

# Individual differences in neural correlates of fear conditioning as a function of 5-HTTLPR and stressful life events

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**Fear learning is a crucial process in the pathogenesis of psychiatric disorders, which highlights the need to identify specific factors contributing to interindividual variation. We hypothesized variation in the serotonin transporter gene (5-HTTLPR) and stressful life events (SLEs) to be associated with neural correlates of fear conditioning in a sample of healthy male adults ( $n = 47$ ). Subjects were exposed to a differential fear conditioning paradigm after being preselected regarding 5-HTTLPR genotype and SLEs. Individual differences in brain activity as measured by functional magnetic resonance imaging (fMRI), skin conductance responses and preference ratings were assessed. We report significant variation in neural correlates of fear conditioning as a function of 5-HTTLPR genotype. Specifically, the conditioned stimulus ( $CS^+$ ) elicited elevated activity within the fear-network (amygdala, insula, thalamus, occipital cortex) in subjects carrying two copies of the 5-HTTLPR S' allele. Moreover, our results revealed preliminary evidence for a significant gene-by-environment interaction, such as homozygous carriers of the 5-HTTLPR S' allele with a history of SLEs demonstrated elevated reactivity to the  $CS^+$  in the occipital cortex and the insula. Our findings contribute to the current debate on 5-HTTLPR x SLEs interaction by investigating crucial alterations on an intermediate phenotype level which may convey an elevated vulnerability for the development of psychopathology.**

**Keywords:** imaging genetics; classical conditioning; fear; 5-HTTLPR; amygdala

## INTRODUCTION

Fear conditioning has been emphasized as a key process in the development of anxiety disorders (Shin and Liberzon, 2010). Thus, current research attempts to identify specific genetic and environmental factors that contribute to individual differences in conditioned fear responses. A well-established method to investigate variation in fear conditioning is the differential conditioning paradigm. This procedure involves pairing of a neutral conditioned stimulus ( $CS^+$ ) with a salient aversive unconditioned stimulus (UCS), while a second stimulus ( $CS^-$ ) is never paired with the UCS (non-UCS). After a few trials, the  $CS^+$  elicits conditioned fear responses (CRs) such as increased skin conductance responses (SCRs), changes in subjective ratings and elevated brain activity (Delgado *et al.*, 2006; De Houwer, 2009). Neural correlates of fear conditioning involve activation of a subcortical fear-network comprising the amygdala, the thalamus and the anterior cingulate cortex (ACC) (Hamm *et al.*, 2003; Delgado *et al.*, 2006; Sehlmeier *et al.*, 2009). The thalamus–amygdala pathway possesses a crucial role for the  $CS$ –UCS association process and mediates CRs (Öhman and Mineka, 2001). In addition, recent studies have identified an extended network including the occipital cortex and the insula, which is considered to convey the evaluation of the current  $CS$  value and the interoceptive processing of CRs (Sehlmeier *et al.*, 2009).

Results from twin studies suggesting CRs to be partly heritable (Hettema *et al.*, 2003) stimulated genetic association studies in this field. A functional polymorphism in the promotor region of the serotonin transporter gene (5-HTTLPR) has been considered a promising candidate, given that pharmacological manipulation of serotonergic neurotransmission relates to substantial alterations in fear conditioning (Inoue *et al.*, 2004; Burghardt *et al.*, 2007; Almada *et al.*, 2009). This 43 bp insertion/deletion polymorphism presumably influences transcriptional activity of the serotonin transporter (5-HTT) gene and comprises a low-expressing short (S allele) and a high-expressing long (L allele) variant (Lesch *et al.*, 1996; Stoltenberg *et al.*, 2002). Numerous neuroimaging studies using a variety of different aversive stimuli have suggested the S allele to be associated with increased reactivity of fear-relevant brain structures (Hariri *et al.*, 2002; Bertolino *et al.*, 2005; Hariri *et al.*, 2005; Munafò *et al.*, 2008), whereas neural correlates of fear conditioning have not yet been investigated. However, several studies have addressed the association between 5-HTTLPR and peripheral physiological measures of CR (Garpenstrand *et al.*, 2001; Crişan *et al.*, 2009; Lonsdorf *et al.*, 2009). For instance, in a well-conducted study, Crişan and colleagues (2009) found increased SCRs towards the  $CS^+$  in S allele compared to homozygous L allele carriers. In contrast, Lonsdorf *et al.* (2009) report increased startle responses in S allele compared to homozygous L allele carriers, whereas no differences in SCRs emerged. Following this line of research, genetic imaging studies are needed to elucidate the underlying neural mechanisms by which 5-HTTLPR genotype biases CRs.

In addition, substantial effort has been undertaken to identify specific combinations of genetic and environmental risk factors which jointly modulate vulnerability to psychiatric disorders. Regarding 5-HTTLPR, the S allele has been suggested to convey an increased risk for depression under conditions of elevated stress (Caspi *et al.*, 2003; Uher and McGuffin, 2008; Caspi *et al.*, 2010). Recently, this issue

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has attracted a considerable controversy raised by meta-analyses with conflicting results (Munafò *et al.*, 2008; Risch *et al.*, 2009; Karg *et al.*, 2011). This debate highlights the need to explore biological alterations associated with 5-HTTLPR x life stress (Caspi *et al.*, 2010), which can be discussed as a potential intermediate phenotypes conveying an elevated vulnerability. Following this line of research, prior studies have identified variation in neural and endocrine threat sensitivity as a function of 5-HTTLPR x stressful life events (SLEs). On a neural level, a significant 5-HTTLPR x SLEs interaction has been reported regarding resting state activity and in response to emotional face stimuli within fear-relevant brain structures (Canli *et al.*, 2006; Williams *et al.*, 2009; Lemogne *et al.*, 2011). Furthermore, we have observed significantly elevated cortisol responses to acute stress in subjects homozygous for the S allele with a history of SLEs (S'S'/high SLEs group) in a recently published study (Alexander *et al.*, 2009). The latter results imply that this specific constellation of genetic and environmental risk factors relates to biological alterations associated with elevated threat sensitivity. Thus, a gene-by-environment ( $G \times E$ ) approach might also contribute to a deeper understanding in genetic association studies on fear conditioning, but has not been applied in this field of research so far.

