

Impact of chronic doxycycline treatment in the APP/PS1 mouse model of Alzheimer's disease

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26 ABSTRACT

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Due to the pathophysiological complexity of Alzheimer's disease, multitarget 28 29 approaches able to mitigate several pathogenic mechanisms are of interest. Previous 30 studies have pointed to the neuroprotective potential of Doxycycline (Dox), a safe and 31 inexpensive second-generation tetracycline. Dox has been particularly reported to slow 32 down aggregation of misfolded proteins but also to mitigate neuroinflammatory 33 processes. Here, we have evaluated the pre-clinical potential of Dox in the APP/PS1 34 mouse model of amyloidogenesis. Dox was provided to APP/PS1 mice from the age of 35 8 months, when animals already exhibit amyloid pathology and memory deficits. 36 Spatial memory was then evaluated from 9 to 10 months of age. Our data demonstrated 37 that Dox moderately improved the spatial memory of APP/PS1 mice without exerting 38 major effect on amyloid lesions. While Dox did not alleviate overall glial reactivity, we 39 could evidence that it rather enhanced the amyloid-dependent upregulation of several 40 neuroinflammatory markers such as CCL3 and CCL4. Finally, Dox exerted 41 differentially regulated the levels of synaptic proteins in the hippocampus and the cortex 42 of APP/PS1 mice. Overall, these observations support that chronic Dox delivery does 43 not provide major pathophysiological improvements in the APP/PS1 mouse model.

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46 **KEYWORDS**

47 Doxycycline; Alzheimer's Disease; mouse model

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51 **INTRODUCTION**

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Alzheimer's disease (AD) is a neurodegenerative disorder defined by the presence of
 two

55 neuropathological lesions: intraneuronal aggregates of hyperphosphorylated Tau 56 proteins and extracellular accumulation of toxic A β peptides, generated by the 57 sequential cleavages of the Amyloid Precursor Protein (APP; for a review see (Müller et 58 al., 2017). In addition, neuroinflammation - characterized by the pathological reactivity 59 of microglial and astrocytic cells as well as the release of various cytokines/chemokines 60 - develops along with AD brain lesions (Heneka et al., 2018; Laurent et al., 2018). 61 These pathological changes promote early synaptic dysfunctions, notably at forebrain 62 glutamatergic synapses (Canas et al., 2014; Dennis J. Selkoe, 2002; Kirvell et al., 2006) 63 and, subsequently, synapse loss leading to the progressive development of cognitive deficits. 64

65 Due to the pathophysiological complexity of AD, multitarget approaches to mitigate at 66 several pathological mechanisms the same time (protein aggregation. 67 neuroinflammation, neuronal loss...) might be of interest. Previous studies have pointed 68 to the neuroprotective effects of tetracyclines, a class of antibiotics, against 69 neurodegenerative and neuroinflammatory conditions (D. Orsucci, V. Calsolaro, M. 70 Mancuso, 2009). Among tetracyclines, doxycycline (6-Deoxy-5-hydroxytetracycline; 71 Dox) is a safe and inexpensive second-generation tetracycline which has demonstrated 72 safe toxicological profile, even after long-lasting administration (Haïk et al., 2014) and 73 good brain penetration (Domercq and Matute, 2004). Dox has been reported to exhibit 74 neuroprotective properties in a large range of neurodegenerative models not only thanks 75 to its ability to slow down aggregation of misfolded proteins but also to mitigate

76 neuroinflammatory processes and enhance neurogenesis (for review see (Santa-Cecília 77 et al., 2019; Sultan et al., 2013). In particular, Dox inhibit AB42 aggregation and disassemble preformed fibrils (Forloni et al., 2001). Accordingly, in Caenorhabditis 78 79 *elegans* and drosophila models of A β toxicity, Dox was found to mitigate A β 80 proteotoxicity, by interacting with A^β assemblies and reducing formation of amyloid 81 aggregates, thereby improving motor behavior (Costa et al., 2011; Diomede et al., 82 2010). Only one study evaluated the potential impact of Dox in an animal model of AD. (Balducci et al., 2018) tested its efficacy in the APP/PS1 mouse model of 83 84 amyloidogenesis. This study showed that subchronic i.p. treatment (20 days or 2 85 months) with the antibiotic restored recognition memory in aged APP/PS1 animals 86 without impacting cortical A^β plaques. This effect on memory was confirmed by the 87 same authors in another mouse model induced by i.c.v injection of A^β oligomers. In 88 both models, the benefit observed was associated with a lower neuroinflammation.

In the present study, we extended this former work and evaluated the impact of chronic Dox delivery in mid-aged APP/PS1 mice and found that while Dox exerted a significant impact in alleviating spatial memory impairments in two spatial memory tasks, without exerting major effect on amyloid lesions, we could evidence paradoxical effects towards inflammatory and synaptic markers in the hippocampus and the cortex. These observations support that the clinical use of Dox should be considered with caution in the context of AD.

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97 MATERIAL AND METHODS

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Animals. Heterozygous male APPswe/PS1dE9 mice (41 in total) were used in this
study (herein referred to as APP/PS1, C57B16/J background; (Jankowsky et al., 2001)

together with littermates controls (WT). All animals were maintained in standard cages
under conventional laboratory conditions (12 h/12 h light/dark cycle, 22°C), with ad
libitum access to food and water. Mice were maintained 5–6 per cage. The animals were
used in compliance with European standards for the care and use of laboratory animals
and experimental protocols approved by the local Animal Ethical Committee (CEEA75,
Lille, France).

