

1 **Relationships between eating behaviours, psychopathology, brain**
2 **maturation and genetic risk for obesity in a longitudinal adolescent cohort**
3 **study**
4

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66

67 **Abstract**

68 Unhealthy eating, a risk factor for eating disorders (EDs) and obesity, often co-exist with
69 emotional and behavioural problems, but the underlying neurobiological mechanisms are
70 poorly understood. Analysing data from the longitudinal IMAGEN adolescent cohort, we
71 investigated associations between eating behaviours, genetic predispositions for high BMI
72 (BMI PGS), and trajectories (ages 14y to 23y) of ED-related psychopathology and brain
73 maturation. Clustering analyses at age 23y ($N = 996$) identified three eating groups:
74 restrictive, emotional/uncontrolled, and healthy eaters. BMI PGS, trajectories of ED
75 symptoms, internalising and externalising problems, and brain maturation distinguished these
76 groups. Decreasing volumes and thickness in several brain regions were less pronounced in
77 restrictive and emotional/uncontrolled eaters. Smaller cerebellar volume reductions uniquely
78 mediated the effects of BMI PGS on restrictive eating. Smaller volumetric reductions across
79 multiple brain regions mediated the relationship between elevated externalising problems and
80 emotional/uncontrolled eating, independently of BMI. These findings shed light on distinct
81 contributions of genetic risk, protracted brain maturation and behaviour in ED
82 symptomatology.

83

84

85 **Main**

86 Eating disorders (EDs) are serious psychiatric disorders with high mortality rates, substantial
87 impacts on quality of life, and economic burdens^{1,2}. Their increasing prevalence^{3,4} particularly
88 during mid-adolescence⁵ highlights the need for early detection and effective interventions.

89

90 Key risk factors for EDs include eating behaviours such as dietary restraint and overeating,
91 which increase the risk for unhealthy weight control behaviours and EDs like bulimia nervosa
92 and binge eating disorder⁶. Assessments of eating behaviours in various population groups^{7,8}
93 indicated that cognitive restraint (CR), the conscious restriction of food intake to control body
94 weight and shape, can lead to episodic overeating and is a strong predictor of disordered eating
95 and negative body image. Conversely, uncontrolled eating (UE) –which refers to eating in
96 response to food palatability, social cues and hunger, resulting in eating episodes– and
97 emotional eating (EE) –eating episodes elicited by negative affect– is associated with higher
98 susceptibility to EDs, hedonically driven food choices, higher BMI, and obesity. Genetic
99 studies indicate that these behaviours may be partially genetically determined, with obesity-
100 associated variants linked to CR, UE and EE⁹, and eating behaviour trajectories in childhood¹⁰
101 and adolescence¹¹.

102

103 Neural factors also play a role in EDs¹², with neurobiological differences observed in clinical
104 samples^{13–17}. Neurobehavioural correlates suggest that the hypothalamic, emotion/memory,
105 and executive systems are involved in eating control^{18,19}. Neural activation of lateral
106 prefrontal structures underpinning self-control and decision-making and striatal reward
107 regions have been shown to underly individual differences in CR²⁰. Longitudinal studies have
108 revealed volumetric brain differences, particularly in striatal and prefrontal regions,

109 suggesting differences in brain maturation as etiological factors for disordered eating
110 behaviours and comorbid depressive symptoms²¹.

111

112 Internalising and externalising psychopathology symptoms may serve as premorbid risk
113 factors for eating disorders²²⁻²⁴. Externalising problems in early adolescence predict the
114 onset¹¹ and persistence of eating pathology²⁵, while generalized anxiety symptoms predict
115 adolescent-onset ED²⁶. Negative affect and functional impairment are found to predict the
116 onset of all eating disorders^{6,27}. However, how trajectories of adolescent maladjustment, as
117 evidenced by internalising and externalising problems, relate to eating behaviours, genetic
118 liability and brain maturation, is not well understood.

119

120 This study aimed to advance our understanding of eating behaviours by analysing
121 longitudinal data from the IMAGEN adolescents' cohort. Applying a multivariate analytical
122 framework, the study identified eating profiles at age 23, characterized by associations with
123 polygenic scores for higher BMI (BMI PGS), and differences in earlier trajectories of
124 disordered eating, internalising and externalising problems, and brain maturation. The
125 mediating roles of brain maturation and BMI PGS were also investigated using multivariate
126 models. The analytical workflow is illustrated in **Fig. 1**.

127

128 **Results**

129 **Identification of groups with distinct eating profiles**

130 A total of 996 participants (478 males and 518 females) with completed Three-Factor Eating
131 Questionnaire (TFEQ) scores at age 23 and had at least one measure from the Strengths and
132 Difficulties Questionnaire (SDQ) available at ages 14, 16, 19 and 23 were included in the
133 current study (**Methods**). Three groups were identified from K-means clustering analysis
134 with distinct eating behaviours (**Table 1** and **Fig. 2a**). Group distribution and the within-
135 group sum of squares are detailed in **Supplementary Fig. S1**. Validity and stability analyses
136 confirmed the three-group solution with Jaccard similarities of 0.83, 0.77, and 0.75,
137 respectively. One group ($N = 423$) scored low on all eating behaviours, indicating healthy
138 eaters (HEs). Another group ($N = 324$) exhibited the highest cognitive restraint, indicating
139 restrictive eaters (REs). Behaviours that differentiated REs most from HEs (Odds Ratios > 5)
140 included consciously eating less to control weight and weight gain, the intensity of restrained
141 eating, consciously eating less than wanted, and not eating foods that made them fat
142 (**Supplementary Information**). This group also scored significantly higher than HEs on
143 uncontrolled eating. The third group ($N = 249$) showed the highest emotional and
144 uncontrolled eating, indicating emotional and uncontrolled eaters (E/UEs). Behaviours that
145 distinguished E/UEs most from HEs included eating/overeating when feeling blue, lonely or
146 anxious (all EE items), inability to stop eating, and frequency of binge eating episodes (UE
147 items). This group also reported significantly higher cognitive restraint than HEs. REs and
148 E/UEs comprised predominantly female participants, contrasting with HEs, which had a
149 higher proportion of males. Consistent with differences in BMI, the BMI PGSs were higher
150 in REs and E/UEs than in HEs (**Table 1**).

151

152 **Group differences in trajectories of eating disorder symptoms across adolescence**

153 Linear mixed models were applied to investigate group differences in trajectories of ED
154 symptoms (dieting, binge eating and purging) from ages 14 to 23. Analyses of age-by-group
155 interactions were used to identify symptom trajectories that differed in the REs or E/UEs
156 groups compared to HEs (**Fig. 2b and Supplementary Tables S1 and S2**). REs were
157 characterised by significantly ($p = 6.29 \times 10^{-3}$) higher overall levels of dieting compared to
158 HEs, with no significant age-by-group interactions. Trends towards increased dieting from
159 ages 14 to 16 and increased binge eating from ages 14 to 19 were observed in this group. In
160 E/UEs, significant age-by-group interactions were observed when compared to HEs, with
161 significantly increased dieting from ages 14 to 16 ($p_{\text{Bonferroni}} = 0.026$), and increased binge
162 eating from ages 14 to 19 ($p_{\text{Bonferroni}} = 0.028$) and 14 to 23 ($p_{\text{Bonferroni}} = 3.024 \times 10^{-7}$). Nominal
163 increases in purging were observed from age 14 to 16, 19 and 23.

164

165 **Group differences in trajectories of internalising and externalising problems**

166 We explored behavioural group differences further, using latent growth curve models
167 (LGCM) to measure trajectories of internalising and externalising problems across (**Fig. 3a**)
168 and within groups (**Fig. 3d**). Between groups univariate analyses revealed that compared to
169 HEs, unhealthy eaters (REs and E/UEs) exhibited significant differences in how their
170 internalising problems developed over time. Both REs ($b = 0.073$, 95% CI = 0.019-0.126, $p =$
171 0.008; **Fig. 3b**) and E/UEs ($b = 0.110$, 95% CI = 0.051-0.169, $p < 0.001$) reported a
172 significant increase (i.e., the slopes of their trajectories; **Supplementary Tables S3 and S4**)
173 in internalising problems with age. Intercepts of internalising problems also differed, with
174 E/UEs already reporting higher levels of internalising problems at age 14 than HEs ($b =$
175 0.712, 95% CI = 0.281-1.144, $p = 0.001$). Regarding externalising problem trajectories, all

176 three groups showed a decrease in these problems over time, but the rate of decrease did not
177 significantly differ between groups (**Fig. 3c**). Nevertheless, the externalising problems
178 reported at age 14 were higher in E/UEs compared to HEs ($b = 0.855$, 95% CI = 0.418-1.292,
179 $p < 0.001$) and REs ($b = 0.743$, 95% CI = 0.278-1.209, $p = 0.002$).

