

Protective Role of Cerebrospinal Fluid Inflammatory Cytokines in Patients with Amnestic Mild Cognitive Impairment and Early Alzheimer's Disease Carrying Apolipoprotein E4 Genotype

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Abstract.

Background: Neuroinflammatory cytokines can play a pivotal role in Alzheimer's disease (AD) contributing to the evolution of degenerative processes.

Objective: We aimed at evaluating the levels of cerebrospinal fluid (CSF) inflammatory cytokines, chemokines, and growth factors in subjects with diagnosis of amnestic mild cognitive impairment and mild AD.

Methods: We evaluated CSF contents of inflammatory cytokines in 66 patients divided according to the NIA-AA research framework and the *APOE* genotype. CSF of a group of cognitively unimpaired individuals ($n = 23$) was evaluated as control. All patients were evaluated for 24 months using Mini-Mental State Examination (MMSE).

Results: We found significant increased levels of IL-4, IL-6, IL-8, and G-CSF in the CSF of A+/T- *APOE4* carriers, respect to A+/T- patients homozygous for *APOE3*, respect to A+/T+ patients, regardless the *APOE* status, and respect to controls. Over a period of 24 months, A+/T- *APOE4* carriers, with increased levels of cytokines, showed a preserved cognitive evaluation when compared to the other subgroups of patients (delta MMSE at 24 months respect to baseline: 0.10 ± 0.35 ; $p < 0.05$).

Conclusion: Our data suggest that during early phases of AD, in *APOE4* carriers, A β pathology likely induces a specific cytokines pattern synthesis associated to cognitive preservation. These data highlight the different role that neuroinflammation can play in AD pathology based on the presence of specific CSF biomarkers and on the *APOE* status.

Keywords: Amyloid- β 42, *APOE*, cognitive decline, G-CSF, interleukins, tau

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INTRODUCTION

Alzheimer's disease (AD) is a multifactorial, chronic neurodegenerative disorder, which main pathological features are the extracellular senile plaques and the intraneuronal neurofibrillary tangles [1]. In last decades, neuropathological analysis of AD brains revealed that neuroinflammation is an important driving force for neurodegeneration and AD progression [2]. During physiological aging and in AD, cytokines levels increase and set neuronal environment in an inflammatory state [3], contributing to the evolution of degenerative process. Neuroinflammation is a complex mechanism mediated by cytokines mainly released by microglial cells and astrocytes, whose activation may have both detrimental or protective role for neurons. Beneficial pro-inflammatory cytokines are protective when involved in the induction and modulation of neuronal growth, cell survival, and modulation of synaptic plasticity mechanisms. Conversely, a prolonged and aberrant pro-inflammatory signaling is responsible for surrounding tissue neurodegeneration [4]. Microglial cells play a key role in the inflammatory process of the central nervous system (CNS) and represent a major focus of neurodegenerative disease research. Microglia could remain in balance between a pro-inflammatory status (M1 phenotype), characterized by the synthesis of inflammatory cytokines such as interleukin 1 β (IL-1 β), IL-6, and tumor necrosis factor (TNF), counteracted by the synthesis and release of anti-inflammatory cytokines (IL-4, IL-8, and IL-10) and neurotrophic factors (M2 phenotype), depending on the specific stimulus the microglia has been exposed to [5]. Thus, in such intricate scenario, the complex role of inflammatory cytokines in both neurodegeneration and neuroprotection is far from completion. In AD amyloid- β (A β) peptides, including both the oligomeric and the senile plaques forms, are considered main trigger for inflammatory signaling [6]. In particular, a prolonged proinflammatory signaling due to A β mis-metabolism, can lead to overproduction of pro-inflammatory cytokines involved in the neurodegenerative pathways signaling [4]. On the other hand, there is evidence that an increased A β production arises as a direct result of prolonged neuroinflammation [7]. It is important to note that in AD most of the modulatory effects of cytokines are related to the amyloid cascade signaling [8], while cognitive dysfunction progression is rather related to neuronal degeneration and tau-pathology [9]. Tau