Based on the above mentioned literature, the present study aimed to investigate neural correlates of fear conditioning as a function of 5-HTTLPR genotype and SLEs. We hypothesized homozygous carriers of the low-expressing 5-HTTLPR alleles (S' group) to demonstrate exaggerated neural activity in the contrast  $CS^+ > CS^-$ . In addition, we assumed that the association of 5-HTTLPR and neural correlates of fear conditioning is further modulated by SLEs. Based on the observation of elevated endocrine threat/stress reactivity in S'S'/high SLEs subjects (Alexander *et al.*, 2009), we expected this group to demonstrate increased neural reactivity to the  $CS^+$ , which would indicate a cross-validation of our previous findings. In order to address these hypotheses, participants were preselected from our stress study regarding 5-HTTLPR genotype and SLEs. This preselection process resulted in a balanced number of participants for different GxE constellations, enhanced statistical power and further ensured sufficient variation regarding the prevalence of SLEs.

## METHODS

### Participants

Forty-eight subjects (mean age: 26.8; s.d.: 3.0) were recruited from our previously published stress study (Alexander *et al.*, 2009) comprising an ethnically homogenous Caucasian sample of 100 healthy male adults. Current or past mental (assessed by structured clinical interviews, Margraf, 1994), chronic physical problems and consumption of psychotropic drugs or drugs exerting influence on endocrine stress reactivity were defined as exclusion criteria. All subjects were right-handed, had normal or corrected-to-normal vision, and received 40 Euro for their participation. Participants signed an informed consent. The study was conducted in accordance with the Declaration of Helsinki and was approved by the ethics committee of the German Psychological Society.

Participants of the present fMRI study were recruited depending on the 5-HTTLPR/rs25531 mini-haplotype and SLEs. Regarding 5-HTTLPR, it has become increasingly common in genetic association studies to additionally account for an A→G single-nucleotide polymorphism (rs25531), which is located upstream of the 5-HTTLPR promoter variant within the greater repeat structure (Nakamura *et al.*, 2000). Prior studies suggested that the  $L_G$  allele is associated with a similar reduction in 5-HTT expression as the S allele (Hu *et al.*, 2006; Praschak-Rieder *et al.*, 2007). For enhanced clarity, the terms S' (S,  $L_G$ ) and the L' ( $L_A$ ) allele are used in the following when

referring to tri-allelic classification of 5-HTTLPR. In previous studies, dominant, co-dominant and recessive models of the 5-HTTLPR S' allele have been proposed without a clear consensus (Uher and McGuffin, 2008). Since we aimed to explore multiple threat-related biological alterations within the same sample, we based our selection process on a recessive model of the S' allele which has provided the best fit in our initial stress study (Alexander *et al.*, 2009) and all other previous studies on endocrine stress reactivity (Gotlib *et al.*, 2008; Dougherty *et al.*, 2010; Mueller *et al.*, 2010; Way and Taylor, 2010). Thus, we preselected 24 subjects homozygous for the low-expressing alleles (S'S' group) and 24 subjects carrying at least one high-expressing allele (L' group) to participate in the present study. Both the S'S' and the L' group comprised 12 subjects with a high number and 12 subjects reporting a low number of SLEs (as defined by median split within the original sample). Due to technical problems, data of one participant (L'/low SLEs) had to be excluded from the analyses.

### Assessment of stressful life events

Information on SLEs was obtained using the Life Events Checklist (LEC) developed by the National Centre for Posttraumatic Stress Disorder (PTSD). This 17-item self-report measure showed good psychometric properties in a sample of undergraduate students and has been associated with PTSD symptoms in a clinical sample (Gray *et al.*, 2004). As a unique feature, the LEC provides information on multiple types of exposure to a wide variety of potentially traumatic experiences (e.g. physical and sexual assault, combat, sudden, unexpected death of a loved one). Subjects respond to the items using a 5-point nominal scale (1 = happened to me, 2 = witnessed it, 3 = learned about it, 4 = not sure; 5 = does not apply).

### Conditioned stimuli (CS)

Two neutral visual stimuli (two squares; one with continuous lines; one with broken lines) served as  $CS^+$  and  $CS^-$ . All stimuli were grey in colour, had identical luminance, and were presented in an  $800 \times 600$  pixel resolution. The stimuli were projected onto a screen at the end of the scanner (visual field =  $18^\circ$ ) using an LCD projector. The two stimuli were counterbalanced as  $CS^+$  across participants. Pictures were viewed through a mirror mounted on the head coil.

### Unconditioned stimuli (UCS)

For the acquisition phase, a set of 20 highly aversive pictures (e.g. mutilations) were presented as UCS. These pictures were successfully used in previous studies, have been rated as highly aversive and have proven to elicit CRs (Libkuman *et al.*, 2007; Klucken *et al.*, 2009a). All pictures were presented in colour and had identical pixel resolution.

