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## Cognitive and serum BDNF correlates of BDNF Val66Met gene polymorphism in patients with schizophrenia and normal controls

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### Abstract

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**Background**—Studies suggest that a functional polymorphism of the brain-derived neurotrophic factor gene (BDNF Val66Met) may mediate hippocampal-dependent cognitive functions. A few studies have reported its role in cognitive deficits in schizophrenia including its association with peripheral BDNF levels as a mediator of these cognitive deficits.

**Methods**—We assessed 657 schizophrenic inpatients and 445 healthy controls on the Repeatable Battery for the Assessment of Neuropsychological Status (RBANS), the presence of the BDNF Val66Met polymorphism and serum BDNF levels. We assessed patient psychopathology using the Positive and Negative Syndrome Scale (PANSS).

**Results**—We showed that visuospatial/constructional abilities significantly differed by genotype but not genotype×diagnosis, and the Val allele was associated with better visuospatial/constructional performance in both schizophrenic patients and healthy controls. Attention performance showed significant a genotype by diagnosis effect. Met allele-associated attention impairment was specific to schizophrenic patients and not shown in healthy controls. In the patient group, partial correlation analysis showed a significant positive correlation between serum BDNF and the RBANS total score. Furthermore, the RBANS total score showed a statistically significant BDNF level × genotype interaction.

**Conclusions**—We demonstrated an association between the BDNF Met variant and poor visuospatial/constructional performance. Furthermore, the BDNF Met variant may be specific to attentional decrements in schizophrenic patients. The association between decreased BDNF serum levels and cognitive impairment in schizophrenia is dependent on the BDNF Val66Met polymorphism.

## Keywords

Schizophrenia; brain-derived neurotrophic factor; cognition; association; genotype

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Cognitive deficits in learning, memory, attention, executive functioning, spatial working memory and cognitive processing speed are core features of schizophrenia (Sharma and Antonova 2003; Kraus and Keefe 2007). These deficits occur prior to the onset of other symptoms of schizophrenia (Lieberman et al 2001; Dickerson et al 2011), and generally persist during the course of the illness (Hughes et al 2003). Indeed, cognitive impairment associated with schizophrenia is a major obstacle to rehabilitation with its severity being a predictor of poor clinical outcome (Kaneda et al 2009; Granholm et al 2008; Harvey et al 2003; Harvey 2009). Although studies have defined cognitive constructs that are impaired in this disease, the pathophysiological mechanisms underlying these cognitive deficits are less clear.

BDNF is a member of the neurotrophin family of growth factors. BDNF is critical in modulating memory-associated neuroplasticity through regulating cell survival, proliferation and synaptic growth in the developing central nervous system (Poo 2001; Egan et al 2003). More specifically, BDNF appears to play an important role in both early long-term potentiation (LTP) and late phase LTP in hippocampal neurons (Lu and Gottschalk 2000), a cellular model of learning and memory (Poo 2001). Inhibition of BDNF signaling by gene knockout or antisense RNA impairs spatial learning and memory (Guzowski et al 2000). Also, BDNF has been prominently implicated in survival and function of midbrain dopamine neurons related to cognitive function (Altar et al 1997). Thus, preclinical evidence suggests that BDNF activity or levels may contribute to alterations in hippocampal function and hippocampal-dependent learning and memory (Hariri et al 2003). Indeed, previous studies indicate that a SNP (rs6265) producing a valine (Val)-to-methionine (Met) substitution in the proBDNF protein at codon 66 (Val66Met) is related to hippocampal-mediated memory performance in humans (Egan et al 2003; Hariri et al 2003). Consistent

with this, both normal controls and schizophrenic patients with Met alleles have significant deficiencies in episodic memory (Egan et al 2003; Hariri et al 2003; Dempster et al 2005). Moreover, imaging studies show that individuals with this allele exhibit abnormal hippocampal activation, and reduced brain volume in regions associated with memory and attention (Szeszko et al 2005; Bueller et al 2006). Taken together, these findings support a role for the *BDNF* Val66Met polymorphism in normal hippocampal-dependent memory function in healthy controls as well as in individuals with schizophrenia.

