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Relationship of the serotonin transporter gene promoter polymorphism (5-HTTLPR) genotype and serotonin transporter binding to neural processing of negative emotional stimuli

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Abstract

Background—The lower-expressing (*S'*) alleles of the serotonin transporter (5-HTT) gene promoter polymorphism (5-HTTLPR) are linked to mood and anxiety related psychopathology. However, the specific neural mechanism through which these alleles may influence emotional and cognitive processing remains unknown. We examined the relationship between both 5-HTTLPR genotype and *in vivo* 5-HTT binding quantified *via* PET with amygdala reactivity to emotionally negative stimuli. We hypothesized that 5-HTT binding in both raphe nuclei (RN) and amygdala would be inversely correlated with amygdala reactivity, and that number of *S'* alleles would correlate positively with amygdala reactivity.

Methods—In medication-free patients with current major depressive disorder (MDD; *N* = 21), we determined 5-HTTLPR genotype, employed functional magnetic resonance imaging (fMRI) to examine amygdala responses to negative emotional stimuli, and used positron emission tomography with [¹¹C]DASB to examine 5-HTT binding.

Results—[¹¹C]DASB binding in RN and amygdala was inversely correlated with amygdala response to negative stimuli. 5-HTTLPR *S'* alleles were not associated with amygdala response to negative emotional stimuli.

Limitations—Primary limitations are small sample size and lack of control group.

Conclusions—Consistent with findings in healthy volunteers, 5-HTT binding is associated with amygdala reactivity to emotional stimuli in MDD. 5-HTT binding may be a stronger predictor of emotional processing in MDD as compared with 5-HTTLPR genotype.

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Keywords

Depression; Genetics; Functional MRI; Biological Markers; Brain Imaging

1. Introduction

Triallelic variation of the upstream, human serotonin transporter-promotor polymorphic region (5-HTTLPR) comprises one short (S) and two long (L_G and L_A) variants, of which the S and L_G variants (together designated as S') have comparably lower transcriptional efficiency *in vitro* compared with L_G (designated L') (Heils et al., 1996; Hu et al., 2006; Lesch et al., 1996; Mortensen et al., 1999). Low-expressing alleles are associated with elevated amygdala reactivity to threat (Hariri et al., 2002) and other negative stimuli (see Murphy et al. (2012) for a review), and are a risk factor for psychopathology such as major depressive disorder (MDD) in context of life stress (Caspi et al., 2003). However, the mechanism by which genotype at this locus confers risk for psychopathology remains unknown.

One hypothesis is that low-expressing 5-HTTLPR polymorphisms alter intra-synaptic serotonin levels, leading to increased amygdala activity (Fisher et al., 2009, 2006; Hariri and Holmes, 2006). Although 5-HTTLPR genotype affects 5-HTT mRNA expression *in vitro* (Heils et al., 1996; Hu et al., 2006; Lesch et al., 1996; Mortensen et al., 1999), *in vivo* studies using positron emission tomography (PET) imaging report both null findings (Guzey et al., 2012; Kobiella et al., 2011; Miller et al., 2013; Murthy et al., 2010; Oquendo et al., 2007; Parsey et al., 2006a; Rhodes et al., 2007; Shioe et al., 2003) and decreased 5-HTT binding associated with the S' allele (Kalbitzer et al., 2009; Praschak-Rieder et al., 2007; Reimold et al., 2007).

Although S' alleles are associated with greater amygdala responses to negative stimuli (reviewed in (Murphy et al., 2012)), less is known about the relationship between 5-HTT protein levels and amygdala reactivity. 5-HTT binding in amygdala is inversely correlated with amygdala responses to negative stimuli in healthy controls and depressed subjects (Rhodes et al., 2007; Ruhe et al., 2014). However, the same studies disagree regarding the relationship between raphe nuclei (RN) 5-HTT binding and amygdala activity. These discrepant findings justify further study of regional 5-HTT binding with respect to amygdala reactivity.

We sought to clarify relationships between 5-HTTLPR genotype, regional 5-HTT binding, and amygdala responses to negative stimuli in currently depressed MDD subjects. Based on the regulatory role of serotonergic projections to the amygdala (Jasinska et al., 2012; Jiang et al., 2009; Rainnie, 1999), we hypothesized that 5-HTT binding in both RN and amygdala would be inversely correlated with amygdala reactivity. Given that the S' allele is associated with lower transcriptional efficiency of 5-HTT *in vitro*, we predicted that S' allele number would be positively correlated with amygdala reactivity. This is the first study, to our knowledge, that combines these genetic and neuroimaging approaches in a study of MDD.

