

Associations between lifestyle, genetic risk for Alzheimer's disease, and longitudinal brain atrophy in UK Biobank (N=2214)

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18 **Abstract**

19 INTRODUCTION: Modifiable behaviours such as smoking, diet, alcohol intake,
20 sedentariness, and physical inactivity are associated with brain health, but longitudinal
21 evidence in adults without dementia is limited.

22 METHODS: Among 2214 UK Biobank participants with two MRI scans over 2.2 years,
23 unweighted lifestyle scores (ranging 0 - 10) were derived from five behaviours and
24 classified as unfavourable, moderate, or favourable. Linear regression examined
25 associations, and *APOE* e4 genotypic interactions, with changes in 16 brain volumetric
26 markers.

27 RESULTS: Compared with favourable lifestyle, unfavourable lifestyle showed
28 significantly greater grey matter (2059 mm³; 0.046 standard deviations [SDs]) and left
29 hippocampal (36.5 mm³; 0.044 SD) atrophy. Moderate-risk drinking was linked to
30 higher total brain atrophy (2075 mm³; 0.063 SD), while high-risk drinking and low
31 physical activity associated with accelerated regional grey matter loss. No effects were
32 found for smoking, diet, or sedentariness. High-risk drinking was associated with
33 greater frontal lobe atrophy, particularly among *APOE* ε4 carriers.

34 DISCUSSION: Lifestyle variables showed modest associations with brain atrophy, with
35 alcohol drinking appearing most influential, with significant modification by *APOE* e4
36 genotype.

37 **KEYWORDS**

38 Brain health; atrophy; longitudinal; lifestyle; *APOE* e4; alcohol intake

39 **1. BACKGROUND**

40 With longer life expectancy, the prevalence of brain impairment and
41 neurodegenerative diseases are increasing, such as dementia, which affected
42 cognitive ability, memory, and the ability to perform daily activities in 57 million people
43 worldwide in 2019.¹⁻³ Dementia is a global public health problem, and the number of
44 people living with this disease is predicted to increase to 153 million by 2050.² Causes
45 of dementia may include multiple diseases or injuries that directly and indirectly lead
46 to brain damage and deterioration in cognitive function; Alzheimer's disease (AD), as
47 the most common form, accounts for 60-70% of cases.³ Apolipoprotein E gene (*APOE*)
48 e4 allele is known as the strongest genetic risk factor for AD.⁴ Although effective and
49 broadly indicated pharmacological treatments for dementia are still lacking, the Lancet
50 Commission on dementia suggests that around 45% of dementia might be preventable
51 by implementing interventions on modifiable risk factors, for example, smoking,
52 excessive alcohol consumption, obesity, and physical inactivity.¹ Understanding the
53 relationships between modifiable risk factors and brain structural changes is a priority
54 to prevent or delay dementia.

55 Brain structure is a relatively sensitive mediator of cognitive health, and a more
56 proximal phenotype to exposures, compared with cognitive test scores which are
57 widely used to explore different abilities,^{5,6} but remains incompletely explored.
58 Associations between several modifiable risk factors and poorer brain health have
59 been demonstrated through structural magnetic resonance imaging (MRI) phenotypes,
60 for example, smoking and smaller grey matter volumes,⁷ unhealthy diet and poorer
61 brain connectivity,⁸ and habitual sedentary behaviour with poorer white matter health,⁹
62 and excessive alcohol drinking with smaller grey matter volumes and poorer white
63 matter tract integrity.¹⁰ Yet, it is difficult to infer potential causality because of limitations

64 in these cross-sectional designs: these findings could reflect either differences in
65 baseline levels, reverse causation, or long-term accumulated effects.¹¹ Longitudinal
66 studies could isolate the change aspects and provide better tests of causality, which
67 so far remain scarce.¹²

68 Though human cognitive function changes continuously throughout life,
69 preventive interventions in early-life might be effective for late-life brain health and
70 cognitive outcomes.¹³ A few studies have provided longitudinal evidence on the
71 associations between lifestyle and brain structural changes, such as, moderate and
72 heavy alcohol drinking with greater decrease in cortical thickness of medial
73 orbitofrontal cortex among 179 participants;¹⁴ null associations between the
74 Mediterranean-DASH Intervention for Neurodegenerative Delay (MIND) diet score
75 and brain structural changes in UK Biobank.¹⁵ These studies were limited by small
76 sample sizes or examining only a single lifestyle variable with brain structural changes,
77 and neglected real-world clustering of risky lifestyle factors within individuals.
78 Convincing evidence that healthy lifestyle variables, either alone or in combination,
79 can protect brain health and prevent dementia is lacking.¹⁶ Thus, large-scale cohorts
80 with repeated neuroimaging data are warranted to elucidate the impacts of daily
81 lifestyle variables on subtle but clinically meaningful brain structural changes over time
82 while accounting for the genetic risk, and their interactions.