### Conditioning procedure

The conditioning procedure contained an acquisition and a short extinction phase. Only data from the acquisition phase is discussed here. Subjects were instructed to pay attention to all stimuli and to figure out a possible contingency between the CS and the UCS (e.g. Schiller *et al.*, 2008; Schiller *et al.*, 2010). Prior to the experiment and immediately after the conditioning procedure, participants rated the  $CS^+$ ,  $CS^-$ , and UCS. Detailed methods and results of the ratings are presented in the Supplementary Data. The acquisition phase consisted of 40 trials (20 per CS). The CS duration was 8 s. The UCS appeared immediately after the  $CS^+$  (100% reinforcement) for 4 s. Two training trials were conducted but excluded from the analyses, since learning could not have yet occurred (Phelps *et al.*, 2004; Klucken *et al.*, 2012).

For each subject, a pseudo randomized stimulus order was used with the following restrictions: (1) no more than two straight presentations of the same CS, and (2) equal distribution of CS presentations within the first and the second half of the acquisition. In an equally distributed interval of 1–2 s after the UCS offset, participants had to react to a simple distractor task (Goldin *et al.*, 2008; see Schweckendiek *et al.*, 2011 for a detailed description). This procedure was chosen to (1) distract the attention from the aversive pictures during the inter trial interval (ITI) and (2) to enhance overall vigilance. No group differences occurred between the four groups regarding the distractor task (false response rates were below 5% in all four groups). The ITI ranged from 12.5 s to 15 s. Throughout the experiment an MRI-compatible video camera was used to control if subjects watched the stimuli.

### Skin conductance measuring

SCRs were sampled simultaneously with MR scans using Ag/AgCl electrodes filled with isotonic (0.05 M NaCl) electrolyte medium, placed hypothenar at the non-dominant (left) hand. SCRs were defined in three analysis windows (Prokasy and Ebel, 1967): the maximum response within the time window 1–4 s after the CS ( $CS^+$  or  $CS^-$ ) onset was counted as the first interval response (FIR), within the time windows 4–8 s as the second interval response (SIR), and within the time window 9–13 s as the unconditioned response (third interval response; TIR). Responses were only registered when the response amplitude was greater than 0.01  $\mu$ S. Three subjects (2  $S'S'/$ high SLEs, 1  $L'/$ high SLEs) did not show SCRs (no responses to the UCS) and were excluded from SCRs analyses. Statistical analyses were performed via analysis of variance (ANOVA) in a 2 (stimulus-type:  $CS^+$  vs  $CS^-$ )  $\times$  2 (5-HTTLPR genotype:  $S'S'$  vs  $L'L'$  group)  $\times$  2 (environment: high vs low number of SLEs) experimental design followed by *post hoc* tests in PASW 18 (SPSS Inc., Chicago, IL, USA).

### Magnetic resonance imaging

#### Imaging parameters

Functional and anatomical images were acquired with a 1.5 Tesla whole-body tomograph (Siemens Symphony with a quantum gradient system) with a standard head coil. Structural image acquisition consisted of 160 T1-weighted sagittal images (MPRage, 1 mm slice thickness). For functional images, a total of 505 images were registered using a T2\*-weighted gradient echo-planar imaging (EPI) sequence with 25 slices covering the whole brain (slice thickness = 5 mm; 1 mm gap; descending slice procedure; TR = 2.5 s; TE = 55 ms; flip angle = 90°; field of view 192  $\times$  192 mm; matrix size = 64  $\times$  64). The orientation of the axial slices was paralleled to the orbitofrontal cortex-bone transition in order to minimise susceptibility artefacts in prefrontal areas. Data were analysed using Statistical Parametric Mapping (SPM8, Wellcome Department of Cognitive Neurology, London UK; 2008) implemented in MATLAB 7.5 (Mathworks Inc., Sherborn, MA, USA).

#### Preprocessing and statistical analyses

Prior to all analyses, data were preprocessed. This included realignment and unwarping (b-spline interpolation), slice-time correction, coregistration of functional data to each participant's anatomical image, and normalization to the standard space of the Montreal Neurological Institute (MNI) brain. Spatial smoothing was executed with an isotropic three-dimensional Gaussian filter with a full width at half maximum of 9 mm to allow for corrected statistical inference. The modelled experimental conditions were  $CS^+$ ,  $CS^-$ , UCS, non-UCS (defined as the time window after  $CS^-$  presentation corresponding to the time window of UCS presentation after the  $CS^+$ ; e.g. Stark *et al.*, 2006; Tabbert *et al.*, 2011), the distractor task, and the button

presses, modelled as events. Regressors were convolved with a hemodynamic response function (hrf) in the general linear model (GLM). The six movement parameters of the rigid body transformation applied by the realignment procedure were introduced as covariates in the model. The voxel-based time series was filtered with a high pass filter (time constant = 128 s). We further conducted analyses to account for the potential problem of collinearity between the regressors, e.g. in order to ensure that brain activity to the  $CS^+$  can be reliably distinguished from brain activity to the UCS. Results revealed no substantial collinearity between regressors (absolute values of cosine of angles between  $CS^+$  and UCS were: .15; between  $CS^-$  and non-UCS: .14), comparable to other previous studies (e.g. Klucken *et al.*, 2009a).

