# Psychopathological Syndromes Across Affective and Psychotic Disorders Correlate With Gray Matter Volumes

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Introduction: More than a century of research on the neurobiological underpinnings of major psychiatric disorders (major depressive disorder [MDD], bipolar disorder [BD], schizophrenia [SZ], and schizoaffective disorder [SZA]) has been unable to identify diagnostic markers. An alternative approach is to study dimensional psychopathological syndromes that cut across categorical diagnoses. The aim of the current study was to identify gray matter volume (GMV) correlates of transdiagnostic symptom dimensions. Methods: We tested the association of 5 psychopathological factors with GMV using multiple regression models in a sample of N = 1069 patients meeting *Diagnostic and* Statistical Manual of Mental Disorders, Fourth Edition (DSM-IV) criteria for MDD (n = 818), BD (n = 132), and SZ/SZA (n = 119). T1-weighted brain images were acquired with 3-Tesla magnetic resonance imaging and preprocessed with CAT12. Interactions analyses (diagnosis × psychopathological factor) were performed to test whether local GMV associations were driven by DSM-IV diagnosis. We further tested syndrome specific regions of interest (ROIs). Results: Whole brain analysis showed a significant negative association of the positive formal thought disorder factor with GMV in the right middle frontal gyrus, the paranoid-hallucinatory syndrome in the right fusiform, and the left middle frontal gyri. ROI analyses further showed additional negative associations, including the negative syndrome with bilateral frontal opercula, positive formal thought disorder with the left amygdala-hippocampus complex, and the paranoid-hallucinatory syndrome with the left angular gyrus. None of the GMV associations interacted with DSM-IV diagnosis. Conclusions: We found associations between psychopathological syndromes and regional GMV independent of diagnosis. Our findings open a new avenue for neurobiological research across disorders,

using syndrome-based approaches rather than categorical diagnoses.

*Key words:* voxel-based morphometry/transdiagnostic/dimensional/major psychiatric disorders

#### Introduction

There is strong evidence that psychotic (schizophrenia [SZ] and schizoaffective disorder [SZA]—both henceforth referred as schizophrenia spectrum disorder [SSD]) and affective disorders (major depressive disorder [MDD] and bipolar disorder [BD]) (henceforth, together referred to as major psychiatric disorders) are overlapping regarding symptoms, course, and outcome.<sup>1,2</sup> Neurobiological research showed that these major psychiatric disorders share familial and molecular genetic risk,<sup>3</sup> environmental risks,<sup>4</sup> structural brain changes,<sup>5-8</sup> and other neurobiological markers.<sup>9</sup> On a phenomenological level, the major psychiatric disorders share many syndromes such as depression, mania, and psychosis.<sup>10-12</sup>

Since research using diagnostic categories might overlook psychopathological mechanisms across disorders, transdiagnostic, dimensional approaches 13–15 can serve as an important addition to traditional approaches comparing diagnostic groups. In studies on a shared psychopathological factor structure across SSD, MDD, and BD, 3–5 factors have been delineated. 16–19 Most commonly paranoid-hallucinatory, depressive, negative, disorganized, and manic dimensions can be identified as separate dimensions. 16,18,20 Confirmatory analyses of 6 competing factor models revealed that symptom dimensions are better represented by factor models including 4 or 5 factors rather than by models with fewer factors. 17

In our own previous study, we cross-validated a 5-factor model comprising depression, negative syndrome, positive formal thought disorder (pFTD), paranoid-hallucinatory syndrome, and increased appetite across the major psychiatric disorders.<sup>2</sup>

For a long time,<sup>21</sup> it has been hypothesized that particular syndromes might share a common brain structural network alteration, independent of the diagnostic category. 1,5-7,22 But up to now, most studies investigating gray matter volume (GMV) alterations across disorders focused on categorical approaches.<sup>6,7</sup> Using a multimodal machine learning approach aiming to classify recent onset psychosis and depression revealed no points of rarity on a brain structural level indicating comparable GMV across psychosis patients with comorbid depressive symptoms and patients with recent onset depression.8 This result is further supported by a study<sup>23</sup> showing that specifically in patients with younger age disorder onset, neuroanatomical disease signatures fail to separate affective and psychotic disorders. Based on these findings, we aim to shed light on brain structural correlates of psychopathological factor dimensions across disorders. Below, we summarize results of previous voxel-based morphometry (VBM) studies on structural correlates of psychopathological factors. We focus on the neural substrates of the 5 psychopathological factors derived from our previous study.2

