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2   **Gene-exercise interaction on brain health in children with**
3   **overweight/obesity: The ActiveBrains randomized controlled**
4   **trial**

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ABSTRACT

We investigated the interaction between a genetic score and an exercise intervention on brain health in children with overweight/obesity. One hundred one children with overweight/obesity (10.0 ± 1.5 years, 59% girls) were randomized into a 20-week combined exercise intervention or a control group. Several cognitive and academic outcomes were measured with validated tests. Hippocampal volume was quantified using magnetic resonance imaging. Six brain health-related polymorphisms (rs6265 [*BDNF*], rs2253206 [*CREB1*], rs2289656 [*NTRK2*], rs4680 [*COMT*], rs429358, and rs7412 [*APOE*]) were genotyped. Cognitive flexibility and academic skills improved significantly more in the exercise than in the control group only in the children with a "favorable" genetic profile (mean z score, 0.41-0.67 [95% CI 0.11 to 1.18], yet not in those with "less favorable" genetic profile. An individual response analysis showed that children responded to exercise in cognitive flexibility only in the "genetically favorable" group (i.e. 62% of them had a meaningful [≥0.2 Cohen *d*] increase in the exercise group compared with only 25% in the control group). This finding was consistent in per-protocol and intention-to-treat analyses (P=0.01 and P=0.03, respectively). The results were not significant or not consistent for the rest of outcomes studied. Our findings suggest that having a more favorable genetic profile makes children with overweight-obesity more responsive to exercise, particularly for cognitive flexibility.

Keywords: Genetics; cognition; pediatrics; physical activity; fitness

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91   **NEW & NOTEWORTHY**: Inter-individual differences have been reported in brain

92   health-related outcomes in response to exercise interventions in adults, which could be

93   partially explained by genetic background differences. However, the role of genetic

94   polymorphisms on brain health-related outcomes in response to exercise interventions

95   remains unexplored in pediatric population. The current study in children with

96   overweight/obesity showed that a genetic score composed of six brain health-related

97   polymorphisms (*BDNF, CREB1, NTRK2, COMT, and APOE*) regulated the exercise-

98   induced response on several brain health comes, yet mainly and more consistently on

99   cognitive flexibility.

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1. Introduction

The prevalence of childhood obesity worldwide increased from 4 to 18% between 1975 and 2016 (1). Excess of body weight/adiposity during childhood is inversely associated with children's brain health indicators, including a poor cognitive and academic performance, and reduced gray matter volume (2–4). Several randomized controlled trials (RCTs) reported that regular exercise counteracts the negative impact of childhood obesity on brain health (5–8). Indeed, our recent RCT showed the positive effects of a 20-week combined exercise intervention on a broad set of brain health indicators such as intelligence, cognitive flexibility, and academic performance in children with overweight/obesity (OW/OB) (9).

A consensus statement reported considerable interindividual differences in the response to exercise interventions on cardiometabolic risk factors and cardiorespiratory fitness (10). Nevertheless, less is known on the interindividual response to exercise interventions on brain health. Yu *et al*. reported interindividual differences brain health indicators in response to an aerobic exercise intervention in older adults (11). Genetic background, including single nucleotide polymorphisms (SNPs), could partially explain these interindividual differences in response to exercise interventions (10, 12). In this regard, Stroth *et al*. reported that 17 weeks of running training improved cognitive function in young adults presenting the Val/Val genotype for the SNPs rs4680 in Catechol-O-methyltransferase (*COMT*) compared to peers with the Met/Met genotype (13). On the other hand, SNPs (rs6265, rs429358, and rs7412) in brain-derived neurotrophic factor (*BDNF*) and apolipoprotein E (*APOE*) did not explained inter-individual differences in motor status and brain volume in response to motor

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rehabilitation therapy after stroke (14). Several genes have an impact on complex phenotypes and they show small effect sizes, which when combined may have an impact (12, 15). Thus, the use of genetic scores could be more powerful than individual SNPs to reveal the influence of genetic constitution on brain health in response to exercise interventions. However, to our knowledge, the interaction between a genetic score based on SNPs genes related to brain health and the effects of an exercise intervention on brain health remains unexplored in pediatric population.

Previous literature has identified several genes that encode proteins related to cognition or brain outcomes (**Table S1**). For instance, BDNF is one of the most investigated neurotrophic factor involved in neuronal physiology and cognition (16, 17). Circulating BDNF levels are reduced in patients with neurodegenerative diseases and obesity (18, 19), while RCTs reported that exercise might increase circulating BDNF protein levels (18, 20). BDNF binds to the neurotrophic tyrosine kinase receptor 2 (NTRK2; also called TrkB) on neurons inducing the activation of downstream pathways such as PI3K-AKT and Ras-MAPK, modulating the neurotransmitter release in hippocampal neurons(16). Importantly, cAMP-responsive element binding proteins (CREB) are a transcription factor family considered the main regulator of BDNF expression at the transcriptional level in cortical neurons (21). Also, APOE plays an important role through the transport of cholesterol and peptides with involved in cognitive function, such as amyloid beta(22), which can inactivate the PKA/CREB pathway (23). COMT is an enzyme that regulates dopamine levels in different brain regions, such as the prefrontal cortex (24). Indeed, dopamine can increase BDNF production in a concentration-dependent manner in hippocampal tissue and may influence cognition (25).

160   This study aimed to examine the interaction between genetic background,

161   specifically brain health-related SNPs in a set of candidate genes (*BDNF*, *CREB1*,

162   *NTRK2*, *COMT* and *APOE*) and the effects of a 20-week exercise intervention on

163   primary brain health outcomes (i.e., intelligence, executive function [cognitive

164   flexibility, inhibition, and working memory], academic performance, hippocampal

165   volume) in children with OW/OB from the ActiveBrains RCT. Thus, we hypothesized

166   that the brain health-related genetic background would play a role in the effect of the

167   ActiveBrains exercise intervention on brain health outcomes.

168

169   **2. Methods**

170   *2.1 Study design and participants*

171   From a total of 109 participants randomized in the ActiveBrains RCT (9, 26), 101

172   children with valid and complete genetic data were included in this study. The primary

173   aim of this project was to study the effects of the exercise program on brain health

174   outcomes (9). The inclusion criteria were: (i) to be children with OW/OB according to

175   the age- and sex-specific World Obesity Federation cut-off points (27); (ii) to be 8 to 11

176   years old; (iii) not to present neurological disorders or physical disabilities; (iv) for

177   girls, not to have started the menstruation at the beginning of the study; (v) for this

178   specific study to have blood samples and valid genotype data.The exclusion criteria

179   were: (i) to take medications that affect the function of the central nervous system; (ii)

180   to have any physical disabilities or neurological disorders that may limit exercise

181   performance; (iii) to be left-handed (due to brain differences in neuroimaging); (iv) to

182   report an attention-deficit/hyperactivity disorder. Body mass index (kg/m^2) was

183   calculated using body weight and height assessed with an electronic scale and a

184 stadiometer (Seca instruments, Germany, Ltd), while peak height velocity was

185 computed as an indicator of maturational status (28). The children's parents or guardians

186 gave their written informed consent for them to take part in the trial. The ActiveBrains

187 trial received approval from the University of Granada's ethics committee and was

188 registered on ClinicalTrials.gov (NCT02295072). This trial adhered to the CONSORT

189 (Consolidated Standards of Reporting Trials) guidelines (9). The entire pre- and post-

190 exercise data was gathered between November 21, 2014, and June 30, 2016. The

191 evaluation of all the brain health outcomes was described in detail previously (9, 26).

192 *2.2 Exercise intervention*

193 The exercise group was instructed to perform at least three of the five provided

194 supervised exercise sessions each week (in total 20 weeks). Sessions lasted 90 minutes

195 (including 60 minutes of aerobic exercise and 30 minutes of resistance training).