On the first level of analysis, the following contrasts were analysed for each subject:  $CS^+ > CS^-$  and UCS > non-UCS. The contrasts were calculated for each subject and introduced as dependent variables in the group analysis (second level analysis). We decided to use full-factorial models in order to avoid potentially biased type I errors in second level analyses due to the use of pooled errors (Boik, 1981; Barcikowski and Robey, 1984). Hence, contrasts from the first level GLM were analysed by full-factorial ANOVAs using partitioned errors (Penny and Henson, 2007). The full-factorial models included the group factors 5-HTTLPR genotype ( $S'S'$  vs  $L'L'$ ) and SLEs (low vs high number of SLEs) implemented in SPM8. In detail, four groups were introduced in the ANOVA ( $S'S'/$ high SLEs,  $S'S'/$ low SLEs,  $L'L'/$ high SLEs,  $L'L'/$ low SLEs). Main effects of 5-HTTLPR genotype, SLEs and interaction effects were analysed for each contrast (e.g.  $CS^+ > CS^-$ ; UCS > non-UCS). Further, appropriate *post hoc* group comparisons were conducted to specify potential interaction effects. Whole-brain analyses were conducted for the contrast  $CS^+ > CS^-$  ( $P < 0.001$ , uncorrected). Regions of interest (ROI) analyses were performed using the small volume correction in SPM8 ( $P < 0.05$  family-wise-error (FWE) corrected;  $k > 5$  voxel). The amygdala constitutes a target region, given that fear potentiated startle responses have been closely linked to this structure (Hamm and Weike, 2005; Weike *et al.*, 2005). Further ROIs derive from recent neuroimaging studies suggesting the 5-HTTLPR genotype to be associated with functional and/or structural alterations in other fear-relevant brain regions, comprising the insula, the thalamus and the occipital cortex (Rao *et al.*, 2007; Munafò *et al.*, 2008; Lemogne *et al.*, 2011). The masks for the amygdala, the insula and the thalamus were taken from the 'Harvard-Oxford cortical and subcortical structural atlases' provided by the Harvard Centre for Morphometric Analysis (Fox and Lancaster, 1994; Nielsen and Hansen, 2002). Since no mask for the occipital cortex is available in the Harvard-Oxford cortical and subcortical structural atlases, the mask for the primary visual cortex was taken from the probabilistic cytoarchitectonic maps and functional imaging data of the SPM anatomy toolbox (Eickhoff *et al.*, 2005). Since the masks were based on a probabilistic approach, a 50% cut-off was used for defining the ROI.

#### Genotyping

DNA was extracted from buccal cells using a standard commercial extraction kit (High Pure PCR Template Preparation Kit; Roche, Mannheim, Germany) in a MagNA Pure LC System (Roche). Subjects were genotyped for the 5-HTTLPR (and rs25531) by means of polymerase chain reaction (PCR) and gel electrophoresis. A detailed protocol is provided elsewhere (Alexander *et al.*, 2009).

## RESULTS

### Group characteristics

Within the original sample (Alexander *et al.*, 2009), there was no significant deviation from Hardy-Weinberg-Equilibrium using diallelic

( $\chi^2_{(1)} = 0.16; P > 0.60$ ) or triallelic ( $\chi^2_{(3)} = 30; P > 0.90$ ) classification of 5-HTTLPR. Genotype frequencies in the preselected sample of the present fMRI study were S/S = 21, S/L<sub>G</sub> = 3, and L<sub>G</sub>/L<sub>G</sub> = 0 for the S'S' group and S/L<sub>A</sub> = 11, L<sub>A</sub>/L<sub>G</sub> = 1, L<sub>A</sub>/L<sub>A</sub> = 11 for the L' group. Due to the preselection, we achieved significant differences ( $t_{(45)} = 7.54; P < 0.001$ ) regarding the number of SLEs in the low (M = 2.3, s.d. = 1.3) and the high (M = 6.3, s.d. = 2.0) SLEs group. As intended, no significant differences between the S'S' and the L' group emerged regarding self-reported SLEs (S'S' group: M = 3.9; s.d. = 2.5; L' group: M = 4.6, s.d. = 2.9;  $t_{(45)} = 0.9; P > 0.05$ ). Furthermore, the four groups depending on 5-HTTLPR genotype and SLEs did not differ with respect to age ( $F(3,44) < 1; P > 0.05$ )

**UCS results**

**Skin conductance responses**

ANOVA revealed a significant main effect of UCS-type ( $F_{(1,41)} = 27.65; P < 0.001$ ), which was not modulated by 5-HTTLPR genotype, SLEs or 5-HTTLPR × SLEs interaction. *Post hoc* tests confirmed greater responses to the UCS as compared to the non-UCS both in the S'S' group as well as in the L' group (Figure 1).

**Hemodynamic responses**

The results showed a significant main effect of task (UCS responses) in all ROIs, including the amygdala, the insula, the thalamus and the occipital cortex. No main effects of 5-HTTLPR genotype and SLEs or 5-HTTLPR × SLEs interactions were observed in the contrast UCS > non-UCS (Table 1).

**CS results**

**Skin conductance responses**

We found a significant main effect of CS-type in the FIR ( $F_{(1,41)} = 15.70; P < 0.001$ ) and in the SIR ( $F_{(1,41)} = 20.77; P < 0.001$ ). *Post hoc* tests revealed higher responses towards the CS<sup>+</sup> as compared to the CS<sup>-</sup>, thus, providing evidence for successful fear conditioning in all groups. No main effect of 5-HTTLPR genotype, SLEs, or its interaction occurred with regard to conditioned SCRs (Figure 2).

