

# Sensorimotor Gating is Associated with *CHRNA3* Polymorphisms in Schizophrenia and Healthy Volunteers

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Attentional gating deficits, commonly measured by prepulse inhibition (PPI) of the acoustic startle response (ASR), have been established as an endophenotype of schizophrenia. Prepulse inhibition is heritable and has been associated with polymorphisms in serotonin and dopamine system genes. Prepulse inhibition can be enhanced by nicotine, and therefore it has been proposed that schizophrenia patients smoke to ameliorate their early attentional deficits. The PPI-enhancing effects of nicotine in rodents are strain dependent, suggesting a genetic contribution to PPI within the nicotinic acetylcholine receptor (nAChR) system. Recent human genetic studies also imply that tobacco dependence is affected by polymorphisms in the  $\alpha3/\alpha5$  subunits of the nAChR (*CHRNA3/CHRNA5*) gene cluster. We, therefore, investigated the impact of two common *CHRNA3* polymorphisms (rs1051730/rs1317286) on PPI, startle reactivity, and habituation of the ASR in two independent samples of 107 healthy British volunteers and 73 schizophrenia patients hailing from Germany. In both samples, PPI was influenced by both *CHRNA3* polymorphisms (combined  $p$ -value = 0.0027), which were strongly linked. Moreover, *CHRNA3* genotype was associated with chronicity, treatment, and negative symptoms in the schizophrenia sample. These results suggest that sensorimotor gating is influenced by variations of the *CHRNA3* gene, which might also have an impact on the course and severity of schizophrenia.

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## INTRODUCTION

Prepulse inhibition (PPI) of the acoustic startle response is defined as the suppression of the startle reflex that occurs when a relatively weak and non-startling sensory event (the prepulse) is presented 30–500 ms before a strong startle-eliciting stimulus (the pulse; Graham, 1975). The phenomenon, PPI, has been firmly established as an operational measure of sensorimotor gating because it is thought to regulate sensory input by filtering or 'gating out' excess or irrelevant stimuli to prevent sensory information overflow (Braff *et al*, 2001). Prepulse inhibition is regulated by a cortico-striato-pallido pontine (CSPP) circuitry including frontal and mediotemporal brain regions, the ventral striatum, the ventral pallidum, and pontine regions of the brainstem (Koch, 1999; Swerdlow *et al*, 2001). Lesion and

pharmacological challenge studies in rodents have revealed that dopamine, serotonin, glutamate, and acetylcholine are involved in the mediation of PPI (Geyer *et al*, 2001; Swerdlow *et al*, 2001). Recent association studies in healthy humans and schizophrenia patients suggest that genetic variations of the serotonin-2A receptor (5-HT<sub>2A</sub>R), the catechol-*O*-methyltransferase (COMT), and the dopamine-D3 receptor (DRD3) may all affect PPI (Quednow *et al*, 2008b, c, 2009; Roussos *et al*, 2008a, b). However, it is most likely that each of these genetic polymorphisms only explains a fraction of the PPI variance and that further genes, including perhaps genes within the glutamate and acetylcholine system, also contribute to the variant expression of PPI.

Several psychiatric and neurological disorders have been reported to present sensorimotor-gating deficits (for reviews, see Braff *et al*, 2001; Quednow, 2008) but the most consistent findings are PPI deficits in schizophrenia spectrum disorder (Braff *et al*, 1978, 1992; Cadenhead *et al*, 1993, 2000; Kumari *et al*, 2000; Ludewig *et al*, 2003; Parwani *et al*, 2000; Perry *et al*, 2002; Quednow *et al*, 2006b; Swerdlow *et al*, 2006). The findings that PPI is heritable (Anokhin *et al*, 2003; Greenwood *et al*, 2007), that it is

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reduced in unaffected relatives of schizophrenia patients (Cadenhead *et al*, 2000; Kumari *et al*, 2005b), that it is influenced by polymorphisms within the dopamine and serotonin system (Quednow *et al*, 2008b,c, 2009; Roussos *et al*, 2008a,b), and that PPI deficits are present during the prodromal stage, before the clinical diagnosis of schizophrenia (Quednow *et al*, 2008a), suggested that PPI is an important and valid candidate as an intermediate or endophenotypic marker in genetic studies of schizophrenia (Braff and Freedman, 2002; Gottesman and Gould, 2003).

Studies in humans (Baschnagel and Hawk, 2008; Della Casa *et al*, 1998; Duncan *et al*, 2001; Hong *et al*, 2008; Kumari *et al*, 1996, 1997) and animals (Acri *et al*, 1994, 1995; Curzon *et al*, 1994; Faraday *et al*, 1999; Schreiber *et al*, 2002; Spielewoy and Markou, 2004) have shown that the nicotinic acetylcholine receptor (nAChR) agonist, nicotine, enhances PPI. These findings, taken together with the idea that schizophrenia patients have a strongly increased likelihood of smoking (for reviews, see Dalack *et al*, 1998; Ripoll *et al*, 2004), have led to the 'self-medication hypothesis' that proposes that schizophrenia patients may attempt to transiently remedy otherwise deficient attentional processes; this effect, however, could vary depending on genetic background (Kumari and Postma, 2005). Prepulse inhibition studies also contribute to the view that neurocognitive dysfunction in schizophrenia could be based on a dysfunctional nAChR system: George *et al*. (2006) demonstrated that tobacco smoking selectively enhances PPI deficits in schizophrenia and more recently it was also shown that nasally administered nicotine increases PPI in schizophrenia patients and healthy controls (Hong *et al*, 2008). Interestingly, the PPI-enhancing effects of nicotine in schizophrenia patients could be dose-dependently blocked by the non-competitive and non-selective nAChR antagonist, mecamylamine, which suggests that this effect is mediated through activation of central nAChR (George *et al*, 2006). Given that mecamylamine antagonizes  $\alpha 3\beta 2$ ,  $\alpha 3\beta 4$ ,  $\alpha 4\beta 2$ , and  $\alpha 7$  nAChRs but that its antagonism at the  $\alpha 7$  subunit is not functional (Papke *et al*, 2001), the PPI-increasing effects of nicotine are likely explained by an activation of  $\alpha 3\beta 2$ ,  $\alpha 3\beta 4$ , and/or  $\alpha 4\beta 2$  nAChRs (Sacco *et al*, 2005). Moreover,  $\alpha 7$  nAChR knock-out mice display normal PPI levels, and neither selective  $\alpha 7$  nAChR agonists (GTS-21 and AR-R-17779) nor a selective  $\alpha 7$  nAChR antagonist (methyllycaconitine) altered PPI level in rodents (Schreiber *et al*, 2002; Suemaru *et al*, 2004), making it unlikely that  $\alpha 7$  nAChRs significantly contribute to the normal regulation of PPI. At present, it is still unclear whether genetic variation in the nAChR system could have an impact on human PPI. However, the PPI-enhancing effects of nicotine in rodents seem to be strain dependent, which points to a strong genetic modulation of this effect within the nAChR system (Curzon *et al*, 1994; Faraday *et al*, 1998, 1999; Schreiber *et al*, 2002; Spielewoy and Markou, 2004).

