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A Network of Psychopathological, Cognitive, and Motor Symptoms in Schizophrenia Spectrum Disorders

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The Genetic Risk and Outcome of Psychosis (GROUP) investigators are listed above the References section.

Schizophrenia spectrum disorders (SSDs) are complex syndromes involving psychopathological, cognitive, and also motor symptoms as core features. A better understanding of how these symptoms mutually impact each other could translate into diagnostic, prognostic, and, eventually, treatment advancements. The present study aimed to: (1) estimate a network model of psychopathological, cognitive, and motor symptoms in SSD; (2) detect communities and explore the connectivity and relative importance of variables within the network; and (3) explore differences in subsample networks according to remission status. A sample of 1007 patients from a multisite cohort study was included in the analysis. We estimated a network of 43 nodes, including all the items from the Positive and Negative Syndrome Scale, a cognitive assessment battery and clinical ratings of extrapyramidal symptoms. Methodologies specific to network analysis were employed to address the study's aims. The estimated network for the total sample was densely interconnected and organized into 7 communities. Nodes related to insight, abstraction capacity, attention, and suspiciousness were the main bridges between network communities. The estimated network for the subgroup of patients in remission showed a sparser density and a different structure compared to the network of nonremitted patients. In conclusion, the present study conveys a detailed characterization of the interrelations between a set of core clinical elements of SSD. These results provide potential novel clues for clinical assessment and intervention.

Key words: network analysis/psychosis/positive symptoms/negative symptoms/cognition/extrapyramidal symptoms

Introduction

Schizophrenia spectrum disorders (SSDs) are complex syndromes with widely different clinical presentations.¹ Psychopathological symptoms are the most visible manifestations and, therefore, used to guide much research. However, 2 other domains—cognitive and motor symptoms—are also impactful. Impairments in social cognition and neurocognition are associated with worse functioning and overall prognosis while also being closely tied to psychopathological symptoms.^{2,3} Motor abnormalities are increasingly regarded as a crucial element of SSD not only as side effects of antipsychotic medication but also as having diagnostic and prognostic value early in the course of illness.^{4,5} Extrapyramidal symptoms (EPS), in particular, are a common motor abnormality that is associated with cognitive deficits already in the early stages of psychotic disorders.⁶ Thus, psychopathological, cognitive, and motor symptoms are interrelated in complex ways in the clinical picture of SSD—possibly reflecting common neurobiological underpinnings⁷—requiring more investigation. A better understanding of how these elements mutually impact each other may

reveal novel insights into the diagnosis, prognosis, and treatment of SSD.

Network models may form a suitable approach to jointly study these areas of clinical manifestations.^{8,9} Network analysis as a tool offers a way to overcome some limitations of other analytical methods that deal with complex interdependencies (eg, need for a priori assumptions in structural equation models or absence of feedback loops in directed acyclical graphs) and generates specific metrics (eg, centrality), which might enrich our knowledge about mental disorders.¹⁰ Moreover, this approach paves the way for a network theory of mental disorders, which asserts that mutually impacting symptoms regulate mental functioning rather than psychological latent constructs. This approach is also aligned with the recent theory that considers health and disease as states of systems that can be studied as networks at multiple levels.¹¹

The most used method for network analysis in psychology research is the Gaussian Graphical Model (GGM), extensively applied to cross-sectional data.¹² In this model, a connection (edge) between 2 variables (nodes) is depicted if they show a pattern of co-occurrence across cases. An edge between 2 nodes can be interpreted as indicative of a potential causal relation.¹³ Through GGM, one can explore a set of clinically relevant variables and gain further insight into, most notably, which elements are crucial in sustaining the network's local and global structure (nodes acting as hubs or bridges between domains are particularly relevant). In less technical terms, this means identifying which symptoms might be responsible for keeping pathological states active.

Several cross-sectional studies used network analysis to investigate clinical variables in psychotic disorders^{14–16}; however, to our knowledge, only 3 included the combination of psychopathological (factor level) and cognitive symptoms.^{17–19} Overall, no previous research looked simultaneously at psychopathology, cognition, and motor symptoms.

We aimed: (1) to estimate a network consisting of psychopathological, cognitive, and motor symptoms in patients with SSD; (2) to describe the network structure by identifying the most important nodes in the overall network and in connecting different variable domains; and (3) to explore network differences between patients in remission vs patients not in remission, given that SSDs frequently are persistent disorders that greatly vary in their clinical presentation according to remission status (in psychotic episodes or between episodes).