protein, part of the neuronal cytoskeleton, is necessary for axonal physiology, for neurite outgrowth, neural plasticity mechanisms, repair of neurons after injuries [10], and even a regulatory role for cell firing has recently been described, giving to this protein a wider function than previously believed [11]. Impaired metabolism of tau protein has been demonstrated to rapidly induce impairment of neurotransmission and synaptic plasticity, all mechanisms responsible for cognitive decline in AD patients [12–15]. However, a clear relationship between tau pathology and neuroinflammation is still unclear. We recently showed that human astrocytes cultures incubated with cerebrospinal fluid (CSF) samples from AD patients were vulnerable in terms of increased apoptosis only in the presence of high levels of tau protein and APOE4 genotype [16]. Such findings led us to hypothesize a major role for tau protein in astrocytes degeneration and likely a proinflammatory role for tau in APOE4 individuals [17]. Interestingly, it has been reported that APOE4 carriers present an unbalanced switching of the microglial phenotype M1–M2 [18]. Moreover, microglial ApoE downstream regulates the microglial homeostatic gene expression, leading to a neurodegenerative-associated phenotype switch, which could further promote AD pathology [19].

Thus, the main objective of this study was to evaluate the levels of 15 cytokines in the CSF of patients with amnesic mild cognitive impairment (aMCI) and mild AD, according to the hallmarks processes of amyloid deposition, tau pathology, and APOE genotype. To reduce possible discrepancies between clinical presentation and CSF biomarkers profile, patients were divided using the NIA-AA research framework [20]. The AT(N) classification divides biomarkers into amyloid deposits (A), neurofibrillary tangles (T), and neurodegeneration (N), determined by measuring CSF levels of A β ₄₂, phosphorylated tau (p-tau), and total tau (t-tau), respectively. Although biomarkers of neurodegeneration (N) provide important pathologic staging information they are not specific for neurodegeneration due to AD; for this reason, A and T biomarkers are commonly used to discriminate patients in the AD continuum, subdivided into AD pathologic change (A+/T-) and AD (A+/T+). Because of the evidence of modulation of ApoE isoforms in neuroinflammation [17–19] and the effect of neuroinflammation on the neurodegenerative processes in several types of dementia [21], we expect to find a different profile of neuroinflammatory cytokines in patients classified by AT and

134 APOE genotype. Furthermore, we expect to find dif- 183
135 ferent rate of disease progression among groups, thus 184
136 patients were evaluated with neuropsychological test- 185
137 ing for a period of two years. 186

138 METHODS 187

139 Subjects 188

140 Sixty-six consecutive patients (range, 58–79 years; 189
141 median, 71) were recruited at the memory clinic of 190
142 the University Hospital Tor Vergata, admitted for 191
143 complaining memory symptoms. The diagnosis of 192
144 probable or possible AD fulfilled the criteria of the 193
145 National Institute on Aging and Alzheimer’s Associa- 194
146 tion (NIA/AA) [22] and all patients had a mild disease 195
147 with Mini-Mental State Examination (MMSE) scores 196
148 ranging 20–24. The aMCI patients were diagnosed 197
149 using the NIA/AA criteria for MCI [23]. All patients 198
150 underwent, for diagnostic purposes, a complete 199
151 clinical investigation in a period not superior to 200
152 60 days, including medical history, neurological 201
153 examination, MMSE, a complete blood screening, 202
154 and neuropsychological assessment [24] including 203
155 the following cognitive domains: general cognitive 204
156 efficiency: MMSE; verbal episodic memory: Rey 205
157 auditory verbal long-term memory (15-Word List 206
158 Immediate and 15 min Delayed recall); visuospatial 207
159 abilities and visuospatial episodic memory: Com- 208
160 plex Rey’s Figure (copy and 10 min Delayed recall); 209
161 and executive functions: phonological word fluency; 210
162 analogic reasoning: Raven’s Colored Progressive 211
163 Matrices. Patients underwent also a neuropsychi- 212
164 atric evaluation, magnetic resonance or computed 213
165 tomography (CT) imaging, positron emission tomog- 214
166 raphy/CT, and lumbar puncture for CSF analysis. 215
167 Exclusion criteria were: cognitive isolated deficits, 216
168 clinically manifest acute stroke in the last 6 months 217
169 showing a Hachinski scale score >4, and a radio- 218
170 logical evidence of ischemic lesions, A β _{1–42} CSF 219
171 values >600 pg/mL. All patients started treatment 220
172 with rivastigmine patch or donepezil and were fol- 221
173 lowed longitudinally with clinical assessments and 222
174 MMSE testing at 6, 12, and 18 months. 223

