

RESEARCH

Open Access



Biologic TNF- α inhibitors reduce microgliosis, neuronal loss, and tau phosphorylation in a transgenic mouse model of tauopathy

Weijun Ou¹, Joshua Yang², Juste Simanauskaite³, Matthew Choi⁴, Demi M. Castellanos², Rudy Chang¹, Jiahong Sun¹, Nataraj Jagadeesan¹, Karen D. Parfitt³, David H. Cribbs⁵ and Rachita K. Sumbria^{1,6*} 

Abstract

Background: Tumor necrosis factor- α (TNF- α) plays a central role in Alzheimer's disease (AD) pathology, making biologic TNF- α inhibitors (TNFIs), including etanercept, viable therapeutics for AD. The protective effects of biologic TNFIs on AD hallmark pathology (A β deposition and tau pathology) have been demonstrated. However, the effects of biologic TNFIs on A β -independent tau pathology have not been reported. Existing biologic TNFIs do not cross the blood–brain barrier (BBB), therefore we engineered a BBB-penetrating biologic TNFI by fusing the extracellular domain of the type-II human TNF- α receptor (TNFR) to a transferrin receptor antibody (TfRMAB) that ferries the TNFR into the brain via receptor-mediated transcytosis. The present study aimed to investigate the effects of TfRMAB-TNFR (BBB-penetrating TNFI) and etanercept (non-BBB-penetrating TNFI) in the PS19 transgenic mouse model of tauopathy.

Methods: Six-month-old male and female PS19 mice were injected intraperitoneally with saline ($n = 12$), TfRMAB-TNFR (1.75 mg/kg, $n = 10$) or etanercept (0.875 mg/kg, equimolar dose of TNFR, $n = 10$) 3 days/week for 8 weeks. Age-matched littermate wild-type mice served as additional controls. Blood was collected at baseline and 8 weeks for a complete blood count. Locomotion hyperactivity was assessed by the open-field paradigm. Brains were examined for phosphorylated tau lesions (Ser202, Thr205), microgliosis, and neuronal health. The plasma pharmacokinetics were evaluated following a single intraperitoneal injection of 0.875 mg/kg etanercept or 1.75 mg/kg TfRMAB-TNFR or 1.75 mg/kg chronic TfRMAB-TNFR dosing for 4 weeks.

Results: Etanercept significantly reduced phosphorylated tau and microgliosis in the PS19 mouse brains of both sexes, while TfRMAB-TNFR significantly reduced these parameters in the female PS19 mice. Both TfRMAB-TNFR and etanercept treatment improved neuronal health by significantly increasing PSD95 expression and attenuating hippocampal neuron loss in the PS19 mice. The locomotion hyperactivity in the male PS19 mice was suppressed by chronic etanercept treatment. Equimolar dosing resulted in eightfold lower plasma exposure of the TfRMAB-TNFR compared with etanercept. The hematological profiles remained largely stable following chronic biologic TNFI dosing except for a significant increase in platelets with etanercept.

Conclusion: Both TfRMAB-TNFR (BBB-penetrating) and non-BBB-penetrating (etanercept) biologic TNFIs showed therapeutic effects in the PS19 mouse model of tauopathy.

*Correspondence: sumbria@chapman.edu

¹ Department of Biomedical and Pharmaceutical Sciences, School of Pharmacy, Chapman University, Irvine, CA 92618, USA
Full list of author information is available at the end of the article



© The Author(s) 2021. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by/4.0/>. The Creative Commons Public Domain Dedication waiver (<http://creativecommons.org/publicdomain/zero/1.0/>) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

Keywords: Biologic TNF- α inhibitor, Alzheimer's disease, Tau, Microgliosis, Transferrin receptor, Molecular Trojan Horse, Blood–brain barrier

Introduction

Alzheimer's disease (AD) is a progressive neurodegenerative disease that was first discovered in 1906 by the German clinical psychiatrist and neuroanatomist, Alois Alzheimer [1]. The neuropathological alterations reported by Alzheimer in 1906 constitute the main pathological hallmarks of AD today: extracellular deposition of amyloid-beta (A β)-containing plaques and intracellular tau-containing neurofibrillary tangles in the brain [2]. While the accumulation of A β plaques in the brain is a characteristic of AD pathology, abnormal intracellular accumulation of tau lesions is observed in several neurodegenerative disorders, known as tauopathies [3].

Apart from the predominant role of A β and tau pathology in AD, neuroinflammation has emerged as a driver and an accelerator of AD pathology [4–6], and therefore provides a broad window of opportunity (from early to late-stage AD) for therapeutic intervention. Among the pro-inflammatory mediators, TNF- α is one of the main inflammatory cytokines involved in initiating and propagating an inflammatory response [7]. Both clinical and preclinical studies support the role of peripheral and central nervous system (CNS) TNF- α in AD, and elevated TNF- α levels were observed in the serum [8, 9] and post-mortem brains of AD patients and AD mouse models [10, 11]. Furthermore, serum TNF- α levels correlated with disease progression in AD patients [9]. Global TNF- α receptor 1 (TNFR-1) knock-out prevented A β generation and learning deficits in the APP23 AD mice [12]. Direct application of TNF- α to cell culture preparations, or CNS administration of TNF- α , increased A β production, tau phosphorylation, and synaptic dysfunction [13–17], and accelerated disease progression and cognitive decline [18]. Similarly, peripheral TNF- α modulated AD pathology in mouse models [19, 20]. Adding to this, TNF- α polymorphisms linked with increased TNF- α production were found to be associated with late-onset AD [21].

Several studies report the effects of the existing FDA-approved biologic TNF- α inhibitors on AD pathology both in humans and preclinical models. For example, infliximab, a chimeric anti-TNF- α antibody, significantly reduced cognitive deficits in a female AD patient and significantly reduced A β - and tau-pathology and memory impairment in AD mouse models, following intrathecal and intracerebroventricular administration [22–24], respectively. Similarly, perispinal etanercept, a fusion protein of the extracellular domain of the type-II TNFR and human IgG1 Fc-domain, improved cognitive deficits

in AD patients [25]. Accordingly, intrahippocampal injection of dominant-negative TNF- α inhibitors reduced A β pathology in the 3xTg mouse model [26]. Recently, intracerebroventricular injection of a nanobody directed against the TNFR-1 was protective in two mouse models of amyloidosis [27]. One commonality among the aforementioned studies is the invasive route of injection used to evaluate the CNS effects of these biologic TNF- α inhibitors due to their limited blood–brain barrier (BBB) penetration. For example, the brain uptake of etanercept was found to be equal to that of a human IgG control [28].

Tau protein belongs to the family of microtubule-associated proteins, has a high binding affinity for microtubules, and plays an important role in microtubule stabilization for optimal neuronal function [29]. Tauopathies, including AD, are characterized by the aggregation of hyperphosphorylated tau protein, which is strongly correlated with neuroinflammation [30, 31]. Recent studies have shown that microglial activation and release of pro-inflammatory cytokines exacerbate tau pathology, independent of A β pathology [31, 32], and that anti-inflammatory strategies can reduce tau pathology in mice [33, 34]. Despite a large body of literature looking at the effects of TNF- α inhibition on AD pathology, the effects of TNF- α inhibition on A β -independent tau pathology have not been reported.

To study the effects of direct CNS TNF- α blockade via a non-invasive injection route, we engineered a BBB-penetrating biologic TNF- α inhibitor by fusing the extracellular domain of the type-II TNFR (decoy TNFR) to a transferrin receptor antibody (TfRMAB) for BBB penetration [35, 36]. The TfRMAB-TNFR fusion protein binds to the BBB TfR and enters the brain via receptor-mediated transcytosis. The resulting brain concentrations of the TfRMAB-TNFR are >40-fold higher than that of a non-TfR targeting IgG [36]. Our previous work showed that this BBB-penetrating TNF- α inhibitor reduces A β plaques, insoluble A β 1-42, BBB-disruption, brain endothelial activation, and cognitive memory deficits in the APP/PS1 double transgenic AD mouse model of amyloidosis [37]. Overall, the BBB-penetrating TNF- α inhibitor had better therapeutic indices than etanercept, a non-BBB-penetrating TNF- α inhibitor [28] expected to modulate only peripheral but not CNS TNF- α , in the mouse model of amyloidosis [37]. However, the effects of these biologic TNF- α inhibitors on tau pathology are currently unknown.

Based on the above, the goal of the current investigation was to examine the effects of peripheral and CNS TNF- α inhibition on tau pathology using the P301S mutant tau transgenic mouse model of tauopathy (line PS19). For this, we first evaluated the effect of the biologic TNF- α inhibitors (etanercept: non-BBB-penetrating and TfrMab-TNFR: BBB-penetrating) on tau phosphorylation, microgliosis, and neuronal health. We further assessed the effect of the biologic TNF- α inhibitors on the hyperactive phenotype of the PS19 mice using the open-field test. To gain insight into the differential therapeutic effects of the two biologic inhibitors, we examined the plasma pharmacokinetics of these biologics in mice. Finally, the overall hematologic safety of chronic biologic TNF- α inhibitor dosing was evaluated by performing a complete blood count in the PS19 mice.

Methods

Fusion protein

TfrMab-TNFR was produced via transient expression in CHO-K1 cells, and cell supernatants were collected for protein A purification followed by size exclusion chromatography (WuXi Biologics, NJ, USA), as described previously [36]. The final protein purity was >96%, and the target protein was verified by Western blot. Etanercept was obtained from International Laboratory USA (CA, USA). The affinity of the TfrMab-TNFR fusion protein to the mouse Tfr and human TNF- α was confirmed through ELISA (Additional file 1: Supplemental Methods, Additional file 2: Fig. S1A, B). TNF- α inhibitory effects were confirmed in vitro using induced pluripotent stem cell (iPSC)-derived human brain microvascular endothelial cells (ihBMECs; Additional file 1: Supplemental Methods, Additional file 2: Fig. S1C, D). Both the TfrMab-TNFR fusion protein and etanercept were formulated in 100 mM glycine, 150 mM NaCl, 28 mM Tris, and 0.01% Polysorbate 80, pH=6.42, sterile filtered, and stored at -80°C until use.

Chronic dosing in a mouse model of tauopathy

Animal studies were performed on hemizygous Tg(P301S)PS19Vle (PS19) mice (6 months old at the start of the study, Jackson Laboratories, ME, USA) in compliance with University Laboratory Animal Resources under protocols approved by the University of California, Irvine, Institutional Animal Care and Use Committee. Mice were provided constant access to food and water and were maintained under a 12 h light/12 h dark cycle. Mice were injected intraperitoneally (IP) three days per week for 8 weeks with saline (PS19-Saline; $n=12$, female=7, male=5), TfrMab-TNFR (PS19-TfrMab-TNFR; 1.75 mg/kg, $n=10$, female=5, male=5), or etanercept (PS19-Etanercept; 0.875 mg/

kg, $n=10$, female=5, male=5). These are equimolar doses of TfrMab-TNFR and etanercept since the TNFR domain comprises ~50% of etanercept and ~26% of the TfrMab-TNFR fusion protein based on amino acid sequence [36, 38]. Age-matched noncarrier wild-type (WT) littermates ($n=12$, female=6, male=6) received equivalent volume of saline via the IP route (Fig. 1A). All the mice were evaluated for signs of immune response (general appearance, spontaneous locomotion, and posture) after each injection, and body weights were checked weekly. Blood was collected at baseline and 8 weeks after treatment for a complete blood count (Molecular Diagnostic Services, Inc., CA, USA). At the end of 8 weeks, mice were subjected to the open-field paradigm that was performed over a week, anesthetized with a lethal dose of Euthazol (150 mg/kg, IP), and perfused with ice-cold phosphate buffer saline (PBS). Brains were harvested, and hemi-brains were fixed in 4% paraformaldehyde (PFA) for immunostaining or were frozen in dry ice for Western blotting (Fig. 1A).