196 Exercise sessions were built around games and other enjoyable activities to increase

197 motivation and adherence. The adherence was recorded as the number of sessions

198 attended vs. the total number of sessions recommended in the program and expressed in

199 percentage (6 participants were removed from the per-protocol analyses for the low

200 attendance to the exercise program, i.e., <70% of the 3 recommended sessions/week)

201 (9). The participants in the control group carried on with their regular activities of daily

202 life. At the beginning of the study, information on healthy eating and physical activity

203 recommendations were given to both the control and exercise groups. The 20-week

204 exercise intervention has been described in detail in our previous study (9).

205 *2.3 Blood sampling and molecular analyses*

206 Blood samples were obtained after an overnight fast of 12 h between 8:30- 10:30

207 AM. at the hospital. Ethylenediamine tetraacetic acid-filled (EDTA) tubes were used to

208 collect blood, which was subsequently centrifuged for 10 minutes at 4°C with 1000 g of

209 force. Leukocytes were isolated, aliquoted, and kept at -80°C for SNPs analyses.

210 *2.3.1 Genotyping*

211 Six brain health-related SNPs (rs6265, rs2253206, rs2289656, rs4680, rs429358,

212 rs7412) located in 5 genes (*BDNF*, *CREB1*, *NTRK2*, *COMT* and *APOE*) were

213 genotyped. DNA genotyping was performed using TaqMan® Genotyping Master Mix

214 (Applied Biosystems, USA) for real-time polymerase chain reaction (RT-PCR). Assay

215 ID for each SNP were: rs6265 no. C__11592758_10, rs2253206 no. C___2859107_10,

216 rs2289656 no. C__15882271_20, rs4680 no. C__25746809_50, rs429358 no.

217 C___3084793_20, rs7412 no. C____904973_10. Allelic discrimination assays were

218 carried out in a QuantiStudio 6 Flex Fast Real-Time PCR System (Applied Biosystems,

219 USA). Results were read using QuantStudio™ Real-Time PCR Software, Version 1.3

220 (Life Technologies, USA). The six SNPs (rs6265, rs2253206, rs2289656, rs4680,

221 rs429358, rs7412) were considered for computing the genetic score. Each SNP was

222 coded as follows: the low-response allele homozygote was assigned 0, heterozygote

223 received 1, and homozygote for the high-response allele was assigned 2 (29). The

224 number was assigned based on information provided by the scientific literature, for

225 details see **Table S1**. The theoretical range of the score was from 0 (no beneficial

226 alleles) to 12 (two copies of the beneficial alleles) (29). In order to have a balanced

227 sample size in different subgroups, the median value of this genetic score, i.e., 6

228 beneficial alleles, was used to classify participants into two subgroups, "favorable" (≥6

229 beneficial alleles) or "unfavorable" (<6 beneficial alleles) genetic profile.

230 *2.4 Intelligence*

9

Intelligence was measured using the Spanish version of the Kaufman Brief Intelligence Test (K-BIT) (9). Experienced evaluators individually administered the K-BIT, and the different scores were calculated. K-BIT includes vocabulary (assess word knowledge using pictures that answer a question or illustrate a word) and matrices sub-tests (evaluate the kid's ability to make visual analogies spatial relationships). Crystallized intelligence score was obtained from the vocabulary sub-tests, while fluid intelligence score was estimated from the matrices sub-tests. A total intelligence score was calculated using crystallized and fluid intelligence scores.

2.5 Executive function

2.5.1 Cognitive flexibility

Cognitive flexibility was measured using the Design Fluency Test and The Trail Making Test (9, 30, 31). The Design Fluency Test is composed of three different conditions (filled dots, empty dots, and switching), each lasting 1 minute (in total 3 minutes). Children were instructed to connect dots using four straight lines to design as many novel shapes/designs as possible during the abovementioned period. The total number of correct designs in the three conditions was computed in one single variable, so that higher values indicate a better cognitive flexibility performance.

The Trail Making Test includes five different conditions, but in the current study only the condition 2 and condition 4 were used (hereinafter called Part A and Part B, respectively). Regarding Part A, children had to draw lines to connect numbers 1–25 following an ascending order and try to be as fast as possible (no more than 2.5 minutes to finish the part A). In Part B, children had to draw a line to connect the numbers (numerically) and the letters (alphabetically), switching each time from a number to a letter in consecutive order (e.g., 1–A–2–B–3–C, and so on). The maximum time to

255 complete the part B was 4 minutes. A smaller part B – part A difference (seconds)

256 indicated better cognitive flexibility. We computed a composite z-score for cognitive

257 flexibility which was calculated as the re-normalized mean of the z-scores for the

258 Design Fluency Test and Trail Making Test.

259 *2.5.2 Inhibition*

260 Inhibition was assessed using a modified version of the Stroop test (9). The

261 condition 1 and condition 3 were used in this study. Condition 1 consisted of naming

262 the color of filled rectangles. Regarding condition 3, color-words were printed in a color

263 that differs from their meaning (e.g., the word "orange" printed in blue), while the task

264 consisted of avoiding reading the word and naming the color of the word (i.e., blue is

265 the correct answer in the above example). The variable inhibition was computed as the

266 difference between the completion time (seconds) in condition 3 and condition 1 (i.e.,

267 completion time in condition 3 – condition 1) (9). For analytical purposes, the variable

268 inhibition was reverted (i.e., it was multiplied by -1), so that higher values were related

269 to better cognitive performance.

270 *2.5.3 Working memory*

271 A modified version of the Delayed NonMatch-to-Sample (DNMS) computerized

272 task was used to evaluate working memory (9). A total of 16 practice trials were

273 presented on a computer screen using E-Prime software (Psychology Software Tools,

274 Pittsburgh, PA), followed by 140 experimental trials in 5 different blocks. Each trial had

275 two phases (choice and sample) and high and low memory loads. The high working

276 memory load (100 trials) was used for the current study. Participants were required to

277 memorize a set of four different sequential stimuli (Pokémon cartoons) as part of the

278 pre-target phase. During the selection phase following the last stimulus, two distinct

279 Pokémon were presented. Participants were instructed to choose the one that had not

280 been exhibited before. Working memory was measured using response accuracy (%)

281 under high load. Higher response accuracy denoted improved working memory

282 capabilities.

283 *2.6 Academic performance*

284   Academic performance was reported using the Spanish version of the

285 Woodcock-Johnson III Tests of Achievement (9). Different academic tests (reading,

286 mathematics, oral language, written language, social sciences, and humanities) were

287 individually performed by children in a session of 100-120 min. All academic tests

288 included in the Woodcock-Johnson III battery were corrected by two independent

289 researchers and processed in the Compuscore and profile software version 3.1

290 (Riverside Publishing Company, Itasca, IL, USA). Academic performance scores for

291 reading, mathematics, writing, academic skills, academic fluency, problem solving, and

292 total academic performance were computed. For more detailed information about

293 different academic components and scores calculations, please see https://n9.cl/zsikj and

294 Ortega *et al.*, (9).

295 *2.7 Hippocampal volume*

296   Hippocampal volume was measured with the FMRIB's Integrated Registration

297 and Segmentation Tool (FIRST) in FMRIB's Software Library (FSL) version 5.0.7. The

298 tool FIRST uses a Bayesian framework from morphological brain models obtained from

299 the Center for Morphometric Analysis, Massachusetts General Hospital, Boston, MA,

300 USA. Brain volume analyses were reported in detail elsewhere (9).

301 *2.8 Statistical analyses*

302   Statistical analyses were carried out using the SPSS software (Version 22.0,

303   IBM Corp., Armonk, NY, USA). Since we were more interested in the efficacy than the

304   effectiveness of our exercise intervention (i.e., in the effects on brain health

305   outcomes when exercise was actually done, that was to attend to at least 70% of

306   program'sd sessions), the main findings were derived from the per-protocol analyses. In

307   addition, we report the results using the intention-to-treat principle which include all

308   participants initially randomized in the analysis. For this purpose, multiple imputation

309   of missing values was applied using the predictive mean matching approach, for more

310   details see our previous publication (9). Overall, dropouts and non-dropouts did not

311   differ in the primary study outcomes (as described in the main article of the trial (9)).

