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1 **The effects of two Alzheimer’s related genes APOE and MAPT in**
2 **healthy young adults: An attentional blink study**

3
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24 **Abstract**

25 **Background** genetic risk factors start to affect the brain and behavior in Alzheimer’s Disease (AD) before
26 clinical symptoms occur. Although AD is mainly associated with memory deficits, attention and executive
27 dysfunctions can present at the early presymptomatic stages in middle age for those with non-modifiable
28 risks. Here, we investigated whether known risk genes for AD already affected attention in young adulthood.

29 **Methods** a total of 392 healthy young adults aged around 20 years underwent genetic testing for risks of
30 dementia (APOE and MAPT) and performed a computerized cognitive test for temporal attention called the
31 Attentional Blink (AB) task, in which patients with dementia tested in previous studies often showed reduced
32 performance. Here, the AB task was analyzed using repeated-measurements analysis of variance for the
33 ability of visual perception, attention deployment and temporal memory encoding/binding performance.

34 **Results** the results showed that all participants exhibited AB effects. Importantly, genetic risk factors had
35 statistically significant influence on temporal attention depending on sex in healthy young adults. APOE4
36 status was associated with enhanced attention deployment in males ($F(1, 124) = 8.285, P = .005, \eta^2 = .063$)
37 but not females, while MAPT AA carriers had poorer performance in AB but only in females ($F(1, 254) =$
38 $4.114, P = .044, \eta^2 = .016$). No genetic effects were found for visual perception and temporal memory binding
39 errors between high and low risk groups.

40 **Conclusions** We provided evidence that both APOE and MAPT start to affect attentional function as early
41 as young adulthood. Furthermore, unlike previous findings in older people, these genes had a differential
42 effect for males and females in young adults.

43

44 **Key words:** Alzheimer’s disease, attentional blink, APOE, MAPT, dementia, genetic risk factors

45 **Introduction**

46 Alzheimer's disease (*AD*) is an age-related neurodegenerative disease that is accompanied by cognitive
47 impairments such as memory loss, impairments of learning, and impairments in executive function in the
48 early stage [1], progressing to speech difficulties, and losing reading and writing skills in the latter stage [2].
49 As family history of AD increases the likelihood of developing the condition, it is believed that genetic risk
50 factors are key drivers of AD [3]. The main genetic risk factors of AD include Apolipoprotein E (*APOE*)- ϵ 4
51 allele [3-6], Microtubule-associated protein tau (*MAPT*) [6-9], and other genetic loci discovered in GWAS
52 studies [10]. Previous research on these genetic risk factors showed that people carrying risk genes were
53 more likely to develop dementia in later life and have shown differences in cognitive performance before
54 clinical symptoms occur and in middle and young age [11], so it is important to study how early the effects
55 of these Alzheimer's related genetic risk factors begin in cognitively healthy young adults.

56 *APOE* is a protein, which functions to transport lipid and cholesterol [12]. Among its three major alleles
57 (ϵ 2, ϵ 3, and ϵ 4), both homozygotes (ϵ 4/4) carriers and heterozygous carriers (ϵ 4/-) of ϵ 4 are regarded to have
58 a higher risk and earlier onset of AD than noncarriers [13]. The mechanism underlying this association
59 remains unclear, but evidence shows that it may be because *APOE* ϵ 4 carriers (denoted as *APOE*4) have an
60 impaired capability to remove amyloid beta ($A\beta$), a pathological hallmark of AD in the brain among other
61 neuropathological mechanisms [12].

62 As another genetic risk factor of AD, the *MAPT* gene encodes the tau protein, a protein that is associated
63 with stabilizing and assembling microtubules. Containing hyperphosphorylated tau protein, neurofibrillary
64 tangles (NFTs) within neurons are regarded as another hallmark of AD besides $A\beta$ [14]. *MAPT* has two
65 haplotypes (H1 and H2) [15], and the H1c sub-haplotype [16]. Several single nucleotide polymorphisms
66 (SNP) contained in H1 haplotype, such as rs242557 AA genotypes, are considered to increase the risk of AD

67 [17].

68 Previous studies have found that these two genetic risk factors are linked with worse cognitive
69 functioning in healthy older individuals. APOE ϵ 4 allele was associated with poorer executive function [18],
70 an increased risk of mild cognitive impairment [19] and faster cognitive decline [19-22] in older people.
71 Interestingly, in younger healthy individuals, some emerging research has found cognitive advantage
72 associated with episodic memory [23], verbal fluency [24], learning [25], and attention [4] in young APOE4
73 carriers. In this respect, an “antagonistic pleiotropy” hypothesis was proposed to explain the apparent
74 “beneficial” effect of APOE4 on cognition in early life and the detrimental effect in later life [26]. However,
75 such advantage of APOE4 on cognition in early life was not found in some other studies [27-30]. This
76 inconsistency may result from the different tasks used in these studies suggesting that the effects of APOE4
77 in early adulthood may vary among different cognitive domains and APOE also interacts with other genetic
78 and environmental factors. So, as discussed later, in our study, we have chosen a robust psychophysics task
79 that can be sensitive to subtle effects of genes.

80 Less studied than APOE, the relationship between MAPT and cognitive performance is also mixed. In
81 an experiment carried out in 350 adults with progressive supranuclear palsy (PSP), the rs242557 A carriers
82 performed better than noncarriers in general cognitive function, executive function, and attention [31]; other
83 research found that H1 homozygosity carriers performed worse in a picture memory task than H2 carriers
84 [32], but a study on 1191 patients with Parkinson’s Disease (PD) found no association between H1 haplotype
85 and the psychometric results [33]. How MAPT relates to cognition in middle age and young adults remains
86 understudied.

