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# COMT polymorphism modulates the resting-state EEG alpha oscillatory response to acute nicotine in male non-smokers

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## **Abstract**

Performance improvements in cognitive tasks requiring executive functions are evident with nicotinic acetylcholine receptor (nAChR) agonists, and activation of the underlying neural circuitry supporting these cognitive effects is thought to involve dopamine neurotransmission. As individual difference in response to nicotine may be related to a functional polymorphism in the gene encoding catechol-O-methyltransferase (COMT), an enzyme that strongly influences cortical dopamine metabolism, this study examined the modulatory effects of the COMT Val158Met polymorphism on the neural response to acute nicotine as measured with resting-state electroencephalographic (EEG) oscillations. In a sample of 62 healthy non-smoking adult males, a single dose (6 mg) of nicotine gum administered in a randomized, double-blind, placebocontrolled design was shown to affect a oscillatory activity, increasing power of upper a oscillations in frontocentral regions of Met/Met homozygotes and in parietal/occipital regions of Val/Met heterozygotes. Peak  $\alpha$  frequency was also found to be faster with nicotine (vs. placebo) treatment in Val/Met heterozygotes, who exhibited a slower  $\alpha$  frequency compared to Val/Val homozygotes. The data tentatively suggest that interindividual differences in brain  $\alpha$  oscillations and their response to nicotinic agonist treatment are influenced by genetic mechanisms involving COMT.

## **Keywords**

Alpha; catechol-(	O-methyltransfer	ase; cognition;	dopamine; e	lectroencepha	alography;	genotype;
nicotine; oscillation	ons; polymorphi	sm; resting state	e			

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The cognitive enhancement properties of nicotinic acetylcholinergic receptor (nAChR) agonists such as nicotine (Heishman *et al.* 2010) are associated with their moderating effects on the dopamine (DA) pathway connecting the ventral tegmental area (VTA) with cortical regions, including the prefrontal cortex (PFC). Agonists effect this enhancement by binding to nAChRs on VTA DA projection neurons, increasing dopamine signaling and processing in cortical networks (Jasinska *et al.* 2013; Livingstone & Wonnacott 2009; Mansvelder *et al.* 2006). A range of evidence has further shown that nicotinic stimulation increases dopamine concentrations in the PFC, where stimulation of presynaptic nAChRs elevates dopamine levels and influences cognitive processes (Wallace & Bertrand 2013).

Studies of cognitive performance (Newhouse et al. 2004; Perkins 1999) and patterns of activation of task-specific neural networks (Bentley et al. 2011; Newhouse et al. 2011) show considerable intersubject response variability to nicotine and nicotinic agonists, often resembling an 'inverted-U' shaped function (persons exhibiting suboptimal performance prior to drug challenge tend to show performance benefits and normalized neural activity with nicotine, while those performing at optimal level or exhibiting task-related neural efficiency tend to show nil or diminished cognitive and neural response to nicotine). While the neurobiological causes underlying this heterogeneity are not well understood, there is an increasing trend to use molecular genetic approaches to assay individual differences in cognitive functions (Greenwood & Parasuraman 2003) and response to pharmacological treatments (Apud & Weinberger 2006; Goldstein et al. 2007), including nicotine (Herman & Sofuoglu 2010). Such approaches focus on allelic variations in the pharmacodynamic and pharmacokinetic properties of neurotransmitter genes involved in the different aspects of cognition (Parasuraman 2009). Of the likely candidate genes influencing response heterogeneity to nicotine, those regulating DA neurotransmission show promise as pharmacological studies in animals (Grannon et al. 2000) and humans (Kimberg et al. 1997; Mattay et al. 2000; Mehta et al. 2000) indicate that the effects of amphetamine and other dopamimetic drugs are baseline dependent. Relatively poor performers on prefrontal cognitive tasks have improved with treatment, whereas high performers have shown no response or response deterioration.

Catabolic flux of synaptic dopamine in the cortex is controlled primarily by the enzyme catechol-O-methyltransferase (COMT) (Huotari *et al.* 2002). The COMT gene contains a single nucleotide polymorphism that produces a valine-to-methionine (Val/Met) substitution at position 158 (Val158Met), producing a trimodal distribution of enzyme activity (Floderus *et al.* 1981; Lachman *et al.* 1996). Met158 homozygotes biotransform dopamine less than Val carriers, showing one third less COMT enzymatic activity in brain. Hence, there are higher extracellular dopamine levels with Met homozygotes (Chen *et al.* 2004). Val/Met heterozygotes exhibit intermediate levels of COMT activity (Weinshilboum *et al.* 1999).

