



## Original article

# Longitudinal Evidence of a Vicious Cycle Between Nucleus Accumbens Microstructure and Childhood Weight Gain



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**Article history:** Received April 30, 2021; Accepted January 4, 2022

**Keywords:** Adolescent health; Pediatric obesity; Diet; Neuroinflammation; Nucleus accumbens; Restriction spectrum imaging

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 A B S T R A C T

**Purpose:** Pediatric obesity is a growing public health concern. Previous work has observed diet to impact nucleus accumbens (NAcc) inflammation in rodents, measured by the reactive proliferation of glial cells. Recent work in humans has demonstrated a relationship between NAcc cell density—a proxy for neuroinflammation—and weight gain in youth; however, the directionality of this relationship in the developing brain and association with diet remains unknown.

**Methods:** Waist circumference (WC) and NAcc cell density were collected in a large cohort of children ( $n > 2,000$ ) participating in the Adolescent Brain Cognitive Development (ABCD) Study (release 3.0) at baseline (9–10 y) and at a Year 2 follow-up (11–12 y). Latent change score modeling (LCSM) was used to disentangle contributions of baseline measures to two-year changes in WC percentile and NAcc cellularity. In addition, the role of NAcc cellularity in mediating the relationship between diet and WC percentile was assessed using dietary intake data collected at Year 2.

**Results:** LCSM indicates that baseline WC percentile influences change in NAcc cellularity and that baseline NAcc cell density influences change in WC percentile. NAcc cellularity was significantly associated with WC percentile at Year 2 and mediated the relationship between dietary fat consumption and WC percentile.

**Conclusions:** These results implicate a vicious cycle whereby NAcc cell density biases longitudinal changes in WC percentile and vice versa. Moreover, NAcc cell density may mediate the relationship between diet and weight gain in youth. These findings suggest that diet-induced inflammation of reward circuitry may lead to behavioral changes that further contribute to weight gain.

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**IMPLICATIONS AND CONTRIBUTION**

Neuroinflammation of reward circuitry both predicts and is predicted by weight gain in youth—a vicious cycle that may be associated with a high-fat diet. Identifying early predictors and consequences of diet-induced neuroinflammation may provide insight into interventions that have the potential to interrupt this cycle prior to obesity onset.

**Data sharing:** ABCD data are publicly available through the National Institute of Mental Health Data Archive (<https://nda.nih.gov/abcd>). The ABCD data used in this report came from the ABCD Data Release 3.0 (DOI: [10.15154/1519007](https://doi.org/10.15154/1519007), November 2020).

**Author contributions:** K.M.R., B.J.C., and R.W. conceptualized the study; K.M.R., A.B.S., and R.W. developed the methodology and performed the analyses; all authors contributed to verification of the results and research outputs; K.M.R., N.

B., A.B.S., B.J.C., and R.W. contributed to writing the original draft; all authors equally contributed to reviewing and editing the final manuscript.

**Conflict of interest:** The authors have no conflicts of interest to disclose.

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More than 18% of youth and 39% of adults are overweight or obese worldwide [1]. Obesity increases the risk of several physical illnesses such as hypertension, type 2 diabetes, coronary heart disease, stroke, cancer, and mortality [2]. Childhood overweight and obesity is a strong predictor of obesity later in life [3] and has been linked to anxiety, depression, lower self-esteem, and lower self-reported quality of life [4]. Given the high prevalence and serious physical and mental health consequences, studies have sought to understand the biological risk factors underlying the development of obesity; however, the neurobiological mechanisms leading to excessive weight gain and obesity in youth remain less well understood.

The mesolimbic dopaminergic pathway, which includes the nucleus accumbens (NAcc) [5], is necessary for motivated behavior [6], such as eating behavior. Previous work has observed varying relationships between obesity-related metrics and NAcc structure and function. For example, NAcc volume is positively associated with genetic risk for obesity in children [7], as well as with body mass index (BMI) and percent body fat—a relationship that may be age-dependent and specific to youth [8,9]. Moreover, NAcc blood oxygen level-dependent (BOLD) responses to reward cues have demonstrated associations with obesity-related behaviors and outcomes (i.e., eating behavior, weight gain) in adolescents and adults [10–13], although findings are mixed and may be dependent on additional factors [14], such as the type of reward cue (e.g., monetary reward; food reward) [15,16], stimulus type (e.g., visual gustatory) [17], and metabolic factors (e.g., insulin sensitivity) [18]. Despite a clear role of the NAcc in reward motivation more generally, the associations between this region and obesity-related behaviors and outcomes appear to be complex and dependent on a variety of factors.

Animal models of diet-induced obesity have revealed microstructural differences in the NAcc indicative of neuroinflammation [19–21]. NAcc inflammation—measured by an increase in glial cells responding to proinflammatory factors and upregulation of proinflammatory genes—has been associated with highly palatable, caloric diets in rodents [20,21], and more specifically, has been linked to a high saturated fat diet and visceral fat accumulation in mice [19]. Critically, diet-induced NAcc inflammation has been shown to modify subsequent eating behavior in rodents, and reducing this inflammation through intervention has been shown to revert increased consumption of highly palatable foods, consequently reverting diet-induced weight gain [19,20]. This work, taken together, suggests that diet, and the consumption of highly palatable foods in particular, plays a crucial role in driving NAcc inflammation, which may, in turn, promote further unhealthy eating behavior and subsequent weight gain.