Several recent studies have consistently shown that multiple single-nucleotide polymorphisms (SNPs) within the  $\alpha 3/\alpha 5$  nAChR subunit (*CHRNA3/CHRNA5*) gene cluster, on chromosome 15q25.1, are strongly associated with nicotine dependence (Berrettini *et al*, 2008; Bierut *et al*, 2008; Caporaso *et al*, 2009; Saccone *et al*, 2009). Berrettini *et al* (2008) observed that 'cigarettes per day' (CPD) was

associated with a common haplotype in the *CHRNA3/CHRNA5* gene cluster in three independent populations of European origin totaling about 15 000 individuals. In particular, SNPs: rs578776, rs1051730, rs1317286, and rs6495308 were associated with CPD. A recent meta-analysis by Caporaso *et al* (2009) further supports the idea that diverse SNPs of the *CHRNA3* and *CHRNA5* genes are associated with CPD, including rs1051730 in *CHRNA3* (combined  $p$ -value =  $5 \times 10^{-32}$ ). Furthermore, a meta-analysis of schizophrenia linkage studies suggested that chromosomal region 15q21.3–15q26.1 may harbor illness-associated genes (Lewis *et al*, 2003), but presently it is unclear whether there are genetic polymorphisms in the nAChR system, which account for both the vulnerability to schizophrenia and smoking behavior. An altered nAChR system may lead to gating dysfunctions, which could be ameliorated by self-administered nicotine, and may at least explain a portion of the variance for the development of schizophrenia and/or smoking. We therefore assume that certain genetic polymorphisms in nAChR subunits similarly account for attentional endophenotypes in schizophrenia and nicotine dependence.

On the basis of findings from Berrettini *et al* (2008) and the meta-analysis by Caporaso *et al* (2009), we hypothesized that SNPs of the *CHRNA3/CHRNA5* gene cluster would have an impact on sensorimotor gating that is disturbed in schizophrenia and enhanced by nicotine. Given that the  $\alpha 3$  nAChR might be involved in the reversal of sensorimotor-gating deficits in schizophrenia (George *et al*, 2006), we focused on SNPs of the *CHRNA3* gene. We have chosen two of the four SNPs (rs1051730 and rs1317286) that were associated with cigarette smoking in the large study conducted by Berrettini *et al* (2008) and that displayed the highest minor allele frequencies of approximately 30–40% in European-ancestry samples (Bierut *et al*, 2008; International HapMap Consortium, 2003; Saccone *et al*, 2009). We investigated two independent European Caucasian samples—a sample of 107 healthy volunteers recruited in London, United Kingdom, and a sample of 73 schizophrenia inpatients recruited in Bonn, Germany, to directly test for the replicability of our findings.

## MATERIALS AND METHODS

### Participants

For the first experiment, 107 healthy Caucasian volunteers were recruited through local advertisements in South London, United Kingdom. Participants between 18 and 43 years of age were screened for the exclusion criteria of DSM-IV Axis I disorders using the Structured Clinical Interview for DSM-IV Disorders (SCID-I). Additional exclusion criteria were: a history of head injuries, any known neurological abnormalities or systemic illness with known neurological complication, a first-degree relative with psychosis or obsessive-compulsive disorder, and a history of substance abuse or dependence. Current smoking behavior was assessed using the Fagerström Test for Nicotine Dependence (FTND; Heatherston *et al*, 1991) and the CPD value.

For the second experiment, 73 schizophrenia inpatients admitted to the psychiatric hospital of the University of

Bonn, Germany, were considered eligible for the study if the following criteria were met: a diagnosis of schizophrenia according to DSM-IV, age between 18 and 65 years, and Caucasian ethnicity. Patients were excluded if they had a history of head injuries, a neurological disease, a history of substance abuse or dependency, or a severe somatic disease. Every patient was evaluated through SCID-I. Clinical symptoms were measured using the Positive and Negative Syndrome Scale (PANSS; Kay *et al*, 1992). A total of 14 patients were unmedicated, 12 patients received a typical antipsychotic drug, 43 patients were treated with an atypical antipsychotic drug, 3 received two atypical drugs, and 1 received a typical and an atypical antipsychotic. Moreover, three patients received antidepressants and five were treated with benzodiazepines that were tapered 24 h before startle testing. Current smoking status was assessed by the question: Do you smoke more than 10 cigarettes per week (yes: smoker/no: non-smoker)?

For both samples, ethical approvals of the local ethics committees were obtained, and all participants provided written informed consent before inclusion into the study.

### Genotyping

In the British healthy volunteers, SNP genotyping assays were run as submicroliter polymerase chain reaction-based assays on Array Tape (<http://www.douglasscientific.com>) at PreventionGenetics (Marshfield, Wisconsin). For the schizophrenia patients hailing from Germany, SNPs were analyzed by TaqMan assays (Applied Biosystems, Foster City, California). Both procedures are described in detail in Supplementary Material 1.

### Startle Response Measurement

The two samples were assessed with slightly different PPI paradigms. Equipment, set up, PPI testing, and data acquisition and scoring procedures for both have previously been described in detail (London sample: Kumari *et al*, 2005a; Bonn sample: Quednow *et al*, 2006a, b).