180 Within-group multivariate LGCM analyses, which included internalising and externalising
181 problems in the same model (**Fig. 3d**), revealed significant within- and between-constructs
182 correlations (**Fig. 3e**). Within-construct correlations revealed that in all groups, higher levels
183 of externalising problems at age 14 were significantly correlated with smaller decreases in
184 these problems over time. This association was especially strong in unhealthy eaters (REs and
185 E/UEs), indicating that higher initial levels of externalising problems were linked to less
186 improvement over time. Similarly, in REs and HEs, higher initial levels of internalising
187 problems were associated with smaller decreases in these problems over time, but this pattern
188 was not observed in E/UEs. Interestingly, our models highlighted notable connections
189 between internalising and externalising problems. Specifically, we found that these problems
190 tended to co-occur, especially in unhealthy eaters (REs and E/UEs), suggesting that
191 individuals who started with higher levels of one type of problem were more likely to have
192 higher levels of the other as well. In healthy eaters, changes in internalising and externalising
193 problems were positively correlated, indicating that as one type of problem decreased, the
194 other tended to decrease as well. Additionally, in contrast to the other groups, in REs, there
195 was a significant positive correlation between the initial levels of internalising problems and
196 the changes in externalising problems over time, suggesting that higher initial levels of
197 internalising problems were related to more subsequent changes in externalising problems.
198 No other significant relationships between internalising and externalising problems were
199 found in these analyses.

200

201 **Group differences in brain maturation across adolescence**

202 Longitudinal analyses were conducted to investigate between-groups differences in brain
203 maturation during adolescence, comparing changes in grey matter volumes (GMV), cortical
204 thickness (CT), and sulcal depth (SD) from age 14 to 23. All primary analyses included sex,
205 recruitment sites, and total intracranial volume (TIV) as covariates.

206 For GMV (**Supplementary Table S6** and **Fig. 4a**), longitudinal VBM analyses were
207 conducted, which revealed significant age-by-group interactions. Compared to HEs, REs
208 showed smaller GMV reductions in the left cerebellum. E/UEs had smaller GMV reductions
209 in 2 subclusters in the left cerebellum and 5 subclusters in the right hemisphere, including the
210 middle frontal gyrus (MFG), putamen, medial superior frontal gyrus (SFGmedial) and
211 postcentral gyrus. Lower GMV increase in the right parahippocampal gyrus was also
212 observed. Differences remained similar after controlling for the joint effects of pubertal status,
213 IQ, educational attainment and age- and sex-adjusted BMI (**Supplementary Table S7**). No
214 significant differences in GMV trajectory were observed between REs and E/UEs.

215 For CT (**Supplementary Table S8** and **Fig. 4b**), mixed linear analyses revealed significant
216 age-by-group interactions in the right frontal pole when comparing E/UEs to REs, with REs
217 experiencing more pronounced CT reduction. In contrast, comparing E/UEs to HEs revealed
218 significant interactions across 9 brain regions. These included the left lingual gyrus, bilateral
219 frontal pole, bilateral rostral MFG, left pericalcarine, left cuneus, right caudal MFG and right
220 caudal anterior cingulate gyrus, with E/UEs displaying less CT reduction than HEs. Most
221 differences remained significant after controlling for the joint effects of pubertal status, IQ,
222 educational attainment (EA), age- and sex-adjusted BMI, Euler's number, and their joint
223 effects (**Supplementary Table S9**).

224 Regarding SD (**Supplementary Table S10** and **Fig. 4c**), REs had a less pronounced
225 reduction in the left frontal pole compared to E/UEs. Moreover, E/UEs displayed larger SD
226 reductions across 9 regions compared to HEs, including the bilateral rostral MFG, left frontal
227 pole, bilateral SFG, right caudal MFG, right pars orbitalis, right pars opercularis, and right
228 pars triangularis gyrus. Most differences remained significant after controlling for all
229 covariates (**Supplementary Table S11**).

230 No significant age-by-group interactions in CT and SD were found when comparing REs to
231 HEs.

232

233 **Brain maturation mediates relationships between adolescent psychopathology and** 234 **eating behaviours**

235 We conducted mediation analyses to examine whether the brain differences identified above
236 mediated the relationships between variations in internalising (IP) and externalising problem
237 (EP) trajectories during adolescence and eating behaviours in young adulthood, as determined
238 by the k-means derived clusters (the “psychopathology-brain maturation-eating behaviours”
239 models). Brain regions with significant group differences in their GMV, CT or SD
240 trajectories were identified as regions of interest (ROIs) and tested for their mediating effects
241 on behavioural group differences between RE or E/UEs compared to HEs. For REs, the
242 cluster in the left cerebellum was used as the ROI. For E/UEs, differences across E/UE-
243 related clusters were combined into a single ROI for each structural brain measure. All
244 primary mediation analyses were adjusted for sex, recruitment sites, and TIV differences to
245 account for overall brain size variations across ages.

246 For REs, who differed from HEs during adolescence by their increasing trajectories of
247 internalising problems (IP slope) and less pronounced GMV reductions in the left cerebellum,

248 differences in cerebellar volume reductions partially mediated the relationship between
249 increased internalising problems and being classified as an RE at age 23 (**Fig. 5a**). This
250 mediation was no longer significant after adjustments for BMI, IQ and pubertal status.

251 Behaviourally, E/UEs differed from HEs in adolescence by their higher levels of internalising
252 and externalising problems (EP and IP intercepts) and increasing trajectory of internalising
253 problems (IP slope). Nominal significance was found when testing the mediation effects of
254 GMV (**Fig. 5b**) and CT (**Fig. 5c**) differences on the relationship between IP slope and being
255 classified as an E/UE at age 23. Mediation in the “IP slope-GMV-E/UE” model remained
256 significant after controlling for covariates, while for the “IP slope-CT-E/UE” model, the
257 mediation remained significant after adjusting for BMI or EA, but no longer when adjusting
258 for pubertal status or IQ. GMV differences in E/UE-related ROIs significantly ($p < .001$) also
259 mediated the associations between EP intercept and E/UEs (**Fig. 5d**). This was unaffected by
260 BMI, IQ, EA or pubertal status. No significant mediating effects were found for SD
261 differences.

262 Altogether, these findings suggest that altered neurodevelopment in the restrictive and
263 emotional/uncontrolled eating groups, as evidenced by protracted GMV and CT reductions,
264 may contribute to ED symptoms, partly by mediating the effects of internalising and
265 externalising problems.

266

267 **Relationships between genetic predispositions, brain maturation, psychopathology and** 268 **eating behaviours**

269 Considering the effects of covarying BMI in the mediation analyses above, we explored the
270 potential contributions of polygenic risk for higher BMI. Simple “genetics-brain maturation-
271 eating behaviours” mediation models indicated that smaller GMV reductions in the RE and

272 E/UE groups (compared to HEs) mediated the effects of BMI PGS on eating behaviour
273 profiles at age 23 in these groups (REs vs. HEs: indirect effect = 0.042, $p = 0.002$, 13.5%
274 mediated; E/UEs vs. HEs: indirect effect = 0.043, $p = 0.026$, 11.7% mediated). For REs, this
275 remained significant after adjusting for IQ, EA or pubertal status, not BMI. For E/UEs, this
276 only remained significant after controlling for IQ or EA. No significant association was found
277 between BMI PGS and CT reductions related to E/UEs. These analyses suggest that genetic
278 predispositions to higher BMI influence BMI and restrictive and emotional/uncontrolled
279 eating partly through their effects on protracted GMV reductions during adolescence.

280 Multivariate mediation analyses including internalising or externalising problems in these
281 models revealed the unique contribution of smaller cerebellar GMV reductions in mediating
282 the effects of BMI PGS on REs, when changes in internalising problems were also
283 considered (**Fig. 5e**). In contrast, smaller GMV reductions in E/UEs mediated the effects of
284 early externalising problems (at age 14) in this group, beyond the effects of BMI PGS (**Fig.**
285 **5f**). These findings suggest a specific role for cerebellar maturation in the control of BMI and
286 restrictive eating, and additional roles for cortical and putamen maturations in mediating the
287 effects of behavioural problems on emotional/uncontrolled eating.