312   Two-way analysis of covariance (ANCOVA) was performed to explore the

313   interaction between the genetic score and the effects of the exercise intervention on

314   brain health indicators. The model included: factor 1, genetic score (0 = unfavorable

315   genetic profile; 1 = favorable genetic profile); factor 2, group (0 = control; 1 =

316   intervention); outcome (post-intervention values); covariable (baseline values of the

317   outcome studied). P-value <0.1 was considered indicative of a potential gene x exercise

318   interaction, which was further explored in separate analyses by genetic sub-groups. This

319   study was powered to test the effects of the intervention (control vs. exercise) in the

320   whole sample; therefore, these gene-group interaction analyses were exploratory, and

321   we considered that there was certain evidence of interaction when P < 0.1.

322   Subsequently, one-way ANCOVA (factor: group [0 = control; 1 = intervention];

323   outcome [post-intervention values]; covariable [baseline values of the outcome studied])

324   was performed to report mean differences of post-intervention values (adjusted by

325   baseline values) of brain outcomes between exercise and control groups (9), separately

326   for each specific genetic profile (i.e., "favorable" and "unfavorable" genetic profile).

327   For intervention effects, we kept the standard 5% alpha error (i.e. P-value <0.05) for

328   consistency with the reporting of the intervention effects in this trial (9), yet we are

329   aware that by splitting the sample into two genetic groups the power was markedly

330   reduced and only relatively large effect sizes will be flagged as significant. In addition,

331   due to the high number of outcomes, we performed multiple hypothesis testing

332   corrections, i.e. false discovery rate [FDR] Benajmini-Hochberg procedure, in line with

333   the primary paper (9). The standardized effects of the exercise intervention on brain

334   health outcomes were presented using Z-scores of change (9). It shows how many

335   standard deviations (SDs) of the postexercise program values changed from the baseline

336   mean and SD values. This effect size can be interpreted as follows: a small effect size

337   (0.2 SDs), a medium effect size (0.5 SDs), and a large effect size (0.8 SDs) (9).

338   In addition, we explored the individual changes in brain health outcomes that

339   showed an interaction P<0.1. We reported the % of children that showed a meaningful

340   change (>0.2 Cohen's d) for brain health outcomes with statistically significant

341   differences of % between subgroups (chi-square test). We performed exploratory sex

342   interaction analyses.

343

3. Results

345   The baseline characteristics of the participants stratified by genetic profile (i.e.,

346   "favorable" or "unfavorable" genetic profile) and group (i.e., exercise or control) are

347   presented in **Table 1**. The genotype frequencies for each SNP were in Hardy-Weinberg

348   equilibrium (**Table 2**). Per-protocol analyses showed an interaction of the genetic score

349   with cognitive flexibility, as measured by Trail Making Test and the composite score,

350   working memory, academic skills, reading and writing (all P<0.1) (**Table 3**). Exercise

351 only increased cognitive flexibility and academic skills in children presenting a

352 "favorable" genetic profile (mean z score, 0.41-0.67 [95% CI 0.11 to 1.18]; **Figure 1**

353 **[Panel A]** and **Table 3**), yet not in the rest of brain health outcomes. Among children

354 presenting an "unfavorable" genetic profile, exercise only improved working memory

355 and writing (mean z score, 0.47 [95% CI 0.04 to 0.90] and mean z score, 0.55 [95% CI

356 0.04 to 1.05]; **Figure 1 [Panel A]** and **Table 3**). An interaction effect was reported for

357 reading (P<0.1; **Table 3**), but the effect of exercise intervention was not statistically

358 significant in both subgroups computed using the genetic score (P>0.05; **Table 3**).

359 Reading showed a trend to improve in the "favorable" genetic profile but not in the

360 "unfavorable" group (**Figure 1 [Panel A]** and **Table 3**). All the results described before

361 were consistent when using intention-to-treat instead of per-protocol analyses (**Figure 1**

362 **[Panel B]**; **Table S2**, except for the no interaction of the genetic score with writing and

363 reading (P>0.1). The signficant effects on cognitive flexibility and academic skills

364 remained consistent after correction for multiple comparisons (FDR<0.05), but the

365 effects on working memory and writing became non-significant (FDR>0.05).

366   Regarding the individual changes in brain health outcomes, only cognitive

367 flexibility showed statistically significant differences in the % of participants that

368 reported a meaningful change among subgroups (**Figure 2**). An individual response

369 analysis among the children with a "favorable" genetic profile showed that 62% of them

370 had a meaningful (≥0.2 Cohen *d*) increase in cognitive flexibility in the exercise group

371 compared with only 25% in the control group (**Figure 2**). This result was consistent in

372 per-protocol and intention-to-treat analyses (P=0.01, P=0.03 respectively) (**Figure 2**).

373 We did not find differences at the individual response level for the other variables that

374 showed a gene*exercise interaction P-value <0.1 (**Figures S1-4**). For exploratory

375 purposes, we tested whether the most consistent and robust gene*exercise interaction

376   observed in cognitive flexibility was also consistent in boys and girls, so we analyzed

377   the sex*gene*exercise interaction on this primary outcome and observed no evidence of

378   sex having a moderating effect (sex interaction P=0.31, P=0.15 for per-protocol and

379   intention-to-treat analyses, respectively). We also explored the sex interactions for the

380   rest of the outcomes studied, finding no interactions, except for a few academic

381   outcomes. However, separate analyses by sex would not be meaningful since the sample

382   would be stratified too much, i.e., in 8 groups (2 sexes * 2 genetic groups * 2

383   intervention groups) having many of them less than 10 participants per group.

384

385   **4. Discussion**

386   This study showed, for the first time, the role of a genetic background, namely a

387   combination score of polymorphisms in brain health-related candidate genes, on the

388   response to a 20-week exercise intervention in a broad set of brain health indicators in

389   children with OW/OB. The genetic score composed of 6 candidate (selected based on

390   evidence) SNPs located in genes that encode proteins with important role in the brain

391   health [*BDNF, CREB1, NTRK2, COMT,* and *APOE*] modulated the exercise-induced

392   response on cognitive flexibility, working memory, and academic performance. The

393   significant effect persisted after correction for multiple comparisons for cognitive

394   flexibility and academic skills, but not for working memory or writing. Our findings

395   suggest that the differential response to exercise according to the genetic predisposition

396   was especially consistent and robust for cognitive flexibility in the analyses conducted

397   both at group and individual level, as well as in per-protocol and intention-to-treat

398   analyses. The beneficial effect of the genetic profile on the response to exercise in

399   cognitive flexibility seems to be also consistent in boys and girls, as evidenced by the

400   non-interaction by sex observed.

401   The ActiveBrains RCT reported an improvement on cognitive flexibility after a

402 20-week exercise intervention in children with OW/OB (9). The current study adds that

403 the abovementioned improvements were observed only in children with a "favorable"

404 genetic profile. Interestingly, studies performed in animal models demonstrated that

405 cognitive flexibility impairment was related to decreased BDNF production in the

406 frontal cortex (32). In this regard, exercise training increases BDNF protein levels in the

407 plasma of children and adults, benefiting brain health (18, 20). Also, a single bout of

408 high-intensity exercise improved cognitive flexibility in healthy young adults,

409 specifically the performance on the Trail Making test, in parallel to an improvement of

410 BDNF protein levels in circulation(33). Also, the expression levels of other proteins

411 such as CREB1 and NTRK2 (BDNF receptor) influenced by SNPs included in our

412 genetic score can regulate BDNF protein expression and function (16, 21), affecting

413 cognitive flexibility. Furthermore, academic skills that are regulated by BDNF (34)

414 improved only after the exercise intervention in children with a "favorable" genetic

415 profile. We can hypothesize that the genetic score can modulate the expression of the

416 abovementioned proteins in response to exercise, contributing to improve cognitive

417 flexibility. Importantly, crystalized intelligence was the brain outcome with the largest

418 effect size in the ActiveBrains RCT (9). In the current study, we did not observe an

419 interaction between the genetic score and crystalized intelligence; however, the effect

420 size was larger in the "favorable" compared to the "unfavorable" genetic subgroup

421 (**Table 3**).

