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Psychotic-like experiences and polygenic liability in the ABCD Study®

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Abstract

Background: Childhood psychotic-like experiences (PLEs) often precede the development of later severe psychopathology. This study examined whether childhood PLEs are associated with several psychopathology-related polygenic scores (PGS), and additionally examined possible neural and behavioral mechanisms.

Methods: Adolescent Brain Cognitive DevelopmentSM Study baseline data from children with European ancestry (n=4,650; ages 9-10; 46.8% female) were used to estimate associations between PLEs (i.e., both total and presence of significantly distressing) and PGS for psychopathology (i.e., schizophrenia, psychiatric cross-disorder risk, PLEs) and related phenotypes (i.e., educational attainment [EDU], birth-weight, inflammation). We also assessed whether variability in brain structure indices (i.e., volume, cortical thickness, surface area), as well as behaviors proximal to PGS (e.g., cognition for EDU), indirectly linked PGS to PLEs using mediational models.

Results: Total and significantly distressing PLEs were associated with EDU and cross-disorder PGS (all % R^2 s=0.202-0.660%; p_{FDR} s<0.006). Significantly distressing PLEs were also associated with higher schizophrenia and PLEs PGS (both % R^2 =0.120-0.216%; p_{FDR} s<0.03). There was evidence global brain volume metrics and cognitive performance indirectly linked EDU PGS to PLEs (estimated proportion mediated= 3.33-32.22%).

Conclusions: Total and significantly distressing PLEs were associated with genomic risk indices of broad-spectrum psychopathology risk (i.e., EDU and cross-disorder PGS). Significantly distressing PLEs were also associated with genomic risk for psychosis (i.e., schizophrenia, PLEs). Global brain volume metrics and PGS-proximal behaviors represent promising putative intermediary phenotypes that may indirectly link genomic risk to psychopathology. Broadly,

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polygenic scores derived from genome-wide association studies of adult samples generalize to indices of psychopathology risk among children.

Keywords

psychotic-like experiences; polygenic; schizophrenia; educational attainment; psychopathology; MRI

Introduction

Psychotic-like experiences (PLEs) are nonclinical schizophrenia-spectrum symptoms that include perceptual abnormalities and mild delusional thoughts. They commonly occur in children (~10% of children and adolescents) and are considered a dimensional, transdiagnostic marker of significant psychopathology risk (e.g., odds ratio ~3), including conversion to adult psychotic disorders in some children (1,2). Indeed, supporting the potential validity of PLEs as markers of psychopathology and psychosis-specific risk, PLEs are associated with a range of risk factors (e.g., family history of psychotic disorders, developmental milestone delays, cognition, and neural correlates) within the Adolescent Brain Cognitive DevelopmentSM (ABCD) Study (3-5). The adverse mental health prognosis of children with PLEs, even beyond those who eventually develop schizophrenia, has inspired efforts to improve our understanding of PLE etiology to ultimately facilitate advances in prevention and treatment.

Building upon twin work documenting the moderate heritability of PLEs, genome-wide association studies (GWASs) have shown that PLEs are highly polygenic, much like other complex behavioral and biological phenotypes (6). Results from well-powered discovery GWASs may be projected onto individuals in an independent sample by averaging common variants weighted by GWAS effect size and number of risk alleles present across the genome to generate polygenic scores (PGS) that represent an individual's genomic predisposition for the discovery GWAS phenotype (7). Initial evidence suggested null associations between schizophrenia PGS and adolescent PLEs using an initial discovery GWAS (cases n=9,394(8)) (9-11). In contrast, recent work leveraging results from a larger GWAS of schizophrenia (cases n=36,989 (12)) has generally linked schizophrenia PGS to PLEs during adolescence and adulthood (e.g., ages 15-19) (13) (although see (14)) and predicted psychosis conversion in individuals at risk for psychosis (15). Further, other work has found associations between later-life PLEs and both schizophrenia and mood disorder PGSs (6).

This study examined associations between childhood PLEs and several polygenic scores associated with risk for psychopathology (e.g., psychosis), including scores that may be putatively associated with PLEs through neural or behavior mechanisms (e.g., inflammation, birthweight). In addition to PLEs being related to schizophrenia risk, associations between PLEs and general psychopathology (6,16) suggest that genomic vulnerability to broad-spectrum psychopathology may confer risk for PLEs in childhood. Further, clues from epidemiological studies linking psychosis spectrum symptoms to low birthweight (17), inflammation (18), and reduced educational attainment (i.e., a measure of both cognitive functioning and non-cognitive factors, including risk taking and household income,

considered risk factors for psychosis (19,20)) (21) raise the possibility that polygenic propensity for these phenotypes may correlate with PLEs independently or through their phenotypic expression. PGS scores for birthweight and inflammation may provide important insights regarding associations between genetic liability for early developmental environmental insults and PLEs. Finally, emerging evidence linking PLEs to lower global brain volume (22) as well as evidence of brain volume indirectly linking risk factors to PLEs in the ABCD Study[®] (23), provide a basis for the possibility that brain structure may indirectly link genomic risk to PLE expression (24).

This study examined data from non-Hispanic children of European ancestries (n=4,650; aged 9-10) who completed the baseline session of the ABCD Study[®]. We tested whether PLEs are associated with genomic liability to schizophrenia, psychiatric cross-disorder risk, PLEs, educational attainment (EDU), birthweight, and inflammation (i.e., c-reactive protein) in school-age children. As childhood PLEs represent potential harbingers of adult psychopathology, it is critical to understand whether, as expected, polygenic liability estimates derived from GWASs of adult phenotypes are associated with their expression in childhood, whether these associations with PGS vary according to PLE severity, and whether variability in brain structure and behavior indirectly link polygenic vulnerability to the expression of PLEs in children.

