

1 **Exploring the association between brain-derived neurotrophic factor (BDNF) levels and**
2 **longitudinal psychopathological and cognitive changes in Sardinian psychotic patients**

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25 **Abstract**

26

27 **Background and Hypothesis:**

28 Schizophrenia is among the most debilitating mental disorders and has complex pathophysiological
29 underpinnings. There is growing evidence that brain-derived neurotrophic factor (BDNF) can play a
30 role in its pathogenesis. The present study investigated the longitudinal variation of serum BDNF levels
31 in a 24-month observational prospective cohort study of Sardinian psychotic patients and its relationship
32 with psychopathological and cognitive changes. Further, we examined whether genetic variation within
33 the *BDNF* gene could moderate these relationships.

34

35 **Study design:**

36 Every six months 105 patients were assessed for their BDNF serum levels, as well as for a series of
37 psychopathological, cognitive, and social measures. We performed a targeted analysis of four tag single
38 nucleotide polymorphisms (SNPs) within the *BDNF* gene that were selected and analyzed using
39 Polymerase Chain Reaction (PCR). Longitudinal data were analyzed using mixed-effects linear
40 regression models (MLRM).

41

42 **Study results:**

43 We observed a declining longitudinal trajectory of BDNF levels in psychotic patients in general, and in
44 relation to the severity of depressive and negative symptoms. BDNF serum levels also declined in
45 patients scoring lower in cognitive measures such as attention and speed of information processing and
46 verbal fluency. The rs7934165 polymorphism moderated the significant association between verbal
47 fluency and BDNF levels.

48

49 **Conclusions:**

50 These findings in patients from real-world settings suggest a plausible role of peripheral BDNF levels
51 as a marker of illness burden in schizophrenia spectrum disorders.

52

53 **Keywords:** BDNF; complex disorders; schizophrenia; biomarkers; longitudinal trajectories

54 Introduction

55 Schizophrenia (SCZ) and schizoaffective disorders (SAD) are relatively heterogeneous psychiatric
56 disorders characterized by an array of symptoms including delusions, hallucinations, psychomotor,
57 social, and cognitive deficiencies [1]. SCZ affects about 1% of the population [1] while SAD has a
58 lifetime prevalence of about 0.3% [2]. SCZ is one of the most debilitating mental disorders. However,
59 our comprehension of its pathophysiological underpinnings remains inadequate. SCZ and SAD have
60 complex pathogenesis that involves the interaction of multiple biological, genetic, and environmental
61 factors [3]. Indeed, they both demonstrate high heritability according to family, twin, and adoption
62 studies [4,5], indicating a major genetic contribution to the illness risk. In fact, in SCZ, genome-wide
63 association studies (GWAS) have successfully identified genetic variants contributing to the risk of
64 developing SCZ, with 287 distinct genomic loci associated with the disorder [6].

65 In this context, the brain-derived neurotrophic factor (BDNF) seemingly plays a relevant role [7]. BDNF
66 is the most prevalent and extensively studied neurotrophin in the human central nervous system (CNS),
67 and is able to cross the blood-brain barrier [8]. BDNF is a key regulator of a wide range of
68 neurophysiological processes including neurogenesis, neuronal differentiation [9], synaptogenesis, and
69 long-term potentiation [10]. Like other neurotrophins, BDNF is synthesized as a precursor form, prepro-
70 BDNF which is further cleaved to produce mature BDNF either intracellularly or extracellularly [11].

71 Even though a strong correlation between peripheral BDNF and BDNF levels in the CNS has been
72 reported [12], studies measuring BDNF levels in the serum as a potential biomarker of SCZ have yielded
73 controversial results. Several studies have found decreased peripheral levels of BDNF in SCZ patients
74 including first-episode psychosis (FEP) patients [13–15] and chronic patients that have been medicated
75 for a substantial period of time [16,17]. Nevertheless, some studies found no difference between serum
76 BDNF levels of SCZ patients and those of healthy controls [18,19]. and some even found an increase
77 [20]. Meta-analytical findings demonstrated that there was a substantial decrease in BDNF serum and
78 plasma levels of SCZ patients in acute episodes, suggesting that decreased peripheral BDNF levels can
79 be considered as a biomarker of disease activity [21].