Recent studies indicate that circulating BDNF may be a biomarker of general cognitive function in healthy adults (Yu et al 2008). The finding that serum BDNF levels are significantly decreased in individuals with diseases associated with progressive cognitive decline, such as mild cognitive impairment (Yu et al 2008), Alzheimer disease (Gunstad et al 2008), and Huntington's disease (Ciammola et al 2007) also supports this conclusion. In fact, higher BDNF levels are associated with better neuropsychological test performance in individuals with these diseases (Yu et al 2008). More recent studies also support the notion that circulating BDNF is a biomarker of memory and general cognitive function in healthy adults (Komulainen et al 2008). However, none have examined this association in patients with schizophrenia. Therefore, the present study was designed to determine the interrelationships of BDNF serum levels with cognitive function and *BDNF* genotype in schizophrenia and healthy controls. The primary aims of this study were: (1) to examine the relationship between serum BDNF levels and cognitive function in schizophrenia and healthy controls; (2) to determine whether the Val66Met polymorphism influences cognitive function in these groups; and (3) to examine whether the relationship between serum BDNF levels and/or cognitive function is dependent on the *BDNF* Val66Met polymorphism.

## MATERIALS AND METHODS

### Subjects

Six hundred and fifty-seven inpatients were recruited from Beijing Hui-Long-Guan Psychiatric hospital, and HeBei Province Veterans Psychiatric Hospital in BaoDing city, 50 miles from Beijing. All patients met the DSM-IV diagnosis of schizophrenia, which was confirmed by two psychiatrists based on the Structured Clinical Interview for DSM-IV (SCID). Their clinical subtypes were: paranoid, 216 (32.9%); undifferentiated, 351 (53.4%); disorganized 53 (8.1%); residual, 34(5.2%) and others, 3 (0.4%). Patients were between 25–75 years old and had a mean duration of hospitalization of  $9.4 \pm 7.5$  years. All patients had been receiving stable doses of oral antipsychotic drugs for at least 12 months.

Four hundred and forty-five control subjects (Male/female=263/182) were also recruited from the local community in Beijing. Their mean age was  $44.9 \pm 13.6$  years (range 25–75), with a mean education of  $9.7 \pm 5.4$  years. The mean body mass index (BMI) of the controls was  $25.1 \pm 4.2$  Kg/m<sup>2</sup> while that of the patients was  $24.7 \pm 4.0$  Kg/m<sup>2</sup>. A research psychiatrist assessed current mental status and personal or family history of any mental disorder in controls by unstructured clinical interviews to exclude potential controls with Axis I disorders. None of them presented a personal or family history of psychiatric disorder.

We obtained a complete medical history, physical examination and laboratory tests from patients and control subjects. Any subjects with test abnormalities or with medical illnesses were excluded. Neither schizophrenic patients nor control subjects suffered from drug or alcohol abuse/dependence. No schizophrenic patients took medications for treating physical diseases. All subjects were Han Chinese, and they gave written informed consent, which was approved by the Institutional Review Board of Beijing Hui-Long-Guan hospital.

## Clinical Measures

We assessed cognitive functioning using the Repeatable Battery for the Assessment of Neuropsychological Status (RBANS, Form A) (Randolph et al 1998). The RBANS was previously translated into Chinese by our group and its clinical validity and test-retest reliability established among controls and schizophrenic patients (Zhang et al 2009).

Four psychiatrists who had simultaneously attended a training session in the use of the Positive and Negative Syndrome Scale (PANSS) rated patients on this scale. After training, repeated assessment showed that the inter observer correlation coefficient was maintained at greater than 0.8 for the PANSS total score. We obtained blood samples to assess BDNF levels at the time of PANSS ratings.