288

289 **Discussion**

290 In this study, we used a longitudinal, multivariate analytical framework to explore the
291 interplay among eating behaviours, genetic factors, ED-related psychopathology, and brain
292 maturation during adolescence. Our analyses revealed a high prevalence of unhealthy eating
293 behaviours (restrictive and emotional/uncontrolled eaters) associated with higher BMI and
294 BMI PGS. Unhealthy eaters exhibited higher levels of externalising problems
295 (emotional/uncontrolled eaters) and increasing trajectories of dieting (restrictive and
296 emotional/uncontrolled eaters), binge eating (emotional/uncontrolled eaters) and internalising
297 problems (restrictive and emotional/uncontrolled eaters). Age-related decreases in volumes
298 and thickness in several brain regions, particularly in the cerebellum and prefrontal cortex,
299 were less pronounced in restrictive and emotional/uncontrolled eaters compared to healthy
300 eaters, suggesting protracted brain maturation. Smaller volumetric reductions in the left
301 cerebellum mediated the effects of BMI PGS on restrictive eating, even after accounting for
302 internalising problems. Reductions within additional brain regions of the right hemisphere
303 uniquely mediated the relationship between externalising problems and
304 emotional/uncontrolled eating, even after accounting for BMI PGS. These findings enhance
305 our understanding of adolescent neurodevelopment related to eating disorder
306 symptomatology.

307

308 Unhealthy eaters were characterised by differing trajectories of ED symptoms during
309 adolescence, with increasing rates of binge eating, dieting and purging in
310 emotional/uncontrolled eaters, suggesting a greater risk for BN, in line with previous
311 research⁶. Unsurprisingly^{11,28}, higher levels of externalising problems in this group also
312 indicated that adolescents with ADHD and conduct disorder symptoms may be more likely to

313 engage in emotional/uncontrolled eating. Internalising problems also increased during
314 adolescence in unhealthy eaters, notably in emotional/uncontrolled eaters, highlighting the
315 close relationships between emotional eating and psychological well-being²⁹. While the
316 decreasing levels of internalising and externalising problems in healthy eaters replicate
317 previous findings³⁰, differing trajectories in unhealthy eaters, already evidenced at age 14,
318 indicate that internalising and externalising problems predate the development of ED
319 symptoms.

320

321 Consistent with cortical development trajectories^{31,32}, declines in structural brain measures
322 (except for increases in sulcal depth) were observed in all eating groups, with evidence of
323 protracted brain maturation in unhealthy eaters. Protracted brain maturation was also related
324 to internalising and externalising problems, corroborating findings from children³³, and
325 findings of delayed trajectories of cortical thinning in children and adolescents with ADHD³⁴.
326 The smaller volumetric reductions in the cerebellum observed in unhealthy eaters support its
327 role in eating pathology. Findings from anatomical, functional, and behavioural studies
328 indicate that the cerebellum is involved in the regulation of feeding behaviours and appetite
329 control^{35,36,37,38}. Disruption of a cerebellum-driven satiety network contributes to excessive
330 eating, difficulties in stopping eating, and weight gain³⁸. Our finding of protracted cerebellar
331 maturation in unhealthy eaters suggests disrupted development of such satiety network, and
332 engagement in restrictive eating as a compensatory mechanism to consciously control weight
333 gain. Additional alterations in reward (right putamen) and prefrontal (e.g., right PFC)
334 circuitry might lead emotional/uncontrolled eaters to also engage in disinhibited eating due to
335 impairments in self-regulation³⁹ and impulsive action control⁴⁰, making them more
336 susceptible to emotional and uncontrolled eating behaviours.

337

338 Genetic and environmental factors⁴¹ contribute to changes in developmental brain trajectories
339 related to eating. Genetic risk for obesity was previously found to correlate with disordered
340 eating and weight control behaviours in adolescents^{42,43}. High BMI was also found to be
341 associated with reduced cortical thinning in adolescence^{44,45}. Our findings extend these
342 analyses, indicating that genetic risk for obesity may influence eating-related weight control
343 behaviours partially via effects on cerebellar maturation, in a BMI-dependant manner.

344

345 Environmental stressors, such as adverse social environments and peer interactions, also
346 modulate brain development⁴⁶. Smaller GMV decreases in the cerebellum, PFC and anterior
347 cingulate, are observed in adolescents disliked by their peers, correlating with callous-
348 unemotional traits found in externalising disorders⁴⁷. This altered brain development may
349 underlie these adolescent adjustment problems and hypersensitivity to peer rejection⁴⁸. Our
350 findings suggest that adjustment problems influence brain maturation beyond genetic
351 predispositions for high BMI, increasing the risk for emotional/uncontrolled eating.

352

353 Our study's key strengths include a well-characterized, deeply phenotyped longitudinal
354 adolescent cohort and an innovative multivariate analytical approach. However, some
355 limitations exist. First, the analysis is based on participants of European ancestry,
356 necessitating future research with more ethnically diverse samples for broader applicability.
357 Second, the use of summary scores to assess eating behaviours may oversimplify complex
358 interactions and variations. Third, some analyses rely on VBM, which can impact GMV
359 estimates⁴⁹. Additionally, while we considered cortical thickness and sulcal depth, we did not

360 analyse surface area, a key volume component with distinct genetic architecture, reducing the
361 resolution of our genetic findings.

362

363 Nonetheless, our study sheds light on how genetic risk for higher BMI, along with increasing
364 internalising and externalising problems experienced during adolescence, distinctly
365 contributes to unhealthy eating through their effects on brain maturation. The implications of
366 these findings underscore the potential benefits of education targeting early maladaptive
367 coping mechanisms and dietary habits to prevent eating disorders while promoting brain
368 health.

369 **Methods**

370 **Participants**

371 Data analysed in this study was collected as part of IMAGEN, a longitudinal genetic ×
372 neuroimaging cohort study of adolescents recruited from eight study centres in England,
373 Ireland, France and Germany⁵⁰. Information on specific ethnic categories was not collected,
374 but the study, aimed at identifying the genetic and neurobiological basis of individual
375 variability in behaviour, was designed to include predominantly participants of European
376 ancestry (White), based on their self-reports. To further account for population stratification,
377 statistical approaches were applied to identify and exclude genetic ancestries other than
378 European, when analysing the genetic data. The IMAGEN study was approved by local
379 research ethics committees at each study site, and informed consent was obtained from
380 participants and their parents/guardians. A detailed description of the study protocol and data
381 acquisition can be found in⁵⁰. Specifically, the Strengths and Difficulties Questionnaire
382 (SDQ) questionnaire data used in this study were acquired at ages 14, 16, 19 and 23 years;
383 neuroimaging data ($N = 949$) were acquired at ages 14 and 23 years, and the Three-Factor
384 Eating Questionnaire (TFEQ) data were obtained at age 23 ($N = 996$).

385

386 **Neuropsychological assessments**

387 *Eating behaviours*

388 The short version (18 items) of the Three-Factor Eating Questionnaire (TFEQ) was used to
389 assess eating behaviours. It contains three subscales: cognitive restraint, which measures the
390 tendency to restrict one's food intake constantly and consciously instead of using
391 physiological cues, hunger and satiety, as regulators of food intake (6 items); emotional

392 eating, which reflects the tendency to eat in response to negative emotions (3 items); and
393 uncontrolled eating, which characterise the tendency to overeat, with the feeling of being out
394 of control (9 items). It has good structural validity and has been used and validated in
395 different European populations^{7,51}, and was found to distinguish different eating patterns in
396 the general population⁵².

397

398 *Eating disorder symptoms*

399 Dieting, binge eating and purging symptoms were assessed using the self-reports from the
400 eating disorder section (section P) of the Development and Well-being Assessment^{21,53}.

401 Dieting symptoms were evaluated based on responses to questions P18a, P18b, and P18c,
402 which asked about eating less at meals, skipping meals, and fasting, respectively. Binge
403 eating symptoms were assessed using the question P15, which inquired about eating a large
404 amount of food and losing control overeating. Purging symptoms were measured using the
405 questions P1c, P18f, and P18g, which asked about self-induced vomiting or taking pills or
406 medicines to lose weight.

407

408 *Emotional and behavioural problems*

409 The Strengths and Difficulties Questionnaire (SDQ) was used to assess emotional and
410 behavioural problems in adolescents. It has five hypothesised subscales, including emotional
411 symptoms, conduct problems, hyperactivity/inattention, peer relationship problems and
412 prosocial behaviours⁵⁴. In low-risk and general population samples, the emotional and peer
413 subscales can be combined into an “internalising” subscale (10 items) and the behavioural
414 and hyperactivity subscales into an “externalising subscale (10 items), respectively⁵⁵. We

415 used self-reported scores at ages 14, 16, 19 and 23 years for internalising problems (IP) and
416 externalising problems (EP) in further analyses.