In the London healthy volunteer sample, each examination began with a 4-min acclimation period of 70-dB background noise that was continued throughout the session. Participants received 49 white-noise sound pulses at an intensity of 115 dB (duration of 40 ms) separated by variable inter-trial intervals (ITIs) between 9 and 23 s (mean = 15 s). In 36 of the trials, the pulse was preceded by a 20-ms, 85-dB white-noise prepulse with stimulus-onset asynchronies (SOAs) of 30, 60, and 120 ms (12 trials each). The initial trial was a pulse-alone (PA) trial, which was separated for further analysis. All following trials were presented in a pseudo-randomized order. The entire test session lasted approximately 16 min.

In the Bonn schizophrenia sample, each examination began with a 4-min acclimation period of 70-dB background noise that was continued throughout the session. Participants received 73 white-noise sound pulses at an intensity of 116 dB (duration of 40 ms) separated by variable ITIs between 8 and 22 s (mean = 15 s). In 36 of the trials, the pulse was preceded by a 20-ms, 86-dB white-noise prepulse with a SOA of 120 ms. The PA initial trial was separated for

further analysis. All following trials were presented in a pseudo-randomized order. The entire test session lasted approximately 20 min.

To ensure that PPI was not influenced by smoking withdrawal, smoking *ad libitum* was permitted before testing in both laboratories (Kumari and Gray, 1999). Trial exclusion and scoring criteria were identical to those used in previous studies (Quednow *et al*, 2006b). Subjects with response rejections >50% were excluded from data analysis (healthy volunteers:  $n=3$ ; schizophrenia patients:  $n=2$ ).

### Statistical Analysis

The calculation of the mean percent PPI and the habituation measures (percent habituation and linear gradient coefficient *b*) have been described in detail elsewhere (Quednow *et al*, 2006a, b). For the assessment of startle habituation, PA trials were divided each into four blocks in the British sample and in six blocks in the German sample. Startle reactivity was assessed by the mean amplitude of the first block of PA trials and the mean amplitude of all PA trials.

We did *a priori*- and *post hoc*-power analyses of both experiments using G\*Power 3.0.3 (Faul *et al*, 2007). The actual power of both experiments ranged from satisfactory to good: Bonn sample 72%; London sample 80%; and total sample 98% (see Supplementary Table A). All demographic data were analyzed by analysis of variance (ANOVA) with the exception of frequency data. Frequency data were analyzed using  $\chi^2$  tests. Given that sex (Swerdlow *et al*, 1997) and smoking (Kumari and Gray, 1999) could affect PPI, these variables were introduced as covariates in analyses of covariance (ANCOVA) of PPI data independent of the statistical significance of the covariates. All other psychophysiological parameters were analyzed through ANOVA. On the basis of significant main effects or interactions, conservative *post hoc* comparisons according to Bonferroni were performed. The confirmatory statistical comparisons were carried out at a significance level set at  $p < 0.05$  (two-tailed). Within the Pearson Product-Moment correlation analyses, the significance level was set at  $p < 0.01$  (two-tailed) to avoid accumulation of  $\alpha$ -error. Effect size calculations between two groups refer to Cohen's *d*. The Armitage's Trend Test was used to analyze associations between CHRNA3 SNPs and schizophrenia. For a meta-analysis of the genotype effects on psychophysiological parameters across both samples, we used Stouffer's *z*-score method (Whitlock, 2005).

## RESULTS

### London Sample (Healthy Volunteers)

The CHRNA3 rs1051730 and rs1317286 SNPs were in strong linkage disequilibrium ( $r^2=0.967$ ) and both genotype frequencies were distributed in accordance with Hardy-Weinberg equilibrium (HWE) (rs1051730:  $\chi^2(1)=0.09$ ;  $p=0.77$ ; rs1317286:  $\chi^2(1)=0.47$ ;  $p=0.51$ ). Genotype frequencies were also very similar between the UK and German populations. Given that the rs1051730 and rs1317286 SNPs were in such strong linkage disequilibrium and that their effects on the psychophysiological parameters

**Table 1** Demographic Data and Psychophysiological Parameters of British Healthy Human Volunteers

CHRNA3 rs1051730 genotype	TT	TC	CC	Total	F/ $\chi^2$	df/df <sub>err</sub>	p	$\eta_p^2$
N	12 (12.5%)	40 (41.7%)	44 (45.8%)	96 (100%)				
Age (years)	24.6 (1.1)	25.6 (1.0)	26.7 (0.9)	26.0 (0.6)	0.80	2/95	0.45	0.02
Years of education	16.8 (0.9)	16.8 (0.4)	17.4 (0.6)	17.1 (0.3)	0.41	2/95	0.67	0.01
Men (%)	50.0	47.5	47.7	47.9	0.02	2	0.99	—
Current smokers (%)	16.7	25.0	25.0	24.0	0.40	2	0.82	—
Current cigarettes smoked per day (only in smokers)	12.2 (7.9), n = 2	8.6 (1.7), n = 10	8.7 (2.1), n = 11	9.0 (1.3), n = 23	0.25	2/22	0.78	0.02
Fagerström nicotine dependence test (score; only in smokers)	2.0 (2.0), n = 2	1.6 (0.7), n = 10	2.0 (0.7), n = 11	1.8 (0.5), n = 23	0.09	2/22	0.92	0.01
First block, amplitude of pulse-alone trials (arbitrary units)	662 (157.2)	722 (60.6)	700 (55.5)	704 (40.3)	0.11	2/95	0.90	0.00
Mean amplitude of pulse-alone trials (arbitrary units)	520 (133.1)	551 (49.7)	606 (55.2)	572 (36.4)	0.40	2/95	0.67	0.01
Mean percent (%) prepulse inhibition <sup>a</sup> (mean across three SOA conditions: 30, 60, and 120 ms)	20.7 (4.3)	32.6 (2.3)	33.3 (2.2)	31.4 (1.6)	3.67	2/95	0.03 <sup>b</sup>	0.08
Percent (%) early habituation of pulse-alone trials (between first and second block)	37.1 (7.4)	26.8 (4.1)	10.5 (3.9)	20.6 (22.8)	7.10	2/95	0.001 <sup>c</sup>	0.13
Percent (%) total habituation of pulse-alone trials (between first and fourth block)	32.0 (9.9)	33.5 (5.6)	23.9 (4.5)	28.9 (3.3)	0.96	2/95	0.39	0.02
Habituation of pulse-alone trials across 4 blocks (linear gradient coefficient <i>b</i> )	-61.8 (28.8)	-76.9 (13.1)	-58.6 (11.2)	-66.6 (8.3)	0.55	2/95	0.58	0.01