Methods

Participants

A sample of 11,875 individuals was obtained from the ABCD Study[®] (Data Release 2.0.1; see Acknowledgements), a large-scale ongoing longitudinal study of children recruited from 22 research sites across the United States (25). The ABCD Study aimed to explore factors associated with development of both healthy behaviors and mental health challenges, and aimed to utilize a multi-stage probability sample of eligible youth (Supplement for study-wide exclusion criteria, power analysis), selecting a stratified, probability sample of schools across the United States designed to capture demographic diversity (26-28). Participants who did not pass quality control metrics and those who were not of European ancestries were removed, leaving a final analytic sample of 4,650 (46.8% female; mean age=9.93±0.63 [range=9.00-10.92] years; Supplemental Figure 1 for sample overlap with European ancestry reference population). Additionally, a sample of individuals with African ancestries was used in exploratory analyses (n=1,201; see Supplement). Caregivers provided written informed consent and all children provided assent.

Measures

Psychotic-like Experiences.—Participants completed the Prodromal Questionnaire-Brief Child Version (PQ-BC), a 21-item questionnaire previously validated for use with school-age children using the ABCD Study[®] sample (5,29), which asks about the occurrence of PLEs (e.g., unusual, thought content, perceptual abnormalities) in the past month. All PQ-BC questions were read to participants by research assistants. The dimensional total score was used to measure PLEs. Total scores index the full dimension of PLEs, including more normative PLEs, although individuals endorsing at least one PLE

generally show greater impairment than those endorsing no PLEs (Supplemental Table 1). To address the large number of 0 PLE values (44.6% of the sample) and obtain a more clinically-relevant PLE metric, we also formed 3 groups based upon PLE endorsement: *Group 1* - reporting 0 PLEs (n=2,076), *Group 2* - reporting 1 PLE but no significant distress associated with PLEs (n=1,601), *Group 3* - reporting 1 PLE with significant distress (i.e., rating a PLE 3 on a five-point scale of distress; n=972; Table 1).

Proximal EDU, Birth-weight, and Cross-Disorder PGS Behaviors.—Total cognition composite scores assessed using the National Institutes of Health Toolbox Cognitive Battery (30), caregiver-reported child birthweight (Supplement for birth complications and gestational age), and a general psychopathology factor (31) created using the Child Behavior Checklist (32), served as proximal behavioral measures of EDU, birthweight, and cross-disorder PGS, respectively.

Brain Structure.—T1- and T2-weighted structural scans (1mm isotropic) were acquired using 3T scanners (either Siemens, General Electric, or Phillips) with 32-channel head coils (Supplement). The following structural MRI metrics were examined: *global*: intracranial, total cortical, and total subcortical volume; total surface area; total cortical thickness; *regional*: 34 Desikan cortical regions for surface area, thickness, and volume, as well as 23 Freesurfer segmentation subcortical volume regions (33).

Polygenic Scores.—Summary statistics from the most well-powered “discovery” GWASs of schizophrenia (N=69,369 cases + 236,642 controls) (34), cross-disorder (N=232,964 cases [anorexia nervosa, attention-deficit/hyperactivity disorder, autism spectrum disorder, bipolar disorder, major depressive disorder, obsessive-compulsive disorder, schizophrenia, and Tourette’s syndrome] + 494,162 controls) (16), PLEs (N=127,966) (6), EDU (N=766,345; Supplement for executive functioning PGS; results generally consistent) (35), birthweight (N=321,223) (36), and inflammation (N=469,856; <http://biobank.ctsu.ox.ac.uk/crystal/field.cgi?id=30710>), were used to generate PGS (n=6) within the ABCD Study® dataset. Summary statistics from a schizophrenia GWAS study with individuals of African ancestries were used for our exploratory analyses (Supplement) (37).¹ PGSs were generated using polygenic risk scores-continuous shrinkage (PRS-CS) (38), which uses a Bayesian regression framework to include all SNPs in PGS calculation by placing a continuous shrinkage prior on SNP effect sizes; simulation studies show that PRS-CS outperforms other PGS methods (38). Analyses using traditional p-value clumping and thresholding produced results consistent with PRS-CS (Supplemental Tables 2-3).

Statistical Analyses

Continuous predictor and outcome variables were Winsorized to ± 3 SD to minimize the influence of extreme values. Analyses nested data with random intercepts for site (n=22) and family (n=3,874; siblings n=616). The following covariates were included in all analyses: age, sex (Supplemental Table 4 for results stratified by sex), genotyping batch,

¹As we are unaware of any other GWAS of African ancestry for the GWAS phenotypes investigated here, no other PGS were generated for this sample. Due to the relatively small sample of other ancestries in the ABCD dataset, we did not explore PGS in any other ancestries.

and the first ten ancestrally-informative principal components (described in the Supplement; Supplemental Table 5 for associations with PLEs). We used ComBat harmonization (<https://github.com/ncullen93/neuroCombat>), with age and sex added as biological covariates to the design matrix, to estimate and remove scanner model effects from MRI measures. Further, intracranial volume was included as an additional covariate in regional brain structure analyses. Financial adversity and highest parental/caregiver education were included as additional covariates in supplemental analyses (Supplement), as these variables are proxies of socioeconomic status (39) and therefore important predictors in risk for psychopathology and cognitive functioning.

First, we used hierarchical linear models (HLMs) to estimate associations between schizophrenia, PLE, cross-disorder, EDU, birth-weight, and inflammation PGS and each PLE metric (i.e., dimensional total score and binary significantly distressing PLEs) as outcomes. Benjamini-Hochberg False Discovery Rate (FDR) correction was used to account for the six PGSs tested.² For cross-disorder, birth-weight, and EDU, *post-hoc* HLMs examined models including measured behaviors most proximal to the PGS, to examine the extent to which available measured outcomes proximal to PGSs accounted for these associations (e.g., the extent to which cognition accounted for the association between EDU PGS and PLEs).

Second, we estimated associations between reported PLEs and MRI-derived brain structure phenotypes. FDR was used to adjust for multiple testing of 5 global MRI metrics (e.g., total cortical thickness), as well as 102 regional metrics (i.e., 34 each for bilateral [i.e., averaged across hemispheres] cortical thickness, cortical surface area, and cortical volume), and 23 tests for bilateral subcortical volumes (e.g., hippocampal volume), for a total of 130 FDR corrections for each PLE metric. Any significant regional association with PLEs was followed up with *post hoc* testing for lateral (i.e., right, left) associations.

