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## Data-driven subphenotypic dissection of the clinical heterogeneity of schizophrenia spectrum disorders

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## CHAPTER 4

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# Long-term cognitive trajectories and heterogeneity in patients with schizophrenia and their unaffected siblings

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## **ABSTRACT**

### **Objective**

This study aimed to assess the heterogeneity and stability of cognition in patients with a non-affective psychotic disorder and their unaffected siblings. In addition, we aimed to predict the cognitive subtypes of siblings by their probands.

### **Methods**

Assessments were conducted at baseline, three and six years in 1,119 patients, 1,059 siblings and 586 controls from the Genetic Risk and Outcome of Psychosis (GROUP) study. Group-based trajectory modeling was applied to identify trajectories and clustered multinomial logistic regression analysis was used for prediction modeling. A composite score of eight neurocognitive tests was used to measure cognitive performance.

### **Results**

Five stable cognitive trajectories ranging from severely altered to high cognitive performance were identified in patients. Likewise, four stable trajectories ranging from moderately altered to high performance were found in siblings. Siblings had a higher risk of cognitive alteration when patients' alteration was mild (OR = 2.21), moderate (OR = 5.70) and severe (OR = 10.07) compared with patients with intact cognitive function. The familial correlation coefficient between pairs of index patients and their siblings was 0.27 (P=0.003).

### **Conclusions**

The cognitive profiles identified in the current study might be suitable as endophenotypes and could be used in future genetic studies and predicting functional and clinical outcomes.

**Keywords:** Schizophrenia, Psychosis, Cognition, Cognitive Impairment

## INTRODUCTION

Schizophrenia spectrum disorders consist of multiple symptom dimensions, caused by the interaction of genetic, environmental, and internal factors.[1] One of these dimensions is cognitive alteration which is present in 75-80% of the patients[2] and which has been demonstrated to be a predictor of symptomatic and functional outcome (e.g. working activity, daily living activity).[3-7] Evidence on whether cognitive alteration precedes schizophrenia or cognitive function deteriorated after the occurrence of schizophrenia is mixed.[8-10] The most affected cognitive functions are episodic and working memory, attention, verbal fluency, executive function, problem-solving, processing speed and social cognition.[11]

Cognitive function can be measured based on the actual change in patient's cognitive performance over time or deviation from the familial cognitive performance.[12] Although cognitive alteration in patients with schizophrenia is heterogeneous, the majority of the studies shows that cognitive alteration within an individual with schizophrenia is stable and persistent over time.[13-15] On the other hand, in patients with first-episode psychosis, studies investigating the trajectories of different cognitive performance suggested an improvement over time.[16,17] Given the substantial difference across patients, clustering methods using latent class analysis, hierarchical mixture model and group-based trajectory modeling have been used to identify cognitive trajectories in a different group of population.[18] To date, two to five cognitive trajectories have been identified in patients with psychosis who differ in terms of demographic and clinical characteristics.[19-23] Cognitive alteration is also common in unaffected siblings, but to a lesser extent, and subtyping is relevant to identify meaningful homogeneous trajectories.[13,24-28] In our previous cross-sectional study, we demonstrated three subtypes of cognitive alteration (i.e., normal, mixed and impaired) in siblings.[23]

Cognitive alteration in patients with schizophrenia can be associated with cannabis use[29], negative and schizotypy symptoms[30,31] and medical comorbidities such as diabetes mellitus, hypertension, dyslipidemia, and obesity.[32] Other influencing factors include decreased gray matter density in the cerebellum.[33] reduced blood omega-3 fatty acids,[34] N-methyl-d-Aspartate receptor (NMDAR) hypofunction[35] and inflammation.[36] In siblings, cognitive alteration can be attributed to shared genetic [37-39] and socio-environmental risk factors with the patients.[40] Cognitive

alteration can be considered an inheritable trait of schizophrenia, which itself has a heritability ( $h^2$ ) 31 to 62%.<sup>[41-43]</sup> Some family-based studies have also found a higher rate of cognitive alteration in unaffected relatives of individuals with schizophrenia than in the general population.<sup>[44]</sup> These findings support the assertion that cognitive dysfunction can be designated an important endophenotype of schizophrenia. This is important for future large-scale twin/family-based and SNP-based heritability studies.

Schizophrenia is invariably acknowledged to be an extremely heterogeneous disorder. Yet, efforts to find meaningful subtypes (cognitive, clinical, or neurobiological) have been manifold but they remain of questionable success. Possible explanations for the unsuccessful subtyping are use of single cognitive test or domain (e.g. intelligence quotient (IQ) based on mean values that do not take into account heterogeneity, uncertainty on whether cognitive alteration is general or specific, limited number of longitudinal studies with short follow-up period, small sample size, and limited use of data from healthy siblings and controls.<sup>[40,45,46]</sup> Therefore, to our knowledge, this is the first large-scale long-term study which included cognitive trajectory modeling for both patients and their unaffected siblings.

The aim of this study was twofold. Firstly, we examined ways to unravel the heterogeneity of neuro-cognition, by classifying patients and siblings respectively into different subgroups based on the course of composite overall cognition scores over time, separately from each other, based on eight neurocognitive test parameters. After determining meaningful subgroups of patients and siblings, we aimed to predict the cognitive subtypes of siblings by subtypes of patients within a family using sibling-patient analyses. Given the advancement of clustering technique and increasing trend of identified subgroups of patients with schizophrenia<sup>[19-23]</sup>, we hypothesized that more meaningful trajectories can be discovered. There has been a substantial controversy regarding the course of cognitive alteration in patients with schizophrenia and their unaffected siblings.<sup>[40]</sup> The accumulated body of evidence<sup>[40]</sup> shows that the course of cognitive function in schizophrenia is inconsistent while some studies show stability or improvement, others support deterioration of cognitive function. In this study, we hypothesized that the cognitive trajectory over time is stable. In addition, we hypothesized the probands cognitive trajectory predict siblings cognitive alteration trajectory.