Given that the rs1051730 and rs1317286 SNPs were in strong linkage disequilibrium ( $r^2 = 0.97$ ), the rs1317286 data are only shown in Supplementary Table B. British healthy human volunteers grouped according to their CHRNA3 rs1051730 genotype (mean  $\pm$  SEM in parentheses, gender and smoking status in frequency data)

<sup>a</sup>ANCOVA, means adjusted by covariates gender and cigarettes per day.

<sup>b</sup>Bonferroni *post hoc* test: TT < CT & CC,  $p < 0.05$ .

<sup>c</sup>Bonferroni *post hoc* test: CC < CT,  $p < 0.05$ ; CC < TT,  $p < 0.01$ .

were identical, we only reported the results of the rs1051730 SNP in the main body (for rs1317286 data, see Supplementary Table B). The genotype groups of both SNPs did not differ regarding age, sex, education, smoking status and severity, and startle reactivity (see Table 1 and Supplementary Table B).

Prepulse inhibition was affected by CHRNA3 genotype (see Table 1 and Supplementary Table B, as well as Figure 1a and b). A  $3 \times 3$  (SOA condition  $\times$  genotype) repeated-measures ANCOVA of the rs1051730 SNP with sex and CPD as covariates revealed significant main effects for the factors: SOA condition ( $F(2,90) = 16.3$ ;  $p < 0.001$ ;  $\eta_p^2 = 0.27$ ), sex ( $F(1,91) = 14.4$ ;  $p < 0.001$ ;  $\eta_p^2 = 0.14$ ), and genotype ( $F(2,91) = 3.7$ ;  $p < 0.05$ ;  $\eta_p^2 = 0.08$ ). Bonferroni *post hoc* tests revealed that homozygous carriers of the TT allele showed significantly lower PPI levels compared with homozygous carriers of CC allele ( $p < 0.05$ ; Cohen's  $d = 0.79$ ) and the heterozygous TC allele carriers ( $p < 0.05$ ;  $d = 0.74$ ). Homozygous carriers of CC allele and heterozygous carriers of TC allele did not differ in PPI levels ( $d = 0.04$ ). Analyses of polynomial contrasts across the three genotype groups revealed a significant linear trend ( $p < 0.01$ ) but no significance for a quadratic trend. The analysis of the rs1317286 SNP revealed highly similar results (see Supplementary Table B, ANCOVA statistics and *post hoc* test are not shown).

The main effect of SOA mirrors the well-known nature of PPI to increase with rising SOA from 30 ms to 120 ms (Blumenthal, 1999). The effect of sex reflects the known fact that women have generally lower PPI levels than men (pooled SOA conditions:  $F(1,95) = 14.0$ ;  $p < 0.001$ ;  $\eta_p^2 = 0.15$ ; Swerdlow *et al*, 1997).

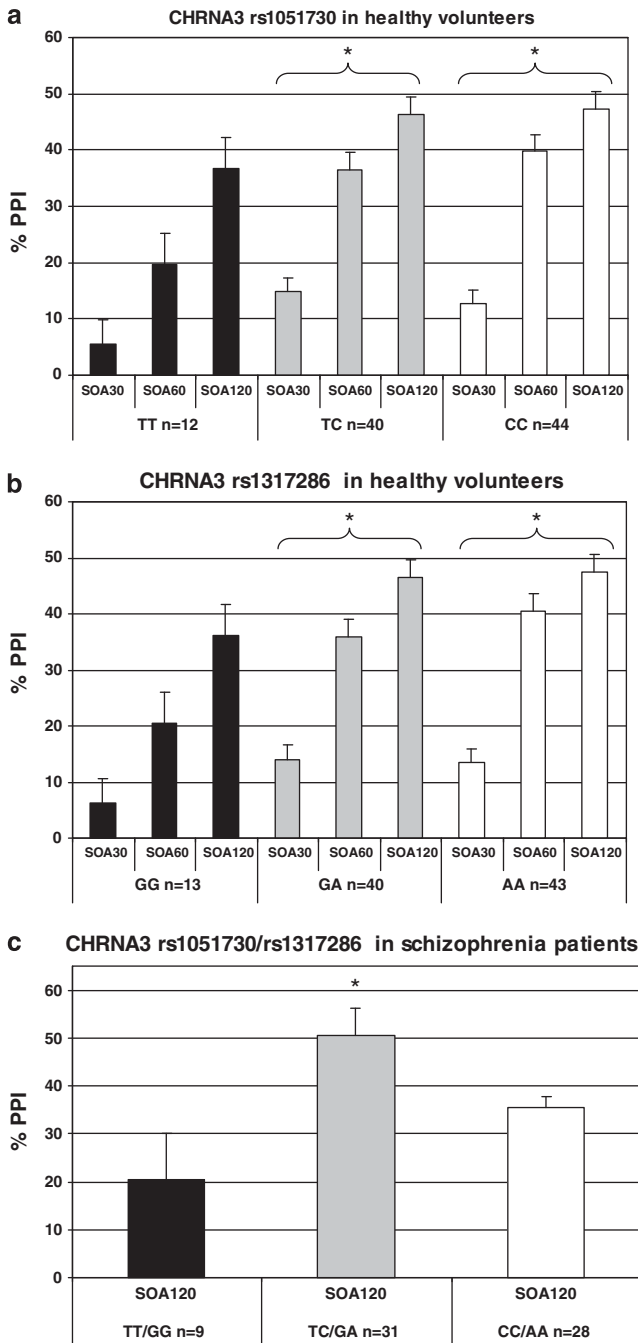
There was also a significant main effect of CHRNA3 genotype on early habituation (see Table 1 and Supplementary Table B) reflecting a diminishing habituation between the first and the second block of PA trials with increasing load of the rs1051730-C and rs1317286-A alleles (polynomial contrasts: linear trend,  $p < 0.002$ ).