Third, we estimated associations between PGS and brain structure phenotypes associated with reported PLEs. Subsequently, we examined whether any brain structure and PGS proximal behavioral phenotypes (e.g., cognition) indirectly linked PGS to reported PLEs using a series of mediation analyses.

Default settings were used to conduct HLMs for total PLEs (lmer) and hierarchical logistic regressions for significantly distressing PLEs (glmer) using the lme4 package (34). MuMIn was used to calculate pseudo-R-squared, with results comparing pseudo-R² for models with and without the predictor of interest (e.g., PRS score) converted to a percentage to create a % R². The lavaan package (40) was used to conduct mediation analyses, with models incorporating clustering and bootstrapping commands.

²Given the strong correlation ($r=.66$) between PLE total score and PLE significant distress group, we did not adjust for multiple testing for these 2 phenotypes.

Results

Polygenic Scores and Psychotic-like Experiences

In our sample ($n=4,650$), 55.33% ($n=2,573$) of children endorsed at least one reported PLE and 20.90% ($n=972$) endorsed at least one significantly distressing PLE. *Total PLEs* were associated with higher cross-disorder and lower EDU PGS (all $|\beta|>0.045$, all $p<0.002$, $p_{FDR}<0.006$; % $R^2>0.20\%$), but not schizophrenia, PLE, or birthweight PGS (all $|\beta| \leq 0.023$, all $p>0.11$); After correction for multiple testing, there was a trend for a positive association between *Total PLEs* and inflammation PGS ($\beta=0.031$, $p=.03$, $p_{FDR}=0.06$, % $R^2=0.092\%$; Table 2; Figure 1a).

Endorsement of *Significantly Distressing PLEs* was associated with higher schizophrenia, PLE, and cross-disorder PGS as well as lower EDU PGS (all $|\beta|>0.032$, all $p \leq 0.02$, $p_{FDR}<0.03$; % $R^2 \leq 0.12\%$; Table 2; Figure 1a), but not birthweight or inflammation PGS (all $|\beta| \leq 0.018$, all $p>0.22$). All associations remained similar when accounting for financial adversity and parental/caregiver education (Supplemental Table 6), or when computing PGS using a traditional clustering and thresholding approach (Supplemental Tables 2-3; Supplement for executive functioning PGS results).

Comparing PLE Groups (i.e., no PLEs, PLEs without significant distress, significantly distressing PLEs) generally revealed a pattern of results suggesting a gradient of severity (Supplement; Table 1); those reporting significantly distressing PLEs showed the greatest divergence from those without PLEs on PGSs, brain structure, and behavior, while those reporting PLEs not associated with significant distress were intermediary between these two groups (Supplemental Table 1).

Consideration of Proximal PGS Behaviors

Measured cognition was negatively associated with *Total PLEs* and endorsement of *Significantly Distressing PLEs*, and positively associated with EDU PGS ($\beta = -0.131$, all $p<2.00 \times 10^{-16}$, % $R^2 \leq 1.55\%$; Supplemental Table 7; Supplemental Figures 2-3). Cognitive performance accounted for 32.3-32.8% of the association between EDU PGS and PLE metrics (Supplemental Table 8), though associations between EDU PGS and reported PLE metrics remained, even when considering cognition ($|\beta|>0.047$, $p<0.002$; % $R^2 \leq 0.58\%$; Supplemental Table 9). The general psychopathology factor was positively associated with *Total PLEs* and endorsement of *Significantly Distressing PLEs*, as well as cross-disorder PGS (all $\beta \leq 0.055$, all $p \leq 0.002$, % $R^2>0.29\%$; Supplemental Table 7; Supplemental Figures 2-3). General psychopathology accounted for 11.3-17.8% of the association between cross-disorder PGS and reported PLEs, though associations between cross-disorder PGS with reported PLEs remained, even when considering general psychopathology (*Total*: $\beta=0.034$, $p=0.048$, $R^2=0.24\%$; *Significantly Distressing PLEs*: $\beta=0.056$, $p=0.002$, % $R^2=0.72\%$; Supplemental Table 9). Although PLEs were not significantly associated with birth-weight PGS (Table 1), reported birthweight was associated with PLEs and birth-weight PGS ($|\beta| \leq 0.047$, all $p \leq .003$, % $R^2>0.25\%$; Supplemental Table 7; Supplemental Figure 2).

Psychotic-like Experiences and Brain Structure

Global Metrics.—Greater reported PLEs (both *Total* and endorsement of *Significantly Distressing PLEs*) were associated with lower intracranial volume, total cortical volume, total subcortical volume, and total surface area (all $|\beta| > 0.058$, all $p < 3.82 \times 10^{-4}$, all $p_{FDR} < 1.24 \times 10^{-2}$; all % $R^2 > 0.29\%$; Table 2; Figure 1b), but not cortical thickness (all $|\beta| < 0.026$, all $p > 0.08$). **Regional Metrics.** When examining individual structural MRI regions for volume, surface area, and thickness, greater *Total PLEs* were associated with lower bilateral ($p_{FDR} = 0.04$) and right inferior parietal cortical thickness (both $|\beta| > 0.047$, both $p < 1.76 \times 10^{-3}$, both % $R^2 > 0.21\%$; Table 2; Figure 1b). No other individual structural MRI regions passed FDR correction (see Supplemental Tables 10-13).

Mediation Analyses

Educational Attainment Polygenic Scores.—EDU PGS was positively associated with all brain structure phenotypes linked to reported PLEs (i.e., intracranial volume, total cortical volume, total subcortical volume, total cortical surface area; all $\beta > 0.069$, all $p < 4.28 \times 10^{-7}$; Table 3), except inferior parietal cortical thickness (all $|\beta| < 0.027$, all $p > 0.08$). A series of individual mediational models examined whether each brain structure phenotype associated with both EDU PGS and PLEs (i.e., intracranial volume, total cortical volume, total subcortical volume, total cortical surface area) indirectly linked EDU PGS to *Total PLEs* alongside cognitive performance in parallel. There was evidence consistent with all volume metrics partially mediating the association between EDU PGS and *Total PLEs* (all indirect effect [path $a*b$] bias-corrected 95% confidence intervals [CI] within -0.012 to -0.001 ; proportion mediated: 3.33%-8.79%; Figure 2a-c). Specifically, evidence was consistent with lower educational attainment PGS being associated with reduced volumes, which were, in turn, associated with higher PLEs. There was no evidence consistent with total surface area indirectly linking EDU PGS to *Total PLEs* (indirect effect 95% CI: -0.004 to 0.002).