417

418 **Structural magnetic resonance imaging acquisition and processing**

419 MRI images were acquired with 3T MRI scanners from different manufacturers (Siemens,
420 Munich, Germany; Philips, Best, The Netherlands; General Electrics, Chalfont St Giles, UK;
421 Bruker, Ettlingen, Germany) from eight IMAGEN recruitment sites. The high-resolution
422 anatomical MRI images acquired included a three-dimensional T1-weighted magnetisation
423 prepared gradient echo sequence (MPRAGE) based on the ADNI protocol
424 (<http://adni.loni.usc.edu/methods/documents/mri-protocols/>), T2 weighted fast-spin echo, and
425 FLAIR scans for visual assessment.

426

427 All raw images were visually inspected to exclude images with movement artefacts, brace
428 artefacts, or field inhomogeneities prior to pre-processing. The pre-processing procedures
429 were then conducted using the Computational Anatomy Toolbox (CAT 12.8 (r1907);
430 <https://neuro-jena.github.io/cat/>) in SPM 12 (Wellcome Department of Cognitive Neurology).
431 We used the “segment longitudinal data” procedure with default settings. Intra-subject co-
432 registration was performed on the baseline (at age 14) and follow-up (at age 23) images. The
433 co-registered images were then realigned across participants and bias-corrected with
434 reference to the mean images computed from each subject’s baseline and follow-up images.
435 Next, the baseline and follow-up images as well as their means images were segmented into
436 grey matter, white matter and cerebrospinal fluid based on the default tissue classification
437 map. Diffeomorphic Anatomical Registration Through Exponentiated Lie Algebra
438 (DARTEL) normalisation was subsequently performed on the segmented mean images using

439 the default DARTEL template. The derived spatial normalisation parameters were then
440 applied to transform the segmented subject baseline and follow-up grey matter images into
441 the standard Montreal Neurological Institute (MNI) space. All normalised grey matter images
442 were finally smoothed with an isotropic Gaussian kernel of 6 mm full width at half maximum
443 (FWHM). The quality measures created during pre-processing for each participant at each
444 time point were examined, and images with sufficient quality (corresponding to grade D or
445 above) were included in further analyses. Changes in grey matter volumes (GMV) were
446 analysed using whole-brain voxel-based morphometry (VBM). Measures of cortical thickness
447 (CT) and sqrt-transformed sulcal depth (SD), which were then resampled to 12 mm in line
448 with the recommendation for surface measures, were also derived. Longitudinal changes in
449 mean CT and SD were extracted for different regions of interest (ROIs) using the Desikan-
450 Killiany atlas (N of ROIs = 68).

451

452 **BMI Polygenic scores (PGS)**

453 A total of 2,087 participants were genotyped with the Illumina Human610-Quad BeadChip
454 and Illumina Human660-Quad BeadChip during the baseline assessments. Stringent quality
455 control procedures were performed prior to imputation (**Supplementary Information**). In
456 brief, multi-dimensional scaling analysis and principal component analysis were conducted to
457 identify genetic ancestry. Participants who were outliers from the European superpopulation
458 were excluded (**Supplementary Fig. S4 and S5**) due to the limited portability across
459 ancestries for polygenic scores. Consequently, 1,899 Participants (49.66% males) who passed
460 genotyping QC and were identified as of European ancestry were selected for generating the
461 BMI PGS. IMAGEN genotype data was integrated into the European ethnicity 1KGP (phase
462 3 release v5) reference panel⁵⁶ for imputation. Summary statistics of genome-wide

463 association study (GWAS) of BMI from ~681,275 individuals of European ancestry⁵⁷ was
464 used to calculate BMI PGS. This was achieved using PRS-CS⁵⁸, which utilises high-
465 dimensional Bayesian regression and a continuous shrinkage prior to SNP effect sizes. The
466 global shrinkage parameter was set to 0.01 as recommended for highly polygenic traits. A
467 total of 905,362 SNPs were used to predict BMI PGS. Participants with available TFEQ
468 scores were included in the subsequent analyses (RE: $N = 255$; E/UE: $N = 194$; HE: $N = 347$).
469 The BMI PGS was residualised for the first 10 PCs and batch effects before being Z-scored
470 for the subsequent analyses.

471

472 **Statistical analyses**

473 *Identification of groups with distinct eating behaviours by K-means clustering*

474 K-means clustering using the TFEQ subscale scores (i.e., cognitive restraint, emotional eating
475 and uncontrolled eating) at age 23 was performed to identify clusters exhibiting different
476 eating behaviours. All continuous variables were transformed into z-scores. We used the
477 “NbClust” package to identify the optimal cluster number and validity of the cluster solution,
478 and the “fpc” package to examine the clustering stability with the Jaccard coefficient and a
479 bootstrap technique ($N = 1000$) in R.

480

481 *Group differences in trajectories of eating disorder symptoms across adolescence*

482 Linear mixed models were used to examine group differences in the trajectories of dieting,
483 binge eating and purging from ages 14 to 23. Age (i.e., 14, 16, 19 and 23) was treated as a
484 categorical variable. The models included age, group, and age-by-group interactions as fixed
485 effects and adjusted for sex. Random intercepts for participants nested within recruitment

486 sites accounted for the dependence of repeated measures. Group-by-age interactions were
487 investigated using healthy eaters (HEs) and age 14 as reference. A Bonferroni correction
488 accounting for 18 tests (3 ED symptoms \times 2 groups comparisons \times 3 age comparisons) was
489 applied (i.e., $p_{\text{Bonferroni}} = 2.78 \times 10^{-3}$).

490

491 *Group differences in trajectories of internalising and externalising problems*

492 The latent growth curve models (LGCM) using the “lavaan” package in R were conducted in
493 these analyses.

494 Univariate LGCM analyses: Latent factors of intercept and slope were estimated for repeated
495 measures (at ages 14, 16, 19, and 23 years) of IP and EP scores separately. Sex, groups and
496 recruitment sites were considered time-invariant covariates. For these analyses, we included
497 only those participants who had TFEQ scores at age 23 and had at least one measure of IP or
498 EP at ages 14, 16, 19 and 23. For both IP and EP, we attempted to fit a quadratic term;
499 however, this specification resulted in a non-positive definite covariance matrix, driven by a
500 correlation greater or equal to one between the linear and quadratic terms. Hence, we decided
501 not to include a quadratic term as the information contained within it was not adding any
502 extra information over the linear term. The full information maximum likelihood estimator
503 was utilised to account for data missing at random. We investigated group differences in
504 intercepts and slopes of IP and EP trajectories, taking HEs as a reference. Bonferroni-
505 corrected p -value threshold of $0.05/(2 \text{ behaviours} \times 2 \text{ measures} \times 2 \text{ groups}) = 6.25 \times 10^{-3}$ was
506 considered statistically significant.

507 Multivariate LGCM analyses within each group were also run to estimate models for IP and
508 EP trajectories simultaneously and investigate covariances between latent factors (i.e., IP

509 intercept, IP slope, EP intercept, and EP slope). Sex and recruitment sites were included as
510 covariates.

511

512 *Group differences in brain maturation: Longitudinal MRI analyses*

513 Participants were excluded from the analysis if they had missing MRI data or failed to meet
514 quality control criteria ($N = 47$; see **Methods** for image pre-processing and QC).

515 Consequently, a total of 949 participants (306 REs, 236 E/UEs, and 407 HEs) were included
516 in the whole-brain VBM analysis and linear mixed models for cortical thickness (CT) and
517 sulcal depth (SD).

518 VBM analysis: longitudinal whole-brain voxel-based morphometry (VBM) analyses were
519 performed using the CAT 12.8 (r1932) toolbox. To identify brain regions reflecting
520 significant changes in grey matter volumes (GMVs) between ages 14 and 23 among the
521 groups identified above, we performed a 2×2 mixed ANOVA on the smoothed images using
522 the “Flexible Factorial” model. The two factors were age (age 14 or age 23; within-subject)
523 and group (i.e., comparison of each of 2 groups, namely REs versus HEs, E/UEs versus HEs,
524 or REs versus E/UEs; between-subject). Intracranial volumes (TIVs) were estimated by CAT
525 12.8 as the sum of the grey matter, white matter and cerebrospinal fluid volume. Analyses
526 were controlled for the effects of participants’ sex, the scanning site and TIV at each time
527 point (at ages 14 or 23). An absolute threshold masking of 0.1 was applied. The grey matter
528 morphological differences showing significant age-by-group interactions were reported after
529 a cluster-level family-wise error correction with a p -value < 0.05 , and a cluster-forming
530 threshold of p -value < 0.001 without correction.