Methods

Subjects

In the current study, data collected in the Genetic Risk and Outcome of Psychosis (GROUP) study were used. The GROUP study is a multisite longitudinal

cohort investigation run across 4 academic centers in the Netherlands (Amsterdam, Groningen, Maastricht, and Utrecht) and including data from patients, siblings, parents, and controls. It is described extensively elsewhere.²⁰ For the current study, only baseline patient data were considered (collected between 2004 and 2007). All the participants had a diagnosis of a nonaffective psychotic disorder according to the *Diagnostic and Statistical Manual of Mental Disorders*, Fourth Edition, criteria²¹ and were aged between 16 and 50 years.

Assessments

Over 30 interviewers per recruitment site were trained for administering the assessments. The Positive and Negative Syndrome Scale (PANSS) was used to assess psychopathological symptoms.²² This instrument is composed of 30 items distributed over 3 subscales: positive, negative, and general psychopathology. The PANSS total score had an intraclass correlation coefficient of 0.946 (95% CI: 0.758–0.996), thus the interrater reliability was considered good.

A battery of neuropsychological tests was used. It covered the fundamental domains affected in schizophrenia as considered in the MATRICS consensus,²³ with the exception of visual learning and memory (not included in the GROUP study): *processing speed*—Digit Symbol Substitution Test from the Wechsler Adult Intelligence Scale-III (WAIS-III)²⁴; *working memory*—arithmetic subtest from the WAIS-III; *reasoning and problem solving*—Response Set-shifting Task,²⁵ accuracy cost score (RST), and the block design subtest from the WAIS-III; *verbal learning and memory*—Word Learning Task,²⁶ correct delayed recall items; *attention/vigilance*—Continuous Performance Test-HQ²⁷, number of correct positives; *social cognition*—Benton Facial Recognition Test,²⁸ total correct items, Degraded Facial Affect Recognition Test,²⁹ percentage of correct items (DFAR), and the Hinting task,³⁰ total score. The WAIS-III information subtest was also included as a general knowledge assessment.

Motor symptoms were assessed with the following scales for different EPS domains: Unified Parkinson's Disease Rating Scale,³¹ motor examination section score (UPDRS); Barnes Akathisia Rating Scale,³² global clinical assessment of akathisia; a rating of dystonia; Abnormal Involuntary Movement Scale,³³ overall score (AIMS). Akathisia and dystonia were graded on an ordinal scale, respectively, from 0 (absent) to 5 (severe) and from 0 (absent) to 4 (severe). All other measures in the analysis were considered continuous.

Subjects with missing data represented 30.1% of the total sample (315/1047). We opted to use a conservative approach and excluded all cases with more than 15% missing values ($N = 40$). Multiple imputation was used for all other cases (*mice* package for R³⁴; [supplementary figure S1](#)).

Network Analysis

Network Estimation A nonparanormal transformation using the *huge* package was applied to all the included variables, except the ordinal variables akathisia and dystonia, in order to relax the normality assumption.³⁵ This step also transformed values through standardization.

A network with 43 nodes was constructed using a GGM and implemented through the R statistical software version 3.6.2,³⁶ with the packages *bootnet* and *qgraph*.^{37,38} Edges between nodes were undirected (no arrowhead) and established while controlling for the influence of every other node in the network. Furthermore, a L1-regularization procedure was used—Graphical Least Absolute Shrinkage and Selection Operator (GLASSO)—in combination with Extended Bayesian Information Criterion (EBIC). GLASSO limits spurious connections between nodes (false-positives) by shrinking small correlations to 0. The EBIC is a goodness-of-fit measure for model selection that is tuned through a hyperparameter γ . Following recommendations in the field, this hyperparameter was tuned to 0.5, ensuring a good balance between sensitivity and specificity.³⁹ Polyserial partial correlations were calculated between the ordinal and the continuous variables; Pearson partial correlations were used in all other cases. The network layout followed the Fruchterman–Reingold algorithm, a method that places strongly related nodes close to each other and repels less related nodes.⁴⁰ In this layout, the edges display their strength (edge-weight) through line thickness and direction through color (blue for positive partial correlations and dashed red for negative ones).

Stability and Accuracy Checks Following recommendations in this field, bootstrapping routines were used to assess the estimated model's stability and accuracy.¹³ Further details about these methods can be found as [supplementary material](#).