83 In the current study, we aim to examine whether and how individual/overall
84 lifestyle(s), including smoking, alcohol drinking, diet, sedentary behaviour, and
85 physical activity, influence atrophy in global brain measures as well as regional grey
86 matter volumes in the frontal lobe, considering *APOE* e4 risk and its interactions
87 among people living without dementia between two-timepoint scans.

88 **2. Methods**

89 2.1 Study design and participants

90 Data for the current study were obtained from the UK Biobank, which is a large
91 community-based cohort with 502,359 participants aged 37-73 years.¹⁷ From 2006 to
92 2010, participants who attended one of 22 assessment centres completed the baseline
93 assessments and questionnaires on physical, lifestyle, sociodemographic and medical
94 characteristics. In 2014, MRI scanning of the heart, brain and abdomen for a sub-
95 group of participants began; this is ongoing until the aim of 100,000 participants being
96 scanned is reached. As of September 2023, 71,091 participants aged 44-85 years had
97 attended the MRI assessment for the first time. Brain structural MRI data being
98 processed at the first scan are available for 46,398 participants, and repeat imaging
99 at the second scan for 4,784 participants. Participants who reported chronic
100 neurological diseases before the first scan which could directly affect cognitive function
101 were excluded (see **Table S1**). We mainly focused on lifestyle variables and covariates
102 collected at the time of first MRI scan as the baseline measures, and brain structural
103 phenotypes in two scans. This research was completed using UK Biobank project
104 number 17689.

105 2.2 Lifestyle

106 *2.2.1 Individual lifestyle variables*

107 Participants self-reported their health related lifestyles at the first scan, including
108 smoking status, alcohol intake, diet, sedentary behaviour duration and physical activity
109 (**Table S2**). The intercorrelations among lifestyle variables are shown in **Table S4**.

110 Three levels (namely optimal, moderate, poor) for each lifestyle variable were
111 defined based on public health guidelines. Smoking status was classified as never,

112 former, and current smoker.⁷ Alcohol drinking was defined here as low, moderate, and
113 high risk, being defined by types, frequencies and units of weekly alcohol intake and
114 grouped based on UK national guidelines: 0–14 units as low risk of disease; 15–35
115 units for females or 15–50 units for males as moderate risk; > 35 units for females
116 and > 50 units for males as high risk.¹⁸ Diet was categorized as healthy, moderate,
117 and unhealthy based on a diet score for cardiovascular diseases sharing similar risk
118 factors with dementia rather than the MIND diet,¹⁹ because information on 24-hour
119 diet recall being used to construct the MIND diet score were not collected when
120 participants attended the first MRI scan. This current score was derived from the food
121 frequency questionnaire, and constructed by the unweighted sum of adherence to the
122 UK dietary guidelines, including: processed meat, red meat individually less than twice
123 per week, total fish more than twice per week, consumption of semi- and skimmed,
124 soya milk, no spread intake, more than five bowls of cereal intake, no salt added to
125 food, more than six glasses of water and more than five servings per day of fruits and
126 vegetables. There is currently no guideline for sedentary behaviour, only being
127 recommended to limit the amount of time being sedentary by WHO; and using the
128 WHO guidelines for physical activity of 150-300 minutes moderate or 75-150 minutes
129 vigorous activity weekly would result in a highly left-skewed distribution.²⁰ Tertiles were
130 therefore used for categories of sedentary behaviour (time [in hours] spent driving,
131 using a personal computer, and watching television), and physical activity (weekly
132 metabolic equivalents task units [METs] minutes of walking, moderate and vigorous
133 activity) as low, medium and high levels²¹.