The paranoid-hallucinatory syndrome has mostly been investigated in studies including SZ patients. There is meta-analytical evidence that auditory verbal hallucinations are negatively correlated with GMV in the left insula and right superior temporal gyrus (STG).<sup>24</sup> Further core regions<sup>25</sup> negatively associated with the paranoid-hallucinatory syndrome are the thalamus,<sup>26–28</sup> the left planum temporale,<sup>29</sup> left anterior cingulate, and the bilateral insulae.<sup>30,31</sup>

pFTD have been frequently associated with neuroanatomical alterations in the left STG, frontal opercula, and left middle temporal gyrus, (ie, Wernicke and Broca area).<sup>32,33</sup> Dimensional analyses of pFTD in SZ patients showed negative associations in the bilateral inferior frontal gyri, the orbito-frontal cortex (OFC), the middle, medial, and superior frontal gyri, the left amygdalahippocampus complex, the precuneus, and the insula.<sup>34,35</sup>

Meta-analyses of negative symptoms in SZ patients reported GMV reductions in the OFC, the anterior cingulate cortex, fusiform gyrus, thalamus, caudate, and amygdala associated with the severity of negative symptoms. However, results are heterogeneous. Some studies reported no association between GMV and negative symptoms on a whole brain level. 38-40

Testing dimensional depressive symptomatology and GMV in MDD, there were correlations in the right OFC, the left hippocampal gyrus, and the right dorsolateral prefrontal cortex.<sup>41</sup> Investigation of subclinical depressive symptoms in healthy populations revealed inconsistent

associations in the anterior cingulate, OCF, and thalamus.<sup>42-44</sup> Nevertheless, several studies found no associations between GMV and psychopathological measures in MDD patients.<sup>45-47</sup>

The psychopathological factor "increased appetite" from our phenomenological study has not been reported in previous factor analytical approaches.<sup>2</sup> This dimension has only been reported within atypical MDD patients.<sup>48,49</sup> One study investigating subtypes of MDD patients indicated that a severely increased appetite MDD subtype showed lower surface area in the anterior insula when compared to a healthy control group.<sup>50</sup> Neurobiological research investigating obesity in otherwise healthy controls indicated the amygdala, hippocampus, insula, OFC, and the striatum to be central regions of appetite behavior.<sup>51,52</sup> A study investigating MDD patients revealed lower temporo-frontal cortical thickness to be associated with obesity (body mass index > 30).<sup>53</sup>

In summary, most transdiagnostic GMV studies are limited by (1) results from studies comparing diagnostic categories (as opposed to dimensional investigations), or (2) dimensional psychopathological investigations restricted to 2 diagnostic groups, and/or by (3) investigating one psychopathological dimension not covering the whole psychopathological spectrum. To overcome these limitations, our aim was to investigate associations of psychopathological dimensions with GMV on a dimensional, transdiagnostic, data-driven level across MDD, BD, and SSD. Based on previous findings, 5,6,8,23 we hypothesize that associations between psychopathological syndromes and local GMV that have previously been detected within one diagnosis, cut across MDD, BD, and SSD, and do not interact with diagnostic categories.

#### **Materials and Methods**

**Participants** 

We included structural magnetic resonance imaging (MRI) data of acute and remitted patients (aged 18–65) meeting Diagnostic and Statistical Manual of Mental Disorders, Fourth Edition (DSM-IV) criteria (SCID-I) for MDD (296.2X, 296.3X: n = 818, F = 530/M = 288), BD (292.5X, 292.6X, 292.7X, 296.0X, 296.4X, 296.5X, 296.6X, 296.7X, or 296.8X: n = 132, F = 71/M = 61), SZ (295.X: n = 74, F = 32/M = 42), and SZA (295.7: n = 45, F = 24/M = 21). This MRI sample is a subgroup of our previous study on psychopathological factors,<sup>2</sup> from which, after quality checks of the brain scans n = 113 patients had to be excluded, leaving N = 1069 participants for the current analyses (see table 1). Patients were part of the DFG FOR2107 consortium<sup>54</sup> and were interviewed and scanned at the Departments of Psychiatry and Psychotherapy Marburg or Münster Universities, Germany. In- and outpatients were recruited from these Universities and local psychiatric hospitals (Vitos Marburg, Gießen, Herborn,