## METHODS

### ***Study setting and population***

The current study was performed within the framework of the Genetic Risk and Outcome of Psychosis (GROUP) project, a longitudinal multi-center cohort study in the Netherlands and Belgium. Patients with non-affective psychotic disorder based on DSM-IV[47], age between 16 and 50 years and good command of the Dutch language were included. Siblings were included if they (i) were between 16 and 50 years, (ii) had a good command of the Dutch language, and (iii) had no lifetime psychotic disorder. Controls were included if they did not have a lifetime diagnosis of psychotic, bipolar or current depressive disorder. In addition, controls were included if they did not have a first or second degree relative with psychotic disorder.[48] The procedure of recruitment, informed consent, ethical approval and population characteristics at baseline have been described in detail elsewhere.[49]

### ***Sample size calculation***

There was no formal sample size calculation for trajectory modeling analysis. However, Formann estimated that the minimum sample size for latent class analysis is  $2^k$  subjects, where  $k$  is the number of variables.[50] Therefore, we need at least  $2^k = 2^8 = 256$  subjects in our study, which we satisfy by large. To establish a power of 93% ( $= 0.05$ ) in a case-control and a casesibling longitudinal study design (standard deviation = 0.15) on cognitive function, at least 1000 patients, 1000 siblings, and 350 controls are needed. In the current study, 1,119 patients, 1,059 siblings and 586 healthy controls were participated at baseline, three-years, and six-years follow-up. [49]

### ***Assessment of neurocognition***

Task selection was based on cognitive domains that have been shown to be altered in schizophrenia.[51] The cognitive battery has been described in detail elsewhere. [28] This study assessed neurocognitive performance using eight assessment tools. Table 1 shows the list of cognitive domains, their corresponding tests, and outcome measures. We calculated a composite neurocognitive score based on the following eight neurocognitive measures: CPT performance, CPT variance, immediate recall, delayed recall, digit symbol substitution, information, arithmetic, and block design.

Subsequently, linear regression analyses were conducted for each time point. The control subjects score was used to obtain age- and gender-stratified neurocognitive tests z-score for both patients and siblings. Finally, the composite score for patients and siblings was computed by averaging z-score of all eight tests.

**Table 1:** Measures of neurocognition

<b>Cognitive domains</b>	<b>Group Tests</b>	<b>Outcome measure</b>
Sustained attention and vigilance	Continuous Performance Test (CPT-HQ) (CPT performance and CPT variance)	An efficiency score [(accuracy/ reaction time) ×1000] was created, in which accuracy was measured as the total number of hits (range 0–28) minus the total number of errors (range 0–28), divided by 28. If this calculation of accuracy was non-positive (i.e. the number of errors equaled or exceeded the number of hits), then the accuracy was set equal to 0.005. This score was referred to as 'CPT performance'. Intra-individual variability in reaction time on the CPT was also evaluated (CPT variance), using the standard deviation score of the subject's mean response time on the hit trials. [23,52]
Verbal learning and memory	Word Learning Task (WLT) (Immediate and Delayed Recall)	Immediate recall (total score of three consecutive trials of 15 words learning) and 'Delayed recall' was assessed after 20 minutes delay.
Global cognitive functioning		
Processing speed	Digit Symbol Substitution	Total raw score (0-133)
Verbal comprehension	Information	Total raw score (0-28)
Working memory	Arithmetic	Total raw score (0-22)
Problem-solving and Visuospatial abilities	Block Design	Total raw score (0-68)

### ***Assessments of socio-demographic and clinical variables***

The socio-demographic variables were age of participants, age of onset of psychosis educational status, gender, and ethnicity. According to Verhage,[53] educational

status was evaluated as a continuous variable. Psychotic disorder was assessed by the Comprehensive Assessment of Symptoms and History (CASH)[54] or Schedules for Clinical Assessment in Neuropsychiatry (SCAN) questionnaire.[55,56] Level of premorbid functioning was assessed using the Premorbid Adjustment Scale (PAS).[57] Positive and negative symptoms of schizotypy were measured using the Structured Inventory for Schizotypy-Revised (SIS-R).[58] The SIS-R is a reliable and valid semi-structured interview for rating schizotypal sign and symptoms administered in unaffected siblings and healthy controls.[58] Frequency and distress of psychotic experiences were measured with the Community Assessment of Psychic Experiences (CAPE) questionnaire.[59] The CAPE is a 42-item reliable and valid self-report questionnaire to assess psychotic-like experiences in the community and in individuals at an increased risk for developing psychosis (i.e. siblings in our study) and in healthy controls.[59] Symptom severity in patients was assessed with the Positive And Negative Syndrome Scale (PANSS).[60,61]

## ***Data analysis and statistical modeling***

### *Descriptive statistics*

Patients, siblings, and controls difference in socio-demographic characteristics at baseline were tested using univariate analyses. For gender and ethnicity, Pearson's Chi-square test was used. Due to the family structure data, linear mixed effects models using family as a random effect were applied to all continuous variables (e.g. socio-demographic, clinical and cognitive tests) to test for differences between the groups. Maximum Likelihood (ML) method was used to estimate the model parameters and Type-III (overall) tests of fixed effects were used to test for differences between groups. Significant differences were followed by pair-wise post-hoc comparisons between groups. Additionally, Pearson's correlation coefficients between cognitive performances were computed for patients and siblings to explore the possible predictive relationship.

### *Subtyping and trajectory modeling*

Subtyping is the level of overall cognitive functioning across time whereas trajectory is the pattern (e.g. stable, fluctuating, decline, or improvement) of overall cognitive functioning or the specific cognitive domain/test. Group-based trajectory modeling (GBTM)[62-65] was conducted to identify clusters of patients and siblings with a

similar pattern of neurocognitive function over time. The neurocognitive composite score was used as a dependent variable and follow-up time (baseline, three and six years) as an independent variable. First order linear and second order quadratic polynomial models were fitted assuming that individual differences in trajectories could be summarized by a finite set of polynomial functions of time. To determine the number of clusters and the best-fitting model, a sequential approach was applied where the number of clusters is increased by one. The less complex model (i.e. less trajectory groups) was compared with the complex model (i.e. more groups) using the Bayesian Information Criterion (BIC)[66] and logged Bayes factor ( $2 \cdot \Delta\text{BIC}$ ), where  $\Delta\text{BIC} = \text{BIC}(\text{complex}) - \text{BIC}(\text{less complex})$ [67,68] and where the logged Bayes factor ( $2 \cdot \Delta\text{BIC}$ ) would indicate trivial (0-2), positive (2-6), strong (6-10) or very strong (>10) evidence for the null hypothesis that the less complex model is the best fit.

