ARTICLE

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Shared polygenic risk for ADHD, executive dysfunction and other psychiatric disorders

Suhua Chang^{1,2}, Li Yang¹, Yufeng Wang¹ and Stephen V. Faraone³

Abstract

Many psychiatric disorders are associated with impaired executive functioning (EF). The associated EF component varies by psychiatric disorders, and this variation might be due to genetic liability. We explored the genetic association between five psychiatric disorders and EF in clinically-recruited attention deficit hyperactivity disorder (ADHD) children using polygenic risk score (PRS) methodology. Genome-wide association study (GWAS) summary data for ADHD, major depressive disorder (MDD), schizophrenia (SZ), bipolar disorder (BIP) and autism were used to calculate the PRSs. EF was evaluated by the Stroop test for inhibitory control, the trail-making test for cognitive flexibility, and the digital span test for working memory in a Chinese ADHD cohort (n = 1147). Exploratory factor analysis of the three measures identified one principal component for EF (EF-PC). Linear regression models were used to analyze the association between each PRS and the EF measures. The role of EF measures in mediating the effects of the PRSs on ADHD symptoms was also analyzed. The result showed the PRSs for MDD, ADHD and BIP were all significantly associated with the EF-PC. For each EF component, the association results were different for the PRSs of the five psychiatric disorders: the PRSs for ADHD and MDD were associated with inhibitory control (adjusted P = 0.0183 and 0.0313, respectively), the PRS for BIP was associated with working memory (adjusted P = 0.0416), and the PRS for SZ was associated with cognitive flexibility (adjusted P = 0.0335). All three EF measures were significantly correlated with ADHD symptoms. In mediation analyses, the ADHD and MDD PRSs, which were associated with inhibitory control, had significant indirect effects on ADHD symptoms through the mediation of inhibitory control. These findings indicate that the polygenic risks for several psychiatric disorders influence specific executive dysfunction in children with ADHD. The results helped to clarify the relationship between risk genes of each mental disorder and the intermediate cognitive domain, which may further help elucidate the risk genes and motivate efforts to develop EF measures as a diagnostic marker and future treatment target.

Introduction

Executive functioning (EF) is a high-order set of cognitive functions that regulate an individual's capacity to change and adjust his or her behaviors according to the shifting demands of complex environments¹. EF

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dysfunction is associated with many psychiatric disorders, such as attention deficit hyperactivity disorder (ADHD), major depressive disorder (MDD), bipolar disorder (BIP), autism (ASD), and schizophrenia $(SZ)^{2-5}$. EF encompasses multiple processes^{6,7}. Although most psychiatric disorders are associated with EF dysfunction, different neuropsychiatric disorders showed disorder-specific EF deficits^{8,9}; for example, the largest effect sizes were observed for response inhibition in individuals with ADHD¹⁰, for verbal and nonverbal memory in individuals with BIP¹¹, and for cognitive flexibility in individuals with SZ¹². These results suggest that specific EF components might be good markers for specific types of

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psychopathology. Understanding the role of separate EF components in psychiatric disorders is a critical step in gaining a better understanding of disease psychopathology. Currently, the specificity of these EF component deficits in different psychiatric disorders is not clear¹³.

EF is heritable. Similar EF performance has been observed in family members and twins. The heritability of EF was calculated to be approximately 20 to 40%¹⁴. The association between EF dysfunction and the psychiatric diseases may, in part, reflect pleiotropy, which is the overlap between the genetic liability of two or more traits, perhaps owing to shared causal pathways¹⁵. Thus, the genes that influence vulnerability to psychiatric disorders may also influence EF deficits.

It has been widely acknowledged that many psychiatric disorders share genetic etiology^{16–18}. Most of the studies investigated shared genetics among a variety of disorders. Only a few studies have discussed the relationship between genetic risk of disorders and intermediate phenotypes. Shared genetic risks have been reported between SZ^{19,20}, depression²¹, ADHD²² and decreased general cognitive ability¹⁷, autism and increased cognitive ability²³, while BIP had mixed results^{17,24}. Polygenic risk score (PRS) of ADHD has been associated with components of EF: working memory, vigilance arousal, but not inhibitory control, verbal-numerical reasoning $^{25-27}$. The research for the genetic association of other disorders with specific component was limited or inconsistent^{28,29}. The Brainstorm project identified several correlations between psychiatric disorders and intelligence, cognition and education¹⁷, but not specific EF components. Benca et al. examined the PRSs of five psychiatric disorders (i.e., autism, ADHD, BIP, MDD, and SZ) with three EF components (mainly involving the shifting component) in a small sample of twins, but failed to find any significant results³⁰.

The reported studies showed inconsistent results for the genetic association between polygenic risk score of psychiatric disorders with cognition. Most of the cognition data of previous studies used population-based samples, whose phenotype distribution may differ with clinical-based samples. Sample difference in ages also yielded inconsistent results. For example, polygenic risk score of ADHD was associated with cognition in children samples³¹ but not in adult sample²³. So, additional data on this topic is warranted to illuminate the genetic associations among psychiatric disorders and EF.

