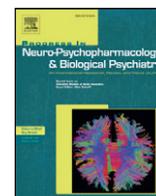




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Brain derived neurotrophic factor gene (*BDNF*) and personality traits: The modifying effect of season of birth and sex



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ABSTRACT

Personality traits are complex phenotypes influenced by interactions of multiple genetic variants of small effect and environmental factors. It has been suggested that the brain derived neurotrophic factor gene (*BDNF*) is involved in personality traits. Season of birth (SOB) has also been shown to affect personality traits due to its influences on brain development during prenatal and early postnatal periods. The present study aimed to investigate the effects of *BDNF* on personality traits; and the modifying effects of SOB and sex on associations between *BDNF* and personality traits. A sample of 1018 young adults (68% women; age range 17–25 years) of Caucasian origin from the Russian Federation was assessed on personality traits (Novelty Seeking, Harm Avoidance, Reward Dependence, Persistence, Self-directedness, Cooperativeness, Self-transcendence) with the Temperament and Character Inventory-125 (TCI-125). Associations between personality traits and 12 *BDNF* SNPs were tested using linear regression models. The present study demonstrated the effect of rs11030102 on Persistence in females only ($P_{FDR} = 0.043$; $r^2 = 1.3\%$). There were significant interaction effects between Val66Met (rs6265) and SOB ($P_{FDR} = 0.048$, $r^2 = 1.4\%$), and between rs2030323 and SOB ($P_{FDR} = 0.042$, $r^2 = 1.3\%$), on Harm Avoidance. Our findings provide evidence for the modifying effect of SOB on the association between *BDNF* and Harm Avoidance, and for the modifying effect of sex on the association between *BDNF* and Persistence.

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1. Introduction

Personality traits are predictors of important life outcomes including well-being, academic achievement, health risk behaviors, and longevity; they are also considered as endophenotypes for major psychiatric disorders (De Beaumont et al., 2013; Duclot and Kabbaj, 2013; Terracciano et al., 2010a).

Personality traits are complex phenotypes affected by interactions of multiple genes of small effect with environmental factors. The estimated

heritability of personality traits variability is 30–40% (Bouchard and Loehlin, 2001; Garcia et al., 2013). However, candidate gene studies, as well as genome-wide association studies (GWAS), often failed to confirm initial findings of specific genetic risk factors for personality traits (de Moor et al., 2012; Shifman et al., 2008; Terracciano et al., 2010a, 2011a). Difficulties in identifying specific genetic risk factors are likely to be related to influences of sex, age, ethnicity, as well as of various environmental factors that can modify the effects of genes. To date, the role of candidate gene approach focusing on genetic factors with known functional role in manifestation of personality traits in the context of gene–environment interactions remains significant.

Brain derived neurotrophic factor gene (*BDNF*) is one of the strong candidate genes for personality traits (Montag, 2014). *BDNF* is involved in the growth and maintenance of several neuronal systems, serves as a neurotransmitter modulator, and participates in use-dependent plasticity mechanisms, such as learning and memory (Nakazato et al., 2003; Rasmusson et al., 2002). Therefore, it has been suggested that *BDNF* can play an important role in anxiety-related personality traits and disorders. In humans, decreased serum *BDNF* levels were associated with depression (Bocchio-Chiavetto et al., 2010; Trajkovska et al., 2008), high Neuroticism (Lang et al., 2004; Terracciano et al., 2011b) and

Abbreviations: *BDNF*, Brain derived neurotrophic factor; SOB, Season of birth; GWAS, Genome-wide association study; SNP, Single nucleotide polymorphism; UTR, Untranslated region; ANOVA, One-way analysis of variance; GxE, Gene–environment interaction; FDR, False discovery rate; PCR, Polymerase chain reaction; HA, Harm Avoidance; NS, Novelty Seeking; RD, Reward Dependence; PS, Persistence; SD, Self-directedness; ST, Self-transcendence; TCI, Temperament and Character Inventory.

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Harm Avoidance (Minelli et al., 2011), while increased BDNF concentrations have been reported after treatment with antidepressants (Shimizu et al., 2003). On the contrary, lower plasma BDNF levels were observed in men who scored lower on depression and vulnerability to stress, higher on Conscientiousness and Extraversion (Terracciano et al., 2010b), and lower on Harm Avoidance (Yasui-Furukori et al., 2013).