Community Detection An Exploratory Graph Analysis (EGA) implemented through the package *EGAnet* was used to investigate how the network was organized in terms of dimensions.⁴¹ This method is based on the GLASSO and uses a random walk algorithm to identify communities in the network. The stability of the EGA's estimation was also assessed through bootstrapping the empirical correlation matrix with 1000 iterations.⁴² This approach allows to quantify the stability of each node regarding their community placement.

Network Metrics—Small-Worldness Index and Centrality Analysis The small-worldness index quantifies how much a network displays small-world properties (*qgraph* package).⁴³ In a small-world network, there is a balance between local robustness of connections (high clustering coefficient) and a short distance between any pair of nodes (short path length). In the context of mental disorders, this means that certain symptoms will be more closely tied and reinforcing each other. They will thus

segregate into communities representing syndromes (eg, negative symptoms) bound to each other by “bridge” symptoms.

Two different types of centrality measures were calculated for each node. *Strength centrality* measures the number of connections of a node considering their weight; it represents the influence of a node considering the local structure of the network (ie, its direct connections with other symptoms). This measure seems to be robust, predictive of subsequent diagnosis in longitudinal studies, and indicative of potential to change the remainder network as reviewed by Robinaugh et al.¹² *Bridge strength centrality* (*networktools* package⁴⁴) measures the weighted connectivity of a node in a given community to other communities in a network.⁴⁵ Communities are sets of highly interconnected nodes in a network. In the medical setting, they might represent clinical syndromes or variable domains. For this study, communities were established through a data-driven approach described in 2.3.3. **Network Comparison Test: Remitted vs. Nonremitted Patients** A binary division of the sample was carried out according to remission status (see [supplementary table S3](#)).⁴⁶ A network was estimated for each subsample (see 2.3.1), and a Network Comparison Test (NCT) between the networks was performed.⁴⁷ The NCT (*NetworkComparisonTest* package) uses a permutation approach to test for differences between the networks' invariant structure and invariant global strength. Moreover, we calculated strength centrality for each node of each subsample's network. We also carried out a stability analysis for each of the subsample's networks and tested for differences between clinical and sociodemographic variables.

Sensitivity Analysis An NCT was performed between the complete sample network and a network estimated without the patients taking typical antipsychotics. This was done to probe the influence on the network's connectivity of patients taking antipsychotics that are more EPS inducing.

Results

Sample Characteristics

Table 1 displays the main demographic and clinical characteristics of the included patients ($N = 1007$; see [supplementary table S1](#) for a detailed description by the diagnostic group). **Table 2** shows the mean score and SD for each variable included as a node in the estimated network. The dystonia score distribution was extremely skewed (96% of subjects rated with absent); therefore, this variable was discarded. The akathisia score was compressed to only vary between 0 and 2 due to few cases with higher scores (scores 3 and 4 converted to 2; no cases scored 5). The 2 subsamples defined according to remission status were composed of 451 patients in remission

Table 1. Demographic and clinical characteristics of the included sample

Variable	Participants (<i>N</i> = 1007)
Age, mean (SD), years	27.3 (7.4)
Females, no. (%)	231 (22.9)
Education, highest achieved	
No education, no. (%)	6 (<1)
Elementary, no. (%)	130 (13)
Secondary, no. (%)	563 (56.4)
University or vocational, no. (%)	299 (30.0)
Married, no. (%)	88/993 (8.9)
Working (full time or part time), no. (%)	485/866 (56.0)
Illness duration, mean (SD), years	5.0 (4.4)
Global assessment of functioning, mean (SD)	51.6 (15.9)
Use of antipsychotic medication, no. (%)	872/876 (99.5)
Use of typical antipsychotics, no. (%)	104/872 (11.9)
Diagnosis	
Schizophrenia, no. (%)	668 (66.5)
Schizoaffective disorder, no. (%)	118 (11.7)
Psychosis NOS, no. (%)	102 (10.1)
Schizophreniform disorder, no. (%)	56 (5.6)
Brief psychotic disorder, no. (%)	20 (2.0)
Other diagnosis, no. (%)	41 (4.1)
In clinical remission, no (%)	451/997 (45.2)

Note: NOS, not otherwise specified.

and 546 not in remission (10 subjects lacked information on remission status).

Network Analysis

Figure 1 displays the estimated network. For the partial correlation matrix used as input, see [supplementary table S2](#). The estimated network was highly interconnected and every node displayed edges to other nodes.