## Serum BDNF measurement

We measured fasting serum BDNF levels by sandwich ELISA using a commercially available kit as described in our previous report (Xiu et al 2009). All samples were assayed by a research assistant blind to the clinical situation. Inter- and intra-assay variation coefficients were 7% and 5%, respectively.

## Genotyping

The genotypes of the *BDNF* Val66Met polymorphisms were identified as reported in our previous study (Zhang et al 2008). A research assistant who was blinded to the clinical status genotyped every subject twice for accuracy of genotyping.

## Statistical analysis

Deviations from Hardy–Weinberg equilibrium (HWE) were assessed using the HWSIM program (Cubells et al 1997). The *BDNF* Val66Met allele and genotype frequencies were compared between schizophrenic patients and healthy controls using  $\chi^2$  tests. Group differences were compared using Student's *t*-test or one-way analysis of variance (ANOVA) for continuous variables and chi-squared for categorical variables. Post-hoc comparisons between subgroups were made using the Fisher's least significant difference (LSD) procedure rather than the more conservative Bonferroni corrections for multiple comparisons. Correlation between variables was studied using Pearson product moment correlations.

For the main models, analyses of covariance (ANCOVA) was constructed with BDNF genotype as the independent variable, and the cognitive scores shown by the RBANS total and 5 index scores as dependent variables, with sex, age, education, illness course, age of onset, body mass index (BMI), smoking, medication type (atypical vs. typical antipsychotics) and dose (chlorpromazine equivalents) and duration of treatment as the covariates. In each model, we tested the main effect of diagnostic group, the main effect of genotype and diagnostic group  $\times$  genotype interaction. The diagnostic group  $\times$  genotype interaction term in the model detects the differential effects that alleles might have on cognitive scores between diagnostic groups. We first contrasted the homozygote groups (Val/Val against Met/Met). When significant differences between homozygote groups were found, we formally assessed whether heterozygotes (Val/Met) had an intermediate level by using a linear regression test. Similarly, we analyzed the main effect of the *BDNF* genotype on serum BDNF levels using ANCOVA.

Lastly, exploratory regression analyses were used to examine whether any relationships between BDNF serum levels and cognitive function differed across *BDNF* genotype groups. Stepwise multiple regression analysis used RBANS total or Index scores as dependent variables with BDNF levels as the independent variable in each BDNF genotype group.

Covariates in these stepwise forward entry models included age, gender, education, BMI, smoking, duration of illness, age of onset, PANSS scores and antipsychotic medication dosage, type (typical vs atypical antipsychotics) and duration.

## RESULTS

Demographic characteristics are summarized in Table 1. A significantly greater proportion of patients were male and smokers (both  $p < 0.01$ ) and older ( $p < 0.001$ ) than controls.

Distributions of the BDNF genotypes were consistent with HWE in both patients and controls (both  $p > 0.05$ ) (Table 1). We found no significant differences in BDNF genotype and allele distributions between patients and controls ( $\chi^2 = 2.57$ ,  $df = 2$ ,  $p > 0.05$ ; and  $\chi^2 = 2.37$ ,  $df = 1$ ,  $p > 0.05$ , respectively).

### Genotype Effects on Cognitive Functioning Between Patients and Normal Controls

RBANS data were available from 575 patients and 405 healthy controls. RBANS total and index scores, and the effects of the *BDNF* Val66Met polymorphism on the RBANS total and index scores are summarized in Table 2.

Cognitive test scores were significantly lower in schizophrenic than control subjects on the total scores and all indexes (all  $p < 0.000$ ) except for Visuospatial/constructional index ( $p > 0.05$ ). There were no significant genotype effects or genotype  $\times$  diagnosis effects for Immediate Memory, Language, Delayed memory and total scores.

