Taylor, M.J. and Freeman, D. and Lundstrom, S. and Larsson, H. and Ronald, Angelica (2022) Heritability of psychotic experiences in adolescents and interaction with environmental risk. JAMA Psychiatry 79 (9), pp. 889-897. ISSN 2168-622X.

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Heritability of Psychotic Experiences in Adolescents and Interaction With Environmental Risk

Mark J. Taylor, PhD; Daniel Freeman, PhD; Sebastian Lundström, PhD; Henrik Larsson, PhD; Angelica Ronald, PhD

IMPORTANCE Genetic risk factors are known to play a role in the etiology of psychotic experiences in the general population. Little is known about whether these risk factors interact with environmental risks for psychotic experiences.

OBJECTIVE To assess etiological heterogeneity and exposure to environmental risks associated with psychotic experiences in adolescence using the twin design.

DESIGN, SETTING, AND PARTICIPANTS This twin study, conducted from December 1, 2014, to August 31, 2020, included a UK-based sample of twin pairs aged 16 years. This investigation evaluated the extent to which the genetic variance underlying psychotic experiences and the magnitude of the heritability of psychotic experiences was moderated by exposure to 5 environmental risk factors (bullying, dependent life events, cannabis use, tobacco use, and low birth weight). Psychotic experiences were assessed by 5 self-reported measures and 1 parent-reported measure. Participants’ exposure to environmental risks was assessed at birth and age 12 to 16 years. Structural equation models were used to assess differences in the variance in and heritability of psychotic experiences across these exposures, while controlling for gene-environment correlation effects. Analyses were repeated in an independent Swedish sample. Data analyses were performed from September 1, 2018, to August 31, 2020.

MAIN OUTCOMES AND MEASURES Primary outcome measures were exposure to environmental factors, as measured by a composite score, and psychotic experiences.

RESULTS A total of 4855 twin pairs (1926 female same-sex pairs, 1397 male same-sex pairs, and 1532 opposite-sex pairs) were included from the Twins Early Development Study (TEDS), and 6435 twin pairs (2358 female same-sex pairs, 1861 male same-sex pairs, and 2216 opposite-sex pairs) were included from the Child and Adolescent Twin Study in Sweden (CATSS). Mean age of twins from TEDS was 16.5 years. Mean age of twins from CATSS was 18.6 years. More exposure to environmental risk factors was associated with having more psychotic experiences. The relative contribution of genetic influences to psychotic experiences was lower with increasing environmental exposure for paranoia (44%; 95% CI, 33%-53% to 38%; 95% CI, 14%-58%), cognitive disorganization (47%; 95% CI, 38%-51% to 32%; 95% CI, 11%-45%), grandiosity (41%; 95% CI, 29%-52% to 32%; 95% CI, 9%-48%), and anhedonia (49%; 95% CI, 42%-53% to 37%; 95% CI, 15%-54%). This pattern was replicated for the measure of psychotic experiences in the independent Swedish replication sample. The heritability of hallucinations and parent-rated negative symptoms remained relatively constant.

CONCLUSIONS AND RELEVANCE Findings of this twin study suggest that environmental factors play a greater role in the etiology of psychotic experiences than genetic factors. The relative importance of environmental factors is even higher among individuals exposed to environmental risks for psychotic experiences, highlighting the importance of a diathesis-stress or bioecological framework for understanding adolescent psychotic experiences.
Psychotic experiences, such as paranoia and hallucinations, are relatively common in adolescence, with an estimated prevalence of 5% in adults. Although they follow a transient course in some individuals, in others they are persistent and associated with increased risk for psychosis, psychiatric disorders, suicide, and physical disorders. Many associations with adverse outcomes are independent of psychiatric disorders, highlighting the need to understand psychotic experiences in their own right.

There has been considerable progress in understanding the etiology of psychotic experiences, including moderate twin heritability (15%-59%) and single-nucleotide variation heritability of 3% to 9%. These estimates indicate that environmental factors play a considerable role in psychotic experiences. Factors found to be associated with psychotic experiences include bullying and childhood maltreatment, life events, cannabis use, and tobacco use. Early exposures, such as obstetric complications, have also been implicated. These exposures are heritable, however, and their association with psychotic experiences is partly attributable to genetic influences. For example, British twin studies found that the associations between psychotic experiences and bullying, life events, and tobacco use were explained by genetics, indicating that some of these associations are not causal.