First, a single quadratic polynomial trajectory model was examined. If the quadratic term was not significant, the model was repeated with a linear trajectory to determine the BIC value. If the quadratic component in one trajectory model was significant, the quadratic two-trajectory model was performed. Next, the BIC value of the appropriate two-trajectory model was compared to the BIC value of the appropriate one-trajectory model. The process was repeated by increasing the number of trajectories until the best-fit model was found.[69] The trajectory modeling was stopped when the  $\Delta\text{BIC}$  became a negative value. GBTM can handle missing data in the cognition score using ML estimation method to create asymptotically unbiased parameter estimates under the assumption of data missingness at random.[70,71] We also checked the drop-out model which includes a logistic model of drop-out probability per period to interrogate whether the attrition rates significantly affected/biased the cognitive trajectory group membership probabilities.[72] For each group, 0 = constant attrition rate, 1 = depends on the previous response (i.e. baseline) and 2 = depends on the two previous responses (i.e. baseline and three years).

### *Comparison among cognitive subtypes*

Differences between the subgroups of patients and siblings in all continuous variables were investigated using linear mixed effect models taking familial relationship into account. Type-III (overall) tests of fixed effects were used to test group differences. If these were significant, pair-wise comparison was done using Dunnett's method (taking the most normal cognitive profile as a reference group). For gender and

ethnicity, the Pearson's Chi-square test was used to test the difference between subgroups of patients and siblings.

### *Predicting cognitive subtypes of siblings using subtypes of probands*

We considered sibling subtype (multi-category) as a dependent variable and patient subtype (multi-category) as an independent variable. The probability of concordance, probability of discordance and Somers' D statistic[73] were computed on the pairs of subtypes of patients and siblings. A pair of patients-siblings subtype was reported concordant if the larger value of subtypes of patients was paired with the larger value of subtypes of siblings, and discordant if the larger value of subtypes of patients was paired with the smaller value of subtypes of siblings. Somers' D of subtypes of siblings, with respect to subtypes of patients, was defined as the difference between the two conditional probabilities of concordance and discordance.

Clustered multinomial logistic regression analysis was conducted taking into account family as a random effect given that siblings and patients belong within the same family. PROC NLMIXED in SAS (Statistical Analysis System) was applied to determine the predictive relationship between subtypes of patients and siblings. An adaptive Gaussian quadrature with ten quadrature points was specified to integrate out the random effect of the likelihood function[74,75] and to estimate the parameters (i.e. subtypes of patients) and their standard errors. The intra-cluster correlation coefficient (ICC) was calculated to estimate the familial correlation between pairs of unaffected siblings and probands in the same family. The ICC was calculated as  $ICC_{\text{Family}} = \text{var}(\text{family}) / (\text{var}(\text{family}) + \pi^2 / 3)$ , where  $\text{var}(\text{family})$  is the variance of random effect and  $\pi$  is 3.14159. A two-tailed test at  $p < 0.05$  was considered as statistical significant throughout the analyses. For pair-wise comparisons, Bonferroni correction for multiple testing was done and statistical significance was determined at  $p < 0.05/20$ . All analyses were performed using SAS version 9.4. In this study, GROUP data version 4.00 was used.

## RESULTS

### *Descriptive of the study population*

At baseline, patients, siblings, and controls significantly ( $P < 0.001$ ) differed by socio-demographic, cognitive and clinical variables. Pair-wise comparisons revealed that the differences were significant between patients and controls as well as siblings and controls. Patients' and siblings' cognitive performance was low compared to controls. Additionally, cognitive performances of patients were found to be significantly lower than the performances of their unaffected siblings (Table 2). Correlation between all cognitive tests was significant in patients and siblings except the correlation between CPT performance and Block Design, Arithmetic and Information in siblings (Supplementary Table S1-S2).

**Table 2:** Comparison of control, sibling, and patient baseline profiles\*

Variable/Group	Group			Overall group difference	Pair-wise group comparison
	Controls (n=586)	Siblings (n=1,059)	Patients (n=1,119)		
Age	30.42 (10.58)	27.84 (8.28)	27.58 (7.94)	F=22.8, P<0.001	Sibs<Ctrs, Pts<Ctrs
Age of onset	...	...	23.69 (7.59)	...	...
Gender, male	45.90	45.51	76.14	$\chi^2=254.1$ , P<0.001	Sibs<Ctrs <Pts
Education (Verhage) <sup>a</sup>	5.41 (1.78)	5.07 (2.11)	4.04 (2.05)	F=128.5, P<0.001	Pts<Sbs<Ctrs
Ethnicity, Dutch	92.12	83.24	79.22	$\chi^2=45.2$ , P<0.001	Pts<Sbs<Ctrs
Estimated IQ <sup>b</sup>	109.75 (15.08)	102.76 (15.60)	94.99 (16.12)	F=185.6, P<0.001	Pts<Sbs<Ctrs
Overall score PAS <sup>c</sup>	1.13 (0.59)	1.13 (0.66)	1.98 (0.88)	F=439.0, P<0.001	Sbs<Pts, Pts<Ctrs
SIS-R <sup>d</sup>					
Positive	0.31 (0.35)	0.38 (0.42)	...	F=15.4, P<0.001	Ctrs<Sbs
Negative	0.24 (0.22)	0.27 (0.26)	...	F=9.0, P=0.002	Ctrs<Sbs

