

Stress-induced activation of the HPA axis predicts connectivity between subgenual cingulate and salience network during rest in adolescents

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Background: Responses to stress vary greatly in young adolescents, and little is known about neural correlates of the stress response in youth. The purpose of this study was to examine whether variability in cortisol responsivity following a social stress test in young adolescents is associated with altered neural functional connectivity (FC) of the salience network (SN) measured during resting-state functional magnetic resonance imaging (rs-fMRI). **Methods:** Forty-nine typically developing young adolescents participated in a social stress test during which they contributed salivary cortisol samples. Following this, they underwent rs-fMRI scanning. We examined the association of FC of the SN [composed of anterior cingulate cortex and bilateral anterior insula regions] with cortisol responsivity. **Results:** Greater cortisol responsivity was significantly positively correlated with higher FC between subgenual anterior cingulate cortex (Cg25) and the SN, controlling for participant age. There were no regions of the brain that showed an inverse relation. **Conclusions:** Brain systems that have been implicated in autonomic arousal and that influence subjective feeling states show altered FC associated with stress responsivity in early life. **Keywords:** Resting-state, adolescents, HPA axis, stress, subgenual cingulate, fMRI, salience network, connectivity.

Introduction

When confronted with threat, humans and other animals show rapid engagement of both limbic brain circuitry [e.g., amygdala, prefrontal cortex (PFC), cingulate, anterior insula] and the hypothalamic–pituitary–adrenal (HPA) axis – a system reliably implicated in stress response. Limbic and HPA stress response systems are mutually regulatory, in part to diminish the stress response when it is no longer necessary (Pruessner et al., 2010). In studies of rodents and in human lesion studies, medial prefrontal cortices (mPFC) have been shown to play a critical role in control over the HPA axis (Buchanan et al., 2010; Figueiredo, Bruestle, Bodie, Dolgas, & Herman, 2003; Tchiteya, Lecours, Elie, & Lupien, 2003). Moreover, using high-resolution positron emission tomography (PET) with rhesus monkeys (Jahn et al., 2010) found elevated subgenual cingulate (Cg25) activity to be associated with heightened HPA axis activity. These results are of great interest given considerable evidence that dysregulated stress responding (see Liberzon et al., 2007) as well as elevated Cg25 activation and altered mPFC activity are involved in the pathophysiology of psychiatric disorders such as depression and anxiety (see meta-analysis by Seminowicz et al., 2004). Moreover, despite strong evidence linking frontolimbic dysregulation to HPA axis dysfunction and to difficulties

in emotion regulation, a hallmark characteristic of depression (Gotlib & Hamilton, 2008), few studies have measured directly how frontolimbic circuits are related to HPA axis responsivity.

In late childhood and early adolescence (ages 9–15), brain limbic circuitry is still developing (Cunningham, Bhattacharyya, & Benes, 2002), and protracted maturational brain changes are likely to affect the effectiveness of emotion regulation in children (Kovacs, Joormann, & Gotlib, 2008). A number of approaches have been used to map the ordering and timing of the development of brain circuitry, including cross-sectional studies of gray matter reduction (Giedd et al., 1999; Pfefferbaum et al., 1994), and synaptogenesis (Huttenlocher, 1979), as well as longitudinal studies mapping cortical thickness and brain growth in children (Sowell et al., 2004). This research has demonstrated that the PFC develops more slowly than do other brain areas (reviewed by Sowell, Thompson, & Toga, 2004). Given that early developmental mechanisms have been posited to influence the tendency of individuals to express maladaptive behaviors in response to threatening stimuli (i.e., heightened anxiety; see Gross & Hen, 2004), it is important to examine the relation between frontolimbic function and HPA axis reactivity in youth.

Recent advances in functional neuroimaging techniques, such as resting-state functional mag-

netic resonance imaging (rs-fMRI), afford new opportunities for studying *in vivo* the development of brain corticolimbic circuitry. By cross-correlating time-series data between regions, it is possible to determine which regions of the brain are functionally connected. Importantly, results obtained using this method are consistent with known anatomical connectivity data from postmortem human studies and anatomical tract tracing data from studies of other mammalian species (cf. Fair et al., 2010); moreover, findings of rs-fMRI studies have revealed important principles of neural organization both in normal and atypical human development (reviewed by Uddin, Supekar, & Menon, 2010).

Rs-fMRI data have reliably indicated that there is a canonical coherent network consisting of anterior cingulate cortex (ACC) and bilateral anterior insula (Seeley et al., 2007; Taylor, Seminowicz, & Davis, 2008). As this network comprises regions of the brain that are critical for interoceptive and emotional awareness, it has been labeled the salience network (SN; Seeley et al., 2007; White, Joseph, Francis, & Liddle, 2010). Regions of the SN have been shown to respond to pain, uncertainty, and other homeostatic challenges (Grinband, Hirsch, & Ferrera, 2006; Peyron, Laurent, & Garcia-Larrea, 2000). In addition, the SN has been found to be robust and reliable across repeat rs-fMRI measurements in children and young adolescents (Thomason et al., 2011). The SN is of particular interest for the present study because it involves regions known to modulate physiological responses to stress (i.e., insula; see Critchley, 2005).