Based on the correlation between psychiatric disorder and executive function at phenotype level, our hypothesis (Fig. 1) is that EF is affected by the genetic risks for multiple psychiatric disorders and it is possible that the genetic risks for different disorders are associated with deficits in specific EF components. In the present study, we examined the genetic associations of five psychiatric disorders (ADHD, MDD, SZ, BIP, ASD) with three executive function components (inhibition, working memory, cognitive flexibility) using polygenic risk scores in a clinically-recruited sample of ADHD children sample. We sought to test our hypothesis and to explore the specific associations between these psychiatric disorders and each EF component. Furthermore, since inhibitory control is a well-replicated deficit of ADHD, we hypothesized that the PRS for inhibitory control would also affect ADHD symptoms and would mediate the effects of disorder-specific PRSs on ADHD symptoms.



	#subjects	Mean (SD)	Skewness (SD)	Kurtosis
Inhibitory control	963	28.72 (17.32)	0.804 (0.079)	3.683 (0.157)
Cognitive flexibility	913	129.98 (97.65)	1.564 (0.081)	3.452 (0.162)
Working memory	1145	4.32 (1.68)	0.632 (0.071)	0.698 (0.143)
ADHD symptoms	1134	11.79 (2.97)	0.176 (0.073)	-0.621 (0.149)

Table 1 Skewness and kurtosis of executive function measures and ADHD symptoms.

Methods

Participants

The participants who underwent EF tests included 1147 ADHD patients (954 boys, 83.2%) aged between 6 and 17 years [average (10.1 ± 2.6) years] who were recruited from the Child and Adolescent Psychiatric Outpatient Department of the Peking University Sixth Hospital. All participants met the DSM-IV diagnostic criteria for ADHD. The clinical diagnosis was first made by a senior child and adolescent psychiatrist based on the parent- and teacher-completed ADHD Rating Scale-IV and was then confirmed by a semi-structured interview with the parents and child, using the Chinese version of the Clinical Diagnostic Interview Scale^{32,33}. Individuals with major neurological disorders (e.g., epilepsy), SZ, pervasive developmental disorder or intellectual disability (IQ < 70) were excluded. Individuals with comorbidities were not necessarily excluded. In this sample, 349 (29.7%) children had oppositional defiant disorder (ODD), 28 (2.4%) had conduct disorder, 79 (6.7%) had a specific phobia, 18 (1.5%) had a social phobia, 19 (1.6%) had generalized anxiety disorder, 13 (1.1%) had MDD, 6 (0.5%) had BIP, and 178 (15.1%) had a tic disorder. IQ was assessed using the third edition of the Chinese version of the Wechsler Intelligence Scale for Children. Most subjects were drug naive. Only 31 (2.6%) patients had used methylphenidate, and 150 (12.8%) had used Chinese herbal medicine. For individuals who had been medicated, the drugs were washed out for at least 1 month before the patient was tested. All these samples were independent from the samples in our previous ADHD GWAS paper³⁴, neither overlapping individuals nor related siblings or cousins. The study was approved by the Institutional Review Board of the Peking University Sixth Hospital. Written informed consent was obtained from the parents of the ADHD probands.

Executive function tests

From the 1,147 subjects with EF measures, the numbers of ADHD patients for each EF test are shown in Table 1. The details of these tests are as follows.

Stroop test for inhibitory control

The child and adolescent psychiatrist monitored all the tests conducted on patients. The task included four sessions as described in³⁵. At the beginning, thirty stimuli were presented in a 10×3 matrix for three cards each sized 21×29.7 cm². In the first session, the research participants were asked to read the names of colors (red, green, yellow, and blue) printed in black ink. In the second session, they were asked to name the colored squares (red, green, yellow, and blue). In the third session, the participants were asked to read the color words printed in different colors. In the fourth session, they were asked to name the colors of the ink. The time required to complete each session was recorded. In this study, we used the word interference time, which equals the time required to complete session 4 minus that for session 2, to assess interference inhibition. Higher score indicates worse performance.

Trail-Making Test for cognitive flexibility

The test consisted of two sections (A and B), as described in³⁶. In section A, numbers from 1 to 25 were randomly scattered on the page, and the participant was asked to connect these numbers sequentially as quickly as possible. In section B, the participant was asked to connect numbers and letters alternately (i.e., 1->A->2->B->3->C, ... L->13). When the participant made an error, the investigator pointed out the error immediately before continuing the test. The time to complete section A, T_A , indicates motor speed and visuo-perceptual abilities, while the time to complete section B, T_B, is sensitive to working memory and cognitive flexibility. The time difference $T_B - T_A$, which was highly related to T_B (r = 0.94), minimizes visuo-perceptual and working memory demands, providing a relatively pure indicator of executive control. Therefore, the shifting time $= T_B - T_A$ was the main measure for assessing cognitive flexibility³⁷. Higher score indicates worse cognitive flexibility.

Digit span test for working memory

The digit span test is a component of the Chinese version of Wechsler Intelligence Scales for Children (C- WISC). In this study, we used the digits backward digit span test to assess the patient's working memory, as it measures the child's ability to manipulate verbal information while it is in temporary storage³⁸. In this test, the child listens to a sequence of numbers and repeats them in reverse order. The length of each sequence of numbers increases as the child responds correctly. Higher score indicates better working memory.