Open-field testing

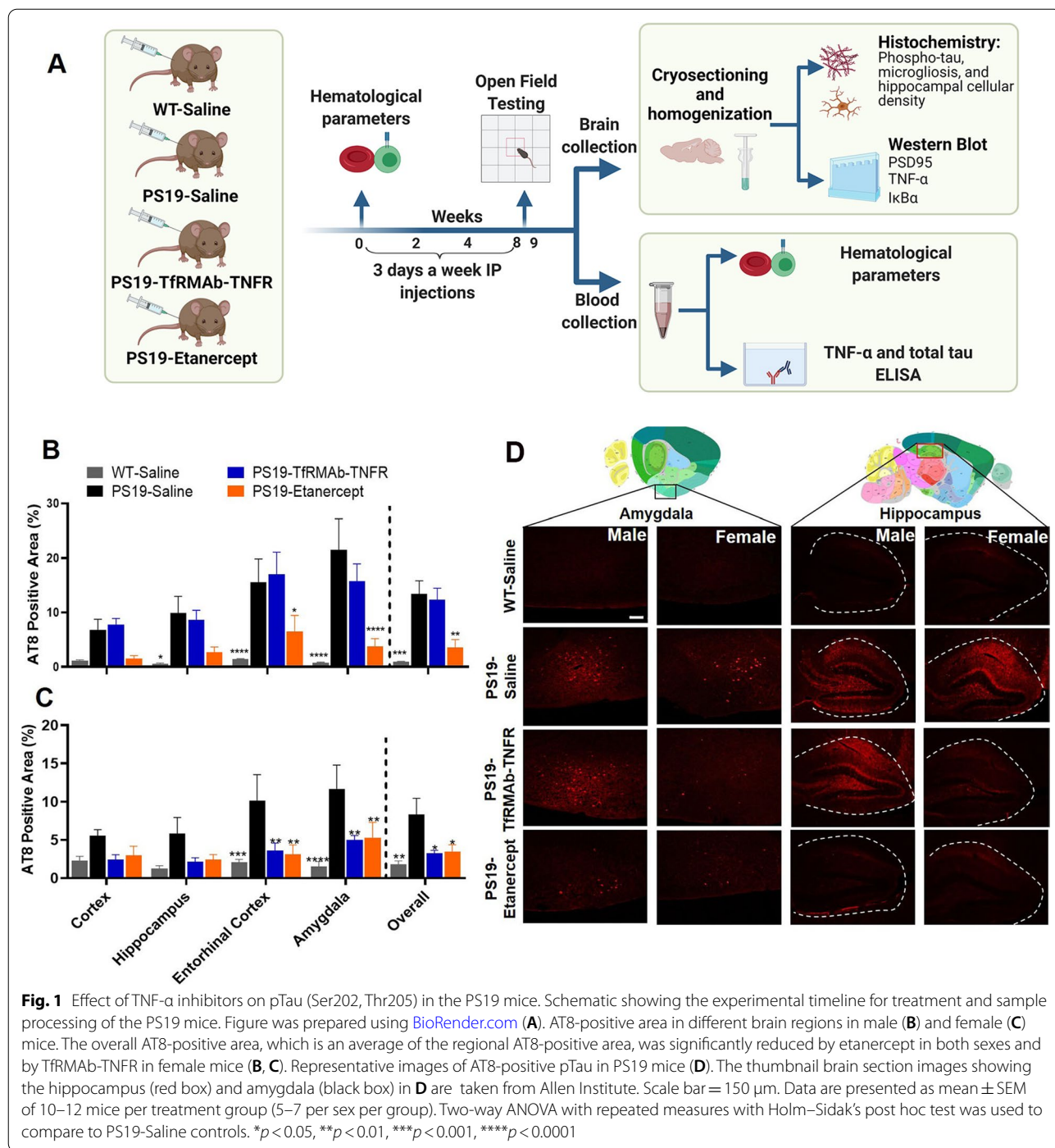
Since the PS19 mice consistently demonstrate a hyperactivity phenotype compared to cognitive deficits [39–41], we used the open-field test to measure locomotion hyperactivity. The open-field test was performed after 8 weeks of treatment (during week 9) as described previously [42–44]. Briefly, mouse movements were recorded for 5 min after placing in a white open box (72 cm x 72 cm with 36 cm walls). Mean speed and total distance were measured for locomotion evaluation. A center square, measuring 36 cm x 36 cm was drawn in the open box to measure time in the center. Time in the center was used as a measure of anxiety in mice. All analysis was performed using the SMART 3.0 Video Tracking Software (Panlab, Harvard Apparatus, MA, USA).

Cryosectioning

The right-cerebral hemi-brain of each mouse was immersion-fixed with 4% PFA in PBS for 24 h, followed by serial incubation in 10%, 20%, and 30% sucrose solution at 4°C for 24 h each, which was followed by freezing. The frozen brain tissues were mounted in Tissue-Tek OCT compound (Fisher Scientific, MA, USA), and sliced into 20- μm -thick sagittal sections at -25°C using a cryostat (Micron Instruments, CA, USA). Five sections (600 μm apart) per mouse were used for immunostaining and histology as described below.

Phospho-tau (pTau; Ser202, Thr205) and Iba-1 immunostaining

Free-floating sagittal brain sections were washed three times for 2 min in PBS and blocked with 0.5% bovine



serum albumin (BSA) in PBS containing 0.3% Triton X-100 (TX100) for 60 min at room temperature. Adjacent tissue sections were incubated with either 0.2% biotin-conjugated pTau monoclonal antibody (AT8 antibody to detect tau phosphorylated at ser202, thr205; Thermo Fisher Scientific, MA, USA) or 0.5 μ g/mL anti-Iba-1 rabbit antibody (to detect microgliosis)

in PBS containing 0.3% TX100 and 0.5% BSA overnight at 4 $^{\circ}$ C. The tissue sections were then washed three times in PBS for 5 min and incubated in the dark with 0.5% AlexaFluor 594 conjugated streptavidin antibody (Biolegend; CA, USA) for pTau detection or with 0.1% AlexaFluor 488 donkey anti-rabbit IgG (Biolegend; CA, USA) in PBS containing 0.3% TX-100 and 0.5%

BSA for 2 h at room temperature. The sections were then washed three times in PBS for 10 min and quickly dipped in distilled water to remove the salts. The sections were mounted onto glass slides, cover-slipped with Vectamount aqueous mounting media (Vector Laboratories, CA, USA), and sealed with nail polish. Slides were stored at 4 °C until imaging.

pTau (Ser202, Thr205) and Iba-1 quantification

The fluorescent staining was analyzed using a Leica TCS SP5 Confocal Microscope (Leica, NJ, USA). For immunostaining, five distinct brain sections (600 μm apart) for each mouse were used for analysis. For each brain section, two regions in the cerebral cortex, one or two regions in the hippocampus, one region in the entorhinal cortex, and one region in the amygdala were examined and imaged at a 10 \times magnification. For Iba-1 immunostaining, for each brain section, three regions in the cortex, two regions in the hippocampus, two regions in the entorhinal cortex, and one region in the amygdala was imaged at a 10 \times magnification with 3 \times digital zoom. The digitized images were analyzed using the NIH ImageJ (version 1.53e, MD, USA) using a threshold setting to calculate tissue area positive for the AT8 or Iba-1. For this, immunofluorescent images were converted to 16-bit and inverted to grayscale. Threshold values of images were manually calibrated to eliminate background noise. After adjusting the threshold, the “analyze” function was used to report the tissue area positive for AT8 or Iba-1. The AT8 or Iba-1-positive area was expressed as a percentage of the brain tissue analyzed area. The stain positive area of each brain region (cortex, hippocampus, entorhinal cortex, and amygdala) was averaged for each mouse to give the ‘overall’ stain positive area. For microglia number quantification, each Iba-1-stained image was manually read to count the number of microglia per image using NIH ImageJ. For each brain region, the microglia counts were summed to give the total number of microglia per region per mouse. The overall number of microglia represents the sum of the microglia in each brain region per mouse. Additionally, microglial soma size in pixel units was determined in Iba-1-stained images of each region using NIH ImageJ by converting images to 8-bit, inverting to grayscale, and using the despeckle function to highlight the soma in the images. A threshold setting of >10-pixel units was used to eliminate background signal detection. The sum of the microglia from all the brain regions was determined for each mouse to calculate the total number of microglial <50- and >50-pixel units per mouse. All the NIH ImageJ analysis was

performed by two readers blinded to the experimental groups.

Hippocampal neuronal loss quantification

Hippocampal neuronal loss was determined in six mice per treatment group (3 mice per sex). Three 20 μm sagittal mouse brain sections (600 μm apart and adjacent to the sections selected for AT8 immunostaining) per mouse were mounted on positive glass slides and allowed to dry overnight. Hematoxylin and Eosin (H&E) staining was performed by consecutive dips in acetal staining jars (Simport, Quebec, Canada) as follows: 30 s in distilled water, 10 min in filtered Mayer’s Hematoxylin (Fisher Scientific, MA, USA), 30 s in tap water, 30 s in Scott’s tap water/bluing reagent (10 g anhydrous MgSO_4 , 2 g NaHCO_3 , 1L tap water), 30 s in tap water, 4 min in Eosin Y (0.5% aqueous solution; Sigma Aldrich, MO, USA), 1 min in 95% ethanol, followed by 1 min in 100% ethanol. The slides received a final dip in xylene (Sigma Aldrich, MO, USA). Slides were cover-slipped with Permount mounting media (Fisher Scientific, MA, USA) and imaged using a light microscope (Motic, British Columbia, Canada). Different hippocampal regions (Dentate Gyrus (DG), CA1, CA2, and CA3) were imaged at a 10 \times magnification, and the area occupied by the H&E-stained hippocampal neurons (μm^2) of the granule cell layer in the DG and pyramidal layer in the CA1-3 regions was manually outlined and quantified using NIH ImageJ (version 1.53e, MD, USA) by a reader blinded to the experimental groups. The total hippocampal neuronal cell area per section was obtained by adding the regional cell areas.

Western blot

Protein was extracted from the left frozen hemi-brains using the radio-immunoprecipitation assay (RIPA) buffer with Pierce Protease Inhibitor (Thermo Fisher Scientific, MA, USA) for Western blot as previously described [44–46]. Briefly, pulverized brains were homogenized in 2 volumes of RIPA (25 mM Tris-HCl, pH 7.6, 150 mM NaCl, 1% NP-40, 1% sodium deoxycholate, 0.1% sodium dodecyl sulfate (SDS) with 5 mM EDTA) and centrifuged at 20,800 \times g for 20 min at 4 °C. The supernatants were collected and processed with 4 \times Laemmli buffer (Bio-Rad, CA, USA) with 10% beta-mercaptoethanol followed by boiling for 10 min. Protein samples (30–50 μg) were separated on 4–20% SDS-ready precast gels (Bio-Rad, CA, USA) and transferred to polyvinylidene fluoride membranes (Bio-Rad, CA, USA). Membranes were sequentially subjected to blocking in Tris-buffered saline (TBS) containing 5% non-fat milk for 1 h at room temperature and probed with the mouse anti-PSD95 primary antibody (a marker of synaptic/neuronal health) (1:1000

in 3% non-fat milk, Santa Cruz Biotechnology, TX, USA) overnight at 4 °C followed by washing with TBS with 0.1% Tween-20 (TBST). Membranes were probed with an anti-mouse IgG kappa, HRP-linked secondary antibody (1:1000 in TBS containing 3% non-fat milk, Santa Cruz Biotechnology, TX, USA) for 1 h at room temperature followed by washing with TBST. Enhanced chemiluminescent (ECL) substrate reagent (Thermo Fisher Scientific, MA, USA) was used for secondary antibody detection and chemiluminescence was imaged using the Azure C500 gel imager (Azure Biosystems, CA, USA). Membranes were probed with an anti- β -actin antibody (1:1000 in TBS containing 3% non-fat milk, Santa Cruz Biotechnology, TX, USA) as a loading control, and membranes were incubated in the secondary antibody and ECL substrate as described above. NIH ImageJ (version 1.53e, MD, USA) was used to quantify the intensity of the Western blot bands and all the values were normalized to PS19-Saline mice.

Pharmacokinetic (PK) study

All animal procedures were approved by the University of La Verne and Pomona College Institutional Animal Care and Use Committee. Eight-week-old male C57BL/6J mice (24–27 g) were obtained from the Jackson Laboratories (ME, USA). Mice had constant access to food and water and were maintained on a 12 h light/12 h dark cycle, and mice were randomly assigned to treatment groups. For the single-dose PK study, TfrMab-TNFR fusion protein (1.75 mg/kg) or etanercept (0.875 mg/kg) were injected via the IP route in the volume of 130–140 μ L per mouse. Each treatment group comprised 4–6 mice. Blood was collected in sodium citrate (9 parts blood and 1-part sodium citrate) at 3, 6, 20, and 24 h following IP administration, and plasma was collected.

For chronic dosing PK study, male heterozygous APP/PS1 mutant mice (strain B6C3-Tg APP^{swe}, PSEN1^{dE9}, 85Dbo/Mmjax, stock 004462, Jackson Laboratories, ME, USA) (~19 months of age) were injected IP three days per week for four weeks with either 1.75 mg/kg TfrMab-TNFR ($n=4$) or an equivalent volume of saline ($n=6$). One week after 4-week dosing, all mice that were chronically treated with saline or TfrMab-TNFR received a final single 1.75 mg/kg dose of the TfrMab-TNFR fusion protein via the IP route to determine the impact of chronic TfrMab-TNFR dosing on plasma PK. Blood was collected at 3, 6, and 24 h after the final injection for plasma collection. For the single and chronic PK studies, plasma TfrMab-TNFR or etanercept concentrations were determined using a TNFR sandwich ELISA described below. Plasma concentration–time profiles were used to perform non-compartmental analysis to determine PK parameters

(maximum plasma concentration (C_{max}) and area under the plasma concentration–time curve (AUC)) using Thermo Scientific Kinetica 5.0 (Thermo Fisher Scientific, MA, USA), as described previously [47–49].