108 Treatment. To avoid potential stress associated with multiple i.p. injections or gavage, 109 animals were orally treated through drinking water with Doxycycline hydrochloride 110 (Thermo Fisher; 0.4 mg/mL in 2.5% sucrose) or with 2.5% sucrose, in control 111 conditions. Animals were randomized into four experimental groups: WT-Sucrose 112 (WT), WT-Dox (WT Dox), APP/PS1-Sucrose (APP), APP/PS1-Dox (APP Dox). 113 Doxycycline solution was kept in dark bottles protected from light and changed 3 times 114 per week. Considering the average intake measured along the treatment and the body 115 weight of mice, the oral Dox treatment corresponded to circa. 50 mg/kg i.e. in the range 116 providing protection to APP/PS1 mice as shown in (Balducci et al., 2018). Treatment 117 started from 8 months of age - at a time amyloid pathology and memory deficits are 118 already present in this strain (Zhou et al., 2015) - i.e. one month before behavioral 119 experiments; and animals were kept under Vehicle or Dox until the end of the 120 experimental procedure i.e. 9-10 months of age.

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Behavioral studies. All the animals underwent all the behavioral tests described belowin a blind manner. An interval of 48h was respected between each task.

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Actimetry. Mice were placed in the center of an infrared Actimeter (45x45x35cm;
Bioseb), composed by a 2-dimensional square frame, and left for 10 min. Spontaneous
behavior of mice was tracked with distance travelled and velocity recorded by Actitrack
software (Bioseb).

129 Y-maze task. Short-term spatial memory was assessed in a spontaneous novelty-based 130 spatial preference Y-maze test. All arms of the Y-maze were 28 cm long, 6.2 cm wide, 131 with 15-cm-high white opaque walls. Different extra maze cues (elements of different 132 colors and shapes such as yellow plastic bag, a black rectangular box and a panel with 133 red and black circles) were placed on the surrounding walls. To avoid intra-maze odor 134 cues, sawdust was placed in the maze and mixed between each phase. Allocation of 135 arms was counterbalanced within each group. During the learning phase, mice were 136 placed at the end of the 'start' arm and were allowed to explore the 'start' arm and the 137 'other (familiar)' arm for 5 min (beginning from the time the mouse first left the start 138 arm). Access to the third arm of the maze ('novel' arm) was blocked by an opaque door. 139 The mouse was then removed from the maze and returned to its home cage for 2 min. In 140 the test phase the mouse was placed again in the 'start' arm of the maze, the door of the 141 'novel' arm was removed, and mouse behavior recorded for one minute (from the time 142 the mouse first left the start arm). The amount of time the mouse spent in each arm of 143 the maze was recorded during both learning and test phases using EthovisionXT 144 (Noldus Information Technology, Wageningen, The Netherlands). For the learning 145 phase, we calculated the percentage of time spent in the 'other' (familiar) arm vs. the 146 'start' arm. For the test phase, a discrimination index [novel arm/(novel+familiar arm)] 147 \times 100 was calculated.

149 Barnes Maze Task. Spatial learning and memory were evaluated using the Barnes maze 150 task. This behavioral apparatus is a white circular PVC open platform surface (120 cm 151 of diameter) with 40 equally spaced holes (5 cm of diameter) located at 5 cm from its 152 circumference and a black escape box located under one of the 40 holes. The platform is 153 placed on a swivel system (to rotate it easily) in the center of the room, elevated 80 cm 154 above the floor, enlighten by a flood light (800 lux) placed above and surrounded by 155 spatial cues. Mice completed a habituation trial to become familiar with the circular 156 platform environment and to practice descending into the escape box. Mice were free 157 for 5 min to explore the area around the escape hole and the escape box. Then, mice 158 completed 4 days of acquisition training (learning phase) with 4 trials per day. For each 159 trial, mice were placed in the start tube in the center of the circular platform (for 5-10 s) 160 and then trained to locate the escape hole (randomized for all mice) using spatial cues 161 (panels with different colors and shapes place on the wall such as a pink heart, yellow 162 face and black and white stripes flag) surrounding the platform. If a mouse did not enter 163 into the escape hole in 3 min it was gently guided to the escape hole. Mice remained 60s 164 in the escape box before returned to its home cage. The inter-trial interval was 15 min. 165 To reduce possible odor cues, the platform surface and escape box were cleaned with 166 70% Ethanol between each trial and the circular platform was rotated clockwise a 167 quarter turn every day. For each trial, path length to enter the escape box was recorded 168 using the Ethovision XT tracking system (Noldus). 24h after the last day of the training, 169 mice completed a 90s probe trial where the escape box was removed. The time spent in 170 the Target quadrant (T) versus average non-target quadrants (O) was determined.

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Sacrifice and Brain Tissue Preparation. To sacrifice animals, they were deeply
anesthetized with a mixture of ketamine/xylazine, then transcardially perfused with cold

174 NaCl (0.9%). The brain was then removed. One hemisphere was post-fixed in 4% 175 paraformaldehyde fixative in PBS (pH 7.4) for 24h at 4°C and transferred to 30% 176 sucrose solution before being frozen at -40°C in isopentane (methyl-butane). Brains 177 were cut in coronal floating sections (35 µm) using a Leica cryostat and stored in PBS-178 azide (0.2%) at 4°C. For each animal, 6-7 sections were collected per animal from 179 bregma -1.46mm to bregma -3.08mm, encompassing the whole hippocampus. The other 180 hemisphere was used for biochemical assays and cortex and hippocampus were 181 dissected out 4°C and stored at -80°C.