2. Methods

2.1. Sample

We studied a convenience sample of depressed subjects who had undergone a multimodal fMRI/PET imaging study including: (1) the negative emotional faces fMRI task employed in previous studies examining amygdala reactivity (Fisher et al., 2009, 2006; Hariri et al., 2002); (2) quantification of 5-HTT binding using PET with [¹¹C]DASB; and (3) genotyping of the 5-HTTLPR. A subset of these subjects had also undergone PET scanning with [¹¹C]WAY-100635 to quantify 5HT_{1A} receptor binding, allowing for accurate localization of raphe nuclei. The average time between MRI and PET scan was $M = 7$, $SD \pm 6$ days.

Subjects ($n = 21$) with current MDD based on the Structured Clinical Interview for DSM-IV, Axis I (SCID-I) (Spitzer et al., 1992) enrolled in this study. Complete eligibility criteria are enumerated in S1.

2.2. Genotyping

Genotype classification followed the method described in Parsey et al. (2006a). S or L_G alleles were classified as an S' allele and the L_A allele (which is higher-expressing *in vitro* (Hu et al., 2005)) was classified as the L' allele. Subjects were classified into 3 functional genotypes: L'L', L'S', S'S'. Correlations between S' allele and demographic variables were calculated in SPSS (SPSS Inc. Released 2007. SPSS for Windows, Version 16.0. Chicago, SPSS Inc) using Pearson and Spearman correlations for normally and non-normally distributed variables.

2.3. PET protocol

Details of the PET protocol have been described elsewhere (Ogden et al., 2007) and are included in Supplemental information.

2.4. MRI image acquisition

MRI scans were acquired on a 3T SignaHDx scanner (General Electric Medical Systems, Milwaukee, WI). Full acquisition parameters are described in S1. EPI acquisition was acquired with a TR = 2000 msec and voxel dimension = $3.75 \times 3.75 \times 5$ mm³.

2.5. fMRI task

The task was modeled after the task reported in Hariri et al. (2002). In this blocked ABAB design, subjects viewed a screen with three pictures of either angry/fearful faces or shapes in a triangular arrangement. They were instructed to determine which of the top two stimuli matches the bottom one. Stimuli were presented for four seconds, with inter-stimulus interval of two to six seconds. Shape stimuli were presented four times per block and face stimuli six times. Five face and four shapes blocks were presented.

2.6. MRI image processing

MRI Image processing is described in S1.

2.7. PET analysis

PET image processing is described further in S1. Because no brain region is devoid of specific binding with [^{11}C]DASB (Parsey et al., 2006b), we used an outcome measure that does not rely on a reference region: V_T/f_p where V_T = volume of distribution in the region of interest and f_p is the plasma free-fraction (Chin et al., 2011; Esterlis et al., 2010; Fujita et al., 2012; Ichise, 2009; Mukhin et al., 2008). [^{11}C]DASB regional V_T values were derived using likelihood estimation in the graphical approach (LEGA) (Ogden, 2003; Parsey et al., 2003). Brain activity was corrected for the contribution of plasma activity assuming a 5% blood volume in regions of interest (Mintun et al., 1984). To facilitate comparison to other [^{11}C]DASB studies using different outcome measures, the following outcome measures were also estimated: BP_F^* ($(V_{T(\text{ROI})} - V_{T(\text{REF})})/f_p$); BP_P^* ($V_{T(\text{ROI})} - V_{T(\text{REF})}$); and BP_{ND}^* ($(V_{T(\text{ROI})} - V_{T(\text{REF})})/V_{T(\text{REF})}$), using cerebellar gray matter as reference region. Asterisks are added to consensus terminology (Innis et al., 2007) to emphasize that the “reference” region does not have measureable specific binding (Parsey et al., 2006b), which violates the assumption underlying estimation of these alternative outcome measures. Time activity curves were generated by plotting measured activity within ROIs over the course of PET acquisition.