TNFR ELISA

TfrMab-TNFR fusion protein binding to TNF- α (Additional file 2: Fig. S1B), and TfrMab-TNFR and etanercept plasma concentrations were quantified by a sandwich ELISA. The human TNF- α (hTNF- α) (Pepro-Tech, Rocky Hill, NJ, USA) was the capture agent and 2 μ g/mL of hTNF- α was plated in 96-well plates, and incubated overnight at 4 °C. The wells were blocked with TBS containing 1% bovine serum albumin (TBSB) (0.01 M Tris/0.15 M NaCl/1% BSA/pH7.4) for 30 min at room temperature. Plasma samples (diluted 1:10 in TBSB) were added to the wells and incubated for 1 h at room temperature followed by washing with TBST. Alkaline phosphatase-conjugated detector agents, goat anti-human IgG-Fc fragment antibody (Bethyl, TX, USA) that binds to the human Fc domain of etanercept and goat anti-mouse light chain (kappa) antibody (Bethyl Laboratories, Inc., TX, USA) that binds to the TfrMab domain of the TfrMab-TNFR, were added to the wells for 1 h at room temperature followed by washing with TBST. Wells were then incubated with P-nitrophenyl phosphate solution (Sigma Aldrich, St. Louis, MO, USA) for 15 min in the dark at room temperature and the reaction was stopped by adding 1.2 M NaOH. Blank corrected absorbance measured at 405 nm was used to calculate plasma concentrations using a standard curve.

Statistical analysis

All numerical variables are represented as mean \pm SEM, and all statistical analysis was performed using GraphPad Prism 8 (GraphPad Software Inc., CA, USA) and as described by others [50]. Outliers were identified using the Grubb's test. Independent two-sample t -test or one-way ANOVA with Holm–Sidak's post hoc test was used to compare the means of two or more than two groups, respectively. To test the effects of two factors (e.g., treatment groups and time, or treatment groups and brain region), two-way repeated-measures ANOVA with Holm–Sidak's multiple comparisons test was used. Open-field data were analyzed using two-way ANOVA with Holm–Sidak's multiple comparisons test. Correlations between two numerical variables were analyzed by the Pearson correlation. A two-tailed $p < 0.05$ was considered statistically significant.

Results

Effect of biologic TNF-α inhibitors on pTau (Ser202, Thr205) in the PS19 mice

To determine the impact of TNF-α inhibitors on tau pathology, we quantified the pTau-positive area in the cortex, hippocampus, entorhinal cortex, and amygdala using the AT8 antibody. We then determined the overall AT8-positive area (average of the regional AT8-positive area). As expected, the overall AT8-positive area was significantly lower in the WT mice compared to the PS19-Saline controls, in both male (93% lower; $p < 0.001$) and female (78% lower; $p < 0.01$) mice (Fig. 1B–D; Additional file 3: Fig. S2A). Etanercept significantly reduced the overall AT8-positive area in both male and female PS19 mice by 73% ($p < 0.01$) and 58% ($p < 0.05$), respectively (Fig. 1B–D; Additional file 3: Fig. S2A). While TfRMAb-TNFR resulted in a 61% decrease

($p < 0.05$) in the overall AT8-positive area in the female PS19 mice (Fig. 1C–D; Additional file 3: Fig. S2A), surprisingly there was no change in the AT8-positive area in the TfRMAb-TNFR-treated male PS19 mice compared to the PS19-Saline mice (Fig. 1B, D; Additional file 3: Fig. S2A).

Effect of the biologic TNF-α inhibitors on microgliosis in the PS19 mice

Microgliosis was assessed in the cortex, hippocampus, entorhinal cortex, and amygdala by measuring the brain tissue area positive for the microglial marker, Iba-1. The overall Iba-1-positive area, which is the average of the regional Iba-1-positive areas, was significantly lower in the WT mice compared with the PS19-Saline controls in both male (35% lower; $p < 0.01$) and female (18% lower; $p < 0.05$) mice (Fig. 2A–C; Additional file 4: Fig.

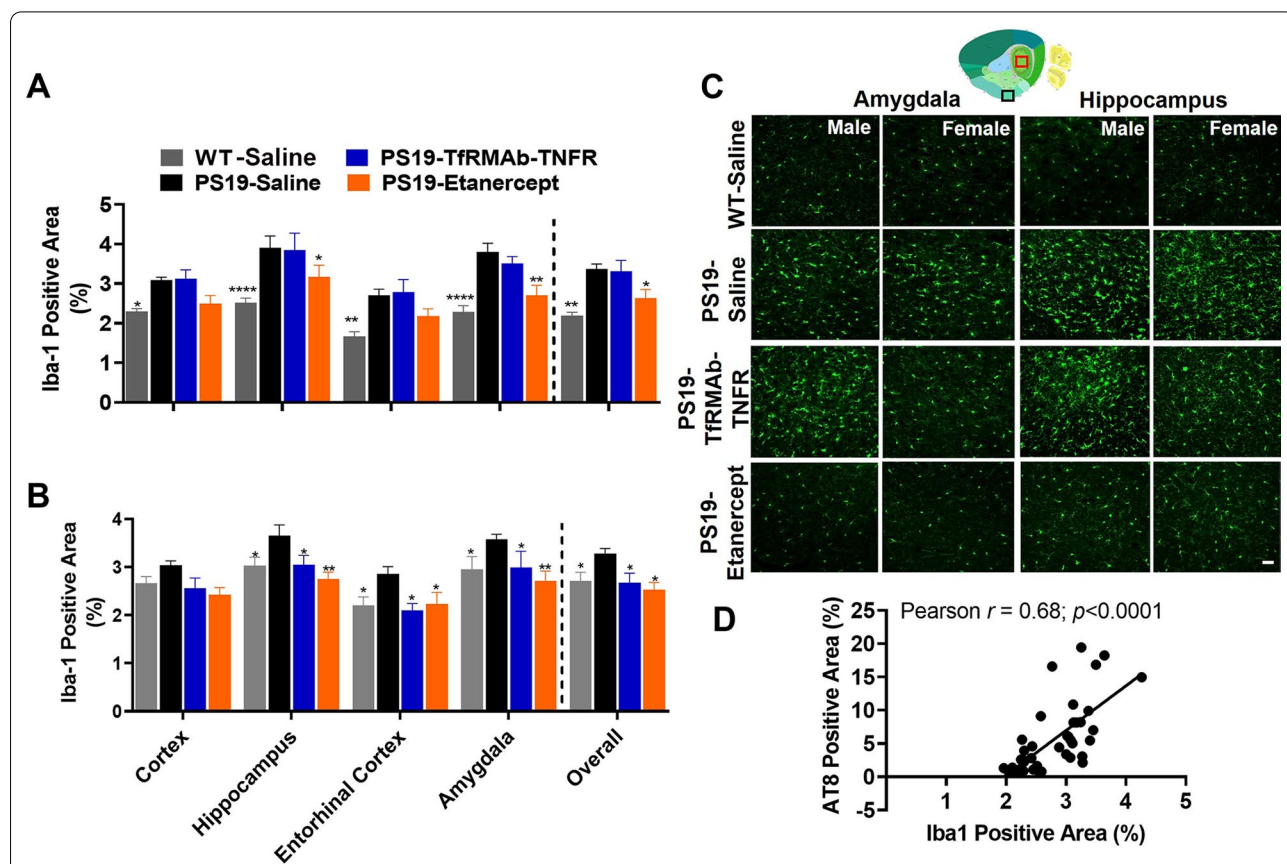
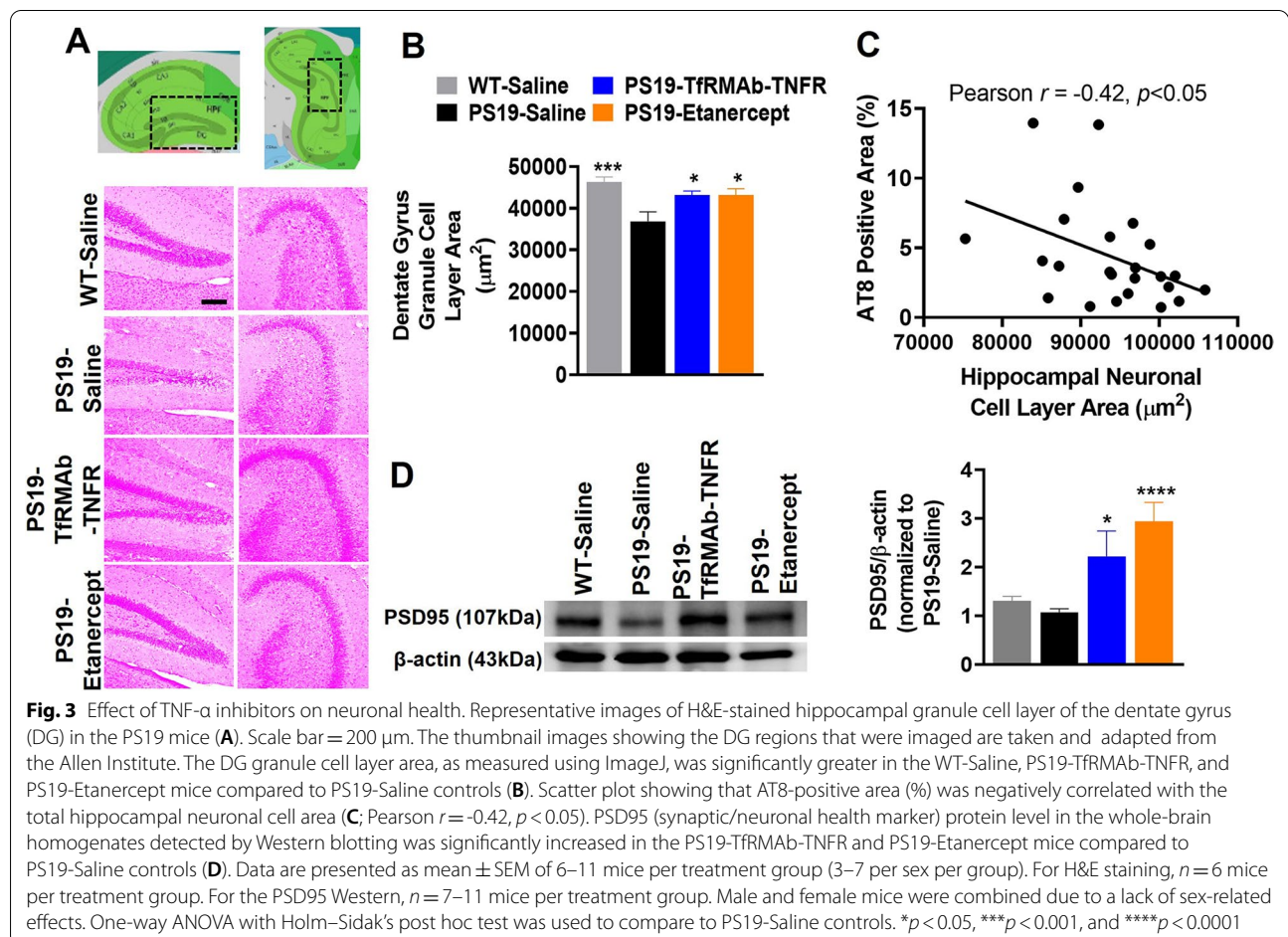


Fig. 2 Effect of TNF-α inhibitors on microgliosis in the PS19 mice. Iba-1-positive area in different brain regions in male (A) and female (B) mice. The overall Iba-1-positive area, which is an average of the regional Iba-1-positive area was significantly reduced by etanercept in both sexes and by TfRMAb-TNFR in female mice (A, B). Representative images of Iba-1-positive microglia in the amygdala (black boxed region representing the piriform-amygdalar region in the thumbnail brain section image) and the hippocampus (red boxed region in the thumbnail brain section image) of the PS19 mice (C). Scale bar = 50 μm and thumbnail image in C was taken from the Allen Institute. Scatter plot showing a strong positive correlation between the overall AT8-positive area (%) and the overall Iba-1-positive area (%) in males and female mice combined (D; Pearson $r = 0.68$, $p < 0.0001$). Data are presented as mean ± SEM of 10–12 mice per treatment group (5–7 per sex per group). Two-way ANOVA with repeated measures with Holm–Sidak’s post hoc test was used to compare to PS19-Saline controls. * $p < 0.05$, ** $p < 0.01$, **** $p < 0.0001$

S3). TfrMab-TNFR treatment significantly ($p < 0.05$) reduced the Iba-1-positive area by 18% in the female PS19 mice compared to PS19-Saline controls (Fig. 2B–C; Additional file 4: Fig. S3). No significant change in the Iba-1-positive area was observed in the TfrMab-TNFR-treated male PS19 mice (Fig. 2A–C; Additional file 4: Fig. S3). Etanercept significantly ($p < 0.05$) decreased the Iba-1-positive area in both male (24% decrease) and female (23% decrease) mice compared to sex-matched PS19-Saline controls (Fig. 2A–C; Additional file 4: Fig. S3). A strong significant positive correlation was observed in the overall AT8-positive and overall Iba-1-positive area in the current study (Pearson $r = 0.68$, $p < 0.0001$; Fig. 2D). To determine if the reduction in the Iba-1-positive area was due to a reduction in microglia number or size, we quantified the number of microglia and estimated the microglial soma size. We found a significant increase ($p < 0.01$) in the total number of microglia in the PS19-TfrMab-TNFR male mice compared to the PS19-Saline male mice (Additional file 5: Fig. S4A), but no difference in the total number of microglia in the female mice

(Additional file 5: Fig. S4B). With respect to the soma size, the number of microglia with a smaller soma size (< 50 -pixel units) was also significantly higher ($p < 0.01$) in the PS19-TfrMab-TNFR male mice compared to the PS19-Saline male mice (Additional file 5: Fig. S4C). The number of microglia with a larger soma (> 50 -pixel units) was significantly higher ($p < 0.0001$) in the PS19 male mice compared to the WT male mice (Additional file 5: Fig. S4E). Etanercept ($p < 0.0001$) and TfrMab-TNFR ($p < 0.01$) treatment significantly reduced the number of microglia with a larger soma in the male PS19 mice (Additional file 5: Fig. S4E). On the other hand, there was a significant reduction in the number of microglia with smaller soma size in the WT-Saline ($p < 0.001$), PS19-TfrMab-TNFR ($p < 0.001$) and PS19-Etanercept ($p < 0.01$) female mice compared with PS19-Saline female mice (Additional file 5: Fig. S4D). The number of microglia with a larger soma size was unchanged in the female mice (Additional file 5: Fig. S4F).