Detecting neuroinflammation via reactive gliosis *in vivo* presents a challenge in the human brain. However, recent methods allow researchers to provide an index of cell density in the brain using magnetic resonance imaging (MRI). More specifically, restriction spectrum imaging (RSI) is a noninvasive imaging technique based on diffusion MRI that separates signal contributions from intracellular (restricted) diffusion and extracellular (hindered) diffusion [22,23]. RSI provides histologically validated measures of cell density in subcortical brain structures [23]. Coupled with behavioral and cellular-level evidence from experiments conducted in animals, RSI may offer complementary insight into microscale properties in the human NAcc

associated with diet and weight gain in youth. For example, RSI has been used to identify a relationship between individual differences in NAcc cell density and weight gain after one year in children [24], which may indicate diet-induced variability in glial cell density akin to those observed in animal models of obesity. Although this study demonstrates an association between NAcc cell density and weight gain *across* individuals, the extent to which weight gain contributes to—or reciprocally influences—changes in NAcc cellularity *within* an individual remains an open question. Moreover, the influence of diet on NAcc cellularity in humans remains untested.

Here we leverage longitudinal modeling in a large cohort of youth to examine directional relationships between two-year changes in NAcc cellularity and weight-related anthropometrics. BMI is a widely used index of body size and provides a clinical standard for classifying individuals according to physical thresholds (i.e., underweight, healthy weight, overweight, or obese) and according to normed growth charts in youth (e.g., BMI percentile). Although BMI percentile provides an important clinical metric for evaluating an individual's body size relative to the general population, recent discussions have highlighted potential limitations of relying on this technique [25]. Relative to BMI, waist circumference (WC) may be more informative for estimating body fat and fat gain in youth [26] and may provide a better indicator of early risk for negative health outcomes such as cardiovascular disease and metabolic dysfunction [27,28], particularly when normed according to age and sex [28].

Longitudinal changes in WC percentile and RSI were evaluated using latent change score modeling (LCSM). LCSM takes advantage of the strengths of structural equation modeling (SEM) to estimate cross-domain coupling—or the contributions of baseline measurements on changes in longitudinal data [29,30]. By explicitly specifying change scores as latent variables, LCSM allows for evaluating individual differences in intra-individual change across time points [29] and has been proposed to provide a powerful and flexible framework for understanding dynamic processes between brain and behavior underlying development [30].

Based on animal models demonstrating a vicious cycle of diet-induced inflammation of the NAcc followed by further unhealthy eating and weight gain [19,20], we hypothesized that NAcc cellularity and WC percentile would mutually influence each other. In other words, we expected NAcc cell density at baseline to predict two-year change in WC percentile and WC percentile at baseline to predict two-year change in NAcc cell density. Given the role of diet in driving neuroinflammation in animal models of obesity, we additionally hypothesized that NAcc cellularity would mediate the relationship between fatty diet and WC percentile in youth.

## Methods

### Data source

The Adolescent Brain Cognitive Development<sup>SM</sup> (ABCD) Study is an ongoing longitudinal study of brain development and child health in the United States, following over 11,000 9–10-year-olds through adolescence [31]. The baseline cohort was recruited from 21 sites using a rigorous epidemiologically informed school-based sampling and recruitment strategy, with the objective of approximating the demographic and socioeconomic

diversity of the U.S. population [32]. Data is collected from consenting parents and assenting children through yearly multimodal assessments, including environmental, behavioral, physical health, and neurocognitive measures—as well as biennial structural and functional MRI scans. ABCD Study recruitment, sample selection, complete battery of assessments, study design, and data collection are detailed elsewhere [33]. Study-wide exclusion criteria for enrollment included a diagnosis of moderate to severe autism spectrum disorder, schizophrenia, moderate to severe intellectual disability, major neurological disorders, or a substance use disorder at recruitment. Children with noncorrectable vision, hearing, or sensorimotor impairments, gestational age less than 28 weeks, birth weight less than 1.2 kg, birth complications requiring more than a 1-month hospitalization, history of traumatic brain injury, and standard MRI contraindications (e.g., implanted metals, claustrophobia, orthodonture) were also excluded. All study procedures were approved by the participating study site Institutional Review Boards and by the ABCD Study centralized Institutional Review Board.

### Participants

Analyses were conducted on data from the ABCD Study 3.0 release, which includes baseline data from 11,875 participants and 6,571 participants at a 2-year follow-up. In addition to study-wide exclusionary criteria, the current analysis excluded participants reporting a history of neurological disorders (e.g., cerebral palsy, seizures), concussion, diabetes, lead poisoning, muscular dystrophy, multiple sclerosis, and substance abuse, as well as participants presently or previously meeting diagnostic criteria for an eating disorder (anorexia nervosa, bulimia nervosa, binge eating disorder), schizophrenia, or psychosis (assessed using the K-SADS-PL) at baseline and/or at Year 2. Moreover, participants identified as having low-quality anatomical images (ABCD NDA name: *fsqc\_qc*) were further excluded from respective analyses at baseline and Year 2. For consistency within the data, only subjects whose data were acquired using MRI scanners from a single vendor (Siemens Healthineers AG, Erlangen, Germany) were included in our analysis ( $n = 14$  of 21 sites). To avoid extreme values due to potential measurement error, participants whose waist circumference, BMI, or NAcc cellularity fell outside of four standard deviations from the group mean at either time point ( $n = 35$ ) were excluded from further analysis. Participants with missing data for any variables of interest or covariates were further excluded, resulting in 2,378 participants with complete data at Year 2 and 2,333 participants with complete data at both baseline and Year 2 (44.6% female; mean [s.d.] age at baseline: 9.97 [.61] years; mean [s.d.] at Year 2: 11.96 [.63] years) (see Table 1 for complete participant demographics).