There was also evidence consistent with cognition uniquely partially mediating the association between EDU PGS and *Total PLEs* in each model (all within 95% CI: -0.038 to -0.020 ; proportion mediated: 28.57-32.22%; Figure 2). Specifically, lower cognitive performance indirectly linked lower EDU PGS to greater PLEs.

For all models, similar results were found when endorsement of *Significantly Distressing PLEs* was the outcome instead of *Total PLEs* (Supplement; Supplemental Figure 4). Mediation models remained consistent when accounting for financial adversity (Supplement). A single parallel mediation model with all neural metrics entered as simultaneous parallel mediators is reported in the Supplement.

Other Polygenic Scores.—Schizophrenia, PLE, birthweight, and inflammation PGS were not associated with global or regional brain structure measures associated with reported PLEs (Supplemental Table 14). Although total surface area was associated with cross-disorder PGS ($\beta = 0.042$, $p = 0.001$, % $R^2 = 0.14$; Table 3), there was not strong evidence consistent with total surface area mediating associations between cross-disorder PGS and reported PLEs (95% CI: -0.005 to 0.000 ; Supplement).

Exploratory Analyses: African-American Ancestral subsample.

There was no association between schizophrenia PGS and PLEs among individuals with African ancestry (Supplement).

Discussion

Here, we show that psychotic-like experiences (PLEs) in middle childhood ($n=4,650$) are associated with GWAS-derived polygenic scores ($\% R^2s=0.120-0.660\%$) as well as putative intermediary neural and behavioral phenotypes that may partially underlie these associations (all $\% R^2s=0.21-2.85\%$). Genomic liability for broad-spectrum psychopathology (cross-disorder PGS) and educational attainment (EDU PGS) were associated with both PLEs measures; however, schizophrenia and late-life PLE PGS were only significantly associated with the presence of distressing PLEs. Consistent with these findings, group contrasts revealed that SCZ PGS among those experiencing significantly distressing PLEs was significantly higher than those reporting PLEs without significant distress (Supplemental Table 1). One possible explanation is that PLEs may portend broad psychopathology vulnerability (1), while significantly distressing PLEs may more specifically foreshadow psychosis risk (41). Finally, reported PLEs were associated with lower global (e.g., intracranial volume) and regional (i.e., inferior parietal thickness was associated with broad PLEs) brain structure metrics (Figure 1b), with evidence that lower volume (i.e., intracranial volume, total cortical and total subcortical) may indirectly link EDU PGS to reported PLEs alongside cognition (Figure 2). Collectively, these results show that polygenic scores derived from adult GWASs can generalize to indices of risk among children and provide incremental predictive usefulness beyond measured proximal phenotypes.

Polygenic Propensity for Education Attainment

Polygenic scores for lower educational attainment was the most robust PGS predictor of reported PLEs.³ These findings suggest that prior reports linking EDU PGS to severe psychosis among clinical patients (42) may generalize to earlier markers of psychosis spectrum symptoms during middle childhood. That cognition accounted for a large portion (28.57-32.22% of variance) of the association between EDU PGS and reported PLEs aligns with evidence that premorbid cognition prospectively predicts PLEs and psychopathology, including schizophrenia (21), and suggests that such vulnerability may be partially genomic in origin. However, other work has failed to find a strong genetic correlation between later-life PLEs and intelligence (6); it is possible that genomic associations between PLEs and cognition may differ across the life course and be more correlated during childhood with divergence in later life (e.g., PLEs related to cognitive decline and/or dementia (43)). Notably, executive functioning PGS generally showed similar patterns compared with models including EDU PGS, although executive functioning PGS showed weaker effects with neural metrics (Supplement). In addition to PGS, it is also likely that there were a number of additional influences on cognitive performance that were not included in this study, including additional pathophysiological factors (e.g., functional connectivity) and environmental influences (e.g., exposure to toxins).

³Additionally, associations between EDU PGS and PLEs remained even when accounting for anhedonia (Supplement).

Supportive of neurodevelopmental models of psychosis spectrum disorders positing that brain differences underlie cognition-related vulnerability for schizophrenia, we found evidence consistent with brain volume accounting for a portion of the association between educational attainment PGS and reported PLEs (3.22-8.79%; Supplemental Table 7; Figure 2). However, given that neural metrics are likely less proximal to EDU PGS than cognition, it is unsurprising that associations for cognition were larger than associations for brain structure (44). Associations between reported PLEs and global volume reductions are consistent with previous research examining volumetric alterations (22). That lower global volume may indirectly link EDU PGS and reported PLEs is consistent with the notion that genomic liability for lower cognitive functioning may be associated with altered neural maturational processes, which may contribute to the development or maintenance of psychosis spectrum symptoms (45).

General Psychopathology, Schizophrenia, and PLE Polygenic Risk

Genomic liability to general psychopathology (i.e., cross-disorder PGS) was associated with broadly defined (i.e., total) and severe (i.e., significantly distressing) PLEs, while polygenic scores for schizophrenia and PLEs were only associated with severe PLEs. These findings align with evidence that polygenic and phenotypic psychopathology associations may be non-specific in middle childhood and potentially become increasingly specific with increased severity (e.g., with more clinically significant PLEs) and/or maturation (e.g., in adolescence) (46).

We did not find strong evidence of associations between broadly defined PLEs and polygenic risk for later life PLEs, although there were associations between later life PLEs PGS and more severe PLEs. Alongside evidence of similar effect size estimates across both PLE measures within PGS, it is plausible that with better powered discovery GWASs and target samples, PLEs defined broadly and with greater severity will show similar relationships.

Genomic Propensity for Birth Weight and Inflammation

Reported birthweight was negatively associated with PLEs during middle childhood, consistent with research in young adults (47), and positively associated with birthweight PGS; however, birthweight PGS were not associated with reported PLEs. Together, this raises the intriguing possibility that environmental factors associated with lower birthweight, as opposed to genetic predisposition to low birthweight, may underlie the association between lower birthweight and PLEs (48).