2.8. fMRI analyses

BOLD data were analyzed using the general linear model to identify voxel-wise parameter estimates to a faces-greater-than-shapes regressor, convolved with a double gamma hemodynamic response function. Activity during the instruction period was covaried as a nuisance regressor. Relative frame displacement for all subjects was less than half the width of a single voxel. Motion parameters were included as nuisance variables.

For all of the following analyses, the minimum Z-score required for significance was set at $z \geq 2.3$ (cluster $p < 0.05$).

2.9. Genotype and 5-HTT

To identify the relationship of 5-HTTLPR genotype to BOLD responses to angry/fearful faces in amygdala, we regressed parameter estimates of task-related activity in the amygdala ROI onto the number of S' alleles. We selected this approach given *in vitro* (Hu et al., 2006) and *in vivo* (Neumeister et al., 2006) evidence that number of triallelic HTTLPR L_A alleles exert a dose response on 5-HTT mRNA expression and downstream effects. Amygdala activity was regressed independently onto V_T/f_p in RN and amygdala. Mean response to faces across the whole brain was calculated.

3. Results

3.1. Sample

Subjects were moderately depressed, with a mean HDRS score of 16.2 ± 6.7 at the time of brain imaging, and a median of 2 previous major depressive episodes. The sample included 5 $L'L'$ homozygotes, 10 $L'S'$ heterozygotes, and 6 $S'S'$ homozygotes. Sample demographics are summarized in Table 1. Number of S' alleles did not correlate with

current depression severity ($r = -0.08$, $p = 0.72$) or number of lifetime depressive episodes ($\rho = 0.37$, $p = 0.09$).

3.2. Amygdala responses to negative stimuli: relationship to 5-HTT binding and 5-HTTLPR genotype

Higher 5-HTT V_T/f_p in the RN was associated with lower right (not left) amygdala responses to angry and fearful faces (Fig. 1, Cluster size 138 voxels, mean z -score -2.7). Higher 5-HTT V_T/f_p in amygdala bilaterally was associated with lower right amygdala response to faces (Cluster size 85 voxels, mean z -score -2.6). The mean neural response to faces across subjects revealed significant clusters in amygdala, midbrain (close to raphe nuclei), occipital and temporal fusiform gyrii, hippocampus, posterior and anterior cingulate (data not shown). A repeated main analyses using alternate PET outcome measures, revealed a predicted inverse relationship between RN BP_F^* and bilateral amygdala BOLD fMRI responses to faces, consistent with V_T/f_p findings. In contrast, RN BP_P^* and BP_{ND}^* were not correlated with amygdala responses to faces.

There were no clusters where 5-HTTLPR genotype was associated with response to faces in either the full 21-subject sample or the 16 subjects with multimodal PET and fMRI data.

4. Discussion

Consistent with our hypotheses, less 5-HTT binding in RN and amygdala was associated with greater amygdala reactivity assessed by fMRI. In contrast, 5-HTTLPR genotype was not associated with amygdala reactivity.

4.1. 5-HTT and neural response to negative stimuli

Our findings indicate that 5-HTT binding levels are related to emotion processing. 5-HTT levels may index serotonergic fiber density or intra-synaptic serotonin concentrations (Descarries et al., 1995; Soucy et al., 1994). Low RN 5-HTT binding may therefore indicate diminished RN serotonergic modulation of GABAergic neurotransmission in the amygdala (Bauman and Amaral, 2005; Jiang et al., 2009; O'Rourke and Fudge, 2006; Rainnie, 1999; Stutzmann and LeDoux, 1999). These data partially replicate and extend previous findings correlating regional [^{11}C]DASB binding to amygdala responses in healthy volunteers. In one study, a negative relationship was observed between right amygdala BOLD fMRI activity and midbrain [^{11}C]DASB BP_{ND} (Kobiella et al., 2011). In another study, a negative relationship was reported between amygdala reactivity and [^{11}C]DASB BP_P in the amygdala, but not in RN (Rhodes et al., 2007). Our study is notable for using within-subject PET images of the 5-HT $_{1A}$ somatodendritic receptor to identify the RN individually, and for its examination of this relationship in currently depressed medication-free subjects with MDD.