**Table 2:** Continued.

Variable/Group	Group Controls (n=586)	Siblings (n=1,059)	Patients (n=1,119)	Overall group difference	Pair-wise group comparison
PANSS 5-factor					
Positive	...	...	13.90 (6.55)	...	...
Negative	...	...	15.00 (6.64)	...	...
Disorganization	...	...	16.77 (6.27)	...	...
Excitement	...	...	12.05 (4.05)	...	...
Emotional distress	...	...	15.82 (5.73)	...	...
CAPE (Positive dimension) <sup>e</sup>					
PE Frequency	0.19 (0.17)	0.21 (0.20)	0.67 (0.49)	F=566.5, P<0.001	Sbs<Pts, Pts<Ctrs
PE Distress	0.43 (0.45)	0.46 (0.48)	1.26 (0.69)	F=531.0, P<0.001	Sbs<Pts, Pts<Ctrs
Cognitive Performance					
CPT performance <sup>f</sup>	246.36 (54.78)	243.70 (57.86)	220.82 (62.14)	F=52.0, P<0.001	Pts<Sbs, Ctrs<Pts
CPT variance (ms) <sup>g</sup>	72.76 (28.28)	75.80 (28.44)	92.99 (36.51)	F=108.4, P<0.001	Sbs<Pts, Pts<Ctrs
Block Design <sup>h</sup>	46.55 (14.16)	44.87 (15.07)	40.42 (16.99)	F=39.2, P<0.001	Pts<Sbs<Ctrs
Digit Symbol <sup>i</sup>	84.01 (14.58)	79.21 (15.39)	65.41(16.27)	F=379.9, P<0.001	Pts<Sbs<Ctrs
Arithmetic <sup>j</sup>	15.32 (4.16)	13.86 (4.43)	12.27 (4.79)	F=93.1, P<0.001	Pts<Sbs<Ctrs
Information <sup>k</sup>	18.84 (4.67)	16.83 (5.22)	16.77 (5.47)	F=31.5, P<0.001	Sbs<Ctrs, Pts<Ctrs
Immediate Recall <sup>l</sup>	28.47 (5.37)	26.89 (5.77)	22.94 (6.09)	F=224.5, P<0.001	Pts<Sbs<Ctrs
Delayed Recall <sup>m</sup>	9.74 (2.70)	9.33 (2.64)	7.53 (2.87)	F=181.6, P<0.001	Pts<Sbs<Ctrs

*\*Table presents means (SD) or %; Empty (..) row or column means no measurements in the respective group or variable; <sup>a</sup>Education (Verhage): range 0 (no education), 3-5 (school diploma) to 8 (university degree); <sup>b</sup>IQ: Wechsler Adult Intelligence Scale-III (WAIS-III), short form; <sup>c</sup>PAS: Premorbid Adjustment Scale; <sup>d</sup>SIS-R: Structured Inventory for Schizotypy - Revised; <sup>e</sup>CAPE: Community Assessment for Psychic Experiences; PE frequency and distress: Frequency of positive psychotic experiences and amount of distress of these PE; <sup>f</sup>CPT performance: Continuous Performance Test HQ, performance index; <sup>g</sup>CPT variance (ms): CPT-HQ variance in reaction time (ms); <sup>h</sup>Block Design: WAIS-III Block Design; <sup>i</sup>Digit Symbol: WAIS-III Digit Symbol Substitution Test; <sup>j</sup>Arithmetic: WAIS-III Arithmetic; <sup>k</sup>Information: WAIS-III Information; <sup>l</sup>Immediate recall: Word Learning Task (WLT) immediate recall; <sup>m</sup>Delayed recall: WLT Delayed recall. For the PAS, higher scores reflect poorer premorbid adjustment. <sup>1</sup>Control, <sup>2</sup>Sibling and <sup>3</sup>Patient; Pair-wise group comparison explains which group has significantly different values on the measurements; <sup>†</sup>p < 0.05/20 was considered as statistically significant.*

### ***Cognitive subtypes and trajectories***

A five-group cognitive trajectory model for patients and four-group trajectory model for siblings were very strongly favoured for overall cognitive scores (z-scores) according to the smallest BIC and the logged Bayes factor ([Supplementary Table S3](#)). In patients, the value of logged Bayes factor was 17.34 (>10), thus favouring the five-cluster model over a six-cluster model. In siblings, the value of logged Bayes factor was 94.62 (>10), favouring the four-trajectory model over a five-cluster model ([Supplementary Table S3](#)). Parameter estimates of linear and quadratic polynomial time functions of trajectory modeling and drop-out model presented in [Supplementary Table S4-S5](#). The drop-out of patients and unaffected siblings did not significantly affect/biased their cognitive trajectory group membership probabilities.

Figures 1a and 1b displays cognitive trajectories of patients and their siblings over six years respectively. The cognitive trajectories in patients were labeled as 'severely altered', 'moderately altered', 'mildly altered', 'normal' and 'high performer'. Similarly, the trajectories in siblings were labeled as 'moderately altered', 'mildly altered', 'normal' and 'high performer'. All initial neurocognitive profiles for patients and siblings were stable over time. Severe and moderate groups were identified based on a broad-based cognitive alteration of, on average, about 1 SD below the normal cognitive trajectory across a range of cognition composite score. We showed 26.7% of patients had a normal cognitive function, 30.4% mild, 28.4% moderate and 10.7% severe cognitive alterations. Likewise, 37.6% of siblings showed a normal cognitive functioning, 25.1% mild and 13.0% moderate cognitive alteration. Three point eight percent of patients and 24.2% of siblings had higher cognitive functioning compared with the mean level of the normal cognitive subtype. Patients in this cluster showed a subtle significant decline over time (Figure 1a) whereas the pattern was stable for siblings (Figure 1b).

### ***Comparison of cognitive subtypes on all baseline characteristics***

Cognitive subtypes of patients were compared based on cognitive measures, demographic characteristics, and clinical variables. Patients with normal cognitive profile differed significantly from other patient subgroups in education, IQ, premorbid functioning and psychotic symptoms. The detailed pair-wise comparisons on cognitive subtypes of patients have been shown in Table 3.

Differences between normal cognitive subtype and other subtypes of siblings were statistically significant with regard to age, gender, education, ethnicity, IQ, premorbid functioning, positive symptoms, frequency of psychotic experiences and all cognitive performances except CPT variance. The detailed pair-wise comparisons on cognitive subtypes of siblings have been shown in Table 4.

