris Water Maze, and showed evidence of reduced levels of both...
neuronal degeneration and CD11b positive microglial cells in comparison to Aβ1-42 alone mice. No differences were evident between mice administered Aβ42-1 with or without drug treatment in any of the measured parameters.

3,6′-Dithiothalidomide, administered daily for 6 weeks, normalized age-associated biochemical, cellular and behavioral features of AD observed in the 3xTg-AD mouse model
Mice containing three transgenes (APPswe + PS1M146V + tauP301L) associated with AD [15] were utilized to assess the effects of 3,6′-dithiothalidomide on two distinct age groups of animals, of 10 and 17 months age at study onset, referred to here as adult and old, respectively. These two age ranges were selected based on prior studies [36,42] demonstrating that the line of 3xTg-AD mice utilized in our study, which had been backcrossed onto a C57BL/6 background for seven generations, presented with brain AD pathology at 16 months age. Our chosen age groups hence can be considered to be prior to and post the onset of evident significant AD pathology, and these 3xTg-AD mice were administered 3,6′-dithiothalidomide or vehicle for 6 weeks thereafter.

Following the final assessment of mouse learning and memory, animals were euthanized and various brain biochemical measures were undertaken. As assessed in the cerebral cortex, a trend to an elevated level in total APP (6E10) was evident when comparing adult vehicle with old vehicle animals (Figure 7A). 3,6′-Dithiothalidomide...
proved to be well tolerated by 3xTg-AD mice, and lowered total APP levels by 19% and 43% in adult and old mice compared to their respective vehicle controls (Figure 7A). Examination of soluble Aβ1–42 levels indicated a marked age effect (Figure 7B), with old vehicle controls expressing levels 143% of adult ones. This elevation was fully inhibited by 3,6′-dithiothalidomide, whose levels were similar to adult control and drug-treated mice (Figure 7B). Total tau protein levels were reduced (23%) in old vehicle animals compared to adult ones, and were unaffected by drug treatment (Figure 7C). However, levels of phosphorylated tau protein presented a strong age-associated rise (176% of adult controls) that, similarly to soluble Aβ1–42 protein, was attenuated by drug treatment in the older group (Figure 7D). Interestingly, the levels of two synaptic markers, SNAP25 and synaptophysin, which showed a trend to decline in the old vehicle control group (90% and 93% of respective adult vehicle levels) were found to be significantly elevated (19% and 37%, respectively) in old drug-treated mice compared to old vehicle ones (Figures 7E and F).

Immunohistochemical staining of insoluble Aβ plaques indicated marked plaque formation only in the old 3xTg-AD mice; this Aβ plaque staining was dramatically reduced by drug treatment (Figure 8A). Memory function, assessed by the Morris Water Maze, illustrated a deficit in learning and acquisition of the location of the

Figure 7 3,6′-Dithiothalidomide (42 mg/kg i.p. once daily × 6 weeks) attenuated age-associated changes in hallmark features of AD in old 3xTg-AD animals. Two cohorts of 3xTg-AD mice, adult and old, were evaluated after a 6-week regimen with daily administration of 3,6′-dithiothalidomide or vehicle. After completion of the Morris Water Maze assessment animals were euthanized, and the levels of hallmark features of Alzheimer’s disease were assessed. (A) Treatment with drug (n = 9) significantly reduced the levels of total APP in the old animals vs. old control animals (n = 7). (B) Cortical soluble Aβ1–42 levels were increased in the old + control animals compared to younger adult + control (n = 7–9, P < 0.01). Drug treatment reduced the soluble Aβ1–42 in the old mice (n = 8), but not in younger adult ones. (C) Levels of total tau protein were lower in old control animals compared to younger adult control mice; however, drug treatment had no effect on total tau levels in either age group. (D) Phosphorylated tau protein levels were elevated in old animals compared to younger adult animals (n = 8–9). Treatment of the old animals with drug reduced levels of phosphorylated tau protein. The levels of SNAP 25 (E) and synaptophysin (F) were elevated in old drug-treated mice (P < 0.01 and P < 0.05, both n = 7–9, respectively), compared to old vehicle animals. Data are presented in percent change from appropriate control terms and are expressed as mean ± SEM of n observations; levels of statistical significance are indicated as follows: *P < 0.05, **P < 0.01, ***P < 0.001.
hidden platform in old vehicle control mice during training (Figure 8B and C). Probe trial data obtained 4 h following their final training session indicated that these old vehicle control animals failed to remember the location of the platform, as illustrated by the low time spent within the target zone as well as by the low number of platform crossings (Figure 8C and D). 3,6'-Dithiothalidomide abolished this age-associated memory deficit (Figure 8B, C and D), with treated mice performing on par with adult vehicle control and drug-treated animals.