175 Control patients ($n=23$) were evaluated for 224
176 headache in the Policlinico Tor Vergata Hospital 225
177 Emergency Department between October 2014 and 226
178 December 2015, and the CSF samples were collected 227
179 in accordance with standard hospital practice. The 228
180 control subjects did not carry a diagnosis of active 229
181 infection and were free of cognitive and primary neu- 230
182 rological disorders other than headache. 231

183 All participants or their legal guardian provided 184
185 written informed consent after receiving an extensive 186
187 description of the study. The study was performed 188
189 according to the Declaration of Helsinki. The ethics 190
191 committee of the Santa Lucia Foundation approved 192
193 this protocol (Prot. CE/AG4/PROG.392-08). 194

189 Biomarkers collection and genotype analysis 190

191 The first 12 mL of CSF were collected in a 192
193 polypropylene tube and directly transported to the 194
195 local laboratory for centrifugation at 2000 g at +4°C 196
197 for 10 min. The supernatant was pipetted off, mixed 198
199 to avoid potential gradient effects and aliquoted 200
201 in 1 mL portions in polypropylene tubes, stored 202
203 at –80°C pending biochemical analyses. CSF t-tau 204
205 and p-tau phosphorylated at Thr181 concentrations 206
207 were determined using a sandwich enzyme-linked 208
209 immunosorbent assay (ELISA; Innostest hTAU-Ag; 209
210 Innogenetics, Gent, Belgium). A β _{1–42} levels were 210
211 determined using a sandwich ELISA (Innostest β - 211
212 amyloid; Innogenetics) [25]. Genotyping for APOE 212
213 were performed by allelic discrimination technology 213
214 (TaqMan; Applied Biosystems). 214

205 CSF cytokines and chemokines determination 206

207 In a group of 89 individuals CSF contents of 208
209 cytokines and chemokines were determined. These 209
210 include IL-1 β , IL-2, IL-4, IL-6, IL-7, IL-8, IL- 210
211 10, IL-12, IL-13, IL-17, tumor necrosis factor-alpha 211
212 (TNF α); granulocyte colony-stimulating factor (G- 212
213 CSF), granulocyte-macrophage colony-stimulating 213
214 factor (GM-CSF); macrophage inflammatory pro- 214
215 teins (MIP)-1a and monocyte chemotactic protein 1 215
216 (MCP-1). To determine cytokines and chemokines, 216
217 the CSF was centrifuged and immediately stored 217
218 at –80°C until analyzed using Bio-Plex Multiplex 218
219 Cytokine Assay (Bio-Rad Laboratories, Hercules, 219
220 CA), according to manufacturer’s instructions. Con- 220
221 centration of analytes were calculated according 221
222 to a standard curve and expressed as picograms 222
223 per milliliter. When the concentrations of the ana- 223
224 lytes were below the detection threshold, they were 224
225 assumed to be 0 pg/ml; a maximum of values below 225
226 the limit of detection of 5% for each cytokine was 226
227 considered acceptable for the analysis. 227

226 Statistical analysis 227

228 Data are presented as mean \pm standard deviation 229
229 (SD). Differences among groups were compared 230
230 231

Table 1
Demographical and clinical data of healthy controls and patients divided using the NIA-AA classification and APOE genotype