### Data acquisition and preprocessing

**Waist circumference.** WC was measured for each participant at baseline and Year 2. Measurements were taken by placing a tape measure along the highest point of the pelvic bone and rounded to the nearest .1 inch. Measurements were collected twice and averaged to maximize accuracy. In accounting for differences in age-specific and sex-specific growth curves, resulting values were converted to percentiles based on data from the US

**Table 1**  
Participant demographics

Measurement	Baseline	Year 2
Waist circumference (%ile)	59.56 (28.88)	62.04 (28.05)
Body Mass Index (%ile)	58.44 (30.52)	60.95 (30.60)
Underweight (%)	4.03	3.78
Healthy weight	68.71	65.14
Overweight	13.46	15.35
Obese	13.80	15.73
NAcc cellularity	0.21 (0.02)	0.22 (0.02)
NAcc volume (cm <sup>3</sup> )	5.80 (.88)	5.76 (.88)
Head motion (mm)	1.29 (.52)	1.16 (.44)
Age (yrs)	9.97 (.61)	11.96 (.63)
Sex (%F)	44.62	—
Puberty (stage)		
Prepuberty	54.35%	22.54%
Early puberty	23.23%	26.07%
Mid puberty	21.69%	33.64%
Late puberty	0.73%	17.75%
Race/ethnicity (%)		
White	64.63	—
Black	10.89	—
Hispanic	16.12	—
Asian	0.69	—
Other	7.67	—
Parent marital status (%M)	73.89	—
Parent income (%)		
< \$50,000	24.13	—
\$50,000 to \$100,000	32.23	—
> \$100,000	43.63	—
Parent education (%)		
No high school diploma	2.10	—
High school diploma or GED	5.14	—
Some college	25.42	—
Bachelor's degree	30.78	—
Postgraduate degree	36.56	—
Dietary fat (g)	—	49.90 (21.80)
Dietary carbohydrates (g)	—	142.08 (55.14)
Dietary protein (g)	—	50.76 (21.48)
Dietary fiber (g)	—	10.36 (4.58)
Dietary caloric intake (kcal)	—	120.32 (46.85)

Descriptive statistics for all variables of interest and covariates collected during baseline and/or Year 2. Values represent mean (s.d.) unless specified otherwise. Genetic ancestry scores were utilized for all analyses, but self-report race and ethnicity data are included here for ease of interpretability.

National Health and Nutrition Survey (NHANES III) [28] using the R package *childsds*.

**Body mass index.** Standing height and weight were measured using a stadiometer and digital scale, respectively. Measurements were collected twice and averaged to maximize accuracy. BMI was calculated for each participant using the following formula:  $\text{weight (kg)} / (\text{height [cm]} / 100)^2$ . The resulting BMI values were converted to CDC-standard percentiles and stratified according to BMI class using the R package *PAutilities*.

**Diet.** Dietary fat consumption was estimated using the Block Kids Food Screener (BKFS). The BKFS is a parent-reported, youth-confirmed assessment measuring food intake over the previous week. Quantity and frequency for each of 39 food items are collected and immediately analyzed by the NutritionQuest database for average daily intake of predetermined dietary variables based on the participant's age-sex group. Dietary intake data was only collected at the Year 2 follow-up and was not collected at baseline.

**Pubertal status.** Child pubertal status was assessed by self-report and parent-report of physical development, yielding a categorical maturation score similar to that of Tanner staging.

**Parent marital status, income, and education.** Parent-reported demographic covariates included total combined family income (*less than \$50,000; between \$50,000 and \$100,000; greater than \$100,000*), marital status (*married; single*); and parental years of education (*No high school diploma; high school diploma or GED; Some college; Bachelor's degree; Post-graduate degree*).

**Genetic ancestry.** Saliva samples were collected at baseline and immediately shipped to Rutgers University Cell and DNA Repository (RUCDR), where the sample was prepared for genotyping. Genotyping data for 733,293 single nucleotide polymorphisms (SNPs) were generated using the Affymetrix NIDA Smokescreen™ array.

**Restriction spectrum imaging.** Diffusion images were acquired at baseline and Year 2 using a spin-echo EPI acquisition with TE/TR = 88/4100 ms, multiband acceleration factor of 3, phase partial Fourier factor of .75, matrix size of 140x140, 81 slices, and an axial acquisition with 1.7-mm isotropic resolution. Diffusion-weighted data were acquired with six directions at  $b = 500$  s/mm<sup>2</sup>, 15 directions at  $b = 1000$  s/mm<sup>2</sup>, 15 directions at  $b = 2000$  s/mm<sup>2</sup>, and 60 directions at  $b = 3000$  s/mm<sup>2</sup>. The RSI model was fitted on a voxelwise basis at baseline and Year 2 using a linear estimation approach [22]. The NAcc was anatomically defined using automated atlas-based segmentation and used to extract NAcc-specific cellularity estimates. Subject-specific estimates of head motion (mean framewise displacement) during diffusion scans were included as a covariate in all RSI analyses.

#### Longitudinal changes

Linear mixed-effects models (*lme4*) were used to quantify two-year changes in body measurements. Fixed effect covariates included age, sex, pubertal stage, genetic ancestry, parental education, income, and marital status. Random effects included subject ID and family ID nested within site. Two-year change in NAcc cellularity was similarly assessed with NAcc volume as an additional fixed covariate and family ID nested with scanner ID (rather than site) as a random effect.