The levels of CD68-positive microglial cells within the subiculum and CA1 brain region were quantified as a marker of the inflammatory microenvironment in the
hippocampus, as these regions are among those showing the highest concentration of Aβ plaque staining (Figure 8A). CD68-positive microglial cells were significantly elevated in number (three-fold) only within old vehicle control mice, and this rise was fully abolished by the drug (Figure 8E). Hence, treatment with 3,6'-dithiothalidomide induced a marked normalization of key biochemical, learning and memory features of AD in old 3xTg-AD mice.

**Discussion**

Here we demonstrate that the TNF-α-lowering agent, 3,6'-dithiothalidomide, ameliorates key aspects of neuroinflammation in multiple acute and longer term CNS rodent models. Importantly, several of these models emulate specific cardinal characteristics of AD, and highlight the complex cyclic interaction among the synthesis of TNF-α, the development of neuroinflammation and impact on disease progression, inducing its advancement. Our data suggest that breaking this cycle by lowering TNF-α generation and neuroinflammation can favorably impact AD, as assessed at both a behavioral and biochemical level, even late during the disease course. Our studies hence reinforce the significant role of neuroinflammation in AD and other degenerative neurological disorders, and highlight the potential for targeting TNF-α.

TNF-α has been implicated in the pathogenesis of a wide number of neurological disorders, including AD, PD, stroke and head trauma [5-12]. Indeed, TNF-α levels have been found to be elevated within the CSF of AD patients by as much as 25-fold [43], in line with substantial elevations in TNF-α synthesis that were rapidly induced in RAW 264.7 cells and animals challenged with LPS (Figure 1A,E,F,G). Studies in subjects with mild cognitive impairment (MCI) that progress to develop AD suggest that increased CSF TNF-α levels are an early event, and their rise correlates with disease progression [44]. In accord with this, Janelisins and colleagues [23] noted an elevated expression of TNF-α transcripts within the entorhinal cortex of 3xTg-AD mice at 2 months, prior to the appearance of amyloid and tau pathology, and this increase correlated with the onset of cognitive deficits in these mice [45]. These studies, together with others demonstrating that (1) TNF-α polymorphisms that elevate TNF-α production may increase AD risk, particularly in patients carrying one or more apolipoprotein E e4 alleles [46-49], and that (2) genetic ablation of TNF-α receptor 1 (TNFR1) in APP23 AD mice [50] or a selective lowering of soluble TNF-α brain levels in 3xTg-AD mice [25] reduces AD progression, support the concept of TNF-α inhibition strategies for treatment of AD [10,51,52].

Protein-based TNF-α inhibitors (etanercept and infliximab) that can effectively regulate circulating TNF-α levels by binding them [53] have provided a means to initially target brain TNF-α in AD, and perispinal etanercept administration followed by Trendelenburg positioning in a small prospective open-label pilot study has been reported to provide a rapid onset of cognitive improvement [54]. TNF-α levels can also be regulated at the level of synthesis, which is tightly controlled at the level of mRNA stability to facilitate rapid responses to exogenous and endogenous stimuli [55], as occurs with LPS and Aβ challenge, respectively, and hence is amenable to regulation by small-molecular-weight drugs. The presence of adenylate-uridylate-rich elements (AREs) within the 3’-UTR of TNF-α mRNA supports the potential for post-transcriptional repression, targeting it for rapid degradation or inhibition of translation. This is mediated through interactions with RNA-binding proteins (RBPs), epitomized by HuR, which binds and stabilizes ARE-containing transcripts and conveys them to translational machinery to upregulate protein synthesis and, conversely, by tristetraprolin, which aids the acceleration and degradation of bound mRNAs [56-59].