Thus, COMT is an excellent candidate gene for modulating dopamine levels and function in the cortex and for determining where on the inverted-U shaped curve of dopamine function an individual lies (Tunbridge *et al.* 2006). However, inconsistent behavioral data and performance meta-analyses support only a weak association between COMT polymorphisms and individual differences in PFC function (Barnett *et al.* 2007; Munafo *et al.* 2005). Nevertheless, investigations of individual differences with intermediate brain-based

phenotypes, more sensitive for detecting gene effects on the brain (Green *et al.* 2008; Parasuraman & Jiang 2012), have found greater cortical processing efficiency in Met158 homozygotes compared to Val158 homozygotes, with heterozygotes displaying intermediate activation levels (Egan *et al.* 2001; Heinz & Smolka 2006). Acute dosing with amphetamine, which elevates synaptic dopamine levels, increased PFC task-evoked cortical efficiency in individuals with the Val/Val genotype, who have presumed low prefrontal synaptic dopamine, and reduced PFC processing efficiency compared to low-activity Met/Met genotypes (Mattay *et al.* 2003).

The limited functional magnetic resonance imaging (fMRI) investigations of the COMT polymorphism's effects on the cerebrovascular activational response to nicotine in smokers have shown mixed results. For example, Val/Val smokers were more prone to cognitive impairment and reduced prefrontal activation during smoking abstinence (Loughead *et al.* 2009) but another study showed Met/Met smokers with significant activation reduction of frontal executive control regions during cessation (Ashare *et al.* 2013). Val/Val genotypes experienced more severe withdrawal symptoms following cessation, with greater subjective effects from acute intravenous nicotine (Herman *et al.* 2013; Lee *et al.* 2013). Because these studies in chronic smokers may simply reflect a 'remediation' of a cortical deficiency during nicotine withdrawal (Ashare *et al.* 2014; Beaver *et al.* 2011; Cole *et al.* 2010), a clearer picture of COMT-mediated response differences to nicotine may be obtained using nicotine-naïve volunteers examined with electrophysiological probes that permit direct, instantaneous detection of neuronal activity.

Electroencephalographic (EEG) studies have linked neuronal oscillations at low and high frequency ranges with specific cognitive functions (Lopes da Silva 2013; Uhlhaas *et al.* 2009; Wang 2010). These oscillations are key to sculpting temporal coordination of neural networks governing cognitive functions such as perception, attention and working memory (Cantero & Atienza 2005; Kaiser & Lutzenberger 2003; Lisman & Buzsaki 2008). The basic building blocks defining these oscillations can be probed with the spectral profiling of EEG recordings during a resting state (Narayanan *et al.* 2014), and have shown that distinct changes in oscillatory activity in low and high frequencies are associated with different drug classes (Knott 2000; Saletu *et al.* 2002), including cognitive-enhancing drugs (Ahnaou *et al.* 2014; Leiser *et al.* 2011) and nicotine (Knott 1990).

In acute smoking, EEG studies have shown a characteristic, stimulant-like pharmaco-EEG profile – accelerating the dominant (a) oscillatory frequency (PAF), increasing power of  $a_2$ ,  $\beta$ , decreasing power of  $\delta$ ,  $\theta$ ,  $a_1$  (Knott 2001; Knott & Venables 1977). Similar patterns are observed in smokers using nicotine replacement products (Knott *et al.* 1999; Lindgren *et al.* 1999; Pickworth *et al.* 1986, 1988; Teter *et al.* 2002). Individual differences in these profiles are reported, with power variations in accord with performances on frontal lobe tasks (Knott *et al.* 1995), presmoking arousal level (Shikata *et al.* 1995), hemisphere dominance (Domino *et al.* 1995a) and personality (Tatsuno 1995). In non-smokers, oscillatory changes due to acute nicotine administration are limited primarily to a rhythms, with increases observed in both PAF (Foulds *et al.* 1994; Harkrider *et al.* 2001) and frontal upper frequency  $a_2$  power (Fisher *et al.* 2012a) during resting states, and increases in anterior  $a_2$  during working memory tasks (Fisher *et al.* 2012b, 2013). Nicotine-induced oscillatory response differences

between smokers and non-smokers may reflect genetic factors, including COMT (Beuten *et al.* 2006; Colilla *et al.* 2005; Guo *et al.* 2007), involved in smoking initiation and progression to dependence (Kendler *et al.* 1999; Maes *et al.* 2004) or individual differences in EEG. Little is known about the genetics underlying EEG traits or pharmacologically modulated EEG, but twin studies show that heritability of resting EEG oscillations is substantial (Stassen *et al.* 1987), particularly for PAF (Posthuma *et al.* 2001; Smit *et al.* 2005, 2006) and *a*-band oscillations with 80–90% heritability estimates (Van Beijsterveldt & van Baal 2002; Van Beijsterveldt *et al.* 1996). The COMT polymorphism contributes to individual differences in brain *a* oscillations, with Val homozygotes exhibiting reduced *a2* and PAF compared to Met/Met carriers (Bodenmann *et al.* 2009a; Enoch *et al.* 2003), who exhibited greater delta, theta and beta (Solis-Ortiz *et al.* 2015).