#### Latent change score modeling

Bivariate LCS modeling was performed using Lavaan in R and utilized publicly available code provided by Kievit et al. [30] (<https://osf.io/4bpmq/>). Two-year changes in WC percentile and NAcc cellularity were modeled as latent change scores to identify contributions of baseline measurements on respective outcomes. To provide a comparison to a clinical standard, two-year changes in BMI percentile were additionally modeled using an identical statistical framework (see Appendix A1). Models were computed using maximum likelihood estimation with robust standard errors and a Yuan-Bentler correction for non-normality.

Time-dependent covariates in the model included age (mean-centered), pubertal stage, head motion (framewise displacement) during respective RSI scans, and NAcc volume at baseline and Year 2. Time-independent covariates included sex, parental education, household income, parent marital status, and continuous genetic estimates of African, American, and Asian

ancestry. Time-independent covariates were included for baseline WC percentile and NAcc cellularity. Time-dependent covariates were modeled as separate regressions with both WC percentile and NAcc cellularity for respective time points and were allowed to covary between time points. Variables were allowed to covary based on known associations (e.g., age and puberty) and as observed in the current dataset (see Figure A1). LCSMs were additionally computed without covariates to rule out the possibility that covariate adjustment led to false or misleading findings [34], without the inclusion of siblings ( $n = 301$ ; randomly selected from each sibling pair) to ensure effects were not influenced by family structure, and with site-wise regressors (dummy coded as 0 or 1) to account for potential differences across sites.

Based on a two-index presentation strategy recommended by Hu and Bentler [35], LCSM fit was assessed using the Comparative Fit Index (CFI) and Root Mean Square Error of Approximation (RMSEA). CFI is a relative measure of model fit, comparing the hypothesized model to an unstructured baseline model and adjusting for sample size. CFI ranges from 0 to 1, with scores greater than 0.95 indicating a good fit. RMSEA is an absolute measure of model fit that compares the hypothesized model to the population covariance matrix, with values ranging from 0 to one and scores less than 0.06 indicating a close fit. Effects of interest were bootstrapped (5,000 iterations) to estimate 95% confidence intervals, and path coefficients were standardized to allow for interpretability across variables included in the model.

#### Mediation analysis

To further probe the relationship between NAcc cellularity and obesity-related outcomes, a mediation analysis was performed. Diet information obtained at the Year 2 follow-up allowed for a secondary analysis of the relationship between diet and WC percentile, mediated by NAcc cell density. Dietary fat was used to assess this relationship based on animal literature demonstrating the role of a high-fat diet in promoting NAcc inflammation [19,20] and subsequent weight gain. Dietary fat was normalized by total caloric consumption to obtain an estimate of relative dietary fat consumed. Nonfat macronutrients (dietary carbohydrates, protein, and fiber—each normalized by total caloric consumption) and total caloric intake were used for comparison to demonstrate specificity to dietary fat.

For consistency across analyses, Lavaan was used to perform the mediation analysis within an SEM that included all time-independent covariates used in the LCSM (i.e., sex, genetic ancestry, parental income, marital status, and education), as well as time-dependent covariates at Year 2 (i.e., age, puberty, head motion, and NAcc volume). Nonimaging covariates were regressed with dietary fat.

## Results

#### Longitudinal change

WC percentile was significantly correlated with BMI percentile at baseline (Pearson's  $r = 0.69$ ; 95% CI: [0.67, 0.71];  $p < .0001$ ) and at Year 2 (Pearson's  $r = 0.76$ ; 95% CI: [0.74, 0.77];  $p < .0001$ ). Two-year change in WC percentile was significantly correlated with two-year change in BMI percentile (Pearson's  $r = 0.34$ ; 95% CI: [0.30, 0.37]). Baseline WC percentile (Figure 1A) and BMI percentile (Figure 1B) significantly increased at the Year 2

follow-up visit (WC:  $\beta = 2.46$ ; SE = 0.47;  $t = 5.26$ ;  $p < .0001$ ; BMI:  $\beta = 2.24$ ; SE = 0.34;  $t = 6.68$ ;  $p < .0001$ ). Likewise, NAcc cell density significantly increased after two years ( $\beta = 0.011$ ; SE = 0.0003;  $t = 38.50$ ;  $p < .0001$ ) (Figure 1C).

#### Latent change score model

The LCSM demonstrated a good model fit (CFI = .98; Root-Mean-Square Error of Approximation [RMSEA] = 0.039; 90% CI: [0.036, 0.041];  $p \leq .05 = 1.0$ ) (Figure 2). Baseline measures of both

WC percentile and NAcc cellularity negatively predicted two-year changes in respective measures (WC:  $\beta = -0.49$ ;  $p < .0001$ ; NAcc cellularity:  $\beta = -0.43$ ;  $p < .0001$ ), such that higher values at baseline corresponded with smaller change scores. Cross-domain paths demonstrated that baseline WC percentile predicted two-year change in NAcc cellularity ( $\beta = 0.10$ ; 95% CI = [0.06, 0.14];  $p < .0001$ ) and that NAcc cellularity predicted two-year change in WC percentile ( $\beta = 0.08$ ; 95% CI = [0.04, 0.12];  $p = .001$ ). Although the effect size for the relationship between WC percentile and change in NAcc cellularity was stronger, bootstrapped confidence intervals demonstrate no difference in strength of cross-domain parameter estimates. Using the likelihood ratio test, model fit significantly decreased when either cross-domain path was constrained to zero (baseline WC percentile:  $\Delta\chi^2(1) = 24.19$ ;  $p < .0001$ ; baseline NAcc cellularity:  $\Delta\chi^2(1) = 17.24$ ;  $p < .0001$ ).

