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Zinc status alters Alzheimer's disease progression through NLRP3-dependent inflammation

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- 1 Zinc status alters Alzheimer's disease progression through NLRP3-dependent
- 2 inflammation.
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Abstract

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Alzheimer's disease is a devastating neurodegenerative disease with a dramatically increasing prevalence and no disease-modifying treatment. Inflammatory lifestyle factors increase the risk of developing Alzheimer's disease. Zinc deficiency is the most prevalent malnutrition in the world and may be a risk factor for Alzheimer's disease potentially through enhanced inflammation, although evidence for this is Here we provide epidemiological evidence suggesting that zinc supplementation was associated with reduced risk and slower cognitive decline, in people with Alzheimer's disease and mild cognitive impairment. Using the APP/PS1 mouse model of Alzheimer's disease fed a control (35mg/kg zinc) or diet deficient in zinc (3mg/kg zinc), we determined that zinc deficiency accelerated Alzheimer's-like memory deficits without modifying amyloid beta (β) plaque burden in the brains of male mice. The NLRP3-inflammasome complex is one of the most important regulators of inflammation and we show here that zinc deficiency in immune cells, including microglia, potentiated NLRP3 responses to inflammatory stimuli in vitro including amyloid oligomers, while zinc supplementation inhibited NLRP3 activation. APP/PS1 mice deficient in NLRP3 were protected against the accelerated cognitive decline with zinc deficiency. Collectively, this research suggests that zinc status is linked to inflammatory reactivity and may be modified in people to reduce the risk and slow the progression of Alzheimer's disease.

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Significance Statement

Alzheimer's disease is a common condition mostly affecting the elderly. Zinc deficiency is also a global problem, especially in the elderly and also in people with Alzheimer's disease. Zinc deficiency contributes to many clinical disorders including immune dysfunction. Inflammation is known to contribute to the risk and progression

of Alzheimer's disease, thus, we hypothesised that zinc status would affect
Alzheimer's disease progression. Here we show that zinc supplementation reduced
the prevalence and symptomatic decline in people with Alzheimer's disease. In an
animal model of Alzheimer's disease, zinc deficiency worsened cognitive decline due
to an enhancement in NLRP3-driven inflammation. Overall, our data suggest that
zinc status affects Alzheimer's disease progression, and that zinc supplementation
could slow the rate of cognitive decline.

Introduction

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The global prevalence of dementia (of which the most common type is Alzheimer's disease), is growing with 152 million people predicted to be living with dementia by 2050 (from 50 million in 2018), and there are no effective treatments (Alzheimers, 2015). Around 92% of people living with dementia also suffer from one or more longterm diseases (co-morbidities) (Browne et al., 2017). It has been established that disease-associated inflammation increases the risk of developing dementia, and includes inflammation associated with several co-morbidities for Alzheimer's disease such as infection, diabetes, atherosclerosis and obesity (Whitmer et al., 2005; Profenno et al., 2010). Zinc is an essential trace element obtained from the diet that regulates the expression and activation of many biological molecules. Maintenance of adequate zinc levels through a balanced diet is therefore important to health. Zinc deficiency affects up to two billion people world-wide and has profound effects on immune system function (Prasad, 2012). Given its prevalence, therefore, it is likely that zinc deficiency represents a significant co-morbidity for Alzheimer's disease that could contribute to poorer quality of life and disease outcome. Previous associations between zinc deficiency and Alzheimer's disease include several studies reporting reduced plasma zinc levels in people with Alzheimer's disease (Ventriglia et al., 2015). Overall, these data suggest that people with Alzheimer's disease are at risk of zinc deficiency, though the potential effect zinc deficiency is having on the condition is currently debated.

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Zinc is a critical micronutrient that plays an essential role in a vast number of physiological processes. Approximately 10% of proteins have zinc binding domains, zinc also plays a role in membrane stabilization and can act directly as a signalling molecule (Sensi et al., 2009; Kambe et al., 2015). Because of its physiological

importance, its cytosolic and extracellular concentrations are tightly controlled by several mechanisms including two main families of transporters, the zinc transporters (ZnT) and the Zrt-Irt-related proteins (ZIP), as well as zinc storage proteins (e.g. metallothioneins) and vesical storage of free zinc (Sensi et al., 2009; Kambe et al., 2015). Typically, ZIP transporters act to increase cytosolic concentrations of zinc, while ZnT act to decrease cytosolic concentrations by movement of zinc out of the cell or into cellular compartments including vesicles, Golgi and endosomes (Sensi et al., 2009; Kambe et al., 2015). The ubiquity and diversity of molecular functions which involve zinc has made researching the interactions between zinc status and Alzheimer's disease difficult.

Several potential pathophysiological links of Alzheimer's disease and zinc status have been investigated. The role zinc binding has in amyloid fibril formation has been a substantial area of interest. Several *in vitro* and *in vivo* studies report that excessive zinc promotes the formation of amyloid oligomers and fibrils through direct zinc-amyloid interactions. Amyloid oligomers and fibrils are one of the pathological hallmarks of Alzheimer's disease and considered a likely driver of the disease (Esler et al., 1996; Brown et al., 1997; Lee et al., 2018). These findings have driven preliminary Alzheimer's clinical trials on compounds, such as PBT2, which increase cellular uptake of zinc thereby reducing extracellular zinc levels. These trials have had largely positive results (Faux et al., 2010).

Another focus of the Alzheimer's disease zinc field is the role of zinc proteases in beneficial and pathological processes. Zinc is an essential co-factor in a range of proteases including proteases that cleave the amyloid precursor protein into nonpathogenic peptides including the a-disintegrin and metalloproteinases (ADAMs) (Gough et al., 2011). Zinc is also a co-factor in proteases that digest the pathological forms of amyloid into non-oligomerising products, including the insulin degrading enzyme (Gough et al., 2011). Finally, the zinc protease matrix metallopeptidase 9 (MMP9) plays a critical role in the healthy functioning of central nervous system, particularly the activation of brain derived neurotrophic factor (BDNF) (Yang et al., 2014; Frazzini et al., 2018). BDNF is an important neurotrophin that plays a role in neuronal plasticity and synapse formation. Previously, the administration of zinc chelators in mice has been shown to lower active BDNF levels impairing cognitive performance (Yang et al., 2014; Frazzini et al., 2018). While zinc and Alzheimer's disease has been an active field of research, little attention has been given to the known effects of zinc status on immune function; particularly neuroinflammation, an established driver of Alzheimer's disease pathology.

We previously reported that zinc depletion of macrophages activates an inflammatory complex called the NLRP3 (NACHT, LRR and PYD domains-containing protein 3) inflammasome driving release of the pro-inflammatory cytokine interleukin-1 β (IL-1 β) from the cell (Summersgill et al., 2014). The NLRP3 inflammasome is formed in macrophages, and microglia in the brain, in response to pathogenic or damage associated stress and drives an inflammatory response. The NLRP3 inflammasome complex is composed of NLRP3, the adaptor protein ASC (apoptosis-associated speck-like protein containing a caspase activation and recruitment domain (CARD)), and the protease caspase-1 which is responsible for cleaving the inactive precursor pro-IL-1 β to an active secreted IL-1 β molecule (Brough et al., 2017; Heneka et al., 2018). The NLRP3 inflammasome is suggested to contribute to the damaging inflammation that worsens Alzheimer's disease (Heneka et al., 2013; Daniels et al., 2016; Dempsey et al., 2017; White et al., 2017;

Heneka et al., 2018). NLRP3 is also suggested to contribute to age-related cognitive decline (Yourn et al., 2013).

This study aimed to test the hypothesis that zinc status can influence the development of Alzheimer's disease. We report here that zinc supplementation in people reduced the prevalence and symptomatic decline of Alzheimer's disease, and that zinc deficiency in an Alzheimer's disease mouse model accelerated cognitive decline through potentiation of NLRP3-dependent inflammation, an effect which was reversible. These data suggest that zinc deficiency is a treatable co-morbidity, which if managed correctly, potentially through the use of zinc supplementation, could slow the rate of cognitive decline in people with Alzheimer's disease.