The phenotype distribution plot showed working memory was normally distributed, but inhibitory control and cognitive flexibility had some outliers. Considering excessive scores on cognitive measures may indicate the participant did not understand the task or something else happened during testing, using the method described in ref.³⁹, we used a loose threshold to remove the samples with inhibitory control or cognitive flexibility below $F_{\rm L}-4 \times (F_{\rm U}-F_{\rm L})$ or above $F_{\rm U}+4 \times (F_{\rm U}-F_{\rm L})$, in which $F_{\rm L}$ and F_{II} were the lower and upper fourths of the phenotype. We removed 1 and 2 outliers for inhibitory control and cognitive flexibility respectively. From the 1134 subjects, we also collected ADHD symptoms according to the Clinical Diagnostic Interview Scale⁴⁰. The phenotype distribution plot and related skewness and kurtosis after removing outliers is shown in Fig. S1 and Table 1. We compared the phenotype data with a small group of controls and the result showed ADHD patients had significantly worse performance in shifting time, inhibitory control and working memory than normal controls (Table S1).

For subjects with all three EF measures, we did exploratory factor analysis (EFA) using SPSS 26. EFA method is to identify the underlying relationships between measured variables and get a latent variable to denote common factor of the measured variables. The KMO and Bartlett's test was significant (P < 0.001), which denoted the three EF measures had high correlation and were suitable for EFA. The scree plot showed the first principal component (EF-PC) had eigenvalue >1. Detailed parameters for the EFA model was shown in Fig. S2. The total variance explained by the first component is 56.523%. We used EF-PC to denote the main component among these three EF measures. To test the correlations among the EF measures (including EF-PC) and correlations with ADHD symptoms, two-tailed Pearson correlation was conducted by using SPSS 26.

Genotyping and imputation

The ADHD patients who underwent executive function tests were genotyped using the InfiniumPsychArray-24 array by CapitalBio Ltd. (Beijing). First, we removed individuals with per-individual autosomal heterozygosity >5 S.D. larger than the mean, individuals without age or sex information, individuals with a per-individual call rate <95% and individuals with the lower call rate in a pair of

individuals with proportion identity by descent (IBD) PI_HAT > 0.185. Next, we removed the SNPs with a per-SNP call rate <95%, a Hardy–Weinberg equilibrium test result with P < 0.001, or a minor allele frequency<1%. After quality control was performed, 1147 samples with 328,390 SNPs remained.

Principal component analyses (PCA) was conducted using the SNPs with low linkage disequilibrium (LD, MAF > 0.35 and r^2 < 0.05 for each pair of SNPs) that were outside the 5 long-range LD regions⁴¹ using the EIGEN-SOFT 4.2 software⁴². The PCA plot for the first two principal components (PCs) was shown in Fig. S3. The top principal component (PC) was significant by the Tracy-Widom test⁴³. Genotype imputation was performed using the pre-phasing/imputation stepwise approach implemented in IMPUTE2⁴⁴ and SHAPEIT⁴⁵. The imputation reference set consisted of 2186 phased haplotypes from the full 1000 Genomes Project Integrated Phase 3 Release⁴⁶. Imputed SNPs with a squared correlation between the imputed and true genotypes (rsq) <0.9 or SNPs with minor allele frequency <0.01 were removed. After imputation, 6,064,858 SNPs were used.

Use of GWAS summary data to derive polygenic risk scores

Summary statistics from GWAS of ADHD⁴⁷, MDD⁴⁸, BIP⁴⁹, SZ, and autism spectrum disorders⁵⁰ were down-loaded from the PGC website. These summary data were used as the discovery set to generate the polygenic risk scores for the EF test results. The sample size and number of SNPs for these data sets are in Table S2. Additional details on the demographics for the GWAS summary data are provided in the Supplementary Information.

Polygenic risk score analysis

The polygenic risk score (PRS) was introduced to summarize the effect of a set of SNPs in a test data set based on the GWAS summary statistics of a discovery dataset. PRSice-2 was used to calculate the results⁵¹. Before generating the scores, clumping was used to obtain SNPs in linkage equilibrium with an $r^2 < 0.1$ within a 250 bp window. PRSs from the GWAS summary data for five psychiatric disorders were created for each cognitive phenotype using the SNPs selected according to the significance of their association. The P-value threshold for significance was set from 0 to 0.5, increasing by 0.00005. The associations between the polygenic profile and the target phenotypes were examined in linear regression models with months (measure of age), sex, IQ and the first two PCs from PCA as covariates. The P-value threshold with the largest Nagelkerke's r^2 (variance explained by the PRS) was considered the best-fit threshold. The P-value for the linear regression was adjusted by using 10,000 label-swapping (randomly shuffle the phenotype) permutations. Adjusted P < 0.05 was considered

	Inhibitory control	Cognitive flexibility	Working memory	EF-PC	ADHD symptom	IQ
Inhibitory control	_					
Cognitive flexibility	0.337**	-				
Working memory	-0.267**	-0.390**	-			
EF-PC	0.718**	0.776**	-0.759**	-		
ADHD symptoms	0.153**	0.106**	-0.184**	0.204**	-	
IQ	-0.120**	-0.301**	0.298**	-0.327**	-0.037	-

Table 2 Pearson correlations among phenotypes.