531 Linear mixed models: For group differences in changes in CT and SD, we performed regions
532 of interest (ROIs)-based linear mixed models, investigating interactions between age and

533 groups. Models included age, groups, and their interactions as fixed effects, with the
534 participant nested within recruitment sites as a random effect and adjusted for sex. For both
535 measures, the Bonferroni correction was applied to adjust for multiple testing ($p = 0.05/68$
536 ROIs \times 3 group comparisons = 2.45×10^{-4}).

537

538 *Mediation analyses*

539 Simple mediation models were performed using the PROCESS v4.0 macro for R to test
540 whether the between-group differences in brain changes mediated the relationships between
541 differences in IP or EP trajectories and eating behaviours. We refer to this model as the
542 “psychopathology-brain maturation-eating behaviours” model). Brain clusters that
543 significantly differentiated REs from HEs, E/UEs from HEs, or REs from E/UEs were
544 considered regions of interest (ROIs). For group comparisons involving several brain
545 clusters, these clusters were combined into a single ROI for each structural measure (GMV,
546 CT or SD). For comparisons between REs and HEs, one mediation model was tested –
547 relating GMV differences in the left cerebellum to differences in IP slope–, therefore, a p-
548 value threshold of 0.05 was considered significant. Comparisons between E/UEs and HEs:
549 these two groups differed behaviourally in IP intercept, IP slope, and EP intercept, and in
550 their changes of GMVs, CTs, and SDs, therefore 9 mediation models were tested. The
551 Bonferroni-corrected significance threshold of $0.05/(3 \text{ trajectory measures} \times 3 \text{ structural}$
552 $\text{brain measures}) = 5.56 \times 10^{-3}$ was applied.

553 Subsequent analyses investigated the potential contributions of the BMI PGS on brain
554 mediation models identified above, referred to as the “genetics-brain maturation-eating
555 behaviours” models. The same brain ROIs were considered as mediators in these models. For
556 models comparing REs to HEs, the significance was set at $p = 0.05/(1 \text{ structural measure}) =$

557 0.05. For models comparing E/UEs to HEs, the significance was set at $p = 0.05/3$ structural
558 measures = 1.67×10^{-2} .

559

560 Multivariate mediation models were conducted using AMOS 29, to explore the unique
561 contributions of brain ROIs, psychopathology (IP and EP trajectories) and BMI PGS to
562 mediation models identified in simple mediation analyses, referred to as the “genetics-
563 psychopathology-brain maturation-eating behaviours” model. Continuous variables were
564 transformed into z-scores for these analyses. Confidence intervals (CIs) for the mediation
565 effect were estimated from 5000 bootstrap samples.

566

567 *Covariates*

568 Covariates for all analyses included sex and recruitment sites. For analyses involving GMV,
569 total intracranial volume (TIV) at the corresponding age was additionally included as a
570 covariate. Since there were no significant group differences in age at each data collection, age
571 was considered as a categorical variable in the linear mixed models, and as time points in
572 repeated measures in the LGCM analysis. For the longitudinal MRI analysis (VBM analysis
573 and linear mixed models), participants nested within recruitment were modelled as a random
574 effect, and sex was considered a fixed effect in the model.

575 *Other covariates:* To examine the robustness of findings from our primary analyses,
576 sensitivity analyses were conducted by including pubertal status, IQ, educational attainment
577 (EA), and age- and sex-adjusted BMI as additional covariates. Pubertal status was assessed
578 using the Puberty Development Scale⁵⁹, an 8-item self-report measure of physical
579 development based on Tanner stages, separate for males and females. IQ was calculated as
580 the average of Perceptual Reasoning Index (PRI) and Verbal Comprehension Index (VCI)

581 scores based on age norms using the Wechsler Intelligence Scale for Children, Fourth Edition
582 (WISC-IV; Pearson Clinical Assessment UK). We administered the block design, matrix
583 reasoning, similarities and vocabulary subtests. Raw scores from each subtest were converted
584 into scaled scores based on age norms. For both the PRI and VCI, we calculated prorated
585 sums of scaled scores and then converted these sums into index scores according to the
586 WISC-IV manual. EA was assessed by self-report of the “average grade at the end of the last
587 term completed”. The age- and sex-adjusted BMI Z-score at age 14 was calculated using the
588 jBmi R package based on the CDC recommendations.

589

590 **Data availability**

591 Access to individual-level data from the IMAGEN project is accessible to bona fide
592 researchers upon reasonable request and approval of a project proposal by IMAGEN
593 consortium PIs. Contact the corresponding author for requests related to this study.

594 Summary statistics from the BMI GWAS, used in this study for computing BMI PGS, are
595 accessible via⁵⁷, and can be downloaded from their website at

596 [https://portals.broadinstitute.org/collaboration/giant/index.php/GIANT_consortium_data_file](https://portals.broadinstitute.org/collaboration/giant/index.php/GIANT_consortium_data_files)

597 s. Data from the 1000 Genomes Project Phase 3 may be accessed from

598 <https://www.internationalgenome.org/category/phase-3/>.

599

600 **Code availability**

601 R (version 4.3.2) was employed to perform ANOVA and mixed linear model analysis.

602 Mediation analysis was performed with the PROCESS R macro (version 4.3) and AMOS

603 (version 20.0). MATLAB (version: 2022b) was employed to process the structural MRI

604 images and perform VBM analysis. The polygenic score was calculated using the publicly
605 available PRS-cs toolbox (<https://github.com/getian107/PRScs>) and LD references
606 constructed using the 1000 Genome Project phase 3 samples with European ancestry. MRICro
607 (version 1.9.1; <https://www.nitrc.org/projects/micro>), BrainNet Viewer (version: 20191031;
608 <https://www.nitrc.org/projects/bnv>) and ENIGMA visualisation tool
609 (<https://github.com/MICA-MNI/ENIGMA/tree/master/enigmatoolbox>) were used to visualise
610 images. Other scripts used to analyse the data of this study are publicly available
611 [https://github.com/XinyangYu918/EatingBehaviours-BrainMaturation-Psychopathology-](https://github.com/XinyangYu918/EatingBehaviours-BrainMaturation-Psychopathology-Genetics)
612 [Genetics](https://github.com/XinyangYu918/EatingBehaviours-BrainMaturation-Psychopathology-Genetics).

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653

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656

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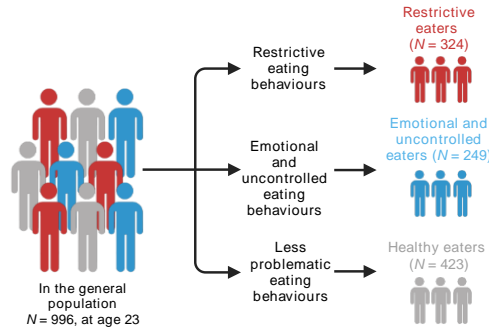
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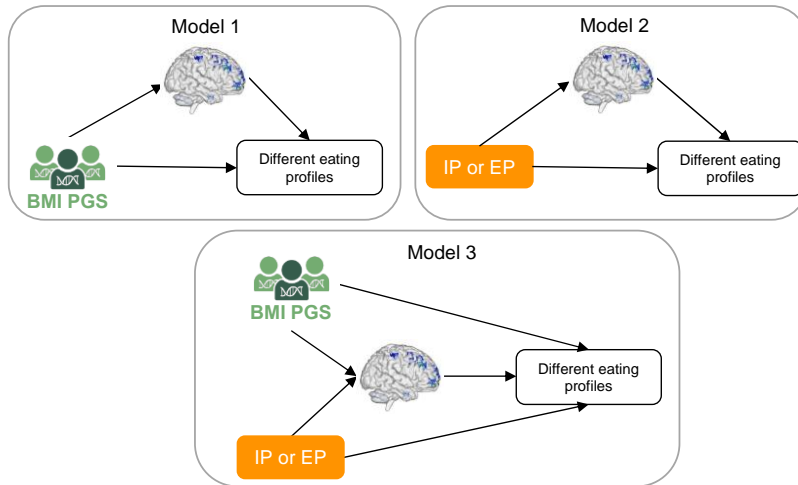
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832 **Figures and figure legends**

Step 1: Identification of distinct eating profiles in the general population



Step 3: Cross-group investigations of “genetics-brain maturation-eating behaviours” and “psychopathology-brain maturation-eating behaviours” models