80 The finding of a decline in BDNF levels of SCZ patients is consistently observed. However, little is
81 known about the temporal trajectory, and the modulators, of this decline. Indeed, several factors,

82 including genetic, treatment, and clinical moderators, might affect the peripheral levels of BDNF in SCZ
83 patients. BDNF levels appear to be affected by Val66Met (rs6265) polymorphism, a single nucleotide
84 polymorphism (SNP) in the *BDNF* gene leading to valine (Val) for methionine (Met) substitution at
85 codon 66 [22]. The Val66Met polymorphism has been associated with intracellular trafficking and
86 activity-dependent secretion of mature-BDNF as well as neurocognitive deficits [23–25]. While some
87 studies have found an association between the functional Val66Met variant and peripheral BDNF levels
88 [26,27], others have not [22,28]. Consistently, two meta-analyses looking at the association between
89 *BDNF* Val66Met polymorphism and several neurocognitive phenotypes found no significant difference
90 between carriers of Met allele and Val/Val homozygotes [29,30].

91 Peripheral BDNF levels might also be affected by treatment factors, and psychopathological and
92 cognitive changes. Cognitive impairments are widely observed in patients with SCZ, and together with
93 symptom severity, they are central to the prediction of the clinical and functional outcomes in SCZ
94 [31,32]. Several studies have shown the strong correlation between reduced peripheral BDNF levels and
95 impaired neurocognitive and psychopathology test scores [33,34].

96 Overall, the analysis of peripheral BDNF levels and their relationship with clinical and treatment factors
97 in schizophrenia spectrum disorders has provided inconsistent results. These discrepancies assume even
98 higher significance in consideration of the possible clinical relevance of using BDNF as
99 diagnostic/prognostic marker in psychiatric disorders. For instance, it is conceivable that patients with
100 a more severe course of illness or with higher genetic predisposition for SCZ might have lower levels
101 of BDNF compared to those with less severe presentation (or with less genetic loading). The present
102 study sought to investigate the longitudinal variation of serum BDNF levels in a 24-month observational
103 cohort study named Longitudinal Assessment of BDNF in Sardinian Psychotic patients (LABSP) [35].
104 Several aims were tested, primarily the assessment of the variation of BDNF serum levels over time
105 and its relationship with psychopathological changes, cognitive function, and social functioning. We
106 hypothesized that BDNF serum levels would decrease in association with longer and more severe
107 clinical course as well as in association with other possible factors such as cognitive decline. Further,
108 we also examined if genetic variation (four tag SNPs) within the *BDNF* gene could moderate these
109 relationships. Finally, as an aside, we performed discriminatory analysis of SCZ and SAD using

110 Receiver Operating Characteristic (ROC) curve, expecting that BDNF serum levels could differentiate
111 individuals affected by the two disorders.

112 **Methods and materials**

113

114 **Sample**

115 An a priori power analysis was conducted using repeated measures and sample size (RMASS) software
116 and the results indicated that the sample size of 59 individuals was sufficient to achieve 90% statistical
117 power to detect significant difference at $\alpha=0.05$. Our sample was comprised of 105 patients with
118 psychosis treated at the community mental health center of the Unit of Psychiatry of the Department of
119 Medical Science and Public Health, University of Cagliari and University of Cagliari Health Agency,
120 Cagliari, Italy. A Structured Clinical Interview for DSM-IV-TR Axis I Disorders, Patient Edition
121 (SCID-I/P) [36] was administered by trained mental health professionals to confirm the diagnosis of
122 SCZ or SAD. To be considered eligible for participation in the LABSP study, patients had to be between
123 18 and 65 years old; diagnosed with SCZ or SAD according to DSM-IV-TR, and with absence of acute
124 psychopathological manifestations for the past six months before recruitment. Patients were excluded
125 from the study if they refused to provide consent; had acute psychopathological symptoms, major
126 unstable medical illness, severe mental retardation, major neurological disorder or a previous head
127 injury, current drug and alcohol dependence, or severe illness-related cognitive impairment which
128 affected their ability to participate in the study. The study was approved by the University of Cagliari
129 Health Agency Ethics Committee and the protocol followed the principles of the Declaration of
130 Helsinki. Written informed consent was obtained from all the patients.