The number of subjects for the correlation analysis among the three EF measures and EF-PC was 911. The number of subjects for the correlation analysis between EF measures and ADHD symptoms was 900. IQ was available for all participants. Asterisks (**) denotes *P*-value <0.01 (two-tailed). EF-PC: principal component for the three EF measures.

significant. To compare the effect of the PRS from different disorders on EF, we did normalization for the PRS under the best *P*-value threshold from PRSice-2 and further ran regression model using the same covariates in R to get the standardized beta coefficient.

Mediation analysis among PRS, inhibitory control, and ADHD symptoms

Since the inhibitory control is the core deficit of ADHD, the hypothesis is the PRS associated with inhibitory control may could affect the ADHD symptoms too. So, two mediation models were analyzed: (1) ADHD PRS associated with inhibitory control as independent variable (X), inhibitory control as mediator (M), ADHD symptoms as outcome (Y); (2) MDD PRS associated with inhibitory control as independent variable (X), inhibitory control as mediator (M), ADHD symptoms as outcome (Y). The standardized PRS was used for this analysis. The analysis was performed using the model 4 in PROCESS⁵². Both the mediator model and the outcome model included months (measure of age), IQ, sex and two PCs as covariates. Nonparametric bootstrapping with n = 10,000 was used to estimate the sampling distribution of the indirect effect. If the confidence interval (CI) of the effect didn't cross zero, the indirect effect was significant. R package "mediation" was further used to for the mediation analysis to double check the result.

Results

Phenotype description for the EF measures and ADHD symptoms

Table 1 shows descriptive statistics for the EF measures and ADHD symptoms. Among the 1,147 individuals, 963, 913, 1145, and 1134 subjects had inhibitory control, cognitive flexibility, working memory and ADHD symptom data, respectively. The distributions of the phenotypes are shown in Fig. S1. One principal component (EF-PC) was extracted from the 911 individuals having all three EF measures. The EF-PC explained 56.5% of the total variance in the exploratory factor model (see Figure S2 for details). All EF measures were significantly correlated with each other (Table 2). Inhibitory control was positively correlated with cognitive flexibility (r = 0.337); working memory was negatively correlated with inhibitory control (r = -0.267) and cognitive flexibility (r =-0.390). As expected, the correlations of EF-PC with the EF measures were higher (r = 0.718, 0.776, and -0.759 for inhibitory control, cognitive flexibility, and working memory respectively). All four EF measures were significantly correlated with ADHD symptoms (r = 0.153, 0.106, and -0.184 for inhibitory control, cognitive flexibility, and working memory respectively) and IQ (r =-0.120, -0.301, and 0.298 for inhibitory control, cognitive flexibility, and working memory respectively), which denoted the EF measures could significantly affect ADHD symptom and IQ; the correlation between ADHD symptoms and IQ was not significant (r = 0.037) (Table 2).

Association of polygenic risk for psychiatric disorders with executive functions

As shown in Table S3, EF-PC was significantly associated with the PRS for MDD (adjusted P = 0.023, $r^2 = 0.581\%$, coefficient = 0.0760 (se = 0.023)), ADHD (adjusted P = 0.036, $r^2 = 0.486\%$, coefficient = 0.0704 (se = 0.023)), and BIP (adjusted P = 7.70E-03, $r^2 = 0.642\%$, coefficient = 0.080 (se = 0.023)).

We further explored the association of each EF component with the genetic risk for psychiatric disorders (Fig. 2, Table S4). The PRS for ADHD was significantly associated with inhibitory control (adjusted P = 0.018, $r^2 =$ 0.923%, coefficient = 1.6922 (se = 0.5225)); the PRS for SZ was significantly associated with cognitive flexibility (adjusted P = 0.034, $r^2 = 0.615\%$, coefficient = 7.7142 (se = 2.6781)); the PRS for MDD was significantly associated with inhibitory control (adjusted P = 0.0313, $r^2 =$ 0.897%, coefficient = 1.6559 (se = 0.5188)); the PRS for BIP was significantly associated with working memory (adjusted P = 0.0416, $r^2 = 0.462\%$, coefficient = -0.1152



(se = 0.0391)); and the association of PRS for ASD with all three EF components did not pass the permutation correction.

Since both the ADHD and MDD PRSs were associated with inhibitory control, we constructed a multivariable regression model with standardized ADHD PRS, MDD PRS, their interaction and other covariates. The model showed P = 2.41E-03 for ADHD PRS and P = 2.86E-03 for MDD PRS, and the interaction was not significant (Table S5). Thus, each of these PRS were uniquely associated with inhibitory control.