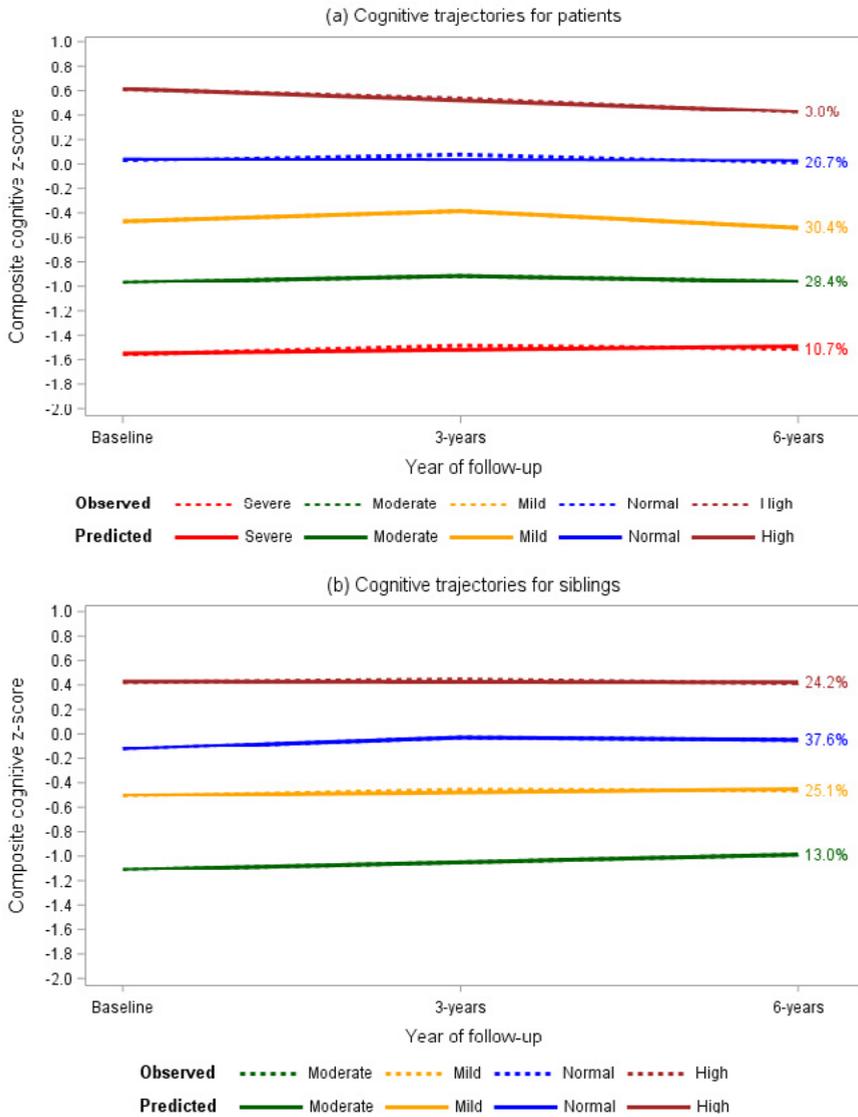
### ***Association of cognitive subtypes of patients and siblings***

In the sibling-patient pair analysis, we generated 1,070 pairs of affected and unaffected siblings. The number of pairs was more than 1,059 because we paired multiple unaffected siblings with their single affected sibling or multiple affected siblings with their single unaffected sibling within a family. The contingency table of the subtypes of patients and siblings is presented in the [Supplementary Table S6](#). Somers' *D* value for the association between cognitive subtypes of patients and subtypes of siblings amongst the 1,070 sib-pairs was 0.29. A positive value of Somers' *D* indicates that the siblings have better cognitive scores than their probands.

Since the cell frequency of moderately altered subtype in the sibling coupled with high cognitive performance in the patient was zero ([Supplementary Table S6](#)), we combined the high cognitive performer group of patients with the normal cognitive group. The combined normal and high performer group of patients was considered as the reference group.

[Supplementary Table S7](#) presents the association between cognitive subtypes of patients and siblings. Overall, cognitive subtypes of patients significantly predicted the siblings' subtypes. Here, we present a risk of an unaffected sibling to be grouped in any of mildly altered, moderately altered and severely altered groups given the cognitive trajectory of their corresponding affected sibling. The familial correlation was 0.27 at a significance level of  $P = 0.003$ .

We observed that unaffected sibling with a severely and moderately altered patients was at risk of having mild alterations with an odds ratio (OR) of 2.56 (95% CI 1.26 - 5.18) and 1.83 (95% CI 1.12 - 2.98) respectively. However, mild cognitive alterations in patients did not predict mild alterations in their unaffected siblings (OR = 0.86,  $P=0.54$ ). Compared with siblings of patients with intact cognitive function, siblings were at risk of moderate alteration if their probands cognitive alteration was mild (OR = 2.21, 95% CI 1.05 - 4.64), moderate (OR = 5.70, 95% CI 2.77 - 11.70) or severe (OR



**Figure 1:** (a) Cognitive trajectories for patients (n=1,119), (b) Cognitive trajectories for siblings (n=1,059)

= 10.07, 95% CI 4.15 - 24.44). Subsequently, siblings had a low likelihood of having high cognitive performance if patients were severely (OR = 0.26, 95% CI 0.09 - 0.63) or moderately (OR = 0.39, 95% CI 0.24 - 0.64) altered and a low likelihood of mild

cognitive alteration (OR = 0.37, 95% CI 0.24 - 0.59) when the probands performance was normal or high. In general, unaffected siblings of patients with severe alteration were likely to develop moderate to mild cognitive alterations.

## DISCUSSION

To our knowledge, this is the first large-scale study to report cognitive trajectories in patients with non-affective psychosis (probands) and their unaffected siblings. We identified five stable and persistent cognitive trajectories within the proband group: 'severely altered', 'moderately altered', 'mildly altered', 'normal' and 'high performer'. Similarly, we identified four stable and persistent trajectories among unaffected siblings: 'moderately altered', 'mildly altered', 'normal' and 'high performer'. The normal subgroup differed significantly from the other subgroups in socio-demographic, symptomatic, and cognitive profiles. Furthermore, the cognitive trajectory of probands significantly predicted the cognitive trajectory of siblings, the intraclass correlation (ICC) between pairs of siblings and probands was 0.27.