87 Although the effects of Alzheimer’s related genetics risk factors such as APOE and MAPT are
88 extensively studied in older patients and controls, their effects on cognitive functioning in younger healthy

89 individuals remains unclear. Also, although memory loss was regarded as the earliest cognitive impairment
90 in AD, and attention and executive dysfunctions being associated with other subtypes of dementia such as
91 dementia with Lewy bodies (DLB), evidence has shown that attention deficits were also an early problem in
92 AD [34, 35]. Some also argued that early memory issues in patients with AD could also be due to poorer
93 attention [36]. Now, it is widely recognized that neuropathology and brain structural and functional changes
94 begin decades before clinical symptoms of AD occur [37]. Early-stage AD provides the best window of
95 opportunity for intervention. However, how genetic risk factors of AD affect key cognitive domains such as
96 attention and executive functions in healthy middle-age or young individuals remain inconclusive. The
97 current study is an extension of the PREVENT-dementia project [38] aimed at establishing a cross lifespan
98 knowledge base for genetic and environmental risk factors of dementia. Previously, we have shown in the
99 same cohort that young APOE4 and MAPT AA carriers have reduced functional connectivity in the default
100 mode network and thinner cortex in areas (temporal and frontal cortices) that typically develop AD pathology
101 including tau [39, 40]. Although research in the field including our own studies have uncovered striking
102 changes in brain cortical morphology and functional connectivity as early as young adulthood and middle-
103 age, sensitive measures of early cognitive changes due to genetic risks of AD is still lacking.

104 Hence, in the present study, we aimed to explore the effects of genetic variants (APOE and MAPT
105 rs242557) in young healthy individuals performing a well-established task called the attentional blink (AB)
106 task [41]. In the AB task, people are asked to detect two targets among a series of distractors in a rapid serial
107 visual presentation (RSVP) stream. In general, people show a deficit in reporting the second target if it
108 appears within 100-500ms after the first target [41]. One theoretical explanation for the AB deficit is the
109 competition of attentional resources between two targets [42], and this task was used to test people's ability
110 to allocate attention and temporally integrate episodic information in temporal processing [43]. In addition,

111 the AB task has been used to detect the cognitive deficit of patients suffering from neurodegenerative diseases.
112 In a previous study, people with PD showed a different pattern of errors from healthy counterparts in the AB
113 task [44]. It is also shown that AB performance was significantly impaired in both AD and DLB although
114 patients with DLB had a more extended and severer reduction in performance than AD, i.e., deeper and longer
115 blinks [45]. However, it is unknown whether healthy young people with genetic risk factors of AD can also
116 show a different performance in AB even, as previous studies may suggest, a ‘beneficial’ effect of these risk
117 factors in younger people.

118 As discussed previously, existing evidence is inconsistent about whether and how genetic risks impact
119 attention in neurodegenerative conditions in young adulthood, although popular views such as the so called
120 ‘the associated antagonistic pleiotropy hypothesis exist [46]. The mixed findings in the literature were partly
121 due to different cognitive tests used in these studies because traditional pen and paper based clinical tools are
122 unlikely to be sensitive to young healthy individuals due to ‘ceiling effects’, and computerized cognitive tests
123 also suffer from many limiting factors such as biases caused by cognitive strategy, level of education,
124 confounding sensory perception functions, cultural and other methodological variations. The attentional blink
125 task chosen here however offers an ideal paradigm to examine cognitive functioning in young people because
126 it is a challenging and robust test that is minimally affected by mental strategy, practice effects, linguistic
127 knowledge and can be easily automated on computers [41]. Previous studies have shown a profound
128 impairment in the AB and rapid visual tests across multiple dementia subtypes [45], as well as several
129 developmental disorders in young people [47, 48].

130 According to the results from previous studies, we hypothesized that APOE4 carriers would have a
131 reduced AB deficit than noncarriers, and for the MAPT (rs242557), people with AA genotypes (the high-risk
132 variant) would have an increased AB deficit than those with G genotype.

133

134 **Method**

135 **Participants**

136 From Chinese students studying undergraduate degrees at the Southwest university in Chongqing,
137 participants from both sexes were recruited using online and offline advertisements on campus. People with
138 severe neurological and psychiatric conditions were excluded. We also excluded anyone majoring in sports,
139 arts, or music because these subjects have a different university entrance requirements from other subjects.
140 The above inclusion / exclusion criteria enhanced the cognitive homogeneity of the participants due to the
141 selection processes and the standard entrance requirements of the university. All participants have given
142 written informed consent and received compensation for their participation. This study was approved by the
143 Ethic Committee of Psychological Research at Southwest University.

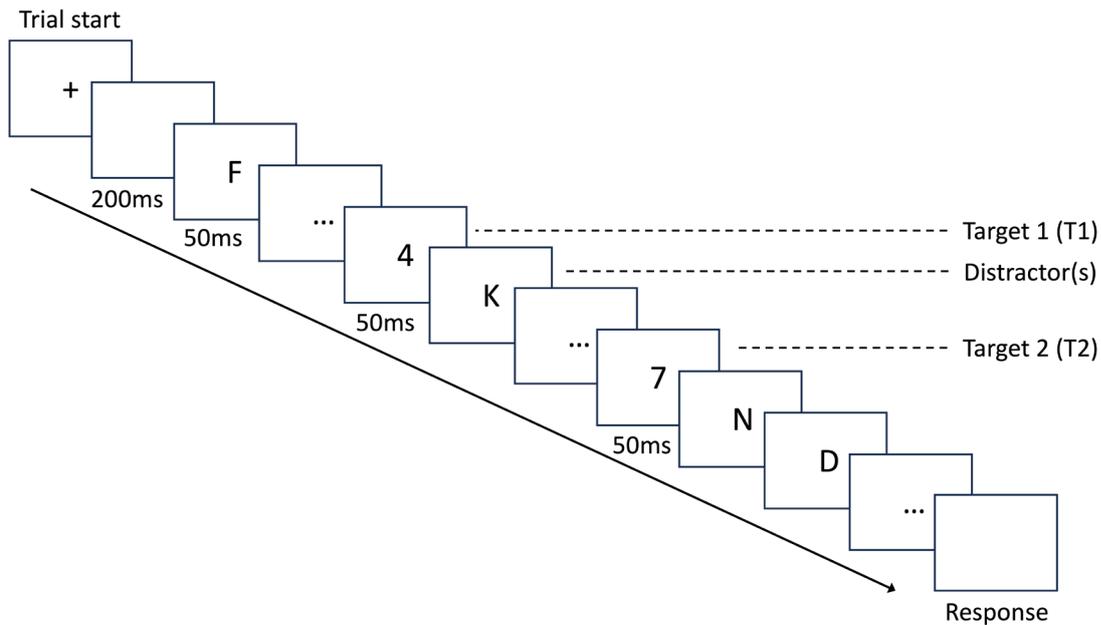
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145 **Procedure**