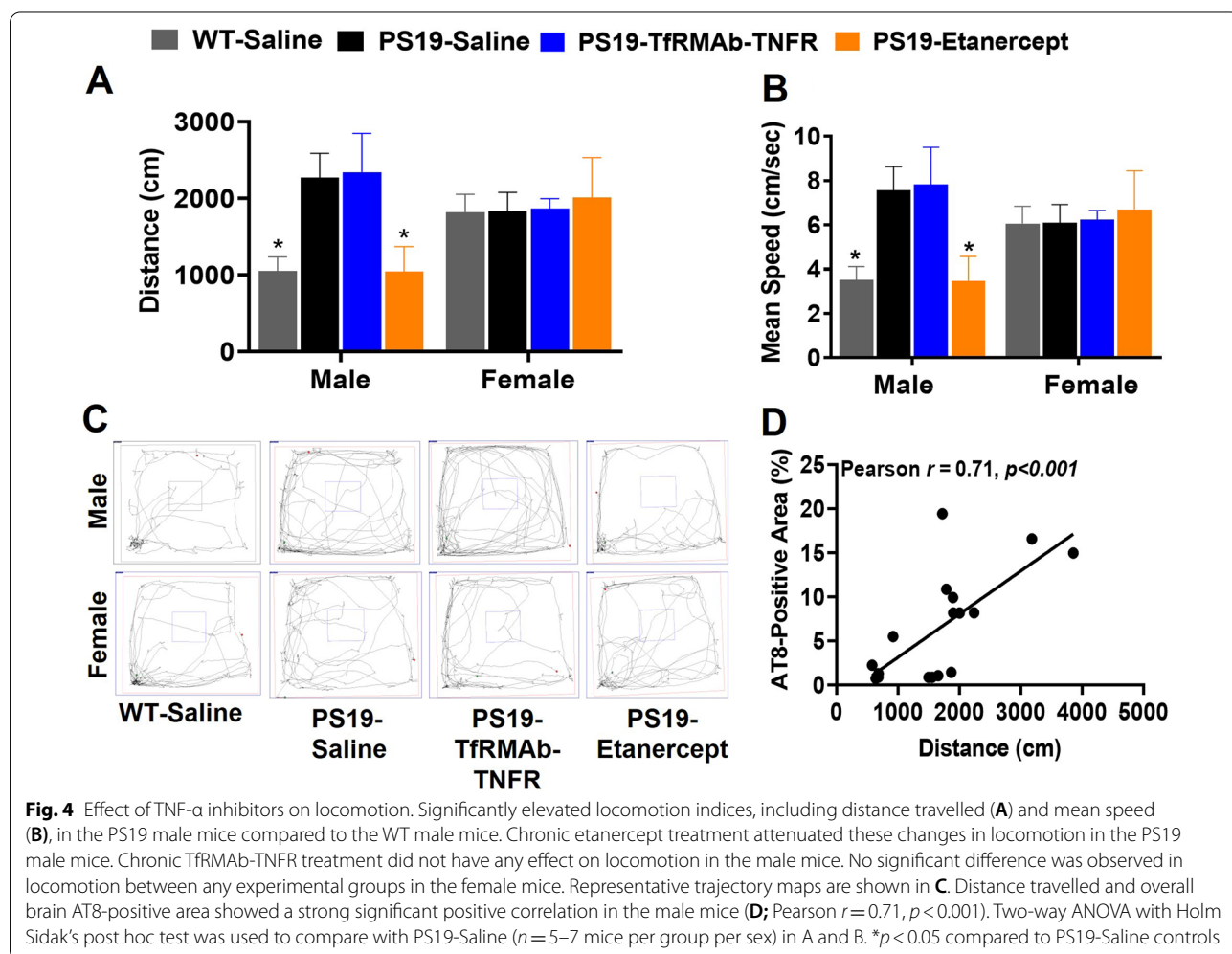
Effect of the biologic TNF- α inhibitors on neuronal health in the PS19 mice

The area occupied by the hippocampal granule cell layer of the DG was significantly smaller ($p < 0.001$) in the PS19-saline mice compared with the WT mice (Fig. 3A, B). The loss of the DG granule cell layer in the PS19 mice was significantly attenuated ($p < 0.05$) by both etanercept and TfRMAB-TNFR treatment (Fig. 3A, B). No significant difference was observed in the hippocampal pyramidal cell layer area of the CA1-3 regions between the experimental groups (data not shown). A small but significant negative correlation was observed in the overall AT8-positive and the total hippocampal neuronal cell area (Pearson $r = -0.42$, $p < 0.05$; Fig. 3C). The protein levels of the post-synaptic marker, PSD95 (a marker of synaptic health), in whole-brain homogenates, were not significantly reduced in the PS19-Saline controls (Fig. 3D). However, TfRMAB-TNFR and etanercept treatment led to a significant increase in the protein levels of

PSD95 in the whole-brain homogenates of PS19 mice (Fig. 3D).

Effect of TNF- α inhibitors on locomotion hyperactivity

The male PS19-Saline controls traveled a greater distance ($p < 0.05$) (Fig. 4A, C) and had a higher mean speed ($p < 0.05$) (Fig. 4B, C) compared to the WT male mice. Interestingly, this increase in distance traveled and mean speed was normalized by chronic etanercept treatment in the PS19 male mice, but not by chronic treatment with TfRMAB-TNFR (Fig. 4A–C). Neither the genotype nor the treatment affected locomotion hyperactivity in the female mice (Fig. 4A–C). A strong significant positive correlation was observed in the overall AT8-positive area in the brain and distance traveled in the male mice (Pearson $r = 0.71$, $p < 0.001$; Fig. 4D). No significant difference was observed in the time spent in the center, an indicator of anxiety-like behavior, between any experimental group in the current study (data not shown).



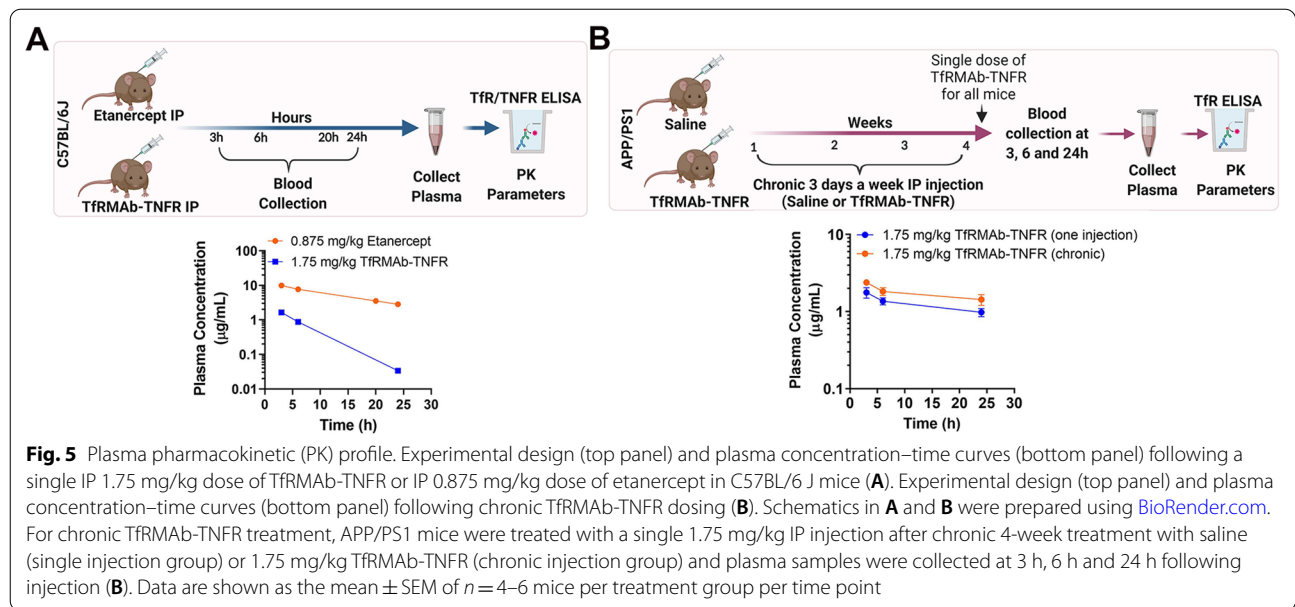


Table 1 Plasma pharmacokinetic (PK) parameters following a single IP injection of 1.75 mg/kg TFRMab-TNFR or 0.875 mg/kg etanercept, and following chronic IP injection of 1.75 mg/kg TFRMab-TNFR (a single 1.75 mg/kg TFRMab-TNFR injection after chronic 4-week treatment with saline or 1.75 mg/kg TFRMab-TNFR)

Route	Group	Dose mg/kg	Dosing frequency	AUC _(0–24 h) µg min/mL	C _{max} µg/mL	AUC _(0–∞) µg min/mL
IP	Single injection PK					
	TFRMab-TNFR	1.75	Once	864 ± 46	1.6 ± 0.08	875 ± 45
	Etanercept	0.875		7947 ± 664****	10.2 ± 1.3***	11,056 ± 843****
Chronic dosing PK	TFRMab-TNFR (single injection)	1.75	Once at the end of 4 weeks	1705 ± 157	1.8 ± 0.3	6114 ± 1268
	TFRMab-TNFR (chronic injection)	1.75	3 days per week for 4 weeks	2352 ± 280	2.4 ± 0.2	6960 ± 2033

Data are shown as the mean ± SEM (n = 4–6 mice per treatment group per time point). Two-sample t test was used to compare PK parameters ***p < 0.001, ****p < 0.0001 compared to TFRMab-TNFR for the single injection PK study.

Plasma PK after IP administration of biologic TNF-α inhibitors in mice

The plasma concentrations following a single IP administration of 1.75 mg/kg TFRMab-TNFR or 0.875 mg/kg etanercept are shown in Fig. 5A, and plasma PK parameters are summarized in Table 1. The plasma C_{max} was 1.6 ± 0.08 µg/mL for the TFRMab-TNFR fusion protein. The plasma C_{max} following an equimolar dose of etanercept was fivefold higher (p < 0.001) at 10.2 ± 1.3 µg/mL (Fig. 5A; Table 1). Accordingly, the plasma AUC over 24 h was eightfold higher (p < 0.001) for etanercept (7947 ± 664 µg•min/mL) compared to the TFRMab-TNFR fusion protein (864 ± 46 µg min/mL) (Table 1).

To study the impact of chronic TFRMab-TNFR dosing on plasma concentrations of the fusion protein, following 4-week chronic treatment with the TFRMab-TNFR or saline, mice received a single 1.75 mg/kg IP injection

of TFRMab-TNFR and plasma samples were collected at 3 h, 6 h, and 24 h following injection. No significant differences were observed in the plasma PK parameters between the 4-week saline-treated mice receiving a single dose of 1.75 mg/kg TFRMab-TNFR and 4-week TFRMab-TNFR-treated mice receiving a single dose of the TFRMab-TNFR (Fig. 5B; Table 1). The C_{max} values were comparable between the groups at 1.8 ± 0.3 µg/mL and 2.4 ± 0.2 µg/mL for the 4-week saline-treated mice receiving a single 1.75 mg/kg dose of TFRMab-TNFR and 4-week TFRMab-TNFR-treated mice receiving a single 1.75 mg/kg dose of TFRMab-TNFR, respectively (Table 1). Similarly, the plasma AUC over 24 h was comparable between the two groups at 2352 ± 280 µg min/mL and 1705 ± 157 µg•min/mL for the 4-week saline-treated mice receiving a single 1.75 mg/kg dose of TFRMab-TNFR and 4-week TFRMab-TNFR-treated mice

receiving a single 1.75 mg/kg dose of TfrMab-TNFR, respectively (Table 1). It should be noted that the plasma concentrations of TfrMab-TNFR following a single injection in Fig. 5A (blue curve) and Fig. 5B (blue curve) show differences that may be attributed to the different age and genotype of the mice used in Fig. 5A (8-week-old C57BL/6J) and Fig. 5B (19-month-old APP/PS1), but not to chronic saline dosing in Fig. 5B based on our previous work [44]. For this reason, the plasma concentrations following single and chronic TfrMab-TNFR dosing were compared in two separate groups of 19-month-old APP/PS1 (in Fig. 5B; blue and orange curves). The single injection data presented in Fig. 5A (blue curve) was not compared with Fig. 5B (blue curve).