In this study, the trajectory modeling demonstrated five cognitive subgroups for patients and four cognitive subgroups for siblings, which confirms the presence of distinct subgroups defined by cognitive functioning and provides further support for cognitive heterogeneity within psychosis patients and healthy siblings. Similarly, previous cluster analysis studies have reported two to five distinct subgroups in patients [19-23] and three subgroups in healthy siblings. [23] In agreement with previous studies [76-78], approximately 69.5% of patients had poor cognitive performance (i.e. severe, moderate, and mild alteration level). On the other hand, 30.5% of patients had an intact cognitive function. This finding is higher than previous studies that reported 19-28% of patients with psychosis found to have an intact cognitive function. [20,79-82] This discrepancy may be due to the difference in duration of follow-up, cognitive assessment tools, use of composite score and level of cognitive function at baseline.

Similar to our earlier cross-sectional study [23], a total of 38% siblings exhibited a stable lower cognitive performance (i.e. both moderate and mild level) over the six years. Siblings in the moderate altered group were more often an ethnic minority, young, less intelligent and with high psychotic experiences as supported by the previous review. [40] In line with the previous study [83], we depicted that patients and siblings with mild to severe cognitive alteration performed at least 1 SD below

normal cognition. In accordance with our hypothesis and previous studies report [40,84,85], we also found cognitive trajectories in patients and their unaffected siblings were stable over the six years follow-up period. This is different from other previous literature [16,17,40], which may be related to the use of a composite score that obliterated the change occurred in individual cognitive domain. Negative and positive symptoms in patients with moderately and severely altered cognition were significantly higher than in the mildly altered and intact subgroups. This finding was consistent with previous studies and supports the notion that psychotic symptoms have been strongly associated with cognitive alteration.[19,20,30,31,86] In line with previous findings[19,20,87], IQ was significantly lower in patients with mild to severe cognitive alteration compared with patients with intact cognitive function.

The positive Somers' *D* value from the sib-pair analysis showed that the proband is more likely to show low cognitive performance than the sibling, which implies the illness itself, medication and motivation may have an impact on cognitive performance. Congruent with our hypothesis, the cognitive subtypes of the patients significantly predicted the cognitive subtypes of siblings within the family. The poorer the cognitive profile of the patient, the better it predicted the profile of a more cognitively altered sibling. Moreover, siblings with a high-performance profile are less predicted when their probands are severely altered. We found a familial correlation of 27% for cognitive function, which is high compared with other complex diseases such as depression or bipolar disorder and can be used in clinical practice. Of note, this correlation represents the familial risk (which involves shared genetic and familial environment), as sibling designs do not allow to estimate the extent of the genetic and shared environmental contribution separately. As is evident from the literature, the neurocognitive alteration is heritable and genetically correlated with schizophrenia.[43,88,89] Indeed, Husted and colleagues showed that neurocognitive traits are highly heritable in patients with known familial schizophrenia ( $h^2= 31$  to 62%). [43]

Using a large patient sample data, we uncovered five cognition trajectories after 10 years, which signifies a high degree of heterogeneity is still evident within non-affective psychosis patients and warrants further investigation. In line with previous studies,[23,42,90] we demonstrated that cognitive performance of siblings was between those of patients and controls indicating a parallel between cognitive performance and familial liability. Subtyping cognitive profile of unaffected siblings

**Table 3:** Patients cognitive trajectory group profiles (n=1,119)<sup>1</sup>

Profiles	Cognitive trajectory group					Overall trajectory group difference	Pair-wise comparison <sup>2</sup>
	1. High (n=31)	2. Normal (n=290)	3. Mild (n=377)	4. Moderate (n=312)	5. Severe (n=109)		
Age	31.4 (8.3)	27.3 (7.2)	28.0 (8.0)	27.3 (8.4)	26.8 (8.0)	F=2.5, P=0.051	2<1
Age of Onset	26.2 (8.1)	23.0 (6.5)	23.2 (7.9)	22.9 (7.9)	23.0 (7.3)	F=14, P=0.253	
Gender, % male	74.2	77.9	74.3	77.2	75.2	$\chi^2=1.6$ , P=0.816	
Education (Verhage) <sup>3</sup>	5.8 (1.9)	4.9 (1.8)	4.3 (2.0)	3.3 (1.9)	2.6 (1.8)	F=49.0, P<0.001	2<1, 2>3,4,5
Ethnicity, % Dutch	90.3	87.2	79.9	75.1	64.5	$\chi^2=30.6$ , P<0.001	2<1, 2>3,4,5
IQ, Estimated <sup>4</sup>	127.8 (10.7)	110.5 (10.4)	95.7 (8.6)	83.3 (7.7)	72.9 (6.7)	F=615.6, P<0.001	2<1,2>3,4,5
PAS, overall score <sup>5</sup>	1.8 (1.0)	1.8 (0.8)	1.9 (0.8)	2.1 (1.0)	2.3 (0.8)	F=9.7, P<0.001	2<4,5
<b>PANSS 5-factor<sup>4</sup></b>							
Positive	14.0 (8.2)	13.4 (5.9)	13.1 (5.9)	15.0 (7.1)	14.8 (7.7)	F=4.0, P=0.003	2<4
Negative	15.1 (6.4)	13.6 (5.9)	14.3 (6.4)	16.1 (7.0)	18.0 (7.1)	F=11.1, P<0.001	2<4,5
Disorganization	16.0 (6.8)	14.8 (4.9)	15.7 (5.6)	18.3 (6.6)	21.3 (7.1)	F=30.5, P<0.001	2<4,5
Excitement	12.8 (5.2)	11.5 (3.5)	11.5 (3.7)	12.7 (4.4)	13.2 (4.5)	F=7.0, P=0.001	2<4,5
Emotional distress	16.0 (5.7)	15.4 (5.4)	15.2 (5.5)	16.7 (5.8)	16.4 (6.8)	F=3.3, P=0.024	2<4
<b>Cognitive Performance</b>							
CPT performance <sup>6</sup>	256.1 (45.7)	240.9 (52.2)	225.8 (59.6)	210.7 (56.5)	170.0 (78.8)	F=30.1, P<0.001	2>3,4,5
CPT variance (ms) <sup>7</sup>	84.2 (35.3)	83.1 (31.9)	90.1 (36.0)	101.8 (37.2)	106.3 (38.9)	F=13.4, P<0.001	2<4,5
Block Design <sup>8</sup>	60.4 (8.8)	53.5 (10.9)	42.2 (14.3)	30.7 (13.8)	20.6 (11.0)	F=197.6, P<0.001	2<1, 2>3,4,5

Table 3: Continued.