	HC (n = 23)	(A+/T-) E3 (n = 20)	(A+/T-) E4 (n = 16)	(A+/T+) E3 (n = 19)	(A+/T+) E4 (n = 11)	p
Age, y (mean ± SD)	66.8 ± 7.6	70.3 ± 6.4	70.7 ± 5.6	69.8 ± 7.1	70.0 ± 6.3	0.34
Female (%)	59%	60%	62%	63%	63%	0.89
Disease duration, m (mean ± SD)	n.a.	11.3 ± 3.5	10.9 ± 3.3	11.3 ± 3.3	11.2 ± 3.9	0.99
Education, y (mean ± SD)	n.a.	8.4 ± 3.6	9.0 ± 3.5	9.1 ± 3.9	9.1 ± 3.5	0.93
MMSE (mean ± SD)	n.a.	24.6 ± 2.1	24.1 ± 1.8	24.2 ± 2.1	23.9 ± 2.1	0.99
CSF total-tau, pg/ml (mean ± SD)	n.a.	244.5 ± 112.1	267.1 ± 90.1	720.1 ± 217.8	707.4 ± 220	<0.01
CSF p-tau, pg/ml (mean ± SD)	n.a.	41.1 ± 15.9	47.2 ± 16.1	89.4 ± 23.4	86.3 ± 27.9	<0.01
CSF Abeta 1-42, pg/ml (mean ± SD)	n.a.	356.6 ± 97.2	377.7 ± 124.6	367.7 ± 99.2	399.6 ± 94.7	0.73
Diabetes (%)	13.1	20.0	18.7	21.1	18.2	0.97
Hypertension (%)	34.8	30.0	31.3	31.6	36.3	0.99
Hyperlipidemia (%)	30.4	30.0	31.3	31.6	27.3	0.98
Arthritis (%)	4.3	5.0	12.5	15.8	9.1	0.68
Thyroiditis (%)	8.7	15.0	12.5	15.8	18.2	0.93
COPD (%)	8.7	10.0	6.3	10.5	9.1	0.99
Cancer (%)	8.7	5.0	0	0	0	0.42
Multimorbidity (%)	26.1	20.0	31.2	31.6	27.3	0.92

n, numbers; y, years; m, months; SD, standard deviation; MMSE, Mini-Mental State Examination; CSF, cerebrospinal fluid; COPD, chronic obstructive pulmonary disease; n.a., not applicable.

229 by univariate analysis using one-way ANOVA for
230 continuous variables and Fisher Exact Test for cat-
231 egorical variables. The Tuckey test was used for
232 *post hoc* multiple comparison. All statistical analy-
233 ses were conducted using GraphPad Prism version
234 8.0 (GraphPad Software, San Diego, CA, USA). A *p*-
235 value (*p*) of less than 0.05 was considered statistically
236 significant.

237 RESULTS

238 Sixty-six consecutive patients were recruited at
239 the memory clinic of the University Hospital Tor
240 Vergata. All patients showed neuropsychological pro-
241 file compatible with a diagnosis of aMCI or mild
242 AD. Based on AT/N classification patients were
243 grouped in (A+/T-) E4, (A+/T-) E3, (A+/T+) E4,
244 and (A+/T+) E3. Groups did not differ in gender,
245 education, age at disease onset, disease duration,
246 MMSE score at baseline, chronic medical conditions,
247 and multimorbidity (defined as the coexistence of
248 two or more chronic conditions in the same individ-
249 ual) as shown in Table 1. Twenty-three cognitively
250 unimpaired subjects, evaluated for headache in the
251 Policlinico Tor Vergata Hospital Emergency Depart-
252 ment, underwent CSF sampling in accordance with
253 standard hospital practice and were used as control
254 subjects.