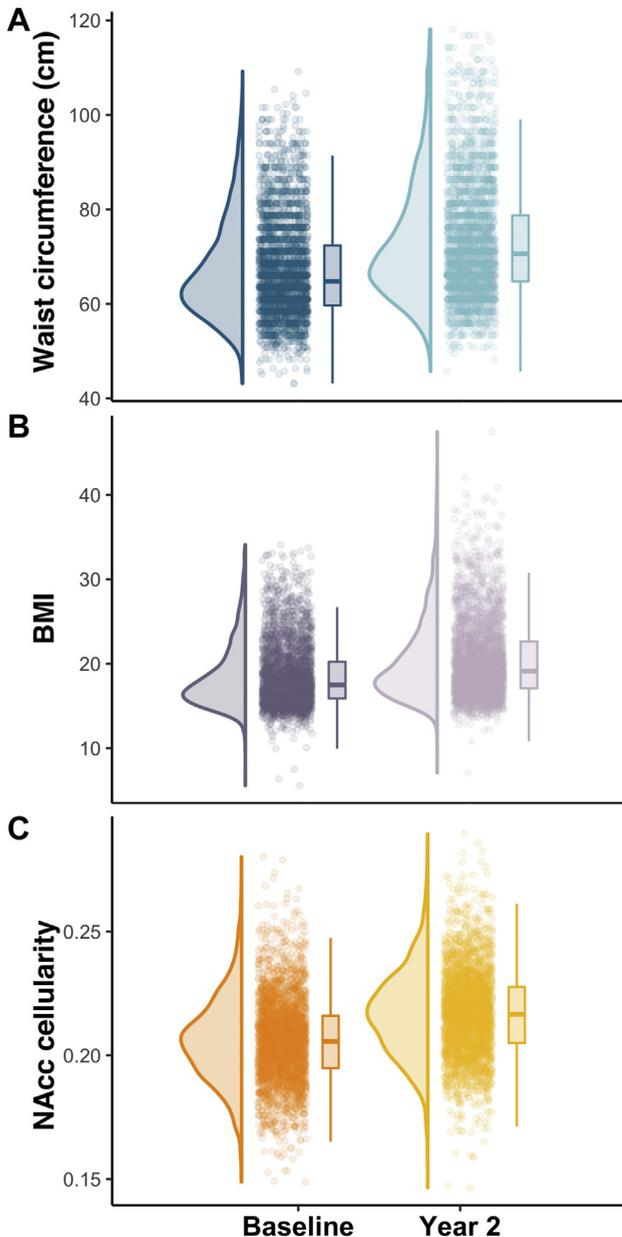
Consistent with previous work [24], WC percentile and NAcc cellularity were significantly associated at baseline ( $\beta = 0.16$ ;  $p < .0001$ ). Moreover, there was a significant association between latent change scores for WC percentile and NAcc cellularity ( $\beta = 0.07$ ;  $p < .001$ ). Excluding all covariates from the LCSM (Figure A3), excluding siblings (Figure A4), or including site-wise regressors (Figure A5) did not affect the interpretation of results, suggesting that the effects observed here are robust to the inclusion of covariates, siblings, and potential site differences. Similar effects were also observed when considering change in BMI percentile (see Appendix A1) regardless of whether covariates were included (Figure A6) or excluded (Figure A7).

#### Association with diet at year 2

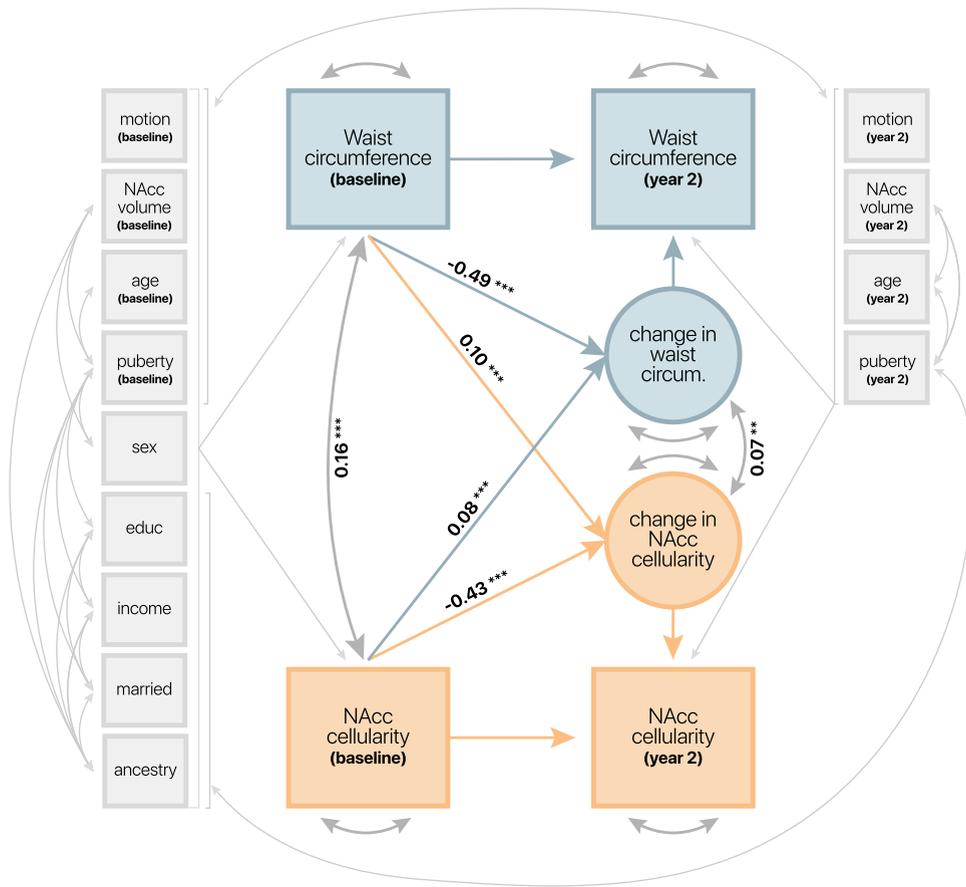
Anthropometrics such as waist circumference and BMI may serve as a proxy for the cumulative impact of diet; however, directly testing the contribution of diet on NAcc cellularity is needed to understand the potential role of diet-induced neuroinflammation on childhood weight gain. Food intake data collected at Year 2 allowed for a more explicit evaluation of the associations between diet, NAcc cellularity, and anthropometrics. An analysis of NAcc cellularity mediating the relationship between diet and WC percentile demonstrated good model fit (CFI = 0.97; RMSEA = 0.035; 90% CI: [0.031, 0.039];  $p \leq .05 = 1.0$ ) (Figure 3).