146 The attentional blink task was implemented in E-Prime 2.0. A typical trial in the AB task is shown in
147 Figure. 1. In each trial, the participant firstly pressed any key on the keyboard to start the trial. They then
148 saw a blank (grey) screen for 200 ms followed by a series of letters at the centre of the screen presented in
149 random order. Among the stream of letters (A, B, C, D, F, H, J, K, L, N, P, R, T, V, X, Y), there were two
150 one-digit numbers (2, 3, 4, 5, 6, 7, 8, 9) positioned pseudo randomly in the RSVP stream. Each letter or
151 digit appeared on the screen for only 50 ms, and the inter-stimulus interval was set to zero. Thus, the
152 stimulus onset asynchrony (SOA) was 50 ms. In each trial, there were 27 letters as distractors and 2 one-
153 digit numbers as targets (represented as Target 1 and Target 2 or T1 and T2 for short). At the end of the
154 trial, the participant was asked to respond by typing in the identities of the two target numbers, in the order

155 they saw them, and to ignore the distractors (letters). To avoid visual confusion about the physical forms of
156 the stimuli that are not related to attentional processing, one-digit numbers resembling letters (e.g., 0, 1)
157 and letters resembling numbers (e.g., I, O, Q and S) were not used.

158



159

160 **Figure. 1** A schematic diagram of a typical trial. Each letter and number were presented for 50
161 milliseconds, with no inter-stimulus interval. T1 appeared at the 6th to 12th positions from the beginning of
162 the trial. T2 was positioned after T1 at 1, 2, 3, 4, 5, 6, 8, 10, or 12 lags. Lag 1 means no intervening
163 distractor between T1 and T2; lag 2 means 1 intervening distractor between T1 and T2; and so on.

164

165 T1 appeared randomly at the 6th to 12th positions from the beginning of the stream. T2 was positioned
166 after T1 by 1, 2, 3, 4, 5, 6, 8, 10, or 12 positions (i.e., the lag between T1 and T2 was 1, 2, 3..., or 12).
167 Combining the different positions of T1 and T2, there were a total of 63 trial types. The computer presented
168 the 63 trial types twice (two cycles) giving 126 trials. Within each cycle, the 63 trials were randomized for
169 each participant. After finishing 50 trials, participants had a break of proximately one minute.

170 Before the start of the actual experiment, the participants received instruction and two practice runs,
171 each of which contained four trials. In the first practice run, a trial was the same as the trial in the actual
172 experiment in all but the following aspects. Each letter or number appeared for a longer duration of 500 ms.
173 Participants were informed that the speed of presentation was slowed down during practice. After each
174 participant's response, feedback of correct (✓) or incorrect (×) was provided on the screen. If the participant
175 could not correctly respond to all 4 trials, they had to repeat the instruction and the first practice run. After
176 the first practice run, a second practice run was conducted, but this time, the stimulus duration was the same
177 as the actual experiment.

178

179 **Genotyping**

180 This study investigated APOE and MAPT rs242557, and the former was determined by rs7412 and
181 rs42935. The genotypes were performed by a Mass Array system (Agena iPLEX assay, San Diego, United
182 States). Approximately 10-20ng of genomic DNA was isolated from saliva samples. The sample DNA was
183 amplified by a multiplex Polymerase Chain Reaction (PCR), then the obtained products were used for locus-
184 specific single-base extension reaction. Finally, the resulting products were desalted and transferred to a 384-
185 element SpectroCHIP array. The alleles were discriminated by mass spectrometry (Agena, San Diego, United
186 States).

187 For rs7412 the PCR primers were ACGTTGGATGGCCCCGGCCTGGTACTG and
188 ACGTTGGATGACCTGCGCAAGCTGCGTAAG; the unextended primer was
189 CCGCTGCCGATGACCTGCAGAAG. For rs429358, the PCR primers were
190 ACGTTGGATGTCGCCGGTACTGCACCA and ACGTTGGATGCTGTCCAAGGAGCTGCAGG; the
191 unextended primer was GACATGGAGGACGTG. For rs242557, the PCR primers were

192 ACGTTGGATGAGACCCTGTGAGATCATCCC and ACGTTGGATGTACAAAGCAGTTGGCTTCGC;
193 the unextended primer was CCATCAGTTGGCTTCGCCAGGGT.

194 Based on genotypes of rs7412 and rs429358, carriers of $\epsilon 4/\epsilon 4$ or $\epsilon 4/\epsilon 3$ were combined into a single
195 APOE4 group. Given that AA was the high-risk genotype at MAPT rs242557, we have categorized rs242557
196 into AA or G carriers in the subsequent analyses. We have also applied Hardy-Weinberg equilibrium tests for
197 each gene in our sample population.

198

199 **Statistical methods**

200 Outcome measures used in the current study included 1) T1 accuracy – the probability of detecting T1
201 primarily reflects perceptual functioning; 2) temporal attentional deployment – a conditional probability of
202 detecting T2 given T1 was detected correctly. So, the performance of temporal attention was calculated as a
203 conditional probability of $P(T2|T1)$; 3) AB index - the magnitude of the AB effect (named as AB index here)
204 by calculating the differences between $P(T2|T1)$ measured at lag 4 (where the performance is lowest when
205 SOA is 50ms) and at lag 12 (where the blink has generally recovered). The AB index is a more sensitive way
206 to measure AB effects, and please note that a larger AB index represent a poorer AB performance; and 4) the
207 swap ratio – reporting both targets identities correctly but in the incorrect order. This represents a deficit in
208 encoding temporal information and binding errors [43].

209 For each group comparison, we firstly performed a repeated-measurements analysis of variance
210 (ANOVA), in which the dependent variable was either the T1 accuracy, $P(T2|T1)$ for attention, AB index, or
211 swap ratio as described above. The independent variables in the ANOVA were the genotype of APOE4 or
212 MAPT, sex, and lag (lag is a within-subject factor while others are between-subject factors). The covariate
213 was centered age (age – averaged age). Then, we have performed secondary group ANOVA tests for each

214 sex separately. This time, the variables were unchanged, except that we removed sex in the ANOVA model.
215 Greenhouse-Geisser corrections (GG corrections for short) were applied to those tests where non-sphericity
216 issues were present.