Stability and Accuracy Checks The bootstrapping routines employed to assess the stability and accuracy of strength centrality and edge weights returned satisfactory results (complete analysis available as [supplementary figures S2–S5](#)).

Community Detection Seven communities were found with the EGA and confirmed through parametric bootstrapping ([figure 1](#)). All the cognitive nodes belonged to one single community, which also encompassed “difficulty in abstract thinking” (N5). The remaining items from the PANSS formed 5 communities. All 3 motor nodes composed a separate single community. Node stability regarding community placement can be retrieved in [supplementary figure S6](#). Notable cases of unstable placement were the nodes P4, P7, G8, G14 (together composing one community), and N5.

Small-Worldness Index and Centrality Analysis The small-worldness index was 1.46, above the threshold value of 1 set by Humphries and Gurney⁴³; hence, the network can be considered to have small-world properties.

Figure 2 displays the different centrality measures per node (*z*-transformed). The node with the highest strength centrality score was distinctively “delusions” (P1). Cognitive nodes belonging to the WAIS were in the upper half of strength centrality scores and the other cognitive nodes were in the bottom half. Parkinsonism node (UPDRS) presented the highest strength centrality for a motor node. Regarding bridge strength centrality, the strongest bridges for each community were “lack of judgment and insight” (G12), “suspiciousness” (P6), “difficulty in abstract thinking” (N5), “excitement” (P4), “motor retardation” (G7), “tension” (G4), and parkinsonism (UPDRS).

NCT: Remitted vs Nonremitted Patients The remitted patients’ network was sparser, even presenting 3 disconnected nodes (DFAR, RST, and AIMS; [figure 3A](#)). The comparison test between the networks of patients in remission vs nonremitters yielded significant results for differences in global strength ($P < .01$) and network structure ($P = .012$). [Figure 3B](#) shows strength centrality for nodes of both subsample networks. Certain nodes changed widely their strength centrality according to remission status, especially “Excitement” (P4), “Emotional withdrawal” (N2), and theory of mind (Hinting task).

Stability analysis of the subsample networks rendered less stable results when compared to the full network, in particular for the network of patients in remission ([supplementary figures S9–S12](#)). Clinical and demographic features varied significantly between the 2 subsamples ([supplementary table S4](#)).

Sensitivity Analysis No significant differences were found between the network of patients not taking typical antipsychotics and the full-sample network (see [supplementary figure S11](#)).

Discussion

To our knowledge, this is the first network analysis study examining associations between psychopathology, cognition, and motor variables in patients with SSD. Overall, our results confirmed that these 3 clinical domains are highly interrelated but, at the same time, organized into separate communities. These communities followed, in general, the expected factor organization of the variables. Nodes related to insight, abstraction capacity, attention, and suspiciousness were the main bridges between network communities. Lastly, different clinical states were represented by different network architectures.

Community Detection

Within the PANSS items, our data-driven network approach replicated the most consensual 5-factor solution—positive, negative, disorganized, excited, and depressed/emotional distress factors.⁴⁸ The main difference found was that “difficulty in abstract thinking” (N5)