Materials and Methods

219 Epidemiological study

220 Data acquisition

Data used in the preparation of this article were obtained from the Alzheimer's Disease Neuroimaging Initiative (ADNI) database (adni.loni.usc.edu). The ADNI was launched in 2003 as a public-private partnership, led by Principal Investigator Michael W. Weiner, MD. The primary goal of ADNI has been to test whether serial magnetic resonance imaging (MRI), positron emission tomography (PET), other biological markers, and clinical and neuropsychological assessment can be combined to measure the progression of mild cognitive impairment (MCI) and early Alzheimer's disease. The subjects were recruited from over 50 sites across the U.S.A and Canada. ADNI has undergone three stages of recruitment each with differences in the imaging and biomarker analyses, these have been named ADNI-1, ADNI-GO and ADNI-2. Collectively these protocols have recruited 1631 adults into

the study consisting of age appropriate cognitively normal individuals, people with early or late MCI, and people with early Alzheimer's disease. The follow up duration of each group is specified in the protocols for ADNI-1, ADNI-2 and ADNI-GO with 120 months as the maximum. Subjects were evaluated upon entry into the study, then at the 6 and 12 month time points, and yearly after this. For up-to-date information, see www.adni-info.org.

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Epidemiological analyses

Epidemiological analyses were performed as described in Rivers-Auty et al. (2020). Briefly, datasets containing the medical history (RECMHIST.csv), recurrent medicines (RECCMEDS.csv) and patient summary data (ADNIMERGE.csv) were downloaded on the 3rd of May 2018. String search methods were applied to identify patients using supplements. Baseline statistics were extracted, and logistic regression was performed on Alzheimer's disease prevalence to evaluate the association between supplement use and Alzheimer's disease risk. Negative binomial generalised linear mixed modelling (GLMM) was applied to the number of failures which occurred during the mini-mental state examination (MMSE). Variables were included in the models both as a fixed effect and as an interaction effect with time. Variables included in the analyses were diagnosis (control, MCI or Alzheimer's disease), age, gender, APOE4 genotype, education level, headaches, arthritis, diabetes, smoking, cardiovascular risk factors, and supplement use. The significance of variable inclusion for the logistic regression and GLMM was evaluated using the log likelihood ratio test, estimated by a Chi-squared distribution. To construct the negative binomial GLMM models the package glmmadmb was used (Fournier et al., 2012; Skaug et al., 2013), logistic regression utilized the LME4 package (Bates et al., 2015), both on R version 3.5.1 (R Core Team 2018) with RStudio version 1.1.453 (RStudio Team 2016). Details of the full analysis can be found in the Extended Data.

Animal experiments

Animals were housed in individually ventilated cages maintained at 21 (± 2) ⁰C with 40-50% humidity on a standard 12h light/dark cycle and free access to water and food. Prior to experiment all animals were on standard rodent chow (BK001 E, Special Diets Services, UK). Genotyping was performed by Transnetyx[™] and these results were not revealed until the end of all behavioural, histological or physiological investigations had been completed. Thus, all experiments were performed blinded to genotype. All behavioural videos, physiological analyses and histological quantifications were recoded to blind the experimenter to both genotype and diet. Diet was randomly allocated by cage using the TrueRandom[™] software. All behavioural experiments were performed with 40db white noise played in surround sound to avoid auditory cues. All animal experiments were carried out in accordance with the United Kingdom Animals (Scientific Procedures) Act 1986 and approved by the Home Office and the local Animal Ethical Review Group, University of Manchester.

The effect of zinc deficiency in APP/PS1 mice

APPswe PSEN1∆E9 (APP/PS1) transgenic mice on a C57BL/6J background were obtained from the Jackson Laboratory™ (#005864). Hemizygous male APP/PS1 mice were bred with C57BL/6J female mice to generate litters of hemizygous APP/PS1 and C57BL/6J (wild-type; WT) littermate controls, of which the males were used in the experiments. At three months of age the animals underwent baseline behavioural testing and were then randomly allocated to a zinc normal (ZN) diet (35

mg/kg, Table 1) or zinc deficient (ZD) diet (3 mg/kg, Table 1). Two thirds of the mice were initially placed on a ZD diet and one third was placed on a ZN diet. After three months on diet another set of behavioural tests were performed. Then half the cages on a ZD diet were randomly allocated to be placed on the ZN diet. All mice remained on diet for another three months, which was followed by another set of behavioural experiments after which mice were sacrificed by exsanguination and cardiac perfusion with saline under isoflurane (2-5%) anaesthesia. A power analysis was performed using the standardized effect size based on an ordinary least squared repeated measures method. A moderate partial η^2 of 0.15 was selected with an α of 0.05 and a β of 0.2. From this it was found an n of 10 was required. To account for attrition and fluctuations in genotype prevalence a target of n = 13 was selected. Final numbers (n) in each group were as follows: WT ZN = 13; WT ZD = 15; WT ZD/ZN = 12; APP/PS1 ZN = 13: APP/PS1 ZD = 11; APP/PS1 ZD/ZN = 11.

The effect of zinc deficiency in aged C57BL/6J mice

At three months (Young) or 20 months (Aged) of age C57BL/6J mice had their baseline memory performance probed utilizing the Y-maze task, before being randomly allocated to a ZD or ZN diet for three months (n = 15 per group). All mice were then evaluated in the Y-maze and culled by exsanguination and cardiac perfusion with saline under isoflurane (2-5%) anaesthesia. Using the same power analysis as above an n of 15 was selected to account for the greater potential of attrition.

The effect of zinc deficiency in APP/PS1 mice deficient in NLRP3

Mice lacking NLRP3 (NLRP3^{-/-}) on a C57BL/6J background mice were provided by

Dr. Vishva Dixit (Genentech) (Mariathasan et al., 2006). These mice were

backcrossed with the APP/PS1 line for 6 generations and then bred into two colonies containing APP/PS1/NLRP3-/- hemizygous male and WT/NLRP3-/- (WT are C57BL/6J) female breeders, and APP/PS1/NLRP3+++ hemizygous male and WT/NLRP3^{+/+} female breeders. These were used to generate the experimental male mice used in this study. At three months of age, cages of mice were randomly allocated onto either a ZD or ZN diet. After six months on diet, memory of all mice was assessed with the Y-maze task. The animals were then sacrificed by exsanguination and cardiac perfusion with saline under isoflurane (2-5%) anaesthesia. The primary outcome of Y-maze performance after 6 months on diet was selected. Using the data obtained in the zinc deficiency experiment above, the mean difference of 13.4 and a pooled standard deviation of 7.6 was used with an α of 0.05 and a β of 0.2. This analysis found an n of 9 mice per group was required for the Sidak corrected post-hoc analysis with 6 comparisons. To account for attrition and fluctuations in genotype prevalence a target n = 10 was selected. Final numbers (n) in each group were as follows: WT/NLRP3 $^{+/+}$ ZN = 12; WT/NLRP3 $^{+/+}$ ZD = 14; $WT/NLRP3^{-/-}$ ZN = 8; $WT/NLRP3^{-/-}$ ZD = 9; $APP/PS1/NLRP3^{+/+}$ ZN = 8: $APP/PS1/NLRP3^{+/+}$ ZD = 11; $APP/PS1/NLRP3^{-/-}$ ZN = 13; $APP/PS1/NLRP3^{-/-}$ ZD = 9.

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Y-maze task

The Y-maze consisted of three connected small rooms each with a unique visual cue on the back wall. Mice were placed in the maze using the tube transfer method and allowed to explore for 8 min. The exploration was recorded, and the videos were then recoded and scored by an observer blinded to the experimental groups. A successful set of alternations was defined as entering all three rooms in succession with no repeated room entries. Performance was expressed as the percentage of

total alternation sets which were successful. The mazes were thoroughly cleaned with 70% ethanol and then dried with paper towels between animals to avoid olfactory cues. Trials in which the mouse completed less than 9 entries were excluded.

Novel smell task

The novel smell task was performed in a cylindrical arena. The mice were first habituated to the arena without olfactory cues by being placed in the arena using the tube transfer method for 8 min. The following day the mice were placed in the arena for 8 min with two olfactory cue dispensers with identical smells. They were then placed in holding cages for 8 min, while the arenas were cleaned with ethanol and dried, and the olfactory dispensers where changed to consist of a novel and familiar smell. The mice were then placed back in the area and recorded for 4 min. The exploration was recorded, the videos were then recoded and timed using the Novel Object Timer program (https://jackauty.com/program/) by an observer blinded to the experimental groups as described in Daniels et al. (Daniels et al., 2016). Trials in which the mouse completed less than 4 sec of exploration were excluded.