Exogenous signals, as arise from exposure to bacterial proteins, potently induce inflammatory responses within the CNS in a manner mimicked by RAW 264.7 cells challenged with LPS [28]. The resulting elevation in TNF-α derives from an LPS-mediated increase in the half-life of TNF-α mRNA, allowing release of its translational repression. By contrast, translational blockade can be induced by small-molecular-weight compounds, such as thalidomide, which induce a shortening of the TNF-α mRNA half-life [55]. In this regard, 3,6’-dithiothalidomide is a more potent TNF-α-lowering thalidomide analogue that acts at the level of the 3’ UTR of TNF-α [27,60]. As is evident in Figure 1, LPS challenge to RAW 264.7 cells or animals activated toll-like receptors (TLR), and induced the generation of TNF-α and nitrite, a stable surrogate marker of highly unstable nitric oxide production. APP levels also were elevated, in line with prior studies [61] that additionally describe rises in interleukin (IL)-1, 6 and 12, and cyclooxygenase-2 [62]. 3,6’-Dithiothalidomide dose-dependently lowered LPS-induced TNF-α, nitrite and APP levels in the absence of cellular toxicity in RAW 264.7 cells, in contrast to thalidomide, which proved ineffective at concentrations up to 30 μM (not shown), but has been reported to lower APP levels in PC12 and SH-SY5Y neuronal cell lines [63]. This action of 3,6’-dithiothalidomide effectively translated to both a systemic LPS challenge in vivo, lowering systemic and central TNF-α expression (Figure 1E-G), as well as to a central LPS challenge.

Administered to brain, LPS reliably induces chronic neuroinflammation associated with the activation of
microglia, which is allied to impaired hippocampal-dependent spatial cognitive function [17,18,21]. In the present study this was achieved by the slow continuous infusion of a low dose of LPS into the fourth ventricle of the brain, which produced microglia activation within the hippocampus and, importantly, induced abnormal Arc expression in response to a simple behavioral task (Figures 2 and 3). The activity-regulated, cytoskeleton-associated IEG Arc is a key regulator of protein synthesis-dependent forms of synaptic plasticity, which are fundamental to memory formation [37,64,65]. In healthy brain, Arc protein functions in a transient manner, and its abnormally elevated sustained expression, as occurs during neuroinflammation, may generate synaptic noise and thereby impair long-term memory formation [64]. Co-administration of systemic 3,6′-dithiothalidomide, which has been reported to readily enter the brain (brain/plasma ratio 1.34) [30] and reversed an acute LPS-mediated increase in TNF-α expression (Figure 1G), fully inhibited LPS-induced activation of microglia and the resulting altered coupling of neural activity with de novo synthesis of Arc (Figures 4 and 3, respectively). A similar dose of 3,6′-dithiothalidomide has recently been described to normalize the expression of Arc and to restore the acquisition and consolidation of spatial memory impairments in a fully established model of neuroinflammation [29], in which LPS was administered to rodents for a full 28 days prior to drug treatment (rather than in parallel with drug treatment, as in the present study). In this study by Belarbi et al. [29], 3,6′-dithiothalidomide normalized LPS-induced elevations in brain TNF-α expression, but not IL-1β, and additionally normalized the expression of specific genes involved within the TLR-mediated signaling pathways (in particular, TLR2, TLR4, Hmgb1 and IRAK1) that are established to lead to elevated TNF-α expression [66].

Aβ, particularly in the form of soluble oligomeric assemblies or Aβ-derived diffusible ligands (ADDLs) [67-69], has been described to target synapses, induce neuronal dysfunction and impair cognition. Its administration into the lateral ventricle of mice has been widely used to model neuroinflammation and induce these Aβ-related impairments [32,70-72], which in the present study resulted in the activation of microglia and neuronal degeneration within the DG, accompanied by a learning impairment in the Morris Water Maze paradigm (Figure 6). Pretreatment of animals with 3,6′-dithiothalidomide markedly inhibited each of these aspects and, together with our prior studies, suggested that the agent could prove of value in Tg models of AD that, like the human condition, increasingly develop neuroinflammation during disease progression [23,24]. This hypothesis was tested in two cohorts of 3xTg-AD mice of 10 and 17 months age, chosen to represent times that in our specific line coincided with the pre- and post-development of amyloid plaques and neurofibrillary tangles, as the presence of activated microglia in close proximity to amyloid plaques is a cardinal feature of AD-affected brain [73].