134 2.2.2 *Unweighted lifestyle score and lifestyle categories*

135 We assigned 0,1, and 2 points for each lifestyle variable based on the adherence to
136 poor, moderate and optimal lifestyle behaviour, respectively. Two points were

137 assigned for membership of each respective optimal level: never smoking; low risk
138 drinking; healthy diet; low duration of sedentary behaviour; high level METs. One point
139 was assigned for each of moderate-level lifestyles, such as former smoking, moderate
140 risk drinking, moderate diet, medium duration sedentariness, and medium METs; and
141 0 for poor ones. An overall lifestyle score was summed across these five three-level
142 lifestyle variables using an unweighted score, ranging from 0 to 10, with a higher score
143 indicating a healthier lifestyle. Based on the distribution of frequencies of participants'
144 lifestyle scores (see **Figure S1**), participants who scored 0, 1, 2 or 3 were classed as
145 unfavourable lifestyle; 4, 5, or 6 as moderate lifestyle; 7, 8, 9, or 10 as favourable
146 lifestyle.¹⁰ For example, someone who never smoked and had low risk drinking (two
147 points each), moderate diet (one point), low sedentariness (two points) and high METs
148 (two points) would receive nine points and be included in the favourable lifestyle group.
149 We used this method which prioritized clinical relevance and interpretability over
150 statistical convenience, rather than tertiles which might mask meaningful biological
151 patterns.

152 2.3 Imaging data

153 Brain structural markers were measured using MRI performed with a 3 Tesla Siemens
154 Skyra scanner (Siemens Healthineers) with VD13 software and a 32-channel head
155 coil since 2014.²² In the current study, we used data from T1-weighted imaging,
156 including global brain volumetric measures of grey matter, white matter, total brain;
157 hippocampal volumes: left hippocampus, right hippocampus, hippocampal asymmetry
158 (left minus right); volumetric measures of grey matter in frontal lobe (including frontal
159 pole; superior frontal gyrus; middle frontal gyrus; inferior frontal gyrus, pars triangularis;
160 inferior frontal gyrus, pars opercularis; frontal medial cortex; frontal orbital cortex;
161 frontal operculum cortex, see **Table S3**); these regions were selected as they have

162 shown association with cognitive deficits.²³ Additionally, we included the volume of
163 white matter hyperintensities from the T2 FLAIR structural scan as this is strongly
164 associated with dementia.²⁴ The primary outcomes of interest were grey matter
165 volumes, white matter volumes, total brain volumes, white matter hyperintensities
166 volumes, left/right hippocampal volumes, hippocampal asymmetry and volumes of
167 grey matter in frontal lobe regions (all uncorrected for head size).

168 2.4 Genetic data

169 UK Biobank genotyping was conducted by Affymetrix using a bespoke BiLEVE Axiom
170 array for ~50,000 participants and the remaining 450,000 on the Affymetrix UK
171 Biobank Axiom array. All genetic data were quality controlled by UK Biobank.²⁵ The
172 *APOE* e genotype is directly genotyped. The two *APOE* e single nucleotide
173 polymorphisms—rs7412 and rs429358—were both in Hardy-Weinberg equilibrium ($p >$
174 0.05) assessed with PLINK V1.90.²⁶ Further information on the genotyping process is
175 available (ukbiobank.ac.uk/scientists-3/geneticdata). For enough granularity to test
176 lifestyle interactions, a dominant *APOE* e4 model - e3/e4 and e4/e4 collated vs. e2/e2,
177 e2/e3 and e3/e3, was used as previous research showed that about 2% of the cohort
178 were e4 homozygotes.²⁷ *APOE* e2/e4 was removed as standard practice because it
179 has potentially both risk and protective alleles.²⁸

180 2.5 Covariates

181 We used information from questionnaires completed at the first scan which recorded
182 sociodemographic factors including age, sex (male or female), Townsend deprivation
183 index (categorized as quintiles);²⁹ health factors on body mass index (BMI) as
184 underweight, normal weight, overweight, or obese;³⁰ sleep durations as short (<7
185 hours), normal (7-9 hours) or long (>9 hours),³¹ social isolation (yes or no),³² history

186 of diabetes (yes or no), hypertension (yes or no), and depression (yes or no). Genetic
187 and imaging covariates were also included, such as genotypic arrays, the first 10
188 principal components for stratification, MRI sites (n=4), intracranial volume (ICV),
189 which was generated by summing the volumes of white matter, grey matter, and
190 ventricular cerebrospinal fluid at the first scan,³³ and time intervals, which were the
191 number of days between the dates of attending two scans. Since these measures were
192 ascertained prior to brain imaging, they cannot be mediators and were treated as
193 confounders (associated with both exposures and outcomes).