131

132 **Assessment procedures**

133 Recruited patients were assessed and evaluated using various measures at 5 different waves. Clinical,
134 cognitive and social performance measures, and blood samples of the patients were collected at the
135 baseline (T0), and at four consecutive time points: 6 months (T1), 12 months (T2), 18 months (T3), and
136 24 months (T4). The details of the assessment and evaluation process including used measures and
137 materials [35], as well as primary findings [37], have been previously published. General

138 psychopathology, the severity of positive and negative symptoms, and clinical status of the patients were
139 assessed using the original 30 item Positive and Negative Syndrome Scale (PANSS) [38] and Clinical
140 Global Impression Scale for Schizophrenia (CGI-SCH) [39]. In addition, we applied the consensus five-
141 factor model of PANSS [40] consisting of 20 items that are categorized into Positive, Negative,
142 Disorganized/Concrete, Excited, and Depressed factors, because previous studies have shown that
143 PANSS-FCTer better characterizes the structure of PANSS data [40,41]. The Brief Assessment of
144 Cognition in Schizophrenia (BACS) scale [42] was used to evaluate changes in cognitive domains
145 including verbal memory, working memory, reasoning, and processing speed. The evaluation of social
146 functioning was carried out using Personal and Social Performance scale [43] that has shown to be a
147 valid and reliable measure for patients with SCZ [44,45].

148

149 **Sample collection and measurement of BDNF**

150 For the assessment of BDNF serum levels, the blood from each patient was drawn at the same time of
151 the day (between 8:00 and 10:00 AM) at each visit. Collected blood samples were kept at room
152 temperature for about 4 hours to allow for clotting, after which they were centrifuged at approximately
153 1000 X g for 15 min. All samples were immediately stored in small aliquots at -20°C until analyzed.
154 Then, the serum BDNF levels were determined using a commercial human enzyme-linked immunoassay
155 (ELISA) kit (Booster Immunoleader, Cat. N° EK0307) following the manufacturer's instructions. This
156 kit is used for the quantitative detection of human BDNF in cell culture supernatants, serum, and plasma
157 with a high sensitivity of <2pg/mL, the measuring interval of 31.2-2000pg/mL, and no detectable cross-
158 reactivity with other relevant proteins. The absorbance was measured using a microplate reader (Thermo
159 Scientific Multiskan FC) set at 450nm within 30 minutes after the final step of the kit procedure.

160

161 **Genetic analysis**

162 We used the Tagger program implemented in the Haploview v4.2 to select SNPs in linkage
163 disequilibrium (LD) ($r^2 \geq 0.8$) and with a minor allele frequency threshold of 0.01. The genotyping of
164 SNPs rs1519480, rs11030104, rs6265 (Val66Met), and rs7934165 was performed using TaqMan probes
165 on demand (C_11592757_20, C_1751792_10, C_11592758_10, C_1197567_10, ThermoFisher

166 Scientific) on a StepOne Plus instrument (ThermoFisher Scientific). The reaction mixture was prepared
167 in a final volume of 10 μ l consisting of 5 μ l of MasterMix (2x), 0.5 μ l of Custom TaqMan® SNP
168 Genotyping Assay (20x) containing primers marked as VIC and FAM to discriminate between alleles,
169 1 μ l of cDNA, and 3.5 μ l of RNase-free water. Polymerase Chain Reaction (PCR) was performed with
170 the following conditions: 30 sec. 60°C, 10 min 90°C, and 40 cycles at 95°C for 15 sec and 60°C for 1
171 min.