As shown in Table 2, the three genotypes significantly differed on the Visuospatial/constructional index ( $F = 4.25$ ,  $p < 0.01$ ), and using Fisher's least significant difference (LSD) to control for the two potential comparisons only the Val/Val vs. Met/Met difference was significant ( $p = 0.03$ ). This difference in performance between Val/Val and Met/Met genotypes held within the schizophrenia ( $p < 0.05$ ) and the controls ( $p < 0.01$ ) based on Fisher's LSD. Including heterozygotes produced a significant linear correlation between the number of Met-66 alleles (e.g. zero, one or two alleles) and the Visuospatial/Constructional Index score within the controls ( $p < 0.01$ ), but not the schizophrenia ( $p = 0.07$ ).

Further analysis showed a significant genotype  $\times$  diagnosis effect on Attention (Table 2) ( $F = 3.15$ ,  $p < 0.05$ ). Based on Fisher's LSD comparison this weak interaction reflected significantly higher attention index scores in patients who were Val homozygous than those who were Met homozygous ( $p < 0.05$ ), while the controls showed no differences.

### Genotype Effects on Serum BDNF Levels Between Patients and Normal Controls

Serum BDNF levels were normally distributed in both patients and controls (Kolmogorov-Smirnov one-sample test: both  $p > 0.05$ ). Table 3 revealed a significant main effect of diagnostic group on BDNF levels ( $F = 15.2$ ,  $df = 664$ ,  $p < 0.001$ ). Male patients had significantly lower BDNF levels than female patients ( $p < 0.01$ ; adjusted  $p < 0.01$ ). However, BDNF levels were not significantly different between men and women in controls ( $p > 0.05$ ).

There was no main effect of genotype on serum BDNF levels ( $p > 0.05$ ) nor genotype  $\times$  diagnosis effect ( $p > 0.05$ ). We observed no significant differences in BDNF levels between genotypic subgroups (Val/Val, Val/Met, and Met/Met) in patients or controls (both  $p > 0.05$ ) (Table 3). When sex and BMI were added as covariates, there were still no significant differences in BDNF levels in genotypic subgroups in patient and control groups (both  $p > 0.05$ ), suggesting that BDNF genotype did not influence serum BDNF levels in the patient or control groups.

## Differential Effects of BDNF Genotype on the Relationships between BDNF Serum Levels and Cognitive Functioning

The patients showed a significant overall main effect of BDNF levels on the RBANS total score and its index scores ( $F=2.59$ ,  $df=6, 411$ ,  $p=0.016$ ). The BDNF level  $\times$  genotype interaction was also statistically significant ( $F=2.48$ ,  $df=6, 412$ ,  $p=0.023$ ), indicating differences between genotypes in the relationships between BDNF levels and cognitive scores. A further partial correlation analysis showed a significant positive correlation between serum BDNF levels and the RBANS total score ( $r=0.21$ ,  $p<0.03$ ), and between BDNF levels and the Immediate Memory index ( $t=3.22$ ,  $p<0.01$ ). However, no significant effects of BDNF levels, *BDNF* genotype, gender and their interactions were found in the control group (all  $p>0.05$ ).

Regression analyses among Val homozygote patients found a significant negative association of BDNF levels with the Delayed Memory index ( $\beta=-0.36$ ,  $t=-2.21$ ,  $p=0.04$ ) and with the PANSS total score ( $\beta=-0.60$ ,  $t=-3.52$ ,  $p=0.002$ ). Among Met homozygote patients increases in BDNF levels were significantly associated with better performance on the Visuospatial/constructional index ( $\beta=0.41$ ,  $t=2.89$ ,  $p=0.006$ ). Among Met/Val heterozygous patients BDNF levels were positively correlated with the Attention ( $\beta=0.59$ ,  $t=3.48$ ,  $p=0.001$ ) and Immediate Memory indices ( $\beta=0.46$ ,  $t=2.56$ ,  $p=0.013$ ), but negatively correlated with smoking behavior ( $\beta=-0.33$ ,  $t=-2.99$ ,  $p=0.004$ ).