Dietary fat was significantly associated with WC percentile at Year 2 (total effect [path c]:  $\beta = 0.07$ ; 95% CI = [0.03, 0.11];  $p < .001$ ) and was fully mediated by NAcc cellularity (direct effect [path c']:  $\beta = 0.036$ ; 95% CI = [-0.002, 0.07];  $p = .06$ ). Dietary fat was associated with NAcc cellularity (path a:  $\beta = 0.049$ ; 95% CI = [0.01, 0.08];  $p = .008$ ), and consistent with previous findings at baseline and Year 1 [24], NAcc cellularity was associated with WC percentile at Year 2 (path b:  $\beta = 0.18$ ; 95% CI = [0.15, 0.22];  $p < .0001$ ). The indirect effect of NAcc cellularity on the relationship between dietary fat and WC percentile was significant ( $a*b$ :  $\beta = 0.01$ ; 95% CI = [0.003, 0.02];  $p < .01$ ; proportion mediated = 0.20), suggesting a role of NAcc cell density in mediating diet-induced weight gain in youth. Similar results were observed using BMI percentile (see Appendix A1; Figure A8).

As a comparison, additional mediation models were tested using nonfat macronutrients (dietary carbohydrates, protein, fiber) and total caloric intake as independent variables in identical models. Among nonfat macronutrients and caloric intake, only dietary fat showed a significant positive relationship with WC percentile that was mediated by NAcc cellularity (see



**Figure 1.** Two-year change in waist circumference and NAcc cellularity. Raincloud plots of waist circumference (A), BMI (B), and RSI-based NAcc cellularity (C) at baseline and at Year 2 follow-up. Sex-specific WC and BMI distributions are visualized in Figure A2.



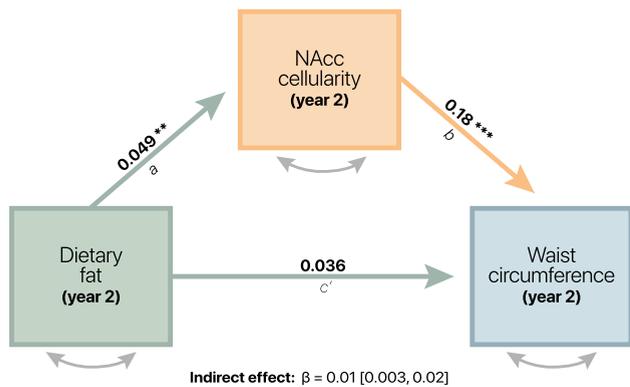
**Figure 2.** Latent change score model demonstrating longitudinal associations between waist circumference percentile and NAcc cellularity. Standardized coefficients plotted for paths of interest. Significance represented as  $p < .0001$  (\*\*\*) ;  $p < .001$  (\*\*). Thin, light gray arrows represent covariate paths of no interest.

Appendix A1). Dietary carbohydrates demonstrated an inverse association with NAcc cellularity and waist circumference (Figure A9), and no significant mediations were observed with dietary fiber, protein, or total caloric intake.

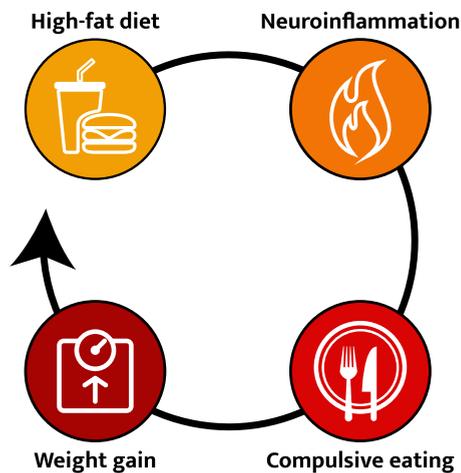
**Discussion**

Consistent with the “vicious cycle” hypothesis of diet-induced brain changes promoting unhealthy eating and weight gain [36], the present study observed reciprocal influences of baseline waist circumference percentile (as well as BMI percentile) and NAcc cellularity on two-year changes in the converse measures. In addition, NAcc cell density was found to mediate the relationship between diet and WC percentile at the Year 2 follow-up. These findings replicate and extend prior work linking NAcc cellularity and weight gain in youth [24] by further demonstrating the longitudinal associations between obesity-related metrics (e.g., WC) and the microstructural properties of the developing brain. Moreover, these findings mirror rodent models of obesity that have demonstrated diet-induced neuro-inflammation of the NAcc, marked by an increase in glial cell proliferation [19,20]. The present study, taken together with previous human and animal literature [19,24], suggests that diet influences NAcc inflammation, which may, in turn, contribute to further unhealthy eating and weight gain (Figure 4). Future work is needed to disentangle potential mechanisms underlying this cycle and to more explicitly test relationships within this framework that were not directly assessed in the present study.