This evidence suggests that genetic and environmental influences on psychotic experiences do not operate independently of one another. Another manner in which this may occur is through gene-by-environment interaction, whereby the importance of genetic influences may vary dependent on environmental exposure and vice versa. Two long-standing theoretical frameworks provide means to illustrate this. The diathesis-stress model posits that genetic susceptibility is required to trigger a response to environmental exposures. In the context of a twin study, this would lead to higher heritability estimates in communities with more environmental exposure compared with communities with less exposure. By contrast, the biocological model posits that genetic factors become more pertinent in more favorable environments, leading to lower heritability in twin studies in the context of environmental exposure. It is unknown whether these models apply to psychotic experiences.

We aimed to test gene-by-environment interaction in relation to psychotic experiences using twin methods. We tested whether the genetic and environmental variance in these phenotypes varied as a function of exposure to environmental risks. After testing for these associations in a British sample of adolescents, we assessed whether our results could be replicated in a Swedish sample. We hypothesized that the genetic and environmental variance in psychotic experiences would fluctuate with exposure to environmental risks. We did not draw specific hypotheses about the direction of such changes owing to the paucity of existing evidence.

### Methods

#### Study Populations

The Twins Early Development Study (TEDS) includes twins born in England and Wales from 1994 to 1996 who participated in the Longitudinal Experiences and Perceptions Study (LEAP) at age 16 years. A total of 5059 of 10874 invited families (47%) participated. The sample is representative of the UK population on various demographic characteristics, including ethnicity and socioeconomic status. We excluded participants with autism, genetic syndromes, chromosomal abnormalities, extremely severe obstetric complications, and those missing first-contact data. Zygosity was ascertained through DNA testing and a questionnaire assessing twin resemblance. TEDS has ethical approval from the King’s College London research ethics committee. TEDS participants provided written informed consent before participation.

The Child and Adolescent Twin Study in Sweden (CATSS) comprises families of Swedish twins who are invited to participate when the twins turn age 9 years. We used data collected from the participants at ages 15 and 18 years, which have respective response rates of 61% and 59%. CATSS is representative of the Swedish population on various characteristics. We excluded participants with chromosomal abnormalities or brain injuries. Zygosity was ascertained using a panel of single nucleotide variations or a questionnaire assessing twin similarity and reconfirmed for genotyped twins. CATSS has ethical approval from the Stockholm County Ethical Review Board. Informed consent was inferred from participant completion of questionnaires. The measures used in this study are described in eTable 1 in the Supplement.

#### Environmental Exposures

We selected 5 exposures, based on prior research: bullying, dependent life events, cannabis use, tobacco use, and low birth weight. We summed them to create an exposure score for each individual. Individuals with 4 or 5 exposures were collapsed into 1 group. The definition of each exposure was designed to maximize statistical power, while simultaneously ensuring that participants who were sufficiently strongly exposed to each risk factor.

#### TEDS Measures

Bullying was measured using the Multidimensional Peer Victimization Scale at age 12 years. This measure comprises...
16 items, including 4 subscales. Scores on each scale ranged from 0 to 8; we considered participants exposed to bullying if they scored more than 6 on at least 1 subscale. Dependent life events are life events that are associated with an individual’s behavior or circumstances, such as the breakdown of a relationship or experiencing a crime. They were measured using the abbreviated Coddington Life Events Record at age 16 years.\(^{26}\) Participants reported whether or not they had experienced 10 dependent life events. We considered them exposed if they experienced at least 3. Cannabis use and tobacco use were both measured using a checklist inquiring about substance use and were assessed by a binary item at age 16 years. If these items were endorsed, participants were considered exposed. Birth weight was reported by the parents at first contact. We defined low birth weight as a birth weight within the lowest 15% of the distribution (<1990 g).

**CATSS Measures**

Bullying was assessed at age 15 years using the Olweus Bully/Victim Questionnaire,\(^{27}\) including 16 questions. We considered participants to have been bullied if they reported at least 1 form of bullying, perpetrated mainly by 1 student, and with a duration of 1 to 2 weeks. Dependent life events were measured by a checklist of 29 items at age 18 years, including 13 about dependent life events. We considered participants to be exposed if they endorsed 3 or more dependent life events. Cannabis use and tobacco use were assessed in the same way as in TEDS, at age 18 years. Birth weight was ascertained from the medical birth register. We considered participants to be exposed if their birth weight was in the lowest 15% of the distribution (<2040 g).