Correlation and mediation analysis with ADHD symptoms

The ADHD PRS had a trend to be associated with ADHD symptoms but did not pass the multiple corrections (*P*-value = 0.0194, adjust P = 0.1828, $r^2 = 0.46\%$, beta = 0.2028 (se = 0.0867)). The ASD PRS was significantly associated with ADHD symptoms (P-value = 2.549E-03, adjust P = 0.038, $r^2 = 0.765\%$, beta = 0.2618 (se = 0.0866)). The PRSs of other disorders were not associated with ADHD symptoms (Table S6). To assess the indirect effect of the PRSs associated with inhibitory control on ADHD symptoms, we conducted mediation analyses. The ADHD PRS associated with inhibitory control had no direct effect on ADHD symptoms, but the indirect effect through the mediation of inhibitory control was significant (effect = 0.0209 (se = 0.0126), BootCI = (0.0027-0.0550)) (Fig. 3A). The MDD PRS associated with inhibitory control also showed an indirect effect on ADHD symptoms through the mediation of inhibitory control (effect = 0.0207)(se = 0.0123),BootCI =(0.0026–0.0524)) (Fig. 3b). The result from the "mediation" R package also obtained the same results (Table S7).

Discussion

This study used polygenic risk score analyses to explore the shared and specific genetic associations between psychiatric disorders and executive functions. The PRSs for MDD, ADHD, and BIP were associated with our overall measure of EF. These results agree with the idea that executive dysfunction is a transdiagnostic phenotype⁵³. On the other hand, we also found that some EF components associated with PRSs for different disorders: the ADHD and MDD PRSs were associated with inhibitory control; the PRS for BIP was associated with working memory, and that for SZ was associated with cognitive flexibility.

Dysfunction of inhibitory control and working memory in individuals with ADHD has been reported by many studies^{40,54}. Some studies found that the PRS for ADHD was associated with working memory²⁵⁻²⁷, but in this study, we did not identify the association of ADHD PRS with working memory, which may indicate the genetic basis of working memory was more complex and the current sample size was not large enough to catch the association of ADHD PRS with working memory. The previous papers did not find the association of ADHD PRS with inhibitory control in community recruited populations^{25–27}, but we found the ADHD PRS was associated with inhibitory control as measured by the variance of word interference time. The divergent results might be due to the different target sample because, compared with population samples, our ADHD sample would be expected to be enriched for the disinhibition. Mediation analyses indicated that the ADHD PRS contributed to ADHD symptoms through the mediation effect of inhibitory control. This result suggests that ADHD risk variants might affect inhibitory control, which leads to ADHD symptoms. This result is consistent with theoretical models that posit inhibitory control as a core deficit of ADHD patients.

Depression and ADHD often co-occur in clinical populations. ADHD symptoms are very common among MDD patients⁵⁵, while ADHD patients have higher rates of MDD⁵⁶. Inhibitory dysfunction has been reported, not only for patients with ADHD, but also for patients with MDD, as measured by the stop signal and Stroop inhibition tasks⁵⁷. The comorbidity between ADHD and MDD can partially be explained by their shared cognitive impairments and shared genetic variants. The genetic correlation between ADHD and MDD has been reported by the Brainstorm Consortium and the ADHD-PGC results^{17,47}. The present study found that the MDD PRS was also associated with inhibitory control. Although prior studies show the genetic risk for MDD to be associated with general cognitive ability²¹, ours is the first study that reports an association with inhibitory control. Both results are in the same direction, with higher genetic



risk associated with worse performance. Furthermore, the mediation analysis showed that the MDD PRS also had an indirect on ADHD symptoms through the mediation of inhibitory control. These results again suggested a shared genetic risk between these two disorders that may be mediated by the inhibitory control.

Although some study identified the association of inhibitory control with SZ and BIP⁵⁸, we didn't find their association but the association of BIP PRS with working memory and SZ PRS with cognitive flexibility. Working memory impairment is a characteristic that BIP shares with ADHD. Brown et al. reported hypoactivity in the frontal and parietal regions in individuals with ADHD and those with BIP compared with controls when they performed a working memory task⁵⁹. CACNA1C, a common risk gene for five major psychiatric disorders, has been associated with spatial working memory in BIP⁶⁰. In this study, the PRS for BIP was associated with decreased working memory, providing further evidence for the genetic correlation between BIP, working memory and ADHD. Impairment of cognitive flexibility has been found in SZ⁶¹. In our study, the PRS for SZ was associated with cognitive flexibility in ADHD children. Although Vitiello et al. did not find that ADHD increased the risk for psychotic symptoms⁶², a significant genetic correlation between ADHD and SZ was reported^{17,47}. Our result suggests that impaired cognitive flexibility has a shared genetic basis in ADHD and SZ, which is in the same direction as the association with general cognitive function.

The relationship of cognition with ASD is complex. Genetic correlation analysis^{17,50} and PRS analysis in the general population²³ showed that the polygenic risk for ASD was associated with *higher* general cognition or

intelligence. In contrast, the PRS analysis for EF showed the opposite effect such that high PRS for general intelligence was associated with worse social problems²⁸. Grove et al. identified a positive genetic correlation between ADHD and ASD⁵⁰ and both of ADHD and ASD were reported to have executive impairments. In this study, the ASD PRS was significantly associated with ADHD symptoms, but not EF. One possible explanation is that EF is not a core impairment in ASD, for which social communication and repetitive behavior were core symptoms that might be directly correlated with the genetic risk of the disorder. ASD may express different features of EF deficits from ADHD⁶³. Another plausible explanation is the relatively low number of genome-wide significant signals in the GWAS of autism⁵⁰.