217 Importantly, since our sample size is large, the data degrees of freedom are large. Such high data degrees
218 of freedom can at least in part drive a significant finding [49]. Accordingly, we will report partial eta squared
219 effect sizes throughout, which are not impacted by sample size in this way [49] and should be accurate when
220 sample size is large.

221

222 **Results**

223 **Genetic tests**

224 Out of 392 participants, 385 participants were successfully genotyped for rs7412. Genotype CC N=344
225 (228 females), CT N=40 (27 females), and TT N=1 (1 females). Three tests of Hardy-Weinberg equilibrium
226 were performed for all, male, and female participants respectively ($\chi^2 = 0.02, 0.04, \text{ and } 0.36$ respectively,
227 and all $P's > 0.05$). So, the genotype frequencies of rs7412 in our samples were consistent with Hardy-
228 Weinberg equilibrium.

229 Out of 392 participants, 390 participants were successfully genotyped for rs429358. Genotype CC N=1
230 person (0 females), CT N=70 (48 females), and TT N=319 persons (213 females). Three tests of Hardy-
231 Weinberg equilibrium were performed ($\chi^2 = 1.97, 2.68, \text{ and } 0.01$ respectively, and all $P's > 0.05$). So, the
232 genotype frequencies of rs429358 in our samples were consistent with Hardy-Weinberg equilibrium.

233 Out of 392 participants, 386 participants were successfully genotyped for rs242557, with genotype AA
234 N=121 (77 females), AG N=193 (136 females), and GG N=72 (45 females). Three tests of Hardy-Weinberg
235 equilibrium were performed for all three genetic groups according to MAPT ($\chi^2 = 0.10, 1.29, \text{ and } 1.12$

236 respectively, and all P 's > 0.05). So, the genotype frequencies of rs242557 in our samples were consistent
 237 with Hardy-Weinberg equilibrium.

238

239 **Demographics**

240 In the total of 392 participants, 261 are females. The mean and standard deviation of the participants'
 241 ages were 19.71 and 0.98 years respectively. The demographic data divided into each genetic group is shown
 242 in Table 1. We found no statistically significant difference in age, sex, and years of education between
 243 different risk groups defined by APOE and MAPT status.

244

Table 1. Demographics by Genotype

Category	APOE4		rs242557	
	Non-carrier	Carrier	G carrier	AA
Gender				
Male	108	20	84	44
Female	214	42	181	77
Education				
U1	174	33	143	66
U2	124	24	101	48
U3	14	3	11	5
U4	10	2	10	2
Age				
Mean	19.67	19.72	19.71	19.71
SD	0.97	0.99	0.99	0.97

Note. U1 to U4: 1st to 4th year undergraduates. Age was calculated as (participating date - birth date)/365.25.

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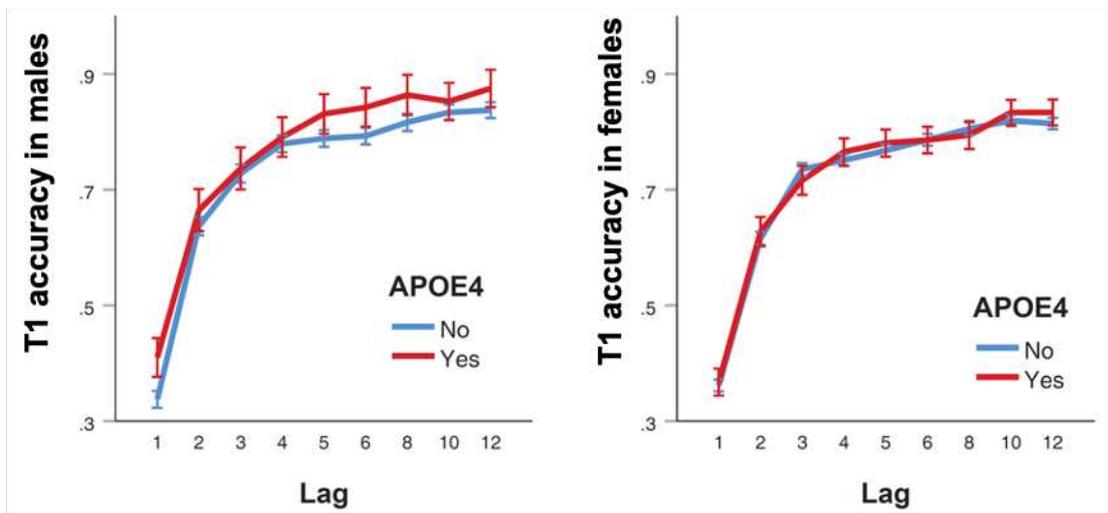
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247 **APOE effects on T1**

248 In the ANOVA test, the main effect of lag was significant: $F(8, 3023) = 306.844, P < .001, \eta^2 = .447$
 249 [after GG correction, $F(7.101, 2691.461) = 306.844, P < .001, \eta^2 = .447$]. No other main effects or

250 interactions were significant. The results remained the same after splitting the males and females into two
 251 separate groups in the secondary ANOVA tests. See Figure. 2 for more details. We noted that the T1 accuracy
 252 was generally low at lag-1 due to the short SOA of 50ms, but this finding is consistent with previous research
 253 [50]. Importantly, there was no statistically significant effect of APOE genotype on visual perception
 254 measured by the AB task.