172

173 **Statistical analysis**

174 We analyzed the longitudinal data using mixed-effects linear regression models (MLRM). MLRM was
175 used particularly because it allows the observation of the effects and interaction of multiple independent
176 variables on a dependent variable while considering repeated measures across participants [46]. To test
177 our hypotheses, we regressed our predictor variables and fixed effects on our log-transformed serum
178 BDNF data. In a preliminary step, PANSS, CGI-SCH, BACS, and PSP scale scores of each subject at
179 each time point were separately regressed on BDNF data to analyze the relationship between them. We
180 fitted regression models while adjusting for age and sex and checked for linearity and homoscedasticity
181 by examining plots of residuals against fitted values. Lastly, *BDNF* gene polymorphisms were added to
182 the MLRM as a covariate to examine the possible moderating effect. The longitudinal data were
183 analyzed using the statistical programming language R [47]. All regression models were fitted using
184 “lme4” package [48]. A significance level of $P < 0.05$ was considered after Holm–Bonferroni
185 corrections for multiple comparisons. ROC analysis, with Sensitivity, Specificity and Predictive Value
186 Analysis of the ability to discriminate between SCZ and SAD in relation to longitudinal BDNF levels
187 was also applied.

188

189 Results**190 Sample characteristics**

191 Table 1 summarizes the demographic and clinical characteristics of the patients that participated in this
192 study. The sample consisted of 105 patients including 64 with a diagnosis of SCZ and 41 with SAD.
193 The mean age of the sample at the baseline was 48.85 ± 10.45 years.

194

195 Associations between psychopathological symptoms and serum BDNF levels

196 The MLRM analysis showed a statistically significant decline in BDNF levels over time ($Z = -4.9$, $p =$
197 9.02×10^{-7}). As shown in Table 2, analysis of the relationship between scores of original three-subscale
198 PANSS and longitudinal BDNF levels yielded no significant association. However, when we examined
199 the association between serum BDNF levels and five-factor PANSS scores we found a significant
200 relationship between longitudinal BDNF levels and negative factor ($Z = -2.245$, $p = 0.025$). MLRM
201 found a significant relationship between CGI depressive symptoms and BDNF levels ($Z = -2.796$, $p =$
202 0.005). This association remained significant when we added age and sex as covariates ($Z = -2.819$, $p =$
203 0.009). The results also showed a significant association between CGI negative symptoms and serum
204 BDNF levels ($Z = -2.057$, $p = 0.039$). In terms of serum BDNF levels and scores of positive symptoms
205 subscale of CGI-SCH, the results did not show a significant relation between them.

206

207 Associations between cognition, social functioning, and serum BDNF levels

208 The analyses showed that the BDNF serum levels decreased in patients scoring lower on symbol coding
209 ($Z = 2.194$, $p = 0.028$) and semantic fluency ($Z = 3.228$, $p = 0.001$) subscales of BACS. The association
210 between semantic fluency and BDNF levels remained significant after correcting for age and sex ($Z =$
211 3.167 , $p = 0.003$). An examination of the relationship between the measure of social functioning and
212 longitudinal BDNF levels yielded no significant associations.

213

214 Moderating effect of genetic variation

215 The moderating effect of the SNPs within *BDNF* gene was analyzed using genotypic and allelic effect
216 models including additive, dominant, and recessive models (Supplementary Table 2). We found a

217 significant moderating effect of rs7934165 on the relationship between BACS subscale for semantic
218 fluency and serum BDNF levels when analyzed using the recessive model ($Z = -2.359$, $p = 0.0367$). This
219 interaction effect remained significant after adjusting for age and sex ($Z = -2.339$, $p = 0.0466$).

220

221 **ROC curve**

222 We performed ROC analysis (Supplementary Figure 1) to examine the ability of our model to
223 discriminate between SCZ and SAD in relation to longitudinal BDNF levels. Area under the curve
224 (AUC) was calculated to evaluate the overall accuracy of the diagnostic test in discriminating between
225 SCZ and SAD patients. Optimal diagnostic cut-off value was 2.8303 with a sensitivity 65.2% and
226 specificity of 50.4%. Algorithm of this model had an AUC of 57.1% (95%, CI: 0.5183-0.624), which
227 indicates poor diagnostic performance.