Compared with previous PRS analyses for cognition and psychiatric disorders, a main difference with this study is the age of our sample and how it was ascertained. Most prior studies used data from community-based populations; we used data from ADHD patients. The distribution of EF may differ in these populations. The samples used in this study likely have ADHD-specific features (e.g., worse executive functions) that differ from the general population, which might facilitate the finding of shared cognition deficit with other psychiatric disorders by using PRS analysis. Another difference was the age of the samples. Most prior studies used adolescent or adult samples, whose cognitive development differs with children's. The association of genetic risk of some disease with children's cognition may not be consistent for adults. For example, ADHD related impairment that exist in childhood can disappear or attenuate in late adolescence or adults⁶⁴.

This study has some limitations. The sample sizes for the EF measures were small. Replication in larger samples is needed. Second, the GWAS summary data for the PRS calculations in this study were from European individuals. Cross-ethnic PRS analyses are affected by the different LD structures and allele frequencies in different ethnicities⁶⁵, which reduces the power to find associations. We also attempted to analyze the association of PRSs for the Chinese ADHD GWAS data³⁴ with EF. Since the sample size of the Chinese ADHD GWAS is small, there was no significant result for all EF components (data not shown). A larger sample size is important for the PRS definition to address a comprehensive genetic component. As estimated by Martin et. al.⁶⁶, applying GWAS summary statistics from the European population to other ancestries will reduce prediction accuracy. If using Chinese population GWAS data with the same sample size, the effect size might be bigger.

In conclusion, by using polygenic risk score analyses, we found that the polygenic risk for different psychiatric disorders (ADHD, BIP, MDD, SZ) showed distinct effects on executive function domains. Although all psychiatric disorders had general cognition dysfunction, the specific impairments of the EF components may be different. The results of this study helped to identify the core cognitive impairment that most associated with the genetic risk for each mental disorder, which may further help elucidate the risk genes and motivate efforts to develop EF measures as a diagnostic marker and future treatment target.

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Conflict of interest

 $\ensuremath{\mathsf{Dr}}$ S.V.F. is a consultant for Genomind. The remaining authors declare no conflict of interest.

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References

- Funahashi, S. & Andreau, J. M. Prefrontal cortex and neural mechanisms of executive function. J. Physiol. 107, 471–482 (2013).
- Baune, B. T. & Malhi, G. S. A review on the impact of cognitive dysfunction on social, occupational, and general functional outcomes in bipolar disorder. *Bipolar Disord*. 17(Suppl 2), 41–55 (2015).
- Bortolato, B., Carvalho, A. F. & McIntyre, R. S. Cognitive dysfunction in major depressive disorder: a state-of-the-art clinical review. *CNS Neurological Disord. Drug Targets* 13, 1804–1818 (2014).
- Kitchen, H., Rofail, D., Heron, L. & Sacco, P. Cognitive impairment associated with schizophrenia: a review of the humanistic burden. *Adv. Ther.* 29, 148–162 (2012).
- Snyder, H. R., Miyake, A. & Hankin, B. L. Advancing understanding of executive function impairments and psychopathology: bridging the gap between clinical and cognitive approaches. *Front. Psychol.* 6, 328 (2015).
- Best, J. R. & Miller, P. H. A developmental perspective on executive function. Child Dev. 81, 1641–1660 (2010).
- 7. Diamond, A. Executive functions. Annu. Rev. Psychol. 64, 135-168 (2013).
- Weissman, A. S., Chu, B. C., Reddy, L. A. & Mohlman, J. Attention mechanisms in children with anxiety disorders and in children with attention deficit hyperactivity disorder: implications for research and practice. J. Clin. Child Adolesc. Psychol. 41, 117–126 (2012).
- Shanmugan, S. et al. Common and dissociable mechanisms of executive system dysfunction across psychiatric disorders in youth. *Am. J. Psychiatry* 173, 517–526 (2016).
- Willcutt, E. G., Doyle, A. E., Nigg, J. T., Faraone, S. V. & Pennington, B. F. Validity of the executive function theory of attention-deficit/hyperactivity disorder: a meta-analytic review. *Biol. Psychiatry* 57, 1336–1346 (2005).
- Bora, E. Neurocognitive features in clinical subgroups of bipolar disorder: a meta-analysis. J. Affect. Disord. 229, 125–134 (2018).
- Essizoglu, A., Kosger, F., Akarsu, F. O., Ozaydin, O. & Gulec, G. Theory of mind and selective attention, response inhibition, cognitive flexibility in patients with schizophrenia. *Noro Psikiyatr. Arsivi* 54, 162–167 (2017).
- White, L. K. et al. An evaluation of the specificity of executive function impairment in developmental psychopathology. J. Am. Acad. Child Adolesc. Psychiatry 56, 975–982 e973 (2017).
- Blokland, G. A. M. et al. Heritability of neuropsychological measures in schizophrenia and nonpsychiatric populations: a systematic review and metaanalysis. *Schizophrenia Bull.* 43, 788–800 (2017).
- Solovieff, N., Cotsapas, C., Lee, P. H., Purcell, S. M. & Smoller, J. W. Pleiotropy in complex traits: challenges and strategies. *Nat. Rev. Genet* 14, 483–495 (2013).
- Consortium. C-DGotPG. Genomic relationships, novel loci, and pleiotropic mechanisms across eight psychiatric disorders. *Cell* **179**, 1469–1482 e1411 (2019).
- Brainstorm, C. Analysis of shared heritability in common disorders of the brain. Science 360, eaap8757 (2018).
- Schork, A. J. et al. A genome-wide association study of shared risk across psychiatric disorders implicates gene regulation during fetal neurodevelopment. *Nat. Neurosci.* 22, 353–361 (2019).
- Lencz, T. et al. Molecular genetic evidence for overlap between general cognitive ability and risk for schizophrenia: a report from the Cognitive Genomics consorTium (COGENT). *Mol. Psychiatry* **19**, 168–174 (2014).
- Hubbard, L. et al. Evidence of common genetic overlap between schizophrenia and cognition. *Schizophrenia Bull.* 42, 832–842 (2016).
- Howard, D. M. et al. Genome-wide meta-analysis of depression identifies 102 independent variants and highlights the importance of the prefrontal brain regions. *Nat. Neurosci.* 22, 343–352 (2019).
- Stergiakouli, E. et al. Association between polygenic risk scores for attentiondeficit hyperactivity disorder and educational and cognitive outcomes in the general population. *Int. J. Epidemiol.* 46, 421–428 (2016).
- Clarke, T. K. et al. Common polygenic risk for autism spectrum disorder (ASD) is associated with cognitive ability in the general population. *Mol. Psychiatry* 21, 419–425 (2016).
- 24. Stahl, E. A. et al. Genome-wide association study identifies 30 loci associated with bipolar disorder. *Nat. Genet* **51**, 793–803 (2019).