There were nominally significant associations between inflammation PGS and total reported PLEs, although this was trend-level after adjusting for multiple testing. Inflammation has been widely associated with psychopathology, with emerging evidence suggesting that inflammation-driven variation in neurodevelopment (i.e., neural pruning during puberty) may play a prominent role in the etiology of schizophrenia (18). It is possible that associations between inflammation PGS and the expression of PLEs may increase following periods of heightened neural development (e.g., adolescence) and/or in interaction with other factors (e.g., infection) that will require larger samples to address.

Limitations

It is important to consider limitations of this study while interpreting these findings. First, the generalizability of these findings is limited, as we restricted most analyses to individuals of European ancestries due to the sample compositions of the discovery GWASs and evidence that polygenic risk does not translate across ancestries (49). Due to this exclusion and exclusion for quality control reasons, a number of participants (n=7,225) were not included in analyses. Excluded participants showed higher scores than included participants on a number of measures (Supplemental Table 15), and therefore, if anything, would have contributed to the clinical severity of this sample. Second, and consistent with expectations from the ABCD Study[®] (28), which uses a heterogeneous sample, and from prior PGS studies (44), the effects reported are generally small (for associations with reported PLEs, $|\beta| < 0.09$, % $R^2 < 0.7\%$) (50). Third, prevalence rates of PLEs (e.g., 55.3% for total PLEs) were higher than some previous estimates (i.e., ~10%) (1), although consistent with others (51). This high rate of endorsement may reflect over-endorsement or transient phenomena related to assessing PLEs in middle childhood that may have diluted the magnitude of associations found in the current study. Total PLEs are unlikely useful as a clinical indicator, although may be useful as a measure of the dimension of PLEs, including developmentally normative experiences and trait-relevant phenomena (e.g., oddness). Regardless, the PQ-BC, including Total scores, has been validated for use with children as young as age 9 (4), and there is evidence that only people with high trait levels of PLEs typically endorse PQ-BC items (52). Fourth, while the non-experimental and cross-sectional nature of our data does not preclude conducting mediation analyses (53), they should not be interpreted by themselves to imply causation. However, these analyses provide some empirical evidence consistent with putative gene-brain-behavior mechanisms underlying childhood PLE risk. Fifth, it is important to note that the GWASs have differential power to detect effects (e.g., the EDU GWAS was based on 766,345 individuals, whereas the PLEs GWAS was based on 127,966 individuals). These differences in power present challenges for interpreting across different PGSs. As “discovery” GWASs continue to grow, it will be critical to acquire additional GWASs datasets across development, including in childhood, to examine genetic correlations for the same phenotype across ages as well as differential associations with psychopathology and structural neural metrics. Future studies using the ABCD Study dataset should further examine the validity of the PGS scores including validity in other populations, and also examine potential clinically applicable thresholds (54). Finally, future research should begin to examine associations between PLEs, PGS, and other factors previously found to be associated with PLEs in the ABCD Study[®] (e.g., environmental toxins) (23).

Conclusions

Broadly, GWAS-based polygenic scores for psychopathology and EDU generated from adult samples are associated with indices of PLEs during middle childhood. Polygenic propensity to EDU was the most robust predictor of reported PLEs, and there was evidence that these associations may be partially mediated by cognitive performance and brain structure. Polygenic score associations mirrored phenotypic evidence that broadly defined PLEs are associated with broad-spectrum psychopathology risk while more severe PLEs may index psychosis liability. Taken together, this study documents that GWAS-derived polygenic

scores index psychopathology and psychosis vulnerability in children. PGSs may support the identification of putative intermediate biological and behavioral mechanisms through which genomic risk for psychopathology emerges. Although there are a number of important ethical considerations with regards to using PGS for prediction purposes, including concerns about early identification efforts and the potential for exacerbating health disparities that must be addressed prior to clinical use (55), more severe PLEs may be important early indicators of psychosis liability. Therefore, future research should begin to examine whether severe PLEs can be utilized as markers for further assessment and potential intervention.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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The ABCD data repository grows and changes over time. The ABCD data used in this report came from DOI [10.15154/1460410](https://doi.org/10.15154/1460410).