Tissue processing

Mice were terminally anaesthetised with 2-5% isoflurane (30% O_2 , 70% N_2O). Plasma was collected via cardiac puncture using citrate-treated needles, followed by trans-cardiac perfusion with 0.9% saline. Femurs were taken for ICP-Mass spectrometry of the bone and one epididymal fat pad was removed and weighed. Brains were dissected into hemispheres. The left hippocampus was extracted and

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snap frozen for RNA-sequencing. RNA was extracted using a Qiagen mini-kit as per instructions and tested for purity and concentration using the nano-drop method and analysed for purity and digestion with a TAPE station. All samples had RNA integrity numbers (RIN) of over 8.5. RNAseq was performed as described in Hoyle et al. (2018) on an Illumina HiSeq4000 instrument. The right hemisphere was immersefixed in 4% paraformaldehyde for 24 h and cryoprotected in 30% sucrose before being snap frozen and sectioned (15 µm) coronally every 90 µm from the rostral end of the cortex (for experiment in Figs 2 and 3A-C), or immerse-fixed in 4% paraformaldehyde for 24 h then dehydrated, paraffin embedded, and sectioned (5 μm) every 100 μm from the central sulcus (for experiment in Fig 5) and mounted onto Superfrost™ Plus slides (VWR). Six sections evenly spaced throughout the cortex were selected for immunohistochemistry. Paraffin sections deparaffinised with xylene emersion and rehydrated in descending concentrations of ethanol. For all sections, antigen retrieval consisted of 30 min immersion in 0.2 mM citrate buffer pH 6 at 96°C followed by 10 min in 90% formic acid. Sections were washed (3x 5 min) with 0.1% tween in phosphate buffered saline (PBST), blocked for 1 h in a 1% bovine serum albumin (BSA, A9647 Sigma-Aldrich), 0.2 M phosphate buffer (PB) solution and then incubated overnight in 1:200 biotinylated 6e10 antibody (for amyloid beta (Aβ) plaques; #SIG-39340-200, Covance) or 1:1000 lba1, (for microglia; Wako Ltd) in a 1% BSA, 0.2 M PB solution at 40C. Sections were then washed (3x 5 min, PBST), incubated with biotinylated anti-mouse secondary (lba1), or/followed by, 1:20 Strep-Avidin (P188503, RnD)/1% BSA, 0.2 M PB solution for 2 h at room temperature, washed and then visualised with a DAB-nickel solution (D0426-50SET, Sigma-Aldrich). Sections were dehydrated in serial ethanol solutions of increasing concentration and then 100% xylene, and then mounted with DPX (DI5319/05, Fisher Scientific). The slides were scanned, and the images recoded. Aβ

plaque burden, approximated as percentage stained area, was calculated utilizing threshold-particle analyses performed on ImageJ. The microglial activation stage was performed as in Daniels et al. (Daniels et al., 2016).

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Laser microdissection

Eight to nine-month-old APP/PS1 and C57B/6J littermate mice were terminally anaesthetised with 2-5% isoflurane (30% O₂, 70% N₂O) followed by trans-cardiac perfusion with 0.9% saline. Brains were harvested, snap frozen using dry ice and stored at -80 °C until analysed. Coronal 30 µm brain sections were taken from the hippocampal region utilising Leica CM3050 S cryostat and mounted onto RNase-free membrane-coated microscopy slides (Molecular Machines & Industries). The slides were stained by immersion in a filtered Congo red solution (0.2% Congo red solution in 20% NaCl. 80% ethanol, pH 9.0) for 20 min at room temperature. Using the laser microdissection (LMD) MMI (Molecular Machines & Industries) SmartCut software, UV laser excised discs of a fixed 100 µm diameter with Congo Red stained amyloidplagues in the centre were cut (Fig. 3H). Plague-free tissue from both the same animals and WT littermate controls were also cut and collected separately. Once cut, microdissected regions were acquired using mechanically operated adhesive MMI Isolation Caps (Molecular Machines & Industries), which were subsequently filled with the lysis buffer component of RNeasy Micro Kit (Qiagen). On completion of LMD, sample-containing microcentrifuge tubes immediately underwent lysis via freeze-thaw cycling and brief (< 5 sec) sonication. Purified RNA was acquired as per RNeasy Micro Kit (Qiagen) instructions with on-column DNase digestion and washing according to manufacturer's guidelines. Images were collected on an Olympus IX83 inverted microscope modified for LMD capture.

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ICP-Mass spectrometry

Inductively coupled plasma mass spectrometry (ICP-MS) (Agilent 7900) was used to assess zinc levels in the plasma and bone. The diets, bedding, drinking water and cage materials were also measured for zinc content prior to the start of the experiments. As a result of these analyses, the bedding was changed to the low zinc nesting material Wood Wool (TAPVEI®). Plasma samples were diluted in 1:10 with 0.1 M hydrochloric acid, centrifuged (2500 g) and then analysed with appropriate standards generated. Whole femurs were taken from the mice and wet-ashed by HNO₃ (0.5 ml, concentrated) immersion at room temperature for 24 h, then dried at 60 °C. After cooling, 0.2 ml of H₂O₂ was added and the samples were incubated for 1 h in a drying oven at 60 °C. The solutions were diluted to 10 ml using deionised water and analysed.

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Quantitative Reverse Transcription PCR

Isolated RNA was reverse-transcribed using Tetro cDNA Synthesis Kit (Bioline), with 427 samples being incubated at 25 °C for 10 min, 45 °C for 30 min and 85 °C for 5 min. 428 429 cDNA was subjected to qPCR using the 7900HT Fast Real-Time PCR detection system (Applied Biosystems) Power SYBR® Green Master Mix (Applied 430 Biosystems). A standard amplification programme (1 x 2 min cycle at 50 °C, 1 x 10 431 min cycle at 95 °C, 50 x 15 sec cycle at 95 °C and 1 x 1 min cycle at 60 °C) was 432 433 used for all amplifications. Primers used for the amplification were as follows: 434 TREM2 (forward; TGGGGACCTCTCCACCAGTT, reverse; GTGGTGTTGAGGGCTTGG), NLRP3 (forward; GCCCAAGGAGGAAGAAGAAG, 435 436 reverse; TCCGGTTGGTGCTTAGACTT), and IL-1β (forward; 437 AACCTGCTGGTGTGACGTTC, reverse; CAGCACGAGGCTTTTTTGTTGT). 438 Standard curves of serial control cDNA dilutions were used to determine the fold differences in target cDNA expression. HMBS was used as the housekeeping gene (forward; GAAATCATTGCTATGTCCACCA, reverse;

441 GCGTTTTCTAGCTCCTTGGTAA).

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Cell isolation and culture

444 **BMDM**

All primary cells were harvested from 8-16-week-old, mixed sex, C57/BL6J mice (Charles River, UK) which were euthanised with rising CO₂ concentration. Murine bone-marrow derived macrophages (BMDMs) were generated as reported previously in Daniels et al. (Daniels et al., 2016). Briefly, BMDMs were collected by flushing bone marrow from femurs and cultured in Dulbecco's Modified Eagle's Medium (DMEM) (Gibco) supplemented with 10% foetal bovine serum (FBS) (Biowest), 1% PenStrep, 2 mM, 1% L-glutamine (Sigma) and 20% M-CSF (macrophage colony-stimulating factor)-conditioned medium from L929 cells. Cells were grown for 7 days and then scraped, counted and seeded for 24 h in 24 to 96 well plates at 1 x 10⁶ cells/ml in complete DMEM. After media change, cells were primed with bacterial endotoxin (lipopolysaccharide (LPS), 1 µg/ml) from Escherichia coli O26:B6, for 4 h. The media was changed to serum free DMEM prior to treatment with pathway inhibitors (MCC950 10 μM or ZnCl₂) or vehicle (DMSO) for 15 min followed by stimuli treatment for 4 h with TPEN (N,N,N',N'-tetrakis(2-pyridinylmethyl)-1,2-ethanediamine), or silica (<15 µm, MIN-U-SIL 15; 300 µg/ml), or 24 h for amyloid 1-42 (Sigma, oligomerised at 37°C for 24 h in PBS prior to treatment. N.B for amyloid, cells without LPS-priming were used). Supernatants were taken for enzyme-linked immunosorbent assay (IL-1β) (ELISA) and western blot. An in-well lysis was performed with 1% Triton for western blot analyses of cytosolic and secreted proteins.