194 2.6 Statistical analysis

195 Characteristics of participants were described, stratified by the dominant *APOE* e4
196 model (presence versus absence), with Chi-square tests for categorical variables, t-
197 tests for normally distributed continuous variables, and Mann-Whitney U tests for not
198 normally distributed continuous variables; MRI volumes were combined across
199 hemispheres (summing left and right) for grey matter in the frontal lobe. For white
200 matter hyperintensities, outlier values greater than 5 standard deviations (SDs) from
201 the sample mean were excluded; data were log-transformed due to a positively
202 skewed distribution.

203 We used the raw change scores to examine the longitudinal associations
204 between lifestyle, *APOE* e4 genotype, and brain atrophy by linear regression, where
205 standardised betas, intuitive unstandardised betas and standard errors (SEs) were all
206 reported. The outcome variables were defined as the differences of raw brain structural
207 volumes (in mm³) between follow-up and baseline scans (e.g. brain volumetric
208 markers in the second scan minus markers in the first scan) and the main predictor
209 was the lifestyle variable in each regression model. Each model was fully adjusted for

210 corresponding brain structural values in the first scan; sociodemographic factors: age,
211 sex, Townsend deprivation categories; genetic and imaging related factors: genotypic
212 arrays, MRI sites, the first 10 principal components, ICV, and time intervals; health and
213 social factors: BMI, sleep duration, isolated status, history of diabetes, hypertension,
214 and depression. Positive betas indicate that the exposure group had less volume loss
215 (i.e., attenuated atrophy) compared to the reference group, whereas negative betas
216 indicate greater volume loss (i.e., accelerated atrophy), see **Figure S2**.

217 We firstly examined the general associations of lifestyle categories (including
218 unfavourable, moderate, and favourable lifestyles) and brain atrophy, additionally for
219 continuous lifestyle scores. To further interpret the association, we computed adjusted
220 marginal means of brain structural changes across lifestyle categories using *margins*
221 in Stata, holding other covariates at their observed values. Then, the intuitive
222 contributions of individual lifestyle variables on raw brain changes were investigated
223 in two phases: 1) we estimated the associations of individual lifestyle variable with
224 brain changes without adjusting for other lifestyle variables; 2) all lifestyle variables
225 were included in one regression model simultaneously, being adjusted mutually in one
226 multiple linear regression model for each brain MRI variable of interest. Interactions
227 between APOE e4 status with lifestyle categories, and individual lifestyle variable, on
228 brain structural atrophy were also tested.

229 Analyses were conducted in Stata v.18 and R (v4.2.3). We report two-sided P-
230 values throughout and false discovery rate (FDR) correction was applied to keep the
231 error rate at 0.05.

232 **3. Results**

233 **3.1 Characteristics of participants**

234 A total of 2214 participants were included with complete information on lifestyle factors,
235 *APOE* e4 genotype, and covariates in the first scan and brain structural markers of
236 two scans (**Figure 1**). The median (IQR) of follow up was 2.25 (0.30) years. The 2214
237 participants had a mean (SD) age of 61.3 (7.3) years in the first scan and 1099 (49.6%)
238 were female (**Table 1**). There were 547 (24.7%) participants carrying at least one copy
239 of the e4 allele. Among e4 carriers, there were 250 (45.7%) participants with moderate
240 lifestyles, followed by 249 (45.5%) with favourable lifestyles and 48 (8.8%) with
241 unfavourable lifestyles. *APOE* e4 carriers were more likely to be younger (Cohen's d= 0.11, P=0.031) and have smaller white matter volume (Cohen's d= 0.10, P=0.036),
242 and left hippocampal volume (Cohen's d= 0.12, P=0.018) in the first scan.
243

244 3.2 Associations between lifestyle, *APOE* e4 status, and longitudinal brain atrophy