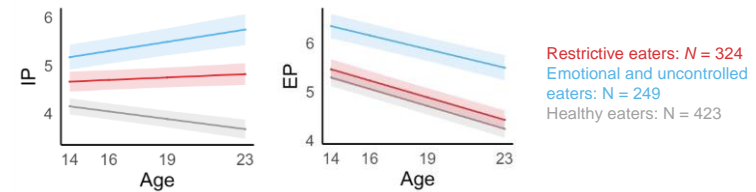


Step 2: Multi-modal characterisation of different eating profiles

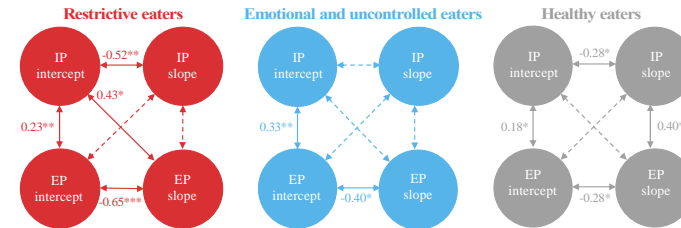
Genetic level: BMI PGS ($N = 1899$)

Psychopathological level: trajectories of internalising (IP) and externalising problems (EP)

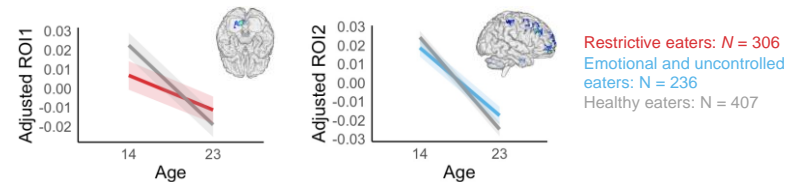
• Between-groups:



• Within-groups:

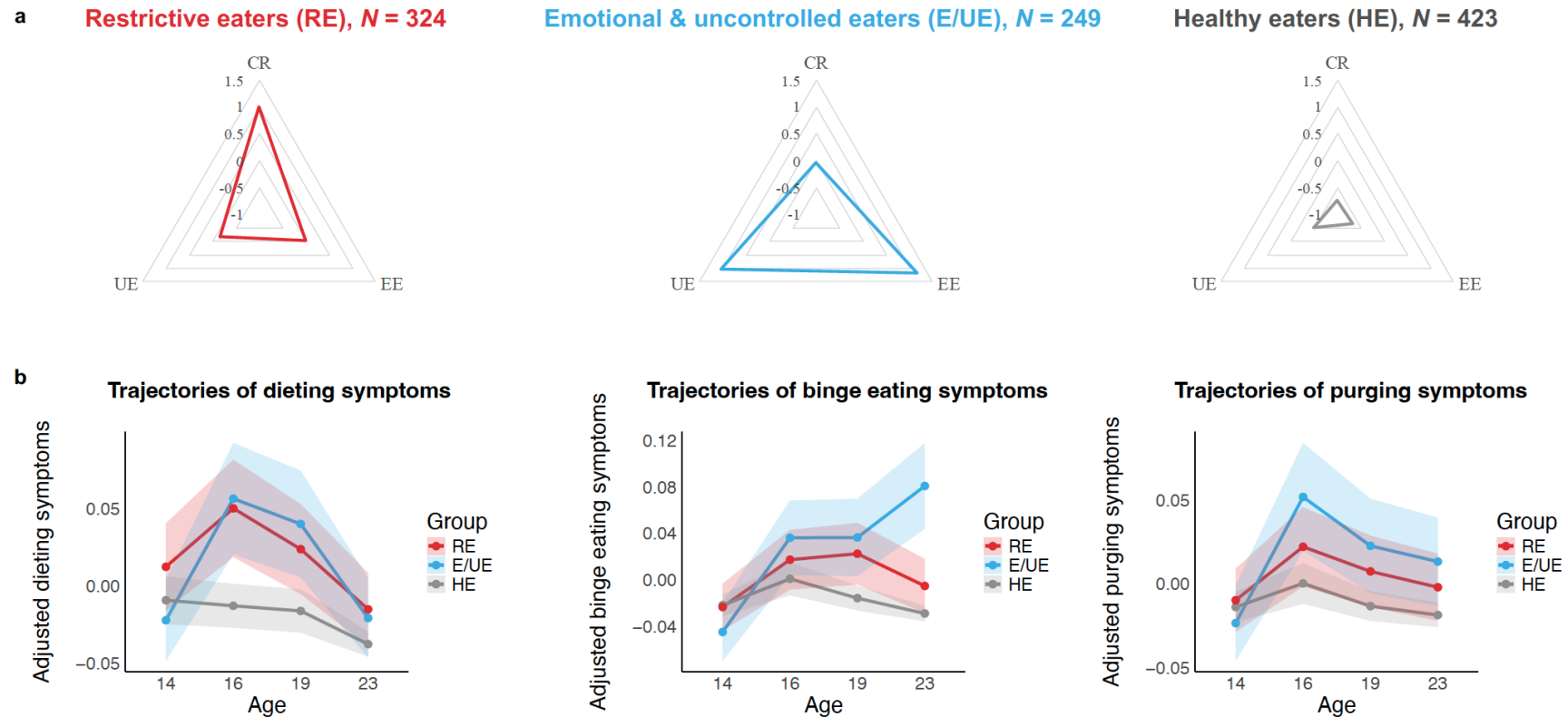


Neurobiological level: age × group interactions in grey matter volume (GMV), cortical thickness (CT), and sulcal depth (SD), between ages 14 and 23, $N = 949$



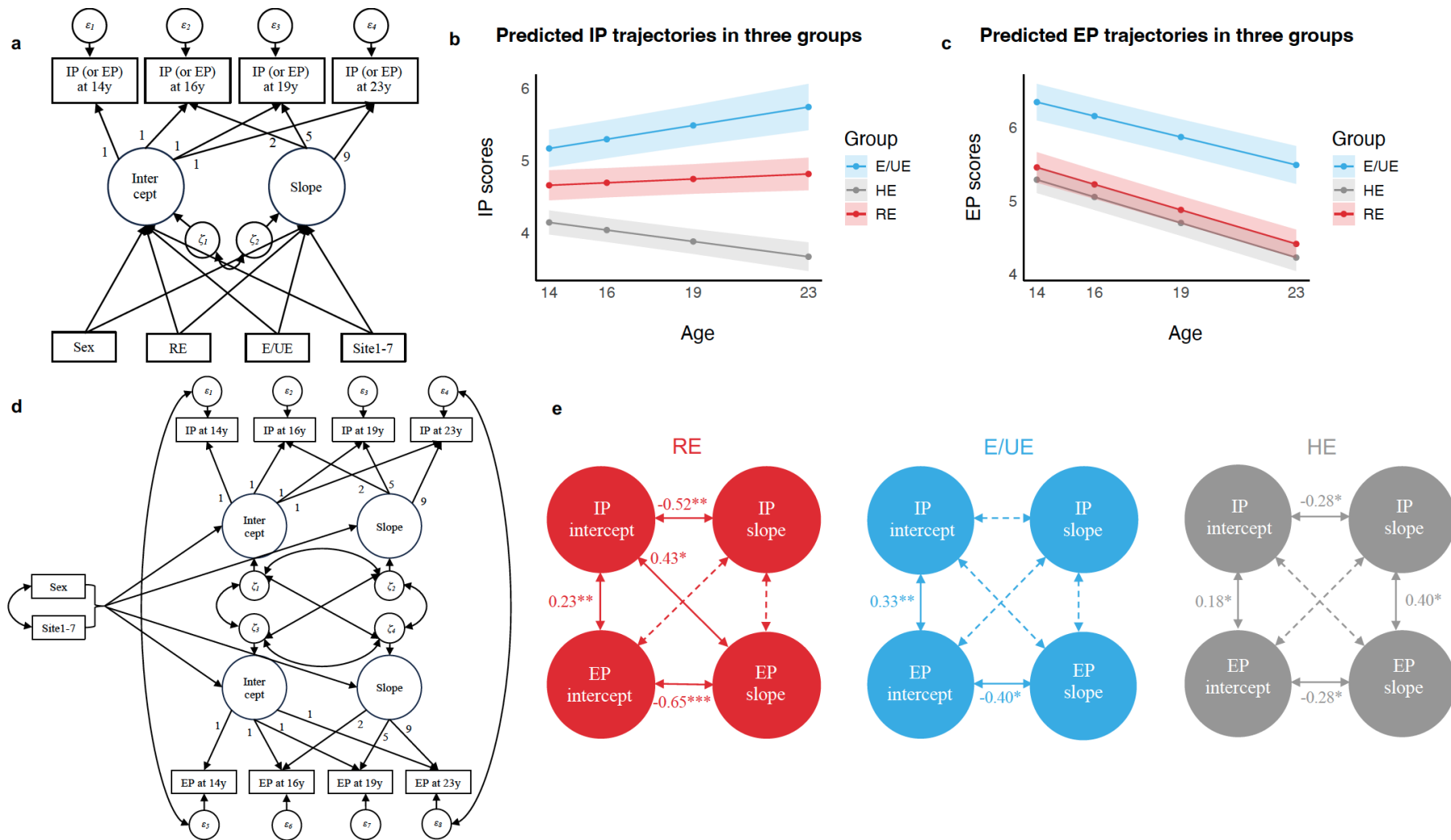
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834 **Fig. 1.** Workflow of research questions and analyses.