255 CSF levels of cytokines, chemokines, and growth 256 factors according to NIA-AA research framework 257 and APOE genotype

258 In this experimental setting cytokines, chemokines
259 and growth factor's levels (IL-1 β , IL-2, IL-4, IL-6,
260 IL-7, IL-8, IL-10, IL-12, IL-13, IL-17, TNF α ; G-
261 CSF, GM-CSF; MIP-1 and MCP-1) were determined
262 in CSF samples of each group of patients and con-
263 trols. Results showed that differences among groups
264 reached statistical significance only for G-CSF
265 ($F=6.463$; $p<0.001$), IL-4 ($F=4.059$; $p=0.004$),
266 IL-6 ($F=4.481$; $p=0.002$), and IL-8 ($F=5.296$;
267 $p<0.001$) (see Table 2). In particular, in the multi-
268 ple comparisons analyses, we found that G-CSF and
269 IL-4 levels were significantly higher in the (A+/T-)
270 E4 group ($p<0.05$ for all comparisons) (Fig. 1A, B).
271 Similarly, we found significant higher levels of IL-6
272 and IL-8 in the CSF of (A+/T-) E4 group respect to
273 (A+/T-) E3 group, (A+/T+) E4 group, and controls
274 ($p<0.05$ for all comparisons), as well as a strong ten-
275 dency compared to the (A+/T+) E3 group ($p=0.06$
276 for IL-6, $p=0.09$ for IL-8) (Fig. 1C, D).

277 Cognitive decline over 24 months

278 Patients were then clinically followed over a period
279 of 24 months in our memory clinic. Results showed a
280 significant difference after 12 months in clinical pro-
281 gression (evaluated as delta MMSE scores respect

Table 2
CSF levels of cytokines and chemokines in healthy controls and patients divided by *APOE* genotype

	HC	(A+/T-) E3	(A+/T-) E4	(A+/T+) E3	(A+/T+) E4	<i>p</i>
IL-1beta	0.06 ± 0.01	0.07 ± 0.01	0.06 ± 0.04	0.09 ± 0.02	0.10 ± 0.05	0.697
IL-2	0.70 ± 0.09	0.61 ± 0.13	0.50 ± 0.17	0.56 ± 0.12	0.57 ± 0.13	0.898
IL-4	0.19 ± 0.02	0.21 ± 0.03	0.37 ± 0.06	0.21 ± 0.03	0.17 ± 0.02	0.004
IL-6	3.51 ± 0.33	4.69 ± 0.83	9.71 ± 2.08	5.32 ± 1.02	4.54 ± 0.97	0.002
IL-7	7.81 ± 0.98	9.70 ± 1.28	11.54 ± 1.94	6.90 ± 1.48	6.96 ± 2.31	0.175
IL-8	20.68 ± 0.85	20.82 ± 1.70	30.11 ± 3.27	23.86 ± 0.97	19.74 ± 1.29	<0.001
IL-10	2.87 ± 0.14	2.81 ± 0.07	2.78 ± 0.14	2.90 ± 0.12	3.07 ± 0.31	0.425
IL-12	1.21 ± 0.23	1.21 ± 0.25	1.76 ± 0.23	1.39 ± 0.25	1.26 ± 0.25	0.476
IL-13	1.99 ± 0.52	1.57 ± 0.26	1.63 ± 0.45	1.46 ± 0.27	1.92 ± 0.56	0.845
IL-17	1.96 ± 0.39	1.65 ± 0.38	1.88 ± 0.36	2.20 ± 0.39	2.07 ± 0.51	0.893
G-CSF	4.14 ± 0.45	5.74 ± 0.77	9.22 ± 1.21	5.011 ± 0.57	4.88 ± 0.83	<0.001
GM-CSF	40.77 ± 5.28	50.49 ± 5.96	51.99 ± 7.00	57.55 ± 5.94	46.48 ± 8.53	0.364
MCP-1	349.9 ± 25.8	320.1 ± 18.2	330.3 ± 27.2	311.5 ± 19.7	322.7 ± 22.7	0.447
MIP-1b	10.15 ± 0.58	9.88 ± 0.64	12.35 ± 1.68	12.55 ± 1.15	12.43 ± 1.02	0.152
TNFα	1.84 ± 0.21	1.52 ± 0.22	1.85 ± 0.39	1.93 ± 0.25	1.86 ± 0.24	0.800

HC, healthy controls; IL, interleukin; G-CSF, granulocyte colony-stimulating factor; GM-CSF, granulocyte-macrophage colony-stimulating factor; MCP-1, monocyte chemoattractant protein 1; MIP-1b, macrophage inflammatory proteins 1b; TNFα, tumor necrosis factor-α.