Despite the evidence supporting weight-related changes in NAcc inflammation observed here, the mechanisms underlying these proposed inflammatory changes remain unclear.



**Figure 3.** Mediation model. NAcc cellularity significantly mediated the relationship between dietary fat and waist circumference percentile at Year 2. Path coefficients are standardized, and significance is represented as  $p < .0001$  (\*\*\*) ;  $p < .01$  (\*\*). Covariates were consistent with those included in the LCSM at Year 2.



**Figure 4.** Cartoon schematic illustrating the proposed cycle of a highly palatable diet contributing to neuroinflammation, particularly within brain regions associated with reward (e.g., the NAcc), subsequently influencing behavioral changes in eating behaviors and increases in weight gain.

The production of proinflammatory cytokines by white adipose tissue is a major source of obesity-related inflammation [37]. Diets that are high in fat and sugar promote abdominal fat accumulation, and thus, greater waist circumference to favor local immune responses that can propagate to the brain [38]. Previous studies have found that prolonged exposure to a highly palatable (i.e., high-fat, high-sugar) diet additionally produces neuroplastic and functional changes in the NAcc that influence behavior. For example, Gutiérrez-Martos et al. [20] examined the NAcc of mice fed a “cafeteria” diet and observed an increase in the expression of inflammatory cytokines (IL-1 $\beta$ , IFN- $\gamma$ ), as well as a morphological change in microglia characteristic of a reactive state. This diet-induced neuroinflammation was associated with increased consumption of calories and a corresponding increase in body weight. However, these behavioral changes reversed when inflammation and microglia activation was reduced via systemic administration of minocycline, a broad inhibitor of peripheral and central inflammation. Importantly, inflammatory responses were accompanied by structural changes in dendritic cell density, which may contribute to functional differences in the rewarding effects of food and food-motivated behaviors.

A number of studies have begun to explore the possibility that the impact of palatable foods on neural plasticity [39], as well as on neuroinflammatory responses [40], may contribute to altered emotional and cognitive processing [41], ultimately giving rise to dysfunctions in learning and memory, mood regulation, and compulsive behaviors. The direct contribution of NAcc inflammation to heightened food-seeking in obesity is underscored by observations of reduced compulsivity for sugar in diet-induced obese mice with targeted genetic inhibition of inflammation in the NAcc [19]. In another study, increased dendritic spine density within the NAcc was associated with the consumption of palatable foods— independent of caloric content— and occurred specifically in mice demonstrating enhanced food-seeking behaviors [42]. Although the consumption of high palatable foods appears to drive NAcc inflammation, different classes of macronutrients have been shown to differentially impact inflammatory responses. For example, several studies have found that a high-fat

diet leads to increased neuroinflammation [20,43–45], and NAcc inflammation has been specifically linked to saturated fats [19]. Moreover, the present study observed an inverse association between WC percentile and dietary carbohydrates—an association that was negatively mediated by NAcc cell density (Figure A9). Future work should aim to further disentangle the contributions of various macronutrients on neuroinflammation, as well as consider a possible protective mechanism whereby certain macronutrients buffer against neuroinflammation.

Another potential mechanism by which this cycle may occur is in interactions with neuroinflammation in the hypothalamus and subsequent differences in hypothalamic neuroendocrine and metabolic regulation. Hypothalamic inflammation accelerates energy imbalance [46], interferes with the ability to regulate food intake [44], and has been suggested to modify hypothalamic circuitry and interfere with outputs to other brain regions—including regions involved in reward-processing [45] and eating behavior [47]. As a result, the hypothalamus has been a key target for investigations of diet-induced neuroinflammation in rodents, which suggests that prolonged exposure to high-fat diets and associated metabolic dysfunction are drivers that sustain neuroinflammation in subcortical structures [43,48]. Methodological challenges have precluded replicable segmentation of the hypothalamus in human neuroimaging, and thus hypothalamus-specific cellularity estimates are not available in the ABCD Study release 3.0 data. However, recent advances in machine learning have allowed for the development of new segmentation algorithms that are capable of automated hypothalamic segmentation using a deep convolutional neural network [49]. As tools such as these become more readily available, future work will be able to examine potential relationships between diet and neuroinflammation of the hypothalamus and NAcc in humans.

Several limitations of the present study warrant further consideration. First, although the study motivation and interpretation are based on prior work conducted in animal models of obesity and neuroinflammation [19], the ABCD Study release 3.0 dataset does not include direct markers of neuroinflammation. Recent work has used diffusion-based spectrum imaging (DBSI) to relate imaging markers of striatal neuroinflammation to self-report emotional eating and obesity in adults [50]. DBSI provides biomarkers of inflammation by characterizing water diffusion properties associated with axon/myelin injury and inflammation [51] and may provide convergent information alongside RSI-based measures of cell density. However, additional work is needed to directly quantify and assess the relationship between eating behavior, obesity, and in vivo neuroinflammation in humans. Human neuroimaging has demonstrated relationships between food reward sensitivity in the NAcc and genetic risk for obesity [7], eating behavior [52], and weight gain [53] in youth, yet it remains unclear how these findings might relate to potential diet-induced inflammatory factors. Future studies are needed to integrate prior work in animals with human studies examining reward-related brain function and behavior.