The pre-pathological upregulation of TNF-α and associated enhancement of activated microglia have been reported in the 3xTg-AD mouse model [23,25], and it has been postulated that these activated immune cells are key in the process of clearing extracellular Aβ [62]. A potential consequence of heightened Aβ exposure, however, is microglia TLR4 stimulation and a resultant upregulation of cytokine production and release [74]. TNF-α as well as IL-1β can correspondingly elevate Aβ generation by stimulating γ-secretase activity [24,75], potentially spawning a self-propagating positive feedback loop of Aβ induction of inflammation and TNF-α signaling that, in turn, may provoke further Aβ generation [5,6,9,10,62]. In our study, in accord with the literature [24], activated microglia were markedly elevated in old versus adult vehicle-treated 3xTg-AD mice (Figure 8E), which additionally presented with a significant elevation in brain Aβ1–42 and phosphorylated tau levels, a decline in total tau and a trend towards elevation of APP levels (Figure 7). A substantial accumulation of extracellular amyloid plaques was clearly evident within the cerebral cortex and hippocampus of old versus adult 3xTg-AD mice, which was accompanied by deficits in learning and memory, as assessed within the Morris Water Maze paradigm (Figure 8). The administration of 3,6′-dithiothalidomide to old 3xTg-AD mice reversed each of these parameters, significantly reducing Aβ1–42, phosphorylated tau and APP levels, lowering levels of activated microglia and fully ameliorating memory deficits (Figures 7 and 8), which were accompanied by an elevation in synaptic protein markers (Figures 7E and F). These drug-induced changes are in line with studies by McAlpine and colleagues [25], demonstrating that blockade of TNF-α signaling (either by viral vector-mediated expression of TNFR constructs or by crossing 3xTg-AD mice with TNFR1 knockout mice) significantly suppressed AD pathology. Importantly, our studies additionally demonstrate that cognitive deficits that accompany the classical pathology of AD appear to be reversible, at least in the 3xTg-AD mouse model.

A caveat with this 3xTg-AD mouse model, like all such models, is that it provides a partial model of the human disease. APP and tau expressions (specifically, human APPswg and human tauP301L) are driven in the 3xTg-AD model by the unnatural mouse Thy1.2 regulatory element [22]. Hence, the possibility that some actions of 3,6′-dithiothalidomide may be mediated via suppression of this unnatural transgene promoter cannot be ruled out. Importantly, however, the action of 3,6′-dithiothalidomide to favorably lower
APP levels as well as neuroinflammation in cellular studies (Figure 1) occurred in cells controlled by their natural endogenous regulatory elements, and wt rodents were used in all other studies.

TNF-α has been shown to regulate numerous cellular processes, not only inflammation and cell death, but also cellular differentiation and survival, and achieves this by binding and activating two cognate receptors, TNFR1 (p55) and TNFR2 (p75) [76]. TNFR1, expressed ubiquitously including on neurons, astrocytes and microglia, possesses an intracellular death domain and contributes to neuronal dysfunction and death following activation by soluble TNF-α [77], whereas TNFR2, principally expressed on cells of hematopoietic origin but also on neurons, has been associated with cell survival [76,78-80] and chiefly responds to membrane-bound TNF-α [81,82]. The engagement of homotrimeric TNF-α to either receptor can activate three major signaling pathways: an apoptotic cascade initiated via the TNF-α receptor-associated death domain, a nuclear factor kappa B (NFkB) signaling pro-survival pathway implemented via NFkB-mediated gene transcriptional actions, and a JNK (c-Jun N-terminal kinase) cascade involved in cellular differentiation and proliferation that is generally pro-apoptotic [62]. In large part, although the contrasting pro-survival versus death-inducing actions of TNF-α plausibly rely on the TNF-α receptor subtype activation, the target cell types involved and their expression ratio of TNFR1/2 and associated coupling proteins, the temporal levels of available soluble and membrane-bound TNF-α [79], and the scale and duration of neuroinflammation combine in determining the eventual physiological consequences of TNF-α receptor activation [5,6,9,10,62]. Consequent to the diverse actions of TNF-α and the influence of the brain microenvironment in which they occur, it is hence not always clear under which conditions TNF-α promotes beneficial versus deleterious neuronal actions, and this, in large part, accounts for how an initially pro-survival response may develop into a pro-apoptotic one.