Table 2. Descriptive statistics of the variables submitted to network analysis

Domains and variables	Mean (SD)
Positive symptoms	
P1—Delusions	2.41 (1.52)
P2—Conceptual disorganization	1.69 (1.05)
P3—Hallucinations	2.11 (1.53)
P4—Excitement	1.33 (0.76)
P5—Grandiosity	1.50 (1.00)
P6—Suspiciousness/persecution	2.27 (1.41)
P7—Hostility	1.32 (0.77)
Negative symptoms	
N1—Blunted affect	2.35 (1.33)
N2—Emotional withdrawal	2.13 (1.20)
N3—Poor rapport	1.72 (1.02)
N4—Passive/apathetic social withdrawal	2.27 (1.31)
N5—Difficulty in abstract thinking	2.02 (1.28)
N6—Lack of spontaneity and flow of conversation	1.90 (1.20)
N7—Stereotyped thinking	1.59 (0.96)
General psychopathology	
G1—Somatic concern	1.68 (1.81)
G2—Anxiety	2.34 (1.27)
G3—Guilt feelings	1.97 (1.25)
G4—Tension	1.84 (1.03)
G5—Mannerisms and posturing	1.30 (0.72)
G6—Depression	2.47 (1.41)
G7—Motor retardation	1.86 (1.11)
G8—Uncooperativeness	1.23 (0.63)
G9—Unusual thought content	2.04 (1.31)
G10—Disorientation	1.19 (0.56)
G11—Poor attention	1.73 (1.02)
G12—Lack of judgment and insight	2.10 (1.39)
G13—Disturbance of volition	1.59 (0.93)
G14—Poor impulse control	1.31 (0.74)
G15—Preoccupation	1.47 (0.89)
G16—Active social avoidance	1.76 (1.08)
Motor symptoms	
Parkinsonism (Unified Parkinson's Disease Rating Scale)	0.21 (0.26)
Dyskinesia (Abnormal Involuntary Movement Scale)	0.07 (0.18)
Akathisia (Barnes Akathisia Rating Scale)	
0 (absent)	796 (79%)
1 (questionable)	109 (11%)
2 (mild or more severe)	102 (10%)
Cognition	
Processing speed: WAIS-III Digit Symbol Substitution Test	7.41 (3.01)
World knowledge: WAIS-III Information	10.65 (3.09)
Working memory: WAIS-III Arithmetic	9.50 (3.26)
Reasoning and problem solving: WAIS-III Block design	9.60 (3.36)
Verbal learning and memory: Word Learning Task, delayed recall	7.54 (2.88)
Attention/vigilance: Continuous Performance Task	26.43/28 (2.77)
HQ	
Reasoning and problem solving: Response Set-shifting Task ^a	0.20 (0.27)
Social cognition (emotional face recognition): Degraded Facial Affect Recognition Test	68.76 (10.53)
Social cognition (facial recognition): Benton Facial Recognition Test	22.77 (2.29)
Social cognition (theory of mind): hinting task	17.52 (2.79)

^aScores were transformed to the symmetrical distribution so that they could have the same meaning as the other cognitive variables (ie, higher score means better performance).

was part of the cognitive community in the current study, whereas it is commonly placed with disorganized symptoms. This was probably due to the simultaneous analysis

of cognitive measures in the present study and not strictly the PANSS as in the factor analytic studies.

As expected, cognitive and motor variables formed 2 separate communities. The relatively small number of nodes in each of these domains might explain why no further communities were detected. Some nodes, notably N5 and the excitement community, had unstable community placements, probably reflecting multidimensionality (supplementary figure S6).

Small-Worldness Index and Centrality Analysis

The estimated network showed high interconnectivity, also extending the finding of small-world properties of psychopathological networks to a broader set of symptoms.⁴⁹ This means that interferences at any node/edge within the network will likely affect many other nodes/edges directly or indirectly within a few steps. At the same time, some nodes might play a more important role in network connectivity, acting as bridges between communities.

Within each of the 5 communities composed of PANSS items, the node with highest strength centrality corresponded to the item that commonly has the highest factor loading—P1 for positive symptoms, N6 for negative symptoms, P2 for disorganization, P4 for excitement, and G2 for emotional distress.⁵⁰ This is in line with recent insights in the field that state that strength centrality is redundant with factor loading.⁵¹ Results from previous network analysis studies using the PANSS were largely replicated, except for N6 (alogia factor), which had been reported as a less central node.^{52,53} A possible explanation for this difference concerns the high amount of connections N6 establishes outside of the negative symptom domain, making it more central in the present study. The same pattern occurred for cognition, where working memory—known to have a great overlap with general cognition⁵⁴—was highlighted. Parkinsonism was the most central motor node possibly due to its higher construct complexity, capturing multiple subcomponents. Previous studies have shown that parkinsonism is a particularly important symptom in SSD for its high prevalence, spontaneous presence in drug-naïve patients, and association with cognition.⁵⁵

Regarding bridge strength centrality, 2 main sets of bridge nodes should be addressed. A first set was composed of metacognitive abilities—such as abstraction capacity (N5) and insight (G12)—the capacity to maintain attention (G11) and movement-related nodes (G7 and UPDRS). A careful look into the edges established by these bridge nodes also reveals proximity to the theory of mind (ToM) node and negative symptom nodes associated with the expressive deficit factor (N1 and N6). Taken together, these results highlight previous findings from different lines of research, for instance: the previously described association between

Fig. 1. Estimated network ($N = 1007$). Nodes were grouped according to the community detection results using Exploratory Graph Analysis.