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183 Biochemical analyses. For all biochemical experiments, tissue was homogenized in 200 184 µL Tris buffer (pH 7.4) containing 10% sucrose, sonicated, and kept at -80°C until use. 185 Protein amounts were evaluated using the BCA assay (Pierce), subsequently diluted 186 with lithium dodecyl sulfate (LDS) 2X supplemented with reducing agents (Invitrogen) 187 and then separated on 4-12% Criterion Bis-Tris Gels (Invitrogen). For total AB (6E10), 188 Aβ42 (21F12) and C-terminal fragments (CTFs), proteins were separated on 16.5% 189 Criterion Gel Tris tricine (Invitrogen). Proteins were transferred to nitrocellulose or 190 PVDF membranes, which were then saturated (5% non-fat dry milk or 5% BSA) in 191 TNT (Tris 15mM pH8, NaCl 140mM, 0.05% Tween) and respectively incubated with 192 primary (see **Table 1**) overnight and corresponding secondary antibodies: peroxidase 193 labeled horse anti-rabbit and anti-mouse (1/5000 and 1/50000 respectively, Vector 194 Laboratories). Signals were visualized using chemiluminescence kits (ECLTM, 195 Amersham Bioscience) and a LAS4000 imaging system (Fujifilm). Results were 196 normalized to β -actin or GAPDH and quantifications were performed using ImageJ 197 software.

199 **ELISA.** Brain levels of human A β 1-40 and A β 1-42 were measured using ELISA kits 200 (Invitrogen, CA, USA; IBL-International, Hamburg, Germany) following 201 manufacturer's instructions. For hippocampal and cortical samples, AB was first 202 extracted as described previously (Faivre et al., 2018). Briefly, 20 µg of protein were 203 diluted in Guanidine/Tris buffer. Samples were then diluted in a Dulbecco's phosphate 204 buffered saline containing 5% BSA and 0.03% Tween-20 (BSAT-DPBS) solution. The 205 homogenates were centrifuged, and supernatants were collected for the analysis of A β 1-206 40 and A β 1-42 by colorimetric immunoassays. To measure levels of human A β 207 oligomers we used the 82E1-specific ELISA kit (IBL-International, Hamburg, 208 Germany). 20 µg of proteins in tris sucrose buffer were used for this analysis. In all 209 cases, absorbances were measured by Multiskan Ascent counter (ThermoLab Systems). 210 The normalized amounts of A β and A β oligomers were expressed as pg/mL or pmol/L, 211 respectively.

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213 Immunohistochemistry and Image Analysis. All antibodies used in this study are 214 detailed in Table 1. For Aß immunohistochemistry (IHC), sections were first pretreated 215 with 80% formic acid for 3 min, after washing them in water, they were permeabilized 216 with 0.2% Triton X-100/sodium phosphate buffer. Sections were then blocked with 217 10% "Mouse On Mouse" Kit serum (Vector Laboratories) for 1 h before incubation with mouse biotinylated anti-AB antibody (6E10) at 4°C overnight. After washing in 218 219 PBS, labelling was amplified by the application of the avidin-biotin-HRP complex 220 (ABC kit, 1:400 in PBS, Vector) prior to addition of diaminobenzidine 221 tetrahydrochloride (DAB, Vector) in Tris-HCl 0.1 mol/l, pH 7.6, containing H₂O₂ for 222 visualization. Brain sections were then mounted, air-dried, dehydrated by passage 223 through a graded series of alcohol (70%, 95%, 100%) and toluene baths, and finally 224 VectaMount (Vector Laboratories). For Iba1 mounted with and GFAP 225 immunohistochemistries, sections were first permeabilized with 0.2% Triton X-226 100/sodium phosphate buffer. Sections were then blocked with 1/100 normal goat 227 serum (Vector Laboratories) for 1 h before incubation with rabbit anti-GFAP and rabbit 228 anti-Iba1 antibody at 4°C overnight. After washing in PBS, the sections were incubated 229 with anti-rabbit secondary biotinylated antibody (Vector BA-1000) during 1h in 0.2% Triton X-100/sodium phosphate buffer. After washing in PBS, the sections were 230 231 incubated with the ABC kit developed using DAB, mounted, and dehydrated as before. 232 For blinded quantification, section imaging was performed using a slide scanner 233 (Axioscan Z1-Zeiss) with a 20× objective. Brain sections were analyzed using Mercator 234 image analysis system (Explora Nova, La Rochelle, France) using hue, saturation, and 235 intensity to distinguish objects in the image field. Thresholds were established manually 236 using identified objects on a set of slides and these segmentation thresholds remained 237 constant throughout the analysis. ROIs were outlined as follows: for the hippocampus, 238 the whole hippocampal signal was measured in each animal all along the rostro-caudal 239 axis and averaged; for the cortex, the signal was measured all along the rostro-caudal 240 axis from the dorsal part of visible cortex (retrosplenial cortex) to the more ventral part 241 of the cortex (excluding the piriform). Results are presented either as per cent marker 242 occupancy (immunoreactive area normalized to the whole area analyzed) or the number 243 of plaques and the average plaque size expressed as percentage of analyzed area.