4.2. 5-HTTLPR and neural response to negative stimuli

5-HTTLPR was not associated with activity in amygdala in response to angry and fearful faces in our study. Our results diverge from some earlier studies in healthy controls that found the S' allele associated with amygdala responses to negative stimuli (Costafreda et

al., 2013; Hariri et al., 2002; Morey et al., 2011; von dem Hagen et al., 2011). Some studies have reported that the relationship between S' allele and amygdala reactivity is specific to healthy volunteers (Friedel et al., 2009; Gillihan et al., 2011; Rao et al., 2007) while others have observed this effect in MDD (Dannowski et al., 2007). The lack of observed genetic effect on amygdala responses may also be related to power limitations given the modest sample size in this study.

The association between RN 5-HTT binding and amygdala activity was limited to the right amygdala. Lateralized amygdala findings may be related to phase encoding direction artifact (Mathiak et al., 2012). However, other analyses in this study showed bilateral amygdala activity (group mean response to faces, and correlations with 5-HTT BP_F*). It is therefore less likely that phase encoding artifact explains left lateralization for one analysis only. Most studies of the amygdala report lateralized findings, which may reflect actual biological differences (Baas et al., 2004).

4.3. Limitations

The absence of genotype associations with amygdala response to negative emotional stimuli may be due to the modest sample size. Statistical power likewise was insufficient for analysis of potential gene-environment interactions that might be related to amygdala activity. While we conducted rigorous correction for multiple comparisons within whole-brain voxelwise analyses, we did not correct across analyses. These analyses should be regarded as hypothesis-generating, and require replication in larger samples. The lack of a healthy volunteer group precluded determination of depression-specific effects.

4.4. Conclusion

We found greater right amygdala reactivity to be associated with less RN 5-HTT binding but not 5-HTTLPR genotype. Future studies could employ more specific tasks and include a healthy volunteer group to further delineate the neural correlates of 5-HTT and 5-HTTLPR linked to depression.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Appendix A. Supplementary material

Supplementary data associated with this article can be found in the online version at <http://dx.doi.org/10.1016/j.jad.2015.10.047>.

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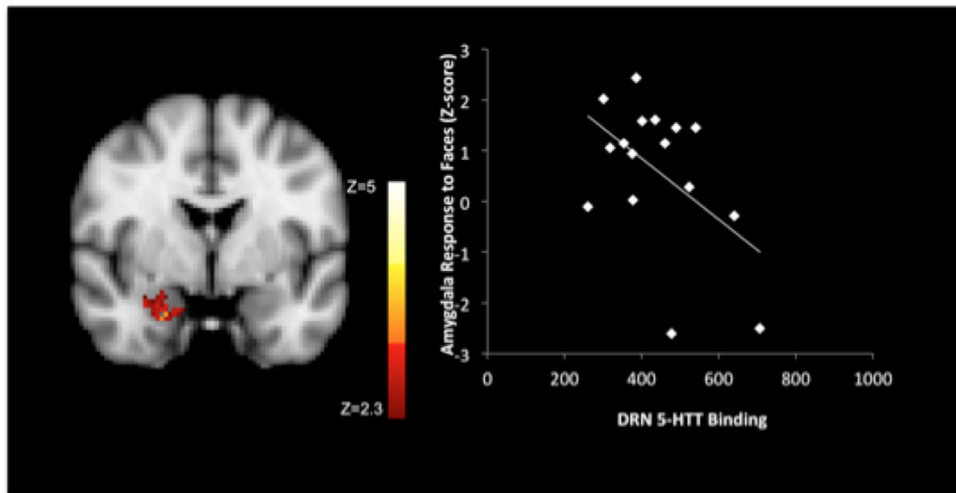


Fig. 1. Cluster of voxels in right amygdala whose response to faces is negatively associated with 5-HTT binding in the RN. Scatterplot shows correlation of average z-score in amygdala cluster with RN 5-HTT binding.

Table 1

Sample characteristics.

	Mean \pm S.D.
HDRS-17	16.19 \pm 6.65
Age	41 \pm 14.24
Education	15.9(3.12)
	Median(range)
Previous Episodes	2(0–109)
Weeks off antidepressant medications	24(0–364)
	Frequency (%)
Female	12(57.1)
Ethnicity	
White	13(61.9)
American Indian	1(4.7)
Black	5(23.8)
Multi-racial	2(9.1)
Medication Naïve	9(42.8)
First episode depression	8(38.1)
Smoking	3(14.2)
Co-morbid Anxiety	8(38.3)
Suicide attempt	4(19.1)