ToM, disorganization symptoms, and metaphor comprehension in SSD^{56,57}; the relation between gestural perception and gestural performance, mediated by frontal lobe dysfunction,⁵⁸ given that motor symptoms are associated with impaired nonverbal communication in psychosis⁵⁹; the association between parkinsonism and disorganization symptoms⁶⁰; and the evidence that, in Parkinson's disease, conceptualization skills are particularly affected within frontal lobe functions.⁶¹ The fact that 2 of the 3 strongest bridge strength centrality nodes belonged to the disorganization community (G11

and G12) reiterates the pivoting role this community plays between psychopathology and cognition in line with findings from previous network analysis studies, including on high-risk subjects.^{17,19,52}

A second set of bridging nodes was related to the “affective” components of the network and had “suspiciousness” as the hub, mostly binding positive symptoms, excitement, and emotional distress communities. This might indicate that suspiciousness is a bridge for the emergence of mood episodes within SSD. In a previous network analysis study, van Rooijen et al¹⁴ also found

Fig. 2. Centrality indices plot (z-scores). Nodes in bridge strength centrality are ordered according to community placement (C1–7).

“suspiciousness” as relevant regarding co-occurrence with depressive symptoms, namely suicidality, in SSD.

Network Comparisons of Subgroups

The remission subsample’s network showed a significantly lower density of connections. This might be caused by the reduction of self-reinforcing feedback loops that contribute to the maintenance of syndromes in line with the network theory of mental disorders.⁸ It could, however, be in part attributed to a floor-effect artifact.⁶²

Network sparsity in remission was conspicuous in the subset of higher-order cognitive functions. This pattern mirrors previously described differences between patients and healthy subjects in terms of cognition factor structure—in patients, cognitive skills load more in one single “g” factor, implying that cognition is less domain specific than in controls.⁶³

Moreover, network structure also differed with respect to remission status and, accordingly, node centrality hierarchy changed too. Interestingly, the highest disparities in centrality were not strictly found in nodes integrated

These results comply with another network analysis study comprising cognitive nodes, which also found

On a cautionary note, 2 aspects limit the interpretation of these findings. First, due to smaller sample size, the estimated networks for each group are less stable, hence possibly failing more to correctly estimate the true edges. Second, these findings might partly be explained by sociodemographical and clinical differences between the 2 subsamples.

Strengths and Limitations

Some of this study's strengths flow from the innovative network approach to psychiatric disorders and include: the joint analysis of many variables, hence dissecting the correlational architecture of the disorder at a system level, and the use of metrics that are specific to this approach and that allow for exploration of results in an objective fashion. In addition, this study benefitted from a large sample size; a thorough stability analysis that corroborated the reliability of results; and the inclusion of a broad set of patients—diagnoses along the nonaffective psychosis spectrum (not limited to schizophrenia) and both males and females.

A first limitation of this study is its group-level approach; thus, extrapolations to the individual level should be performed carefully. Studies using longitudinal data could tackle this limitation in the future. In addition, the cross-sectional nature of the data precludes causal interpretation of associations. Second, important elements might be lacking from the cognitive and motor domains (eg, visuo-spatial memory or neurological soft signs) and, concerning the latter, the inclusion of more objective measures (eg, actigraphy) could have reduced bias caused by poorly defined variables (eg, “G7—motor retardation”).⁶⁹ Third, a node representing antipsychotic medication was not included due to a lack of data on the prescribed dosages. However, the available information on the type of medication used and a further sensitivity analysis showed that only a minority of patients was taking typical antipsychotics and that this variable did not seem to affect the network's connectivity.

Conclusions

In this study, we showed that cognitive and motor symptoms also played an important role in a network of core clinical variables of SSD. Moreover, we found that a group of metarepresentational abilities (abstraction, insight, and theory of mind), expressive deficit negative symptoms, and motor disturbances composed an important aggregating set of symptoms. We believe that these findings have implications (1) for phenomenology and cognitive science in that they reiterate the fundamental role of motor behavior and specific higher-order skills, respectively; (2) for neuroscientific research, encouraging research on brain systems that integrate cognitive, motor, and affective functions⁷⁰; and, eventually, (3) for treatment approaches, emboldening metacognitive and embodied therapies as they target what seem to be the most unifying elements of the estimated network. Lastly, our findings endorse the integration of remission status into future network analysis studies of mental disorders with oscillatory dynamics (eg, SSD and mood disorders).

Supplementary Material

Supplementary material is available at *Schizophrenia Bulletin*.

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