422   The ActiveBrains RCT has shown that working memory and writing did not

423 change after a 20-week exercise intervention (9). Interestingly, in the current study, we

424 observed that exercise improved working memory and writing in children classified as

425 having "unfavorable" genetic profiles. However, no significant effects were observed

426 after correction for multiple comparisons, nor significant differences were observed in

427 the individual response to exercise in both genetic groups for these two outcomes,

428 which suggest these findings are not consistent or robust, and not much attention should

429 be paid on them. Importantly, another aspect that elucidates the complex relationship

430 between genetic variants, exercise, and cognition is that a given allele for the same

431 candidate SNP could be beneficial for some aspects of cognition and hampering for

432 others. As an illustration, the *COMT* gene encodes an enzyme (Catechol-O-

433 methyltransferase) that regulates dopamine levels and time of action in the prefrontal

434 cortex. The change of Met for Val allele [COMT rs4680] results in a three- to four-fold

435 decreased activity of COMT activity that contribute to an extended dopamine action in

436 the prefrontal cortex (24, 35). Interestingly, high levels of dopamine in prefrontal regions

437 (observed in carriers of the Met allele in *COMT* rs4680) may be beneficial for working

438 memory but a disadvantage for cognitive flexibility (24, 36, 37).

439   Our study has some limitations that should be acknowledged. Complex

440 phenotypes such as brain health indicators are influenced by several genes with small

441 effect sizes, which in concert may exert an effect (12, 15). Thus, our target approach

442 would need to be tested by whole genome-wide analyses and in larger RCTs.

443 Importantly, we carefully selected six SNPs based on the scientific literature, although

444 most of the studies relating these genetic variants to brain health indicators were

445 performed in adults, older adults and patients with neurological diseases. Therefore,

446 there was almost inexistent evidence on relevant genes derived from studies in pediatric

447 populations. Future RCTs should explore the genome unbiasedly by performing whole

448 genome-wide analyses integrated with proteomics data in larger cohorts of children.

449 Furthermore, studies with larger sample size and power should confirm or contrast our

450 findings.

5. Conclusion

Our findings revealed that the studied genetic score using brain health-related polymorphisms (selected based on previous scientific literature) influenced the response to exercise in cognitive flexibility and academic skills. Notably, the impact of the genetic score was more consistent and pronounced on cognitive flexibility compared to other outcomes, showing that children with a more favorable genetic profile improved more their cognitive flexibility as a result of the exercise intervention that their peers with a "less favorable" genetic profile. To further enhance our understanding of the genetic factors influencing brain health in response to exercise training, future randomized controlled trials should employ whole genome analyses without bias, yet that requires very large sample size and power, or alternatively aggregation of data from different trials.

SUPPLEMENTAL MATERIAL: The supplementary tables and figures were deposit to a public access data repository

(figshare, https://doi.org/10.6084/m9.figshare.23884197)

Supplemental Table S1
Supplemental Table S2
Supplemental Figure S1
Supplemental Figure S2
Supplemental Figure S3
Supplemental Figure S4

DATA AND RESOURCE AVAILABILITY

We did not obtain children's parents consent to widely share the data nor was it included in the IRB protocol.

DISCLOSURES

The authors declare that they have no competing interests.

AUTHORS' CONTRIBUTIONS

APF, IEC, JMG, LVT, FJO, SA, FBO participated in the manuscript design/conception; APF, IEC, JMG, LVT, JGC, SA, FBO data acquisition, APF and FBO data analysis, APF, JMG and JGC figure generation, APF first drafting of the manuscript; APF, IEC, JMG, LVT, FJOP, JGC, SRA, IL, JRR, SA, FBO critically revised the manuscript for important intellectual content. All authors have read and approved the final version of the manuscript, and agree with the order of presentation of the authors.

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FIGURE LEGENDS

Figure 1. Per-protocol (A) and intention-to-treat (B) effects of the ActiveBrains exercise intervention on the intelligence, executive function, academic performance, and brain structure by genetic profile. To simplify the interpretation of the results, the Stroop Color-Word Test and Trail Making Test were inverted, i.e., a higher value indicates a better performance. * indicates a significant gene x exercise interaction (p-value < 0.1). Number of participants with valid data for each variable pre-and post-intervention in the per-protocol analyses: Intelligence outcomes, Cognitive flexibility 1, Inhibition (N=84; genetic "favorable" [21exercise and 28 control] and genetic "unfavorable" [22 exercise and 13 control]), Cognitive flexibility 2, Cognitive flexibility composite z-score (N=79; genetic "favorable" [21 exercise and 24 control] and genetic "unfavorable" [22 exercise and 12 control]), working memory (N=81; genetic "favorable" [19 exercise and 27 control] and genetic "unfavorable" [22 exercise and 13 control]), Executive function composite z-score (N=77; genetic "favorable" [19 exercise and 24 control] and genetic "unfavorable" [22 exercise and 12 control]), academic performance outcomes (N=83, genetic "favorable" [21 exercise and 27 control] and genetic "unfavorable" [22 exercise and 13 control]), hippocampal volume (N=77; genetic "favorable" [20 exercise and 23 control] and genetic "unfavorable" [21 exercise and 13 control]).

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705 **Figure 2**. Individual change distribution in cognitive flexibility (Panels A and B per-

706 protocol analyses and panels C and D intention-to-treat analyses) for both control and

707 exercise groups, and by genetic profile. Dashed lines indicate a meaningful increase

708 regarding baseline levels. P value from the chi-squared test. Number of participants

709 with valid data for cognitive flexibility composite z-score pre-and post-intervention in

710 the per-protocol analyses (N=79; genetic "favorable" [21 exercise and 24 control] and

711 genetic "unfavorable" [22 exercise and 12 control]). The standardized score of change

712 indicates how many standard deviations have the post-intervention values changed with

713 respect to the baseline mean and standard deviation. E.g., a 0.70 Z-score means that the

714 value at post-intervention is 0.70 standard deviations higher than the mean value at

715 baseline, indicating a positive change, with negative values indicating the opposite.

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Table 1. Descriptive baseline characteristics of the *ActiveBrains* participants by genetic profile and type of intervention.

		All		Favorable genetic profile		Unfavorable genetic profile		Control group		Exercise group
	N	Mean ± SD	N	Mean ± SD	N	Mean ± SD	N	Mean ± SD	N	Mean ± SD
Age (years)	101	10.03 ± 1.51	59	9.91 ± 1.24	42	10.19 ± 1.00	50	10.09 ± 1.16	51	9.96 ± 1.16
Sex										
Girls (n %)	41	59%	26	44%	15	36%	23	46%	18	35%
Boys (n %)	60	41%	33	56%	27	64%	27	54%	33	65%
Weight (kg)	101	55.93 ± 10.99	59	55.61 ± 10.72	42	55.38 ± 11.47	50	55.96 ± 9.42	51	55.90 ± 12.43
Height (cm)	101	143.91 ± 8.55	59	143.37 ± 8.55	42	144.68 ± 8.59	50	145.22 ± 7.99	51	142.63 ± 8.94
Body mass index (kg/m^2)	101	26.79 ± 3.51	59	26.83 ± 3.24	42	26.74 ± 3.89	50	26.42 ± 2.96	51	27.16 ± 3.97
Peak height velocity (years)	101	–2.28 ± 0.99	59	–2.32 ± 1.09	42	–2.23 ± 0.85	50	–2.13 ± 1.07	51	–2.43 ± 0.91
Wave of participation (%)										
First (n %)	16	15%	7	12%	9	21%	7	14%	9	18%
Second (n %)	45	45%	32	54%	13	31%	23	46%	22	43%
Third (n %)	40	40%	20	34%	20	48%	20	40%	20	39%
Intelligence										
Crystallized intelligence (typical punctuation)	101	103.15 ± 13.26	59	103.64 ± 14.07	42	102.45 ± 12.15	50	102.58 ± 12.00	51	103.71 ± 14.48
Fluid intelligence (typical punctuation)	101	97.87 ± 13.17	59	97.08 ± 13.57	42	98.98 ± 12.65	50	98.84 ± 12.23	51	96.92 ± 14.08