245 3.2.1 *Lifestyle categories*

246 As shown in **Figure 2** (with supporting information in **Table S5**), compared with
247 favourable lifestyle, unfavourable lifestyle was significantly associated with greater
248 brain atrophy in grey matter and left hippocampal volumes, but not with other brain
249 structural phenotypes. Participants with favourable lifestyle experienced a mean
250 atrophy of 7883.8 mm³ atrophy in grey matter and 63.6 mm³ in left hippocampus
251 independently, over a median follow-up of two years. Compared with the favourable
252 lifestyle group, unfavourable lifestyle was associated with 2059.2 mm³ more atrophy
253 in grey matter, equivalent to 0.046 standard deviation (SD) greater decline (95% CI =-3966.5 to -152.0, P=0.034), and 36.5 mm³ more atrophy in left hippocampus,
254 equivalent to 0.044 SD greater decline (95% CI =-72.5 to -0.4, P=0.048). Additionally,
255 a one-point increment in lifestyle score (out of 10 points) was associated with less grey
256 matter loss (unstandardised β = 318.1, 95% CI =32.9 to 603.2, standardised β = 0.047,

258 P=0.029), and less hippocampal asymmetry (unstandardised β = 6.4, 95% CI =0.2 to
259 12.6, standardised β = 0.044, P=0.042) in **Table S6**. However, none of these
260 associations remained significant after correction for multiple testing ($p_{(FDR)} > 0.05$).

261 *3.2.2 Individual lifestyle variable*

262 **Figure 3** presents the heatmap of associations of individual and simultaneous lifestyle
263 variables on longitudinal brain atrophy. In **Figure 3A** with supporting information in
264 **Table S7**, we observed significant impacts of single lifestyle variables on certain
265 investigated brain markers. For example, for alcohol drinking, compared with low-risk
266 drinking, high-risk drinking was associated with greater atrophy of 57.3 mm³ in left
267 hippocampal volume, corresponding to 0.062 SD greater loss (P=0.004), and
268 accelerated decreased volumes of 48.9 mm³ in inferior frontal gyrus, pars triangularis,
269 corresponding to 0.048 SD greater loss (P=0.023), while moderate drinking was linked
270 to accelerated total brain atrophy of 2075.4 mm³, corresponding to 0.063 SD greater
271 loss (P=0.003), especially in grey matter volumes with 1336 mm³ more atrophy,
272 corresponding to 0.054 SD greater loss (P=0.012). For physical activity, compared
273 with participants with high METs, those with low METs showed greater atrophy of
274 1255.4 mm³ in grey matter volumes, equivalent to 0.05 SD greater decrease (P=0.035),
275 and those with medium METs attenuated white matter atrophy of 1413.7 mm³,
276 equivalent to 0.05 SD greater decrease (P=0.035). These significant associations of
277 each lifestyle variable on brain imaging markers remained largely consistent even
278 when we included all lifestyle variables in one simultaneous regression model (**Figure**
279 **3B** with supporting information in **Table S8**), although high-risk drinking's effect on left
280 hippocampus reduced slightly from -0.062 SD to -0.055 SD. After FDR correction, only
281 moderate drinking and total brain atrophy remained statistically significant ($p_{(FDR)}$
282 =0.045); all other associations became non-significant ($p_{(FDR)} > 0.05$). No significant

283 associations were observed for smoking, diet or sedentary behaviour with brain
284 structural atrophy in either univariable or multivariable models (all $P>0.05$).

285 3.3 Interaction of lifestyle and *APOE* e4 status on brain structural phenotypes

286 Two-way interaction tests of lifestyle categories \times *APOE* e4 status for each brain
287 imaging marker did not reveal significant interactions in **Table S9**. However, in two-
288 way interaction tests of each individual lifestyle variable \times *APOE* e4 status on brain
289 structural changes in **Table S10**, significant interactions were found between high-risk
290 drinking and *APOE* e4 status on regional grey matter atrophy in frontal lobe as well as
291 moderate drinking and sedentary behaviour with regional brain changes. After
292 correcting for multiple testing, only the interaction of high-risk drinking and *APOE* e4
293 status on regional grey matter atrophy in frontal lobe survived, including frontal pole,
294 superior frontal gyrus, inferior frontal gyrus, pars triangularis, and frontal operculum
295 cortex ($p_{(FDR)}<0.05$). Based on this finding, we then performed subgroup analyses
296 stratified by *APOE* e4 status (**Table S11**), which showed among e4 non-carriers,
297 moderate drinking subtly accelerated atrophy in grey matter in frontal lobe regions
298 except for frontal operculum cortex (standardised β ranges from -0.064 to -0.051,
299 $P<0.05$), but among e4 carriers, we observed greater decreased volumes
300 (standardised β ranges from -0.093 to -0.072, $P<0.05$) between high drinking and
301 frontal lobe volumes. None of these associations remained significant when correcting
302 for multiple testing.