- Nigg, J. T. et al. Working memory and vigilance as multivariate endophenotypes related to common genetic risk for attention-deficit/hyperactivity disorder. J. Am. Acad. Child Adolesc. Psychiatry 57, 175–182 (2018).
- Hagenaars, S. P. et al. Shared genetic aetiology between cognitive functions and physical and mental health in UK Biobank (N=112 151) and 24 GWAS consortia. *Mol. Psychiatry* 21, 1624–1632 (2016).
- Martin, J., Hamshere, M. L., Stergiakouli, E., O'Donovan, M. C. & Thapar, A. Neurocognitive abilities in the general population and composite genetic risk scores for attention-deficit hyperactivity disorder. J. Child Psychol. Psychiatry Allied Discip. 56, 648–656 (2015).
- Torske, T. et al. Autism spectrum disorder polygenic scores are associated with every day executive function in children admitted for clinical assessment. *Autism Res.* 13, 207–220 (2020).
- Mistry, S., Escott-Price, V., Florio, A. D., Smith, D. J. & Zammit, S. Investigating associations between genetic risk for bipolar disorder and cognitive functioning in childhood. *J. Affect. Disord.* **259**, 112–120 (2019).
- Benca, C. E. et al. Predicting cognitive executive functioning with polygenic risk scores for psychiatric disorders. *Behav. Genet.* 47, 11–24 (2017).
- Nigg, J. T., Willcutt, E. G., Doyle, A. E. & Sonuga-Barke, E. J. Causal heterogeneity in attention-deficit/hyperactivity disorder: do we need neuropsychologically impaired subtypes? *Biol. Psychiatry* 57, 1224–1230 (2005).
- Yang, L., Wang, Y. F., Qian, Q. J., Biederman, J. & Faraone, S. V. DSM-IV subtypes of ADHD in a Chinese outpatient sample. *J. Am. Acad. Child Adolesc. Psychiatry* 43, 248–250 (2004).
- Barkley, R. Attention-Deficit Hyperactivity Disorder: A Clinical Workbook. 2nd edn. (Guilford, New York, 1998).
- Yang, L. et al. Polygenic transmission and complex neuro developmental network for attention deficit hyperactivity disorder: genome-wide association study of both common and rare variants. *Am. J. Med. Genet. B Neuropsychiatr. Genet.* 162B, 419–430 (2013).
- Yang, L. et al. A new locus regulating MICALL2 expression was identified for association with executive inhibition in children with attention deficit hyperactivity disorder. *Mol. Psychiatry* 23, 1014–1020 (2018).
- Zhang, K. et al. Genetic analysis for cognitive flexibility in the trail-making test in attention deficit hyperactivity disorder patients from single nucleotide polymorphism, gene to pathway level. World J. Biol. Psychiatry 20, 1–10 (2017).
- Anderson, V. Assessing executive functions in children: biological, psychological, and developmental considerations. *Pediatr. Rehabilitation* 4, 119–136 (2001).
- Trampush, J. W., Jacobs, M. M., Hurd, Y. L., Newcorn, J. H. & Halperin, J. M. Moderator effects of working memory on the stability of ADHD symptoms by dopamine receptor gene polymorphisms during development. *Dev. Sci.* 17, 584–595 (2014).
- Hoaglin, D. C., Iglewicz, B. & Tukey, J. W. Performance of some resistant rules for outlier labeling. J. Am. Stat. Assoc. 81, 991–999 (1986).
- Barkley, R. Attention-deficit Hyperactivity Disorder (third edition): A Handbook for Diagnosis and Treatment (The Guilford Press, New York, 2006).
- 41. Chen, J. et al. Genetic structure of the Han Chinese population revealed by genome-wide SNP variation. *Am. J. Hum. Genet.* **85**, 775–785 (2009).
- 42. Price, A. L. et al. Principal components analysis corrects for stratification in genome-wide association studies. *Nat. Genet.* **38**, 904–909 (2006).
- Tracy, C. A. & Widom, H. Level spacing distributions and the Bessel kernel. CommunMath Phys. 161, 289–309 (1994).
- Howie, B., Marchini, J. & Stephens, M. Genotype imputation with thousands of genomes. G3 1, 457–470 (2011).