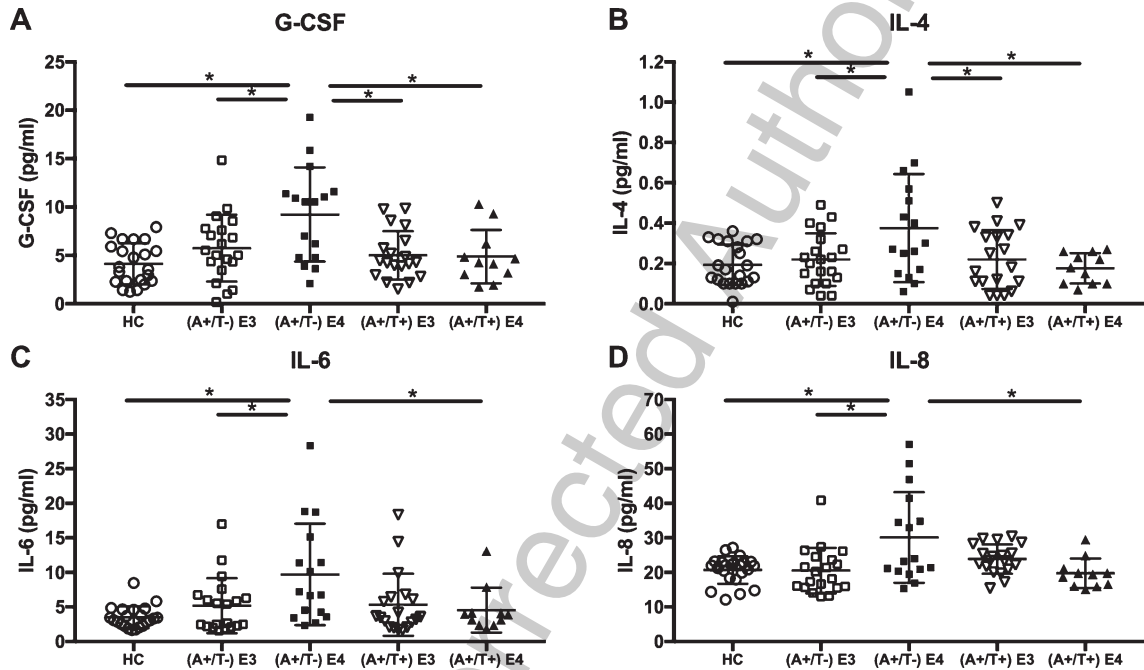


Fig. 1. Multiple comparisons of G-CSF (A), IL-4 (B), IL-6 (C), and IL-8 (D) CSF levels among controls and patients' groups according to *APOE* genotype. **p* < 0.05.

to baseline) for (A+/T-) E4 patients (0.86 ± 0.44), with respect to (A+/T+) E3 (-1.41 ± 0.61 ; $p = 0.048$) and to (A+/T+) E4 (-2.70 ± 0.90 ; $p = 0.005$), but not to (A+/T-) E3 patients (-0.72 ± 0.64 ; $p = 0.276$) (Fig. 2A). Similarly, at 24 months (A+/T-) E4 patients showed a stable MMSE (0.10 ± 0.35) respect to clinical progression showed by (A+/T+) E3 (-3.74 ± 0.91 ; $p = 0.006$), (A+/T+) E4 (-5.06 ± 1.48 ; $p = 0.001$) but not (A+/T-) E3 patients (-1.67 ± 0.61 ; $p = 0.400$). At 24 months we found also a statisti-

cally significant difference for clinical progression between (A+/T-) E3 and (A+/T+) E4 ($p = 0.044$) (Fig. 2B).