303 **4. Discussion**

304 In this longitudinal study, we showed that lifestyle and genetic risk for AD were
305 associated with accelerated structural changes in a range of brain phenotypes over a
306 period of up to 7 years. Longitudinal associations were nominally found between an

307 overall unfavourable lifestyle, and grey matter and left hippocampal atrophy. Among
308 individual lifestyle variables, alcohol drinking showed the most consistent effects on
309 brain atrophy: moderate drinking was associated with total brain atrophy, as well as
310 high drinking and declined volumes in the left hippocampus and frontal lobe at the
311 follow-up scan. Moreover, we observed evidence of interaction between *APOE* e4
312 status and alcohol intake, with e4 carriers showing greater brain atrophy to the adverse
313 effects of high-risk drinking.

314 This is the first study to demonstrate that unfavourable lifestyles accelerate
315 longitudinal grey matter and left hippocampus atrophy nominally, although
316 associations were no longer significant after correcting for multiple comparisons.
317 However, for each lifestyle variable, they had impacts on certain brain volumetric
318 changes.

319 Alcohol drinking was the most consistent lifestyle factor associated with brain atrophy
320 in our study. Moderate drinking was associated with greater total brain atrophy after
321 multiple testing, compared with low risk drinking. Our finding is also partly supported
322 by a Mendelian randomization study reporting that moderate alcohol consumption was
323 associated with higher brain iron, whose accumulation represented a potential
324 mechanism for alcohol-related cognitive decline.³⁴ High-risk drinking was associated
325 with greater atrophy in left hippocampus and grey matter in inferior frontal gyrus, pars
326 triangularis, generally in line with prior studies.^{35,36} This might be because of functional
327 lateralization, which meant the left hippocampus was more involved in verbal memory
328 processing and it was often negatively impacted by high alcohol consumption on
329 cognitive function, making the impact on the left hippocampus more obvious.³⁷ When
330 including all lifestyle factors in one regression model, high-risk drinking was associated
331 with less atrophy in left hippocampus based on the reduced beta. This might be

332 because the detrimental effects of high alcohol consumption may be mitigated through
333 other lifestyle modifications, such as maintaining a balanced diet, engaging in regular
334 physical activity, and avoiding smoking and long duration sedentariness. Evidence
335 suggests that these healthy habits can enhance metabolic resilience, and reduce
336 systemic inflammation, thereby counteracting alcohol-induced oxidative stress.³⁸

337 We also observed that physical inactivity was associated with grey matter atrophy.
338 However, compared with high level METs, medium levels of physical activity were
339 related to less white matter atrophy, namely, better brain ageing. This could be
340 explained by moderate exercise leading to increased cerebral blood flow
341 and neurotrophic factors, which may have a positive impact
342 on angiogenesis and neurogenesis.³⁹ These effects translate to beneficial effects on
343 structural brain measures associated with cognitive function.⁴⁰ High activity levels
344 might be associated with metabolic strain, oxidative stress, and inflammatory
345 responses,⁴¹ which could counteract the neuroprotective effects. We found that
346 medium and high sedentariness were not associated with longitudinal brain atrophy.
347 This is inconsistent with other studies which reported the relationship between
348 sedentary behaviour and reduced white matter health.^{42,43} This might be due to overall
349 time use patterns, as sedentary behaviour per se may not be detrimental when
350 individuals compensate by engaging in sufficient physical activity during their non-
351 sedentary time. We still cannot ignore the possibility that the multiple detrimental
352 cardiovascular and metabolic effects led by sedentary behaviour can be intermediate
353 risk factors for dementia, such as hypertension.⁴⁴ Other metabolic effects, including
354 insulin resistance, and blood lipids, also have a negative impact on white matter health,
355 which in turn might increase the risk of dementia.⁴⁵