We have examined the COMT polymorphism's moderating effects on the EEG oscillatory response to acute nicotine administration in non-smokers with upper alpha ( $\alpha_2$ ) and PAF as primary endpoints. Assuming that DA neuro-transmission innervates the  $\alpha$  oscillatory component of the nicotine-modulated EEG response, and that COMT impacts prefrontal cortical DA signaling, we hypothesized that a single dose of nicotine to non-smokers would act as a pharmacologic probe of dopaminergic tone, enhancing  $a_2$  and accelerating PAF, with the strongest effects in Met/Met individuals with higher levels of cortical dopamine and the weakest effects in Val/Val individuals. These same oscillatory changes should also differentiate COMT polymorphisms per se, with Met/Met homozygotes registering greater a<sub>2</sub> power and a higher PAF than Val/Val homozygotes that exhibit higher COMT activity. Secondary study endpoints included low frequency ( $\delta$ ,  $\theta$ ,  $\alpha_1$ ) and  $\beta$  oscillations as they are consistently modulated by smoking/nicotine (Knott 2001). Given the increasing attention of cortical oscillatory synchrony in the  $\gamma$  frequency range and its association with cognitive processes (Basar 2013; Herrmann et al. 2010; Merker 2013), we studied, for the first time in humans, resting-state  $\gamma$  oscillation response to nicotine and its moderation by the COMT polymorphism.

# **Methods**

# Study participants

The sample of volunteers consisted of 62 right-handed, healthy, non-smoking males between 18 and 34 years of age (mean age = 22.4 years) who were recruited primarily from local universities. Male, and not female, volunteers were chosen to avoid any potential confounding effects of menstrually related hormonal changes on nicotine response. All were screened for medical history, personal psychiatric history using the structured Clinical Interview, Non-Patient version for DSM-IV (SCID-NP; First *et al.* 2002) and family psychiatric history (first-degree biological relatives) with the Family Interview for Genetic Studies (FIGS; Maxwell 1992). Volunteers were included in the study if they were Caucasian, reported no personal or family psychiatric history including substance/alcohol abuse or dependence and had no significant medical issues and were medication free. Non-smokers were defined as those who had consumed no more than 100 cigarettes in their lifetime and had not smoked a cigarette over the past year. Non-smoking status was confirmed by expired carbon monoxide levels, which were <3 parts per million, a level

consistent with that of non-smokers (Cropsey *et al.* 2006). All volunteers signed a consent form prior to participation in the study, which was approved by the Research Ethics Board of the Royal Ottawa Health Care Group. Each participant received \$60 CAD for his involvement in the study.

# **Experimental design**

Each participant was assessed in two test sessions within a randomized, double-blind, placebo-controlled design. The two test sessions, involving nicotine or placebo treatment, were counterbalanced so that half of the participants received nicotine in their first session and placebo in their second session, while the remaining half received treatments in the reverse order. A minimum 2-day interval separated tested sessions.

## **Testing procedures**

Testing was carried out between 0900 and 1630 h, with participants being required to abstain from caffeine, alcohol, drugs and medication and food for a minimum of 8 h prior to their scheduled testing, and abstain from liquids (with the exception of water) for 2 h prior in order to avoid interference with nicotine absorption. Sessions were carried out in a dimly lit, sound-attenuated chamber situated adjacent to the control room housing the monitoring and testing computers. Nicotine/placebo was administered concurrently with EEG electrode hook-up while participants were seated in a large semi-reclining chair.

Electroencephalogram was recorded 30 min after nicotine/placebo administration, the time for nicotine to reach peak level in the blood. Vital signs and adverse events were assessed before and after nicotine administration.

## **Nicotine administration**

Oral administration of nicotine was in the form of two pieces (4 mg + 2 mg) of cinnamon-flavored Nicorette® gum (Johnson & Johnson Inc., Markham, Ontario, Canada).

Administering a 6 mg dose was intended to result in a similar nicotine level as achieved by smokers smoking a single cigarette of average nicotine yield, producing a nicotine blood concentration of approximately 15–30 ng/ml (Hukkanen *et al.* 2005). Peak blood nicotine levels are achieved approximately 30 min after the beginning of the gum chewing, and the elimination half-life of nicotine is ~120 min. Gum chewing was in accordance with the manufacturer's guidelines, which specified a chewing time of 25 min, biting twice every minute (as cued by an audio recoding) and 'parking' the gum between teeth and cheeks between bites. Placebo gum pieces were cinnamon-flavored and were similar in size, color and texture. Participants were blindfolded and wore nose plugs throughout the gum administration in order to reduce any possible sensory differences between nicotine and placebo. Prior to removing the nose plug after the chewing period, participants chewed a mint-flavored gum for 1–2 min in order to remove any lingering taste differences.