Second, waist circumference measurements are prone to measurement error. Variation in WC measurements may be due to inconsistencies across experimenters, differences in lean muscle and bone mass, and difficulty locating anatomical landmarks in overweight participants [54]. The present study sought to identify potential measurement errors by excluding individuals with waist circumference values (and BMIs) outside of four standard deviations; however, this exclusion criteria does

not identify inaccurate values within four standard deviations. Moreover, waist circumference and BMI—even when accurately measured—may not adequately capture adiposity levels relative to more direct measures (e.g., dual-energy X-ray absorptiometry) [55]. Future work should consider additional anthropometrics that more directly and accurately measure fat mass associated with negative health outcomes.

Third, to test a specific hypothesis regarding dietary fat based on prior work in animal models [19], the present study utilized self-report dietary intake data to estimate the relative amount of dietary fat an individual consumes daily. However, individuals tend to poorly estimate their energy intake and commonly underestimate the number of calories they consume [56]. Although unavailable in the current ABCD Study data release (3.0), future work should consider assessing dietary metabolites and corresponding blood markers directly (e.g., LDL cholesterol levels increased by the consumption of saturated and trans fats) as an alternative to self-report intake data.

Fourth, the present study examined relationships between NAcc cellularity and nonfat macronutrients (carbohydrates, fiber, protein) to consider the specificity of the observed relationship to dietary fat. We found that NAcc cellularity fully mediated the relationship between WC percentile and dietary fat intake only and did not mediate relationships with dietary fiber, protein, or overall caloric intake. However, we additionally observed a partial negative mediation with dietary carbohydrates (Appendix A1) such that carbohydrate intake was inversely associated with NAcc cellularity and WC percentile at Year 2 (Figure A9). Although these findings suggest that dietary fat intake may increase neuroinflammation and weight gain—and that dietary carbohydrates may buffer against these effects—additional work is needed to further examine these relationships. Macronutrients are not consumed in isolation, and combinations of different food groups can have synergistic effects on an individual's health. Thus, caution is necessary in interpreting these single-nutrient results. In addition to measuring blood markers of diet and health, future work should consider evaluating an index of the overall pattern of diet, which may provide a more informative indicator of diet quality.

Finally, the interpretation of our results is constrained by the selection of variables and inclusion criteria of participants. For example, the present study does not incorporate other factors that may influence dietary intakes—such as physical activity, access to healthy foods, and medications that affect appetite. In addition, the present study exclusively examines data from participants scanned using a single MRI manufacturer (Siemens). Although excluding MRI data from other scanner manufacturers (Philips and GE) reduces potential confounds due to the scanner platform, doing so may inadvertently bias the sample demographics such that participants from underrepresented groups or with limited access to resources are disproportionately excluded. Future work should take into consideration additional covariates, inclusion criteria (e.g., non-Siemens data), and exclusionary criteria (e.g., medications) that may impact individual differences in diet and how these relate to neuroinflammation in youth.

The present study, taken together, builds on previous literature in animals and in humans to understand the relationship between neuroinflammation of the NAcc and diet-induced weight gain. We observed longitudinal changes in NAcc cellularity to be reciprocally influenced by changes in WC percentile, and further demonstrated a role of NAcc cell density in mediating

the relationship between dietary fat intake and WC percentile. These findings extend prior work linking NAcc cell density with weight gain in youth [24] and suggest that diet plays an important role in neurodevelopmental changes that may influence eating behavior. Given that adolescence is characterized by neurodevelopmental changes, as well as a heightened sensitivity to learning [57] and the formation of habits [58] that can affect later health outcomes, it is crucial to understand the impact of diet on the developing brain, as well as the behavioral consequences. Understanding the role of diet-induced neuroinflammation on the developing brain may provide key insight into interventions that can mitigate pediatric obesity.

## Acknowledgments

Data used in the preparation of this article were obtained from the ABCD study (<https://abcdstudy.org>), held in the National Institute of Mental Health (NIMH) Data Archive (NDA). This is a multisite, longitudinal study designed to recruit more than 10,000 children ages 9 to 10 and follow them over 10 y into early adulthood.

## Funding Sources

The ABCD Study is supported by the NIH and additional federal partners under award numbers U01DA041048, U01DA050989, U01DA051016, U01DA041022, U01DA051018, U01DA051037, U01DA050987, U01DA041174, U01DA041106, U01DA041117, U01DA041028, U01DA041134, U01DA050988, U01DA051039, U01DA041156, U01DA041025, U01DA041120, U01DA051038, U01DA041148, U01DA041093, U01DA041089, U24DA041123, and U24DA041147. A full list of supporters is available at <https://abcdstudy.org/federal-partners.html>. A listing of participating sites and a complete listing of the study investigators can be found at [https://abcdstudy.org/consortium\\_members/](https://abcdstudy.org/consortium_members/). ABCD consortium investigators designed and implemented the study and/or provided data but did not necessarily participate in the analysis or writing of this report. This manuscript reflects the views of the authors and may not reflect the opinions or views of the NIH or ABCD consortium investigators. The ABCD data repository grows and changes over time. The ABCD data used in this report came from NIMH Data Archive Digital Object Identifier DOI 10.15154/1519007. DOIs can be found at <https://nda.nih.gov/study.html?id=901>. This work was supported in part by U01 DA041174 (to B.J.C.) and R01DK097399-01 (to K.M.R.). The funders had no role in study design, data collection, and analysis, decision to publish, or preparation of the manuscript.

## Supplementary Data

Supplementary data related to this article can be found at <https://doi.org/10.1016/j.jadohealth.2022.01.002>.

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