Profiles	Cognitive trajectory group					Overall trajectory group difference	Pair-wise comparison <sup>**</sup>
	1. High (n=31)	2. Normal (n=290)	3. Mild (n=377)	4. Moderate (n=312)	5. Severe (n=109)		
Digit Symbol <sup>b</sup>	85.2 (14.6)	76.2 (14.6)	66.5 (12.5)	57.8 (12.9)	47.5 (11.7)	F=138.7, P<0.001	2<1; 2>3,4,5
Arithmetic <sup>c</sup>	18.8 (2.5)	16.2 (3.2)	13.0 (3.8)	9.1 (3.3)	6.7 (2.7)	F=264.22, P<0.001	2<1; 2>3,4,5
Information <sup>d</sup>	24.6 (2.3)	21.1 (3.4)	17.4 (4.3)	13.6 (4.3)	9.9 (3.5)	F=243.9, P<0.001	2<1; 2>3,4,5
Immediate Recall <sup>e</sup>	32.4 (3.8)	27.1 (4.7)	23.4 (4.7)	19.9 (4.8)	15.9 (5.0)	F=173.2, P<0.001	2<1; 2>3,4,5
Delayed Recall <sup>f</sup>	12.1 (1.8)	9.4 (2.4)	7.6 (2.4)	6.2 (2.2)	4.6 (2.0)	F=142.7, P<0.001	2<1; 2>3,4,5

<sup>\*\*</sup>Table presents mean (SD standard deviation) or %; <sup>\*\*</sup>Only significantly different trajectory groups are reported based on Dunnett's adjustment. The reference group is normal trajectory group; <sup>a</sup>Education (Verhage): range 0 (no education), 3-5 (school diploma) to 8 (university degree); <sup>b</sup>IQ: Wechsler Adult Intelligence Scale-III (WAIS-III), short form; <sup>c</sup>PAS: Premorbid Adjustment Scale; <sup>d</sup>PANSS: Positive and Negative Syndrome Scale (higher scores in PAS and PANSS reflect poorer outcomes); <sup>e</sup>CPT performance: Continuous Performance Test HQ, performance index; <sup>f</sup>CPT variance (ms): CPT-HQ variance in reaction time (ms); <sup>g</sup>Block Design: WAIS-III Block Design; <sup>h</sup>Digit Symbol: WAIS-III Digit Symbol Substitution Test; <sup>i</sup>Arithmetic: WAIS-III Arithmetic; <sup>j</sup>Information: WAIS-III Information; <sup>k</sup>Immediate recall: Word Learning Task (WLT) immediate recall; <sup>l</sup>Delayed recall: WLT Delayed recall; Cognitive trajectory: high performance (1) to severely altered (5); <sup>†</sup>p < 0.05/20 was considered as statistically significant.

may also provide insight to identify familial groups with high-risk for developing psychosis and initiate preventive strategies targeting healthy siblings. Although not investigated in the present study, adding genetic markers may unravel additional cognitive subtypes given that cognitive traits are heritable. Neurocognitive measures have been proposed as one of the reliable endophenotypic markers of liability for schizophrenia, as cognitive deficits are transmitted within families of a patient with schizophrenia.[41,42] Therefore, the subtyping of cognitive trajectories in our study may provide insight for determining possible endophenotypic markers associated with schizophrenia and future genetic analyses. Identifying distinctive cognition subtypes may also be helpful to predict variation in clinical and functional outcomes. In addition, identifying meaningful subtypes lends support to develop group-based diagnostic criteria for cognitive alteration. Moreover, given that not all patients or siblings develop cognitive alteration (30.5% patients versus 61.8% siblings), identifying subgroups of individuals is helpful for personalized medicine to provide individualized evidence-based interventions rather than "one-size-fits-all" approach, such as cognitive remediation therapy, cognitive adaption training, and medication therapy to improve cognitive performance.[19] In this study, we proved that about one-third of siblings cognitive trajectory predicted by patients cognitive trajectory which can be applied in clinical utility on an individual dyad level though further investigation is required on genetic, sociodemographic and environmental factors that affect cognitive performance. Furthermore, the present study showed that clinical characteristics (i.e. subclinical psychotic experiences in siblings and psychotic symptoms in patients) and premorbid functioning were poor in moderately to severely altered subgroups, which shades light to understand the path ophysiologic mechanisms of cognitive alteration in schizophrenia.

Cognitive performance of a large number of patients with non-affective psychosis, their unaffected siblings, and healthy controls was followed for six years using a comprehensive neurocognitive battery, which can be useful to ensure the clinical and cognitive validity of trajectories. We included patients and more than one siblings within the same family to investigate the joint cognitive trajectory of patients and siblings. Moreover, the study provided considerable evidence that siblings' cognitive alteration may be predicted by the different cognitive profiles of their probands. A methodological strength of the current study was that we used drop-out modeling within the same model as the group-based trajectory modeling [71] to identify the