Human molecular genetic studies of the *BDNF* gene can provide further evidence for the role of this protein in personality traits. Human *BDNF* gene (11p13) consists of eleven exons and tissue- and brain-region specific nine functional promoters. The replacement of Val-allele by Met-allele in *BDNF* gene (Val66Met, or rs6265) disrupts cellular processing, trafficking, and activity-dependent secretion of BDNF (Hong et al., 2011). The *BDNF* Met-allele has been associated with gray matter volume deficits especially in the hippocampus, prefrontal cortex (Hajek et al., 2012; Pezawas et al., 2004), and in the right amygdala (Montag et al., 2009). Moreover, Met-allele has been associated with reduced hippocampal activation (Kambeitz et al., 2012), deficient intracellular transport of BDNF to dendrites and reduced magnitude of long term potentiation (Kleim et al., 2006).

Animal studies demonstrated that Met/Met mice showed increased anxiety-related behaviors in stressful conditions (Chen et al., 2006). In humans, a number of studies have reported association between Met-allele and depression that was modified by the presence of stressful life events (Brown et al., 2013; Hosang et al., 2014), or enhanced reactions to external stressful stimuli (Colzato et al., 2011). However, a recent meta-analysis failed to support association between Val66Met and depression (Gyekis et al., 2013). One possible explanation for this inconsistency is that *BDNF* gene might be involved in variation of anxiety-related traits rather than in depression itself. As it has been demonstrated, *BDNF* Met-allele carriers have higher Harm avoidance (Jiang et al., 2005; Montag et al., 2010), Reward Dependence and Extraversion (Itoh et al., 2004) as compared with Val/Val homozygotes. However, associations between Met-allele and lower Harm Avoidance (Ando et al., 2012) and Neuroticism (Sen et al., 2003) have also been reported. A recent GWAS of personality traits has confirmed an association of Met-allele and lower Extraversion, however, together with the meta-analyses has provided no evidence for the effect of Val66Met on anxiety-related traits (Frustaci et al., 2008; Terracciano et al., 2010a, 2010c). Such an inconsistency across the studies could be explained by epistatic effect between *BDNF* Val66Met and other polymorphisms, for example 5-*HTTLPR* (linked polymorphic region in serotonin transporter gene) as demonstrated by Terracciano et al. (2010c). This study showed that 5-*HTTLPR* L/L homozygotes scored lower on Neuroticism in the presence of *BDNF* Val-allele, but scored higher on Neuroticism in the presence of *BDNF* Met-allele (Terracciano et al., 2010c).

The majority of previous studies of the *BDNF* gene in personality traits have focused on the role of a single *BDNF* polymorphism – Val66Met. However, other genetic variants could be involved in regulation of the *BDNF* gene expression. It has been reported that *BDNF* expression is regulated by a group of miRNAs and that common genetic variants (i.e., rs11030100 and rs11030099 in 3'-UTR) influence miRNA targeting and participate in expression modulation (Caputo et al., 2011). A number of other *BDNF* SNPs, such as rs11030102, rs11030107, rs10835211, have also been shown to be associated with serum BDNF level (Terracciano et al., 2013).

A sex-specific effect of the *BDNF* gene on cortisol level has been reported (Shalev et al., 2009). Moreover, animal studies demonstrated that female *BDNF* conditional knockouts displayed an increase in depression-like behaviors, while male knockouts reported normal depression-related behaviors (Monteggia et al., 2007).

Environmental factors may also modify the effect of the *BDNF* gene on personality traits. Season of birth (SOB) can influence anxiety-related personality traits and psychiatric disorders (Antonsen et al., 2012; Chotai et al., 2009). For example, the effect of SOB was demonstrated on Novelty Seeking (Chotai et al., 2009), hyperthymic personality

(characterized with high Novelty Seeking and low Harm Avoidance), and depressive temperament (Rihmer et al., 2011). The findings suggest that people born in spring/summer are more likely to have lower anxiety-related traits (i.e., Harm Avoidance) and higher approach-related traits (i.e., Novelty Seeking) than those born in winter.

The present study aims to explore whether the *BDNF* gene is involved in anxiety-related traits, (i.e., Harm avoidance). In addition, the study aims to investigate whether Val66Met and other *BDNF* SNPs are associated with Novelty Seeking that is correlated with Extraversion. Moreover, since both sex and SOB can affect personality traits (Chotai et al., 2009), the present study aims to test whether associations between the *BDNF* gene and personality traits are modified by sex and SOB.

2. Materials and methods

2.1. Sample

In total, 1018 young adults (68% women; mean age \pm SD: 19.81 \pm 2.65 years, age range: 17–25 years), enrolled at the Universities in the Russian Federation. Socio-demographic data including sex, ethnicity, and date of birth were obtained from all the participants. All participants were of Caucasian origin: Russians (N = 409), Tatars (N = 290), Bashkirs (N = 130) and Udmurts (N = 189). Exclusion criteria were self-reported individual and/or family (of a first and/or second degree relative) history of any psychiatric disorders. The study was approved by the Biological Ethics Committee of Institute of Biochemistry and Genetics (Ufa, Russia), and written informed consent was obtained from all the participants after the procedure had been explained to them. All the participants were informed about the voluntary and confidential nature of their participation.