## Electroencephalographic acquisition

Electroencephalograms were recorded during a vigilance-controlled, 3-min eyes-closed resting-state condition. Electrical activity was sampled from an electrode cap (Electro-Cap International, Eaton, OH, USA) that positioned  $Ag^+/Ag^+$   $CI^-$  electrodes at eight scalp sites;

frontal midline (Fz), left (F3) and right (F4); central midline (Pz), left (C3) and right (C4); midline parietal (Pz) and midline occipital (Oz). An electrode on the nose served as a reference, and an electrode positioned anterior to the Fz site was the ground. Additional electrodes were placed on the supraorbital and suborbital ridges of the right eye, and on the external canthus of both eyes to monitor vertical (VEOG) and horizontal (HEOG) electro-oculographic activity. Electrode impedances were kept below 5 k $\Omega$ . Electroencephalograms were acquired (500 Hz sampling rate) with the eight-channel BrainVision V-Amp® amplifier (bandpass filters set at 0.1–120 Hz) and BrainVision Recorder® software (v1.1, Brain Products, Gilching, Germany). Digital recordings were stored for later off-line analysis.

# Electroencephalographic processing

Electroencephalographic analysis was carried out with BrainVision Analyzer® software (Brain Products). This included bandpass filtering (0.1–70 Hz; 24 dB/octave roll-off), epoch segmentation (2000 ms), ocular correction (Gratton *et al.* 1983) and artifact rejection (excluding ocular-corrected EEG epochs with voltages exceeding  $\pm 100~\mu V$ ). For each test session, a minimum of 45 2-second artifact-free epochs were subjected to a fast fourier transform (FFT) algorithm (with a high-pass, autoregressive filter, weighted by a 5% cosine taper) for computation of absolute amplitude ( $\mu V$ ) at each scalp site for:  $\delta$  (0.5–5.5 Hz),  $\theta$  (6–8 Hz),  $\alpha_1$  (8.5–10 Hz),  $\alpha_2$  (10.5–12 Hz),  $\alpha$  total (8.5–12 Hz),  $\beta_1$  (12.5–18 Hz),  $\beta_2$  (18.5–20.5 Hz),  $\beta_3$  (21–30 Hz) and  $\gamma$  (40–60 Hz) frequency bands. Relative amplitude (%) was also computed for each band by expressing amplitude in each band as a percent of total amplitude across all bands. As with Bodenmann *et al.* (2009b), PAF was determined visually from each individual's average spectra and was defined as the frequency bin (0.5 Hz resolution) in the  $\alpha$  range with the maximal power (0.5 Hz resolution).

# **COMT** genotyping

A sample of each participant's saliva was collected using Oragene DNA Self-Collection Kits (DNA Genotek Inc., Ottawa, Ontario, Canada). The genetic analysis was blinded to the results and provided by an external lab (Dr Paul Albert, Ottawa Hospital Research Institute). Extracted genomic DNA was assessed by real-time polymerase chain reaction (PCR) (Rotor-Gene RG-3000) to determine allele frequencies of the COMT Val158Met polymorphism (rs#4680), using 0.1× Taqman Drug Metabolism Genotyping Assay Kit (Applied Biosystems, USA, Assay ID# C\_25746809\_50) and template DNA with 1× Taqman master mix (4304437). The Rotor-gene 3000 (Corbett Research) real-time PCR apparatus was used with PCR cycling parameters, which included an initial 10-min denaturation at 95°C, 40 cycles of denaturation (15 seconds at 92°C) and annealing/extension (90 seconds at 60°C).

# Statistical analysis

Statistical analysis of the data was carried out with the Statistical Package for Social Sciences (SPSS Inc., Chicago, IL, USA). Mixed analysis of variance (ANOVA) was used to analyze each band, with separate ANOVAs for log-transformed absolute and relative power in each band consisting of two within-subject factors – treatment (two levels: nicotine and placebo) and scalp region (eight levels: Fz, F3, F4, Cz, C3, C4, Pz and Oz), and one between-subject factor, genotype (three levels: Val/Val, Val/Met and Met/Met curriers). Similar ANOVAs were run for PAF but as peak spectral frequency was less apparent at

lateral hemisphere recordings, the site factor in the ANOVA contained only four levels (Fz, Cz, Pz and Oz). Regardless of ANOVA significance for  $\alpha$ -band measures, planned comparison testing (correlated t-tests) study hypotheses were carried out and were mainly limited to placebo–nicotine contrasts within each genotype, as well as genotype contrasts in the placebo condition. For non- $\alpha$ -band measures, significant main or interaction effects (P< 0.05) were followed up with Bonferroni-corrected pairwise comparisons. Significant region effects were not examined in any of these comparisons unless they interacted with treatment and/or genotype.