DISCUSSION

The adoption of the NIA-AA consensus guidelines associated with the *APOE* genotype allowed us to reveal surprising results on the role of neuroinflam-

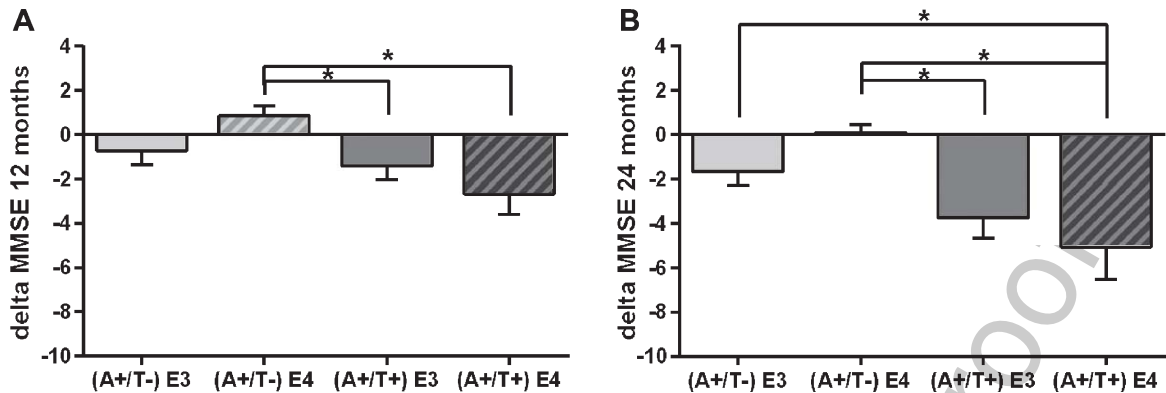


Fig. 2. Clinical progression evaluated as delta MMSE score at 12 and 24 months with respect to baseline. A) At 12 months, A+/T- patients showed substantial clinical stability. In particular, a significant difference was found between (A+/T-) E4 patients and both A+/T+ patients whatever the APOE genotype. B) At 24 months, (A+/T-) E4 patients still showed clinical stability as opposed to A+/T+ patients. Interestingly, (A+/T-) E3 patients showed a slight clinical worsening, with a significant difference respect to (A+/T+) E4 patients. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$.

299 mation in AD. Our study showed that CSF cytokines' levels in A+/T+ patients are similar to that of controls, regardless the APOE genotype. Moreover, we found that a specific pattern of AD related pathology, the (A+/T-) in APOE4 carriers was associated with significantly increased levels of CSF IL-4, IL-6, IL-8, and G-CSF. Cognitive decline progression of this subgroup of patients, measured over a period of 24 months, appeared significantly more preserved than that observed in the other groups. Our data lead us to suggest a relationship between APOE4 status and A β pathology in the absence of tau-related neurodegeneration possibly linked to a subset of cytokines exerting a protective action on the progression of cognitive symptoms.

314 ApoEs are lipoprotein produced and released by astrocytes, mainly involved in lipid transport to neurons and useful to support neuronal metabolism, synaptic plasticity, and neuronal repair in cases of injuries. In humans, APOE4 is the major risk factor for developing AD [26] and in healthy individuals is associated with a reduced A β clearance and to a potential development of pathological changes responsible for cognitive decline [27]. Experimental AD settings have shown that ApoE4 increases A β synthesis, reduces its clearance, and increase A β dependent apoptosis of neurons [28, 29]. Such conditions inevitably lead to increase the A β burden and favor the hampering of cortical neurotransmission. However, our data suggest that APOE4 genotype, associated with an isolated A β pathology, favors the synthesis and release of cytokines from astrocytes and microglial cells that could sustain

the physiological mechanisms of synaptic transmission, thus preserving from cognitive decline. Indeed, cytokines could have beneficial effects reducing A β burden and potentiating synaptic transmission. IL-6 is a pleiotropic cytokine able to influence synaptic functions through IL6R located on neurons [30] of glutamatergic [31], catecholaminergic and cholinergic transmission [32]. Through its interaction with excitatory pathways, IL-6 can participate to the clearance of A β peptides [31] reducing its presence at synapses. IL-4 is a cytokine with anti-inflammatory activity, influencing astrocytes in the synthesis of neurotrophic growth factors [33]. Moreover, in experimental settings, IL-4 has been shown to promote microglial clearance of A β oligomers [34]. IL-8 is a chemokine that protects neurons by both paracrine or autocrine loop [35]. In cases of A β pathology, IL-8 is able to inhibit A β -induced apoptosis and promotes synthesis and release of brain-derived neurotrophic factor protecting neuronal survival [36]. G-CSF is a growth factor involved in stimulation and maturation of blood cells. Besides that, G-CSF plays a key role in neurogenesis and differentiation during brain development and a direct influence on synaptic plasticity [37]. G-CSF can also influence neuronal activity through its receptors expressed in hippocampus and frontal cortices [38].