RNA extraction and quantitative real-time PCR analysis (RT-qPCR). Total RNA was
extracted from hippocampi and cortex and purified using the RNeasy Lipid Tissue Mini
Kit (Qiagen, France). 500 nanograms of total RNA was reverse-transcribed using the
Applied Biosystems High-Capacity cDNA reverse transcription kit. Quantitative real-

time RT-PCR analysis was performed on an Applied Biosystems Prism 7900 System using Power SYBR Green PCR Master Mix. The thermal cycler conditions were as follows: hold for 10 min at 95°C, followed by 45 cycles of a two-step PCR consisting of a 95°C step for 15 s followed by a 60°C step for 25 s. Sequences of primer used are given in **Table 2**. Cyclophilin A was used as a reference housekeeping gene for normalization. Amplifications were carried out in triplicate and the relative expression of target genes was determined by the $\Delta\Delta$ CT method.

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Statistics. Results were expressed as means ± SEM. Differences between groups were
determined using unpaired Student's t-test, One-sample t-test or Two Way-analysis of
variance (ANOVA) followed by a Tukey's post-hoc multiple comparison test using
GraphPad Prism Software. P values < 0.05 were considered significant.

261

262 **RESULTS**

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264 **Chronic doxycycline prevents spatial memory in APP/PS1 mice.** Eight month-old 265 WT and APP/PS1 mice were orally treated with Dox through drinking water (0.4 266 mg/mL circa 50 mg/kg), 4 weeks before behavioral evaluations, at a time APP/PS1 267 mice have been shown exhibit amyloid pathology and memory deficits (Zhou et al., 268 2015).

We first evaluated spontaneous locomotion using actimetry (**Figures 1A**) and found no significant effect of genotype (F(1,32)=1.82; P=0.18; Two-way ANOVA) and Dox treatment (F(1,32)=0.26, P=0.61); Similarly, distance travelled (**Figures 1B**) was not affected (Genotype: F(1,32)=1.88, P=0.18; Treatment: F(1,32)=0.31, P=0.58; Two-way ANOVA).

274 Then, we determined the impact of the treatment on both short and long-term spatial 275 memory, using Y-maze and Barnes-maze tasks, respectively. In the Y-maze task, during 276 the learning phase, all groups explored the maze equally, spending a similar amount of 277 time in the two available arms (Genotype: F(1,33)=0.56, P=0.45; Treatment: 278 F(1,33)=0.72, P=0.40; Two-way ANOVA; Figure 1C). During the test phase, using 279 Two-Way ANOVA, we could not evidence any difference in the discrimination index 280 regarding genotype (F(1,33)=0.51, P=0.48), treatment (F(1,33)=2.34, P=0.13) or their 281 interaction (F(1,33)=1.58, P=0.21; Figure 1C). It was however notable that the 282 discrimination index was significantly above chance (i.e. 50%; Figure 1C, dashed line) 283 for WT (P=0.05; One sample t-test), WT Dox (P<0.001) and APP Dox (P<0.001) but 284 not APP (P=0.43), suggesting that Dox might improve short-term memory in APP/PS1 285 mice.

286 Next, we evaluated long-term spatial memory using the Barnes maze. During the 287 learning phase, all groups showed a decrease in path length across trials (Time: F 288 (3,108)=59.66, P<0.0001) in a similar manner between groups with no difference 289 among genotypes (F(1,37)=0.88, P=0.35) nor genotype x treatment x time interaction 290 (F(3,108)=1.81, P=0.14; RM Two-way ANOVA; Figure 1E), indicative of a proper 291 spatial learning abilities in all groups. 24 hours following acquisition, a probe trial was 292 performed to assess spatial memory. The time spent in the target quadrant was found 293 different between genotypes (F(1,36)=6.86, P=0.01), with a significant difference 294 observed between WT and APP mice (P < 0.01; Two-way ANOVA followed by Tukey's 295 post-hoc test), indicating memory impairment of APP/PS1 mice. While the post-hoc 296 analysis could not show a significant difference between APP and APP dox animals 297 (P=0.17), a significant genotype x treatment interaction could be observed 298 (F(1,36)=6.05; P=0.02) and the percentage of time spent in the target quadrant by APP

299 Dox mice was found similar to WT (P=0.55, Two-Way ANOVA followed by Tukey's 300 post-hoc analysis). In the Barnes Maze, proper memory is acknowledged by a 301 percentage of time spent in the target quadrant significantly above 25%. Interestingly, 302 while APP mice did not reach that cut-off (P=0.46; One-Sample t-test, Figure 1F, 303 dashed lines), WT (P<0.0001), WT Dox (P<0.01) but also APP Dox (P<0.01) animals 304 exhibited a percentage of time spent in the target quadrant largely above chance, 305 supporting a beneficial influence of Dox towards memory of APP/PS1 animals. 306 Altogether, behavioral assessments support a moderate but beneficial impact of Dox to 307 the memory deficits of APP/PS1 mice.