## Results

# Allelic distribution

Catechol-O-methyltransferase genotyping resulted in the following distribution within the study sample: Met/Met = 24.19% (N= 15); Val/Met = 48.38% (N= 30) and Val/Val = 27.42% (N= 17). As evident with  $\chi^2$  statistics, no significant deviation from Hardy–Weinberg equilibrium was shown in the sample ( $\chi^2$  = 3.76, P= 0.05). No significant age differences were observed between genotypes.

# Alpha oscillations

Analysis of absolute amplitude in total a (F= 183.16, df = 7/413, P< 0.0001) and  $a_1$  bands (F= 121.12, df = 7/413, P< 0.0001) yielded significant effects for region but not for treatment, genotype or their interaction. Upper alpha ( $a_2$ ) analysis resulted in significant treatment (F= 6.75, df = 1/2, P< 0.015), region (F= 160.15, df = 7/413, P< 0.0001) and treatment × region × genotype interaction effects (F= 2.93, df = 14/143, P< 0.020). Genotypes did not differ with respect to absolute  $a_2$  amplitude during the placebo session but treatment comparisons within each genotype group (Fig. 1) found increases in absolute  $a_2$  amplitude with nicotine (vs. placebo) in Met/Met carriers at left (F3: P< 0.05), right (F4: P< 0.03) and mid-frontal (Fz: P< 0.015) regions as well as at left (F3: F< 0.03), right (F4: F< 0.03) and mid-central (F3: F< 0.02) regions. In carriers of the Val/Met allele, nicotine-induced absolute  $a_2$  amplitude increases were limited to mid-parietal (F2: F< 0.01) and mid-occipital (F3: F< 0.003) recording regions.

Relative amplitude analysis for total (F= 66.23, df = 7/413, P< 0.0001) and lower ( $a_1$ ) alpha (F= 42.34, df = 7/413, P< 0.0001) showed only significant region effects, whereas relative upper alpha ( $a_2$ ) amplitude was significantly affected by treatment (F= 3.86, df = 1/2, P< 0.05), region (F= 68.95, df = 7/413, P< 0.0001) and a treatment × region × genotype interaction (F= 2.44, df = 14/413, P< 0.05). Similar to the effects shown with absolute amplitude, no significant genotype differences were observed in the placebo session but in the Met/Met carriers, nicotine increased amplitude in all frontal (F3: P< 0.03; F4: P< 0.04; Fz: P< 0.02) and central scalp regions (F3: F3: F4: F5: F5: F8.003 but not at posterior regions.

The PAF analysis resulted in significant treatment (F = 5.92, df = 1/2, P < 0.02), region (F = 18.59, df = 73/197, P < 0.0001) and treatment × genotype (F = 3.41, df = 2/59, P < 0.04) interaction effects. The PAF was shown to be progressively higher from frontal to posterior

regions and in the placebo session, Val/Met carriers displayed the slowest PAF at Fz, which was significantly different from PAF of Val/Val (P< 0.04) but not Met/Met carriers (Fig. 2). Treatment comparisons within genotypes found significant PAF acceleration in the Val/Met carriers with nicotine (vs. placebo) at frontal (Fz: P< 0.01), central (Cz: P< 0.02) and occipital (Oz: P< 0.01) regions, and at the parietal (Pz: P< 0.05) region in Met/Met carriers (Fig. 3).

## Non-alpha oscillations

Absolute amplitude analysis yielded significant region effects for  $\delta$  (F= 80.56, df = 7/413, P < 0.0001),  $\theta$  (F= 43.54, df = 7/413, P< 0.0001),  $\beta$ <sub>1</sub> (F= 55.51, df = 7/413, P< 0.0001),  $\beta$ <sub>2</sub> (F= 15.72, df = 7/413, P< 0.0001),  $\beta$ <sub>3</sub> (F= 12.72, df = 7/413, P< 0.0001) and  $\gamma$  (F= 6.29, df = 7/413, P< 0.0001) but no treatment, genotype or interaction effects were observed.

Similar region effects were found for relative amplitude in  $\delta$  (F= 9.36, df = 7/413, P< 0.0001),  $\theta$  (F= 10.92, df = 7/413, P< 0.0001),  $\beta$ <sub>1</sub> (F= 5.53, df = 7/413, P< 0.0001),  $\beta$ <sub>2</sub> (F= 6.12, df = 7/413, P< 0.0001),  $\beta$ <sub>3</sub> (F= 69.68, df = 7/413, P< 0.0001) and  $\delta$  (F= 63.76, df = 7/413, P< 0.0001) bands, which were not affected by treatment, genotype or their interaction.