228

229 **Discussion**

230

231 In this study we sought to clarify whether longitudinal BDNF serum levels of psychotic patients were
232 correlated with treatment-related, psychopathological, cognitive, and social changes. Previous analysis
233 of LABSP data examined the impact of antipsychotics on BDNF serum levels and found a significant
234 longitudinal increase in those treated with depot/long-acting injectables, but not oral antipsychotics [37].
235 Our current study had several main findings. First, we found an overall decline in the trajectory of serum
236 BDNF levels over time (Supplementary Figure 3). The results revealed that this decline was more
237 pronounced in patients with more severe depressive and negative symptoms. In addition, BDNF serum
238 levels were more declined in patients with lower scores in two cognitive domains including speed of
239 processing and verbal fluency. Finally, when we examined the possible moderating effect of genetic
240 polymorphisms within *BDNF* gene on these statistically significant associations, we found that
241 rs7934165 polymorphism had a significant moderating effect on the association between verbal fluency
242 and BDNF serum levels.

243 As mentioned before, the reason for the temporal decline in BDNF serum levels is unknown, but it might
244 be hastened by disease progression or associated with other factors such as drug treatments and severity
245 of clinical symptoms. A meta-analytic study by Fernandes et al. [49] established that peripheral BDNF

246 levels of SCZ patients were moderately decreased in comparison to healthy controls and the decline was
247 associated with the temporal course of the disease. Results of a recent meta-analysis by Rodrigues-
248 Amorim et al. [50] showed that BDNF levels of both drug-naïve and medicated schizophrenia patients
249 were reduced throughout the disease course. Indeed, the reduced BDNF expression have also been
250 associated with neuroinflammation [51], increased cortisol levels [52], while enriched environment has
251 been shown to increase BDNF levels in psychiatric and neurodegenerative disorders [53]. Nevertheless,
252 the interaction between these factors should be better understood.

253 We did not find an association between BDNF levels and psychopathological symptoms when measured
254 by 30 item three-subscale PANSS. Consistent with our results some studies also did not observe any
255 significant association between PANSS scores and peripheral BDNF levels [15,18,54]. Considering that
256 five-factor PANSS has shown to be better at representing the dimensional structure of PANSS data, we
257 utilized this model for our analysis as well. Surprisingly, we found a significant relationship between
258 negative factor of the five-factor scale and reduced serum BDNF levels. This is a rather interesting
259 outcome, as we also found a significant association between the severity of depressive and negative
260 symptoms and reduced BDNF serum levels when we regressed CGI-SCH subscales on BDNF data. To
261 our best knowledge, this is the first longitudinal study assessing the relationship between BDNF levels
262 of psychotic patients and the severity of psychopathological symptoms using CGI-SCH. In their recent
263 clinical study Fang et al. [55] observed a significant association between reduced plasma BDNF levels
264 and depressive symptoms in SCZ patients. Similar results were observed in another study where a
265 significant negative association between depressive symptoms and BDNF serum levels of chronic SCZ
266 patients was found [56].

267 A possible explanation for the observed correlation between negative and depressive symptoms and
268 peripheral BDNF levels with CGI-SCH and not PANSS might be because CGI-SCH could be a more
269 reliable measure than the 30-item PANSS for monitoring the longitudinal course of psychopathology in
270 SCZ and SAD. [57] The five-factor/20-item PANSS model have also demonstrated to be a better fitting
271 model for the symptoms of SCZ [41,58,59]. which could explain the reason the association between
272 negative factor and serum BDNF levels was detected by this model and not the entire 30-item PANSS
273 scale.

274 Cognitive impairment is a core symptom of SCZ and a number of studies have found an association
275 between neurocognitive deficits and peripheral BDNF levels [33,60,61]. We found a significant
276 correlation between BDNF levels and two cognitive domains including symbol coding and semantic
277 fluency as measured by BACS. A recent meta-analysis examining the relationship between
278 neurocognitive deficits and BDNF levels revealed that higher peripheral BDNF levels were associated
279 with better performance on reasoning and problem-solving tasks in people with SCZ [29]. There is
280 growing evidence associating BDNF to cognitive dysfunctions in psychotic patients at different stages
281 of the disease. According to these findings, it can be assumed that peripheral BDNF levels can be
282 considered as a potential biomarker for neurocognitive deficits in psychosis.

283 This study did not show any significant association between serum BDNF levels and Val66Met
284 polymorphism within the *BDNF* gene. Our findings are in agreement with the results of meta-analysis
285 and GWAS analysis of the Sardinian sample conducted by Terracciano et al. [62], where no association
286 was found between Val66Met polymorphism and serum BDNF levels.