2.2. Measures

2.2.1. Personality traits

Personality traits were assessed using the Russian version of the Temperament and Character Inventory (TCI-125). The TCI-125 evaluates four temperament traits: Novelty Seeking, Harm Avoidance, Reward Dependence, Persistence, and three character traits: Self-directedness, Cooperation and Self-transcendence (Cloninger et al., 1993). Cronbach's alpha reliability, which measures internal consistency of test items, was high for all seven personality scales (Novelty Seeking: $\alpha = 0.76$; Harm Avoidance: $\alpha = 0.81$; Reward Dependence: $\alpha = 0.67$; Persistence: $\alpha = 0.69$; Self-directedness: $\alpha = 0.82$; Cooperation: $\alpha = 0.76$; Self-transcendence: $\alpha = 0.84$) as well as for the TCI-125 in total ($\alpha = 0.87$).

2.2.2. Season of birth (SOB)

Since all the participants were born in the northern hemisphere, SOB was classified according to traditional Russian definition of the four seasons: March, April and May represented spring (26.9% of all the participants); June, July and August represented summer (24.0%); September, October and November represented autumn (24.2%); and December, January and February represented winter (24.9%). We also used astronomical criterion of SOB taking in account the equinoxes (i.e., March 22–June 21 represented spring; June 22–September 21 – summer; September 22–December 21 – autumn; December 22–March 21 – winter). These two definitions of the four seasons were used since some of the previous studies of the effects of SOB on personality traits have used the traditional criterion (Hori et al., 2012; Martínez-Ortega et al., 2011), while others used the astronomical criterion (Hori et al., 2012; Rihmer et al., 2011; Shuman et al., 2010).

2.3. SNP selection and genotyping

Genomic DNA was isolated from the whole blood using a standard phenol–chloroform technique. In total, 12 *BDNF* SNPs (MAF > 10%)

were selected using the Tagger algorithm implemented in the Haploview 4.1 (Barrett et al., 2005).

Genotyping of the 12 SNPs was performed using a PCR-RFLP method. PCR primers for each polymorphism were designed in Primer 3 (<http://bioinfo.ut.ee/primer3-0.4.0/>). PCR was performed in total volume of 15 μ l with 20–50 ng of genomic DNA, Taq polymerase (Silex, Russia). Subsequently, for allele detection PCR products were accomplished by overnight incubation with 3U of corresponding restriction endonuclease (Fermentas, Canada) according to manufactures recommendations, resolved in 7% polyacrylamide gel (PAAG) and visualized by staining with ethidium bromide.

2.4. Statistical analysis

Genotype and allele frequencies of the investigated SNPs, as well as Hardy-Weinberg equilibrium, calculations were performed in a total sample using PLINK v.1.07 (Purcell et al., 2007). Haplotype blocks were delineated using the confidence interval method of Gabriel et al. (2002), and measures of linkage disequilibrium (LD) between markers were obtained using Haploview 4.1. The extent of disequilibrium was demonstrated by the standardized D' characteristic multiplied by 100 in the LD illustration generated in Haploview v.4.1. Haplotypes with a frequency less than 1% were excluded from the further analysis.

Since personality traits scores were distributed normally, the main effects of the individual *BDNF* SNPs and haplotypes, sex and SOB, as well as the effects of gene-by-sex and gene-by-SOB interactions on personality traits were investigated using linear regression models in a total sample with PLINK v.1.07. First, each of the *BDNF* SNPs, SOB, and sex were entered into the model as independent variables, and each of personality traits – as dependent variables. For the categorical variables with the number of categories higher than two, a matrix of dummy variables was constructed that were later used for the linear regression analysis. Additive genetic model was used to estimate the effect of a minor allele of each of the *BDNF* SNPs on personality traits. Information on minor alleles of each *BDNF* SNPs is presented in Table 1. Empirical P -value permutations were run. Effect sizes were calculated for all statistical models. The effect sizes were reported as r^2 , which describes the proportion of variance in personality traits that is accounted

for the differences in genotype or haplotype controlling for sex and season of birth.

Second, the interaction effects of *BDNF* SNPs with SOB and sex were tested in STATA v.11. According to Keller (2014), in order to control for potential confounders in GxE analysis, along with all main effects and gene \times environment interaction effect, it is necessary to enter all covariate \times environment and covariate \times gene interaction terms. So, our linear regression models included the main effects of *BDNF* SNP, SOB, and sex, as well as interaction terms: SNP \times SOB, SNP \times sex, and SOB \times sex. For the models demonstrating interaction effect of a specific SNP or distinct haplotype modulated by SOB or sex on personality traits, we conducted stratification analyses to clarify the direction of effect. For the interaction effects, those with P -value less than 0.10 were considered for stratification analysis.