**Hemodynamic responses**

**Main effect of task**

With respect to our regions of interest (all FWE-corrected), we found a significant main effect of task in the left thalamus ( $z = 3.33; x/y/z = -9/-12/12; P < 0.05$ ), the right thalamus ( $z = 3.49; x/y/z = 12/-6/12; P < 0.05$ ), and the right insula ( $z = 3.17; x/y/z = 36/18/0; P < 0.05$ ). Trends were found in the left ( $z = 3.39; x/y/z = -6/-78/12; P = 0.075$ ) and the right ( $z = 3.34; x/y/z = 27/-66/9; P = 0.082$ ) occipital cortex.

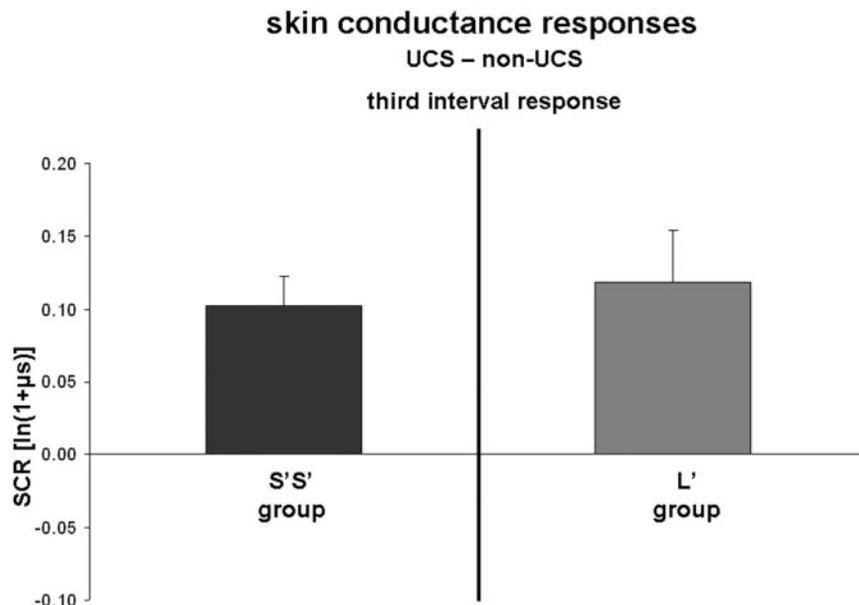
**Main effects of 5-HTTLPR genotype and SLEs**

Whole-brain analysis (all uncorr.) revealed CS<sup>+</sup>/CS<sup>-</sup> differentiation in the middle frontal gyrus ( $z = 3.91; P < 0.0001; x/y/z = -30/41/-15$ ) and in the left lateral orbitofrontal cortex ( $z = 3.52; P < 0.0001; x/y/z = -30/51/-15$ ) in the L' group. In the S'S' group we additionally

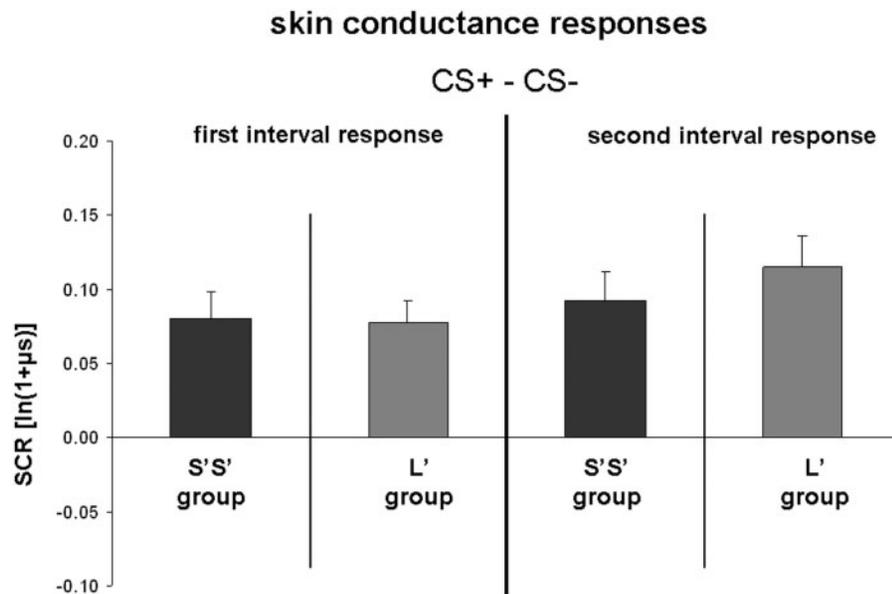
**Table 1** Neural activations for the main effect of UCS-type for the S'S' group and the L' group separately

Group	Structure	Side	x	y	z	Z <sub>max</sub>	P <sub>corr</sub>
S'S' group	Amygdala	L	-18	-6	-12	4.16	0.001
	Amygdala	R	21	-3	-15	3.37	0.010
	Insula	L	-33	21	0	4.60	<0.001
	Insula	R	33	24	-3	5.03	<0.001
	Thalamus	L	-9	-30	-3	5.06	<0.001
	Thalamus	R	6	-15	3	6.11	<0.001
	Occipital cortex	L	-12	-96	-3	6.52	<0.001
	Occipital cortex	R	21	-93	0	7.53	<0.001
	L' group	Amygdala	L	-21	-6	-12	4.36
Amygdala		R	27	0	-15	4.26	<0.001
Insula		L	-33	21	0	5.81	<0.001
Insula		R	33	24	-3	5.31	<0.001
Thalamus		L	-9	-30	-3	5.61	<0.001
Thalamus		R	21	-30	0	6.01	<0.001
Occipital cortex		L	-12	-99	-6	6.96	<0.001
Occipital cortex		R	18	-93	0	7.51	<0.001

The threshold was  $P < 0.05$  (FWE-corrected; small volume correction according to SPM8). All coordinates are given in MNI space. L: left hemisphere, R: right hemisphere.