**Psychotic Experiences**

In TEDS, psychotic experiences were measured using the Specific Psychotic Experiences Questionnaire (SPEQ)\(^{17}\) at age 16 years. The SPEQ was developed from preexisting measures adapted for use in adolescent samples and includes 5 self-reported subscales: paranoia, hallucinations, cognitive disorganization, grandiosity, and anhedonia and a parent-reported negative-symptoms subcale. In CATSS, twins completed the Adolescent Psychotic-Like Symptom Screener (APSS)\(^{28}\) at age 18 years, which includes 7 items.

**Statistical Analyses**

Before analysis, participants with missing data on the environmental exposure were excluded because they could not be included in moderation twin models. Participants with missing data on the psychotic experience measures were addressed using maximum-likelihood estimation, which is robust to missing data. We tested the phenotypic associations between cumulative exposure to environmental risks and psychotic experiences using linear regression models, implemented as generalized estimating equations to account for related individuals in the samples. Second, we fitted univariate twin models to the APSS in CATSS and environmental exposure variables in both samples. We did not repeat univariate analyses of the SPEQ, as these have been published.\(^{8}\)

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**Figure 1. Diagram of a Moderation Model**

The path diagram is shown for 1 twin only. The Adolescent Psychotic-Like Symptom Screener (APSS) and the Specific Psychotic Experiences Questionnaire (SPEQ) were the 2 measures used to assess psychotic experiences. The variance in the exposure is decomposed into additive genetic (A), shared environmental (C), and nonshared environmental (E) components, which are derived from the path coefficients (labeled \(a_u, c_u,\) and \(e_u\) in the path diagram). The paths connecting the exposure with SPEQ/APSS are covariance paths (\(a_{cu}, c_{cu},\) and \(e_{cu}\)). The residual variance in SPEQ/APSS is also decomposed into A, C, and E, based on path coefficients \(a_e, c_e,\) and \(e_e\). Moderation effects are also estimated for the covariance (\(\beta_{auM}, \beta_{cuM},\) and \(\beta_{euM}\)) and residual (\(\beta_{au}, \beta_{cu},\) and \(\beta_{eu}\)) paths. These moderation effects can be used to calculate the value of each variance component (both the raw variance and the proportion of variance explained) at different levels of the moderator variable.

We tested for gene-environment interaction using moderation models.\(^{19}\) These models estimate the genetic and environmental variance in a trait at different levels of a measured environmental exposure (ie, moderator variable). This approach can yield false-positive results in the presence of gene-environment correlation.\(^{30}\) Prior studies in TEDS have reported genetic correlations between some of our exposures and psychotic experiences\(^{17,19}\); therefore, we used the full bivariate moderation model here.\(^{30}\) This model accounts for gene-environment correlation and thus the only model that estimates true moderation effects.\(^{30}\) Figure 1 shows a diagram of this model. The environmental exposure composite and psychotic experiences were included as manifest variables. The variance in each of these was decomposed into genetic and environmental components, which were then used to calculate the proportion of variance explained by each component. Covariance paths between them are included. The exposure was also included as a moderator; coefficients that correspond to moderation effects for each variance component were estimated. Using these parameters, we calculated the genetic (A) and environmental (C and E) variance associated with each level of exposure. The statistical significance of these effects was then tested by constraining the moderation effects to be equal across exposure groups, first for each variance component separately and then for all components. The statistical significance of these effects was tested using the likelihood-ratio test. We present the results here as the proportions of variance explained by each component for ease of interpretation. Phenotypic analyses were conducted using the drgee package of R (R Foundation).\(^{31}\) Twin analyses were conducted using the OpenMx package of R (R Foundation).\(^{32}\) Data were analyzed from September 1, 2018, to August 31, 2020.
Results

Descriptive statistics are in Table 1. A total of 4855 twin pairs (1926 female same-sex pairs, 1397 male same-sex pairs, and 1532 opposite-sex pairs) were included from TEDS, and 6435 twin pairs (2358 female same-sex pairs, 1861 male same-sex pairs, and 2216 opposite-sex pairs) were included from CATSS. Mean age of twins from TEDS was 16.5 years, and the mean age of twins from CATSS was 18.6 years. In TEDS, 43.3% of participants (4209 of 9710) had at least 1 environmental exposure, compared with 52.1% of participants (8932 of 17136) in CATSS.