Power analysis to detect SNP associations and GxE associations with personality traits was conducted with Quanto v.1.2.4 (Gauderman, 2002) with a type I error rate of 5%.

As multiple positive findings were expected, false discovery rate (FDR) procedure (Simes procedure; Benjamini and Hochberg, 1995) was carried out and P -value thresholds were calculated to quantify the joint probability of multiple findings reflecting true associations as opposed to false positives, taking into account all comparisons performed to test our hypotheses. This procedure provides an effective control of type I error rate in the context of multiple correlated tests and has good agreement with the permutation test (Uher et al., 2009). Corrected P -values (P_{FDR}) are shown for all the tests. The number of independent tests was: 1) 12 for ANOVA (analysis of 12 SNPs); 2) 16 for haplotype analysis (analysis of 16 haplotypes); 3) 12 for GxE analysis (analysis of 12 SNPs modified by SOB and sex). The multiple comparison-corrected significance thresholds were then calculated as $(k * 0.05)/m$, where m – the number of statistical tests, k – the order of the tested hypothesis.

3. Results

3.1. Effects of sex and SOB on personality traits

The main effect of sex on personality traits was observed with females scoring significantly higher on Harm Avoidance ($P = 0.016$),

Table 1
The investigated *BDNF* SNPs.

SNP	Chromosomal position, bp ^a	Location in gene	Minor allele	Genotype frequency			P_{HWE}
rs1519479	27624107	Intron (<i>BDNF-AS</i>)	C	C/C 0.241	C/T 0.504	T/T 0.255	0.913
rs2203877	27627486	Intron (<i>BDNF-AS</i>)	T	T/T 0.263	T/C 0.469	C/C 0.268	0.198
rs7124442	27633617	3'-UTR (<i>BDNF</i>) intron(<i>BDNF-AS</i>)	G	G/G 0.097	G/A 0.451	A/A 0.452	0.521
rs6265 (V66M)	27636492	Exon (<i>BDNF</i>) exon(<i>BDNF-AS</i>)	A	A/A 0.021	A/G 0.248	G/G 0.731	1.000
rs11030102	27638172	Intron (<i>BDNF</i>) intron(<i>BDNF-AS</i>)	G	G/G 0.040	G/C 0.354	C/C 0.606	0.447
rs10835211	27657941	Intron (<i>BDNF</i>) intron (<i>BDNF-AS</i>)	A	A/A 0.032	A/G 0.361	G/G 0.607	0.080
rs2030323	27685115	Intron (<i>BDNF</i>)	T	T/T 0.035	T/G 0.303	G/G 0.662	1.000
rs10767665	27690434	Intron (<i>BDNF</i>)	G	G/G 0.246	G/A 0.507	A/A 0.247	0.815
rs1491850	27706301	5' near gene (<i>BDNF</i>)	C	C/C 0.124	C/T 0.496	T/T 0.380	0.185
rs985205	27715568	5' near gene (<i>BDNF</i>)	T	T/T 0.215	T/A 0.531	A/A 0.254	0.162
rs7483883	27723114	5' near gene (<i>BDNF</i>)	C	C/C 0.105	C/T 0.436	T/T 0.459	0.982
rs2172229	27733198	5' near gene (<i>BDNF</i>)	G	G/G 0.181	G/A 0.480	A/A 0.339	0.723

^a According to NCBI36 genome build 36.3. P_{HWE} – P -value for Hardy-Weinberg equilibrium test. Location in both *BDNF* and *BDNF-AS* genes is shown.

Novelty Seeking ($P = 0.037$) and Reward Dependence ($P < 0.001$) as reported previously (Kazantseva et al., 2008). We also tested for the main effect of SOB on personality traits, but no significant differences in personality traits were revealed in individuals with different SOB.

3.2. Main effects of the BDNF gene on personality traits

The distributions of genotype frequencies for the 12 BDNF SNPs were consistent with Hardy-Weinberg equilibrium (Table 1). The analysis of pair-wise linkage disequilibrium revealed the presence of two haplotype blocks in the BDNF gene and neighboring regions spanning 82 and 17 kb ($D' > 0.73$) (Fig. 1). D' coefficients, as well as haplotype structure are shown on Fig. 1. There were nine haplotypes in block 1 and six haplotypes in block 2 with haplotype frequencies higher than 1% (Table 2).

While testing for the main effects of the BDNF SNPs on personality traits, we observed trends in carriers of rs11030102 G-allele to score lower on Persistence ($P = 0.022$; $r^2 = 0.91\%$; power = 0.85) and in carriers of rs1491850 C-allele to score lower on Harm Avoidance ($P = 0.021$; $r^2 = 0.94\%$; power = 0.88); however, these associations became non-significant after FDR-correction (Table S1). Results of linear regression analysis for the associations between the BDNF SNPs and character traits are reported in Supplementary material (Table S2).