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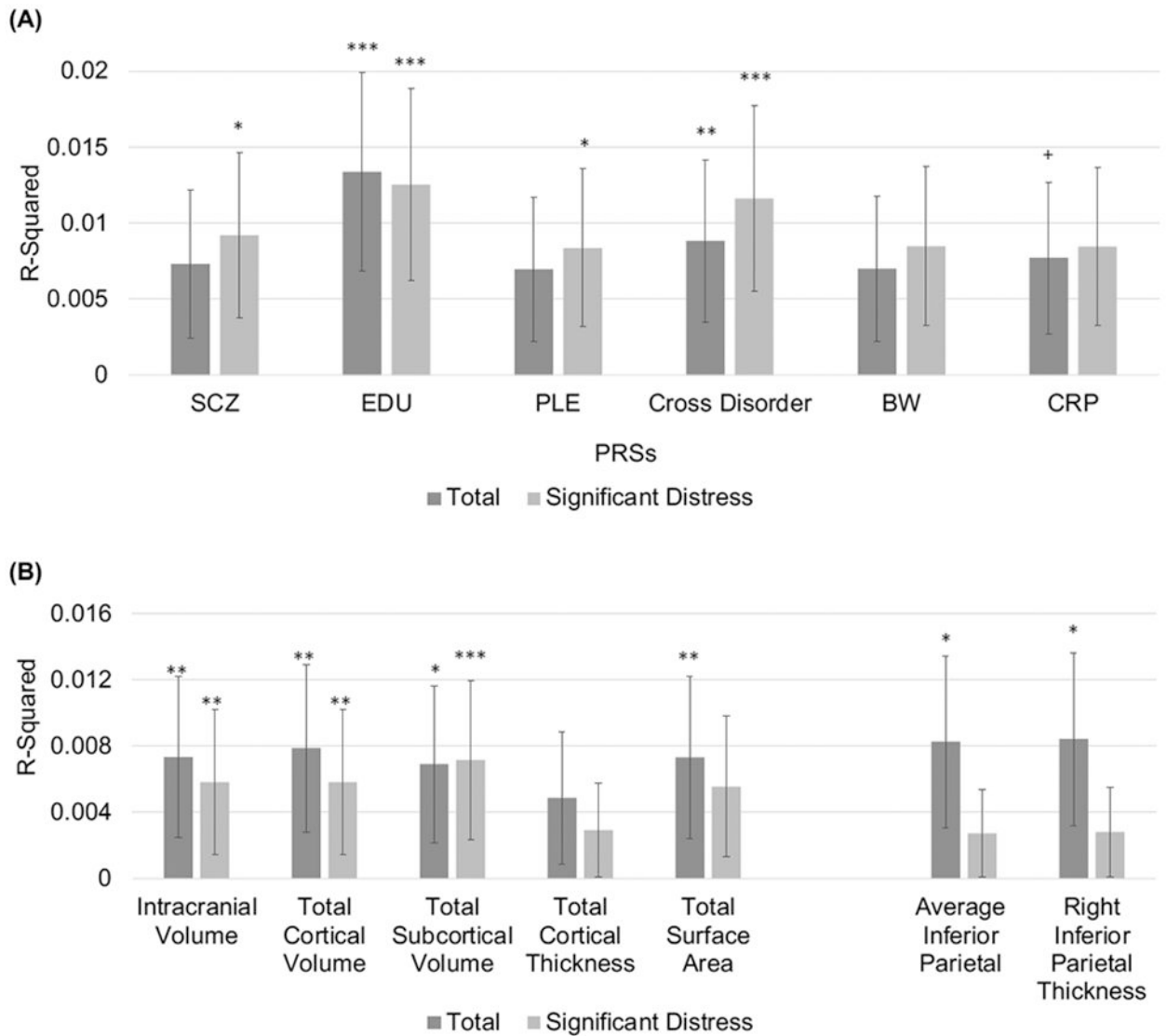


Figure 1.

Proportion of variance explained (R-squared) by (A) each of the different PGS scores (SCZ=Schizophrenia; EDU =Educational Attainment; PLE=Psychotic-like experiences; CROSS=Cross Disorder; BW= Birth weight; CRP = Inflammation) for both Total and Significantly Distressing PLEs, and (B) each of the significant global (intracranial volume, total cortical volume, total subcortical volume, surface area) and regional (inferior parietal thickness) MRI metrics for both Total and Significantly Distressing PLEs. Error bars represent the 95% confidence interval, * $p_{FDR}<0.05$, ** $p_{FDR}<0.01$, *** $p_{FDR}<0.001$.

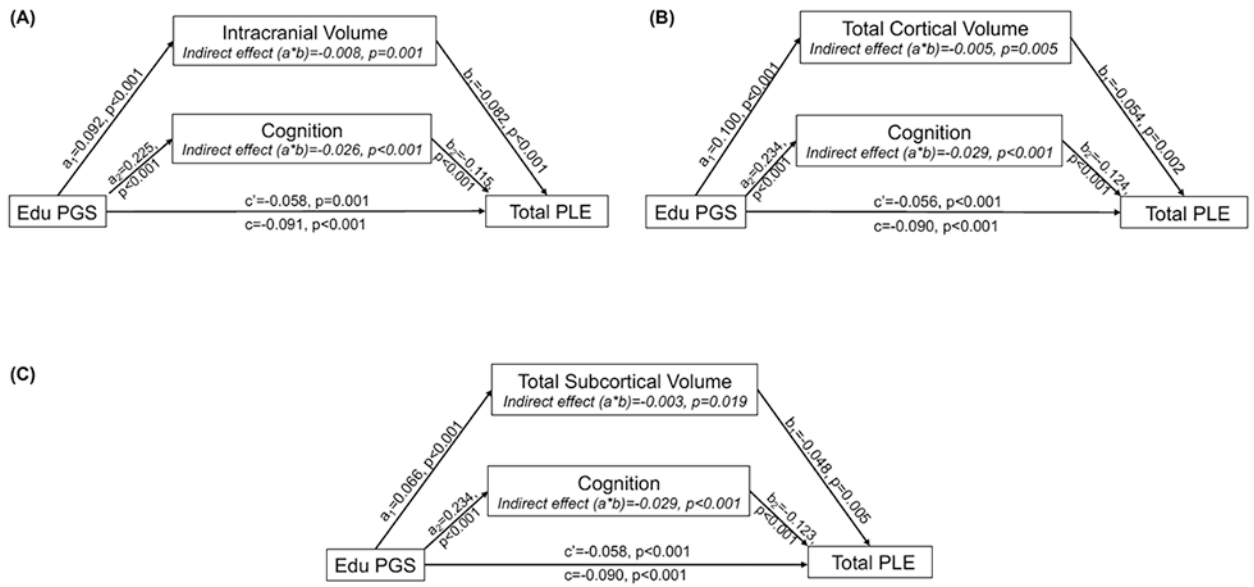


Figure 2.

Depiction of a series of parallel mediation models, examining evidence for each neural metric ((A) intracranial, (B) cortical volume, and (C) subcortical volume) indirectly linking Educational Attainment PGS (EDU PGS) and total psychotic-like experiences (Total PLEs) alongside cognitive performance in parallel. Covariates (i.e., age, sex, genotyping batch, and the first ten ancestrally-informative PCs) were included in all models. Each parallel mediation model depiction includes unstandardized regression coefficients, showing the association between EDU PGS, each neural metric and cognition, and Total PLEs.

Table 1.