Total intelligence (typical punctuation)	101	98.22 ± 12.65	59	98.10 ± 13.56	42	98.38 ± 11.39	50	98.42 ± 11.85	51	98.02 ± 11.49
Executive function										
Cognitive flexibility 1 (total correct designs)	101	19.75 ± 6.47	59	19.53 ± 6.03	42	20.07 ± 7.10	50	20.08 ± 6.95	51	19.43 ± 6.01
Cognitive flexibility 2 (sec)	101	90.99 ± 43.36	59	91.19 ± 39.46	42	90.72 ± 48.83	50	94.78 ± 44.91	51	87.27 ± 41.91
Cognitive flexibility composite z-score	101	0.01 ± 1.00	59	-0.04 ± 0.93	42	0.02 ± 1.11	50	-0.03 ± 1.06	51	0.00 ± 0.96
Inhibition (sec)	101	40.81 ± 17.41	59	43.46 ± 19.51	42	37.10 ± 13.27	50	41.16 ± 19.60	51	40.48 ± 15.15
Working memory (% response accuracy)	101	65.43 ± 16.58	59	63.35 ± 16.22	42	69.75 ± 16.30	50	62.50 ± 18.09	51	68.30 ± 14.56
Executive function composite z-score	101	0.01 ± 1.00	59	-0.14 ± 0.97	42	0.18 ± 1.101	50	−0.09 ± 1.13	51	0.07 ± 0.85
Academic performance (standard score)										
Academic skills	101	118.65 ± 15.31	59	115.07 ± 14.17	42	123.67 ± 15.59	50	116.86 ± 14.70	51	120.39 ± 15.83
Academic fluency	101	103.38 ± 11.42	59	102.41 ± 11.42	42	104.74 ± 11.41	50	102.52 ± 12.81	51	104.22 ± 9.92
Problem solving	101	99.56 ± 9.37	59	99.10 ± 10.06	42	100.21 ± 8.38	50	97.29 ± 9.04	51	101.78 ± 9.23
Reading	101	107.94 ± 12.58	59	106.14 ± 11.44	42	110.46 ± 13.77	50	105.71 ± 11.69	51	110.12 ± 13.14
Mathematics	101	101.80 ± 10.86	59	101.14 ± 11.32	42	102.73 ± 10.25	50	99.49 ± 10.51	51	104.06 ± 10.83
Writing	101	113.89 ± 12.31	59	110.91 ± 12.56	42	118.09 ± 10.75	50	113.45 ± 13.51	51	114.33 ± 11.13
Total academic performance	101	109.16 ± 11.66	59	107.17 ± 11.32	42	111.96 ± 11.70	50	107.19 ± 11.71	51	111.10 ± 11.40
Hippocampal volume (mm^3)	101	6997.62 ± 619.27	59	7015.80 ± 643.85	42	6972.07 ± 589.73	50	6967.46 ± 661.74	51	7027.18 ± 579.70

Values are expressed as means ± standard deviations (SD), unless otherwise indicated. BDNF = Brain-derived neurotrophic factor. Intelligence outcomes (i.e., Crystallized, Fluid, and Total Intelligence) were measured by the Kaufman Brief Intelligence Test. Cognitive flexibility 1 was measured by the Design Fluency Test and expressed as number of total correct designs of the three conditions. Cognitive flexibility 2 was measured by the Trail Making Test and expressed as the total completion time (sec) of Part A subtracted from the total completion time (sec) of Part B. A smaller B – A difference score (sec) indicated better cognitive flexibility. Cognitive flexibility composite z-score was calculated as the re-normalized mean of the z-scores for Cognitive flexibility 1 and Cognitive flexibility 2. Inhibition was measured by the Stroop Color-Word Test. The inhibition score was obtained by subtracting condition 3 completion time – condition 1 completion time (sec). The lower the difference between tests' times, the better the performance was considered. Working memory was measured by the Delayed Non-Match-to sample task. Executive function composite z-score was calculated as the re-normalized mean of the z-scores for Cognitive flexibility, Inhibition, and Working memory. Academic performance was measured by the Spanish version of the Woodcock Johnson III Test of Achievement. Academic skills are the sum of components based on basic skills such as reading decoding, mathematics calculation, and spelling. Academic fluency is the sum of the components based on reading, calculation, and writing fluency. Problem solving is the sum of the components based on solving academic problems in reading, mathematics, and writing. Total academic performance is the overall measure of the academic performance based on reading, mathematics, and writing.

Table 2. Genotype and allele frequencies in genes analyzed in 101 children with overweight/obesity.

SNP	Gene	Genotype frecuencies	Allele frecuencies	X^2 Hardy-Weinberg equilibrium	Computation of the genotype to calculate the genetic score (for more details see **Table S1**)
rs6265	*BDNF*	CC (62; 61%) CT (32; 32%) TT (7; 7%)	p (C allele; 0.77) q (T allele; 0.23)	0.99	0 = TT; 1 = CT; 2 = CC
rs2253206	*CREB1*	GG (36; 36%) AG (53; 52%) AA (12; 12%)	p (G allele; 0.62) q (A allele; 0.38)	1.27	0 = GG; 1 = AG; 2 = AA
rs2289656	*NTRK2*	GG (69; 36%) AG (26; 52%) AA (6; 12%)	p (G allele; 0.81) q (A allele; 0.19)	2.49	0 = GG; 1 = AG; 2 = AA
rs4680	*COMT*	GG (31; 31%) AG (49; 48%) AA (21; 21%)	p (G allele; 0.55) q (A allele; 0.45)	0.04	0 = AA; 1 = AG; 2 = GG
rs429358	*APOE*	CC (0; 0%) CT (15; 15%) TT (86; 85%)	p (T allele; 0.93) q (C allele; 0.07)	0.65	0 = CC; 1 = CT; 2 = TT
rs7412	*APOE*	TT (1; 1%) CT (14; 14%) CC (86; 85%)	p (C allele; 0.93) q (T allele; 0.07)	0.25	0 = CC; 1 = CT; 2 = TT

BDNF, Brain-derived neurotrophic factor; CREB1, cAMP responsive element binding protein 1; NTRK2, tyrosine kinase receptor 2; COMT, Catechol-Omethyltransferase; APOE, apolipoprotein E; SNP, Single nucleotide polymorphism. The term p represents the frequency of the homozygous dominant genotype, while the term q indicates the frequency of the homozygous recessive genotype. X^2 > 0.05 shows that genotype distributions in children with overweight/obesity were in Hardy-Weinberg equilibrium

Table 3 Effects of the ActiveBrains exercise intervention (per-protocol analyses) on z-score post-intervention outcomes (Z-score of change from baseline) by genetic favourable/unfavourable profiles.

	Favorable genetic profile		Unfavorable genetic profile		Intervention vs. Control differences by genetic profile		Gene x exercise interaction (p-value)
	Intervention group	Control group	Intervention group	Control group	Genetic favourable	Genetic unfavourable	
Intelligence							
Crystallized intelligence	0.74 (0.46, 1.02)	-0.10 (-0.34, 0.15)	0.42 (0.21, 0.62)	-0.15 (-0.42, 0.11)	**0.84 (0.47, 1.21)**	**0.57 (0.24, 0.90)**	0.31
Fluid intelligence	0.40 (0.13, 0.79)	0.12 (-0.22, 0.45)	0.39 (0.01, 0.78)	0.24 (-0.27, 0.75)	0.28 (-0.23, 0.80)	0.15 (-0.50, 0.81)	0.73
Total intelligence	0.71 (0.40, 1.02)	0.01 (-0.27, 0.27)	0.57 (0.32, 0.82)	0.01 (-0.31, 0.34)	**0.70 (0.29, 1.12)**	**0.56 (0.15, 0.97)**	0.53
Executive function							
Cognitive flexibility 1	0.63 (0.38, 0.89)	0.09 (-0.13, 0.32)	0.70 (0.31, 1.09)	0.39 (-0.12, 0.91)	**0.54 (0.19, 0.88)**	0.31 (-0.35, 0.96)	0.44
Cognitive flexibility 2	0.55 (0.18, 0.92)	-0.12 (-0.47, 0.22)	0.37 (0.03, 0.70)	0.61 (0.16, 1.07)	**0.67 (0.16, 1.18)**	-0.24 (-0.82, 0.32)	0.02*
Cognitive flexibility composite z-score	0.30 (0.04, 0.55)	-0.37 (-0.61, -0.14)	0.24 (-0.09, 0.56)	0.18 (-0.26, 0.62)	**0.67 (0.32, 1.02)**	0.06 (-0.49, 0.61)	0.05*