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309 Impact of Doxycycline on amyloid pathology in the hippocampus and the cortex of 310 **APP/PS1 transgenic mice.** Age-dependent spatial memory impairment in APP/PS1 has 311 been shown to correlate with increased brain amyloid burden (Garcia-Alloza et al., 312 2006; Savonenko et al., 2005; Zhang et al., 2012). We therefore evaluated the potential 313 impact of Dox treatment on amyloid load and APP metabolism in APP/PS1 mice. First, 314 the concentrations of AB1-42 and AB1-40 were found similar in the hippocampus of 315 APP/PS1 mice treated with vehicle or Dox (Aβ1-42: P=0.37; Aβ1-40: P=0.44; Aβ1-42/ 316 Aβ1-40: P=0.87 in APP/PS1 Dox vs. Vehicle; Student's t-test; Figure 2A). Also, we 317 could not observe difference between these groups in both the cortex (A β 1-42: P=0.65; 318 Aβ1-40: *P*=0.65; Aβ1-42/ Aβ1-40: *P*=0.61) and the plasma (Aβ1-42: *P*=0.96; Aβ1-40: 319 P=0.37; A β 1-42/ A β 1-40: P=0.11). We also quantified A β oligomers in the 320 hippocampus and the cortex of both APP/PS1 groups (Figure 2B). Although no change 321 was observed in the hippocampus (P=0.72 in APP/PS1 Dox vs. Vehicle; Student's t-322 test; Figure 2B), A β oligomers were found significantly enhanced in the cortex of 323 APP/PS1 treated with Dox (P<0.001 in APP/PS1 Dox vs. Vehicle; Student's t-test;

324 Figure 2B). In addition, we analyzed hippocampal and cortical amyloid plaques using 325 6E10 immunohistochemistry (Figures 2C). The area covered by plaques in the 326 hippocampus and the cortex remained similar between APP and APP Dox animals 327 (Hippocampus: P=0.99; Cortex: P=0.21, APP/PS1 Dox vs. Vehicle using Student's t-328 test; Figures 2D). In the hippocampus, chronic treatment with Dox did not significantly 329 alter the density of amyloid plaques of low (50-250µm) or large (250-500µm) size 330 (Figure 2D; Hippocampus: F (1,14)=0.005, P=0.93, Cortex: F(1,14)=2.13, P=0.16 vs. 331 APP/PS1 using RM Two-Way ANOVA). Finally, we evaluated the APP metabolism in 332 both APP/PS1 groups. In the hippocampus, no difference could be observed between 333 APP/PS1 vehicle and Dox regarding levels of APP, APP carboxy-terminal fragments 334 (CTFs), total A β (6E10) or A β_{42} (21F12) levels) (Figure 2E). In the cortex, APP level 335 was slightly increased in the Dox group (P=0.02, in APP/PS1 Dox vs. Vehicle; 336 Student's t-test; Figure 2F) while CTFs, total A β and A β_{42} were not modified (Figure 2F). These data indicated that, overall, Dox is ineffective to alter amyloidogenesis in 337 338 both cortex and hippocampus of APP/PS1 mice.

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340 Effect of chronic Doxycycline treatment on neuroinflammatory markers in the 341 hippocampus and the cortex of APP/PS1 mice. Amyloid pathology is known to 342 promote microglial and astrocytic reactivity and a chronic pro-inflammatory 343 environment, favoring the development of memory deficits (Heneka et al., 2018). 344 Moreover, Dox was previously demonstrated to exhibit anti-neuroinflammatory 345 properties, particularly in an amyloid context (Balducci et al., 2018). Therefore, we first 346 evaluated the outcome of Doxycycline treatment towards glial cell activation through 347 IHC immunoreactivity for Iba1 (microglia) and GFAP (astrocytes), in the hippocampus 348 and the cortex of WT and APP/SP1 mice treated with vehicle and Dox (Figure 3A-B).

349 While 9m-old APP/PS1 mice exhibit reactivity of microglia (Cortex: F(1,23)=30.60, 350 $P \le 0.001$; Hippocampus: F(1,22)=29.74, $P \le 0.001$; Figure 3A) and astrocytes (Cortex: 351 F(1,22)=25.01, P<0.001vs. WT; Figure 3B) as compared to WT animals, Dox modified 352 none of these glial activations. In addition, we also evaluated glial and 353 neuroinflammatory markers in the hippocampus and the cortex of all mouse groups 354 using qPCR. Overall, data clearly showed the upregulation of most of the glial and 355 neuroinflammatory markers studied in both the hippocampus and the cortex of APP/PS1 356 mice as compared to WT animals (Figures 4A-B). In the hippocampus (Figure 4A), 357 while most markers (GFAP, CD68, TLR2 C1qa, TNFa) were not significantly modified by the Dox treatment in APP/PS1 mice, several of them were surprisingly found 358 359 upregulated by the tetracycline such as CCL3 (F(1,35)=9.70, P=0.004; P<0.01, APP vs. 360 APP Dox using Two-way ANOVA followed by Tukey's post-hoc test), CCL4 361 (F(1,36)=10.34; P=0.003; P<0.01, APP vs. APP Dox), Clec7a (F(1,35)=14.46, 362 *P*<0.001; *P*<0.001, APP vs. APP Dox) and Itgax (F(1,33)=24.30, *P*<0.001; *P*<0.0001, 363 APP vs. APP Dox). In the cortex (Figure 4B), most of the glial and inflammatory 364 makers were also upregulated in APP/PS1 mice as compared to WT animals but were 365 not significantly altered by Dox, with the exception of CCL4, upregulated as compared 366 with APP/PS1 animals (F(1,28)=6.369, P=0.02; P<0.01, APP vs. APP Dox). Overall, 367 our data suggest that Dox potentiate the amyloid-induced upregulation of neuro-368 inflammatory markers in APP/PS1 mice, with a most effective impact at the 369 hippocampal level.