Sample Characteristics^a

Variable (n)	No PLEs (n=2076)	1 PLE but no Significant Distress (n=1,601)	Endorsed 1 Significantly Distressing PLEs (n=972)	Total Sample (N=4650)
Age in years (4650)	9,960±0.627	9,933±0.627	9,882±0.613	9,934±0.625
Sex, % female (4650)	999 (48.1%)	723 (45.2%)	454 (46.7%)	2176 (46.8%)
Scanner Type (4532)				
Siemens, %	1431 (68.9%)	977 (61.0%)	576 (59.3%)	2984 (64.2%)
Philips, %	241 (11.6%)	240 (15.0%)	125 (12.9%)	606 (13.0%)
General Electric, %	354 (17.1%)	340 (21.2%)	247 (25.4%)	942 (20.3%)
PLEs (4649)				
Total PLEs	0±0	2,548±2.101	6,180±3.686	2,170±3.142
Significantly Distressing PLEs	0±0	0±0	2,158±1.470	0.451±1.105
PGS (4650)				
Schizophrenia	4.867x10 ⁻⁷ ±1.768x10 ⁻⁷	4.800x10 ⁻⁷ ±1.753x10 ⁻⁷	5.033x10 ⁻⁷ ±1.756x10 ⁻⁷	4.879x10 ⁻⁷ ±1.762x10 ⁻⁷
EDU	-1.387x10 ⁻⁷ ±9.044x10 ⁻⁸	-1.553x10 ⁻⁷ ±9.349x10 ⁻⁸	-1.638x10 ⁻⁷ ±9.203x10 ⁻⁸	-1.497x10 ⁻⁷ ±9.238x10 ⁻⁸
PLE	-4.498x10 ⁻⁹ ±6.846x10 ⁻⁸	-4.711x10 ⁻⁹ ±6.717x10 ⁻⁸	3.910x10 ⁻¹⁰ ±6.879x10 ⁻⁸	-3.553x10 ⁻⁹ ±6.810x10 ⁻⁸
Cross-disorder	3.030x10 ⁻⁷ ±8.376x10 ⁻⁸	3.029x10 ⁻⁷ ±8.658x10 ⁻⁸	3.180x10 ⁻⁷ ±8.447x10 ⁻⁸	3.061x10 ⁻⁷ ±8.508x10 ⁻⁸
Birth-weight	-3.611x10 ⁻⁹ ±8.670x10 ⁻⁸	-3.612x10 ⁻⁹ ±8.536x10 ⁻⁸	-7.710x10 ⁻⁹ ±8.250x10 ⁻⁸	-4.478x10 ⁻⁹ ±8.537x10 ⁻⁸
Inflammation	2.355x10 ⁻⁸ ±1.135x10 ⁻⁷	2.722x10 ⁻⁸ ±1.113x10 ⁻⁷	3.132x10 ⁻⁸ ±1.127x10 ⁻⁷	2.641x10 ⁻⁸ ±1.126x10 ⁻⁷
Global Brain Metrics				
Intracranial Volume (3914)	1.550x10 ⁶ ±1.270x10 ⁵	1.550x10 ⁶ ±1.313x10 ⁵	1.534x10 ⁶ ±1.344x10 ⁵	1.547x10 ⁶ ±1.302x10 ⁵
Total Cortical Volume (4520)	6.147x10 ⁵ ±5.143x10 ⁴	6.122x10 ⁵ ±5.218x10 ⁴	6.070x10 ⁵ ±5.328x10 ⁴	6.122x10 ⁵ ±5.216x10 ⁴
Total Subcortical Volume (4520)	6.140x10 ⁴ ±4.550x10 ³	6.134x10 ⁴ ±4.612x10 ³	6.070x10 ⁴ ±4.824x10 ³	6.123x10 ⁴ ±4.637x10 ³
Total Cortical Thickness (4520)	2.801±0.089	2.798±0.092	2.794±0.089	2.798±0.090
Total Surface Area (4520)	1.907x10 ⁵ ±1.662x10 ⁴	1.902x10 ⁵ ±1.648x10 ⁴	1.885x10 ⁵ ±1.655x10 ⁴	1.901x10 ⁵ ±1.658x10 ⁴
Proximal PGS Behaviors				
Cognitive Scores (4581)	90.076±7.659	88.412±7.848	86.774±7.968	88.814±7.893

Variable (n)	No PLEs (n=2076)	1 PLE but no Significant Distress (n=1,601)	Endorsed 1 Significantly Distressing PLEs (n=972)	Total Sample (N=4650)
General Psychopathology Scores (3120)	-0.064±0.868	0.056±0.867	0.292±0.938	0.050±0.892
Birthweight in ounces (4546)	114.275±23.502	113.244±22.946	111.077±24.362	113.258±23.518

Abbreviations: PLEs=psychotic-like experiences; PGS=polygenic scores; EDU=Educational Attainment.

^aMeans ± standard deviations, unless otherwise noted.

Table 2.