**Fig. 1** Mean (± SE) conditioned skin conductance responses (UCS–non-UCS) for the S'S' group and the L' group in the TIR.



**Fig. 2** Mean ( $\pm$ SE) of the conditioned skin conductance responses ( $CS^+ - CS^-$ ) for the  $S'S'$  group and the  $L'$  group in the FIR and in the SIR.

**Table 2** Neural activations for the main effect of 5-HTTLPR genotype (upper part), and the 5-HTTLPR genotype  $\times$  SLEs interaction effects in the contrast  $CS^+ > CS^-$  (lower part)

Analyses	Structure	Side	x	y	z	$Z_{max}$	$P_{corr}$
Main effect	Amygdala	R	21	0	-18	3.26	0.015
5-HTTLPR	Insula	L	-39	-3	-12	3.15	0.044
genotype $S'S' > L'$	Insula	R	36	9	-15	4.04	0.002
	Thalamus	L	-9	-27	-3	3.65	0.015
	Occipital cortex	L	-12	-78	6	3.65	0.031
	Occipital cortex	R	21	-96	-3	3.77	0.025
5-HTTLPR	Insula	R	36	0	12	3.37	0.023
genotype $\times$ SLEs interaction	Occipital cortex	L	-12	-78	3	3.60	0.036

The threshold was  $P < 0.05$  (FWE-corrected; small volume correction; according to SPM8). All coordinates are given in MNI space. L: left hemisphere, R: right hemisphere.

found  $CS^+/CS^-$  differences in the left precentral gyrus ( $z = 6.4$ ;  $P < 0.0001$ ;  $x/y/z = -42/-3/30$ ) and in the cerebellum ( $z = 5.9$ ;  $P < 0.0001$ ;  $x/y/z = -27/-57/-33$ ). Further, the  $S'S'$  group showed significant ROI-activations (all FWE-corrected) in the right amygdala ( $z = 3.22$ ;  $x/y/z = 18/-6/-12$ ;  $P < 0.05$ ), the left insula ( $z = 3.33$ ;  $x/y/z = -39/-12/-3$ ;  $P < 0.05$ ), the right insula ( $z = 3.38$ ;  $x/y/z = 39/9/-15$ ;  $P < 0.05$ ), the left thalamus ( $z = 3.93$ ;  $x/y/z = -3/-24/3$ ;  $P < 0.01$ ), the right thalamus ( $z = 4.02$ ;  $x/y/z = 9/-18/0$ ;  $P < 0.01$ ), the left occipital cortex ( $z = 3.59$ ;  $x/y/z = -9/-81/6$ ;  $P < 0.05$ ), and the right occipital cortex ( $z = 3.73$ ;  $x/y/z = 18/-93/-3$ ;  $P < 0.05$ ).

We observed a significant main effect of 5-HTTLPR genotype in the contrast  $CS^+ > CS^-$  for each ROI. According to our a priori hypothesis, *post-hoc* tests confirmed significantly greater differentiation in the contrast  $CS^+ > CS^-$  in the  $S'S'$  group as compared to the  $L'$  group in the right amygdala, the insula (bilateral), the occipital cortex (bilateral) and the left thalamus (see Table 2; Figure 3 for details). No main effect for SLEs occurred.

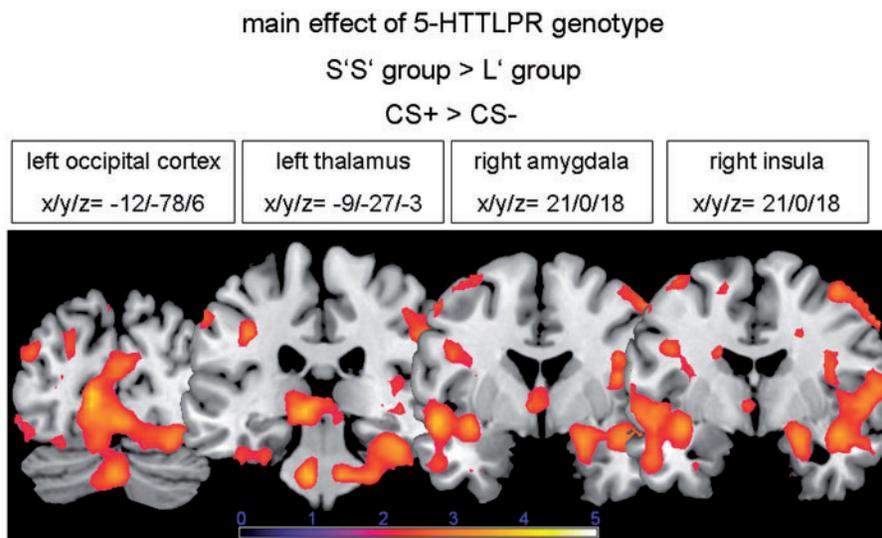
#### Gene $\times$ environment interaction

We observed a significant 5-HTTLPR genotype  $\times$  SLEs interaction effect in the right insula and the left occipital cortex for the contrast

$CS^+ > CS^-$  (Table 2, Figure 4). According to the a priori hypothesized direction of effects, *post hoc* testing revealed that the  $S'S'$ /high SLEs group showed significantly elevated neural responses in the left occipital cortex as compared to the  $L'$ /high SLEs ( $P = 0.016$ ) and the  $S'S'$ /low SLEs group ( $P = 0.019$ ); whereas no significant difference occurred compared to the  $L'$ /low SLEs group ( $P > 0.05$ ). Regarding the right insula activity, *post hoc* tests revealed a similar pattern with  $S'S'$ /high SLEs subjects showing highest activity. Specifically, elevated insula activity in the contrast  $CS^+ > CS^-$  reached significance when comparing the  $S'S'$ /high SLEs to the  $S'S'$ /low SLEs ( $P = 0.011$ ), to the  $L'$ /high SLEs group ( $P = 0.002$ ), and to the  $L'$ /low SLEs group ( $P = 0.018$ ). Pairwise comparisons did not reveal any significant differences between the remaining groups (all  $P > 0.05$ ).