Inhibition	0.38 (0.01, 0.76)	0.27 (-0.05, 0.59)	0.64 (0.42, 0.87)	0.89 (0.60, 1.19)	0.11 (-0.38, 0.61)	-0.25 (-0.62, 0.13)	0.28
Working memory	-0.31 (-0.66, 0.02)	0.02 (-0.27, 0.30)	0.35 (0.09, 0.62)	-0.12 (-0.46, 0.23)	-0.33 (-0.78, 0.11)	**0.47 (0.04, 0.90)**	0.01*
Executive function composite z-score	0.01 (-0.34, 0.36)	-0.33 (-0.64, -0.02)	0.31 (0.08, 0.53)	0.24 (-0.06, 0.54)	0.34 (-0.13, 0.80)	0.07 (-0.30, 0.45)	0.40
Academic performance							
Academic skills	0.26 (0.04, 0.48)	-0.15 (-0.34, 0.05)	0.37 (0.14, 0.61)	0.46 (0.16, 0.77)	**0.41 (0.11, 0.71)**	-0.09 (-0.48, 0.29)	0.05*
Academic fluency	0.24 (-0.02, 0.49)	0.22 (-0.01, 0.44)	0.26 (0.01, 0.51)	0.12 (-0.21, 0.46)	0.02 (-0.33, 0.37)	0.14 (-0.29, 0.56)	0.58
Problem solving	0.29 (0.02, 0.56)	-0.06 (-0.29, 0.18)	0.48 (0.24, 0.72)	0.09 (-0.22, 0.41)	0.35 (-0.02, 0.71)	0.39 (-0.01, 0.79)	0.65
Reading	0.14 (-0.08, 0.35)	-0.11 (-0.30, 0.08)	0.32 (0.12, 0.53)	0.59 (0.33, 0.86)	0.25 (-0.04, 0.54)	-0.27 (-0.61, 0.07)	0.04*
Mathematics	0.24 (-0.04, 0.51)	-0.14 (-0.38, 0.10)	0.47 (0.17, 0.78)	0.33 (-0.07, 0.73)	0.38 (0.01, 0.75)	0.14 (-0.36, 0.54)	0.41
Writing	0.36 (0.16, 0.57)	0.27 (0.09, 0.45)	0.42 (0.11, 0.72)	-0.13 (-0.53, 0.27)	0.09 (-0.18, 0.37)	**0.55 (0.04, 1.05)**	0.09*
Total academic performance	0.24 (0.05, 0.43)	0.01 (-0.17, 0.17)	0.46 (0.26, 0.65)	0.37 (0.12, 0.62)	0.23 (-0.03, 0.50)	0.09 (-0.23, 0.40)	0.62

Brain structure

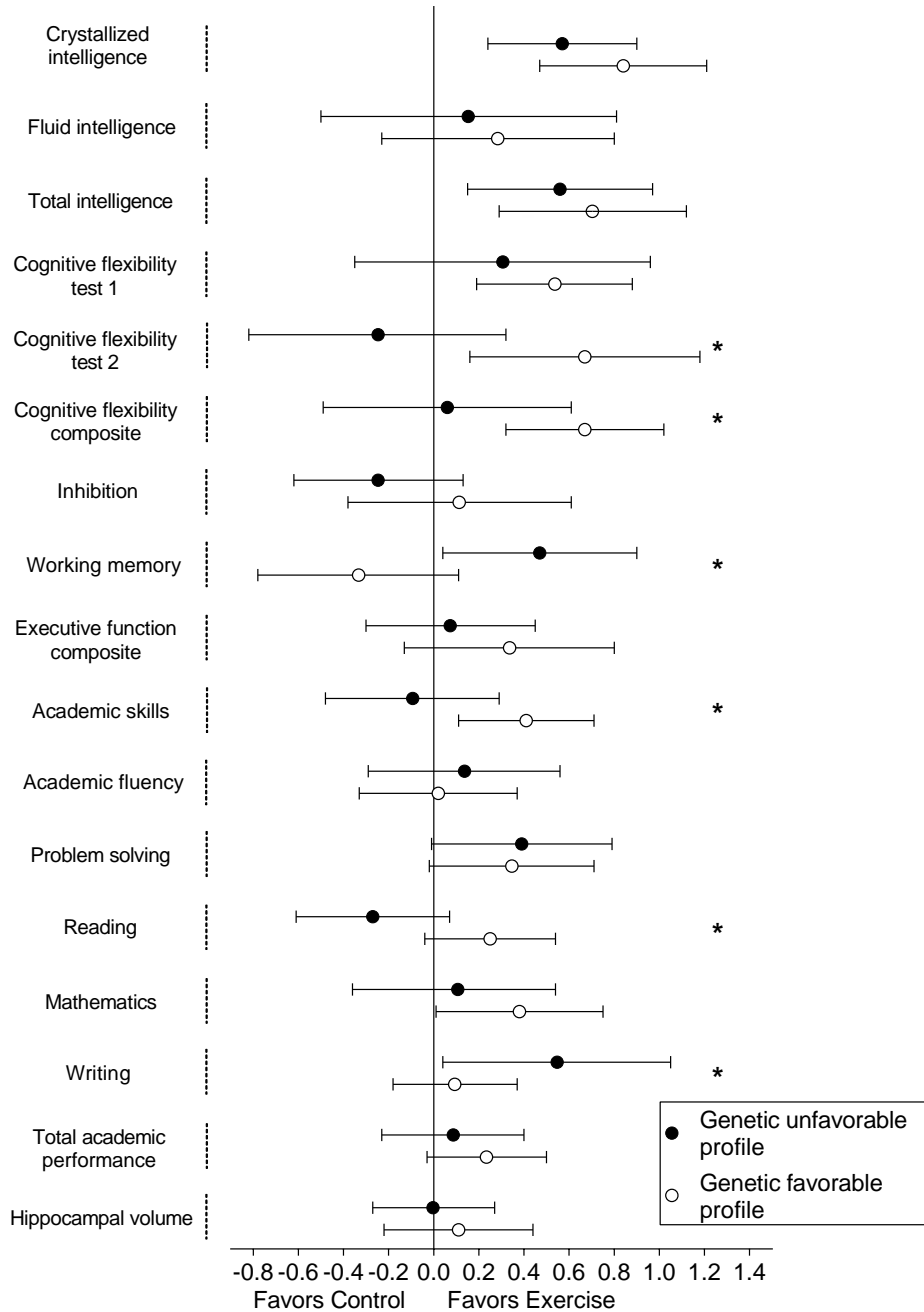|							
Hippocampal volume	0.37 (0.13, 0.61)	0.26 (0.04, 0.49)	0.11 (-0.05, 0.29)	0.12 (-0.09, 0.33)	0.11 (-0.22, 0.44)	-0.01 (-0.27, 0.27)	0.62