## Discussion

Our results indicate that the COMT Val158Met polymorphism contributes to individual differences in resting-state EEG oscillations and their response to acute nicotine. This points to a heritable dopaminergic mechanism in electrocerebral rhythmic activities that may reflect the cognitive response variability to nicotinic agonists. Earlier EEG research showed that in non-smokers only resting- and task-associated oscillations in the  $\alpha$  band varied with COMT polymorphism (Bodenmann et al. 2009a) and were sensitive to nicotine (Fisher et al. 2012a, 2012b, 2013; Foulds et al. 1994; Harkrider et al. 2001). Our EEG findings were limited to PAF and  $a_2$  oscillations, thus linking dopamine signaling and COMT variation with amediated oscillatory functions, possibly supporting cognitive enhancement associated with nicotine. Although cognitive processing was not directly assessed in this study, the case that resting neural changes affected by COMT polymorphism and nicotine interaction may meaningfully influence cognition is based on studies showing significant correlations between EEG during rest (pretask baseline) and behavioral performance. With respect to alpha, resting oscillatory activity in this band has detected both trait and state differences in cognitive functioning (Angelakis et al. 2004a,2004b) and interindividual differences in resting-state PAF and alpha power have been shown to be linearly associated with behavioral performance in healthy (Zhou et al. 2012; Zunini et al. 2013) and pathological populations (Dubbelink et al. 2013; Dubovik et al. 2012, 2013; Velikova et al. 2011).

Nicotine effects on EEG oscillations were evident only in the  $\alpha$ -frequency range and, consistent with our study hypothesis and previous non-smoker studies, PAF was accelerated and amplitude in  $\alpha_2$  was enhanced with single-dose treatment. A sizable body of research on the functional roles of brain oscillations has implicated  $\alpha$  in sensory processing and cognition (Basar 2012; Basar & Guntekin 2012), and during working memory (Roux & Uhlhaas 2014), where it may play a role in maintaining information through active inhibition

of task-irrelevant information (Klimesch *et al.* 2007; Sauseng 2009). Acetylcholine is thought to modulate the efficiency of the cortical processing of sensory or associational input (Basar & Guntekin 2012; Sarter *et al.* 2005); damage to the basal forebrain, the main source of acetylcholine, reduces resting EEG  $\alpha$ , which also diminishes in a number of cognitive disorders (schizophrenia and Alzheimer's disease) which, as our lab demonstrated, exhibit  $\alpha$  increases with smoking (Knott *et al.* 1995) and nicotine (Knott *et al.* 2000).

This is, to our knowledge, the first time that nicotine influences on resting EEG have been shown to be modulated by COMT polymorphism. Because these effects were assessed in non-smokers, they cannot be attributed to a 'remediation' of withdrawal-induced alterations in neural networks observed in smokers during abstinence (Ashare et al. 2014; Beaver et al. 2011; Cole et al. 2010). The Val158Met polymorphism of COMT was shown to modulate the resting-state response to nicotine, with  $\alpha$ -band oscillatory changes confirming our hypothesis, namely that such modulations occurred with carriers of the Met allele and not Val/Val. For  $\alpha_2$ , nicotine was found to increase oscillatory power in both Met/Met and Val/Met carriers but in different brain regions, with Met/Met in the frontocentral and Val/Met in the posterior cortical region. These COMT polymorphisms also exhibited PAF acceleration following nicotine administration but this time with Met/Met showing PAF increments in the posterior and Met/Val in the frontocentral regions. Because faster PAF and greater upper- $\alpha$  activity are linked to greater cognitive and memory performance (Klimesch 1997, 1999), altered PAF and  $\alpha_2$  oscillatory states induced by nicotine might be expected to preferentially enhance associated cognitive processes, presumably by increasing dopamine signaling in a regionally specific manner.

In line with these resting-state findings, previous fMRI studies on acute nicotine dosing in healthy volunteers (Herman et al. 2013), dopamimetics (Hamidovic et al. 2010; Mattay et al. 2003) and COMT inhibitors (Apud et al. 2007; Farrell et al. 2012; Giakoumaki et al. 2008) generally support the inverted-U model of dopamine function. By enhancing PFC dopamine in the higher COMT activity of homozygotes (Val/Val, positioned left of Met genotypes on the curve) these drugs shifted Val genotypes rightward, closer to the increased subjective responses as well as enhanced cortical efficiency and performance, while reducing cortical processing in Met/Met genotypes, presumably beyond the peak of the inverted-U. The actions of DA in the PFC are concentration- and receptor subtype-dependent, and while a balance is assumed between D1 and D2 dopamine receptor activation during resting states, working and memory tasks produce low to moderate increases in dopamine (Phillips et al. 2004) and D1 dopamine receptor activation (thought to enhance PFC glutamatergic and GABAergic currents) (Seamans et al. 1998). This is in contrast to high phasic dopamine levels, which are thought to activate D2 receptors and reduce these currents (Seamans & Yang 2004). Interestingly, in our study with smokers, smoking-induced  $\beta_2$  increases appear to be mediated by D2 receptor activation as they are blocked by haloperidol, a D2 receptor antagonist (Walker et al. 2001).