255



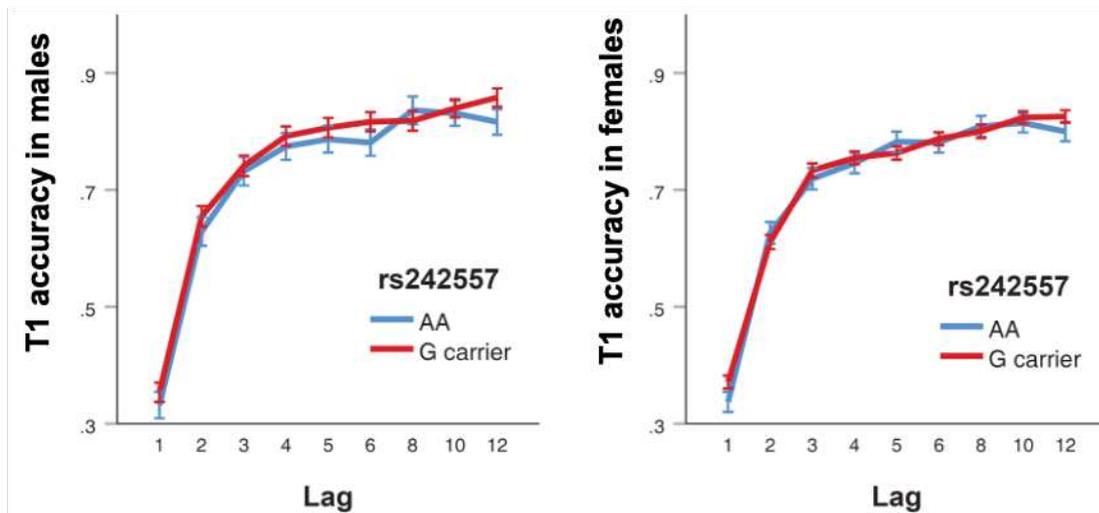
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257 **Figure. 2** The effects of APOE4 on T1 accuracy for males (left) and females (right) respectively. Error bars
 258 represent standard errors. X-axis represents lag and Y-axis represents probability of detecting T1. We found
 259 no group difference for APOE status suggesting that there was no significant effect of APOE on T1 accuracy.

260

261 **MAPT effects on T1**

262 In this ANOVA test, the main effect of lag was also significant: $F(8, 3048) = 534.303, P < .001, \eta^2 = .584$
 263 [after GG correction, $F(7.112, 2709.828) = 534.303, P < .001, \eta^2 = .584$]. No other main effects nor
 264 interactions were significant. The results remained the same after splitting the males and females into two
 265 separate analyses. See Figure. 3 for more details. Like APOE, MAPT also did not change perception
 266 functioning in this task.



268

269 **Figure. 3** The effects of MAPT (rs242557) on T1 accuracy for males (left) and females (right) respectively.

270 Error bars represent standard errors. X-axis represents lag and Y-axis represents probability of detecting T1.

271 Similar to the previous results in APOE, MAPT also did not change perception functioning in this task.

272

273 APOE effects on temporal attention

274 Unsurprisingly, in the ANOVA test, the main effect of lag was statistically significant: $F(8, 3016) =$

275 $47.543, P < .001, \eta^2 = .112$ [after GG correction, $F(6.313, 2380.005) = 47.543, P < .001, \eta^2 = .112$], suggesting

276 that there was an AB effect as expected. We found that the main effect of APOE4 was statistically significant:

277 $F(1, 377) = 10.545, P = .001, \eta^2 = .027$. The interaction of APOE4 and sex only showed an insignificant

278 trend: $F(1, 377) = 3.415, P = .065, \eta^2 = .009$. The interaction of APOE4, sex and lag was significant: $F(8,$

279 $3016) = 2.454, P = .012, \eta^2 = .006$ [after GG correction, $F(6.313, 2380.005) = 2.454, P = .021, \eta^2 = .006$].

280 No other main effect or interaction was significant.

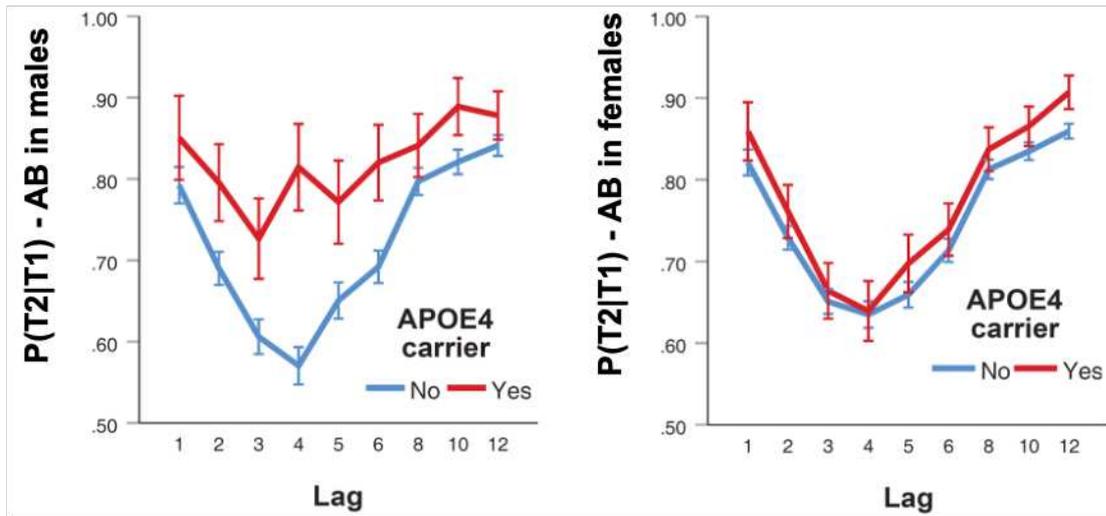
281 In the secondary ANOVA tests, for males, the main effect of APOE4 was significant: $F(1, 124) = 8.285,$

282 $P = .005, \eta^2 = .063$. The interaction between APOE4 and lag was significant: $F(8, 992) = 2.907, P = .003, \eta^2$

283 $= .023$ [after GG correction, $F(6.267, 777.113) = 2.907, P = .007, \eta^2 = .023$]. The main effect of lag was also

284 significant. For females, the main effect of lag was significant, but no other main effect or interaction was
 285 statistically significant. As shown in Figure. 4, the APOE4 effect on attention deployment was only
 286 significant in male participants.