Phenotypic Analyses

Figure 2 shows regression coefficients from the phenotypic analyses. In TEDS, all levels of exposure were associated with paranoia (1 exposure: \( \beta = 2.20; 95\% \text{ CI}, 1.70-2.71; 2 \) exposures: \( \beta = 3.72; 95\% \text{ CI}, 2.89-4.54; 3 \) exposures: \( \beta = 5.25; 95\% \text{ CI}, 3.99-6.59).
Twin Analyses

Univariate results for the APSS and environmental composites are displayed in eTables 2 to 5 in the Supplement. The APSS showed heritability of 23% in boys and 40% in girls. The heritability of the environmental composite was 37% (95% CI, 27%-46%) in TEDS and 24% (range, 19%-29%) in CATSS. Figure 3 shows the estimates from the moderation models (eTables 6 and 7 in the Supplement). Table 2 shows the fit statistics. Statistically significant differences between 2 models indicate statistically significant moderation effects. We observed moderation effects for 5 out of 6 measures (83.3%) in TEDS: paranoia, hallucinations, cognitive disorganization, grandiosity, and anhedonia (of which 4 were statistically significant). Their heritability decreased with increasing environmental exposure. However, fluctuations in the underlying variance components differed across these measures. Paranoia heritability changed from 44% (95% CI, 33%-53%) to 38% (95% CI, 14%-58%) with increasing exposure. Cognitive disorganization heritability changed from 47% (95% CI, 38%-51%) to 32% (95% CI, 11%-45%) across groups, because the total genetic variance decreased, whereas nonshared environmental
variance increased. The same was true for grandiosity, where heritability changed from 41% (95% CI, 29%-52%) to 32% (95% CI, 9%-48%), owing to decreasing genetic variance and increasing nonshared environmental variance. For anhedonia, the change in heritability from 49% (95% CI, 42%-53%) to 37% (95% CI, 15%-54%) was attributable to decreasing genetic variance and stable nonshared environmental variance. For hallucinations, heritability was constant, from 32% (95% CI, 21%-42%) in individuals with no exposure to 31% in all 4 exposure groups. Underlying this apparent stability were increases in both genetic and environmental variance. The CATSS sample yielded statistically significant moderation effects, with heritability of psychotic experiences changing from 35% (95% CI, 23%-43%) to 31% (95% CI, 9%-52%) across groups.

Discussion

In this cohort study, we tested whether the heritability of adolescent psychotic experiences changes with exposure to environmental risks associated with psychotic experiences. Results suggest a gene-by-environment interaction for paranoia, hallucinations, cognitive disorganization, grandiosity, and anhedonia. The findings were replicated in an independent Swedish sample, thus lending robustness to our results. Our study thus suggests differences in heritability of certain psychotic experiences that may be associated with environmental exposures.

Specifically, there was an observed reduction in heritability of psychotic experiences in the presence of environmental exposures. These results are consistent with a biocological framework, which would predict that more favorable environments would lead to higher heritability.²⁰ Our results run contrary to a diathesis-stress pathway to psychotic experiences, which would predict that environmental risks trigger a genetic susceptibility to a given disorder and would thus lead to higher heritability of a phenotype in the presence of environmental risks.²⁰ It is also important to put our results in the context of prior studies of gene-by-environment interaction in relation to...
One study\(^3\) reported that the association between environmental risks and psychotic experiences increased among individuals with a family history of psychosis. Although this finding also supports gene-by-environment interaction, the results are somewhat different than what would be expected from our analyses because our analyses suggest that genetic factors were less salient in the presence of environmental exposures. Moreover, other studies\(^4,5\) have found no evidence of gene-by-environment interaction for psychotic experiences. Methodological differences may underlie these discrepancies. First, family history is not the same as genetic influence because family history includes a combination of