Haplotype analysis revealed a trend for associations of BDNF TCTGCC GAC-haplotype (Block 1) with higher Persistence ($P = 0.010$); TTG-haplotype (Block 2) ($P = 0.030$) and ATA-haplotype (Block 2) with Novelty Seeking ($P = 0.007$), which became non-significant after FDR-correction (Table S3). No statistically significant effects of BDNF SNPs or haplotypes on character traits were observed (Supplementary Tables S2 and S4).

3.3. Interaction effects between the BDNF gene and sex on personality traits

We tested for interaction effects of sex and the BDNF SNPs on personality traits. There was an interaction effect between sex and rs11030102 on Persistence ($P = 0.039$) (Table 3). Subsequent stratified analysis revealed that female-carriers of rs11030102 G-allele had lower Persistence as compared with C/C-genotype carriers ($P = 0.003$,

Table 2 Haplotype frequencies in the total sample, males and females.

Haplotype	Total	Males	Females
<i>BDNF block1</i>			
TCAGCGGAT	0.435	0.395	0.460
CTGGGAGGC	0.157	0.161	0.153
CTAACGTGC	0.132	0.135	0.131
CTGGCGGGT	0.110	0.126	0.098
CTGGGAGGT	0.030	0.043	0.022
CTAGCGTGC	0.025	0.023	0.027
TCAGCGGAC	0.024	0.026	0.023
TCAGCGGGT	0.011	0.013	0.011
<i>BDNF block2</i>			
TTA	0.443	0.475	0.426
ATA	0.088	0.075	0.095
ACG	0.273	0.251	0.285
ATG	0.115	0.136	0.103
ACA	0.046	0.037	0.051
TTG	0.031	0.026	0.034

BDNF block1 consists of rs1519479, rs2203877, rs7124442, rs6265 (Val66Met), rs11030102, rs10835211, rs2030323, rs10767665, rs1491850; block 2 – of rs985205, rs7483883, rs2172229, respectively. The most frequent haplotypes are shown in bold. Haplotypes with the frequencies less than 1% are not shown.

$P_{FDR} = 0.043$, $r^2 = 1.3\%$). The power to detect the interaction effect between sex and rs11030102 on Persistence was 0.83.

3.4. Interaction effects between the BDNF gene and SOB on personality traits

There were interaction effects between SOB (based on traditional criterion) and Val66Met ($P = 0.037$), as well as between SOB (based on astronomic criterion) and rs1491850 ($P = 0.052$), on Novelty Seeking (Table 3). However, when stratifying our sample according SOB, there were no effects of these SNPs on Novelty Seeking. The power to detect interaction effects between rs6265 and SOB, and between rs1491850 and SOB, on Novelty Seeking was 0.82 and 0.85, respectively.

There were interaction effects between SOB and Val66Met ($P = 0.006$ for traditional criterion of SOB, and $P = 0.009$ for astronomic criterion