- Delaneau, O., Marchini, J. & Zagury, J. F. A linear complexity phasing method for thousands of genomes. *Nat. Methods* 9, 179–181 (2012).
- Sudmant, P. H. et al. An integrated map of structural variation in 2,504 human genomes. *Nature* 526, 75–81 (2015).
- Demontis, D. et al. Discovery of the first genome-wide significant risk loci for attention deficit/hyperactivity disorder. *Nat. Genet.* **51**, 63–75 (2019).
- Wray, N. R. et al. Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. *Nat. Genet.* 50, 668–681 (2018).
- Bipolar, D., Schizophrenia Working Group of the Psychiatric Genomics Consortium. Genomic dissection of bipolar disorder and schizophrenia, including 28 subphenotypes. *Cell* **173**, 1705–1715 e1716 (2018).
- Grove, J. et al. Identification of common genetic risk variants for autism spectrum disorder. *Nat. Genet* 51, 431–444 (2019).
- Euesden, J., Lewis, C. M. & O'Reilly, P. F. PRSice: polygenic risk score software. Bioinformatics 31, 1466–1468 (2015).
- Hayes, AF. Introduction to Mediation, Moderation, and Conditional Process Analysis A Regression-Based Approach (Guilford Press, 2013).
- McTeague, L. M., Goodkind, M. S. & Etkin, A. Transdiagnostic impairment of cognitive control in mental illness. J. Psychiatr. Res. 83, 37–46 (2016).
- Nigg, J. T., Blaskey, L. G., Stawicki, J. A. & Sachek, J. Evaluating the endophenotype model of ADHD neuropsychological deficit: results for parents and siblings of children with ADHD combined and inattentive subtypes. J. Abnorm. Psychol. 113, 614–625 (2004).
- Bron, T. I. et al. Prevalence of ADHD symptoms across clinical stages of major depressive disorder. J. Affect. Disord. 197, 29–35 (2016).
- Biederman, J. et al. New insights into the comorbidity between ADHD and major depression in adolescent and young adult females. J. Am. Acad. Child Adolesc. Psychiatry 47, 426–434 (2008).
- Disner, S. G., Beevers, C. G., Haigh, E. A. & Beck, A. T. Neural mechanisms of the cognitive model of depression. *Nat. Rev. Neurosci.* 12, 467–477 (2011).
- Tsujii, N., Mikawa, W., Adachi, T., Hirose, T. & Shirakawa, O. Shared and differential cortical functional abnormalities associated with inhibitory control in patients with schizophrenia and bipolar disorder. *Sci. Rep.* 8, 4686 (2018).
- Brown, A. et al. Working memory network alterations and associated symptoms in adults with ADHD and Bipolar Disorder. J. Psychiatr. Res. 46, 476–483 (2012).
- Zhang, Q. et al. The effects of CACNA1C gene polymorphism on spatial working memory in both healthy controls and patients with schizophrenia or bipolar disorder. *Neuropsychopharmacology* **37**, 677–684 (2012).
- Fellini, L., Kumar, G., Gibbs, S., Steckler, T. & Talpos, J. Re-evaluating the PCP challenge as a pre-clinical model of impaired cognitive flexibility in schizophrenia. *Eur. Neuropsychopharmacol.* 24, 1836–1849 (2014).
- Vitiello, B. et al. Psychotic symptoms in attention-deficit/hyperactivity disorder: an analysis of the MTA database. J. Am. Acad. Child Adolesc. Psychiatry 56, 336–343 (2017).
- Xie, R., Sun, X., Yang, L. & Guo, Y. Executive function of high functioning autistic adults: a systematic review and meta-analysis. *Autism Res.* https://doi.org/ 10.1002/aur.2304 [Online ahead of print] (2020).
- Shaw, P. et al. Attention-deficit/hyperactivity disorder is characterized by a delay in cortical maturation. *Proc. Natl Acad. Sci. USA* 104, 19649–19654 (2007).
- Vilhjalmsson, B. J. et al. Modeling linkage disequilibrium increases accuracy of polygenic risk scores. Am. J. Hum. Genet. 97, 576–592 (2015).
- Martin, A. R. et al. Clinical use of current polygenic risk scores may exacerbate health disparities. *Nat. Genet.* **51**, 584–591 (2019).