287



288

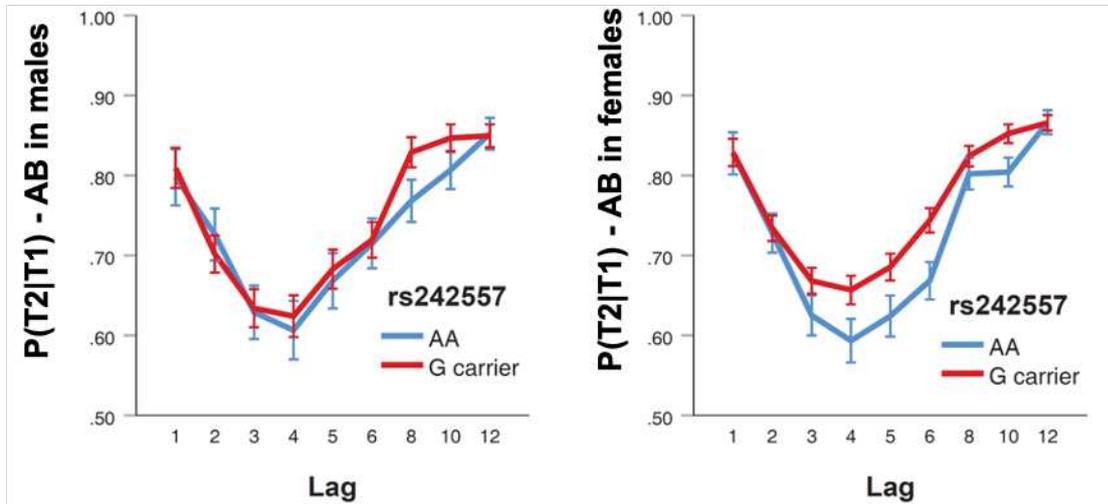
289 **Figure. 4** The effects of APOE4 on T2 accuracy conditional on T1 for males (left) and females (right)
 290 respectively. Error bars represent standard errors. X-axis represents lag and Y-axis represents conditional
 291 probability of detecting T2 condition on T1. Male but not female APOE4 carriers had better AB
 292 performance, i.e., carriers were more likely to report T2 correctly after correctly reporting T1 than noncarriers.

293

294 **MAPT effects on temporal attention**

295 In this ANOVA test, the main effect of lag was significant: $F(8, 3032) = 93.272, P < .001, \eta^2 = .197$
 296 [after GG correction, $F(6.327, 2397.845) = 93.272, P < .001, \eta^2 = .197$]. No other main effect or interaction
 297 was significant. In addition, in the secondary ANOVA tests, for males, the main effect of lag was significant,
 298 but no other main effect or interaction was significant. For females, the main effect of MAPT rs242557 was
 299 significant: $F(1, 254) = 4.114, P = .044, \eta^2 = .016$. The interaction between MAPT rs242557 and lag showed
 300 an insignificant trend: $F(8, 2032) = 1.817, P = .069, \eta^2 = .007$ [after GG correction, $F(6.260, 1590.097) =$

301 1.817, $P = .089$, $\eta^2 = .007$]. The main effect of lag was significant, but no other main effect or interaction was
 302 significant. As shown in Figure. 5, MAPT only significantly affected female participants in this task.
 303



304 **Figure. 5** The effects of MAPT (rs242557) on T2 accuracy for males (left) and females (right) respectively.
 305 Error bars represent standard errors. X-axis represents lag and Y-axis represents conditional probability of
 306 detecting T2 conditional on T1. Female but not male MAPT AA carriers had poorer AB performance than
 307 MAPT G carriers.
 308

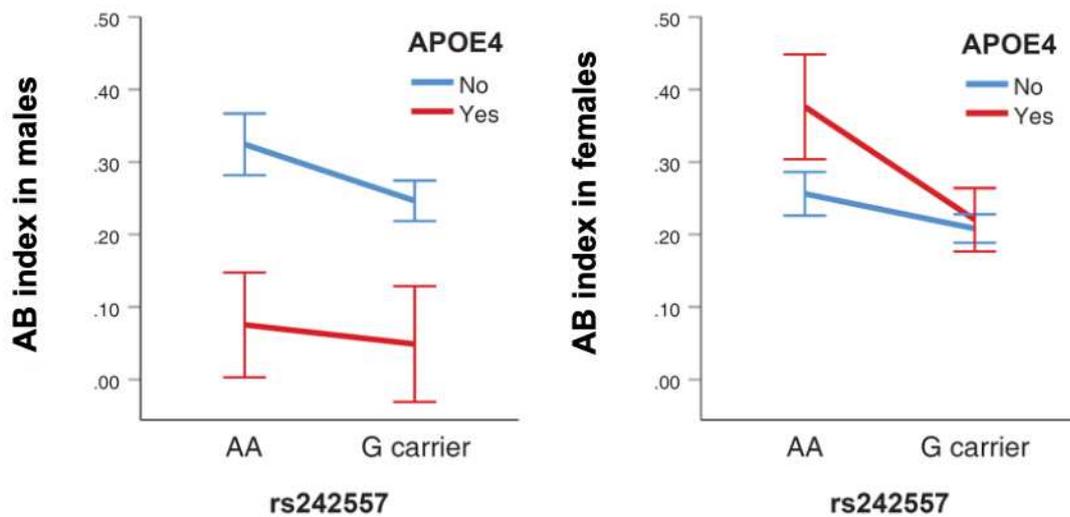
309
 310 **Interaction between APOE and MAPT on the AB index**

311 We reanalysed the attention effect using the AB index. As previously discussed, it is more sensitive and
 312 powerful to test the interaction between two genetic risks. In this ANOVA test (lag is no longer an independent
 313 variable), the main effect of sex was significant: $F(1, 370) = 5.902$, $P = .016$, $\eta^2 = .016$. The main effect of
 314 APOE4 was significant: $F(1, 370) = 4.370$, $P = .037$, $\eta^2 = .012$. The main effect of MAPT rs242557 was
 315 significant: $F(1, 370) = 4.201$, $P = .041$, $\eta^2 = .011$. The interaction among APOE4 and sex was statistically
 316 significant: $F(1, 370) = 14.813$, $P < .001$, $\eta^2 = .038$.