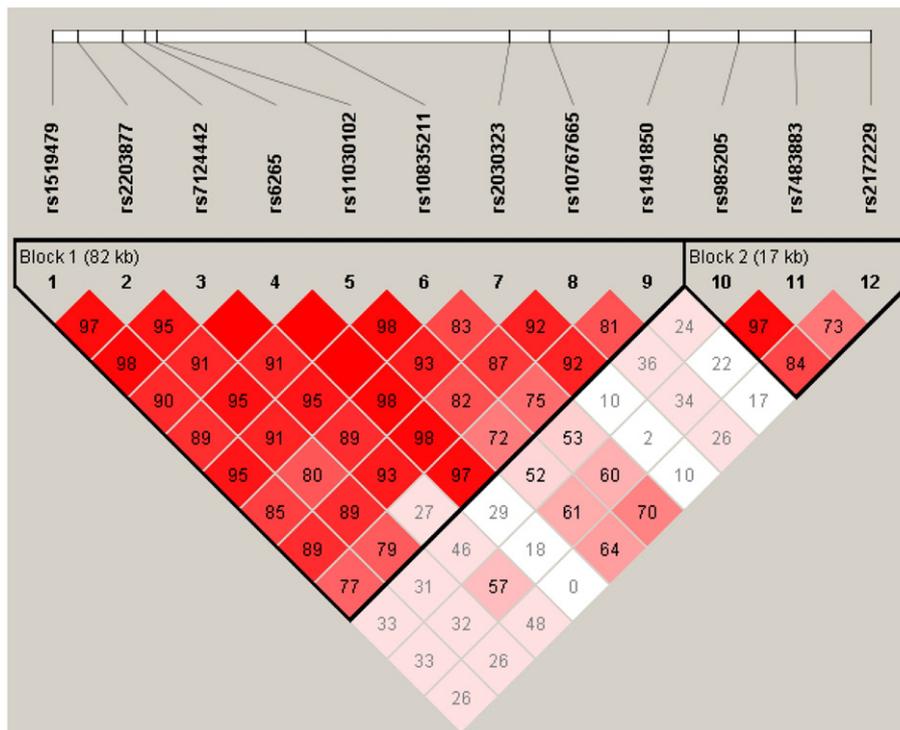


Fig. 1. Haplotype structure and D' -coefficients (multiplied by 100) in the BDNF gene in the total sample.

Table 3
Significant effect modifications by sex or season of birth of *BDNF* genetic associations with personality traits in the combined GxE models (according to Keller, 2014).

Personality trait	Items in linear regression model	β	<i>P</i>	<i>r</i> ²
Novelty Seeking	rs6265	−1.39	0.222	0.035
	SOB (spring)	−0.33	0.803	
	sex	0.93	0.302	
	rs6265*SOB	1.44	0.037	
	Sex*SOB	−0.96	0.129	
	Sex* rs6265	0.43	0.514	
Harm Avoidance	rs1491850	−1.53	0.140	0.037
	SOB _{ast} (summer)	1.67	0.286	
	sex	−0.25	0.803	
	rs1491850* SOB _{ast}	−1.29	0.052	
	Sex* SOB _{ast}	0.29	0.670	
	Sex* rs1491850	0.93	0.122	
Harm Avoidance	rs6265	−0.09	0.944	0.014
	SOB (spring)	2.84	0.068	
	sex	0.55	0.599	
	rs6265*SOB	−2.20	0.006*	
	Sex*SOB	−0.18	0.808	
	Sex*rs6265	0.21	0.778	
Harm Avoidance	rs2030323	0.32	0.791	0.013
	SOB (spring)	2.27	0.141	
	sex	0.86	0.401	
	rs2030323*SOB	−1.90	0.011*	
	Sex*SOB	−0.05	0.950	
	Sex* rs2030323	−0.06	0.935	
Harm Avoidance	rs6265	−1.20	0.366	0.019
	SOB _{ast} (autumn)	−2.59	0.150	
	sex	0.65	0.528	
	rs6265* SOB _{ast}	2.21	0.009	
	Sex*SOB _{ast}	−0.58	0.479	
	Sex*rs6265	0.17	0.822	
Persistence	rs11030102	0.532	0.182	0.013
	sex	0.581	0.093	
	rs11030102*sex	−0.481	0.039*	

SOB_{ast} – season of birth according to astronomic criterion; SOB – season of birth according to traditional criterion. Significant *P*-values are in bold. Statistically significant *P*-values after FDR-correction are marked with asterisk.

of SOB), as well as between SOB (traditional criterion) and rs2030323 ($P = 0.011$), on Harm Avoidance (Table 3). Stratified analysis by SOB demonstrated that Val66Met Met-allele carriers born in spring had lower Harm Avoidance than Val/Val homozygotes ($P = 0.004$, $P_{FDR} = 0.048$, $r^2 = 1.4\%$; Fig. 2, A), whereas associations in other SOB groups were non-significant. Among those born in spring (but not in other seasons), rs2030323 T-allele was associated with lower Harm Avoidance ($P = 0.007$, $P_{FDR} = 0.042$, $r^2 = 1.3\%$; Fig. 2, B). The power to detect interaction effects between rs6265 and SOB, and between rs2030323 and SOB, on Harm Avoidance was 0.96 and 0.91, respectively.

4. Discussion

In the present study, we observed interaction effect between sex and *BDNF* rs11030102 on Persistence, as well as interaction effects between SOB and *BDNF* Val66Met and rs2030323 on Harm Avoidance. Our results showed that alleles associated with a decreased *BDNF* level (Val66Met Met-allele and linked rs2030323 T-allele) were associated with lower Harm Avoidance, but only in those born in spring.