Associations between PLEs and both PGS and Structural Neural Metrics^a

Metric	Total PLEs						Significant Distress Group					
	b [95% CI]	β	t	p	p FDR	% R ²	OR [95% CI]	β	Z	p	p FDR	% R ²
PGS												
Schizophrenia	4.17x10 ⁵ [-9.35x10 ⁴ ,9.27x10 ⁵]	0.023	1.598	.11	.17	0.052%	1.130 [1.050,1.220]	0.043	3.166	.002	.01	0.216%
Educational Attainment	-2.76x10 ⁶ [3.74x10 ⁶ ,-1.79x10 ⁶]	-0.081	-5.561	2.85x10 ⁻⁸	1.71x10 ⁻⁷	0.660%	0.825 [0.766,0.890]	-0.072	-4.988	6.09x10 ⁻⁷	3.65x10 ⁻⁶	0.587%
PLEs	6.21x10 ⁵ [-6.93x10 ⁵ ,1.94x10 ⁶]	0.013	0.925	.36	.36	0.019%	1.090 [1.010,1.170]	0.032	2.259	.02	.03	0.120%
Cross-disorder	1.67x10 ⁶ [6.18x10 ⁵ ,2.73x10 ⁶]	0.045	3.099	.002	.006	0.202%	1.190 [1.110,1.290]	0.066	4.600	4.23x10 ⁻⁶	1.27x10 ⁻⁵	0.505%
Birth weight	-5.17x10 ⁵ [-1.57x10 ⁶ ,5.40x10 ⁵]	-0.014	-0.96	.34	.36	0.021%	0.954 [0.885,1.030]	-0.016	-1.23	.22	.22	0.037%
Inflammation	8.62x10 ⁵ [6.54x10 ⁴ ,1.66x10 ⁶]	0.031	2.12	.03	.06	0.092%	1.050 [0.973,1.130]	0.018	1.228	.22	.22	0.036%
Neural Metrics												
Global Metrics												
Intracranial Volume	-1.56x10 ⁻⁶ [-2.36x10 ⁻⁶ ,-7.59x10 ⁻⁷]	-0.065	-3.815	1.38x10 ⁻⁴	5.98 x10 ⁻³	0.332%	0.841 [0.770,0.919]	-0.063	-3.851	1.18E-04	5.11x10 ⁻³	0.371%
Total Cortical Volume	-4.12 x10 ⁻⁶ [-6.06x10 ⁻⁶ ,-2.18x10 ⁻⁶]	-0.068	-4.169	3.12x10 ⁻⁵	4.06 x10 ⁻³	0.388%	0.845 [0.776,0.920]	-0.062	-3.898	9.72E-05	5.11x10 ⁻³	0.370%
Total Subcortical Volume	-3.93x10 ⁻⁵ [-6.09x10 ⁻⁵ ,-1.76x10 ⁻⁵]	-0.058	-3.555	3.82x10 ⁻⁴	1.24 x10 ⁻²	0.291%	0.823 [0.756,0.895]	-0.071	-4.519	6.21E-06	8.07x10 ⁻⁴	0.503%
Total Cortical Thickness	-0.908 [-1.918,0.105]	-0.026	-1.760	.08	.42	0.085%	0.936 [0.868,1.009]	-0.024	-1.723	.08	.50	0.079%
Total Surface Area	-1.22x10 ⁻⁵ [-1.85x10 ⁻⁵ ,-5.97x10 ⁻⁶]	-0.065	-3.824	1.33x10 ⁻⁴	5.98 x10 ⁻³	0.330%	0.847 [0.777,0.924]	-0.061	-3.727	1.94E-04	6.31x10 ⁻³	0.343%
Regional Metrics												
Average Inferior Parietal Thickness	-1.270 [-2.040, -0.492]	-0.048	-3.206	1.35x10 ⁻³	.04	0.226%	0.926 [0.853,1.006]	-0.027	-1.810	.07	.50	0.078%
Right Inferior Parietal Thickness	-1.150 [-1.880, -0.432]	-0.047	-3.131	1.76x10 ⁻³		0.214%	0.927 [0.854,1.007]	-0.027	-1.798	.07		0.077%

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Abbreviations: PLEs=psychotic-like experiences; PGS=polygenic scores; b=unstandardized beta coefficient; 95% CI=95% confidence interval; β =standardized beta coefficient; |t|-test test statistic; p =p-value; p FDR=False Discovery Rate-corrected p ; % R^2 =percentage change in marginal proportion of variance explained, calculated as the difference in pseudo- R^2 between the current model versus a model excluding the predictor of interest, converted to a percentage; OR=odds ratio; $Z=Z$ statistic.

p FDR<.05 models are in bold. Tests are two-tailed. All PGS (i.e., schizophrenia, educational attainment, cross-disorder, PLE) results remained consistent with accounting for financial adversity and parental education.

Table 3.

Associations between PGS and Structural Neural Metrics ^a

Metric	Educational Attainment PGS					Cross-disorder PGS				
	b [95% CI]	β	t	p	% R ²	b [95% CI]	β	t	p	% R ²
Global Structural Metrics										
Intracranial Volume	1.27x10 ¹¹ [9.10x10 ¹⁰ , 1.63x10 ¹¹]	0.090	6.908	5.61x10⁻¹²	0.819%	2.21x10 ¹⁰ [-1.69x10 ¹⁰ , 6.10x10 ¹⁰]	0.014	1.112	.27	0.010%
Total Cortical Volume	5.49x10 ¹⁰ [4.01x10 ¹⁰ , 6.99x10 ¹⁰]	0.097	7.219	6.14x10⁻¹³	0.949%	1.45x10 ¹⁰ [-1.68x10 ⁹ , 3.06x10 ¹⁰]	0.024	1.759	.08	0.040%
Total Subcortical Volume	3.45x10 ⁹ [2.12x10 ⁹ , 4.79x10 ⁹]	0.069	5.064	4.28 x10⁻⁷	0.460%	4.70x10 ⁸ [-9.79x10 ⁸ , 1.91x10 ⁹]	0.009	0.637	.52	0.000%
Total Cortical Thickness	-1.71x10 ³ [-3.05x10 ⁴ , 2.72 x10 ⁴]	-0.002	-0.116	.91	0.000%	-1.42x10 ⁴ [-4.54x10 ⁴ , 1.70x10 ⁴]	-0.013	-0.889	.37	0.017%
Total Surface Area	1.72x10 ¹⁰ [1.26x10 ¹⁰ , 2.19x10 ¹⁰]	0.096	7.343	2.48 x10⁻¹³	0.923%	8.15x10⁹ [3.16x10⁹, 1.31x10¹⁰]	0.042	3.207	.001	0.142%
Regional Metrics										
Average Inferior Parietal Thickness	-1.56x10 ⁴ [-5.36x10 ⁴ , 2.23x10 ⁴]	-0.012	-0.806	.42	0.013%	-5.73x10 ³ [-4.68x10 ⁴ , 3.54x10 ⁴]	-0.004	-0.273	.79	0.001%
Right Inferior Parietal Thickness	-3.68x10 ⁴ [-7.73x10 ⁴ , 3.73x10 ³]	-0.027	-1.776	.08	0.068%	-1.40x10 ⁴ [-5.79x10 ⁴ , 2.99x10 ⁴]	-0.009	-0.626	.53	0.008%

Abbreviations: PLEs=psychotic-like experiences; PGS=polygenic scores; b=unstandardized beta coefficient; 95% CI=95% confidence interval; β =standardized beta coefficient; t=t-test test statistic; p =p-value; % R²=percentage change in marginal proportion of variance explained, calculated as the difference in pseudo-R² between the current model versus a model excluding the predictor of interest, converted to a percentage; OR=odds ratio; Z=Z statistic.

^a pFDR<.05 models are in bold. Tests are two-tailed.