317 In the secondary ANOVA tests, for males, the main effect of APOE4 was significant: $F(1, 121) = 15.495$,

318 $P < .001$, $\eta^2 = .114$. No other main effect or interaction was significant. For females, the main effect of MAPT
 319 rs242557 was significant: $F(1, 248) = 4.700$, $P = .031$, $\eta^2 = .019$. No other main effect or interaction was
 320 significant. As shown in Figure. 6, findings in this more sensitive analysis were consistent with those from
 321 the two previous analyses although no gene-gene interaction was found for the AB index.

322



323

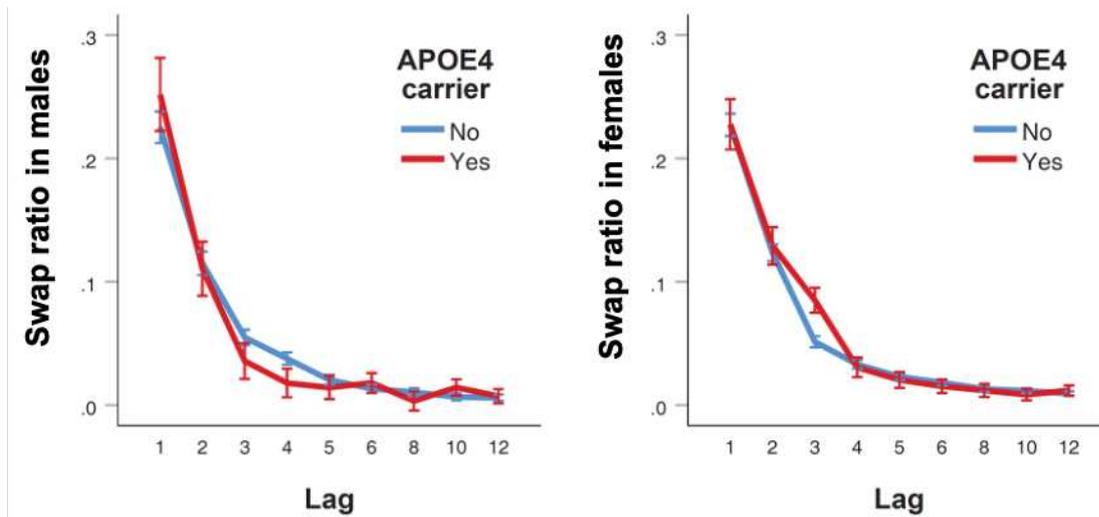
324 **Figure. 6** The interaction effects of APOE4 and MPAT (rs242557) on attentional blink index for males (left)
 325 and females (right) respectively. Error bars represent standard errors. Y-axis represents the difference between
 326 conditional probabilities of detecting T2 conditional on T1 at lag 4 and lag 12. We found no APOE-MAPT
 327 interaction.

328

329 APOE effects on swaps

330 In the ANOVA test, the main effect of lag was significant: $F(8, 3032) = 250.988$, $P < .001$, $\eta^2 = .398$
 331 [after GG correction, $F(3.437, 1302.522) = 250.988$, $P < .001$, $\eta^2 = .398$] but no other main effects nor
 332 interactions were significant. See Figure. 7 for more details.

333



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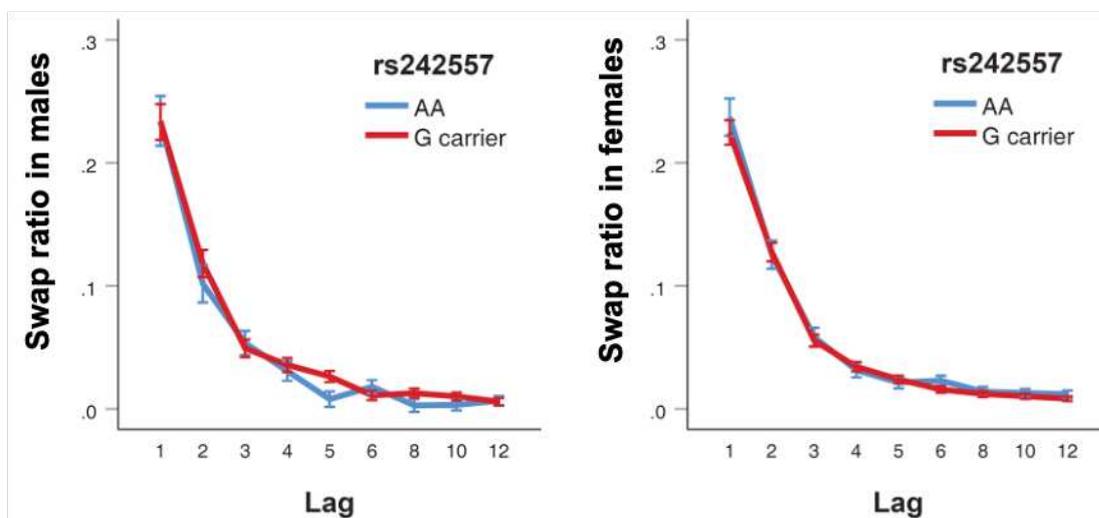
335 **Figure. 7** The effects of APOE4 on swap ratio for males (left) and females (right) respectively. Error bars
 336 represent standard errors. X-axis represents lag and Y-axis represents probability of reporting T1 and T2 in
 337 the incorrect order.

338

339 **MAPT effects on swaps**

340 Like APOE, in this ANOVA test, the main effect of lag was also significant: $F(8, 3048) = 399.328, P$
 341 $< .001, \eta^2 = .512$ [after GG correction, $F(3.420, 1302.915) = 399.328, P < .001, \eta^2 = .512$], but no other main
 342 effects nor interactions were significant. See Figure. 8 for more details.

343



344

345 **Figure. 8** The effects of MAPT (rs242557) on swap ratio for males (left) and females (right) respectively.
346 Error bars represent standard errors. X-axis represents lag and Y-axis represents probability of reporting T1
347 and T2 in the incorrect order.