Previous association studies of the *BDNF* gene in personality traits mainly focused on Val66Met due to its functional significance (Egan et al., 2003). Animal and human studies demonstrated that *BDNF* Met/Met-genotype (Chen et al., 2006; Montag et al., 2010) and Met-allele (Jiang et al., 2005) was associated with higher anxiety-related traits as compared with Val/Val homozygotes. However, some other studies are in agreement with the present findings showing that Met-allele is

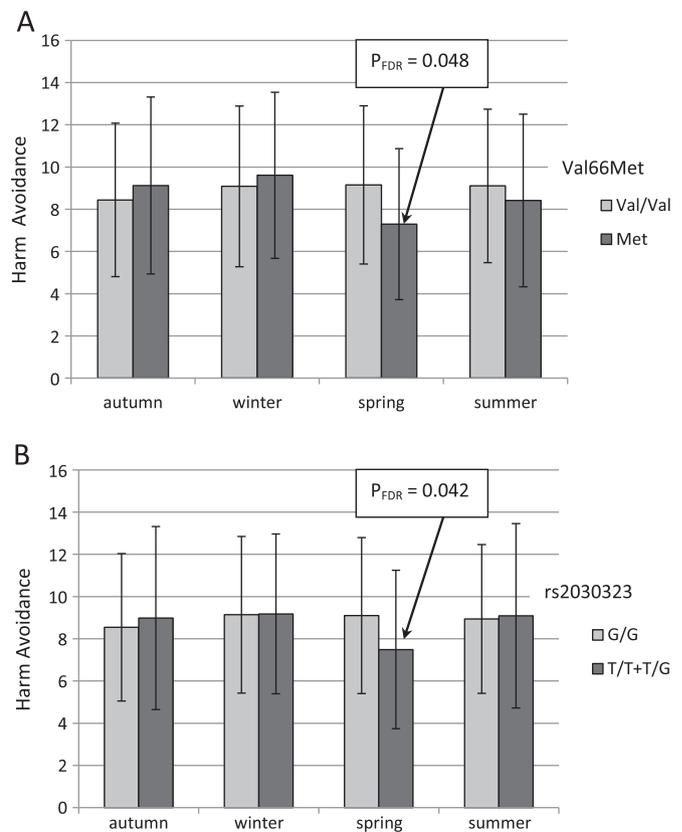


Fig. 2. Interaction effects between *BDNF* Val66Met and SOB (A), and rs2030323 and SOB (B) on Harm avoidance. Error bars stand for standard deviation.

associated with lower Neuroticism (Lang et al., 2005; Sen et al., 2003) and Harm Avoidance (Ando et al., 2012).

Our results demonstrated that the association between the *BDNF* gene and Harm Avoidance was modified by SOB. Variations in personality traits can be influenced by prenatal differences in photoperiod, behavioral rhythms, nutrition, infections, stress and lifestyle (Chotai et al., 2009). According to published studies, downregulation of hippocampal *BDNF* via epigenetic modification might be explained by neonatal iron (Blegen et al., 2013) and zinc deficiency (Chowanadisai et al., 2005), as well as by deficiency of micronutrients involved in one-carbon metabolism (folic acid, vitamin B (Chotai and Adolfsson, 2002), and docosahexaenoic acid (DHA)) (Dhobale and Joshi, 2012). For example, maternal nutrition is sufficiently enriched with micronutrients within the first pregnancy trimester (summer) of individuals born in spring, and this period is known to be important for active brain and nervous system formation, including *BDNF*. This could be an explanation for lower behavioral inhibition (i.e., lower Harm Avoidance) in individuals born in spring. Serum *BDNF* concentrations are shown to be increased in spring-summer as compared to autumn–winter (Molendijk et al., 2012b); this might also explain why lower Harm Avoidance was associated with *BDNF* level in spring borns.

To our best knowledge, the present study is the first one to explore the modifying effect of SOB on genetic association of the *BDNF* gene with personality traits. Previously, interaction effects between the dopamine D4 receptor gene (*DRD4*) and SOB on personality traits (Roussos et al., 2010) and on preference for reciprocal fairness in economic game (Zhong et al., 2010) have been reported. At the same time, the influence of SOB on adult monoamine neurotransmitter turnover is well known (Chotai and Adolfsson, 2002) with a number of studies indicating the involvement of serotoninergic system genes in anxiety-related traits variation (Munafò et al., 2009). Several studies have indicated *BDNF* Val66Met \times 5-HTTLPR interaction effect on anxiety-related traits

and disorders. In a Spanish sample of psychiatrically healthy individuals without familial history of mental disorder, those with *BDNF* Met/Met and *5-HTTLPR* S/S genotypes had higher Harm Avoidance as compared with *BDNF* Met/Met and *5-HTTLPR* L-allele carriers (Arias et al., 2012). In a Russian sample of unaffected parents of patients with major psychosis, carriers of *BDNF* Val/Val and *5-HTTLPR* S/S genotypes scored higher on Depression and Psychasthenia scales (Golimbet et al., 2009). Hiio et al. (2011) revealed that *BDNF* Met-allele carriers with *5-HTTLPR* S/S-genotype scored lower on Conscientiousness. A recent study demonstrated, in agreement with our findings, the effect of *BDNF* Met-allele (in the presence of *5-HTTLPR* S/S-genotype) on higher postnatal depression scores for those born in autumn/winter (Comasco et al., 2011). In the present study, *BDNF* Met-allele carriers showed lower Harm Avoidance only if they were born in spring. Another recent study revealed interaction effect of solar activity and glucocorticoid receptor gene (*NR3C1*) on Neuroticism (anxiety-related trait) (Montag et al., 2013). They reported that *NR3C1* rs41423247 C/C-genotype carriers grown in the womb under the influence of high sun radiation (high solar activity) showed both the highest hippocampal volume in the left hemisphere and lowest Neuroticism scores.