**Table 1.** Sample Characteristics

	Major Depressive Disorder ( $n = 818$ )	Bipolar Disorder $(n = 132)$	Schizophrenia Spectrum Disorder $(n = 119; SZ n = 74, SZA n = 45)$	$P(F/\chi^2)$
Age	36.95 (13.19)	41.03 (11.94)	38.13 (11.81)	.003ª (5.82)
Sex	M = 288, $F = 530$	M = 61, F = 71	M = 63, F = 56	< .001 (17.53)
Years of education	13.19 (2.74)	14.02 (2.8)	12.61 (2.68)	< .001 <sup>b</sup> (8.07)
Age of onset	26.12 (12.62)	24.26 (11.29)	22.46 (9.4)	$.005^{\circ}(5.39)$
Life time cumulative duration of hospitalizations (months)	11.68 (17.84)	33.23 (33.59)	38.46 (38.91)	< .001 <sup>d</sup> (96.71)
Duration of current episode	22.84 (46.46)	12.92 (35.61)	30.2 (56.93)	.093 (2.39)
(months)				
Verbal IQ	112.67 (13.78)	114.98 (15.62)	111.82 (14.79)	.161 (1.83)
Psychopathological factors				
Depression (F1)	0.69 (1.02)	-0.33(0.95)	-0.29(0.88)	< .001e (13.83)
Negative syndrome (F2)	-0.06(0.47)	-0.14(0.36)	0.33 (0.74)	$< .001^{\rm f}(35.42)$
Positive formal thought disorder (F3)	-0.04 (0.1)	0.04 (0.21)	0.19 (0.37)	< .001g (107.14)
Paranoid-hallucinatory syndrome (F4)	-0.07 (0.13)	-0.03 (0.13)	0.47 (0.71)	< .001 <sup>h</sup> (219.5)
Increased appetite (F5)	-0.01(0.53)	0.09 (0.71)	-0.04 (0.51)	.12 (2.13)

*Note*: SZ, schizophrenia; SZA, schizoaffective disorder. Values indicate means and SD (in brackets). Post hoc differences between groups. <sup>a</sup>Bipolar disorder (BD) > major depressive disorder (MDD).

and Haina, LWL Münster, Germany) and via posting in local newspapers and flyers. Exclusion criteria were any history of neurological (head trauma or unconsciousness) and medical condition, substance dependence, current use of benzodiazepines, and IQ ≤80. The assessment of psychopathological symptoms and MRI data acquisition was performed within the same week. n = 341 patients (31.9%) did not receive any psychotropic medication, 53.5% received antidepressants, 12.1% mood stabilizers, and 29.6% antipsychotics at time of data collection. Based on DSM-IV criteria, n = 12 MDD (296.24, 296.34) patients and n = 6 BD (295.04, 296.44, 296.54) patients presented with psychotic features. A small number of participants were diagnosed with a past alcohol (n = 52) or substance (n = 26) abuse. Patients gave written informed consent to study protocols approved by the local Ethics Committees according to the Declaration of Helsinki.

#### MRI Data Acquisition and Preprocessing

MRI data acquisition was done according to an extensive quality assurance protocol.<sup>55</sup> In Münster, a 3T MRI scanner (Prisma, Siemens, Germany) and a 20-channel head matrix Rx-coil were used. MRI data in Marburg were obtained using a 3T MRI scanner (Tim Trio, Siemens, Germany) and a 12-channel head matrix Rx-coil. At