### Table 2. Twin Model Fit Statistics

| Model                | \(-2LL^a\)  | Parameters | \(df\)  | \(\Delta\chi^2b\) | \(\Delta df^c\) | \(P\) value\(^d\) |
|----------------------|--------------|------------|---------|------------------|----------------|------------------|
| **Paranoia**         |              |            |         |                  |                |                  |
| Full bivariate       | 49.291.69    | 17         | 19.292 | NA               | NA             | NA               |
| Drop A moderation    | 49.293.00    | 16         | 19.293 | 1.32             | 1              | .25              |
| Drop C moderation    | 49.291.70    | 16         | 19.293 | 0.01             | 1              | .91              |
| Drop E moderation    | 49.292.59    | 16         | 19.293 | 0.90             | 1              | .34              |
| Drop all moderation  | 49.299.02    | 14         | 19.295 | 7.33             | 3              | .06              |
| **Hallucinations**   |              |            |         |                  |                |                  |
| Full bivariate       | 49.578.99    | 17         | 19.302 | NA               | NA             | NA               |
| Drop A moderation    | 49.579.16    | 16         | 19.303 | 0.16             | 1              | .69              |
| Drop C moderation    | 49.579.00    | 16         | 19.303 | 0.00             | 1              | .95              |
| Drop E moderation    | 49.582.82    | 16         | 19.303 | 3.82             | 1              | .05              |
| Drop all moderation  | 49.588.62    | 14         | 19.305 | 9.63             | 3              | .02              |
| **Cognitive disorganization** |          |            |         |                  |                |                  |
| Full bivariate       | 49.476.53    | 17         | 19.291 | NA               | NA             | NA               |
| Drop A moderation    | 49.476.87    | 16         | 19.292 | 0.34             | 1              | .56              |
| Drop C moderation    | 49.476.53    | 16         | 19.292 | 0.00             | 1              | <.99             |
| Drop E moderation    | 49.488.19    | 16         | 19.292 | 11.66            | 1              | .001             |
| Drop all moderation  | 49.494.14    | 14         | 19.294 | 17.61            | 3              | .001             |
| **Grandiosity**      |              |            |         |                  |                |                  |
| Full bivariate       | 49.509.94    | 17         | 19.246 | NA               | NA             | NA               |
| Drop A moderation    | 49.510.30    | 16         | 19.247 | 0.37             | 1              | .55              |
| Drop C moderation    | 49.511.65    | 16         | 19.247 | 1.71             | 1              | .19              |
| Drop E moderation    | 49.513.53    | 16         | 19.247 | 3.59             | 1              | .06              |
| Drop all moderation  | 49.518.32    | 14         | 19.249 | 8.38             | 3              | .04              |
| **Anhedonia**        |              |            |         |                  |                |                  |
| Full bivariate       | 49.541.41    | 17         | 19.249 | NA               | NA             | NA               |
| Drop A moderation    | 49.541.43    | 16         | 19.250 | 0.01             | 1              | .92              |
| Drop C moderation    | 49.541.53    | 16         | 19.250 | 0.11             | 1              | .74              |
| Drop E moderation    | 49.548.50    | 16         | 19.250 | 7.09             | 1              | .008             |
| Drop all moderation  | 49.553.57    | 14         | 19.252 | 12.16            | 3              | .007             |
| **Negative symptoms**|              |            |         |                  |                |                  |
| Full bivariate       | 47.326.21    | 17         | 19.327 | NA               | NA             | NA               |
| Drop A moderation    | 47.326.96    | 16         | 19.328 | 0.75             | 1              | .39              |
| Drop C moderation    | 47.326.21    | 16         | 19.328 | 0.00             | 1              | .99              |
| Drop E moderation    | 47.326.65    | 16         | 19.328 | 0.44             | 1              | .51              |
| Drop all moderation  | 47.332.04    | 14         | 19.330 | 5.83             | 3              | .12              |
| **APSS**             |              |            |         |                  |                |                  |
| Full bivariate       | 90.237.22    | 17         | 43.692 | NA               | NA             | NA               |
| Drop A moderation    | 90.237.77    | 16         | 43.693 | 0.55             | 1              | .46              |
| Drop C moderation    | 90.237.65    | 16         | 43.693 | 0.43             | 1              | .51              |
| Drop E moderation    | 90.244.17    | 16         | 43.693 | 6.95             | 1              | .008             |
| Drop all moderation  | 90.258.61    | 14         | 43.695 | 21.39            | 3              | <.001            |

Abbreviation: NA, not applicable.