Hematological profile

The hematologic profiles following a single 1.75 mg/kg TfrMab-TNFR IP injection in C57BL/6 J mice and chronic 8-week treatment with 1.75 mg/kg TfrMab-TNFR or 0.875 mg/kg etanercept in the PS19 mice are shown in Fig. 6A, B, respectively. A single injection of 1.75 mg/kg TfrMab-TNFR induced a significant 92% reduction ($p < 0.001$) in reticulocyte counts compared to saline controls (Fig. 6A), while the other hematologic parameters remained unchanged (Fig. 6A). Interestingly, the suppression of reticulocyte count following a single injection normalized after chronic treatment with 1.75 mg/kg TfrMab-TNFR for 8 weeks (Fig. 6B). Chronic treatment of TfrMab-TNFR did not result in significant changes in other hematological parameters compared to PS19-Saline controls (Fig. 6B). Chronic 8-week treatment with 0.875 mg/kg of etanercept led to a significant 58% increase ($p < 0.01$) in the platelet counts compared to PS19-Saline controls, and no change in any other hematological parameter was observed (Fig. 6B). All the mice survived the duration of the study and compared to the baseline body weights, no significant change in the body weights of mice was observed throughout the study for any experimental group (Fig. 6C).

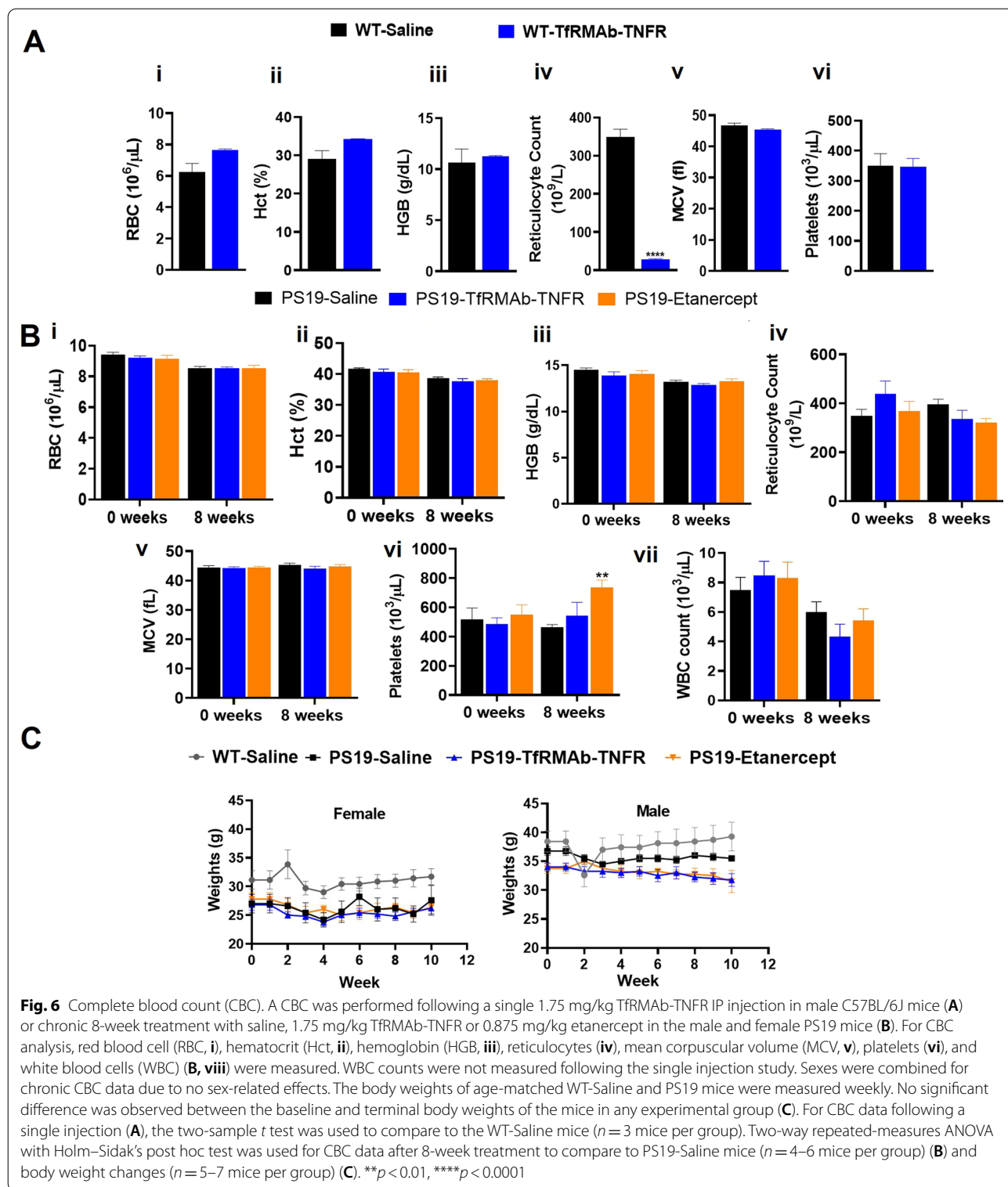
Discussion

The results of the current study show that both the BBB-penetrating biologic TNF- α inhibitor (TfrMab-TNFR) and the non-BBB-penetrating biologic TNF- α inhibitor (etanercept) exert protective effects in the female PS19 mice, with differential therapeutic effects in male PS19 mice. First, we show that etanercept significantly reduced pTau (ser202 and thr205) in both the male and female PS19 mice, and TfrMab-TNFR reduced pTau in female PS19 mice. Second, pTau was significantly correlated with microgliosis, and etanercept significantly reduced microgliosis in both male and female PS19 mice. TfrMab-TNFR reduced microgliosis in the female

PS19 mice. Third, both TfrMab-TNFR and etanercept improved neuronal health. Fourth, the locomotion hyperactivity of the PS19 male mice was attenuated with chronic etanercept treatment. Fifth, equimolar dosing of TfrMab-TNFR and etanercept resulted in significantly different plasma concentrations, which may explain the differential therapeutic effects of these biologic TNF- α inhibitors in male mice. Finally, chronic 8-week treatment with these biologic TNF- α inhibitors had no major impact on the hematologic parameters of the PS19 mice.

The link between microglial activation-mediated neuroinflammation and tau pathology in AD and other tauopathies has been extensively studied [51, 52]. Accumulating research in this area postulates that microglial activation is an early event in tauopathies that precedes tau phosphorylation and tangle formation, eventually causing neuronal cell death and neurodegeneration [53, 54]. An important event in this cascade of microglia-mediated abnormal tau lesion accumulation is the release of cytokines that directly augment tau phosphorylation and the formation of tau tangles both in vivo and in vitro. For example, direct application of IL-1 β or IL-6 to neuronal and microglial cultures increased tau phosphorylation and neuronal loss [55, 56]. Similarly, sustained brain levels of IL-1 β and TNF- α increased tau phosphorylation and neuronal cell death in vivo [15, 57]. Accordingly, CNS TNF- α inhibition with the intracerebroventricular injection of a TNF- α antibody, and peripheral TNF- α modulation significantly reduced tau phosphorylation in a mouse models of amyloidosis [22, 58].

Consistent with these findings, we found a significant reduction in the AT8-positive immunoreactive area, which measures pTau at sites ser202 and thr205, with chronic treatment with both the biologic TNF- α inhibitors. We also observed sex differences in the efficacy of the two biologic TNF- α inhibitors in the current study. The BBB-penetrating TfrMab-TNFR significantly reduced pTau in the female PS19 mice; interestingly, etanercept, which is shown not to cross the BBB [28], reduced pTau in both the male and female PS19 mice (Fig. 1; Additional file 3: Fig. S2). No change in total tau levels in the plasma and whole-brain homogenates was observed between the PS19-Saline and the biologic TNF- α inhibitor-treated PS19 mice (Additional file 3: Fig. S2). Further, the reduction in tau phosphorylation with the biologic TNF- α inhibitors paralleled the reduction in microgliosis in the current study, and the AT8-positive area showed a strong positive correlation with the Iba-1-positive area (Fig. 2). A recent report showed similar results wherein blockage of microglial proliferation with JNJ-40346527, a brain-permeant CSF1R inhibitor, reduced TNF- α and tau pathology in the spinal cords of the PS19 mice [59]. Together, these mouse



studies reinforce the association between pTau lesions and microglial activation seen in human AD brains [51, 60] and in rodent models, including the PS19 mouse model used in the current study [31, 61].

One key characteristic of microglia is an activation-induced change in their morphology [62–64]. Under physiological conditions, microglia are said to be in a resting/quiescent/surveillance state that is typically

characterized morphologically by the presence of highly ramified microglia with a small soma. However, once activated, the microglia transition into an amoeboid shape with an enlarged soma [62, 63], and soma size is a significant correlate of microgliosis [65]. Accordingly, the number of microglia with a larger soma size was significantly higher in the PS19 male mice compared to the WT mice suggesting the presence of more activated microglia in the PS19 male mice. The impact of the biologic TNF- α inhibitors on the microglial morphology varied by sex and the inhibitor. For example, the lack of change in microgliosis by TfrMAB-TNFR in the male mice (Fig. 2) appears to be driven by an increase in the number of microglia with a smaller soma size and an accompanying decrease in the number of microglia with a larger soma size (Additional file 5: Fig. S4). However, the reduction in microgliosis by etanercept appears to be driven by a reduction in the microglia with a larger soma size. In the female mice, the reduction in microgliosis appears to be driven by the reduction in the microglia with a smaller soma size rather than microglia with a larger soma size (Additional file 5: Fig. S4). Overall, our results suggest that the effect of the biologic TNF- α inhibitors on microgliosis is largely modulated by changes in microglia morphology rather than by changes in total microglia number. Future work using different markers for homeostatic (e.g., *P2ry12*) versus disease-associated (e.g., *Axl*, *ApoE4*) microglia will help elucidate the specific microglial phenotype modulated by the biologic TNF- α inhibitors [66].

Abnormal tau lesions correlate with neuronal loss and cognitive decline in AD [67], and a recent study showed that both tau and microglial activation can predict cognitive decline in AD [68]. In line with this, we wanted to determine if the attenuation in the pTau and microglial activation with the TNF- α inhibitors is associated with improvement or preservation of neuronal health. The PS19 mice exhibited a significant hippocampal neuronal loss in the DG, and we found a small but significant negative correlation between the AT8-positive area and hippocampal neuronal cell area in the PS19 mice (Fig. 3). These findings are consistent with the progressive hippocampal neuronal loss seen in the PS19 mouse model between 6- to 12-months of age [31]. We also measured the levels of PSD95, a post-synaptic protein that is widely used as a marker of synaptic health and is known to be modulated by TNF- α [69]. Notably, both the TfrMAB-TNFR and etanercept improved hippocampal and whole-brain tissue neuronal health in the current study (Fig. 3).

PS19 mice exhibit a hyperactive phenotype which is a characteristic of AD patients [39, 40]. Consistent with previous work with the PS19 mice [39], we found increased locomotion hyperactivity in the PS19 male

mice (Fig. 4). Interestingly, the locomotion hyperactivity was associated with increased pTau lesions in the male mice in the current study, showing a direct relationship between these parameters in these mice. Locomotor hyperactivity was attenuated by chronic etanercept treatment but not TfrMAB-TNFR treatment, paralleling the reduction in AT8-positive tau lesions with these TNF- α inhibitors in the male mice. Though the PS19 mice display a hyperactive phenotype, these mice may also display deficits in hippocampal-dependent learning [39–41], and future work will investigate the effects of the biologic TNF- α inhibitors using different hippocampal-dependent memory testing paradigms.

Both TfrMAB-TNFR and etanercept can modulate peripheral TNF- α following systemic administration, while only the TfrMAB-TNFR can enter the brain to exert direct CNS anti-TNF- α effects [36]. Using radio-labeled TfrMAB-TNFR in mice, we have previously shown that a single IP injection of TfrMAB-TNFR for doses between 1–3 mg/kg results in a brain concentration of ~0.5% injected dose/gram brain at 24 h after injection [35]. Based on this, the predicted brain concentration of TfrMAB-TNFR following a single 1.75 mg/kg dose is 0.3 μ g/g brain (2 nM) at 24 h following a single IP injection, assuming a 30 g mouse. The brain uptake of etanercept, on the other hand, is reported to be 20- to 13-fold lower than that of the brain-penetrating TNF- α inhibitor; furthermore, etanercept is found to distribute only to the brain blood volume, and not enter the brain parenchyma [28]. An alternate route for etanercept entry into the brain is via an impaired BBB in the PS19 mice. However, we did not observe changes in BBB tight junction proteins (ZO-1 and claudin-5) in the PS19 mice compared to the WT mice (Additional file 6: Fig. S5) consistent with previous work showing no change in BBB permeability in 8–10-month-old PS19 mice [70]. We measured TNF- α levels in the plasma and whole-brain homogenates using ELISA and Western blotting, respectively, but we did not see any significant change with the biologic TNF- α inhibitor treatment (Additional file 7: Fig. S6). However, we did see an increase in the whole-brain protein levels of I κ B α , which is degraded following TNF- α stimulation [71] and is therefore expected to increase after TNF- α inhibition, with TfrMAB-TNFR treatment ($p < 0.01$), and a similar trend was observed with etanercept treatment ($p = 0.08$; Additional file 7: Fig. S6). Taken together, the reduction in pTau, microgliosis, and improvement in neuronal health with both the TfrMAB-TNFR and etanercept in the current study indicates that these protective effects are modulated by peripheral, rather than direct CNS effects of these inhibitors in the PS19 mice. These findings are in agreement with recent reports showing that peripheral TNF- α alone can modulate microgliosis,

neuronal health, and A β -dependent AD pathology in the 5xFAD mice [19, 20]. Accordingly, inhibition of peripheral TNF- α and modulation of the innate immune system were shown to reduce A β -associated tau-phosphorylation in the 3xTg mouse model [58].