both sites, we used a fast gradient echo MP-RAGE sequence with a slice thickness of 1.0 mm consisting of 176 in Marburg and 192 in Münster sagittal orientated slices and a field of view of 256 mm. Parameters were differing across sites: Marburg: time of repetition [TR] = 1.9 seconds, echo time [TE] = 2.26 milliseconds, inversion time [TI] = 900 milliseconds, flip angle = 9°; Münster: TR = 2.13 seconds, TE = 2.28 milliseconds, TI = 900 milliseconds, flip angle = 8°. Before preprocessing, all scans were visually inspected regarding artifacts and anatomical abnormalities by a senior clinician (U.D.). Structural MRI data were preprocessed<sup>56</sup> using default parameters as implemented in the CAT12-Toolbox (Computation Anatomy Toolbox for SPM, build 1184, Structural Brain Mapping group, Jena University Hospital, Germany; http://dbm.neuro.uni-jena.de/cat/) building on SPM12 (Statistical Parametric Mapping, Institute of Neurology, London, UK) providing bias-corrected, tissue classified, and normalized data ratings. During preprocessing, images were segmented<sup>57</sup> into gray matter, white matter, and cerebrospinal fluid. Images were spatially registered, segmented, and normalized<sup>58</sup> using a DARTEL algorithm. All scans underwent the automated quality assurance, using the CAT12 "check data quality using covariance" procedure. After preprocessing and the described quality assurance protocols, we excluded n = 113 patients, due to major artifacts or abnormalities not accomplishing

<sup>&</sup>lt;sup>b</sup>BD > MDD, schizophrenia spectrum disorder (SSD).

 $<sup>^{</sup>c}MDD > SSD.$ 

 $<sup>^{</sup>d}SSD > MDD$ , BD > MDD.

cMDD > SSD, BD.

fSSD > MDD, BD.

gSSD > BD, MDD; BD > MDD.

 $<sup>^{</sup>h}SSD > BD, MDD.$ 

the CAT12 quality criteria, leaving N = 1069 for the current study. MRI data sets were spatially smoothed with a Gaussian kernel of 8 mm full width at half maximum.

## Statistical Analyses

Multidimensional Factors. Using a cross-validation approach within 2 samples, we had performed an exploratory and confirmatory psychopathological factor analysis in our previous study.2The scale for the assessment of negative symptoms,59 scale for the assessment of positive symptoms,60 Young mania rating scale,61 Hamilton anxiety rating scale,62 and the Hamilton depression scale63 with a total of 104 symptoms were used to identify multidimensional, psychopathological factors across diagnosis. Psychopathological data were obtained during a clinical interview and were rated immediately afterwards by clinically trained psychologists (for detailed information see, Stein et al.<sup>2</sup>). Interrater reliability achieved excellent values of >.86 in all scales. Summarizing the procedures, we divided the total sample and conducted a varimax-rotated principal axis factor analysis in the first sample. To validate the explorative factor solution, we performed a confirmatory factor analysis in the second sample using Mplus<sup>64</sup> (MLR model estimation) showing a good fit:  $\chi^2 = 1287.842$ , df = 571, P < .0001, comparative fit index = 0.932, root mean square error of approximation = 0.036. The following factors were found in our previous study: depression, negative syndrome, pFTD, paranoid-hallucinatory syndrome, and increased appetite. Latent, standardized factor scores for each patient of the current study were used from this previous study,<sup>2</sup> to test whether the previously established dimensional factors were associated with GMV.

Voxel-Based Morphometry Analyses: Whole Brain Level. We used smoothed GMVs and standardized latent factor scores for each patient to perform separate linear regression models for each transdiagnostic factor (not for each diagnostic group). Analyses were carried out using SPM12 (https://www.fil.ion.ucl.ac.uk/spm/). To avoid potential confounders, we applied several covariates of no interest: age, sex, site and total intracranial volume, and the change of one gradient coil.54,55 As previously reported in our factor analytic study,<sup>2</sup> increased appetite was not correlated to antipsychotics, only to intake of serotoninnorepinephrine reuptake inhibitors and noradrenergic and specific serotonergic antidepressants with negligible effects. Therefore, we used 3 dummy-coded covariates accounting for the intake of at least one antidepressant, mood stabilizer, and antipsychotic. As recommended for VBM analyses, absolute threshold masking with a threshold value of 0.1 was used. Cluster labeling was applied using the dartel space Neuromorphometrics atlas (http://www. neuromorphometrics.com/).

For each psychopathological factor, associations with GMV (whole brain) at peak-level threshold P < .05,

family-wise error (FWE) corrected, and cluster extend threshold k=10 were investigated. To investigate if our transdiagnostic brain correlates were driven by DSM-IV diagnostic categories, we performed ANCOVA interaction analyses (diagnostic category  $\times$  factor) in SPM on whole brain level.