## DISCUSSION

Our results demonstrate that the BDNF Val66Met polymorphism may not contribute directly to the susceptibility to schizophrenia, although two previous studies reported such an association (Neves-Pereira et al., 2005; Rosa et al., 2006). Like our study, several studies did not replicate these earlier associations (Naoe et al., 2007; Varnas et al., 2008), including studies in Han Chinese populations (Xu et al 2007; Zhou et al 2010; Yi et al 2011). One possible explanation for the inconsistency is a difference in ethnic background, since we found the allele frequency distribution of Val66Met varied significantly between Chinese and Caucasian subjects. The Met allele frequency was around 50% in Chinese (Xu et al 2007; Zhou et al 2010; Yi et al 2011), but around 20% in Caucasian subjects (Rosa et al., 2006; Naoe et al., 2007; Varnas et al., 2008). In our current study, the Met allele frequency was 49.7% in control subjects, which is similar to other studies in Chinese (Xu et al 2007; Zhou et al 2010; Yi et al 2011). Thus, interethnic differences in the genotype frequencies of the BDNF Val66Met polymorphism may play an important role in accounting for the inconsistent results across the different populations. Several other factors may also account for these divergent results, for example, small gene effects, heterogeneity of the schizophrenia diagnosis, and population stratification.

### BDNF in human cognition and cognitive dysfunction among schizophrenia

We found significant differences in cognitive scores in nearly all of the post-hoc pairwise comparisons between schizophrenics and controls except for the Visuospatial/Constructional index. These data replicate numerous studies indicating that patients with schizophrenia perform worse than controls on a range of cognitive tasks (Sharma and Antonova 2003; Kraus and Keefe 2007). In fact, we found that the absolute difference between the mean RBANS total scores of the two groups was more than 16 points (78.4 vs. 61.9), which is similar to a result by Dickerson et al (Dickerson et al 2004).

We found that Met allele carriers, regardless of whether they were healthy controls or patients with schizophrenia, consistently performed worse than their Val homozygous counterparts on the Visuospatial/constructional index. Moreover, we found a significant



linear correlation between the number of Met66 alleles and the Visuospatial/Constructional Index score in both patients and controls. The underlying mechanisms that are responsible for the Met66 allele's influence on this cognitive measure are unknown. However, studies have shown that the *BDNF* Val66Met polymorphism is related to hippocampal-mediated memory performance in humans (Egan et al 2003; Hariri et al 2003). Hippocampal neurons that are transfected with the Met variant of BDNF show less depolarization-induced BDNF secretion (Chen et al 2004). The Met variant results in inefficient trafficking of BDNF to secretory granules, reduced activity-dependent BDNF release, and poorer hippocampal-mediated memory (Egan et al 2003). Overall, Met allele carriers, regardless of whether they are controls or patients with schizophrenia, or unaffected siblings, demonstrate poorer episodic memory than their respective Val-homozygous counterparts (Egan et al 2003; Dempster et al 2005). The patients who are Met allele carriers have lower hippocampal function while performing a declarative memory task as assessed with functional MRI blood oxygenation level responses (Hariri et al 2003). These Met allele patients also have reduced hippocampal and prefrontal gray matter (Szeszko et al 2005; Bueller et al 2006). Ho et al found the *BDNF* 66Met variant correlates with reduced gray matter volumes within temporal and occipital lobes, brain regions known to participate in verbal memory and visuospatial abilities in both normal controls and patients with schizophrenia (Ho et al 2007). The convergence of these cognitive and brain morphology findings suggest that the *BDNF* Val66Met polymorphism influences specific aspects of human cognition. Our current study also indicated the association between the *BDNF* Met-66 variant and poor Visuospatial/Constructional performance, but this genotype status did not appear to impact total cognitive function, or affect language, immediate and delayed memory performances.