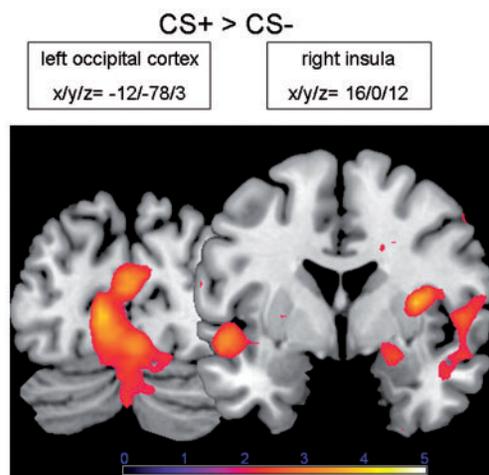
#### DISCUSSION

The present study is the first to investigate neural correlates of fear conditioning as a function of 5-HTTLPR genotype and SLEs. Our results indicate a significant association between 5-HTTLPR and BOLD-responses during fear conditioning in the right amygdala, the left thalamus, the bilateral insula and the bilateral occipital cortex. As hypothesized,  $S'S'$  subjects showed increased  $CS^+/CS^-$  differentiation within these fear-related structures compared to the  $L'$  group. Thus, the present study extends previous findings on peripheral physiological measures of CR by linking 5-HTTLPR genotype to individual differences in neural correlates of fear conditioning. Furthermore, our results point to a potential neural mechanism explaining why 5-HTTLPR genotype has been differentially associated with conditioned startle and skin conductance responses. In the study by Lonsdorf and colleagues (2009), S allele carriers showed increased startle potentiation, whereas consistent with our own findings, no such 5-HTTLPR genotype dependent differences appeared with regard to conditioned SCRs. A possible explanation for this selective effect is provided by studies suggesting conditioned startle and skin conductance responses to at least partly involve different neural circuits (Hamm and Weike, 2005; Tabbert et al., 2006). Whereas startle modulation is primarily mediated by connections from the amygdala to the brainstem (Davis and Whalen, 2001; Hamm and Weike, 2005), conditioned SCRs have



**Fig. 3** Neural activations for the main effect of 5-HTTLPR genotype in the contrast CS<sup>+</sup> > CS<sup>-</sup> in the region of interests (also see Table 2). For illustration reasons, the data were thresholded with a *T* 2.0 (see colour bar for exact *t*-values).

**5-HTTLPR genotype x environment  
interaction effect**



**Fig. 4** Neural activations for the 5-HTTLPR genotype × SLEs interaction effect in the contrast CS<sup>+</sup> > CS<sup>-</sup> in the region of interests (also see Table 2). For illustration reasons, the data were thresholded with a *T* 2.0 (see colour bar for exact *t*-values).

been reported to be largely dissociated from amygdala activity (Weike *et al.*, 2005; Tabbert *et al.*, 2006; Klucken *et al.*, 2009b). Thus, the observation of altered amygdala reactivity to the CS<sup>+</sup> as a function of 5-HTTLPR raises the notion that this genetic variant may predominantly modulate amygdala-dependent CR. However, it has to be acknowledged that associations between 5-HTTLPR and SCRs have been observed in other studies. For example, Crişan *et al.* (2009) conducted a conditioning paradigm (observational learning) and found increased SCRs towards the CS<sup>+</sup> in S allele carriers. These differences in design and statistical analyses (cf. Garpenstrand *et al.*, 2001) might explain the inconsistencies.

Moreover, the presented results may relate to the consistently reported finding of elevated amygdala reactivity to fearful stimuli in carriers of the 5-HTTLPR S allele (Hariri *et al.*, 2002; Hariri *et al.*, 2005; Munafò *et al.*, 2008). The ability to respond in a fast and adaptive

manner when confronted with an aversive stimulus has been suggested to be a key function of fear conditioning (Domjan, 2005). Thus, it is tempting to speculate that amygdala hyperreactivity observed as a function of 5-HTTLPR genotype may at least partly result from increased fear learning in the past. Even though this potential mechanism is supported by our data, only longitudinal studies will provide insights into the causal relationship between altered fear conditioning and processing in 5-HTTLPR S' allele carriers.