Age, smoking parameters, and years of education did not correlate with any of the psychophysiological parameters. In particular, neither the Fagerström score nor 'cigarettes per day' were correlated with PPI (see Supplementary Table C).

### Bonn Sample (Schizophrenia Patients)

The CHRNA3 rs1051730 and rs1317286 SNPs were in complete linkage disequilibrium ( $r^2 = 1.0$ ) and both genotype frequencies were distributed in accordance with HWE ( $\chi^2(1) = 0.05$ ;  $p = 0.82$ ). The genotype groups did not differ regarding age, sex, education, and smoking status (see Table 2). Within the clinical data, there was a significant main effect of genotype on the number of psychotic





**Figure 1** The effects of *CHRNA3* genotype on percent prepulse inhibition (PPI) of the acoustic startle response in healthy human volunteers and schizophrenia patients (means  $\pm$  SEM, adjusted for sex and smoking): (a) *CHRNA3* rs1051730 polymorphism in 96 British healthy human volunteers (Bonferroni *post hoc* test vs TT allele group: \* $p < 0.05$ ), (b) *CHRNA3* rs1317286 polymorphism in 96 British healthy human volunteers (Bonferroni *post hoc* test vs GG allele group: \* $p < 0.05$ ), and (c) the completely associated *CHRNA3* rs1051730 and rs1317286 polymorphisms in German schizophrenia patients (Bonferroni *post hoc* test vs TT/GG allele group: \* $p < 0.05$ ).

significantly higher doses of antipsychotics compared with the two other genotype groups. In contrast, homozygous carriers of the CC/AA allele revealed less negative symptoms than heterozygotes. Further clinical data were not affected by *CHRNA3* genotype.

In contrast to all other startle parameters, PPI was again significantly influenced by *CHRNA3* genotype (ANCOVA controlled for sex and smoking status (yes/no):  $F(2,67) = 4.3$ ,  $p < 0.05$ ,  $\eta_p^2 = 0.13$ ; see Table 2 and Figure 1c). Bonferroni *post hoc* tests revealed that homozygous carriers of the TT/GG allele displayed significantly lower PPI levels compared with heterozygous carriers of the TC/GA genotype ( $p < 0.05$ ,  $d = 1.09$ ). Both homozygous groups did not significantly differ from each other with respect to PPI; however, the difference showed a moderate effect size ( $d = 0.55$ ). Prepulse inhibition in the CC/AA and the TC/GA group did not differ significantly but again the difference was present with a moderate effect size ( $d = 0.55$ ). Analyses of polynomial contrasts across the three genotype groups revealed a significant quadratic trend ( $p < 0.01$ ) but no significance for a linear trend. If homozygous carriers of TT/GG allele were compared with a merged group of carriers of the C/A allele (TC/GA + CC/AA), both groups differed significantly regarding PPI (ANCOVA controlled for sex and smoking status:  $F(1,64) = 3.94$ ,  $p < 0.05$ ,  $\eta_p^2 = 0.06$ ). Further ANCOVAs of the three genotype groups introducing chlorpromazine equivalents, number of psychotic episodes, and PANSS negative symptoms as further covariates still revealed significant main effects for the factor genotype.

Correlational analysis of startle parameter and patient characteristics revealed that age was negatively correlated with startle reactivity (mean amplitude PA trials:  $r = -0.38$ ,  $p = 0.001$ ; first block of PA trials:  $r = -0.38$ ,  $p = 0.002$ ), indicating that older patients displayed decreased startle reactivity. Moreover, the startle reactivity was negatively associated with the PANSS positive score (mean amplitude PA trials:  $r = -0.33$ ,  $p = 0.007$ ; first block of PA trials:  $r = -0.36$ ,  $p = 0.003$ ), negative score (mean amplitude PA trials:  $r = -0.37$ ,  $p = \text{n.s.}$ ; first block of PA trials:  $r = -0.31$ ,  $p = 0.01$ ), and total score (mean amplitude PA trials:  $r = -0.38$ ,  $p = 0.002$ ; first block of PA trials:  $r = -0.40$ ,  $p = 0.001$ ). These findings indicate that low startling patients displayed more severe psychotic symptoms. Finally, the number of psychotic episodes was inversely correlated with startle reactivity (mean amplitude PA trials:  $r = -0.33$ ,  $p = 0.007$ ; first block of PA trials:  $r = -0.29$ ,  $p = \text{n.s.}$ ), reflecting that more chronic patients also showed less startle reactivity. Prepulse inhibition did not correlate with any clinical or demographic data. Smoking status was also not correlated with PPI (see Supplementary Table C).

Owing to the methodological differences, German schizophrenia patients and the British controls could not be directly compared regarding startle parameters and PPI. However, in line with many previous studies, the total group of schizophrenia patients displayed significantly decreased PPI levels when compared with an equivalent group of healthy controls ( $t(114) = 3.2$ ,  $p < 0.003$ , controls: 56.9% PPI (SEM 3.2),  $N = 46$ ), who were assessed in Bonn with the same PPI paradigm in our previous studies (Quednow et al, 2006b, 2008a).

episodes, daily chlorpromazine equivalents, and negative symptoms measured with the PANSS (see Table 2). Homozygous carriers of the TT/GG allele displayed significantly more psychotic episodes than heterozygotes and received