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371 Effect of chronic Doxycycline treatment on hippocampal and cortical synaptic
372 markers in APP/PS1 mice. Finally, we evaluated, in APP/PS1 mice, the impact of Dox
373 treatment on the level of various synaptic markers (AMPA and NMDA receptors,

pre/post-synaptic markers). In the hippocampus, we did not observe change in the 374 375 amount of NMDA receptor subunits (NR1, NR2A, NR2B) as well as for GluR2 and 376 PSD95. Interestingly, GluR1 was found increased in the hippocampus of APP/PS1 after Dox treatment (+27.6± 4.5% vs APP/PS1, P<0.001 using Student's t-test; Figure 5A). 377 378 Dox treatment also enhanced the level of the pre-synaptic SNAP25 protein 379 $(+46.2\pm17.1\% \text{ vs APP/PS1}, P=0.05 \text{ using Student's t-test; Figure 5A})$ as well as the 380 post-synaptic Cofilin protein (+114.7± 2.6% vs APP/PS1, P<0.001 using Student's t-381 test; Figure 5A). However, Munc18 was found downregulated in mice treated with Dox 382 $(-33.4 \pm 7.9\% \text{ vs APP/PS1}, P=0.02 \text{ using Student's t-test; Figure 5A})$. In the cortex, 383 while Dox treatment did not change the levels of most markers, we observed a 384 significant decreased in NR2B protein levels as well as of its phosphorylation at Y1472 385 (NR2B: -28.7±5.3% vs. APP/PS1, P=0.02; pY1472 NR2B: -36.9±7.4% vs APP/PS1, 386 P=0.02 using Student's *t*-test; Figure 5B). These data suggest that Dox enhances the 387 level of several synaptic markers in the hippocampus while remains largely ineffective 388 in the cortex.

389

390 **DISCUSSION**

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392 Doxycycline is a safe second-generation tetracycline which has been reported to exhibit 393 neuroprotective properties in various neurodegenerative models through, in particular, 394 its ability to slow down aggregation of misfolded proteins and to mitigate 395 neuroinflammatory processes (Santa-Cecília et al., 2019; Sultan et al., 2013). So far, 396 only one study tested Dox in a mouse model of amyloidogenesis which showed that 397 acute or subchronic Dox improved recognition memory and lower neuroinflammation 398 without altering plaques in APP/PS1 mice or animals injected i.c.v. with Aβ oligomers

399 (Balducci et al., 2018). In the present study, we reinvestigated Dox potential in 400 APP/PS1 mice and showed that Dox administered chronically and orally during 2 401 months provide a benefit, even moderate, to 9 month-old APP/PS1, particularly in the 402 long-term spatial memory. No major impact on the amyloid load has been observed. 403 However, memory improvement was associated with an enhancement of some synaptic 404 markers, in the hippocampus only. Puzzlingly, Dox also enhanced the mRNA 405 expression of various neuroinflammatory markers responsive to amyloid pathology. 406 Overall, the present data question the potential of Dox in the context of AD.

407 Our data, showing that oral Dox improves spatial memory abilities in APP/PS1 mice 408 (essentially in Barnes task), are in line the previous study of (Balducci et al., 2018) 409 which demonstrated that the acute or subchronic i.p. delivery of Dox mitigate A β -410 induced memory deficits. This is in accordance with the improvement seen in 411 invertebrate Aβ models (Costa et al., 2011; De Luigi et al., 2008; Diomede et al., 2010). 412 However, in sharp contrast with other studies (Forloni et al., 2001; Paldino et al., 2020) 413 which showed that Dox is prone to disassemble $A\beta$ fibrils, the mouse studies of 414 (Balducci et al., 2018) were not able to observe an effect on amyloid plaques, even the 415 levels of AB oligomers in the hippocampus seemed reduced in APP/PS1 mice treated 416 with the antibiotic. However, we do not confirm the beneficial impact upon oligomers 417 and our data globally indicate that Dox does not impact amyloid load in APP/PS1 mice. 418 Surprisingly, spatial memory improvements of Dox-treated APP/PS1 animals was 419 associated with the enhancement of several hippocampal neuroinflammatory markers. 420 This is at odds with other reports supporting anti-inflammatory properties of Dox (Balducci et al., 2018). Dox particulalry enhances CCL3, CCL4, Clec7a and Itgax that 421 422 are aged and AD associated genes, which suggest anyhow an immunomodulatory 423 impact of Dox (Benmamar-Badel et al., 2020). However, while CCL3 and CCL4 were

thought to significantly affects hippocampal plasticity and related memory (Carvalho et
al., 2021; Marciniak et al., 2015), although we cannot exclude that chemokines might
also exert beneficial effects on plasticity and memory as suggested by others (Ajoy et
al., 2021; Couturier et al., 2016; Kuijpers et al., 2010). It is therefore difficult to link
neuroinflammatory changes observed to spatial memory effects of Dox in APP/PS1.