Mixed glia

Murine mixed glia cells were prepared from the brains of 2–4-day-old C57BL/6J mixed sex mice. Briefly, the cerebral hemispheres were dissected, and the meninges were then removed. The remaining tissue was homogenized in DMEM containing 10% FBS and PenStrep via repeated trituration. The resulting homogenate was centrifuged at 500 x g for 10 min, and the pellet was resuspended in fresh culture medium, seeded directly into 24-well plates and incubated at 37 °C, 90% humidity, and 5% CO₂. After 5 days, the cells were washed, and fresh medium was placed on the cells. The medium was then replaced every 2 days. Cells were used after two weeks. Cells were primed with LPS (1 μ g/ml, 3 h) in DMEM containing 10% FBS and PenStrep. For zinc chelation experiments, cells were washed once with serum-free DMEM prior to treatment with serum-free DMEM containing vehicle (DMSO) or TPEN (20 μ M, 4 h) with or without silica (300 μ g/ml) or MCC950 (10 μ M). For the amyloid experiment LPS-primed cells were made zinc deficient (TPEN, 10 μ M) and incubated in amyloid oligomers (10 μ g/ml, 4 h). The supernatant was centrifuged at 12,000 x g for 10 min prior to analysis for IL-1 μ content by ELISA.

ELISA and Western blot

IL-1 β release was quantified by ELISA (DuoSet, R&D Systems) according to the manufacturer's instructions. IL-1 β , gasdermin D and caspase-1 cleavage was determined by western blot. All supernatants or in-well lysates were separated by 12 well, 1.5 mm Tris-glycine SDS/PAGE and then transferred onto nitrocellulose or PVDF membranes at 25 V using a semidry Trans-Blot Turbo system (Bio-Rad). Membrane blocking was performed using in 2% bovine serum albumin (BSA) in 1% (vol/vol) Tween 20/PBS (PBST). Membranes were then incubated with primary

antibodies at 4 °C overnight (IL-1β AF-401,R&D; gasdermin D ab209845, Abcam; and caspase-1 EPR16883, abcam), washed and then labelled with HRP-conjugated secondary antibodies (2 h at room temperature). Target proteins were then visualized with Amersham ECL detection reagent (GE Healthcare) using the G:Box Chemi XX6 (Syngene) system.

ASC Oligomerization Assay.

Cells were lysed in-well with 1% Triton X-100 and protease inhibitor cocktail. Cell lysates were centrifuged at 6,800 x g for 20 min at 4 °C to separate the Triton X-100 soluble and insoluble fraction. The Triton X-100 insoluble fraction was chemically crosslinked with 2 mM disuccinimidyl suberate (DSS) (Thermo Fisher) for 30 min at room temperature. Crosslinked pellets were precipitated by centrifugation at 6,800 x g for 20 min and resuspended in Laemmli buffer. Oligomerised ASC were then detected by western blot.

ASC Speck Formation.

Adult microglia were extracted from 8-16-week-old, mixed sex, C57/BL6J mice expressing ASC-citrine on a CAG promoter using MACS® MicroBeads on digested brains as per manufacturer's instructions. Microglia were seeded for 2 h into 96-well plates at 0.5×10^5 /ml. They were primed with LPS (1 µg/ml, 2 h). All cells were treated with the pan-caspase inhibitor Z-VAD-FMK (50 µM) prior to simulation to prevent caspase dependent cell death. Without media change, NBC19 (10 µM), MCC950 (10 µM) or vehicle (DMSO) were spiked into the appropriate wells, followed by TPEN administration (final concentration 10 µM). At 15-min intervals images were captured using a 20×/0.61 S Plan Fluor objective for a total of 12 h. Speck formation

was quantified using an IncuCyte ZOOM System (Essen Bioscience). All experiments performed in serum free OptiMEM.

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Experimental statistics

All data are presented as mean values ± standard error of the mean (SEM) unless stated. For repeated measures analyses, linear mixed modelling was used to evaluate the significance of the effect of independent factors (genotype, diet and time) on the dependent variable. All factors and interactions were modelled as fixed effects and mouse identifier as a random variable. A within-subject design with random intercepts was used for all models. The significance of inclusion of an independent variable or interaction terms was evaluated using log-likelihood ratio (Bates et al., 2015). Holm-Sidak post hocs were then performed for planned pairwise comparisons using approximated least square means (Lenth, 2016). For the remaining data, statistical analyses were unpaired Welch corrected Student's t test, or one-way, two-way or three-way analysis of variance (ANOVA) test depending on the number of experimental groups and independent variables in the analyses. Holm-Sidak post hocs were then performed for planned pair-wise comparisons. Homoscedasticity and normality were evaluated graphically using predicted verses (Pearson) residuals and Q-Q plots, respectively, and transformations and/or corrections were applied where necessary. All analyses were performed using R (version 3.3.3) (R Core Team; RStudio Team.). Accepted levels of significance were *p < 0.05, **p < 0.01, and ***p < 0.001.

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Results

Zinc supplementation use is associated with reduced prevalence of Alzheimer's disease and slower cognitive decline during disease development.

The ADNI dataset was used to investigate the effects of mineral supplementation on both the prevalence of cognitive dysfunction including early and late MCI, and Alzheimer's disease, as well as the progression of cognitive decline over time. Investigating the six most common mineral supplements (calcium, iron, magnesium, multi-vitamin, selenium, zinc), and utilizing logistic regression adjusting for common confounding variables, we found that calcium, iron, magnesium and zinc supplement use was associated with significantly reduced Alzheimer's disease prevalence compared to those not taking any supplements (Fig. 1*A*,*B*). To further investigate the potential effects of mineral supplementation, innovative statistical methods for analysing cognitive decline, as measured by the MMSE, were employed. From this it was found that zinc supplementation was associated with significantly less cognitive decline (Fig. 1*C*,*E*). Other supplements, including the divalent cations calcium and magnesium, were not associated with significant changes in cognitive decline (Fig. 1*D*).

Subclinical zinc deficiency accelerates memory deficits in an animal model of

Alzheimer's disease without effecting plasma zinc or weight gain

We then sought to establish that the association between zinc supplementation and improved Alzheimer's disease outcome was causal by using the APP/PS1 mouse model of Alzheimer's disease. WT control, or APP/PS1 mice were fed a diet containing the recommended zinc intake levels for mice (35mg/kg, zinc normal, ZN) or a diet that would induce a mild subclinical zinc deficiency (3mg/kg, zinc deficient, ZD) (Fig. 2A). Animals were put on diet from three months of age and maintained for six months with memory tests (Y-maze and novel smell test) performed at three-month intervals (Fig. 2A). To determine if any effects of zinc deficiency were reversible, one group of mice was maintained on a zinc deficient diet for three

months and then placed on a normal zinc diet for three months (Fig. 2A). APP/PS1 mice do not show altered cognitive behaviour in short term memory tasks such as the Y-maze, T-maze or novel object recognition before nine months of age (Webster et al., 2014). At the end of the experimental period animals were sacrificed zinc levels measured in the plasma and bone, and a number of histological and physiological measures taken including RNAseq on hippocampal tissue, and analyses on Aß plague burden and microglia morphology. There was no change in plasma zinc between all groups (diet main effect: F(2,68)=0.86, p=0.43) but a decrease in zinc content in the bone was seen in WT and APP/PS1 mice fed a zinc deficient diet indicating a change in long-term zinc status (Fig. 2B,C), (diet main effect: F(2,64)=4.26, p=0.018). Severe zinc deficiency can lead to weight loss, but the marginal zinc deficient diet used here had no effect on body weight compared to normal diet (Fig. 2D), (diet main effect: F(2,43)=0.49, p=0.616) or fat mass (data not shown, diet main effect: F(2,74)=0.14, p=0.866) of the WT or APP/PS1 mice. These data suggest that we had modelled a sub-clinical level of zinc deficiency (Beattie et al., 2012).