COMT studies involving fMRI have typically assessed Blood-oxygen-level dependent (BOLD) activation during executive (e.g. working memory) task performance. There is increasing evidence, however, that the resting-state activity and deactivation of these neural networks, such as the default mode network (DMN) (regions that are active during non-task

conditions and are suppressed by goal-directed cognitive demands; Raichle *et al.* 2001), determine the ability of task-positive networks to perform tasks, as measured by task-related fMRI (De Luca *et al.* 2006) and vary across individuals to predict behavioral performance (Kelly *et al.* 2008). There are a number of studies of dopaminergic modulation of task-induced changes in the DMN with transient dopamine depletion (Carbonell *et al.* 2014; Nagano-Saito *et al.* 2008), administration of dopamine receptor agonists and antagonists (Cole *et al.* 2013), levodopa (Delaveau *et al.* 2010), apomorphine (Nagano-Saito *et al.* 2009) and with pharmacological blockade (Minzenberg *et al.* 2011) and natural variation in dopamine transporter binding (Tomasi *et al.* 2009). All of these suggest that higher dopamine transmission is associated with augmented task-induced DMN deactivation. The COMT gene variation has similarly affected the fMRI–DMN response (Liu *et al.* 2010) with *intermediate* levels of COMT activity associated with increased medial PFC connectivity, which in turn correlates with increased task-induced DMN deactivation (reduced BOLD) and better performance (Dang *et al.* 2013).

Although there are inconsistencies (Neuner *et al.* 2014), both EEG (Cannon & Baldwin 2012; Chen *et al.* 2008, 2013; Knyazev 2013; Knyazev *et al.* 2011) and simultaneous EEG–fMRI studies (Chang *et al.* 2013; Jann *et al.* 2009; Liu *et al.*, 2014; Mo *et al.* 2013) suggest that alpha and upper a, in particular (Jann *et al.* 2009), positively correlate with the fMRI–DMN. As ongoing oscillatory alpha activity is associated with processing internal stimuli and can determine stimulus detection and attention (Hanslmayr *et al.* 2011; Klimesch 1999, 2012; Klimesch *et al.* 2007), the apparent nicotine-induced enhancement of the EEG–DMN network (evidenced by  $a_2$  increases in frontocentral regions in Met/Met carriers and in parietal/occipital regions in Val/Met carriers) may reflect the differential effects of dopamine neurotransmission on inhibitory and selection processes in anterior and posterior hubs of the DMN. Both of these are sensitive to individual differences (Knyazev *et al.* 2012) but appear to exhibit distinct relationships with dopamine transporter availability (Tomasi *et al.* 2009) and levodopa treatment (Asanuma *et al.* 2006).

Genotype differences in the placebo condition were expressed only with PAF, which, contrary to our hypothesis, was found to be faster in Val homozygotes than heterozygotes. This is in direct contrast to a previous report of a slower PAF in Val/Val (vs. Met/Met) genotype (Bodenmann et al. 2009a) and another showing no effect of COMT polymorphism on PAF (Veth et al. 2014). Although this may be due to differences in EEG methodology or PAF measurement techniques (Bazanova & Vernon 2014), COMT-PAF relationships require further study as this electrophysiological index is thought to reflect the influence of individual genes on the underlying neural mechanisms generating  $\alpha$  activity (Hughes et al. 2011; Lopes da Silva 1991; Steriade & Timofev 2003; Steriade et al. 1990). The behavioral significance of PAF is not clear but does provide a mechanism for searching and identifying encoded information (Angelakis et al. 2007; Bazanova & Aftanas 2006, 2008; Bazanova & Vernon 2014; Bodenmann et al. 2009b; Klimesch et al. 1993; Zoefel et al. 2011). Posterior PAF increases with increasing cognitive demands (Haegens et al. 2014) and artificially induced increases in a power above one's PAF have resulted in improved cognitive performance (Hanslmayr et al. 2005; Klimesch et al. 2003). Interestingly, given the impaired executive performance in Val allele carriers, performance and brain activation are more efficient and emotional (anxiety) disorders less prevalent in this same genetic group

compared to Met allele carriers (Heinz & Smolka 2006; Mier *et al.* 2010). The negative emotionality associated with low COMT activity (high cortical/subcortical tonic dopamine) in the Met allele may be related to the inflexibility of neural networks in processing information related to emotion, which in the Val allele is more flexible due to decreased tonic dopamine cortically and subcortically (Bilder *et al.* 2004). Alpha oscillations are particularly sensitive to the processing of negative emotional stimuli (Guntekin & Basar 2007) and as PAF is reduced in patients with anxiety disorder and increased with treatment (Saunders *et al.* 2015), the faster PAF in the Val carriers may be a neural marker of the dopaminergic regulation of the stability/flexibility of brain networks related to emotional processing.