Visuospatial performance is subserved by large scale, distributed neuronal networks. Object perception involves the ventral occipito-temporal whereas processing spatial information involves the dorsal occipito-parietal pathway. Therefore, brain regions that mediate visuospatial abilities include the secondary visual cortices, inferior temporal regions, and the parietal heteromodal association cortices (Ho et al 2007). Patients with schizophrenia and healthy controls who carry the Met allele have smaller temporal and occipital lobar GM volumes than their respective Val-homozygous counterparts, and those patients with the *BDNF* Met-66 allele are significantly more impaired in visuospatial abilities than patients who are Val homozygous (Ho et al 2007).

The *BDNF* 66Met variant also may have a specific role in attentional dysfunction among schizophrenics. Unlike controls, patients with schizophrenia who carry the Met allele consistently performed worse than their Val-homozygous counterparts on attentional performance. In our current studies this common SNP showed different effects on cognitive function in patients with schizophrenia than in healthy controls. However, the mechanisms by which the *BDNF* Val66Met variant affects attention are not well understood. Hence, the differential effects of the *BDNF* Val66Met genotype on attention in patients with schizophrenia and healthy controls deserves further investigation.

Our cognitive index test findings about the differences between schizophrenia and controls across the three *BDNF* Val66Met genotypes remained significant in key comparisons between the two homozygous genotypes using Fisher's LSD for group comparisons. However, the heterozygous genotype did not differ from either homozygous group among the schizophrenia suggesting that the effects of the *BDNF* Val66Met genotype on some aspects of cognitive function may be moderate. Moreover, the genotype by diagnosis interaction on attention was difficult to evaluate especially with out modest sample size. Therefore, our results should be interpreted with caution.

Previous studies reported associations of the Met allele with deficits in memory functions (Egan et al 2003; Hariri et al 2003; Dempster et al 2005; Ho et al 2007), but we found associations with visuospatial and attention more than memory. These other aspects of neuropsychological functioning are clearly not independent of memory and cultural aspects of learning. One possible explanation for discrepant association findings between Met allele with cognitive functioning is differences in population genetic structure between patient samples. For example, a recent population-based genetic study showed that the derived Met allele of the Val66Met polymorphism increased in frequency from 0.55% to 19.9% and 43.6% frequency in Sub-Saharan Africa, Europe, and Asia, respectively (Petryshen et al. 2010). Furthermore, this study examined haplotypes comprised of the 12 BDNF SNPs and observed considerable haplotype diversity among global populations (Petryshen et al. 2010), suggesting substantial population diversity at the *BDNF* locus. Thus, interethnic differences in the genotype frequencies of the BDNF gene Val66Met polymorphism may play an important role that may account for the inconsistent results between Met allele with cognitive functioning seen in different samples from the different populations. On the other hand, quantitative and molecular genetic studies have established that there are additive genetic contributions to different aspects of cognitive ability--especially general intelligence, which involves several genes of small effects that interact in order to express the different aspects of cognition (Deary et al 2010). However, whether some cognition-related candidate genes are related to different aspects of cognition in different population is still unclear. Hence, whether this discrepancy may be related to the interethnic differences in the allelic frequencies of the BDNF gene polymorphisms between different populations or complex genetic mechanisms related to different aspects of cognition deserve further investigations.

#### **Decreased serum BDNF levels and cognitive impairment: relationship to BDNF Val66Met genotype**

We showed decreased serum levels of BDNF in patients with a chronic form of schizophrenia, which replicates our previous studies (Xiu et al 2009; Zhang et al 2008) and is consistent with others (Rizos et al 2008) but not all (Vinogradov et al 2009). Numerous factors may have contributed to these differences, such as the clinical status of patients, the subtypes of schizophrenic patients recruited, and the type, dosage and length of administration of antipsychotic drugs (Rizos et al 2008).

Only one recent study has shown that serum BDNF levels were increased in chronic schizophrenic patients after cognitive training, and BDNF levels were correlated with cognitive performance (Vinogradov et al 2009). In our present study, BDNF was associated with cognition in schizophrenics but not in healthy controls, suggesting that the effects of serum BDNF on cognitive function differs between patients with schizophrenia and healthy control. Furthermore, the *BDNF* Val66Met polymorphism affects the correlation between BDNF levels and different aspects of cognitive function in schizophrenic patients.