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The mild zinc deficiency cause reversible cognitive decline in the APP/PS1 mice while having no effect on the cognitive performance of the WT mice (Y-maze task 3-way interaction: $\chi^2(6)$ =19.33, p=0.004). In WT mice there was no effect of zinc deficient diet on memory over 6 months as analysed by the Y-maze (Z=-0.16, p=1.000) and novel smell tests (Z=-0.16, p=0.743), (Fig. 2E,G). As expected, the APP/PS1 mice showed no significant reduction in cognitive performance over the course of the experiment on the zinc normal diet in the Y-maze (Z=-1.81, p=0.281) or novel smell tests (Z=-0.05, p=1.000), (Fig. 2E,H). However, APP/PS1 mice on a zinc deficient diet showed accelerated memory loss at three months in the Y-maze test

(Z=-2.81, p=0.024), and this was worse after six months on the zinc deficient diet (Z=-3.75, p=0.001), (Fig. 2F). This cognitive impairment measured by Y-maze of APP/PS1 mice fed a zinc deficient diet for three months was reversed by returning the mice to a zinc normal diet for three months (Z=-3.93, p<0.001), (Fig. 2F). Similar, but less dramatic, trends were observed in the novel smell test in response to a zinc deficient diet in the APP/PS1 mice (Fig. 2H), (3-way interaction: χ^2 (6)=14.53, p=0.024). These data suggest that zinc deficiency is causally accelerating cognitive decline in Alzheimer's disease and that these cognitive deficits are reversible. They also place our epidemiological observations in people in context, suggesting the association of improved cognitive outcome in patients taking zinc supplements is due to a direct causal relationship between zinc status and Alzheimer's disease progression.

Zinc deficiency does not alter plaque burden, microglia activation or agerelated cognitive decline

The effects of zinc deficiency on memory were not because of altered A β plaque burden as plaque burden was not significantly different between APP/PS1 mice fed a zinc normal or a zinc deficient diet (Fig. 3A,B), (2-way interaction: F(3,61)=1.93, p=0.154). Similarly, levels of microglia with activated morphology, typically associated with plaques, were not altered between treatment groups (Fig. 3C), (2-way interaction: F(3,49)=0.06, p=0.940). We then hypothesised that the effects of zinc deficiency may be common with NLRP3-dependent age-related cognitive decline (Youm et al., 2013). Thus, we placed 20-month-old C57BL/6J mice on a zinc deficient diet for three months, with 3-month-old mice as young controls. No acceleration in cognitive decline after a zinc deficient diet was observed at either age in the Y-maze task (3-way interaction: $\chi^2(7)$ =7.34, p=0.394) or Morris water maze

task (interaction term: F(1)=0.284, p =0.570), indicating the effect of zinc deficiency
seen in the APP/PS1 mice was specific to amyloidopathy and its downstream effects
(Fig. $3D$, E). To elucidate the mechanism of zinc deficiency-induced cognitive decline
in APP/PS1 mice, RNAseq was performed on the whole hippocampal homogenate.
However, no substantial differences in transcriptome were observed between
genotype or diet (Fig. $3F$, G). We hypothesised therefore that transcriptional
differences could be occurring primarily in the periplaque regions, which were
masked in the hippocampal RNAseq data (Fig. $3F$, G) by homogenising the whole
tissue. To investigate this potential locality of transcriptional changes, we utilised
laser capture microdissection to probe gene expression in the periplaque regions of
8-9-month-old APP/PS1 mice (Fig. $3H$). Performing qPCR on the established the
Alzheimer's disease linked gene TREM2 and the inflammatory genes for NLRP3 and
IL-1 β , we confirmed that transcriptional changes were largely confined to the
periplaque regions (Fig. 3/), (periplaque main-effect: TREM2 $\chi^2(1)$ =28.40, p <0.001;
IL-1 $\chi^2(1)$ =32.21, p <0.001; NLRP3 $\chi^2(1)$ =13.93, p <0.001). Given that NLRP3
activation has been strongly linked to memory impairment in Alzheimer's disease
(Heneka et al., 2013; Daniels et al., 2016) and that NLRP3 is activated by zinc
deficiency (Summersgill et al., 2014), we hypothesised that accelerated memory
impairment in the APP/PS1 mice on the zinc deficient diet is due to a periplaque
NLRP3 response (Heneka et al., 2013; Daniels et al., 2016; Dempsey et al., 2017;
Venegas et al., 2017; White et al., 2017).

Zinc deficiency activates the NLRP3 inflammasome and potentiates NLRP3 responses to stimuli, while zinc supplementation inhibits NLRP3 activation

To investigate the potential regulation of NLRP3 by zinc status *in vitro* methods were

employed. We previously established that primary BMDMs have similar NLRP3

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responses as microglia, mixed glial cultures and ex vivo brain tissue slices, therefore, represent a robust, and high throughput method for investigating NLRP3 activation (Daniels et al., 2016; Hoyle et al., 2020). BMDMs were primed with LPS and then treated with the zinc chelator TPEN (1-30 µM, 4 h) to cause zinc depletion. Analysis of the supernatant showed TPEN-induced zinc depletion induced-1ß release in a concentration dependent manner (F(4,20)=25.28, p<0.001), (Fig. 4A). Zinc depletion (TPEN 10 µM, 4 h) in LPS-primed BMDMs also caused oligomerisation of ASC, caspase-1 activation, and gasdermin-D cleavage, and this was NLRP3 dependent as it was inhibited by the selective NLRP3 inhibitor MCC950 (Fig. 4Bi). Conversely, zinc supplementation (with ZnCl₂) in LPS-primed BMDMs blocked established NLRP3 activators, including silica, by inhibiting IL-1β release (F(5,12)=122.0, p<0.001), ASC oligomerisation, caspase-1 and gasdermin D cleavage (Fig. 4Bii, C). We then investigated potential interactions between Aβ and zinc depletion. Utilizing BMDMs we found that only Aβ42 oligomer treatment followed by TPEN was sufficient to induce IL-1β secretion, which was inhibited by MCC950 (Fig. 4D). The effects of TPEN were then confirmed in LPS-primed mixed glial cultures, and again TPEN treatment induced IL-1β release which was inhibited by MCC950 (F(5,12)=20.18, p<0.001), (Fig. 4E). Using LPS-primed mixed glia we also found that TPEN was also able to exacerbate Aβ42 oligomer-induced IL-1β secretion (F(3,8)=8.06, p=0.008), (Fig. 4F). To investigate if TPEN-induced zinc depletion caused inflammasome specks in microglia, ASC-citrine expressing microglia were isolated from adult mice (Tzeng et al., 2016). Zinc depletion resulted in speck formation which was observed with live cell imaging and the NLRP3 inhibitors MCC950 (Coll et al., 2015) and NBC19 (Baldwin et al., 2017) were found to inhibit ASC speck formation (Fig. 4G,H), (MCC950 Z=-6.34, p<0.001; NBC19 Z= -26.80, p<0.001). Collectively, these experiments indicate that zinc status can influence

NLRP3 activity. Zinc deficiency acts as an NLRP3 activating and potentiating stimuli in macrophages, microglia and mixed glia, while zinc supplementation is inhibitory of NLRP3 activation. Critically, we found that the combination of zinc deficiency and amyloidopathy are sufficient to induce NLRP3 activation and IL-1β release in mixed glia (and macrophages).

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Zinc deficiency accelerates cognitive decline in the APP/PS1 mouse model of

Alzheimer's disease through an NLRP3 dependent mechanism

As we had established that zinc deficiency could potentiate NLRP3 responses in vitro we hypothesised that zinc deficiency accelerated cognitive impairment in the APP/PS1 mouse via the NLRP3 inflammasome. To test this hypothesis, we crossed WT and APP/PS1 mice with NLRP3-1- mice to generate four experimental groups of mice which were, WT, NLRP3-/-, APP/PS1, and APP/PS1/NLRP3-/-, Three-month-old animals were fed either a zinc normal or zinc deficient diet for six months before memory assessment using the Y-maze. As above, there was no effect of a zinc deficient diet on the memory of WT mice (t(72)=-0.33, p=1.000), or in NLRP3^{-/-}mice (t(72)=-0.81, p=1.000), (Fig. 5A). Also, as previously, the zinc deficient diet caused memory impairment in APP/PS1 mice (t(72)=-8.06, p<0.001), (Fig. 5A). However, the zinc deficiency-induced memory impairment in APP/PS1 mice was abolished in APP/PS1/NLRP3^{-/-} mice (t(72)=4.54 p<0.001), (Fig. 5A). These data suggest that the effects of zinc deficiency on memory depend on NLRP3. These protective effects of NLRP3 deficiency on memory after zinc deficiency were independent of plaque burden or microglia activation as there was no significant difference in these parameters in APP/PS1 mice with or without NLRP3 (3-way interaction term: plaque F(1,46)=1.15, p=0.290; microglia F(1,46)=0.49, p=0.488), (Fig. 5B-D).