356 Notably, although multiple significant cross-sectional associations between smoking,
357 and diet and lower brain volumes have been reported,^{7,8} we found null associations
358 between smoking or diet with longitudinal brain atrophy in the investigated brain
359 structural markers. Of them, current or former smoking was not associated with brain
360 changes in the current study. Our finding is inconsistent with previous research stating
361 that smoking was associated with decreased structural integrity of multiple brain
362 regions among elderly people without dementia.⁴⁶ This might be explained by the
363 impacts of smoking on brain changes being distributed by other factors, or that the
364 measurement of smoking is less precise than other, i.e. non-misclassification bias, and
365 thus associations could not be detected. We cannot neglect the impact of chronic
366 exposure of noxious agents in cigarette smoking, which may directly compromise
367 neuronal and cellular membrane functions of cerebral tissue and lead to
368 cardiovascular and cerebrovascular disease.⁴⁶

369 Furthermore, we found no significant associations of moderate or unhealthy diet with
370 brain atrophy. We used a developed diet score from the food frequency questionnaire
371 at the first scan designated for cardiovascular diseases which shared similar risk
372 factors with cerebrovascular diseases.¹⁹ Although the MIND diet is well-known to slow
373 cognitive decline and lower risks of all-cause dementia,⁴⁷ data on 24-hour diet recall
374 which is used to construct the MIND diet score was not collected at the first MRI scan
375 in the UK Biobank, which meant we were unable to explore its impacts on brain
376 changes properly. However, a prior study showed no significant associations between
377 MIND diet and longitudinal brain structural changes by accessing the first wave of
378 imaging data being taken long after dietary assessments.¹⁵

379 Significant interactions were found between high-risk drinking and *APOE e4* status on
380 grey matter volume decline in frontal lobe. Non-low risk drinking was significantly

381 associated with decreased volumes of grey matter in frontal lobe regardless of *APOE*
382 e4 status. However, *APOE* e4 carriers may be more vulnerable to the effects of
383 alcohol-related neurodegeneration due to more brain atrophy compared with non-
384 carriers. Our finding is supported by one study showing that heavy drinking history
385 was associated with an earlier onset of AD by 2–3 years among *APOE* e4 carriers.⁴⁸
386 However, it contrasted with another study stating *APOE* e4 did not moderate the
387 relationship between alcohol use and cognitive performance.⁴⁹ This study was limited
388 by a smaller sample size (N=818) and missing covariates of medical conditions. The
389 findings remained unclear and further research on interaction tests are needed.

390 Major strengths were the relatively large sample size with long-term follow-up (up to 7
391 years), repeated brain MRI data for conducting longitudinal analyses, and
392 simultaneous inclusion of multiple daily lifestyle factors that allowed us to investigate
393 their individual and combined effects on brain changes. Moreover, the use of raw brain
394 changes, adjusted for baseline brain phenotypes, yielded intuitive and robust results.

395 However, limitations remain. First, our findings might be underestimated potentially
396 because of healthy volunteer bias. We have previously observed prevalent healthy
397 bias in the imaging sub-sample from the already biased UK Biobank baseline
398 sample.⁵⁰ Second, lateralized effects of lifestyle factors on longitudinal brain ageing
399 might exist as we investigated the summed left/right phenotypes in volumes of grey
400 matter in the frontal lobe in analyses. For future studies, it might be interesting to
401 examine the associations of specific lifestyle factors and left/right phenotype changes
402 to explore potential underlying mechanisms. Third, while these associations highlight
403 the neuroimaging consequences of specific lifestyle factors, the interplay between
404 individual factors and white matter tracts remains underexplored. Future research
405 should prioritize longitudinal investigations into how lifestyle factors influence white

406 matter microstructural changes to elucidate their independent and synergistic effects.
407 Fourth, despite excluding participants with at least one neurological condition in the
408 current study, those with cognitive impairments were not specifically identified in this
409 population-based sample. Thus, the results might be overestimated. Fifth, despite the
410 longitudinal design, and the adjustment of a comprehensive list of confounders,
411 residual confounding cannot be ruled out as with all other observational studies. Last
412 but not least, potential reverse causality might exist; although we have shown
413 evidence consistent with causal effects, we cannot conclude that the lifestyle factors
414 themselves.