Due to the unbalanced distribution of DSM-IV categories, in further confirmatory analyses, we used whole brain clusters of the total sample and tested them as regions of interest (ROIs) in an equally distributed sample matched for age and sex (n=357) (matching was performed using the "MatchIt" package<sup>65</sup> in R<sup>66</sup>). We also performed ANCOVA interaction analyses in the matched sample. For the matched sample, significance level was set at  $\alpha$  <.05, false discovery rate (FDR) corrected.

Clinical variables, ie, life time cumulative duration of hospitalizations,<sup>67,68</sup> duration of current episode,<sup>69</sup> years of education,<sup>70,71</sup> and verbal IQ<sup>72</sup> were tested for potentially moderating effects on the associations between brain structure and psychopathological factor. Therefore, eigenvariates (weighted mean) as an approximation of mean value inside the clusters were extracted. Moderator models (PROCESS macro v3.3 for SPSS,<sup>73</sup> model number 1) were corrected for the same covariates as VBM analyses.

Voxel-Based Morphometry: ROI Analyses. We tested whether our 5 psychopathological factors<sup>2</sup> were associated with GMV across diagnoses, using a ROI approach. To objectively select ROIs, we performed a comprehensive literature search using MEDLINE (PubMed.gov interface) and additionally went through references in the articles identified. For inclusion of a ROI for our analyses, the following literature selection criteria were applied: (1) meta-analyses published after 2010, or if a meta-analysis for a factor was not available review article in a high ranking journal; (2) investigating at least one of the 5 syndromes in patients with MDD, SSD, or BD dimensionally (ie, correlating psychopathological scores with GMV; studies performing a mere patient-healthy control design were not included); and (3) the ROIs had to be replicated in 2 individual original studies, published after 2005. Based on these criteria, the literature search revealed the meta-analyses and ROIs listed in table 2.

Masks for the ROIs were created using the "dartel space neuromorphometrics" atlas (http://www.neuromorphometrics.com/) in CAT12. Using the batch mode, the search space for each factor and the selected ROIs was restricted beforehand. We accounted the same covariates as for whole brain analyses using a P < .05, peak-level FWE corrected, and k = 10 threshold. Since all ROIs were reported to be negatively associated with the symptom dimensions, we performed one sided t tests. Furthermore, interactions analyses were performed to test if ROIs were driven by DSM-IV diagnostic categories.

**Table 2.** Psychopathological Factors Derived From Our Patients,<sup>2</sup> Meta-Analysis or Review Article on Structural Brain Correlates of These Syndromes and Selected Regions for Our Region-of-Interest Analysis

Psychopathological Factor	Literature	Region  Right middle frontal gyrus Left hippocampus Bilateral superior frontal gyri Left middle frontal gyrus Bilateral thalami Bilateral insulae			
Depression (F1)	Schmaal et al. (2017) <sup>46</sup>				
Negative syndrome (F2)	İnce and Üçok (2018) <sup>37</sup>				
Positive formal thought disorder (F3)	Sumner et al. (2018) <sup>34</sup> ; Cavelti et al. (2018) <sup>35</sup>	Bilateral frontal opercula Bilateral orbitofrontal cortices Left superior temporal gyrus Left planum temporale			
Paranoid-hallucinatory syndrome (F4)	Mucci et al. (2019) <sup>25</sup>	Left amygdala-hippocampus complex Left anterior cingulate gyrus Bilateral superior temporal gyri Bilateral amygdalae Bilateral thalami proper Bilateral insulae Bilateral planum temporale Left anterior cingulate gyrus Left insula			
Increased appetite (F5)	Gibson et al. (2010) <sup>52</sup>	Bilateral superior temporal gyri Left angular gyrus Left postcentral gyrus Bilateral nuclei accumbens Bilateral amygdalae			

#### Results

# Whole Brain Analyses

In the total sample, the pFTD factor (F3) was negatively associated with GMV in the right middle frontal gyrus (MFG) (k = 30 voxels, x/y/z = 34/46/16, T = 4.85, Z = 4.82, P = .01, FWE). The paranoid-hallucinatory syndrome factor was negatively correlated with GMV in the right fusiform gyrus (k = 24 voxels, x/y/z = 38/-21/-20, T = 5.08, Z = 5.05 P = .005, FWE) and in the left MFG (k = 27 voxels, x/y/z = -32/62/-3, T = 4.79, Z = 4.76 P = .02, FWE) (see figure 1). No FWE-corrected association was present for the other factors. Interaction analyses of diagnostic group × factor in the total sample (N = 1069) showed no significant results.