**Table 2** Demographic Data and Psychophysiological Parameters of German Schizophrenia Patients

<i>CHRNA3</i> rs1051730 genotype	TT	TC	CC	Total	F/ $\chi^2$	df/df <sub>err</sub>	p	$\eta_p^2$
<i>CHRNA3</i> rs1317286 genotype	GG	GA	AA					
N	9 (13.2%)	31 (45.6%)	28 (41.2%)	68 (100%)				
Age (years)	35.3 (2.5)	33.8 (1.6)	35.5 (2.4)	34.7 (1.3)	0.20	2/67	0.81	0.01
Years of education	15.1 (0.8)	14.8 (0.5)	13.9 (0.7)	14.5 (0.3)	0.86	2/67	0.43	0.03
Men (%)	77.8	71.0	64.3	69.1	0.67	2	0.72	—
Current smokers (%)	22.2	54.8	64.3	54.4	4.86	2	0.09	—
Patients with a first-episode (%)	33.3	71.0	64.3	63.2	4.27	2	0.12	—
Age of onset (years)	28.2 (2.5)	31.4 (1.7)	30.1 (2.0)	30.4 (1.2)	0.40	2/67	0.68	0.01
Duration of illness (years)	7.1 (3.0)	2.5 (0.9)	5.8 (2.0)	4.5 (1.0)	1.76	2/67	0.18	0.05
Number of episodes	3.3 (0.9)	1.6 (0.2)	2.1 (0.4)	2.0 (0.2)	3.81	2/67	0.03 <sup>a</sup>	0.11
Medication status (%) (unmedicated/ typical/atypical antipsychotic drug)	0/22/78	16/16/68	32/18/50	21/18/62	5.30	4	0.26	—
Daily chlorpromazine equivalents <sup>b</sup>	463 (68.0)	246 (32.2)	228.8 (42.8)	266 (26.0)	4.67	2/67	0.01 <sup>c</sup>	0.14
PANSS positive	18.7 (2.3)	19.1 (1.5)	17.3 (1.4)	18.3 (0.9)	0.42	2/67	0.66	0.01
PANSS negative	21.4 (1.6)	23.0 (1.4)	17.5 (1.4)	20.5 (0.9)	4.40	2/67	0.02 <sup>d</sup>	0.13
PANSS general	43.1 (4.8)	43.1 (2.4)	36.8 (2.5)	40.5 (1.7)	1.83	2/67	0.17	0.06
PANSS total	83.2 (8.1)	85.6 (4.7)	71.5 (4.9)	79.5 (3.2)	2.25	2/67	0.11	0.07
First block, amplitude of pulse-alone trials (arbitrary units)	280 (97.7)	278 (27.6)	248.5 (36.8)	266 (23.1)	0.20	2/67	0.82	0.01
Mean amplitude of pulse-alone trials (arbitrary units)	238 (89.9)	221 (28.0)	179.2 (29.2)	206 (20.9)	0.61	2/67	0.55	0.02
Mean percent prepulse inhibition <sup>e</sup> (SOA: 120 ms)	20.3% (9.8)	50.6% (5.6)	35.5% (2.3)	41.9% (3.3)	4.26	2/63	0.02 <sup>f</sup>	0.13
Percent early habituation of pulse-alone trials (between first and second block)	24.0% (9.2)	23.4% (4.9)	22.6% (5.1)	23.2% (3.3)	0.01	2/67	0.99	0.00
Percent total habituation of pulse-alone trials (between first and sixth block)	42.2% (10.7)	37.3% (6.0)	40.4% (5.5)	39.3% (3.8)	0.12	2/67	0.89	0.00
Habituation of pulse-alone trials across six blocks (linear gradient coefficient b)	-13.6 (4.6)	-16.0 (3.5)	-17.8 (3.7)	-16.4 (2.3)	0.18	2/67	0.84	0.01

*CHRNA3* rs1051730 and rs1317286 polymorphisms were in complete linkage disequilibrium.

German schizophrenia patients grouped according to their *CHRNA3* rs1051730 and rs1317286 genotype (mean  $\pm$  SEM values in parentheses, gender and smoking status in frequency data)

<sup>a</sup>Bonferroni *post hoc* test: TT/GG > TC/GA,  $p < 0.05$ .

<sup>b</sup>Unmedicated patients received the value zero.

<sup>c</sup>Bonferroni *post hoc* test: TT/GG > TC/GA & CC/AA,  $p < 0.05$ .

<sup>d</sup>Bonferroni *post hoc* test: CC/AA < TC/GA,  $p < 0.05$ .

<sup>e</sup>ANCOVA, means adjusted by covariates gender and smoking status.

<sup>f</sup>Bonferroni *post hoc* test: TT/GG < TC/GA,  $p < 0.05$ .

### Meta-Analysis of Genotype Effects on Startle Parameter

To assess the total effect of *CHRNA3* genotype on psychophysiological parameters across both samples, we combined the  $p$ -values in a meta-analytic approach by using Stouffer's  $z$ -score method (see Table 4). Prepulse inhibition was the only parameter still showing a highly significant effect of genotype ( $p = 0.0027$ ) even after Bonferroni correction for multiple phenotypes.

### Association Between *CHRNA3* SNPs and Schizophrenia

There was no association of the tested *CHRNA3* SNPs with a diagnosis of schizophrenia *per se* (Table 3). However, given that the control sample and the patient sample were rather small and have low power to detect common, low-risk

variants, and that the samples were recruited in different European countries, this result should not be over interpreted.

### DISCUSSION

This study is, to the best of our knowledge, the first that investigates whether sensorimotor gating depends on polymorphisms in the  $\alpha 3$  subunit nAChR gene *CHRNA3*. We have convincingly demonstrated that the TT genotype of rs1051730 and the GG genotype of rs1317286 were associated with decreased PPI levels in both healthy volunteers and in schizophrenia patients (combined  $p$ -value 0.0027). The *CHRNA3* rs1051730 T allele and rs1317286 G allele have been firmly established as risk alleles for tobacco

**Table 3** Frequencies of *CHRNA3* rs1051730 and rs1317286 Genotypes

<i>CHRNA3</i> genotype	Controls/cases	T allele	C allele	TT (n)	TC (n)	CC (n)	Armitage's trend test
rs1051730 (%)	Controls	33.8	66.2	12.1 (12)	43.4 (43)	44.4 (44)	Odds ratio = 0.952/1.050 $\chi^2 = 0.048$ ; $p = 0.827$
	Schizophrenia patients	35.0	65.0	12.9 (9)	44.3 (31)	42.9 (30)	
		G allele	A allele	GG (n)	GA (n)	AA (n)	
rs1317286 (%)	Controls	35.4	64.6	14.1 (14)	42.4 (42)	43.4 (43)	Odds ratio = 0.975/1.026 $\chi^2 = 0.004$ ; $p = 0.948$
	Schizophrenia patients	35.0	65.0	12.9 (9)	44.3 (31)	42.9 (30)	