\(^a\) \(-2LL = fit statistic, -2 \times \log likelihood of the data.

\(^b\) \(\Delta\chi^2 = -2LL\) discrepancy between models, distributed \(\chi^2\).

\(^c\) The difference in degrees of freedom between the 2 models is equivalent to the difference in number of parameters between 2 models.

\(^d\) Significant values indicate that a nested model fits statistically significantly more poorly than the model it is being compared to, supporting the statistical significance of the parameter(s) dropped from the model.

psychotic experiences. One study\(^3\) reported that the association between environmental risks and psychotic experiences increased among individuals with a family history of psychosis. Although this finding also supports gene-by-environment interaction, the results are somewhat different than what would be expected from our analyses because our analyses suggest that genetic factors were less salient in the presence of environmental exposures. Moreover, other studies\(^4,5\) have found no evidence of gene-by-environment interaction for psychotic experiences. Methodological differences may underlie these discrepancies. First, family history is not the same as genetic influence because family history includes a combination of...
genetic and shared environmental factors. Second, we focused on a composite exposure score comprising 5 environmental factors; prior studies have focused on more specific factors, including childhood physical abuse and trauma. Third, we focused on 6 specific psychotic experiences here, whereas prior studies used single measures. Finally, it is important to note that prior studies focus specifically on the interaction between genetic risk for psychotic experiences based on family history, by contrast with the current study that used the twin design. In our study, we tested whether heritability differed dependent on exposure to environmental risks. Heritability is a population statistic, and as such, an approach based on calculating heritability is somewhat different from an approach based on using family history as a proxy for individual genetic risk.

On a clinical level, it is first important to clarify that we focused on a young sample, who were aged 16 or 18 years. Many of these individuals will be too young to have been diagnosed with a psychotic disorder, but it is known that psychotic experiences in this age group can lead to severe clinical conditions in some individuals. Nonetheless, clinically it is often noted that many individuals with schizophrenia do not have a family history of schizophrenia. Indeed, although the relative risk for schizophrenia is increased among relatives of individuals diagnosed with schizophrenia, most relatives of individuals with schizophrenia do not develop schizophrenia. Our results extend these observations to adolescent psychotic experiences and indicate that they may develop in a variety of contexts. Specifically, our results suggest that psychotic experiences may be prevalent in populations with a high degree of exposure to environmental risks associated with psychotic experiences. Indeed, there is substantial conjecture that psychotic experiences can be reached through multiple pathways, such as pathways that are more based on genetic propensity and others that are more down to environmental risks; however, to our knowledge, this is one of the first studies to provide replicable empirical findings on this topic.

The previously mentioned arguments should be tempered, however, for certain types of psychotic experiences. Although we observed that the heritability differed according to environmental exposure for some psychotic experiences, the heritability was more consistent for hallucinations and negative symptoms. This is important from a clinical perspective, given that negative symptoms are thought to be particularly predictive of subsequent mental illness. As such, these results lend further weight to the argument that the etiology of psychotic experiences may differ according to specific subtypes of psychotic experience.

Strengths and Limitations

Strengths of our study included the use of 2 representative, population-based samples in different countries. In 1 sample, we measured 6 different specific psychotic experiences, enabling us to consider gene-by-environment interactions for different psychotic experiences. There were several limitations, however. Although we employed data from both the UK and Sweden, we still only focused on 2 European countries. Many of our exposures may differ in prevalence across the world, and it would therefore be useful to assess whether similar results emerge in countries with more environmental variability than the UK and Sweden. Our measures of tobacco and cannabis use were more brief than our other measures. Future studies should use more detailed measures. We also created a composite score that involved counting the number of exposures each participant had undergone, which included summing exposures that may have different etiologies or mechanisms underlying their association with psychotic experiences. Further, although our model has controlled for gene environment correlation, we recognize that birth weight is a complex phenotype influenced by parent and child genetics and prenatal environment.