To rule out potential effects of differences in the number of patients per diagnoses, we performed multiple regression and ANCOVA interaction analyses again in the age- and sex- matched subsample (n = 357). We replicated the negative association of the right MFG Cluster and pFTD in the matched sample, too (k = 30 voxels, x/y/z = 33/46/16, T = 4.25, Z = 4.2  $P = 1.198*10^{-7}$ , FDR). The negative association of the paranoid-hallucinatory syndrome and the right fusiform gyrus (k = 24 voxels, x/y/z = 38/-21/-20, T = 4.14, Z = 4.09  $P = 1.384*10^{-4}$ , FDR), as well as the left MFG cluster (k = 27 voxels, x/y/z = -33/57/0, T = 4.73, Z = 4.65  $P = 9.022*10^{-6}$ , FDR) were also present in the matched sample. Interaction analyses of diagnostic group × factor showed no significant results.

Post hoc moderator analyses of illness variables showed a significant moderation of life time cumulative duration of hospitalizations on the association of F4 "paranoidhallucinatory syndrome" and the right fusiform cluster  $(R^2 = .018, F = 20.76; P < .001)$  and the left MFG cluster  $(R^2 = .009, F = 10.01; P = .002)$ . The duration of current episode moderated associations of GMV clusters and F4 (right fusiform gyrus:  $R^2 = .01$ , F = 6.54; P = .012; left MFG:  $R^2 = .011$ , F = 7.23; P = .007). Both life time cumulative duration of hospitalizations ( $R^2 = .002$ , F = 1.86; P = .173) and the duration of the current episode ( $R^2 = .002$ , F = 0.12; P = .727) did not moderate the right MFG cluster (F3). Years of education moderated the association of pFTD and the right MFG ( $R^2 = .007$ , F = 8.16; P = .004). There was no moderating effect of verbal IQ.

## ROI Analyses

Based on our literature search, we applied separate ROI analyses for each factor, in the total sample. We found FWE peak-level corrected GMV negative associations in the selected ROIs for the negative syndrome, pFTD, and the paranoid-hallucinatory syndrome (see table 3). The selected ROIs for the depression and increased appetite factor revealed no significant association. Interaction analyses (diagnoses  $\times$  psychopathological factor) revealed no interaction in all ROIs in both the total sample and in the age- and sex- matched sample (same n per diagnosis).

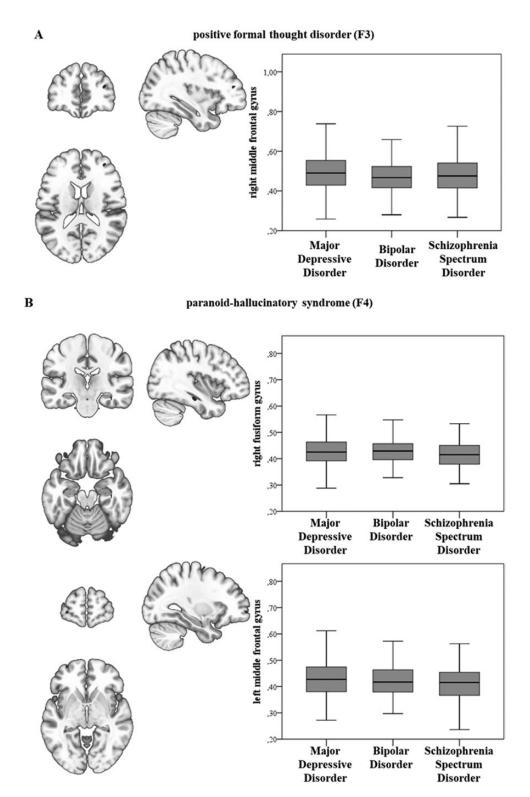


Fig. 1. (A, B) Negative correlations between gray matter volume for (A) positive formal thought disorder and the right middle frontal gyrus and (B) between the paranoid-hallucinatory syndrome and the right fusiform and left middle frontal gyrus, family-wise error peak-level corrected. Bar graphs on the right represent extracted eigenvariates of the clusters for each diagnostic group.