Frequencies of *CHRNA3* rs1051730 and rs1317286 genotypes in 99 British Caucasian healthy controls and 70 German Caucasian schizophrenia patients (full sample inclusive of cases that dropped out for the electrophysiological analyses due to low startle data quality)

**Table 4** Meta-Analysis of *CHRNA3* Genotype Effects on Startle Parameters

	London sample mean $p^a$	Bonn sample $p$	Combined $p$ -value <sup>b</sup>
First block, amplitude of pulse-alone trials	0.95	0.82	0.97
Mean amplitude of pulse-alone trials	0.63	0.55	0.64
Mean percent prepulse inhibition	0.025	0.02	0.0027
Percent early habituation of pulse-alone trials	0.002	0.99	0.15
Percent total habituation of pulse-alone trials	0.37	0.89	0.67
Habituation slope of pulse-alone trials (b)	0.52	0.84	0.73

Meta-analysis performed across both investigated samples (London healthy volunteers ( $n = 96$ ) and Bonn schizophrenia patients ( $n = 68$ ))

<sup>a</sup>Averaged between rs1051730 and rs1317286 genotype.

<sup>b</sup>Stouffer's z-score method.

dependence (Berrettini et al, 2008; Bierut et al, 2008; Caporaso et al, 2009; Saccone et al, 2009). In our patients, these *CHRNA3* SNPs were also associated with number of previous psychotic episodes, daily chlorpromazine dose, and negative symptoms—all indicators for severity of the illness. Thus, further investigation of the *CHRNA3* SNPs regarding risk, course, and treatment of schizophrenia in larger samples is warranted. There was no indication of an association with schizophrenia *per se*, but the sample size and design of this study obviously do not allow drawing conclusions from that. To the best of our knowledge, there are no published studies that have investigated *CHRNA3/CHRNA5* gene variants for association with schizophrenia (<http://www.schizophreniaforum.org/res/sczgene>; Allen et al, 2008), despite the location of these genes within

a region on chromosome 15q possibly linked with schizophrenia (Lewis et al, 2003). The demonstration in this study that *CHRNA3* gene variants are associated with an established endophenotype of schizophrenia might motivate a closer look at these genes in large-scale association studies of schizophrenia. Moreover, we initially found a significant *CHRNA3* effect on early habituation in the London sample, but this effect could not be replicated in the Bonn sample. In the combined analysis of both samples (Table 4), we did not find an overall effect of *CHRNA3* genotype on any habituation measure.

At which level of the brain might mutations in the  $\alpha 3$  nAChR subunits affect PPI?  $\alpha 3$  nAChR mRNA is expressed in high densities in the prefrontal cortex (PFC), the motor and the entorhinal cortex, and in different nuclei of the thalamus, and in lower densities also throughout the hippocampus (Gotti and Clementi, 2004; Paterson and Nordberg, 2000)—all structures that are also implicated in the mediation of PPI (Koch, 1999; Swerdlow et al, 2001). Moreover, there is evidence that  $\alpha 3$  nAChR subunits regulate dopamine neurotransmission in the striatum (Quik et al, 2005; Salminen et al, 2004), which is a key element of the CSPP circuitry (Koch, 1999; Swerdlow et al, 2001). Thus, the effects of *CHRNA3* genotype on PPI could be either direct within the CSPP circuitry or indirect via the striatal dopamine system.

Our findings support the view that the cholinergic system has a key role in pre-attentional and attentional mechanisms. Recent evidence indicates that the cholinergic system has an integrating or synchronizing function in the central nervous system, especially in the PFC (Mansvelder et al, 2009). nAChRs seem to function as gatekeepers in neural transmission as pre-synaptic nAChRs tune inhibitory and excitatory inputs of the neuron (Mansvelder et al, 2009). This is true for modulating glutamatergic transmission (Vidal and Changeux, 1993) and for dopaminergic neurotransmission (Salminen et al, 2004), both of which are relevant for PPI (Koch, 1999; Swerdlow et al, 2001).

Our study showed that PPI is likely to be mediated by the  $\alpha 3$  subunit of the nAChR, and the results of the mecamylamine study by George and colleagues (2006)

would be consistent with that. Given that schizophrenia patients display PPI deficits, that PPI and symptom severity are modulated by *CHRNA3* polymorphisms, and that the nAChR system is an important neuromodulator, a future promising target of schizophrenia research might be the development of new medications for treating schizophrenia or at least cognitive deficits in schizophrenia, which affect the  $\alpha 3$  subunits of the nAChR in the central nervous system. It is possible that abnormally reactive cholinergic responsiveness is sufficient to cause an altered dopaminergic transmission (Briand *et al*, 2007), hence stabilizing the cholinergic system should also regulate the dopaminergic system that is well-known to be affected in schizophrenia (Howes and Kapur, 2009).