Z-score values indicate how many standard deviations have the post-intervention values changed with respect to the baseline mean and standard deviation. E.g., a 0.70 Z-score means that the mean value at post-intervention is 0.70 standard deviations higher than the mean value at baseline, indicating a positive change, with negative values indicating the opposite. Values are expressed as mean (95% CI). Analyses were adjusted for baseline values. Gene *x* exercise interaction p-value indicates the interaction between genetic predisposition profile and the effects induced by exercise intervention (ANCOVA analyses, factor 1: genotype profile [0 unfavorable genetic profile; 1 favorable genetic profile]; factor 2: group [0 control; 1 intervention]; outcome: post-intervention values; covariable: baseline outcomes). An asterisk (*) indicates a significant gene *x* exercise interaction (p-value < 0.1). Bold numbers indicate P < 0.05 for the difference between intervention and control group for a specific genetic profile.
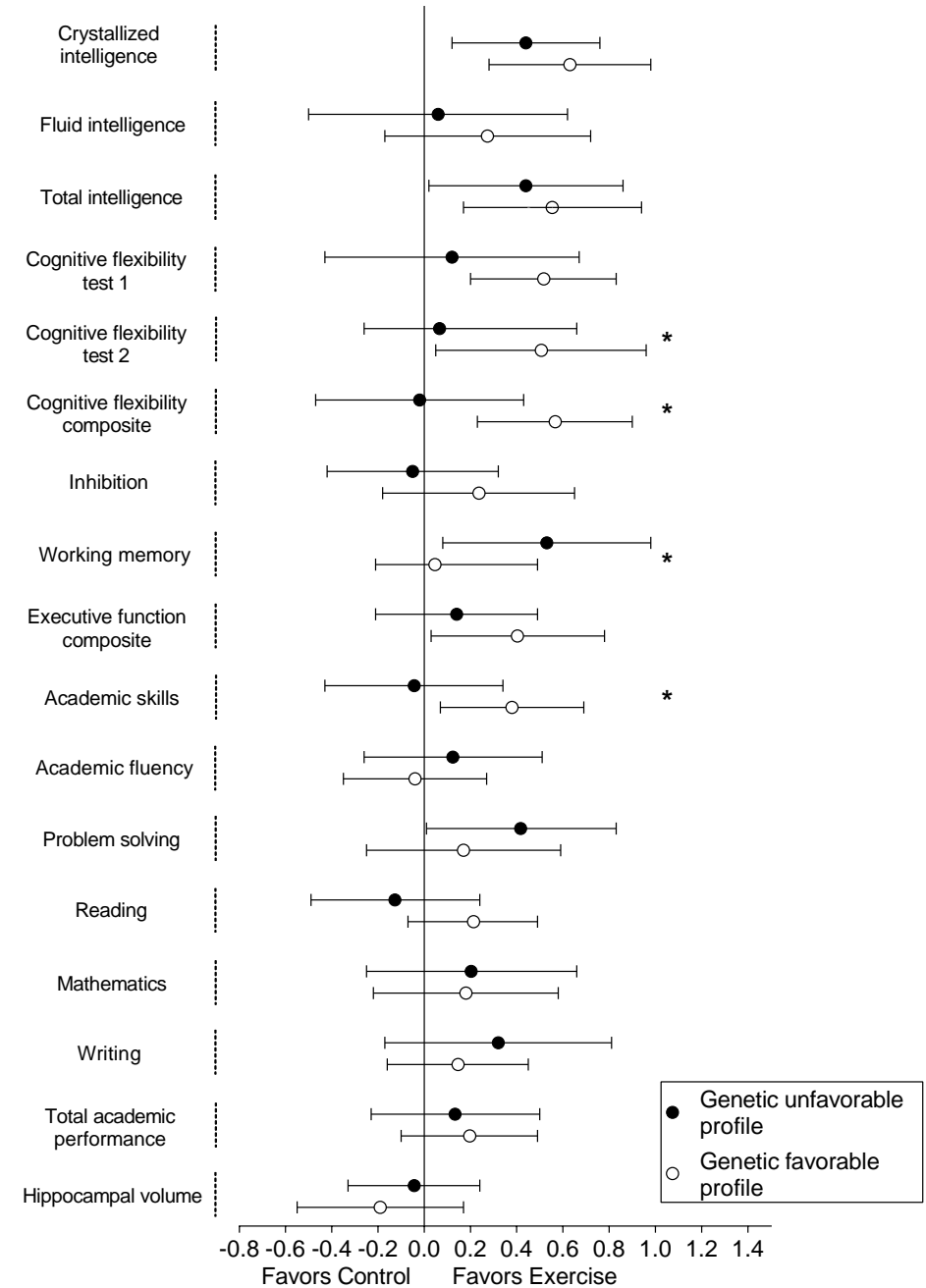
Intelligence outcomes (i.e., Crystallized, Fluid, and Total Intelligence) were measured by the Kaufman Brief Intelligence Test. Cognitive flexibility 1 was measured by the Design Fluency Test and expressed as number of total correct designs of the three conditions. Cognitive flexibility 2 was measured by the Trail Making Test and expressed as the total completion time (sec) of Part A subtracted from the total completion time (sec) of Part B. A smaller B – A difference score (sec) indicated better cognitive flexibility (to simplify the interpretation of the results, the Trail Making Test was inverted, i.e., a higher value indicates a better performance). Cognitive flexibility composite z-score was calculated as the re-normalized mean of the z-scores for Cognitive flexibility 1 and Cognitive flexibility 2. Inhibition was measured by the Stroop Color-Word Test. The inhibition score was obtained by subtracting condition 3 completion time – condition 1 completion time (sec). The lower the difference between tests' times, the better the performance was considered (to simplify the interpretation of the results, the Stroop Color-Word Test was inverted, i.e., a higher value indicates a better performance). Working memory was measured by the Delayed Non-Match-to sample task. Executive function composite z-score was calculated as the re-normalized mean of the z-scores for Cognitive flexibility, Inhibition, and Working memory. Academic performance was measured by the Spanish version of the Woodcock Johnson III Test of Achievement. Academic skills are the sum of components based on basic skills such as reading decoding, mathematics calculation, and spelling. Academic fluency is the sum of the components based on reading, calculation, and writing fluency. Problem solving is the sum of the components based on solving academic problems in reading, mathematics, and writing. Total academic performance is the overall measure of the academic performance based on reading, mathematics, and writing. Number of participants with valid data for each variable pre-and post-intervention: Intelligence outcomes, Cognitive flexibility 1, Inhibition (N=84; genetic "favorable" [21exercise and 28 control] and genetic "unfavorable" [22 exercise and 13 control]), Cognitive flexibility 2, Cognitive flexibility composite z-score (N=79; genetic

"favorable" [21 exercise and 24 control] and genetic "unfavorable" [22 exercise and 12 control]), working memory (N=81; genetic "favorable" [19 exercise and 27 control] and genetic "unfavorable" [22 exercise and 13 control]), Executive function composite z-score (N=77; genetic "favorable" [19 exercise and 24 control] and genetic "unfavorable" [22 exercise and 12 control]), academic performance outcomes (N=83, genetic "favorable" [21 exercise and 27 control] and genetic "unfavorable" [22 exercise and 13 control]), hippocampal volume (N=77; genetic "favorable" [20 exercise and 23 control] and genetic "unfavorable" [21 exercise and 13 control]).

A) Per-protocol analyses

B) Intention-to-treat analyses

Crystallized intelligence

Fluid intelligence

Total intelligence

Cognitive flexibility test 1

Cognitive flexibility test 2

Cognitive flexibility composite

Inhibition

Working memory

Executive function composite

Academic skills

Academic fluency

Problem solving

Reading

Mathematics

Writing

Total academic performance

Hippocampal volume

-0.8 -0.6 -0.4 -0.2 0.0 0.2 0.4 0.6 0.8 1.0 1.2 1.4

Favors Control   Favors Exercise

● Genetic unfavorable profile

○ Genetic favorable profile

Individual responses on cognitive flexibility

Favorable genetic profile

A) Per-protocol

$P = 0.01$

25%

62%

C) Intention-to-treat

$P = 0.03$

25%

56%

Unfavorable genetic profile
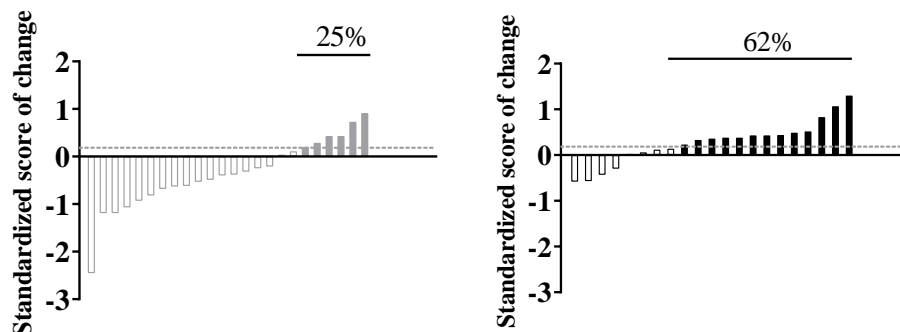
B) Per-protocol