835

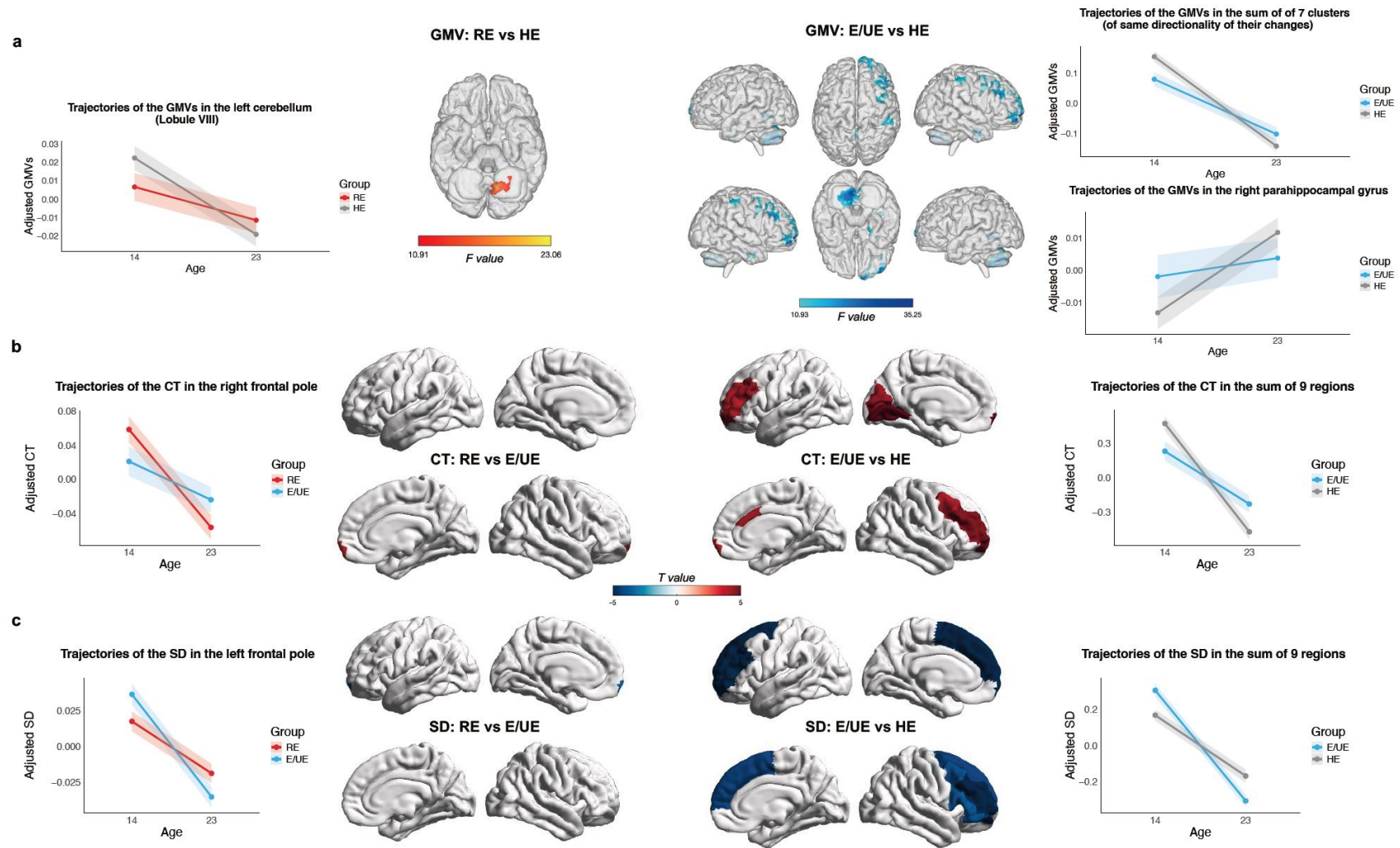
836 **Fig. 2. a**, Three groups of distinct eating behaviour profiles were identified by K-means clustering at age 23. CR, cognitive restraint; EE,
 837 emotional eating; UE, uncontrolled eating. The radar charts display the average standardised scores for CR, EE, and UE across these groups. **b**,
 838 Trajectories of eating disorder symptoms (dieting, binge eating and purging symptoms) from ages 14 to 23 across the three identified groups.
 839 The lines represent the mean symptom scores, and the shaded areas show the 95% confidence intervals. Analyses were adjusted for sex and
 840 recruitment sites. The y-axis indicates the adjusted symptoms scores after regressing out the effects of sex and recruitment sites.



841

842 **Fig. 3.** Psychopathological characterisation across groups using internalising and externalising problems. **a**, Path diagram for the conditional
 843 linear latent growth curve model for individual IP and EP trajectories among three groups. Two dummy variables, RE and E/UE, were included

844 to represent three groups (HE was considered the reference group in the model). Gender and seven dummy variables for different recruitment
845 sites were included as covariates in the analysis. Post-hoc analyses were conducted to examine the differences between the RE and E/UE groups.
846 **b**, Predicted trajectories of IP across three groups. **c**, Predicted trajectories of EP across three groups. **d**, Path diagram for the multivariate latent
847 growth curve model for each group separately. **e**, Significant within-construct and cross-construct correlations between IP and EP trajectories
848 were found within each group. The numbers indicate standardised covariances (i.e., correlations) values. *, $p < 0.05$, **, $p < 0.01$, ***, $p <$
849 0.001.