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Discussion

Using patient data, animal models, and cell culture methods, we have provided evidence that zinc status is important in Alzheimer's disease progression, and specifically that zinc deficiency accelerated cognitive decline in an Alzheimer's disease model through potentiating inflammatory responses mediated by the NLRP3 inflammasome.

Our epidemiological evidence is supported by numerous previous studies which report that people with Alzheimer's disease have lower zinc serum levels (Ventriglia et al., 2015). Ventriglia et al. (2015) performed a meta-analysis resulting in a pooled sample of 1,064 people with Alzheimer's disease and 1,894 healthy controls and found that the serum or plasma zinc levels were lower in people with Alzheimer's disease with a standard mean difference of -0.39, providing initial evidence of a cause association of zinc deficiency and Alzheimer's disease. The present study takes this result further, demonstrating reduced cognitive decline over time in people with MCI and Alzheimer's disease in individuals who are taking zinc supplements. While we acknowledge that this association could be explained by an unaccounted hidden variable including life-style factors or unknown disease modifying medicines, most other mineral supplements were not associated with an improvement in cognitive decline. To probe this association and establish if zinc status had a causal effect on cognitive decline, we conducted *in vivo* experiments using animal models of amyloidopathy.

There is clinical and preclinical evidence that excessive zinc supplementation can lead to health complications by being directly toxic, inhibiting copper absorption and through an interaction with amyloidopathy mechanisms (Fosmire, 1990; Yang et al.,

2013; Flinn et al., 2014; Craven et al., 2018). There is also robust evidence for high zinc deficiency prevalence among the elderly and in particular people with Alzheimer's disease, hence, we hypothesise that the association of zinc supplementation with improved cognitive outcomes reported here was likely due to the mitigation of zinc deficiency, rather than zinc supplementation exceeding recommended daily intake values (Briefel et al., 2000; Ventriglia et al., 2015). Therefore, we designed the *in vivo* experiments to investigate the effects of mild subclinical zinc deficiency on cognitive decline in the APP/PS1 mouse model of Alzheimer's disease and addressed whether any effects could be reversed by normalising the levels of zinc. From this we found that zinc deficient APP/PS1 mice had accelerated cognitive decline, but these deficits could largely be reversed by returning the mice to a diet which contained the recommended daily intake of zinc.

The literature describing the effects of zinc on amyloid deposition is complex (Esler et al., 1996; Brown et al., 1997; Garai et al., 2007; Rezaei-Ghaleh et al., 2011; Abelein et al., 2015; Lee et al., 2018). We report no changes in plaque deposition induced through zinc deficiency and, therefore, we hypothesised zinc was modulating the response to the Alzheimer's amyloidopathy modelled by the APP/PS1 mice and not the amyloidopathy itself. Using the same animal model, Stoltenberg et al. (2007) reported a mild increase in plaque size in zinc deficient animals. However, the zinc deficiency induced by Stoltenberg et al. was more severe, resulting in a 20% reduction in serum zinc levels. The present study induced a mild subclinical zinc deficiency, which caused no change to serum zinc levels and was only detected through reduced zinc levels in the bone. Brain levels of zinc were not measured here due to the complication of amyloid plaques sequestering zinc.

Indeed Stoltenberg *et al.* could not detect changes in cortical zinc even with the more severe zinc deficiency likely due to this sequestration effect.

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We saw an increase in NLRP3 expression in the periplaque region in the hippocampus of the APP/PS1 mice. This observation and previous research supports an importance of NLRP3 in the response to amyloidopathy (Heneka et al., 2013; Daniels et al., 2016). Using cellular models, we demonstrated that zinc deficiency potentiated NLRP3 responses to stimuli including amyloid oligomers and that zinc inhibited NLRP3 responses to activators such as silica. The concentration of amyloid used was supraphysiological in order to observe measurable inflammatory responses in the short time periods possible with the culture methods utilised. This limitation was addressed by translating these results in vivo. We established NLRP3 as the critical mechanism by which zinc deficiency accelerated cognitive decline in the APP/PS1 mouse model through crossing NLRP3-1- and APP/PS1 mouse lines. We demonstrated that the cognitive decline induced by zinc deficiency was completely abated in the APP/PS1 mice deficient in NLRP3. Interestingly, we saw no changes in amyloid deposition in the APP/PS1/ NLRP3-/mice. Heneka et al., (Heneka et al., 2013) generated the same mouse strain and did show reduced plaque burden, however, in that study 16-month-old mice were used. Therefore, it may be the later stages of the pathology in which NLRP3 plays a role in plaque deposition. Given the excellent research linking NLRP3 to the ageing process (Yourn et al., 2013), the age difference between Heneka et al. (Heneka et al., 2013) and the present study could explain this difference.

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Previous research has demonstrated that zinc chelation with clioquinol impedes the MMP9 processing of pro-BDNF, perturbing synapse formation and cognitive

performance (Frazzini et al., 2018). Furthermore, zinc supplementation in the 3xTg mouse model of Alzheimer's disease was found to prevent cognitive decline through increasing BDNF levels and mitigating tau, amyloid and mitochondrial Alzheimer's associated pathology (Corona et al., 2010). The RNAseq performed here found no effects of zinc deficiency on expression of genes downstream of BDNF signalling (e.g. neuropeptide Y). One explanation for this incongruence is the time frame and severity of zinc deficiency. Zinc chelation likely induced a more severe reduction in bioavailable zinc than what was observed in the present study. Additionally, Corona et al. (2010) administered zinc supplementation to mice for 11 months, starting at 1-month old mice. This was both a longer period than the present study and, began at an age where neurodevelopment is ongoing. So, while in the present study we see no evidence of BDNF modulation, it is likely this pathway is affected in other modalities of zinc deficiency and supplementation.

Managing Alzheimer's disease is currently one of the greatest societal problems. An increasing prevalence linked to an ageing population, with limited treatment options, is creating a global crisis. The NLRP3 inflammasome has been identified as a therapeutic target for the treatment of Alzheimer's disease and there are many pharmaceutical and academic initiatives underway to develop new small molecule inhibitors of the NLRP3 inflammasome (e.g. MCC950 and NBCs (Coll et al., 2015; Baldwin et al., 2017)). The possibility of drug repurposing has also been identified for targeting NLRP3 in Alzheimer's disease (Daniels et al., 2016). Here, we show that coupled to these initiatives, the nutritional status, particularly of zinc, could be assessed and used to treat groups of people with Alzheimer's disease shown to be at risk of zinc deficiency. In theory zinc supplementation or dietary improvements could be a cheap, rapid, low risk and easy enhancement to patient care.

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803	J Rivers-Auty: Data curation, formal analysis, investigation, methodology, project
804	administration, and writing-original draft and review and editing.
805	VS Tapia: Investigation
806	CS White: Investigation
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810	HG Spence: Investigation
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812	JP Green: Investigation
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816	AE Mather: Data curation and formal analysis
817	R Peters: Data curation and formal analysis
818	T Tzeng: Generating and providing ASC-citrine mice and advising on their
819	establishment and use.
820	MJ Gordon: Investigation
821	JH Beattie: Funding acquisition, conceptualization and methodology
822	D Brough: Conceptualization, funding acquisition, methodology, project
823	administration, resources, supervision, and writing-original draft and review and
824	editing.

825	CB. Lawrence: Conceptualization, funding acquisition, methodology, project
826	administration, resources, supervision, and writing-original draft and review and
827	editing.
828	
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Figure legends

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Figure 1: Zinc supplementation use was associated with reduced prevalence of Alzheimer's disease and slowed cognitive decline during disease development. A) Table of baseline statistics by supplement showing nominal differences in nuisance variables such as age, gender, APOE4 status, and significant differences in the prevalence of cognitive dysfunction. Shown are the number of participants for each variable with percentage of total in brackets, or for numerical variables the means and standard deviations (SD) are shown. *p < 0.05, **p < 0.01, ***p < 0.001 compared to no supplement use. 'Statistics' column reports the omnibus tests. B) A stacked bar chart showing the proportions of cognitive diagnosis. Zinc supplement use was associated with the lowest proportion of Alzheimer's disease. This was significantly different to 'no supplement' use group as reported in (A). Dashed lines represent proportions of cognitive diagnoses in the 'no supplement' group. C) Summary table of the final negative binomial generalised linear mixed modelling (GLMM) with mini-mental state examination (MMSE) failures as the dependent variable. Zinc supplement use was associated with significantly slowed cognitive decline. Shown are the maximum likelihood estimates ("Estimate" columns) with Laplace estimates of the standard error ("Standard error" columns), Zvalue and p-value from the Wald approximation ("p-value" columns), as well as, the significance of inclusion of the variable in the model evaluated using the loglikelihood ratio test - chi squared ("Statistics of inclusion" columns). D) and E) The effect of supplement use on predicted cognitive decline of a LMCI, 70-year-old female as measured by MMSE scores. Plots model the effect of zinc only supplement use (E) and any supplement (combined) use (D) with no supplement use modelled as the control group in both plots. CN = cognitively normal, EMCI and

LMCI = early and late mild cognitive impairment, AD = Alzheimer's disease, ADAS =

Alzheimer's disease assessment scale, MMSE = mini-mental state examination.