287 We first observed a significant moderating effect of Val66Met on the relationship between BDNF levels
288 and BACS subscale for symbol coding, as well as BDNF levels and CGI depressive symptoms.
289 However, these associations did not survive correction for multiple comparisons. Nevertheless, we
290 found that, when using recessive model, rs7934165 polymorphism within the *BDNF* gene moderated
291 the significant association between serum BDNF levels and verbal fluency cognitive domain. SNPs in
292 the *BDNF* gene have been previously linked to peripheral BDNF levels as well as psychopathological
293 and cognitive aspects of SCZ [26,27,63,64]. One possible interpretation of these findings is that the
294 genetic variants are to some extent incorporating the effect of the predictors. This would be consistent
295 with the putative influence exerted by genetic variation on BDNF levels.

296 Our results should be interpreted considering several limitations. A major limitation of this study is the
297 lack of control group. In addition, the sample size for this study is moderate, especially regarding the
298 analysis of genetic variations of the *BDNF* gene. However, this is compensated by the longitudinal
299 design and the presence of 5 time points for assessment. Moreover, the sample was rather heterogeneous
300 regarding the duration and stage of illness. Indeed, some patients were in the early years of their illness
301 course while others had in some instances decades of clinical history. Hence, we tested the interaction

302 effect of duration of untreated psychosis and duration of illness on BDNF levels but did not find any
303 significant effect (Supplementary Table 3). Likewise, in a recent meta-analysis by Rodrigues-Amorim
304 et al [50], they regressed the duration of illness on serum BDNF levels and did not find a significant
305 effect. Nonetheless, larger sample size and more homogenous sample will be required for future studies
306 to overcome these limitations. The relatively small sample size did not make possible to perform
307 subgroup analyses and led to the inclusion of a limited number of covariates in MLRM models to
308 prevent saturation. Even though there is a substantial genetic overlap between SCZ and SAD [5], further
309 research should be undertaken to explore the differences between these subgroups. In addition to the
310 latter point, not all confounding variables could be added to the same model to avoid overfitting and
311 saturation. Finally, even considering the longitudinal design, and the MLRM modelling applied, it is
312 not possible to exclude that some of the time-varying variables were not entirely captured by our
313 analysis.

314 Another limitation of our study is that only serum BDNF levels of the patients were collected, while
315 plasma BDNF levels were not assessed. We are not sure about the extent to which serum BDNF reflects
316 the processes in CNS. While some authors have proposed that plasma BDNF is a more reliable proxy
317 of what happens in CNS [49], others consider serum BDNF to be a better correlate of cortical BDNF
318 levels [65]. Finally, the inability of ELISA kits to distinguish between pro and mature BDNF is another
319 limitation of our study. Unlike mature BDNF, pro-BDNF plays role in inducing apoptosis, reducing
320 dendritic spines, and other processes that may contribute to long-time depression (LTD) [66]. Being
321 able to measure BDNF by differentiating between these two isoforms is essential as they may have
322 opposing effects, and future studies should consider using newly developed specific mBDNF and pro-
323 BDNF ELISA assays [67] when investigating the proposed associations.

324 **Conclusion**

325 Even considering these limitations, our study identified a longitudinal trajectory of decline of BDNF
326 levels associated with decline in some cognitive domains and higher severity of depressive and negative
327 symptoms in patients affected by SCZ and SAD. These findings in a real-world patient sample suggest
328 a plausible role of peripheral BDNF levels as a marker of illness burden in schizophrenia spectrum
329 disorders

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333 illness.

334

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337

338 **Conflicts of Interest**

339 All authors declare none.

340

341 **Data availability**

342 The data that support the findings of this study are available from the authors upon request in
343 anonymized form.

344

345 **References**

346

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561 **Table 1.** Main Demographic and Clinical Characteristics of LABSP Sample.