Despite the ability of both TfrMab-TNFR and etanercept to modulate peripheral TNF- α , we observed a significant difference in their therapeutic effects in the male PS19 mice. To elucidate the possible reason behind this difference in the male mice, we studied the plasma pharmacokinetics of these two inhibitors when given at equimolar doses to male mice. Furthermore, previous work with TfrMab-based fusion proteins shows no difference in plasma pharmacokinetics between male and female mice [72]. Similarly, etanercept pharmacokinetics show only minor insignificant differences between males and females and were reported to remain largely consistent between sexes [73, 74]. Therefore, female mice were not used for plasma pharmacokinetic studies.

Etanercept is 50% TNFR and TfrMab-TNFR is 25% TNFR based on amino acid sequence [36, 38]. As a result, the dose of etanercept was half that of TfrMab-TNFR in the current study. However, despite equimolar dosing, the plasma C_{max} was >500% higher and the plasma AUC was >800% higher for etanercept compared to the TfrMab-TNFR fusion protein after a single IP dose over 24 h (Fig. 5; Table 1). Plasma concentrations of TfrMab-based therapeutics may be further altered following chronic dosing. For example, our recent work reported a significant increase in plasma clearance, and thereby a significant decrease in the plasma concentrations of TfrMab (not fused to a therapeutic partner), following repeated administrations [47]. On the contrary, when the TfrMab was fused to glial-derived neurotrophic factor (GDNF), no change in plasma clearance or concentrations of the TfrMab-GDNF fusion protein was observed following chronic repeated dosing [72]. Based on this, we hypothesized that the plasma PK of TfrMab-based fusion proteins is highly regulated by the fusion partner [47]. To determine the impact of chronic TfrMab-TNFR dosing on plasma PK which may potentially impact the therapeutic efficacy of the fusion protein, we determined plasma concentrations of the fusion protein after repeated dosing. We found no change in the plasma concentrations of TfrMab-TNFR following repeated dosing (Fig. 5; Table 1). Overall, the TfrMab-TNFR fusion protein had a lower plasma concentration than etanercept, which is attributed to the Tfr-mediated peripheral clearance of the TfrMab-TNFR fusion protein from the circulation. The pTau load was 63% higher in the male mice compared to the female PS19 mice (Fig. 1), and it is conceivable that the higher pTau load seen in the male PS19 mice requires higher sustained plasma exposure

of the TNF- α inhibitor, or a higher dose to produce a therapeutic effect. The low plasma exposure of TfrMab-TNFR and the higher pTau load in the male PS19 mice may explain the lack of an effect of the TfrMab-TNFR on pTau and microgliosis in the male PS19 mice in the current study. Notably, our previous work comparing the TfrMab-TNFR and etanercept at a twofold higher dose (3 mg/kg) showed more robust effects of the TfrMab-TNFR than etanercept in the male APP/PS1 mouse model of amyloidosis [42].

TfrMab-based fusion proteins are associated with acute TfrMab-mediated reticulocyte suppression, and TNF- α inhibitors including etanercept are potent immune modulators that can cause leukopenia and increase the risk of infections [47, 75, 76]. Accordingly, we found a significant reduction in reticulocytes with a single dose of the TfrMab-TNFR, a TfrMab-based therapy (Fig. 6). However, this suppression was acute and no reduction in reticulocytes or any other hematologic parameter was observed following 8-week chronic dosing of the TfrMab-TNFR fusion protein. This is consistent with previous reports showing that reticulocyte suppression is short-lived and acute after administration of TfrMab-based fusion proteins [44, 47, 75]. Mice treated with etanercept, on the other hand, had significantly elevated platelet counts following chronic dosing, which has been previously reported with etanercept use [77], with no other change in the hematologic parameters. Further, no change in weight along with a largely stable hematologic profile showed that chronic treatment with the biologic TNF- α inhibitors was safe in the PS19 mice [78].

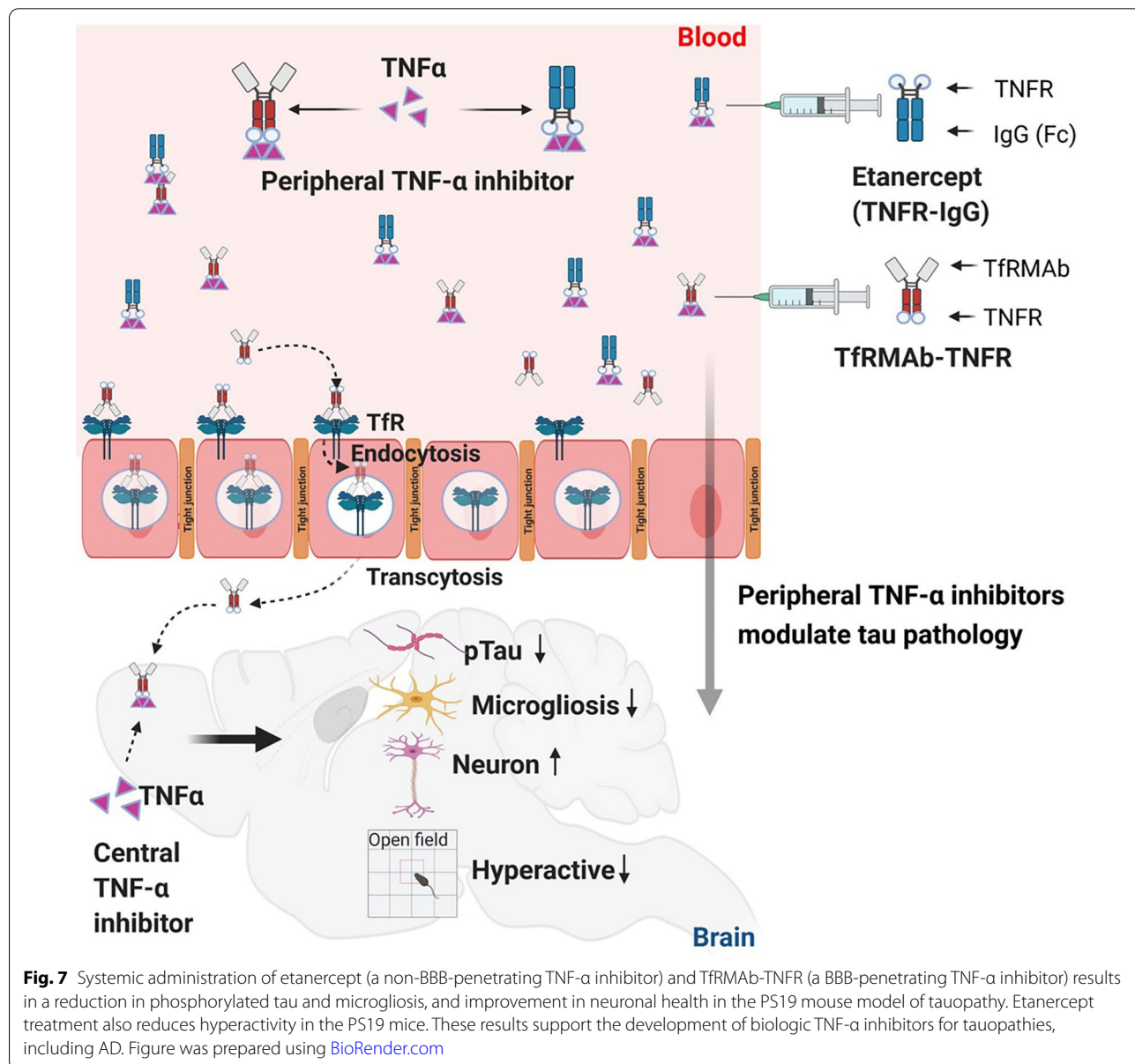
The current study has some limitations. First, the absence of thioflavin-S-positive tau tangles (data not shown) in the brains of the 8-month-old PS19 mice in the current study precluded our ability to explore the effect of the biologic TNF- α inhibitors on tau aggregates. Second, we did not study the impact of these biologic TNF- α inhibitors on tau pathology and microgliosis in the locus coeruleus (LC), which is one of the first brain structures to develop tau pathology and is implicated in tau spreading [79]. A previous study showed accumulation of tau tangles in the LC of PS19 mice, a process that is highly variable and dependent on the motor deficits in these mice [79], and future work will be needed to elucidate the effects of these biologic TNF- α inhibitors on tau pathology and microgliosis in the LC. Third, we have only studied the effect of the TNF- α inhibitors on tau phosphorylated at ser202 and thr205; the effect on tau phosphorylation at other sites was not studied. Fourth, we used whole-brain homogenates to study the levels of PSD95, and we did not see a reduction in PSD95 levels in PS19 mice compared to WT mice. Since the hippocampus is the primary site of neuronal loss in the PS19 mice

[31], the absence of a significant difference in the post-synaptic marker PSD95 between the PS19 and WT mice may be attributed to the use of whole-brain homogenate preparations in the current study rather than hippocampal brain homogenates to measure PSD95. Further, our data showing a reduction in microgliosis and pTau does not show causation. Whether the reduction in pTau reduces microgliosis or a reduction in microgliosis drives the reduction in pTau, is unclear. A recent study showed that amyloid potentiates microgliosis, which in turn drives tau pathology in the human AD brains [80]. Future work looking at the effects of these biologic TNF- α inhibitors in mouse models that combine both amyloid and

tau pathology, along with the use of CSF1R inhibitors to deplete microglia, will allow us to elucidate the role of microglia-derived TNF- α in the protective effects of these biologic TNF- α inhibitors on both amyloid and tau pathology.

Conclusion

Overall, the data presented in this in vivo study in the PS19 mouse model of tauopathy are the first to report significant therapeutic effects of biologic TNF- α inhibitors on A β -independent tau pathology following systemic administration (Fig. 7). Our results show that both the BBB-penetrating TfrMab-TNFR and



non-BBB-penetrating TNF- α inhibitor modulate the microglia-tau-neurodegeneration axis, suggesting the involvement of peripheral TNF- α in these processes. Future work replicating these findings in other models of tauopathies and models combining A β and tau pathology may yield important preclinical data supporting the use of biologic TNF- α inhibitors for AD and other tauopathies.

Abbreviations

AD: Alzheimer's disease; A β : Amyloid beta; BBB: Blood-brain barrier; BSA: Bovine serum albumin; CNS: Central nervous system; DG: Dentate Gyrus; GDNF: Glial-derived neurotrophic factor; Hct: Hematocrit; H&E: Hematoxylin and eosin; HGB: Hemoglobin; hTNF- α : Human TNF- α ; MCV: Mean corpuscular volume; PFA: Paraformaldehyde; PK: Pharmacokinetic; PBS: Phosphate buffer saline; pTau: Phospho-tau; RIPA: Radio-immunoprecipitation assay; SDS: Sodium dodecyl sulfate; TNFIs: TNF- α inhibitors; TNFR: TNF- α receptor; TNFR-1: TNF- α receptor 1; TFRMAB: Transferrin receptor antibody; TX100: Triton X-100; TNF- α : Tumor necrosis factor- α ; WBC: White blood cells; WT: Wild type.

Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s12974-021-02332-7>.

Additional file 1: Supplementary methods.

Additional file 2: Fig. S1 High-affinity binding of the TFRMAB-TNFR to the TFR (A) and TNF- α (B). iPSC-derived brain endothelial cells (ihBMECs) cultured in Transwell inserts were treated with 1 μ g/mL TNF- α and 7 μ g/mL TFRMAB-TNFR for up to 3 days. TNF- α reduced TEER (C) and increased sodium fluorescein passage across the brain endothelial monolayer (D). These effects were normalized by TFRMAB-TNFR treatment. All the experiments were repeated three times independently and data were expressed as mean \pm SEM. Two-way repeated-measures ANOVA or one-way ANOVA with Holm Sidak's post hoc test was used in C and D, respectively. * p <0.05, **** p <0.001, **** p <0.0001. OD: optical density.