348

349 **Discussion**

350 In our study, we found a significant reduction in the AB effect or improved temporal attentional
351 performance for APOE4 carriers, which is consistent with previous evidence of the antagonistic pleiotropy
352 hypothesis supporting the improved attentional function in young adults [4]. However, we only found this
353 beneficial effect in males but not in females. In most previous studies on older people, researchers found that
354 APOE4 confers a greater risk of dementia and mild cognitive impairment in females than in males [51-53].
355 Opposite to our results, in a study on executive function and processing speed, the cognitively beneficial
356 effect of APOE4 was found in young females rather than males [54]. However, a male-specific advantage of
357 APOE4 on short-term memory was found in middle-aged individuals in another study [55]. In one of our
358 own studies, sex differences in path integration and spatial navigation ability were found in middle-aged at-
359 risk individuals without cognitive impairment [56]. We speculate that sex differences in neurotransmitter
360 systems critical to attentional processing, such as acetylcholine (ACh) in the brain, where females have overall
361 higher release than males, and different experimental methodologies discussed earlier may also play a role
362 on these inconsistent sex differences [57].

363 For the other genetic risk factor of AD, MAPT rs242557 AA genotype, we found an increased AB effect
364 or poorer attentional function in carriers but this time only in females. To the best of our knowledge, there is
365 no known sex-specific cognitive bias associated with MAPT rs242557 genotype in the AB task, and as
366 previously discussed, the effect of rs242557 genotype in general cognitive function, executive function, and

367 attention in older adults is inconclusive in the literature too. In addition, although not statistically significant,
368 there seems a nonsignificant trend suggesting an interaction between APOE and MAPT in AB especially in
369 females (Figure. 6), i.e., carriers of both APOE4 and MAPT AA had the deepest blink in AB (larger AB index)
370 suggesting a stronger interference between T1 and T2 or more limited attentional resources to correctly
371 encode both targets. However, this finding should be interpreted with caution. Finally, we did not see any
372 genetic effect on T1 accuracy or swaps (perceptual functioning and temporal memory encoding/binding),
373 which was reported in older patients with dementia [45], suggesting that the early effects of these genetic
374 risk factors are not merely due to limited cognitive resource but a variation in specific attentional control
375 mechanisms.

376 We of course acknowledge that resource limitation is not the only possible explanation of the attentional
377 blink phenomenon. The Simultaneous Type - Serial Token (STST) model [58] is a prominent theory of the
378 AB, which has explained a large number of AB and related phenomena [59-61]. The theory also offers the
379 possibility to provide a theoretical explanation of our findings. In the STST framework, a key driver of blink
380 depth is the time to encode the T1. Specifically, the attentional blink is postulated to be caused by the process
381 of encoding T1 “closing the gate” for working memory encoding on T2 (by withholding a transient attentional
382 enhancement called the *Blaster*). This gate closing ensures that the encoding of T1 is not “contaminated” by
383 the T2 entering that same encoding process, which would yield binding errors, such as features from T2
384 migrating to T1. Thus, the theory raises the possibility that a male APOE4 carrier encodes stimuli more
385 rapidly into working memory than a male noncarrier, and similarly for a female G-carrier. Of course, such a
386 hypothesis needs further investigation in other experimental paradigms, before it can be seen as more than
387 speculative.

388 The current study was limited by its cross-sectional design. Thus, it is unable to reveal the dynamic

389 changes of attentional function transitioning from early adulthood to midlife, during which distinctive
390 cognitive impairments may start to occur. As age is one of the biggest risk factors of AD, and it is known that
391 visual attention task performance is often age dependent [62, 63], future longitudinal studies will be able to
392 address this limitation. The current study samples were heavily skewed towards participants with younger
393 age because they present the majority in the university. Thus, future research should include older adults
394 including those in middle age. Another limitation of the study was using a single test, the AB, although as
395 previously discussed, it is a robust test for attention, nonetheless, sensitive and specific screening tests based
396 on other visual tasks [64] and navigation tasks [56] have provided novel candidate tools for early detection
397 of dementia and practical solutions for screening in multiple clinical and community settings. As evidence
398 points towards AD pathology occurring before the symptom onset, this raises the question of whether these
399 pathological changes can be detected in this pre-symptomatic stage. With the development of novel disease-
400 modifying treatments for AD, such early detection would be highly advantageous by allowing for treatment
401 to begin sooner before more irreversible neurodegeneration occurs thus improving outcomes. Additionally,
402 detection of AD and its risk factors related changes earlier in life would allow individuals to make healthier
403 lifestyle choices in order to reduce their risk of developing AD [65], so detection in early adulthood may also
404 be beneficial. Accordingly, it is predicted that early detection and intervention could drastically reduce the
405 worldwide AD burden, with models suggesting that delaying disease onset by as little as 1 year could reduce
406 the global incidence of AD by up to 9.2 million in the year 2050 [66].

407 This study is also limited by the candidate gene approach with limited number of SNPs genotyped. This
408 may miss other critical genetic factors influencing attention such as the tonic level of neurotransmitters
409 including cholinergic systems. Moreover, only Chinese students were recruited into the study. Although this
410 increased genetic similarity among participants, it also limited the generalizability of the our findings to other

411 ethnic groups. In conclusion, we have found an ultra-early effect of AD related genetic risk factors (APOE
412 and MAPT) that are linked to two cornerstones of AD pathology (amyloid-beta and tau) using a temporal
413 attention task. The findings are generally consistent with previous research suggesting a 'beneficial' effect of
414 APOE4 on attention and executive control, but only in males. The less studied gene MAPT however had an
415 opposite effect than APOE in females. Finally, the study also highlighted sex differences in genetic risks as
416 an important topic for future research.

417

418 **Authors' contribution**

419 JZ and LS designed the study and secured the funding. JZ did the programming. JZ, XH and HQ collected
420 the data. JZ conducted the analyses. JZ, ZG and LS drafted the manuscript. CR, LS and JZ co-designed the
421 PREVENT-Dementia study. XX, YL and DZ provided feedback on the data analysis and the manuscript. YH,
422 HB, JOB provided feedback on the manuscript. LS provided oversight for the study.

423

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430

431 **Conflict of Interest**

432 The authors have no conflict of interest to report.

433

434 **Data Availability**

435 The data supporting the findings of this study are available on request from the corresponding authors. The
436 data are not publicly available due to privacy or ethical restrictions.

437

438 **Ethics approval**

439 This study was approved by the Ethic Committee of Psychological Research at Southwest University.

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