We found higher serum BDNF levels associated with better cognitive function in schizophrenia, which is consistent with two recent studies in healthy older adults showing higher serum BDNF levels associated with better neuropsychological function (Gunstad et al 2008; Komulainen et al 2008). The exact mechanisms responsible for these findings are unknown. One possible explanation is the neuroprotective effects of BDNF. BDNF plays a key role in modulating synaptic transmission and plasticity (Shimizu et al 2003) and has been hypothesized to be an important factor in the induction of long-term potentiation (LTP), a persistent strengthening of synapses associated with learning and memory (Lu et al 2008). Preclinical studies show that LTP is impaired in mice lacking the BDNF gene (Tyler et al 2002), and LTP can be restored with adenovirus-mediated transfection of these mutant mouse CA1 cells with the BDNF gene. Taken together, these results support a mechanism linking cognitive function and BDNF levels.



Yet, it is noteworthy that the source of circulating BDNF remains unknown. Although BDNF is highly concentrated in the nervous system, it is also present in peripheral blood with high concentrations in serum (Radka et al 1996). A recent study found a significant correlation between BDNF levels in plasma and cerebrospinal fluid (CSF) of unmedicated schizophrenic patients (Pillai et al 2009). Accordingly, if low peripheral BDNF levels reflect low levels in the brain, this could affect cognitive function in patients with schizophrenia. Indeed, we showed that increased BDNF serum levels were correlated with better performance on Visuospatial/constructional ability for the Met/Met genotype, as well with better performance on Attention and Immediate memory abilities for the Met/Val heterozygous. In contrast, increased BDNF serum levels were associated with poor Delayed Memory performance for the Val/Val genotype. The clinical significance of these divergent serum BDNF-cognitive function relationships across *BDNF* genotypes in schizophrenia remains unclear. Further studies are needed to determine the association of peripheral BDNF levels and cognitive function in patients with schizophrenia and possibly other psychiatric disorders or subtypes of schizophrenia (Lencz et al 2009).

Like many others we found that the serum BDNF levels were lower among the schizophrenia patients, but were not related to the val66met genotype. For example, the Val66Met polymorphism was unrelated to the concentration of BDNF in whole blood in a healthy European sample (n=78) (Trajkovska et al., 2007) and was also unrelated to plasma or serum concentrations of BDNF in several healthy Asian samples (Yu et al 2008; Jianget al., 2009; Zou et al 2010). However, a recent healthy European sample (n=114) showed an association between the Val66Met genotype and serum BDNF concentrations, such that individuals with the Met allele had higher BDNF levels (Lang et al., 2009). Interestingly, other variants rather than the Val66Met variant in the BDNF gene might regulate the level of serum BDNF, as identified in a recent Italian community-based sample (N = 2054) (Terracciano et al 2011). The BDNF Val66Met polymorphism may indeed have an association with serum BDNF levels only among some ethnic groups with Asians not having this association. Moreover, the Met variant has been hypothesized to decrease the activity dependent but not the constitutive BDNF secretion, which our studies assess (Tramontina et al 2007). Finally, we are assessing serum, not brain BDNF levels, although Klein et al. (2011) recently reported that blood and plasma BDNF levels reflect brain-tissue BDNF levels.

In conclusion, we have provided new evidence to support the association between the *BDNF* 66Met variant and poor visuospatial/constructional performance in both patients with schizophrenia and healthy controls, suggesting a specific role of the *BDNF* 66Met variant in some aspects of poor cognitive function. The present study indicates that peripheral BDNF levels may be a biomarker of general cognitive function in schizophrenia. The association between decreased serum BDNF levels and the degree of cognitive impairment in schizophrenia patients appears to be dependent on the presence of *BDNF* Val66Met polymorphism.