429 Our data indicate that the antibiotic enhances the hippocampal level of several synaptic 430 proteins important for plasticity, notably GluR1, SNAP25 and Cofilin. The enhanced 431 level of the GluR1 subunit of AMPA receptors likely fits with memory improvement 432 (Mitsushima et al., 2011). Enhanced SNAP25 was likely associate with memory 433 improvement in models of AD (Ahmad et al., 2017; Tweedie et al., 2012). However, the 434 level of SNAP25 might also results in impaired memory (McKee et al., 2010) and, in 435 AD patients, higher SNAP25 has been associated with worse memory (Pereira et al., 436 2021). The rise of Cofilin level remains also hard to firmly interpret. Indeed, actin 437 depolymerizing factor (ADF)/cofilin family of actin-associated proteins is important for 438 dendritic spine plasticity. (Gu et al., 2010) showed that cofilin-mediated actin dynamics 439 regulate AMPA receptor trafficking during synaptic potentiation, thereby improving 440 plasticity. In line, cofilin-1 conditional forebrain knock-out mice showed impaired late 441 phase LTP and LTD (Rust et al., 2010) and cofilin finely tunes synaptic function (Ben 442 Zablah et al., 2020). However, when levels of cofilin are elevated, they can also drive 443 the formation of cofilin-actin rods, as seen in the AD brain, which impairs synaptic 444 function. Then, genetic reduction of ADF/cofilin rescues neurodegeneration but also 445 LTP and contextual memory in APP/PS1 mice (Woo et al., 2015, 2012). Therefore, it 446 will be important to interpret these biochemical data in the light of future 447 electrophysiological recordings to clearly estimate the synaptic impact of Dox in 448 APP/PS1 mice and further argue on the therapeutical potential of this compound.

It is also noteworthy that Dox is largely used to regulate gene transcription in conditional Tet-On and Tet-Off transgenic mouse models (Sprengel and Hasan, 2007). Several Tet-On models of neurodegeneration are particularly used (Jankowsky and Zheng, 2017; Khlistunova et al., 2006; Liu et al., 2015)(). Our data therefore highlight that activating the expression of neuropathologically-relevant genes in the brain with Dox might potentially interferes with the pathological processes by themselves and those careful controls are warranted to address that issue.

456 While Dox has not been tested as a monotherapy in AD patients, two clinical trials 457 associating Dox with another antibiotic, rifampin, provided different outcomes. A first 458 trial suggested positive cognitive outcomes (Loeb et al., 2004) while the second failed 459 to reproduce a benefit (Molloy et al., 2013). While reasons for such discrepancies 460 remain unclear (dose, association therapy....), our findings are not overall arguing for a 461 clear potential of Dox in AD. In conclusion, while in a reliable model of AD, Dox is 462 prone to improve memory, the molecular and neuropathological changes induced by 463 this antibiotics need further investigation to define its real potential for further clinical 464 development in the AD context.

465

466	FIGUR	E LEGE	ENDS

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Figure 1. Impact of chronic doxycycline on the spatial memory of APP/PS1 mice.
(A-B) Actimetry. No impact of Dox on spontaneous locomotion was in both WT and
APP/PS1 mice. (C-D). Y-maze task. (C) During learning phase, mice spent a similar
percentage of time in the familiar arm vs the start arm. (D) In the test phase, the
discrimination index was found similar regardless genotype and treatment (Two-Way
ANOVA). However, all groups, except the APP mice, exhibited a percentage of time in

474 the novel arm significantly higher that the theoretical mean of 50% (dashed line). (E-F) 475 Barnes Maze. (E) Evaluation of the path length needed to find the hidden platform 476 during the spatial learning phase. During the 4 days of training, there was no significant 477 genotype x treatment x time difference (RM Two-Way analysis). (F) Spatial memory 478 was assessed 24h after the last training session. The percentage of time spent in the 479 target quadrant by APP mice was significantly lower as compared to WT mice 480 (*p<0.05, Two-Way ANOVA followed by Tukey's multiple comparison test) while it 481 remained similar to WT for APP dox animals (NS). Moreover, the percentage of time 482 spent in the target quadrant was found significantly different from the theoretical mean 483 (ie. 25%, dashed line) for all group but APP. Results are expressed as mean ±s.e.m. 484 N=7-13 animals per group.

485

486 Figure 2. Impact of Doxycycline on hippocampal and cortical amyloid pathology. 487 (A) Hippocampal, cortical, and plasmatic levels of A β 1-40 and A β 1-42, assessed by 488 Elisa, showing no difference between APP/PS1 and APP/PS1 Dox groups. (B) $A\beta$ 489 oligomers levels quantified by Elisa. Levels were increased in the cortex of Dox-treated 490 APP/PS1 mice vs. control APP/PS1 animals. ### P<0.001 vs APP/PS1 mice using 491 Student's t-test. (C) Representative pictures of 6E10-positive amyloid plaques in cortex 492 and hippocampus of APP/PS1 mice treated or not by Dox (Scale bars = $500\mu m$). (D) 493 (a,c) Global A β load in the hippocampus and the cortex of APP and APP dox mice, 494 evaluated by the % of 6E10 surface stained. (b,d) Distribution of low vs large size 495 amyloid plaques in both structures. No effect of Dox could be observed (E-F) Western 496 blot evaluation of APP, C-terminal fragments (CTFs), total AB (6E10) and AB1-42 497 (21F12) in the hippocampus (E) and cortex (F) of APP/PS1 mice treated or not by Dox. 498 No change could be observed except a slight increase of APP (APP C-17) in the cortex

499 in Dox-treated animals. *P < 0.05 vs APP/PS1 mice using Student's t-test. Results are 500 expressed as mean \pm s.e.m. N=7-9 animals per group.

501

502 Figure 3. Impact of Doxycycline on hippocampal and cortical glial activation. 503 Effect of doxycycline on glial activation was determined through Iba1 (microglia; A) 504 and GFAP (astrocytes; B) immunoreactivities. Graphs showed an increased level of percentage of immunoreactive area for Iba1 and GFAP in APP/PS1 mice with no 505 506 impact of the Dox treatment. ** P<0.01, *** P<0.001 vs WT mice using Two-Way 507 ANOVA followed by Tukey's post hoc test. Scale bars = $200\mu m$. Results are expressed 508 as mean ±s.e.m of the average quantification of 4-7 sections per animals with N=6-8 509 animals per group.