415 Our findings have several important implications. First, alcohol drinking consistently
416 showed adverse effects, suggesting that this lifestyle variable may be a key modifiable
417 target for preserving brain health. Notably, our findings identified associations of
418 moderate intake with greater longitudinal brain atrophy: there has been a long-
419 standing perception of moderate alcohol consumption as safe, underscoring the need
420 for cautious public health recommendations. Second, the significant interaction
421 between *APOE* e4 status and alcohol use highlights the potential role of genetic
422 susceptibility in modifying lifestyle effects on brain ageing. This supports the notion
423 that prevention strategies should increasingly consider gene–environment interactions
424 and be tailored to vulnerable subgroups such as *APOE* e4 carriers. Together, these
425 implications emphasise the value of integrating lifestyle modification with personalised
426 risk assessment in efforts to mitigate neurodegeneration and promote healthy brain
427 ageing.

428 Our findings showed an overall unfavourable lifestyle was associated with greater
429 longitudinal atrophy in grey matter and left hippocampus over time. Moderate and
430 high-risk drinking accelerated brain atrophy, compared with low-risk drinking. Among

431 *APOE e4* carriers, high-risk drinking may be an important modifiable risk factor for
432 brain atrophy, indicating a gene - lifestyle interaction in brain ageing. These findings
433 advocate for public health strategies prioritizing relevant lifestyle interventions to delay
434 cognitive aging and prevent dementia.

435

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443

444 **AUTHOR CONTRIBUTIONS**

445 Study concept: YY, FKH, DML, CEH.
446 Obtained data: DML.
447 Design: YY, FKH, DML, CEH.
448 Conducted analyses: YY.
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451

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454 **CONFLICT OF INTEREST STATEMENT**

455 The authors have no conflicts of interest to disclose.

456

457 **CONSENT STATEMENT**

458 All human subjects provided informed consent.

459

460 **SUPPORTING INFORMATION**

461 Supplementary information is available at website.

462

463 **FIGURE LEGENDS**

464 **FIGURE 1. Flowchart summarising participants selection and inclusion**
465 **processes, including the reasons for exclusion of participants**

466 **FIGURE 2. Margin plots for predicted brain structural volumes by lifestyle**
467 **categories**

468 Unstandardized betas and 95% confidence intervals were calculated using linear
469 regression models (in mm³), where lifestyle categories are regressed on raw changes
470 of grey matter volumes (A), white matter volumes (B), total brain volumes (C), left
471 hippocampal volumes (D), right hippocampal volumes (E), white matter hyperintensity
472 volumes (F), hippocampal asymmetry (G), frontal pole (H), superior frontal gyrus (I),
473 middle frontal gyrus (J), inferior frontal gyrus, pars triangularis (K), inferior frontal gyrus,
474 pars opercularis (L), frontal medial cortex (M), frontal orbital cortex (N), frontal
475 operculum cortex (O) with the reference of people having favourable level lifestyle,
476 adjusted for corresponding baseline brain structural volumes, age, sex, genotypic
477 array, four MRI assessment centres, first ten genetic principal components, *APOE4*
478 presence, deprivation index, BMI, sleep duration, isolated status, history of diabetes,
479 hypertension, depression, intracranial volume among N=2214 participants. * denotes
480 significant associations (P<0.05).

481 **FIGURE 3. Heatmap of longitudinal associations of individual lifestyle variable**
482 **individually (A.) and simultaneously (B.) modelled on brain structural markers**

483 Standardized betas were calculated using linear regression models, where each three-
484 level lifestyle factor regressed on raw changes of brain structural markers individually
485 (A), then an overall lifestyle factors regressed on brain metrics simultaneously (B)
486 among N=2214 participants with the reference of people having healthy lifestyle, such

487 as never smoking, low risk drinking, healthy diet, low sedentariness duration and high
488 METs. Models were fully adjusted for corresponding baseline brain structural volumes,
489 age, sex, genotypic array, four MRI assessment centres, first ten genetic principal
490 components, *APOE4* presence, deprivation index, BMI, sleep duration, isolated status,
491 history of diabetes, hypertension, depression, intracranial volume. Stronger
492 associations are indicated by darker shades, light/dark red indicates a positive
493 association ($P<0.05$), light/dark purple indicates a negative association ($P<0.05$). *
494 denotes significant associations ($P<0.05$).

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