#### Discussion

Given the limitations of meta-analytic studies pooling single DSM diagnoses, our study investigated the relationship between regional brain volumes and dimensional, transdiagnostic psychopathological syndromes in one large sample comprising patients with MDD, BD, and SSD. Building on our previous work, we had derived 5 psychopathological factors (ie, depression, negative syndrome, pFTD, paranoid-hallucinatory syndrome, and increased appetite).2 On whole brain level, we found negative associations for pFTD with the right MFG and the paranoid-hallucinatory syndrome with the right fusiform gyrus and the left MFG. Using ROI analyses, we were able to confirm a number of previous results across diagnoses: The negative syndrome was negatively associated with the bilateral frontal opercula, pFTD with the left amygdala-hippocampus complex and the paranoid-hallucinatory syndrome with the bilateral thalami proper, the left postcentral gyrus, the left posterior cingulate gyrus, and the left angular gyrus. Based on these findings, we can draw 3 new insights. First, there was no interaction effect of DSM-IV diagnostic categories × psychopathological factors in all GMV associations on whole brain level and within the ROI analyses in both the total sample and in the matched sample containing the same n per diagnosis, strengthening our assumption of shared transdiagnostic, psychopathological factor-local GMV associations<sup>74-77</sup> within MDD, BD, and SSD. Additionally, whole brain clusters found in the total sample could be replicated in the matched sample, too. This mirrors meta- or mega-analytic results from brain structural, 5,6,8,23 molecular genetic genome-wide association studies,<sup>3,78</sup> immunology,<sup>9</sup> and environmental factors, 4,79 showing large biological overlapping across these disorders. In contrast to these previous pooling studies, we included only patients from one large study. Interestingly, important clinical variables which have previously been related to brain structural alterations, such as life time cumulative duration of hospitalizations<sup>67,68</sup> and duration of the current illness episode, <sup>69</sup> did indeed moderate the extracted GMV clusters in our study. This validates our findings and confirms the significance of illness aspects other than the clinical diagnosis. Based on these converging findings across modalities, we hypothesize that genetic and environmental factors impact the developing brain at different times and intensities across individuals affected regional brain structure, with the particular location/network depending on the individual factors. The involved network during development determines the psychopathological syndrome predominant later in life after disorder onset. Second, the approach applied here allowed us to investigate symptom complexes such as the paranoid-hallucinatory syndrome across disorders, which has previously been mostly investigated in SZ patients. But it is well known that MDD patients also show elevated psychotic symptoms. 10,11 Regression analyses were performed independent of the state of patients (ie, acute, chronic, and remitted). Thus, our approach is less prone to subgroup effects that may arise when applying categorical approaches. Third, by using whole brain and ROI-based approaches, we exploratively tested transdiagnostic brain structures on whole brain level and identified anatomical signatures across disorders previously reported to be associated with psychopathological syndromes within 1 or 2 disorders.

We did not find associations between GMV and the depression factor, a finding in line with the vast majority of previous studies. 45,46,80 Structural alterations in patients with MDD are not correlated with current depressive symptomatology but rather chronicity and cumulative severity. 45,47,81,82

Within the ROI analysis, the negative syndrome was associated with the bilateral frontal opercula volumes. This is a well-known finding from structural imaging studies investigating SZ patients. 83–87 GMV reductions in the frontal opercula correlated with persistent negative symptoms. 37,83,88

 Table 3. Significant Gray Matter Volume Reductions in the ROIs, FWE Peak-Level Corrected

Factor	ROI	Coordinates	P FWE	k	T	Z
Negative syndrome (F2)	Bilateral frontal opercula	[-48; 18; 21]	.031	21	3.512	3.501
Positive formal thought disorder (F3)	Left amygdala-hippocampus complex	[-34.5; -22.5; -15]	.018	99	3.386	3.376
Paranoid-hallucinatory syndrome (F4)	Bilateral thalami proper	[-6; -12; 12]	$.006^{a}$		3.995	3.979
		[-13.5; -25.5; 10.5]	.007	1201	3.943	3.928
		[9; -3; 1.5]	.013		3.760	3.747
	Left angular gyrus	[-55.5; -64.5; 27]	.026	15	3.627	3.615
	Left postcentral gyrus	[-61.5; -19.5; 31.5]	.003a		4.267	4.248
		[-58.5; -21; 39]	$.005^{a}$	405	4.172	4.154
		[-57; -22.5; 25.5]	.007		4.076	4.059
	Left posterior cingulate gyrus	[-13.5; -40.5; 3]	.011	11	3.633	3.621

*Note*: FWE, family-wise error; ROI, region of interest. ROIs tested are listed in table 2. <sup>a</sup>Significant difference after Bonferroni correction for multiple ROI testing.