Variable (continuous)	N	Mean	SD
BDNF serum levels, ng/ml	105	25.45	13.67
Age, years	105	48.85	10.45
Age of onset, years	105	21.77	9.30
Duration of illness, months	105	308.51	134.33
Age at first treatment, years	105	24.23	8.95
Duration of untreated illness, months	105	29.07	54.60
Antipsychotics, chlorpromazine equivalents, mg/day	103	378.92	272.03
Variable (categorical)	N	%	
Sex (male)	74	70.5	
Presence of family history of mental disorders	64	61.0	
Presence of family history of schizophrenia	31	29.5	
Presence of family history of bipolar disorder	8	7.6	
Presence of family history of major depressive disorder	19	18.1	
Presence of family history of anxiety disorders	10	9.5	
Diagnosis of schizophrenia (SCID-I)	64	61.0	
Diagnosis of schizoaffective disorder (SCID-I)	41	39.0	
Diagnosis of obsessive-compulsive disorder (SCID-I)	5	4.8	
Diagnosis of cluster A personality disorders (SCID-II)	2	1.9	
Diagnosis of cluster B personality disorders (SCID-II)	2	1.9	
Diagnosis of cluster C personality disorders (SCID-II)	2	1.9	
Diagnosis of personality disorder NOS (SCID-II)	1	1.0	

Abbreviations: LABSP, longitudinal assessment of BDNF in Sardinian psychotic patients; BDNF, brain-derived neurotrophic factor; SD, standard deviation; SCID-I, Structured Clinical Interview for DSM-IV Axis I Disorders (SCID-I); SCID-II, Structured Clinical Interview for DSM-IV Axis II Disorders (SCID-II).

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563

564 **Table 2.** Results of Mixed Effects Linear Regression Models

Independent Variables	Unadjusted Models			Adjusted Models*		
	Estimated Coefficient	Z value	P value	Estimated Coefficient	Z value	P value
PANSS, total score	-0.001525	-1.198	0.231	-0.001686	-1.342	0.359
PANSS, positive symptoms	-	-0.194	0.846	-0.001650	-0.342	0.912
	0.0009322					
PANSS, negative symptoms	-0.005553	-1.590	0.112	0.005246	-1.494	0.270
PANSS, general psychopathology	-0.002449	-1.083	0.279	-0.002637	-1.177	0.478
PANSS-FCTcr, positive factor	-0.0007404	-0.101	0.92	-0.001647	-0.224	0.895
PANSS-FCTcr, negative factor	-0.009772	-2.245	0.0248	-0.009337	-2.118	0.068
PANSS-FCTcr, disorganized/concrete factor	-0.008265	-1.089	0.276	0.008029	-1.016	0.406
PANSS-FCTcr, excited factor	-0.005327	-0.742	0.458	-0.005763	-0.794	0.772
PANSS-FCTcr, depressed factor	-0.005875	-0.646	0.518	-0.006804	-0.755	0.771
CGI-SCH, severity positive symptoms	0.008801	0.487	0.626	0.005490	0.294	0.89
CGI-SCH, severity negative symptoms	-0.048374	-2.057	0.0397	-0.049882	-2.072	0.077
CGI-SCH severity depressive symptoms	-0.058876	-2.796	0.0052	-0.059671	-2.819	0.0096
CGI-SCH severity cognitive symptoms	-0.034471	-1.819	0.0689	-0.0340535	-1.733	0.166
CGI-SCH, global severity	-0.037967	-1.498	0.127	-0.039493	-1.559	0.244
BACS, verbal memory	0.001736	0.154	0.878	-0.0003715	-0.032	1
BACS, digit sequencing task (number of correct responses)	0.004699	1.034	0.301	0.004188	0.906	0.730
BACS, digit sequencing task (longest sequence recalled correctly)	0.00972	0.739	0.46	0.008199	0.612	1
**BACS, verbal fluency (controlled oral word association test)	0.011660	3.228	0.0013	0.0113920	3.167	0.003
BACS, attention and speed of information processing (symbol coding)	0.004147	2.194	0.0282	0.004306	2.142	0.064
BACS, executive functions, Tower of London	0.004319	1.024	0.306	0.0037557	0.846	0.795
PSP, total score	0.002010	1.251	0.211	0.001436	1.206	0.456

565 Brief Assessment of Cognitive in Schizophrenia; PSP, Personal and Social Performance Scale

566 *Adjusted for age and sex

567 ** BACS, verbal fluency (category instances) variable was not included in the table as there were no sufficient observations to support the

568 model

569 Significant P values in bold

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