$P = 0.72$

50%

59%
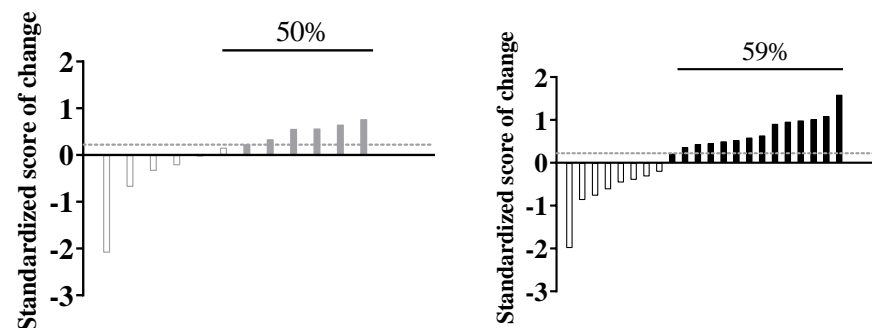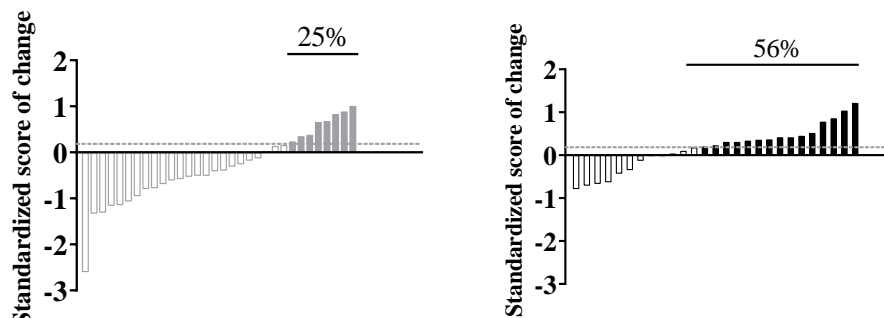
D) Intention-to-treat

$P = 0.76$

50%

58%
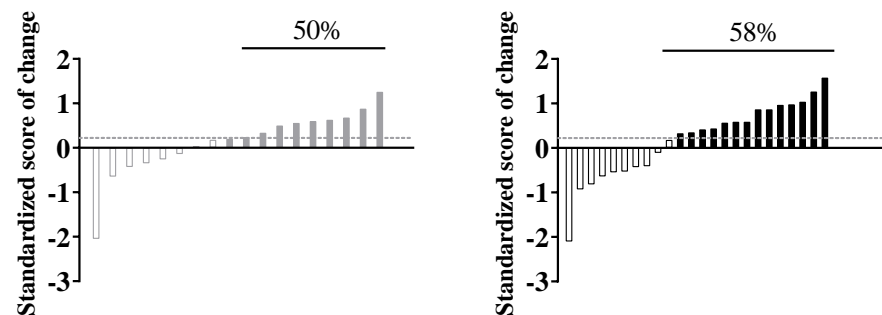
Control

Exercise

Standardized score of change
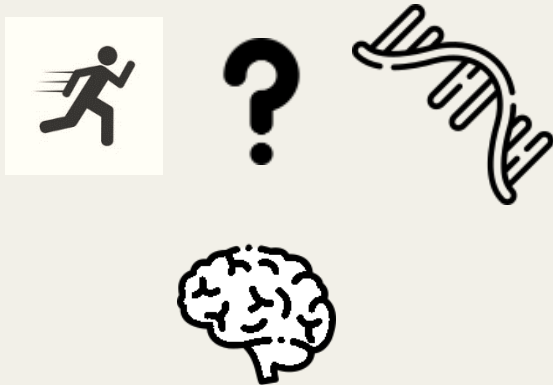
Meaningful increase (Cohen's $d \geq 0.2$)

Non-meaningful change (Cohen's $d < 0.2$)

Gene-exercise interaction on brain health in children with overweight/obesity: The ActiveBrains randomized controlled trial
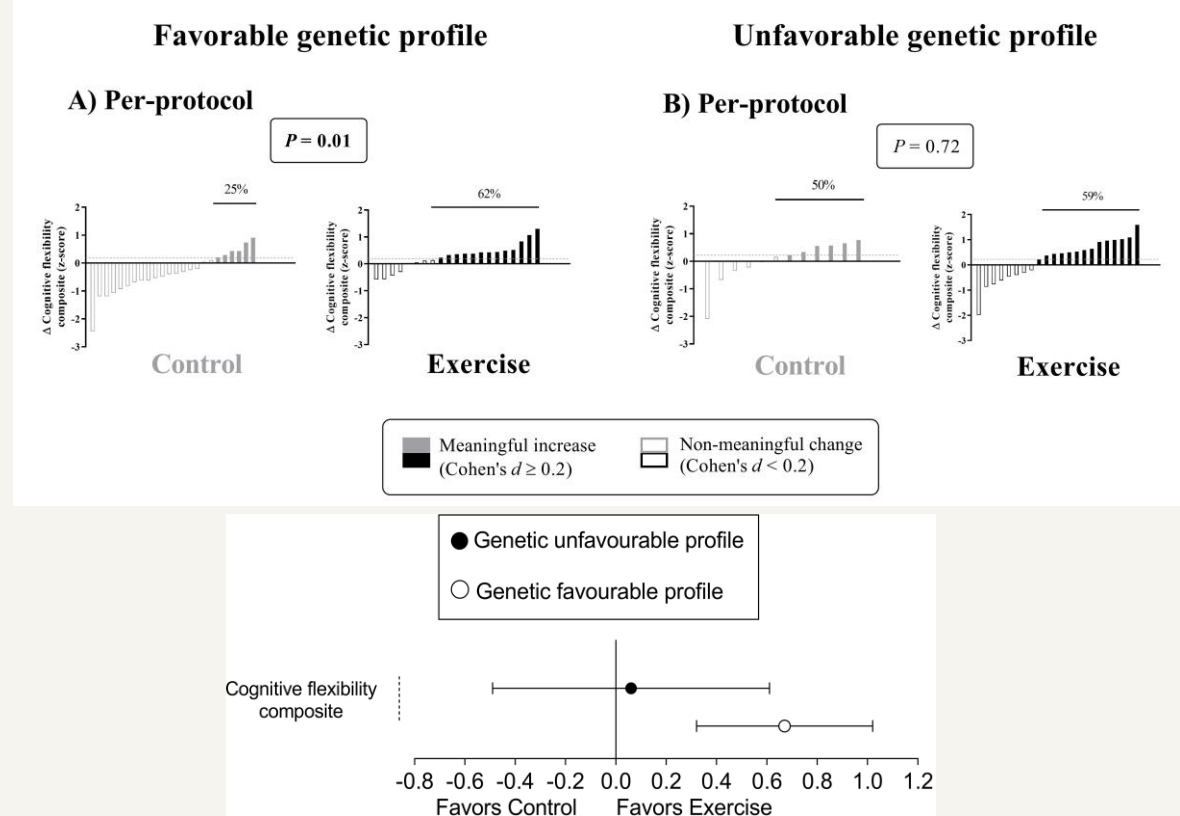
METHODS

101 children with overweight/obesity (8-11 years old) were randomly allocated to the exercise and control groups

We aimed to study the interaction between a genetic score and the effects of a 20-week exercise intervention on brain health outcomes

OUTCOME: Cognitive flexibility

CONCLUSION Our findings suggest that having a more "favorable" genetic profile makes children with overweight-obesity more responsive to exercise, particularly for cognitive flexibility.