#### Limitations

The present findings on COMT-nicotine interactions with EEG need to be interpreted within the limitations of the study, which include the assessment of a relatively small sample and one that was all male and was nicotine naive, thus limiting generalization to smokers – individuals who comprise a proportion of the population with psychiatric disorders (e.g. schizophrenia) and are purported to use smoking/nicotine for their cognitive enhancing properties (Kumari & Postma 2005). Dose- and time-response effects were not examined with nicotine and the neural effects associated with the relatively slow absorption of nicotine via gum may be dissimilar from those obtained with the rapid nicotine delivery associated with cigarette smoking. Blood nicotine levels were not assessed and as  $a_2$  increments with smoking are not observed with blood levels of nicotine below 10 ng/ml (Domino et al. 1995b), it is uncertain whether our observed genotype differences reflect different nicotine absorption levels. Our EEG recording montage, containing only eight recording sites, did not allow for a comprehensive assessment of electrocerebral activity and the use of larger electrode arrays in future studies would not only permit the localization of sources contributing to COMT and nicotinic influences on scalp EEG activity but also on interregional connectivity, which has been shown to be modulated by COMT polymorphism (Lee et al. 2011). Oscillatory measures included power and frequency but not phase characteristics, which, for a rhythms, is associated with unique sensory and cognitive processes (Bazanova & Vernon 2014). Electroencephalogram was assessed only during rest and not during task engagement which, if incorporated in future studies, would allow for clearer understanding of the cognitive implications of the COMT-modulated resting electrocortical response to nicotine.

# Conclusion

This study tentatively suggests that COMT polymorphism, nicotine and their interaction affect resting-state electro-cerebral rhythms. That the nicotine-induced electrocortical changes were evident in individuals with the COMT Met allele infers that the acute, nicotine-modulated spontaneous oscillations reflect tonic cortical dopamine level and its potential functional role in cognitive tasks. These spectral EEG findings were observed with respect to  $\alpha$  oscillatory activity, which is associated with cognitive processes and therefore may be of relevance to people diagnosed with schizophrenia – a disorder invariably linked with COMT polymorphism (Gupta *et al.* 2011; Lewandowski 2007; Sagud *et al.* 2010;

Williams *et al.* 2007), and is also associated with excessive smoking (Winterer 2010), aberrant dopamine neuro-transmission (Brisch *et al.* 2014), abnormalities in  $\alpha$  rhythms (Knott *et al.* 2001; Narayanan *et al.* 2014; Venables *et al.* 2009; Wix-Ramos *et al.* 2014), fMRI– (Whitfield-Gabrieli *et al.* 2009) and electrophysiologic–DMN (Kim *et al.* 2014) and cognition (Nuechterlein *et al.* 2014), with the latter being shown to improve both with smoking and nicotine administration (D'Souza & Markou 2012). Schizophrenia patients have shown increases in  $\alpha_2$  with the smoking of a single cigarette (Knott *et al.* 1995), but the role of COMT in these oscillatory changes is not yet known and the cognitive changes accompanying these cortical rhythms require addressing.

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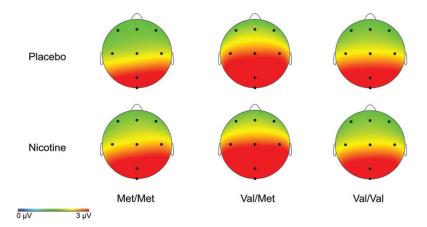


Figure 1. Grand averaged topographic EEG maps of  $a_2$  during placebo and nicotine treatment in Met/Met (M/M), Val/Met (V/M) and Val/Val (V/V).

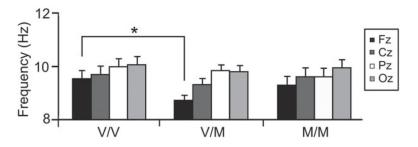


Figure 2. Mean ( $\pm$ SE) placebo peak alpha frequency in Val/Val (V/V), Val/Met (V/M) and Met/Met (M/M) genotypes at each electrode site.

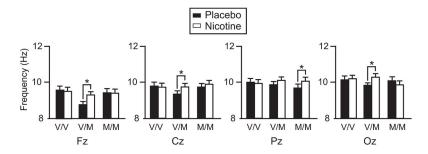


Figure 3. Mean ( $\pm$ SE) peak alpha frequency of placebo and nicotine in Val/Val (V/V), Val/Met (V/M) and Met/Met (M/M) carriers at each midline electrode site.