The pFTD factor was negatively associated with the right MFG on whole brain level, and the left amygdala-hippocampus complex in the ROI analysis. Previously, pFTD in SZ has been associated with neuroanatomical alterations in Wernicke's and Broca's language areas, as well as the middle frontal gyri and the left amygdala-hippocampus complex. <sup>32–35</sup> We could replicate these findings in our transdiagnostic sample. As opposed to the left, we found the right MFG homologue associated with pFTD, an area which is involved in language processing in SZ patients. <sup>32,34,35</sup> Furthermore, there is evidence that the hippocampus plays a major role in word generation tasks which coincides with our findings. <sup>89,90</sup>

On a whole brain level, the paranoid-hallucinatory syndrome was negatively associated with the right fusiform gyrus and the left MFG. Previous studies<sup>24,25,27,31,40,91</sup> showed that medial temporal regions are correlated with "positive" symptomatology, which mainly comprises the paranoid-hallucinatory syndrome. Besides the inferior temporal gyri, the bilateral fusiform gyri have been reported as a significant correlate of positive symptoms (measured with Positive and Negative Syndrome Scale [PANSS]).40,91 A meta-analysis including 4474 paranoidhallucinatory SZ patients showed that reductions in both the right fusiform gyrus and the bilateral inferior temporal gyri have the largest effect sizes. 92 Several studies have shown negative associations of GMV and the paranoid-hallucinatory syndrome factor in the overall frontal volume<sup>93</sup> as well as GMV reductions in the left pars orbitalis and the left superior frontal gyrus.<sup>40</sup> The left MFG has been shown to be deactivated in preceding auditory hallucinations.94 The ROI analyses revealed significant associations between the paranoid-hallucinatory syndrome and the bilateral thalami proper, <sup>27,28,95</sup> the left angular gyrus, the postcentral gyrus, and the left posterior cingulate gyrus, <sup>25,27</sup> confirming many previous studies.

The factor "increased appetite" was, independent of medication, a dimension in our psychopathological factor analytic study² that has been shown to emerge if psychopathological scales are used that capture vegetative symptoms. 48,96 We did not find associations with GMV neither on whole brain level nor within the ROI analyses. This may indicate that biological influences other than GMV have an impact on the manifestation of increased appetite. Nevertheless, increased appetite is a relevant syndrome across disorders that deserves further investigation.

## Limitations

Some limitations must be noted. First, patient groups were unequally distributed which potentially biased our results since the MDD group was the largest. However, we were able to confirm the results of our interaction analyses (total sample) in an age- and sex-matched subsample (same *n* in each diagnosis). Second, the MDD group only marginally presented with psychotic symptoms compared

to BD and SSD patients resulting in restricted variance found for psychotic symptoms (ie, factor 4 "paranoidhallucinatory syndrome"). Third, the extracted factors were based on current psychopathology. Correlating current state measures with brain structure misses past psychopathology. However, specific current symptoms are an indication for a particular neuroanatomical, stateindependent alteration that outlasts current symptoms. Still, it would be of great interest to investigate the stability of both factor dimensions and their relationship to brain structure validating the present cross-sectional results. Fourth, pharmacological treatment was considered in our models using 3 dummy-coded variables accounting for the intake of antidepressants, mood stabilizers, and antipsychotics. However, the influence of both dosage and duration of intake on GMV is not considered.

#### Conclusion

In sum, our findings provide a novel anatomical mapping of psychopathological symptom dimensions across disorders. The main strength of the present study is the use of a large transdiagnostic cohort, innovative data-driven psychopathological factors, and the inclusion of patients across psychotic and affective disorders. Our findings give evidence for shared and diagnosis-independent GMV reductions associated with symptom dimensions. We try to overcome a current deadlock in scientific approaches which has one origin in a misguided reification of DSM diagnoses.

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