According to published data the *BDNF* Met-allele was shown to be associated with gray matter deficits in hippocampus and prefrontal cortex (Pezawas et al., 2004) – brain regions associated with assigning meaning to social stimuli and stimulus evaluation (Cunningham and Zelazo, 2007). In the present study, there was a sex-specific effect of *BDNF* rs11030102 on Persistence – trait characterizing the maintenance of behavior despite frustration, fatigue, and intermittent reinforcement. Previously, in a large family-based cohort this SNP was associated with serum *BDNF* level (Terracciano et al., 2013). This SNP is located 1.6 kb apart from the functional Val66Met with G-allele being linked with Val66Met Val-allele, reported to be associated with lower Reward Dependence in females (Itoh et al., 2004). Since Persistence was the subscale of Reward Dependence (Cloninger et al., 1993), our findings of a sex-specific association of the *BDNF* gene with a reward-related personality trait are congruent with the study by Itoh et al. (2004). In a sample of women with premenstrual dysphoric disorder, *BDNF* Met-allele was associated with lower fronto-cingulate cortex activation in the luteal phase (characterized by increased progesterone levels). It could be suggested that progesterone might modify the effect of *BDNF* gene on personality traits, possibly via the GABAergic system (Comasco et al., 2014).

Another gene, *BDNF* antisense (*BDNF-AS* or *BDNFOS*), overlaps with the *BDNF* gene and is transcribed in reverse orientation. Inhibition of non-coding *BDNF-AS* transcript upregulates *BDNF* mRNA by two- to sevenfold, alters chromatin marks at the *BDNF* locus, leads to increased protein levels and induces neuronal outgrowth and differentiation both *in vitro* and *in vivo* (Modarresi et al., 2012). Since Val66Met and rs11030102 reside in both *BDNF* and *BDNF-AS*, these SNPs may also be involved in *BDNF-AS* expression regulation on epigenetic level. Moreover, DNA methylation within the promoter/exon IV (Perroud et al., 2013) could be another mechanism of epigenetic regulation of *BDNF* gene. *BDNF* rs2030323 located in intron 3 (3 kb apart from exon IV) may be in linkage disequilibrium with some functional SNP in this region.

4.1. Strengths and limitations

The present study has a number of methodological strengths including homogeneity of the sample in respect to age and education. Our sample had a sufficient power (0.82 – 0.99) to detect the main effects of the investigated SNPs and the proposed GxE interaction effects under the type I error rate of 0.05. Sex as a potential confounder and/or modifier was controlled for in all the statistical models as recommended for GxE interaction studies (Keller, 2014).

However, the study has a number of limitations. First, the use of self-reports for the assessment of personality traits may result in over-

or underreporting some behavior due to social desirability. To minimize these biases, in the present study, individuals were not allowed to discuss questions or answers with anyone. On the other hand, TCI self-report measures were shown to be the strong predictors of self-reported personality by both peer-report measures and ratings by non-acquainted judges (Gruca and Goldberg, 2007). Second, multiple tests have been performed in the present study that may increase the type I error. However, in order to minimize the possibility of false positive results we performed correction for multiple testing using the false discovery rate (FDR) procedure (Simes procedure) (Benjamini and Hochberg, 1995). This procedure provides an effective control of type I error rate in the context of multiple correlated tests and has good agreement with the permutation test (Uher et al., 2009). It is worth noting, that FDR-correction resulted in very few positive findings: only three interaction effects remained statistically significant. Finally, we did not use genomic control to test for genetic homogeneity of our sample; however, the risk for population stratification in our study is likely to be low since all the participants are of Caucasian origin.

5. Conclusion

The present study revealed the interaction effects between the *BDNF* SNPs and SOB on Harm Avoidance, and the interaction effect between the *BDNF* gene and sex on Persistence, in a large sample of Russian young adults. Future studies are necessary in order to replicate these findings in independent samples. Moreover, to get insight into plausible mechanisms of the interaction between the *BDNF* gene and SOB, samples of individuals born in different seasons with detailed information on prenatal and early postnatal influences, as well as on epigenetic markers at birth, are needed.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at <http://dx.doi.org/10.1016/j.pnpbp.2014.08.001>.

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