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**Table 1**

Demographic Characteristics and *BDNF* Allele and Genotype Distributions in Normal Controls and Patients with Schizophrenia

Variable	Normal Controls (n=445)	Schizophrenia (n=657)	Statistic (P Value)
Sex (Male/Female)	263/182	578/79	122.4 (<0.001)
Age (years)	44.9±13.6	48.4±13.7	25.4 (<0.001)
Education (years)	9.7±5.4	9.0±5.6	2.1 (0.15)
Body Mass Index (kg/m <sup>2</sup> )	25.1±4.2	24.7±4.0	2.3 (0.13)
Allele frequency (%)			
Val	50.3%	53.7%	2.37(0.12)
Met	49.7%	46.3%	2.57(0.28)
Genotype frequency (%)			
Val/Val	104 (23.6%)	175 (27.0%)	
Met/Val	236 (53.5%)	347 (53.4%)	
Met/Met	101 (22.9%)	127 (19.6%)	



**Table 2**  
Comparisons of Total and Index Scores on the RBANS by Diagnostic and Genotype Groupings

Cognitive Index	Schizophrenia				Normal Controls				Genotype, F (P value)	Genotype×Diagnosis, F (P value)
	Val/Val (n=155)	Val/Met (n= 316)	Met/Met (n=104)	Val/Val (n=103)	Val/Met (n=204)	Met/Met (n=98)	Genotype, F (P value)	Genotype×Diagnosis, F (P value)		
Immediate Memory	57.2±16.3	55.6±14.3	56.4±12.9	72.1±16.7	74.0±18.5	70.8±15.1	0.32 (0.72)	1.24 (0.29)		
Attention	68.6±17.6	66.9±17.2	63.7±15.5	83.6±19.0	86.2±20.7	82.9±19.1	0.09 (0.92)	<b>3.15 (&lt;0.05)</b> <sup>b</sup>		
Language	80.2±16.8	78.5±15.5	80.7±15.0	92.2±13.1	94.3±13.1	92.5±13.7	0.03 (0.97)	1.52 (0.22)		
Visuospatial/Constructional	77.8±17.3	76.0±20.1	74.2±17.9	79.6±14.6	77.8±15.1	74.4±14.6	<b>4.25 (&lt;0.01)</b> <sup>a</sup>	1.29(0.27)		
Delayed Memory	65.6±20.1	64.5±18.7	63.8±15.1	84.8±13.7	85.2±15.9	83.6±15.6	0.32 (0.73)	0.13 (0.88)		
Total	63.4± 15.4	60.9±13.3	62.3±12.8	77.5±14.1	78.9±15.5	75.4±13.6	0.51 (0.60)	2.03 (0.13)		

**Note:**

<sup>a</sup>There were significant genotype effects on Visuospatial/constructional index (F=4.25, p<0.01). Further, there was a linear negative correlation between the number of Met 66 alleles and the Visuospatial/Constructional Index score in controls (p<0.01) and in patients (p=0.07).

<sup>b</sup>There was a significant genotype × diagnosis effect on Attention (F=3.15, p<0.05). Patients who were Val homozygous had significantly higher Attention index scores than patients who were Met homozygous (p<0.05). There were no genotype differences in Attention among healthy controls (F=0.82, p>0.05).

**Table 3**

Serum BDNF Levels (ng/ml) in Normal Controls and Schizophrenia by Genotype Groupings

	Val/Val	Met/Val	Met/Met
Patients (n=424)	9.6±3.1	9.5±2.9	9.8±2.7
Controls (n=422)	11.8±2.6	11.8±2.7	11.9±2.1
P value <sup>a</sup>	<0.01	<0.01	<0.01

**Note:**

<sup>a</sup> indicates the comparison between patients and controls by genotype groupings. There was no main effect of genotype on serum BDNF levels ( $p>0.05$ ) nor genotype  $\times$  diagnosis effect ( $p>0.05$ ).