It is further important to be aware that the environmental composite is not identical across TEDS and CATSS. The life events scale, for example, includes items about increasing numbers of quarrels with parents in CATSS, which are not covered in TEDS. Percentile-based cutoffs were used to define low birth weight; birth weight was lower in TEDS than CATSS on average, and therefore, this led to heavier twins being captured in CATSS. Exclusion criteria also differed between samples; individuals with extreme obstetric complications were excluded from TEDS but not CATSS. The fact that we observed similar results between CATSS and TEDS gives us confidence that these differences did not strongly influence our results; however, they should nevertheless be interpreted with these differences in mind. Finally, twins are generally born lighter than singletons. We included birth weight as an exposure here, and hence, individuals with very low birth weight may have been overrepresented in our sample. However, studying birth weight in twins here is unlikely to create any issues for generalizability for 2 reasons. First, our modeling analyses were focused on variance rather than mean differences. Second, twins were compared with twins in the design (not singletons); as such, modest mean differences between singletons and twins in birth weight did not affect the findings.

Conclusions

To our knowledge, this twin study was the first with results that suggest that environmental factors play a greater role in the etiology of psychotic experiences than genetic factors. For clinicians who may be aware that psychotic disorders are very highly heritable, it is an important message that early manifestations of psychotic experiences during adolescence are not so strongly heritable, especially in the context of higher environmental exposure. Psychotic experiences are likely to manifest in adolescents both with and without a family history of such challenges and further highlight that genetic and environmental risks for psychotic experiences do not operate in isolation from one another.
Heritability of Psychotic Experiences in Adolescents and Interaction With Environmental Risk

Original Investigation Research

Author Affiliations: Department of Medical Epidemiology and Biostatistics, Karolinska Institutet, Stockholm, Sweden (Taylor, Larsson); Department of Psychiatry, University of Oxford, Oxford, United Kingdom (Freeman); Gillberg Neuropsychiatry Centre, Centre for Ethics, Law and Mental Health, University of Gothenburg, Gothenburg, Sweden (Lundström); School of Medical Sciences, Örebro University, Örebro, Sweden (Larsson); Genes Environment Life-span Laboratory, Centre for Brain and Cognitive Development, Department of Psychological Sciences, Birkbeck University of London, London, United Kingdom (Ronald).

Author Contributions: Dr Taylor had full access to all of the data in the study and takes responsibility for the integrity of the data and the accuracy of the data analysis.

Concept and design: Taylor, Freeman, Ronald. Acquisition, analysis, or interpretation of data: All authors. Drafting of the manuscript: Taylor, Ronald. Critical revision of the manuscript for important intellectual content: All authors. Statistical analysis: Taylor. Obtained funding: Lundström, Larsson, Ronald. Administrative, technical, or material support: Lundström. Supervision: Larsson, Ronald.

Conflict of Interest Disclosures: Dr Larsson reported receiving grants from Shire/Takeda and personal fees from Shire/Takeda, Evolan, and Medici. Dr Ronal reported receiving editor honorarium from the Association for Child and Adolescent Mental Health, and grants from the Genetics Society Special interest group. No other disclosures were reported.

Funding/Support: This work was supported in part by grants MR/V01287B/1 and MR/022475/1 (Dr Ronal) and grant G1000559 (Dr Ronald) from the UK Medical Research Council, with additional support from grant AG046938 from the US National Institutes of Health.

Role of the Funder/Sponsor: The funders had no role in the design and conduct of the study; collection, management, analysis, and interpretation of the data; preparation, review, or approval of the manuscript; and decision to submit the manuscript for publication.

Disclaimer: The views expressed are those of the authors and not necessarily those of the National Health Service, the National Institute for Health Research, or the Department of Health and Social Care.

Additional Contributions: We thank the participants in the Twins Early Development Study (TEDS) and the Child and Adolescent Twin Study in Sweden (CATSS) for their ongoing participation; Robert Plomin, PhD, Social, Genetic and Developmental Psychiatry Centre, Institute of Psychiatry, Psychology and Neuroscience, Kings College London, and Paul Lichtenstein, PhD, Department of Medical Epidemiology and Biostatistics, Karolinska Institutet, Stockholm, Sweden, for allowing us to use TEDS and CATSS data in this project; and the ongoing contribution of the participants in TEDS and CATSS and their families. No one was financially compensated for their contribution.

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