A measure of sensory gating is the suppression of the auditory-evoked potential P50 response to repeated stimuli (for a review, see Turetsky *et al*, 2007). Both P50 suppression and PPI are modulated by the nAChR agonist, nicotine, which suggests that brain cholinergic systems are involved in gating processes common to both paradigms (Turetsky *et al*, 2007). However, sensory gating measured by P50 suppression and sensorimotor gating measured by PPI seem to be largely independent phenomena because they are not correlated (Braff *et al*, 2007; Oranje *et al*, 2006; Schwarzkopf *et al*, 1993). Our results further support that P50 suppression and PPI are likely to be two different inhibitory mechanisms. Although P50 suppression seems to depend on  $\alpha 7$  subunits of the nAChR (Freedman *et al*, 1997), PPI is very likely to be influenced by  $\alpha 3$  nAChR functioning as shown here. Animal studies further support this hypothesis:  $\alpha 7$  nAChR knock-out mice exhibit normal PPI levels and neither selective  $\alpha 7$  nAChR agonists (GTS-21 and AR-R-17779) nor a selective  $\alpha 7$  nAChR antagonist (methyllycaconitine) altered PPI level in rodents (Schreiber *et al*, 2002; Suemaru *et al*, 2004). On the basis of the seminal findings of Freedman *et al* (1997),  $\alpha 7$  agonists were suggested as cognitive enhancer in schizophrenia, but the  $\alpha 7$  nAChR partial agonist, DMXB, failed to improve cognitive deficits in schizophrenia in a recent clinical trial (Freedman *et al*, 2008). However,  $\alpha 7$  and  $\alpha 4\beta 2$  nAChRs developed receptor insensitivity (tachyphylaxis) if there were repeatedly stimulated (Kuryatov *et al*, 2000), making these receptors less suitable targets for a continuous drug treatment (Harvey, 2009). In contrast, other nAChRs, including the  $\alpha 3$  subunit, are largely resistant to inactivation after repeated stimulation (Kuryatov *et al*, 2000) and, thus,  $\alpha 3$  subunits of the nAChR might be more promising targets for potential cognitive enhancers in schizophrenia.

Interestingly, effects of specific pharmacological challenges on PPI in rodents have been replicated in human genetic studies. Previously, it was consistently shown that dopaminergic, serotonergic, glutamatergic, and acetylcholinergic challenges have an impact on PPI in rodents (for review, see Geyer *et al*, 2001). In recent years, it was demonstrated that polymorphisms within the human dopamine and serotonin systems affect PPI (Quednow *et al*, 2008b, c, 2009; Roussos *et al*, 2008a, b). This study extends these findings to the nAChR system, as we have clearly shown that SNPs within the  $\alpha 3$  subunit of the nAChR also influence PPI in humans. Thus, these recent developments predict that polymorphisms within the glutamatergic system might also explain a portion of the PPI variance in the population.

One limitation of this study is that the two independent samples were assessed using slightly different PPI paradigms. This precludes a comparison of the genetic effects on PPI between samples. However, the replication across samples and study sites despite some procedural differences underscores the robustness of our findings. A second limitation is that we did not assess severity of smoking behavior in the schizophrenia sample assessed in Bonn. However, in contrast to some previous studies (Duncan *et al*, 2001; Swerdlow *et al*, 2006), the method that we used in Bonn was less susceptible to the effects of smoking status on PPI, whereas it was proven sensitive to psychopathological modulations of PPI (Quednow *et al*, 2006a, b, c, 2008a, b). This might have two reasons: (1) smoking withdrawal can decrease PPI (George *et al*, 2006; Kumari and Gray, 1999), but our subjects were allowed to smoke *ad libitum* before testing; (2) heavy smokers with schizophrenia display higher PPI levels than light- or non-smoking patients (Swerdlow *et al*, 2006), which is in line with the finding that nicotine itself could enhance PPI (Kumari *et al*, 1997; Postma *et al*, 2006). As we did not assess smoking severity, the effect of smoking status alone was possibly too small to detect in our sample. Moreover, the investigation of smoking  $\times$  genotype associations was not the main focus of this study. Thus, we believe that the lack of detailed smoking data in the schizophrenia sample does not impair our main results presented here. A further minor limitation is that the low number of smokers in the London healthy controls did not allow for separate analyses of *CHRNA3* genotype effects on PPI in smokers and nonsmokers. Thus, the power to assess smoking status effects on PPI, as a function of *CHRNA3* genotype, was restricted because of the small number of smokers in the healthy controls (and the limited sample size in the schizophrenia sample). However, controlling for smoking status in statistical analyses, and excluding the 23 smokers in the British healthy volunteers did not alter the results (genotype effect on PPI (corrected for gender): rs1051730:  $F(2, 69) = 3.98$ ;  $p < 0.05$ ;  $\eta_p^2 = 0.10$ ; rs1317286:  $F(2, 69) = 4.28$ ;  $p < 0.05$ ;  $\eta_p^2 = 0.11$ ), suggesting that the reported association is not mediated or confounded by current nicotine use. In addition, 17 of the 23 smokers had a Fagerström score of 0–2, reflecting very low addiction in most of the smoking healthy controls. Future studies might systematically investigate the interaction of chronic smoking and acute nicotine administration with genetic polymorphisms in their effect on sensory gating and attentional functions. Extending the self-medication hypothesis, we would hypothesize stronger nicotine effects on PPI in carriers of gene variants associated with smoking or with schizophrenia. Finally, we have only investigated two SNPs of the  $\alpha 3$  subunits of the nAChR. Hence, mutations in other subunits and of nAChR haplotypes should be examined in future studies.

One might be surprised that we did not replicate the association of smoking severity and *CHRNA3* genotype (Berrettini *et al*, 2008; Bierut *et al*, 2008; Caporaso *et al*, 2009; Saccone *et al*, 2009). However, recent studies showing an association of smoking behavior with *CHRNA3* genotype all included relatively strong smokers (eg, Caporaso *et al*, 2009: 18.5–22.0 CPD), whereas we had only seven smokers in the total healthy control sample ( $n = 96$ ) that smoked



more than 15 cigarettes per day. Moreover, in the schizophrenia sample, we did not assess smoking severity and none of the previous studies reported an association of smoking status *per se* and *CHRNA* genotype. Lastly, in contrast to the previous association studies on smoking and *CHRNA3*, we have investigated relatively small samples, indicating that the *CHRNA* effect on PPI seems to be much greater than on smoking severity. Thus, our samples were not ideal for testing smoking  $\times$  *CHRNA3* genotype associations, which was, however, also not the focus of our study.

In conclusion, our results confirmed previous pharmacological studies that proposed an impact of the nAChR system on sensorimotor gating. We have observed that polymorphisms of the  $\alpha 3$  subunits of the nAChR explained approximately 8–13% of the PPI variance in two independent samples, suggesting a major role of  $\alpha 3$  subunits of the nAChR in the mediation of PPI. Finally,  $\alpha 3$ -containing nAChRs might be explored as a new pharmacological target for cognitive enhancement in patients with schizophrenia.

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## DISCLOSURE

The authors declare no conflict of interest.

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