Figure 2: Subclinical zinc deficiency accelerates memory deficits in an animal model of Alzheimer's disease without effecting plasma zinc or weight gain. A) Timeline of the zinc deficiency experiment in APPswe/PSδE9 (APP/PS1) and wildtype (WT) mice. B. C) ICP-Mass spectrometry revealed that six months on a zinc deficient diet (ZD) (3 mg/kg) induced sustained subclinical zinc deficiency as measured by reduced bone zinc levels (B) but not plasma zinc levels (C). D) ZD diet did not affect growth of the animals with no effect on body weight change during the course of the study. E, F) The Y-maze memory task revealed that a ZD diet induced a significant memory deficit in the APP/PS1 mice compared to mice on a zinc normal diet (ZN, 35 mg/kg). These deficits were reversed by returning these mice onto a ZN for three months (ZD/ZN group; F). G, H) The novel smell recognition (NSR) task revealed significant deficits in the APP/PS1 ZD group compared to the ZN group (H). Animals returned to the ZN diet at six months old did not return to cognitive normal performance. In WT mice, no deficits were observed in the Y-maze or the NSR tasks on any diet combination (E, G). * p < 0.05, ***p < 0.001 compared to the ZN group within the same genotype. Holm-Sidak corrected post-hoc analysis. Data are presented as mean \pm SEM. WT ZN, n = 13; WT ZD, n = 15; WT ZD/ZN, n = 12; APP/PS1 ZN, n = 13; APP/PS1 ZD, n = 11; APP/PS1 ZD/ZN, n = 11.

Figure 3: The effects of zinc deficiency alters the local response to the plaque without altering plaque burden or age-related cognitive decline. *A*) Six months on a zinc deficient (ZD, 3 mg/kg) diet did not alter amyloidβ (Aβ) plaque burden in APPswe/PSδE9 (APP/PS1) mice compared to zinc normal (ZN, 35 mg/kg) controls.

The zinc recovery group was also unchanged (ZD diet for 3 months followed by a ZN diet for three months; ZD/ZN). **B)** Example images of A β plaque burden. **C)** No differences were observed in microglia morphology between any diet combination, measured as percentage of microglia in an amoeboid phenotype. For **A-C** Sections were incubated with 6e10 (A β) or Iba1 (microglia) antibodies, visualised with DAB nickel (3,3'-diaminobenzidine, nickel sulphate) staining averaged across 6 coronal hippocampal sections per mouse, scale bars are 500µm. ***p < 0.001 main effect of genotype (2-way ANOVA). **D,E)** Unlike in the APP/PS1 mouse, zinc deficiency did not accelerate age-related cognitive decline, as measured by the Y-maze, in 20-month-old C57B/6J mice placed on a ZD diet for three months. Similarly, no changes where observed in the 3-month-old 'young' C57B/6J controls. This suggested zinc deficiency was altering an amyloid specific response such as NLRP3-dependent inflammation.

(*F*, *G*) Zinc content in the diet had no effect on the transcriptome in whole hippocampal tissue of APP/PS1 and wild-type (WT) mice. *F*) a principal component analysis (PCA) of the transcriptome data showing no clustering of the WT or APP/PS1 mice or the ZD and ZN diet. *G*) Volcano plot of genes comparing APP/PS1 mice on a ZN or ZD diet showing minimal effects of diet with only one gene found to be significantly different following false discovery rate (FDR) adjustment (n = 6/group). *H & I*) To establish if there was a local response in APP/PS1 mice, periplaque regions (100 μm diameter) and plaque free tissue from 8-9-month-old APP/PS1 mice were isolated using laser capture (*H*) and probed for the plaque-associated gene TREM2 (*I ii*) and the inflammatory genes NLRP3 (*I ii*) and IL-1β (*I iii*), and from this it was found that the NLRP3/IL-1β response is very tightly associated to the peri-plaque region (n = 3, for all groups). **p < 0.01 and ***p < 0.001 versus plaque containing region. Plaques were identified with Congo red

staining. Holm-Sidak corrected post-hoc analysis. Data are presented as mean \pm SEM. For **A-C**, WT ZN, n = 13; WT ZD, n = 15; WT ZD/ZN, n = 12; APP/PS1 ZN, n = 13; APP/PS1 ZD, n = 11; APP/PS1 ZD/ZN, n = 11. For **D,E**, Young ZN, n = 15, Young ZD, n = 15, Aged ZN, n = 11, Aged ZD, n = 12 (<15 due to attrition

Figure 4: Zinc deficiency activates the NLRP3 inflammasome and potentiates NLRP3 responses to stimuli, while zinc supplementation inhibits NLRP3 activation. A-D) Primary bone marrow derived macrophages (BMDMs) experiments demonstrating zinc deficiency activates the NLRP3 inflammasome and zinc supplementation inhibits NLRP3 inflammasome activation. A) Enhanced IL-1β release from zinc deficient macrophages (BMDM) which were primed with lipopolysaccharide (LPS, 1 µg/ml, 4 h) and then treated with increasing concentrations of the zinc chelation agent TPEN for 4 h. B) BMDMs were primed (LPS 1 µg/ml, 4 h) and treated with TPEN (10 µM) with and without NLRP3 inhibition by MCC950 (10 μ M) (i) or silica (300 μ g/ml) with and without ZnCl₂ (200 μ M) supplementation (ii); the cells were lysed in the media and probed for ASC oligomerisation, IL-1β cleavage (17kd), caspase 1 activation (p10) and gasdermin D cleavage. C) Zinc supplementation inhibits IL-1β release; BMDMs were primed (LPS 1 μg/ml, 4 h) and treated with silica (300 μg/ml) in the presence of increasing concentrations of ZnCl2. D) BMDMs treated with TPEN (10 µM) and amyloid oligomers (10 μg/ml, N.B no LPS priming) for 24 h release IL-1β which is inhibited by MCC950 treatment (10 μ M). *p < 0.05, **p < 0.01 and ***p < 0.001 versus no treatment or LPS control. ###p < 0.001 versus activated group (Amyloid+TPEN). E & F) Confirmation experiments in mixed glial cultures demonstrating that LPS-primed BMDMs and mixed glial cultures have similar NLRP3 responses to zinc deficiency. E) Zinc deficient primary mixed glia release IL-1β in an NLRP3 dependent manner;

primary murine mixed glia were primed with LPS (1 μ g/ml, 4 h) and then treated with TPEN (10 μ M) or silica (300 μ g/ml) for 4 h with and without the NLRP3 inhibitor MCC950 (10 μ M). *F*) LPS (1 μ g/ml, 4 h) primed mixed glia release more IL-1 β in response to amyloid oligomers (10 μ g/ml, 4 h) when zinc deficiency is induced (TPEN, 10 μ M). **p < 0.01 versus LPS alone control. #p < 0.05 versus activated group (LPS/silica or LPS/TPEN). *G* & *H*) Zinc deficient (TPEN 10 μ M) primary ASC-citrine expressing microglia (primed with LPS; 1 μ g/ml, 2 h) form inflammasome specks over time (*G*,*Hii*, white arrows) which is inhibited by the NLRP3 inhibitors MCC950 (10 μ M)(*Hiii*) and NBC19 (2 μ M) (*Hiv*). ***p < 0.001 versus vehicle control (LPS alone). ###p < 0.001 versus activated group (LPS/TPEN). Holm-Sidak corrected post-hoc analysis following one-way (*A*,*C*,*D*), two-way (*E*,*F*) or repeated measures (*G*) ANOVA. Data are presented as mean \pm SEM of 4 independent experiments.