**Table 4:** Siblings cognitive trajectory group profiles (n=1,059)<sup>a</sup>

Profiles	Cognitive trajectory group				Overall trajectory group difference	Pair-wise comparison <sup>b</sup>
	1. High (n=254)	2. Normal (n=413)	3. Mild (n=260)	4. Moderate (n=132)		
Age	28.6 (7.7)	27.9 (8.0)	27.8 (8.9)	26.3 (8.9)	F=5.6, P=0.001	2<1
Gender, % male	55.5	43.6	41.5	40.2	$\chi^2=14.1$ , P=0.003	2<1, 2>4
Education (Verhage) <sup>a</sup>	6.1 (1.8)	5.3 (2.0)	4.5 (2.0)	3.5 (2.1)	F=62.2, P<0.001	2<1, 2>3, 4
Ethnicity, % Dutch	87.4	85.7	79.5	74.8	$\chi^2=14.1$ , P=0.003	2<1, 2>3, 4
IQ, Estimated <sup>b</sup>	120.7 (9.9)	104.9 (9.5)	92.1 (7.9)	81.9 (6.7)	F=684.7, P<0.001	2<1, 2>3, 4
PAS, overall score <sup>c</sup>	0.9 (0.6)	1.1 (0.7)	1.2 (0.6)	1.3 (0.7)	F=13.6, P<0.001	2<4, 2>1
<b>SIS-R<sup>d</sup></b>						
Positive	0.3 (0.4)	0.4 (0.4)	0.4 (0.4)	0.5 (0.5)	F=4.2, P=0.006	2<4
Negative	0.3 (0.3)	0.3 (0.2)	0.3 (0.3)	0.3 (0.3)	F=2.5, P=0.108	
<b>CAPE (Positive dimension)<sup>e</sup></b>						
PE Frequency	0.2 (0.2)	0.2 (0.2)	0.2 (0.2)	0.3 (0.3)	F=4.3, P=0.006	2<4
PE Distress	0.4 (0.4)	0.5 (0.5)	0.5 (0.5)	0.6 (0.5)	F=2.4, P=0.074	
<b>Cognitive Performance</b>						
CPT performance <sup>f</sup>	254.9 (58.9)	245.3 (49.8)	242.4 (59.7)	220.9 (68.3)	F=9.3, P<0.001	2>4
CPT variance (ms) <sup>g</sup>	70.9 (26.0)	74.6 (26.7)	80.0 (30.3)	80.2 (32.3)	F=4.8, P=0.003	2>1, 2<3, 4
Block Design <sup>h</sup>	56.3 (8.4)	48.2 (12.4)	37.1 (13.8)	28.1 (11.9)	F=208.3, P<0.001	2<1, 2>3, 4
Digit Symbol <sup>i</sup>	89.7 (11.9)	81.4 (13.0)	72.7 (14.4)	65.5 (13.8)	F=121.7, P<0.001	2<1, 2>3, 4

**Table 4:** Continued.

Profiles	Cognitive trajectory group				Overall trajectory group difference	Pair-wise comparison <sup>**</sup>
	1. High (n=254)	2. Normal (n=413)	3. Mild (n=260)	4. Moderate (n=132)		
Arithmetic <sup>d</sup>	177 (2.6)	149 (3.2)	115 (3.6)	82 (3.1)	F=318.5, P<0.001	2<1; 2>3,4
Information <sup>e</sup>	218 (3.5)	175 (4.0)	141 (3.8)	106 (3.6)	F=299.9, P<0.001	2<1; 2>3,4
Immediate Recall <sup>l</sup>	314 (4.4)	273 (4.9)	250 (5.0)	207 (4.8)	F=159.0, P<0.001	2<1; 2>3,4
Delayed Recall <sup>m</sup>	113 (2.1)	95 (2.4)	85 (2.3)	66 (1.8)	F=113.2, P<0.001	2<1; 2>3,4

<sup>a</sup>Table presents mean (standard deviation) or %; <sup>\*\*</sup>Only significantly different trajectory groups are reported based on Dunnett's adjustment. The reference group is normal trajectory group; <sup>b</sup>Education (Verhage): range 0 (no education), 3-5 (school diploma) to 8 (university degree); <sup>c</sup>IQ: Wechsler Adult Intelligence Scale-III (WAIS-III), short form; <sup>d</sup>PAS: Premorbid Adjustment Scale; <sup>e</sup>SIS-R: Structured Inventory for Schizotypy - Revised (higher scores in PAS and SIS-R reflect poorer outcomes); <sup>f</sup>CAPE: Community Assessment for Psychic Experiences; PE: frequency and distress; Frequency of positive psychotic experiences and amount of distress of these PE; <sup>g</sup>CPT performance: Continuous Performance Test HQ, performance index; <sup>h</sup>CPT variance (ms): CPT-HQ variance in reaction time (ms); <sup>i</sup>Block Design: WAIS-III Block Design; <sup>j</sup>Digit Symbol: WAIS-III Digit Symbol Substitution Test; <sup>k</sup>Arithmetic: WAIS-III Arithmetic; <sup>l</sup>Information: WAIS-III Information; <sup>m</sup>Immediate recall: Word Learning Task (WLT) immediate recall; <sup>n</sup>Delayed recall: WLT Delayed recall; Cognitive trajectory: high performance (1) to moderately altered (4); <sup>p</sup>p < 0.05/20 was considered as statistically significant.

long-term cognitive trajectories. Other studies did not take into account the drop-out modeling of the dependency of cognitive functioning with the function of several time points.[76,84,85]

Some limitations of this study should also be mentioned. First, even though the drop-out modeling analysis showed absence of dependency of cognitive functioning at previous assessments both in patients and unaffected siblings, attrition bias may still be evident. Second, we found that the prediction of the high cognitive performer group of patients on a moderate group of siblings was ambiguous and unstable due to low frequency. Third, strong associations between the patient's cognitive performance and their siblings' cognitive performance may not be claimed based on the sib-pair analysis. Fourth, generating cognition composite score instead of using multivariate cognitive tests may have an impact on finding meaningful trajectory. Fifth, this study used eight neurocognitive tests; therefore, including other tests may lead to different trajectories with different predictions. Finally, the effect of cognitive trajectories on clinical and functional outcomes was not investigated.

In conclusion, our findings confirmed that cognitive functioning in patients with psychotic disorder and their unaffected siblings is heterogeneous. We demonstrated that the cognitive performance of unaffected siblings is situated between that of the patients and the healthy controls. We identified five distinct cognitive trajectories in patients and four trajectories in siblings, which remained stable during six-year follow-up. These trajectories are validated by observing the associations with external factors, such as socio-demographic, (subclinical) psychotic experiences, and premorbid functioning. Moreover, patients cognitive subtypes significantly predicted the sibling subtypes highlighting the heritability of cognition. The study also supports neurocognitive trajectories as a valuable endophenotypic marker. This profiling approach warrants further evaluation in future molecular studies as well as in studies predicting functional and clinical outcomes.

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