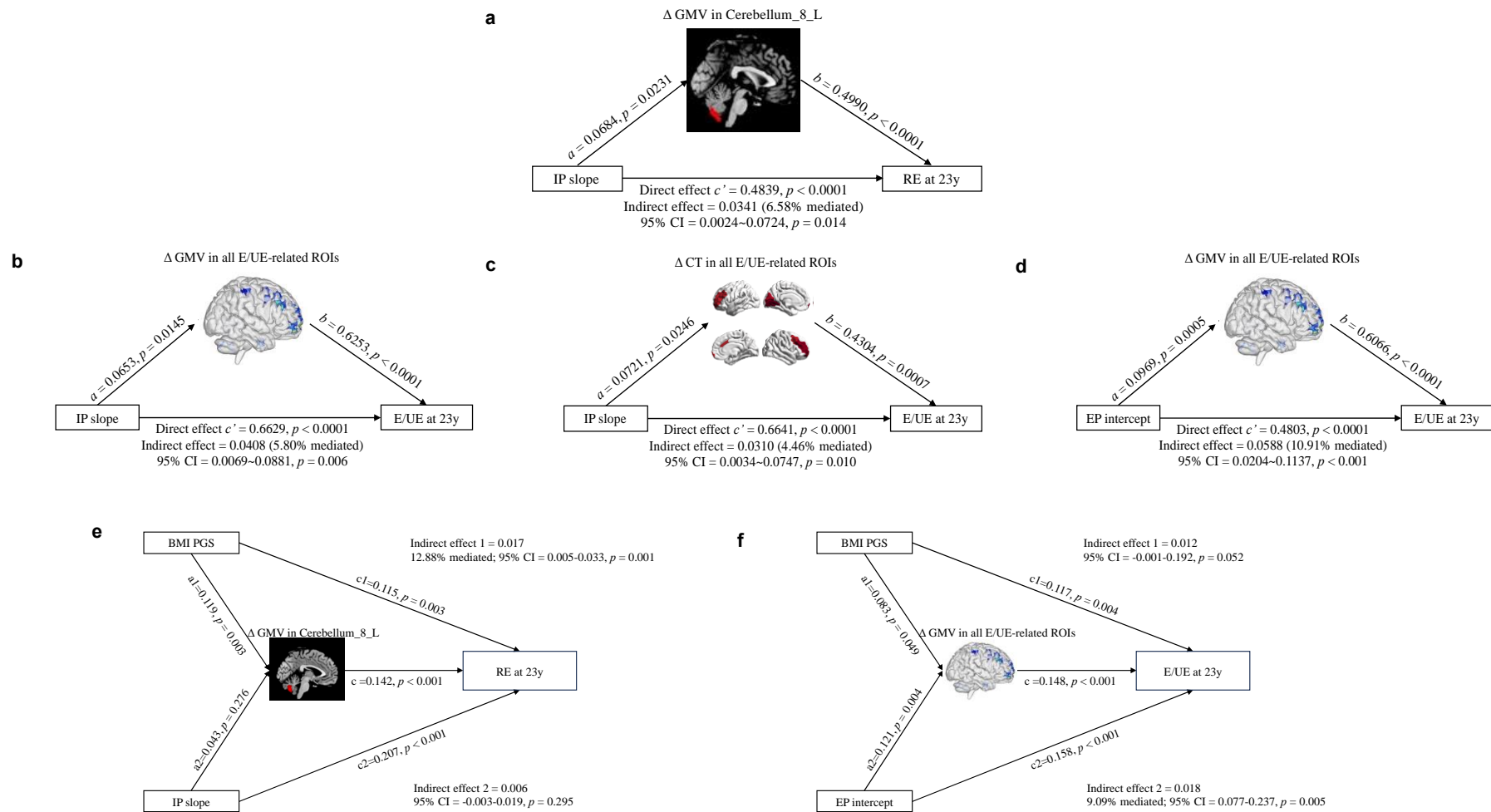


850

851 **Fig. 4.** Significant age-by-group interactions were identified in various brain morphological features, including grey matter volume (GMV; **a**),
 852 cortical thickness (CT; **b**) and sulcal depth (SD; **c**). **a**, A significant age-by-group interaction was observed between REs and HEs, ages 14 to 23,

853 indicating that REs experienced a smaller volumetric reduction in the left cerebellum compared to HEs over time. E/UEs exhibited less
854 volumetric reductions in multiple brain regions, including the left cerebellum, right middle frontal gyrus, right medial superior frontal gyrus,
855 right postcentral gyrus, and right putamen compared to HEs. However, E/UEs had volumetric increases in the right parahippocampal gyrus
856 compared to HEs. **b**, Comparisons of CT trajectories between REs and E/UEs showed that REs experienced more pronounced CT reductions in
857 the right frontal pole. CT trajectory comparisons between HEs and E/UEs revealed that HEs experienced more pronounced CT reductions across
858 a wide range of brain regions, including the frontal and occipital gyri, and the anterior cingulate gyrus, compared to E/UEs. **c**, SD trajectory
859 comparisons between REs and E/UEs indicated that REs had less pronounced SD reductions in the left frontal pole. Comparisons of SD
860 trajectories between HEs and E/UEs revealed that HEs had less pronounced SD reductions in the frontal and temporal regions. All analyses were
861 adjusted for sex, recruitment sites, and total intracranial volume. The lines in the figures represent mean values for brain morphological features
862 and the shaded areas represent the corresponding 95% confidence intervals. The y-axis indicates the adjusted symptom scores after regressing
863 out the sex, recruitment sites, and total intracranial volume.

864



865

866 **Fig. 5.** The “psychopathology-brain maturation-eating behaviours” and “BMI PGS-brain maturation-eating behaviours” models. **a**, Mediation
 867 effect of GMV reductions in the left cerebellum (ages 23-14) on the relationship between age-related IP (IP slope) and RE at age 23. **b**,

868 Mediation effects of GMV reductions on the relationship between age-related IP changes (IP slope) and E/EU at age 23, considering all E/UE-
869 related ROIs (except right parahippocampal gyrus, due to the different directionality of its changes compared to other ROIs). **c**, Mediation
870 effects of CT reductions on the relationship between age-related IP changes (IP slope) and E/EU at age 23, considering all E/UE-related ROIs. **d**,
871 Mediation effects of GMV reductions on the relationship between EP intercept and E/EU at age 23, considering in all ROIs (except right
872 parahippocampal gyrus, due to the different directionality of its changes compared to other ROIs). **e**, The unique contribution of smaller
873 cerebellar GMV reductions in mediating the effects of BMI PGS on restrictive eating, when IP trajectory was also included in the model. **f**, The
874 unique contribution of GMV changes in all E/UE-related ROIs (except right parahippocampal gyrus) in the relationship between EP intercept
875 and E/UE at age 23, beyond the effects of BMI PGS. All mediation models were adjusted for gender, recruitment sites, and TIV differences
876 (between ages 14 and 23).

877

878

Table 1. Sample sizes and demographic characteristics of the three groups of participants with distinct eating profiles at age 23.

	Restrictive eaters, REs (<i>N</i> = 324)	Emotional and uncontrolled eaters, E/UEs (<i>N</i> = 249)	Healthy eaters, HEs (<i>N</i> = 423)	<i>F</i> / χ^2	<i>p</i>	Post hoc tests (Bonferroni corrected)
Age at data collection: Mean (SD)						
Baseline	14.56 (0.43)	14.49 (0.41)	14.49 (0.42)	4.93	0.027	<i>NS</i>
Follow-up 1	16.46 (1.15)	16.26 (1.66)	16.44 (1.39)	0.00	0.997	-
Follow-up 2	19.37 (0.95)	19.29 (1.01)	19.30 (0.98)	0.70	0.403	-
Follow-up 3	22.75 (0.74)	22.68 (0.72)	22.66 (0.73)	2.81	0.094	-
Male/female (% females)	133/191 (58.95%)	92/157 (63.05%)	253/170 (40.19%)	42.10	< 0.001	REs vs HEs: <i>p</i> < 0.001; E/UEs vs HEs: <i>p</i> < 0.001
Eating behaviours: Mean (SD)						
Cognitive restraint	17.55 (3.03)	12.90 (3.70)	9.78 (2.42)	623.94	< 0.001	REs > E/UEs: <i>p</i> < 0.001; REs > HEs: <i>p</i> < 0.001; E/UEs > HEs: <i>p</i> < 0.001
Emotional eating	5.47 (1.76)	8.34 (2.13)	3.87 (1.22)	567.89	< 0.001	E/UEs > REs: <i>p</i> < 0.001; E/UEs > HEs: <i>p</i> < 0.001, REs > HEs: <i>p</i> < 0.001
Uncontrolled eating	18.84 (3.53)	25.15 (4.25)	17.06 (4.52)	307.45	< 0.001	E/UEs > REs: <i>p</i> < 0.001; E/UEs > HEs: <i>p</i> < 0.001, REs > HEs: <i>p</i> < 0.001
Developmental stage: Mean (SD)						
Pubertal status, at age 14	3.03 (0.53)	2.99 (0.53)	2.79 (0.58)	34.47	< 0.001	REs > HEs: <i>p</i> < 0.001; E/UEs > HEs: <i>p</i> < 0.001
Cognition: Mean (SD)						
IQ ^a	109.91 (12.51)	108.51 (12.93)	112.12 (12.85)	6.04	0.014	E/UEs < HEs: <i>p</i> = 0.0018
Educational attainment	2.78 (1.32)	3.02 (1.62)	2.66 (1.18)	1.88	0.170	-
Anthropometric variable: Mean (SD)						
BMI-Zscore, at age 14 ^b	0.43 (0.77)	0.27 (0.90)	-0.17 (0.93)	83.34	< 0.001	REs > HEs: <i>p</i> < 0.001; E/UEs > HEs: <i>p</i> < 0.001
N (%) of available MRI data for longitudinal MRI analysis ^c	306 (94.44%)	236 (94.78%)	407 (96.22%)			
Polygenic scores: Mean (SD)						
N (%) of available genotyping data of European ancestry	285 (87.96%)	220 (88.35%)	376 (88.89%)			
BMI PGS-Zscore ^d	0.03 (1.02)	0.08 (0.88)	-0.24 (0.96)	13.56	< 0.001	REs > HEs: <i>p</i> = 0.001; E/UEs > HEs: <i>p</i> < 0.001

^aIQ was calculated as the average of the Perceptual Reasoning Index and Verbal Comprehension Index scores based on age norms using the Wechsler Intelligence Scale for Children, Fourth Edition (WISC-IV; Pearson Clinical Assessment UK). ^bAge- and sex-adjusted BMI-Zscore was calculated using the jBMI R package (<https://github.com/jbirstler/jBmi>) based on the CDC recommendations. ^cParticipants were excluded from the analysis if they had missing MRI data or failed to meet quality control criteria ($N = 47$; see Methods for image pre-processing and QC). ^dParticipants who passed genotyping QC (see Supplementary Information) and were identified as of European ancestry were selected for generating the BMI PGS. BMI polygenic score (BMI PGS) was calculated using the publicly available PRSCs and 1000 Genome Project 3 European LD panels (<https://github.com/getian107/PRSCs>) and adjusted for the first 10 principal components of genetic ancestry and batch effects. The scores were Z-scored prior to analysis.