Figure 5: Zinc deficiency accelerates cognitive decline in the APP/PS1 mouse model of Alzheimer's disease through an NLRP3 dependent mechanism. *A*) APP/PS1 and C57BL/6J (wild-type; WT) littermate mice were crossed with NLRP3^{-/-} mice. Three-month-old mice from each genotype combination were placed on a zinc deficient (ZD, 3 mg/kg) or zinc normal (ZN, 35 mg/kg) diet for six months and were then assessed using the Y-maze memory task. Memory deficits were seen in the APP/PS1 mice on a ZD diet but not on a ZN diet. However, APP/PS1/ NLRP3^{-/-} mice on a ZD diet did not have memory deficits. *B,D*) No effect of diet or genotype was seen on Aβ plaque burden in APP/PS1 mice, or on *C*) microglia morphology measured as percentage of microglia in an amoeboid phenotype in adjacent sections. For *B* and *C* sections were incubated with 6e10 (Aβ) and Iba1 (microglia) antibodies, respectively, visualised with DAB nickel (3,3'-diaminobenzidine, nickel

sulphate) staining averaged across 6 sagittal hippocampal sections per mouse, scale bars are 250 μ m. ***p <0.001 versus WT littermates of the same NLRP3 genotype and diet. ##p < 0.01 versus NLRP3-/- group of the same diet and APP/PS1 genotype conditions. Holm-Sidak corrected post-hoc analysis following three-way ANOVA (NLRP3 genotype, diet, APP/PS1 genotype). Data are presented as mean \pm SEM. WT/NLRP3+/- ZN, n = 12; WT/NLRP3+/- ZD, n = 14; WT/NLRP3-/- ZN, n = 8; WT/NLRP3-/- ZD, n = 9; APP/PS1/NLRP3+/- ZD, n = 8; APP/PS1/NLRP3+/- ZD, n = 9.

Extended data. The extended data contains the annotated R code and output of the epidemiological analysis of the Alzheimer's disease neuroimaging cohort study.

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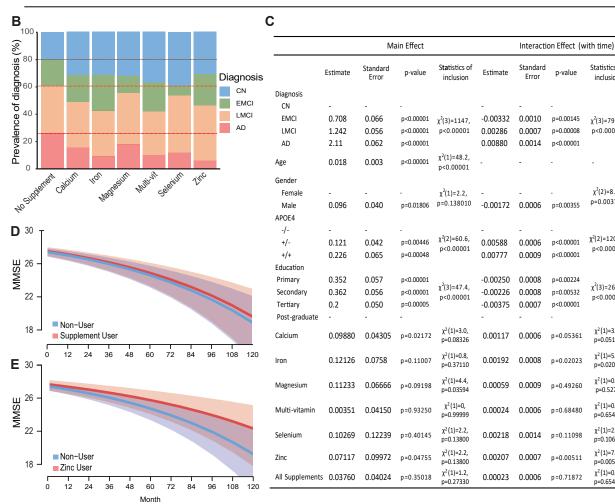
Table 1. Diet composition

D'atana and	
Diet component	g/kg
Egg white solids	212
Maltodextrin	75
Sucrose	327
Corn Starch	186.5
Corn Oil	100
Cellulose	46.236
Mineral Mix (zinc absent)	35
Calcium phosphate, dibasic	3
Chromium Potassium Sulfate, dodecahydrate	0.01
Vitamin Mix AIN 93	10
Biotin	0.004
p-Aminobenzoic Acid	0.11
Vitamin C, ascorbic acid, coated (97.5%)	1.017
Choline Bitartarate	3
Inositol	0.11
Zinc chloride (3mg/kg) (Zinc deficient)	0.0052
Zinc chloride (35mg/kg) (Zinc normal)	0.0719
Biotin p-Aminobenzoic Acid Vitamin C, ascorbic acid, coated (97.5%) Choline Bitartarate Inositol Zinc chloride (3mg/kg) (Zinc deficient)	0.004 0.11 1.017 3 0.11 0.0052 0.0719

The composition of the zinc normal (35 mg/kg) and zinc deficient (3 mg/kg) diets were identical apart from the

1023 amount of zinc.

	No supplement	Calcium	Iron	Magnesium	Multi-vitamin	Selenium	Zinc	Statistics
Diagnosis								
CN	131 (20%)	185 (31%)	35 (32%)	60 (37%)	161 (31%)	17 (40%)	20 (31%)	
EMCI	129 (20%)	116 (20%)	29 (26%)	32 (20%)	70 (13%)	3 (7%)	15 (23%)	X ² (18)=92.5
LMCI	219 (33%)	199 (33%)	37 (33%)	54 (33%)	199 (31%)	18 (42%)	26 (40%)	p<0.0001
AD	176(26%)	89 (15%)***	10 (9%)**	16 (10%)***	94 (18%) ^{ns}	5 (12%) ^{ns}	4 (6%)**	
ADAS (mean,	SD) 18.9 (10.1)	15.8 (9.1)***	14.8 (8.3) ns	14.1 (9.0)**	17.0 (9.3) ^{ns}	14.7 (7.0) ns	14.2 (6.9) ^{ns}	X ² (6)=48.7 p<0.0001
MMSE (mear	1, SD) 26.4 (2.8)	27.3 (2.6)***	27.2 (2.6) ns	28.0 (2.4)*	27.7 (2.2) ^{ns}	27.6 (2.4) ns	27.4 (2.5) ^{ns}	X ² (6)=50.0 p<0.0001
Gender								
Male	232 (35%)	385 (65%)	56 (50%)	91 (56%)	200 (38%)	14 (33%)	25 (38%)	X ² (6)=143.0
Female	423 (65%)	204 (35%)	55 (50%)	71 (44%)	324 (62%)	29 (67%)	40 (62%)	p<0.0001
Age (mean, SI	73.7 (7.5)	73.8 (7.0)**	74.3 (7.1) ^{ns}	72.9 (6.7)*	74.4 (7.1)*	74.5 (6.2) ns	73.4 (7.1) ^{ns}	F(6)=1.46 p=0.1874
APOE4								
-/-	315 (48%)	321 (54%)	68 (61%)	86 (53%)	292 (56%)	23 (55%)	37 (57%)	X ² (12)=15.3
+/-	261 (40%)	209 (35%)	32 (29%)	62 (38%)	185 (35%)	16 (36%)	25 (38%)	p=0.223
+/+	79 (12%)	59 (10%)	11 (10%)	14 (9%)	47 (9%)	4 (9%)	3 (5%)	•
Education								
Primary	223 (34%)	200 (34%)	37 (33%)	57 (35%)	193 (37%)	26 (60%)	22 (34%)	
Secondary	197 (37%)	155 (26%)	26 (23%)	48 (30%)	151 (29%)	10 (23%)	24 (37%)	X ² (18)=29.3
Tertiary	115 (15%)	136 (23%)	30 (27%)	30 (19%)	102 (20%)	4 (9%)	10 (15%)	p=0.045
Post-gradua	te 120 (14%)	98 (17%)	18 (16%)	27 (17%)	78 (17%)	3 (7%)	9 (14%)	
Headache	42 (6%)	65 (11%)	10 (9%)	14 (8%)	48 (9%)	5 (11%)	4 (6%)	X ² (6)=9.5 p=0.1495
Arthritis	213 (33%)	261 (44%)	57 (51%)	70 (43%)	226 (43%)	19 (44%)	31 (48%)	X ² (6)=29.8 P<0.0001
Diabetes	42 (4%)	37 (4%)	14 (8%)	9 (4%)	47 (6%)	2 (3%)	7 (7%)	X ² (6)=10.7 p=0.0980
Smoker	176 (27%)	140 (24%)	30 (27%)	44 (27%)	129 (24%)	11 (24%)	11 (16%)	X ² (6)=4.6 p=0.600
Cardiovascula risk factors	r 413 (63%)	346 (59%)	70 (63%)	91 (56%)	322 (61%)	25 (58%)	40 (61%)	X ² (6)=4.4 p=0.6267
Total	655	509	111	162	524	43	65	



 $\chi^2(3) = 79.8$

p<0.00001

 $\chi^2(2)=8.4$

p=0.003752

χ²(2)=120.6,

 $\chi^2(3)=26.6$

n<0.0001

p=0.02014

χ²(1)=0.4,

p=0.5271

χ²(1)=0.2,

p=0.65470

χ²(1)=2.6,

χ²(1)=7.4, p=0.00540

 $y^2(1)=0.2$