Furthermore, our study provides only preliminary evidence for a significant 5-HTTLPR genotype × SLEs interaction on neural correlates of fear conditioning, given that effects were not consistently observed across crucial structures of the fear-network. In accordance with the a priori hypothesized direction of effects, the S'S'/high SLEs group appeared to be most reactive, as indicated by increased neural activation to the CS<sup>+</sup> as compared to the CS<sup>-</sup> in the insula and the occipital cortex. However, the fact that SLEs were found to selectively moderate the association between 5-HTTLPR and reactivity to the CS<sup>+</sup> as compared to the CS<sup>-</sup> in specific brain areas seems unexpected at first and needs further discussion. Besides the amygdala, increased insula activation to the CS<sup>+</sup> is one of the major and most stable results reported in fMRI studies on fear conditioning (Sehlmeyer *et al.*, 2009). Notably, it has been suggested that the insula is not only involved in interoceptive bodily awareness, but also modulates the evaluation of future emotional states (Nitschke *et al.*, 2005; Paulus and Stein, 2006) by conveying a cortical representation of fear to the amygdala (Phelps *et al.*, 2001). Following this line of argumentation, it could be speculated that the effects of SLEs are most pronounced within brain areas crucially involved in fear representations, which have been acquired on the basis of (aversive) experiences in the past. A similar interpretation may apply for the observation of elevated neural reactivity to the CS<sup>+</sup> compared to the CS<sup>-</sup> within the occipital cortex in S'S'/high SLEs subjects. Enhanced occipital activation has not solely been observed during acute presentation of emotional material, but also occurs during anticipation of aversive stimuli and has often been referred to as increased motivated attention (Bradley *et al.*, 2003; Ueda *et al.*, 2003). These findings support the assumption that activation within the occipital cortex is not only stimulus driven but at least partly results from altered top-down processes, which may be influenced by environmental adversity in the past.

Moreover, it is interesting to note that S'S'/high SLEs subjects were not only characterized by an increased neural activity to the CS<sup>+</sup>, but were also found to exhibit markedly elevated cortisol reactivity to psychosocial stress in our previous study (Alexander *et al.*, 2009). Taken together, these findings suggest a broad network of neural and endocrine alterations associated with stress/threat sensitivity within subjects at high risk for psychopathology. Importantly, changes in cortisol reactivity and fear conditioning may very likely reflect interrelated biological processes, given the crucial role of glucocorticoids in fear learning (for review, see Rodrigues *et al.*, 2009). More precisely, several studies indicate stress exposure to significantly alter subsequent fear acquisition in a sex-specific manner. In males, previous exposure to a psychosocial stressor as well as elevated endogen release of glucocorticoids has been repeatedly associated with facilitated fear conditioning, while no such association or the opposite pattern has been observed in women (Jackson *et al.*, 2006; Zorawski *et al.*, 2006 but see Stark *et al.*, 2006; Merz *et al.*, 2010). The latter findings are thus consistent with the coexistent observation of elevated cortisol reactivity (Alexander *et al.*, 2009) and increased neural activation to the CS<sup>+</sup> observed within the present study in male S'S'/high SLEs subjects, but also imply that obtained effects may not be generalized to a female population.

Regarding potential clinical implications, our findings add to the discussion whether neural correlates of fear conditioning represent a potential intermediate process bridging the gap from 5-HTTLPR genotype to anxiety-related phenotypes. Elevated levels of anxiety-related personality traits in S allele carriers represent a well-established finding in genetic association studies on 5-HTTLPR, which has been confirmed by several independent meta-analyses (e.g. Munafò *et al.*, 2005; Munafò *et al.*, 2009). As suggested by our findings, enhanced neural reactivity in fear-related brain circuits can be discussed as a potential mechanism mediating this association. Indeed, a recent neuroimaging study reports trait anxiety to be positively correlated with the magnitude of amygdala reactivity to the CS<sup>+</sup> (Indovina *et al.*, 2011), once again highlighting the potential role of altered fear conditioning as a possible intermediate phenotype. Moreover, increased neural reactivity to the CS<sup>+</sup> as compared to the CS<sup>-</sup> in healthy S'S' subjects observed in our study may represent a premorbid risk factor. In line with this argumentation, a recent longitudinal study identified neuroticism to significantly mediate the relationship between genetic risk factors and symptoms of anxiety and depression (Kendler and Gardner, 2011). Taken together, altered neural responding to the CS<sup>+</sup> can be discussed as a possible mechanism conveying elevated levels of trait-anxiety in healthy S'S' subjects, thereby possibly increasing vulnerability to anxiety and affective disorders. Regarding the moderating role of SLEs, it is of note that evidence for significant 5-HTTLPR × SLEs interaction is not limited to depression susceptibility (Uher and McGuffin, 2008; Caspi *et al.*, 2010), but has also been reported with regard to anxiety-related personality traits (Stein *et al.*, 2008). Consistent with our findings, S'S'/high SLEs were found to exhibit elevated levels of anxiety sensitivity (AS) (Stein *et al.*, 2008), a well-established risk factor for the development of anxiety disorders (Lang *et al.*, 2002; Schmidt *et al.*, 1999). Again, elevated neural fear reactivity within this group suggested by our study may account for this association, given that increased insula activation during the processing of aversive stimuli was also found in subjects with elevated levels of AS (Stein *et al.*, 2007).

In conclusion, our study is the first to report significant variation in neural correlates of fear learning as a function of 5-HTTLPR and SLEs. While associations of 5-HTTLPR genotype and altered reactivity to the CS<sup>+</sup> as compared to the CS<sup>-</sup> robustly appeared in key structures of the fear network, evidence for a significant G×E interaction is much weaker. Given the limited sample size, our results remain preliminary

until independent replication is available, especially with respect to findings on 5-HTTLPR × SLEs interaction. Beside this limitation, our study significantly contributes to the current debate on the role of 5-HTTLPR and SLEs on psychiatric disorders by investigating alterations on an intermediate phenotype level, which may represent a premorbid risk factor that conveys elevated susceptibility for the development of psychiatric disorders.

## SUPPLEMENTARY DATA

Supplementary data are available at SCAN online.

## Conflict of Interest

None declared.

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