In this view, it is likely to suppose that ApoE4 and A β can interact positively in the synthesis of neuroprotective cytokines. This interaction is specific for APOE4, since APOE3 patients with isolated A β pathology did not show the same increase of CSF cytokines. This is likely because APOE3 is gener-

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ally associated per se with higher neuronal resilience and protection from the risk to develop AD pathology [39]. Interestingly, in A+/T+ patients we did not observe any change in cytokines' level, neither pro-inflammatory nor anti-inflammatory, regardless the APOE genotype. Thus, in case of association between A β and tau pathology the possible protective effect of both APOE4 and APOE3 [38, 40] is hampered. In particular, our data showed a more rapid cognitive decline in A+/T+ respect to A+/T- patients. We hypothesize that in presence of persistent noxious stressors (A β and tau protein) the increased synthesis of ApoE by neurons may induce a neuron-specific proteolytic pathway responsible for the production of ApoE fragments with neurotoxic effects, such as mitochondrial energy impairment, increase of tau phosphorylation and cytoskeletal disruption [29]. Moreover, among A+/T+ patients, APOE4 carriers did not exhibit higher levels of CSF cytokines, but rather a marked cognitive decline, even worse than that shown by APOE3 patients. Indeed, it is likely that apolipoprotein E3, with a specific binding site for tau, can be protective against the excess of tau phosphorylation, which is deleterious for neuronal survival [11, 41]. Conversely, apolipoprotein E4 does not have such binding site, and therefore patients are more exposed to neurodegeneration and cognitive decline [40].

In conclusion, even if previous evidence suggests a detrimental role of neuroinflammation in AD [42], our findings indicate that the specific condition of isolated amyloidosis (A+/T-) with APOE4 status is associated in the CNS to an increase level of cytokines able to support the physiological mechanisms of neurotransmission and to reduce the A β deposition [30–38], which in our patients is expressed by a significant cognitive preservation over a period of 24 months. However, in A+/T+ condition the upregulation of cytokines and chemokines is hampered regardless the APOE genotype, probably because in an advanced stage of neurodegeneration, neuroinflammation is no longer able to support synaptic functioning. In agreement with our results, Taipa and colleagues recently reported a significant correlation between elevated levels of proinflammatory cytokines in the CSF of patients with AD and the cognitive status, suggesting that a stronger inflammatory response leads to a better clinical progression [21]. These findings at a first glance may seem to be in conflict with previous literature, reporting a pathological chronic activation of the innate immune system, with altered production of cytokines [43, 44]

associated with the neurodegenerative processes of dementias. Intriguingly, these conflicting findings, as well our results, do nothing but reinforce the concept of neuroinflammation as a dynamic process that can act differently as a protective or harmful mechanism depending on the stage of disease and the genetic substrate (e.g., APOE).

Our study has some limitation, first of all the small sample size. Larger samples of patients and controls are needed to detect other significant difference in cytokines levels. Nevertheless, our study has the merit to measure cytokine's contents directly in the CSF of patients which have a robust diagnosis of AD pathology, supported by CSF biomarkers (A β , tau, p-tau), and long clinical follow-up periods.

In summary, although several studies suggest the modulation of pro-inflammatory cytokines production as a therapeutic target in AD [45, 46], the present work suggests that caution must be taken on modulate neuroinflammatory signaling to ensure that protective pathways are not compromised. Future studies are needed to disentangle the intricate role of neuroinflammation in AD to provide valuable cues for the development of more selective therapeutic strategies.

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