Additional file 3: Fig. S2. Representative AT8-stained images of the cortex and entorhinal cortex. Images were acquired at 40X. Scale bar = 30 μ m (A). No significant change in the plasma (B) and brain total tau levels was detected using ELISA (C). A heat map showing the correlation between plasma total tau, brain total tau and overall AT8-positive area (%) in male and female mice combined (D). Total tau in plasma and brain shared a strong positive correlation (Pearson r = 0.76, p <0.0001). AT8-positive area showed a modest positive correlation with total tau in the plasma (Pearson r = 0.43, p <0.01) and brain (Pearson r = 0.46, p <0.05). Data are presented as mean \pm SEM of n = 6-11 mice per treatment group in B and C. Male and female mice were combined due to a lack of sex-related effects. One-way ANOVA with Holm-Sidak's post hoc test was used to compare to PS19-Saline controls in B and C. Pearson correlation was used for correlation analysis in D.

Additional file 4: Fig. S3. Representative Iba-1-stained images of the cortex and entorhinal cortex. Images were acquired at 10X with a digital 3X zoom. Scale bar = 25 μ m.

Additional file 5: Fig. S4. Data from male mice are shown in A, C and E, and data from female mice are shown in B, D and F. A significantly higher number of total microglia were observed in PS19-TFRMAB-TNFR male mice compared to the PS19-Saline male mice (A). There was no significant difference in the total number of microglia in the female mice (B). The overall number of microglia is the sum of the microglia in the cortex, hippocampus, amygdala, and the entorhinal cortex. There was a significant increase in the overall number of microglia with a smaller soma size (< 50-pixel units) in the PS19-TFRMAB-TNFR male mice (C) and a significant decrease in the overall number of microglia with a larger soma size (soma size > 50-pixels) in the PS19-TFRMAB-TNFR, PS19-Etanercept, and WT-Saline

male mice compared to PS19-Saline male mice (E). There was a significant decrease in the overall number of microglia with a smaller soma size (< 50-pixel units) in the PS19-TFRMAB-TNFR, PS19-Etanercept and WT-Saline female mice compared to the PS19-Saline female mice (D). There was no change in the number of microglia with a larger soma size (soma size > 50-pixels) in the female mice (F). Data are presented as mean \pm SEM of n = 5-7 per treatment group. Two-way ANOVA with repeated measures with Holm-Sidak's post hoc test was used to compare to PS19-Saline controls. * p <0.05, ** p <0.01, **** p <0.0001.

Additional file 6: Fig. S5. The protein levels of the BBB tight junction proteins, ZO-1 (A) and claudin-5 (B), in whole-brain homogenates measured using Western blotting, were not significantly changed in the PS19-Saline controls compared to the WT-Saline group. Data are presented as mean \pm SEM of n = 6-8 per treatment group. A Student's t-test was used to compare the two groups.

Additional file 7: Fig. S6. Levels of TNF- α in the plasma (A) and whole-brain homogenates (B) were not significantly different between PS19-Saline and PS19-TFRMAB-TNFR and PS19-Etanercept mice, respectively. We attribute the lack of change in plasma and brain TNF- α levels with biologic TNF- α inhibitors to the time of plasma and brain sample collection. There was a 10-day lag between the last treatment dose (8 weeks after treatment initiation) and sample collection due to time for open-field testing (during week 9; Figure 1A). The plasma elimination half-lives of TFRMAB-TNFR and etanercept are ~ 4 h and ~13 h, respectively (Figure 5). Since it takes 6 elimination half-lives (24 h and 78 h for TFRMAB-TNFR and etanercept, respectively) for a drug to be completely eliminated from the blood circulation, we do not expect any circulating biologic TNF- α inhibitor at the time of sacrifice. Another potential reason for the lack of difference in plasma and brain TNF- α is the use of immunoassays. Since both the TFRMAB-TNFR and etanercept bind to TNF- α , detection of TNF- α using immunoassays may be complicated if the TNF- α is still bound to the biologic TNF- α inhibitor in the brain at the time of mouse sacrifice. Though this is unlikely considering the elimination half-lives of the drugs, to rule out any such interference, we measured the protein levels of I κ B α , which is degraded following TNF- α stimulation and is therefore expected to increase following TNF- α inhibition, in whole-brain homogenates using Western blotting (C). The TFRMAB-TNFR-treated PS19 mice had significantly higher (p <0.01) brain I κ B α (an indirect measure of the attenuation of TNF- α signaling). Similarly, PS19 mice treated with etanercept showed a trend (p =0.08) towards an increase in brain I κ B α compared to saline-treated PS19 mice, but this data did not reach statistical significance. Data are presented as mean \pm SEM of n = 7-11 mice per treatment group. Male and female mice were combined due to a lack of sex-related effects. One-way ANOVA with Holm-Sidak's post hoc test was used to compare to PS19-Saline controls. ** p <0.01.

Acknowledgements

We would like to thank Abheerava Koka of Oak Park High School and Sreya Chilukuri of La Salle High School for helping with AT8 ImageJ quantification. Mark Sbertole for assistance with animal care, and Jessica M. Phan for genotyping of transgenic mice. The authors also thank Maria Melville of the Keck Graduate Institute for her contribution to the confocal imaging.

Authors' contributions

WO designed and performed the experiments, analyzed data, prepared the figures, and wrote the paper. JY performed the PK studies and the confocal imaging. JS and MC helped with the H&E experiments and ImageJ analysis. DMC and RC designed and performed the immunostaining and tau ELISA. JS performed the in vitro studies and helped with the in vivo experiments. NJ performed the microglial counts and Western blotting. KP and DHC provided the mouse resources and assisted with the PK studies. RKS conceived the study, designed and coordinated the experiments, acquired the funding, and helped in the drafting and editing of the manuscript. All authors contributed to manuscript writing, and have read and approved the final manuscript.

Funding

Research reported in this publication was supported by the National Institute of Aging of the National Institutes of Health under award number

R01AG062840 awarded to R.K.S. Approximately, \$250k of federal funds supported the effort (89%) on this project. The content is solely the responsibility of the authors and does not necessarily represent the official views of the National Institutes of Health. Approximately \$30k (11%) of non-federal funds from the Joseph H. Stahlberg Foundation awarded to R.K.S also supported the effort on this project.

Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Declarations

Ethics approval and consent to participate

All animal procedures were approved by the University of California Irvine, University of La Verne or Pomona College Institutional Animal Care and Use Committee and were carried out in compliance with University Laboratory Animal Resources regulations. This work does not involve any applicable consent to participate.

Consent for publication

None.

Competing interests

The authors declare that they have no competing interests.

Author details

¹Department of Biomedical and Pharmaceutical Sciences, School of Pharmacy, Chapman University, Irvine, CA 92618, USA. ²Henry E. Riggs School of Applied Life Sciences, Keck Graduate Institute, Claremont, CA 91711, USA. ³Department of Neuroscience, Pomona College, Claremont, CA 91711, USA. ⁴Keck Science Department, Claremont McKenna College, Claremont, CA 91711, USA. ⁵MIND Institute, University of California, Irvine, CA 92697, USA. ⁶Department of Neurology, University of California, Irvine, CA 92868, USA.

Received: 9 May 2021 Accepted: 26 November 2021

Published online: 31 December 2021

References

- Hippus H, Neundörfer G. The discovery of Alzheimer's disease. *Dialogues Clin Neurosci*. 2003;5(1):101–8.
- Perl DP. Neuropathology of Alzheimer's disease. *Mt Sinai J Med*. 2010;77(1):32–42.
- Spillantini MG, Goedert M. Tau pathology and neurodegeneration. *Lancet Neurol*. 2013;12(6):609–22.
- Parbo P, Ismail R, Sommerauer M, Stokholm MG, Hansen AK, Hansen KV, et al. Does inflammation precede tau aggregation in early Alzheimer's disease? A PET study. *Neurobiol Dis*. 2018;117:211–6.
- Wyss-Coray T. Inflammation in Alzheimer disease: driving force, bystander or beneficial response? *Nat Med*. 2006;12(9):1005–15.
- Tarkowski E, Andreasen N, Tarkowski A, Blennow K. Intrathecal inflammation precedes development of Alzheimer's disease. *J Neurol Neurosurg Psychiatry*. 2003;74(9):1200–5.
- Zelova H, Hosek J. TNF-alpha signalling and inflammation: interactions between old acquaintances. *Inflamm Res*. 2013;62(7):641–51.
- Alvarez A, Cacabelos R, Sanpedro C, Garcia-Fantini M, Aleixandre M. Serum TNF-alpha levels are increased and correlate negatively with free IGF-I in Alzheimer disease. *Neurobiol Aging*. 2007;28(4):533–6.
- Paganelli R, Di Iorio A, Patricelli L, Ripani F, Sparvieri E, Faricelli R, et al. Pro-inflammatory cytokines in sera of elderly patients with dementia: levels in vascular injury are higher than those of mild-moderate Alzheimer's disease patients. *Exp Gerontol*. 2002;37(2–3):257–63.
- Akiyama H, Barger S, Barnum S, Bradt B, Bauer J, Cole GM, et al. Inflammation and Alzheimer's disease. *Neurobiol Aging*. 2000;21(3):383–421.
- Patel NS, Paris D, Mathura V, Quadros AN, Crawford FC, Mullan MJ. Inflammatory cytokine levels correlate with amyloid load in transgenic mouse models of Alzheimer's disease. *J Neuroinflammation*. 2005;2(1):9.
- He P, Zhong Z, Lindholm K, Berning L, Lee W, Lemere C, et al. Deletion of tumor necrosis factor death receptor inhibits amyloid beta generation and prevents learning and memory deficits in Alzheimer's mice. *J Cell Biol*. 2007;178(5):829–41.
- Yamamoto M, Kiyota T, Horiba M, Buescher JL, Walsh SM, Gendelman HE, et al. Interferon-gamma and tumor necrosis factor-alpha regulate amyloid-beta plaque deposition and beta-secretase expression in Swedish mutant APP transgenic mice. *Am J Pathol*. 2007;170(2):680–92.
- Liao YF, Wang BJ, Cheng HT, Kuo LH, Wolfe MS. Tumor necrosis factor-alpha, interleukin-1beta, and interferon-gamma stimulate gamma-secretase-mediated cleavage of amyloid precursor protein through a JNK-dependent MAPK pathway. *J Biol Chem*. 2004;279(47):49523–32.
- Janelins MC, Mastrangelo MA, Park KM, Sudol KL, Narrow WC, Oddo S, et al. Chronic neuron-specific tumor necrosis factor-alpha expression enhances the local inflammatory environment ultimately leading to neuronal death in 3xTg-AD mice. *Am J Pathol*. 2008;173(6):1768–82.
- Ralay Ranaivo H, Craft JM, Hu W, Guo L, Wing LK, Van Eldik LJ, et al. Glia as a therapeutic target: selective suppression of human amyloid-beta-induced upregulation of brain proinflammatory cytokine production attenuates neurodegeneration. *J Neurosci*. 2006;26(2):662–70.
- Rosenberg PB. Clinical aspects of inflammation in Alzheimer's disease. *Int Rev Psychiatry*. 2005;17(6):503–14.
- McGeer EG, McGeer PL. Inflammatory processes in Alzheimer's disease. *Prog Neuropsychopharmacol Biol Psychiatry*. 2003;27(5):741–9.
- Paouri E, Tzara O, Kartalou GI, Zenelak S, Georgopoulos S. Peripheral tumor necrosis factor-alpha (TNF- α) modulates amyloid pathology by regulating blood-derived immune cells and glial response in the brain of AD/TNF transgenic mice. *J Neurosci*. 2017;37(20):5155–71.
- Kalovyra N, Apokotou O, Boulekou S, Paouri E, Boutou A, Georgopoulos S. A 3'UTR modification of the TNF- α mouse gene increases peripheral TNF- α and modulates the Alzheimer-like phenotype in 5XFAD mice. *Sci Rep*. 2020;10(1):8670.
- Collins JS, Perry RT, Watson B Jr, Harrell LE, Acton RT, Blacker D, et al. Association of a haplotype for tumor necrosis factor in siblings with late-onset Alzheimer disease: the NIMH Alzheimer disease genetics initiative. *Am J Med Genet*. 2000;96(6):823–30.
- Shi JQ, Shen W, Chen J, Wang BR, Zhong LL, Zhu YW, et al. Anti-TNF- α reduces amyloid plaques and tau phosphorylation and induces CD11c-positive dendritic-like cell in the APP/PS1 transgenic mouse brains. *Brain Res*. 2011;1368:239–47.
- Shi JQ, Wang BR, Jiang WW, Chen J, Zhu YW, Zhong LL, et al. Cognitive improvement with intrathecal administration of infliximab in a woman with Alzheimer's disease. *J Am Geriatr Soc*. 2011;59(6):1142–4.
- Kim DH, Choi SM, Jho J, Park MS, Kang J, Park SJ, et al. Infliximab ameliorates AD-associated object recognition memory impairment. *Behav Brain Res*. 2016;311:384–91.
- Tobinick E, Gross H, Weinberger A, Cohen H. TNF-alpha modulation for treatment of Alzheimer's disease: a 6-month pilot study. *MedGenMed*. 2006;8(2):25.
- McAlpine FE, Lee JK, Harms AS, Ruhn KA, Blurton-Jones M, Hong J, et al. Inhibition of soluble TNF signaling in a mouse model of Alzheimer's disease prevents pre-plaque amyloid-associated neuropathology. *Neurobiol Dis*. 2009;34(1):163–77.
- Steeland S, Gorlé N, Vandendriessche C, Balus S, Brkic M, Van Cauwenberghe C, et al. Counteracting the effects of TNF receptor-1 has therapeutic potential in Alzheimer's disease. *EMBO Mol Med*. 2018;10(4).
- Boado RJ, Hui EK, Lu JZ, Zhou QH, Pardridge WM. Selective targeting of a TNFR decoy receptor pharmaceutical to the primate brain as a receptor-specific IgG fusion protein. *J Biotechnol*. 2010;146(1–2):84–91.
- Wang Y, Mandelkow E. Tau in physiology and pathology. *Nat Rev Neurosci*. 2016;17(1):5–21.
- Laurent C, Buée L, Blum D. Tau and neuroinflammation: what impact for Alzheimer's disease and tauopathies? *Biomed J*. 2018;41(1):21–33.
- Yoshiyama Y, Higuchi M, Zhang B, Huang SM, Iwata N, Saido TC, et al. Synapse loss and microglial activation precede tangles in a P301S tauopathy mouse model. *Neuron*. 2007;53(3):337–51.
- Maphis N, Xu G, Kokiko-Cochran ON, Jiang S, Cardona A, Ransohoff RM, et al. Reactive microglia drive tau pathology and contribute to the spreading of pathological tau in the brain. *Brain*. 2015;138(Pt 6):1738–55.