Under appropriate conditions TNF-α signaling, primarily via TNFR2, can mediate homeostatic actions, epitomized by its role in AMPA receptor surface expression and synaptic scaling to impact LTP [83], as well as neuroprotective ones [9,10]. The genetic ablation of TNFR1 and -R2 in 3xTg-AD mice has been described to increase the progression of AD pathology [84]. Furthermore, TNF-α has a reported role in hippocampal development and function [85] and, with the expression of both TNFR1 and -R2 on neuronal progenitor cells, it can modulate neurogenesis within the hippocampal neurons under pathological conditions [86-89]. The finding that adult 3xTg-AD mice were not detrimentally impacted by 3,6′-dithiothalidomide suggests that such homeostatic actions of TNF-α signaling were largely unimpaired, although clearly substantial classical preclinical toxicological studies are warranted before the agent can be considered for clinical use.

Conclusion
In synopsis, the present data demonstrate the key role of TNF-α in acute and chronic neuroinflammatory models that have relevance to neurodegeneration and, in particular, to AD and its progression. The TNF-α synthesis inhibitor 3,6′-dithiothalidomide, administered at doses that compare favorably to those of thalidomide in human studies (100 to 1,200 mg daily [90]), appears to advantageously reset the delicate balance between the pro- versus anti-apoptotic actions of this signaling cascade in the brain. This resulted in an inhibition of neuroinflammation and reduced AD progression in 3xTg-AD mice, and thereby supports the feasibility of targeting TNF-α as a potential treatment strategy for AD and other neurological disorders involving a neuroinflammatory component.

Abbreviations
AREs: adenylyl-uridylyl-rich elements; AD: Alzheimer’s disease; ADDLs: amyloid-β-derived diffusible ligands; AB: amyloid-β peptide; αCSF: artificial cerebrospinal fluid; CSF: cerebral spinal fluid; CNS: central nervous system; DG: dentate gyrus; DMSO: dimethyl sulfoxide; GFAP: glial fibrillary acidic protein; IEG: immediate early gene; IL: interleukin; IP: interperitoneal; IV: intracerebroventricularly; JNK: c-Jun N-terminal kinase; LPS: lipopolysaccharide; LTP: long-term potentiation; MCI: mild cognitive impairment; MW: molecular weight; NFkB: nuclear factor kappa B; PD: Parkinson’s disease; RBPs: RNA-binding proteins; sAPP: secreted amyloid precursor protein; TNF-α: tumor necrosis factor-alpha; TNFR1: TNF-α receptor 1; TNFR2: TNF-α receptor 2; TLR: Toll-like receptors; WT: wild type.

Competing interests
NHG and HWH declare that they are co-inventors of the original 3,6′-dithiothalidomide patent. Having assigned all their rights to the US government, they declare that they have no ownership, financial interest or any other competing interests. All other authors declare no competing interests.

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Authors’ contributions

DT contributed to the design of the study, and undertook experimental studies and biochemical analyses in Figures 1, 7 and 8, and the associated statistical analyses. RAF and KF contributed to the experimental studies and analyses in Figures 2, 3, 4 and 5. KAF, HVP, HWH and YL contributed to the experimental design, and experimental studies and analyses in Figures 1, 7 and 8. WL undertook synthetic chemistry, chemical characterization and stability assessments. LC, IR, SB and FB contributed to the experimental design and undertook experimental studies and analyses in Figure 6. BR and DKL undertook tissue preparations, Western blot and ELISA analyses in Figure 7. FB, DKL, SR and NHG contributed to the study conception and design, and experimental studies, and wrote the manuscript. All the authors read and approved the final manuscript.

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