510

511 Figure 4. Impact of Doxycycline on hippocampal and cortical neuroinflammation. 512 Hippocampal and cortical neuro-inflammation was evaluated in all experimental groups 513 using quantitative PCR for markers of astrocytic activation (GFAP), microglial 514 activation (CD68 and C1qa), chemokines/cytokines (CCL3, CCL4 and TNFa), and 515 inflammatory receptors (TLR2, Clec7a, Itgax). All markers except TNFa were found 516 significantly upregulated in APP/PS1 mice as compared to WT animals (WT vs. APP 517 comparisons represented by *). In the hippocampus, rise of Clec7a, Itgax, CCL3 and 518 CCL4 in APP/PS1 mice was further enhanced by Dox (APP vs. APP Dox comparisons 519 represented by #). In the cortex, CCL4 was also increased by Dox treatment in APP/PS1 520 mice. *P<0.05, **P<0.01, ***P<0.001 vs WT mice, ##P<0.01 and ###P<0.001 vs 521 APP/PS1 mice using Two-Way ANOVA followed by Tukey's post hoc test. Results are 522 expressed as mean ±s.e.m. N=7-9 animals per group.

524 Figure 5. Impact of Doxycycline on hippocampal and cortical synaptic markers. 525 Western blot evaluation of NMDA receptor (NR1, NR2A, NR2B, p1472 NR2B), 526 AMPA receptors (GluR1, GluR2), pre-synaptic (Munc18, SNAP25) and post-synaptic 527 (PSD95, Cofilin) markers as well as NSE neuronal marker in the hippocampus of 528 APP/PS1 treated or not by Dox. In the hippocampus, GluR1, SNAP25, Cofilin were 529 found increased in APP/PS1 mice treated with Dox while Munc18 was rather 530 decreased). In the cortex, Dox treatment led to a decrease of NR2B and p-Y1472NR2B. # P<0.05 and ## P<0.01 vs APP/PS1 mice using Student's t-test. Results are expressed 531 532 as mean ±s.e.m. N=4-8 per group

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549 AUTHOR CONTRIBUTIONS

550

V.Gomez, K.Carvalho, B.Thiroux, R.Caillierez M.Besegher and E.Faivre performed
experiments, analyzed data, and corrected the manuscript. E. Faivre, N. Sergeant,
L.Buée, and D.Blum supervised the work, analyzed data, and wrote the manuscript.

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В

Distance (cm)



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40

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В



CD68



Antibodies used				
Abbreviation	Origin	Provider	Reference	Dilution
β-actin	Rabbit	Sigma	A5441	1/10000
GAPDH	Mouse	Santa Cruz Biotechnology	25778	1/10000
NR2B	Rabbit	Millipore	06-600	1/1000
pY1472 NR2B	Rabbit	Cell Singaling	4208S	1/1000
NR2A	Rabbit	Cell Signaling	4205	1/1000
SNAP25	Mouse	Santa Cruz	376713	1/1000
Cofilin	Rabbit	abcam	ab134963	1/5000
NR1	Rabbit	Cell Signaling	5804	1/1000
Munc18	Rabbit	Sigma	M2694	1/2000
NSE	Rabbit	Life Science Enzo	BML-NA1247-0100	1/50000
GLUR1	Rabbit	Millipore	4855	1/5000
PSD95	Rabbit	Cell Singaling	25075	1/1000
6e10	Mouse	Covance Biolegend	10LC0282	1/1000
21F12	Mouse	Homemade		1/2000
C-ter part of APP, CTFs	Rabbit	Homemade		1/5000
GluR2	Rabbit	Abcam	20673	1/5000
GFAP	Rabbit	Dako	Z03334	1/1000
Iba1	Rabbit	Wako	1919741	1/1000

MOUSE								
Primers	Accession number	Forward primer	Reverse primer	Amplicon length				
Gfap	NM_001131020.1	CGCGAACAGGAAGAGCGCCA	GTGGCGGGCCATCTCCTCCT	104				
Cd68	NM_009853.1	GACCTACATCAGAGCCCGAGT	CGCCATGAATGTCCACTG	95				
Tlr2	NM_011905.3	GGGGCTTCACTTCTCTGCTT	AGCATCCTCTGCGATTTGACG	110				
Clec7a	NM_020008.2	ATGGTTCTGGGAGGATGGAT	GCTTTCCTGGGGAGCTGTAT	72				
Itgax	NM_021334.2	ATGGAGCCTCAAGACAGGAC	GGATCTGGGATGCTGAAATC	62				
Ccl3	NM_011337.2	TGCCCTTGCTGTTCTTCTCT	GTGGAATCTTCCGGCTGTAG	112				
Ccl4	NM_013652.2	GCCCTCTCTCTCCTCTTGCT	GAGGGTCAGAGCCCATTG	72				
Tnfa	NM_013693.2	TGCCTATGTCTCAGCCTCTTC	GAGGCCATTTGGGAACTTCT	116				
Cyclophilin A	NM_008907.1	AGCATACAGGTCCTGGCATC	TTCACCTTCCCAAAGACCAC	126				
Taqman probe	Accession number	Assay ID	Assay Design	Amplicon length				
C1qa	NM_007572.2	Mm00432142_m1	Probe spans exons	80				
Cyclophilin A	NM_008907.1	Mm02342430_g1	Probe spans exons	148				