33. Garwood CJ, Cooper JD, Hanger DP, Noble W. Anti-inflammatory impact of minocycline in a mouse model of tauopathy. *Front Psychiatry*. 2010;1:136.
34. Yoshiyama Y, Kojima A, Ishikawa C, Arai K. Anti-inflammatory action of donepezil ameliorates tau pathology, synaptic loss, and neurodegeneration in a tauopathy mouse model. *J Alzheimers Dis*. 2010;22(1):295–306.
35. Sumbria RK, Zhou QH, Hui EK, Lu JZ, Boado RJ, Pardridge WM. Pharmacokinetics and brain uptake of an IgG-TNF decoy receptor fusion protein following intravenous, intraperitoneal, and subcutaneous administration in mice. *Mol Pharm*. 2013;10(4):1425–31.
36. Zhou QH, Boado RJ, Hui EK, Lu JZ, Pardridge WM. Brain-penetrating tumor necrosis factor decoy receptor in the mouse. *Drug Metab Dispos*. 2011;39(1):71–6.
37. Chang R, Knox J, Chang J, Derbedrossian A, Vasilevko V, Cribbs D, et al. Blood–brain barrier penetrating biologic TNF- α inhibitor for Alzheimer's disease. *Mol Pharm*. 2017;14(7):2340–9.
38. Hassett B, McMillen S, Fitzpatrick B. Characterization and comparison of commercially available TNF receptor 2-Fc fusion protein products: letter to the editor. *MAbs*. 2013;5(5):624–5.
39. Dumont M, Stack C, Elipenahli C, Jainuddin S, Geroges M, Starkova NN, et al. Behavioral deficit, oxidative stress, and mitochondrial dysfunction precede tau pathology in P301S transgenic mice. *Faseb j*. 2011;25(11):4063–72.
40. Takeuchi H, Iba M, Inoue H, Higuchi M, Takao K, Tsukita K, et al. P301S mutant human tau transgenic mice manifest early symptoms of human tauopathies with dementia and altered sensorimotor gating. *PLoS ONE*. 2011;6(6):e21050.
41. Patel H, Martinez P, Perkins A, Taylor X, Jury N, McKinzie D, et al. Pathological tau and reactive astrogliosis are associated with distinct functional deficits in a mouse model of tauopathy. *Neurobiol Aging*. 2022;109:52–63.
42. Chang R, Knox J, Chang J, Derbedrossian A, Vasilevko V, Cribbs D, et al. Blood–brain barrier penetrating biologic TNF- α inhibitor for Alzheimer's disease. *Mol Pharm*. 2017;14(7):2340–9.
43. Chang R, Al Maghribi A, Vanderpoel V, Vasilevko V, Cribbs DH, Boado R, et al. Brain penetrating bifunctional erythropoietin-transferrin receptor antibody fusion protein for Alzheimer's disease. *Mol Pharm*. 2018;15(11):4963–73.
44. Sun J, Yang J, Whitman K, Zhu C, Cribbs DH, Boado RJ, et al. Hematologic safety of chronic brain-penetrating erythropoietin dosing in APP/PS1 mice. *Alzheimers Dement (N Y)*. 2019;5:627–36.
45. Pan B, Zhang H, Cui T, Wang X. TFEB activation protects against cardiac proteotoxicity via increasing autophagic flux. *J Mol Cell Cardiol*. 2017;113:51–62.
46. Pan B, Li J, Parajuli N, Tian Z, Wu P, Lewno MT, et al. The calcineurin-TFEB-p62 pathway mediates the activation of cardiac macroautophagy by proteasomal malfunction. *Circ Res*. 2020;127(4):502–18.
47. Castellanos DM, Sun J, Yang J, Ou W, Zambon AC, Pardridge WM, et al. Acute and chronic dosing of a high-affinity rat/mouse chimeric transferrin receptor antibody in mice. *Pharmaceutics*. 2020;12(9):852.
48. Yang J, Sun J, Castellanos DM, Pardridge WM, Sumbria RK. Eliminating Fc N-linked glycosylation and its impact on dosing consideration for a transferrin receptor antibody-erythropoietin fusion protein in mice. *Mol Pharm*. 2020;17(8):2831–9.
49. Sun J, Boado RJ, Pardridge WM, Sumbria RK. Plasma pharmacokinetics of high-affinity transferrin receptor antibody-erythropoietin fusion protein is a function of effector attenuation in mice. *Mol Pharm*. 2019;16(8):3534–43.
50. Pan B, Lewno MT, Wu P, Wang X. Highly dynamic changes in the activity and regulation of macroautophagy in hearts subjected to increased proteotoxic stress. *Front Physiol*. 2019;10:758.
51. Sheffield LG, Marquis JG, Berman NE. Regional distribution of cortical microglia parallels that of neurofibrillary tangles in Alzheimer's disease. *Neurosci Lett*. 2000;285(3):165–8.
52. Ishizawa K, Dickson DW. Microglial activation parallels system degeneration in progressive supranuclear palsy and corticobasal degeneration. *J Neuropathol Exp Neurol*. 2001;60(6):647–57.
53. Leys CEG, Holtzman DM. Glial contributions to neurodegeneration in tauopathies. *Mol Neurodegener*. 2017;12(1):50.
54. Zilka N, Kazmerova Z, Jadhav S, Neradil P, Madari A, Obetkova D, et al. Who fans the flames of Alzheimer's disease brains? Misfolded tau on the crossroad of neurodegenerative and inflammatory pathways. *J Neuroinflammation*. 2012;9:47.
55. Quintanilla RA, Orellana DI, Gonzalez-Billault C, Maccioni RB. Interleukin-6 induces Alzheimer-type phosphorylation of tau protein by deregulating the cdk5/p35 pathway. *Exp Cell Res*. 2004;295(1):245–57.
56. Li Y, Liu L, Barger SW, Griffin WS. Interleukin-1 mediates pathological effects of microglia on tau phosphorylation and on synaptophysin synthesis in cortical neurons through a p38-MAPK pathway. *J Neurosci*. 2003;23(5):1605–11.
57. Sheng JG, Zhu SG, Jones RA, Griffin WS, Mrak RE. Interleukin-1 promotes expression and phosphorylation of neurofilament and tau proteins in vivo. *Exp Neurol*. 2000;163(2):388–91.
58. Gabbita SP, Johnson MF, Kobritz N, Esami P, Poteshkina A, Varadarajan S, et al. Oral TNF α modulation alters neutrophil infiltration, improves cognition and diminishes tau and amyloid pathology in the 3xTgAD mouse model. *PLoS ONE*. 2015;10(10):e0137305.
59. Mancuso R, Fryatt G, Cleal M, Obst J, Pipi E, Monzon-Sandoval J, et al. CSF1R inhibitor JNJ-40346527 attenuates microglial proliferation and neurodegeneration in P301S mice. *Brain*. 2019;142(10):3243–64.
60. DiPatre PL, Gelman BB. Microglial cell activation in aging and Alzheimer disease: partial linkage with neurofibrillary tangle burden in the hippocampus. *J Neuropathol Exp Neurol*. 1997;56(2):143–9.
61. Bellucci A, Westwood AJ, Ingram E, Casamenti F, Goedert M, Spillantini MG. Induction of inflammatory mediators and microglial activation in mice transgenic for mutant human P301S tau protein. *Am J Pathol*. 2004;165(5):1643–52.
62. Paasila PJ, Davies DS, Kril JJ, Goldsburly C, Sutherland GT. The relationship between the morphological subtypes of microglia and Alzheimer's disease neuropathology. *Brain Pathol*. 2019;29(6):726–40.
63. Karperien A, Ahammer H, Jelinek HF. Quantitating the subtleties of microglial morphology with fractal analysis. *Front Cell Neurosci*. 2013;7:3.
64. Wang J, Zhao D, Pan B, Fu Y, Shi F, Kouadir M, et al. Toll-like receptor 2 deficiency shifts PrP¹⁰⁶⁻¹²⁶-induced microglial activation from a neurotoxic to a neuroprotective phenotype. *J Mol Neurosci*. 2015;55(4):880–90.
65. Kozlowski C, Weimer RM. An automated method to quantify microglia morphology and application to monitor activation state longitudinally in vivo. *PLoS ONE*. 2012;7(2):e31814.
66. Deczkowska A, Keren-Shaul H, Weiner A, Colonna M, Schwartz M, Amit I. Disease-associated microglia: a universal immune sensor of neurodegeneration. *Cell*. 2018;173(5):1073–81.
67. Nelson PT, Alafuzoff I, Bigio EH, Bouras C, Braak H, Cairns NJ, et al. Correlation of Alzheimer disease neuropathologic changes with cognitive status: a review of the literature. *J Neuropathol Exp Neurol*. 2012;71(5):362–81.
68. Malpetti M, Kievit RA, Passamonti L, Jones PS, Tsvetanov KA, Rittman T, et al. Microglial activation and tau burden predict cognitive decline in Alzheimer's disease. *Brain*. 2020;143(5):1588–602.
69. Poon VY, Choi S, Park M. Growth factors in synaptic function. *Front Synaptic Neurosci*. 2013;5:6.
70. Bien-Ly N, Boswell CA, Jeet S, Beach TG, Hoyte K, Luk W, et al. Lack of widespread BBB disruption in Alzheimer's disease models: focus on therapeutic antibodies. *Neuron*. 2015;88(2):289–97.
71. Iwai K. Diverse ubiquitin signaling in NF- κ B activation. *Trends Cell Biol*. 2012;22(7):355–64.
72. Zhou QH, Boado RJ, Hui EK, Lu JZ, Pardridge WM. Chronic dosing of mice with a transferrin receptor monoclonal antibody-glia-derived neurotrophic factor fusion protein. *Drug Metab Dispos*. 2011;39(7):1149–54.
73. Zhou H. Clinical pharmacokinetics of etanercept: a fully humanized soluble recombinant tumor necrosis factor receptor fusion protein. *J Clin Pharmacol*. 2005;45(5):490–7.
74. Korth-Bradley JM, Rubin AS, Hanna RK, Simcoe DK, Lebsack ME. The pharmacokinetics of etanercept in healthy volunteers. *Ann Pharmacother*. 2000;34(2):161–4.
75. Couch JA, Yu YJ, Zhang Y, Tarrant JM, Fuji RN, Meilandt WJ, et al. Addressing safety liabilities of Tfr bispecific antibodies that cross the blood-brain barrier. *Sci Transl Med*. 2013;5(183):183ra57 (1–12).
76. Hyrich KL, Silman AJ, Watson KD, Symmons DP. Anti-tumour necrosis factor alpha therapy in rheumatoid arthritis: an update on safety. *Ann Rheum Dis*. 2004;63(12):1538–43.
77. Pathare SK, Heycock C, Hamilton J. TNF α blocker-induced thrombocytopenia. *Rheumatology (Oxford)*. 2006;45(10):1313–4.

78. Propson NE, Roy ER, Litvinchuk A, Köhl J, Zheng H. Endothelial C3a receptor mediates vascular inflammation and blood–brain barrier permeability during aging. *J Clin Invest.* 2021;131(1).
79. Iba M, Guo JL, McBride JD, Zhang B, Trojanowski JQ, Lee VM. Synthetic tau fibrils mediate transmission of neurofibrillary tangles in a transgenic mouse model of Alzheimer’s-like tauopathy. *J Neurosci.* 2013;33(3):1024–37.
80. Pascoal TA, Benedet AL, Ashton NJ, Kang MS, Therriault J, Chamoun M, et al. Microglial activation and tau propagate jointly across Braak stages. *Nat Med.* 2021;27(9):1592–9.

Publisher’s Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Ready to submit your research? Choose BMC and benefit from:

- fast, convenient online submission
- thorough peer review by experienced researchers in your field
- rapid publication on acceptance
- support for research data, including large and complex data types
- gold Open Access which fosters wider collaboration and increased citations
- maximum visibility for your research: over 100M website views per year

At BMC, research is always in progress.

Learn more biomedcentral.com/submissions