The present study was designed to examine the association between functional interrelations of SN brain circuitry and HPA axis reactivity in children and adolescents. Connectivity of the SN has not been examined in the context of HPA axis reactivity. We applied a model-free independent component analysis (ICA) to resting-state data to define the brain SN in 49 children and young adolescents. The participants also underwent a social stress interview during which cortisol was sampled to measure HPA axis function. We hypothesized that the strength of the association between prefrontal cortical and SN functional connectivity (FC) would be influenced by HPA axis activity. As the mPFC expresses high levels of glucocorticoid receptors and is therefore involved in negative feedback control of the HPA axis, we hypothesized that significant group differences in connectivity would be observed in midline, prefrontal cortical regions. We further predicted that the Cg25, a structure that has been found to be involved in the experience of emotion in healthy individuals (Maddock, Garrett, & Buonocore, 2003) and that is both structurally and functionally anomalous in depression (Gotlib et al., 2005; Greicius et al., 2007; Liotti, Mayberg, McGinnis, Brannan, & Jerabek, 2002), may be more strongly coupled to SN function in individuals with higher levels of HPA system stress response.

Methods

Participants

Participants were 49 children and young adolescents (22 females) between the ages of 9 and 15 years ($M = 11.8$, $SD = 1.9$). They were recruited through their mothers via online forums, advertisements, and parent networks; each mother-child pair was compensated with \$25 per hour. All participants had no reported history of brain injury, no behavioral indications of possible mental impairment, no past or present Axis I disorder, were right-handed, were fluent in English, and had no learning disorder. Parents and participants gave informed consent and assent, respectively, as approved by the Stanford University Institutional Review Board.

Laboratory assessments

Interviews. Participants were administered structured interviews to assess current and lifetime psychopathology. Trained interviewers assessed the diagnostic status of the young adolescents by administering the Schedule for Affective Disorders and Schizophrenia for School-Age Children-Present and Lifetime version (K-SADS-PL) (Geller, Williams, Zimmerman, & Frazier, 1996; Geller et al., 2001), which has been shown to generate reliable and valid psychiatric diagnoses (Kaufman et al., 1997). During this session, participants and parents also viewed a video to prepare them for the MRI scan session.

Cortisol collection and stress task. Participants underwent a 15-min, controlled, laboratory stress interview that consisted of a 3-min serial subtraction task and the 12-min Ewart Social Competence Interview (Ewart, Jorgensen, Suchday, Chen, & Matthews, 2002), a semistructured interview developed to induce emotional stress and arousal in adolescents by discussing details of stressful life situations. Salivary cortisol was collected at regular intervals during and following the interview. During the serial subtraction task, children/adolescents were instructed to begin at 400 and count backward by sevens as quickly and accurately possible. If they made a mistake, they were interrupted by the experimenter and were told to start over. Further details on administration and analysis of the cortisol measurements and stress test are provided in the online Supporting Information.

Physiological responses during scanning

We wanted to examine whether physiological responses measured during fMRI scanning (i.e., heart rate and breathing rate) are related to measures of stress reactivity measured outside the scanner, particularly given that previous work has shown that the scanning environment can induce anxiety in youth (Eatough, Shirtcliff, Hanson, & Pollak, 2009), and that physiological responses are altered in response to anxiety provocation. If these measures were related, it would be necessary to test the relative contributions of physiological response during scanning vs. stress responsivity (HPA axis activation) to observed differences in brain FC. Thus, it was

critical to test the relation between these parameters in the present study and to correct fMRI data for physiological contributions to signal during data reconstruction (as described before). Bivariate correlation statistics were used to test whether physiological measures were related to cortisol responsivity. In addition, given that heart rates (HR) and respiration rates decrease with age, we tested the relations between participant age and physiological measures; based on our past work (Thomason, Burrows, Gabrieli, & Glover, 2005), we predicted a negative relation between age and HR, and between age and respiration rate. In an exploratory analysis, we examined whether participant gender was related to physiological parameters, given that recent studies have reported differences between women and men in both neural (Koch et al., 2007) and autonomic (Buchanan et al., 2010) responses to emotional material.

fMRI preprocessing

Participant data were preprocessed using Statistical Parametric Mapping software (SPM8; <http://www.fil.ion.ucl.ac.uk/spm/software/spm8/>). Preprocessing included image realignment and coregistration of functional and anatomical images. Functional images were normalized to the Montreal Neurological Institute (MNI) template, using the participant-specific transformation parameters created by fitting mean functional images to the single reference Echo Planar Imaging (EPI) standard SPM template. The default setting in SPM is to resample image data to $2 \times 2 \times 2$ mm during the normalization step of preprocessing; for normalization of this sample, however, data were not resampled, and thus retained the native resolution ($3.44 \times 3.44 \times 4$ mm). Following normalization, all participant images were visually inspected, and we determined that all normalizations had proceeded satisfactorily. Images were smoothed with a 6-mm Gaussian kernel to decrease spatial noise.

Independent component analysis

Data from all participants were concatenated and submitted to a group ICA procedure (<http://icatb.sourceforge.net>) implemented in Matlab (<http://www.mathworks.com>) using GIFT (Calhoun, Adali, Pearlson, & Pekar, 2001). Infomax was used to estimate 26

components. Our decision to derive 26 components was made by taking the mean number from three published reports that used similar analysis approaches (Stevens, Pearlson, & Calhoun, 2009; Thomason et al., 2011; White et al., 2010). Then, a spatial template-matching technique was used (as described in Greicius et al., 2007) to automatically identify the component corresponding to the SN. The SN template used in the present study was derived from an independent set of data (Seeley et al., 2007). Finally, using the GIFT processing toolbox, SN maps were back reconstructed for each participant. These ICA-derived individual participant SN maps were then carried into second-level random effects and regression analyses. Subject-specific SN maps derived by this method are therefore similar in overall spatial features but each has variance in spatial topography that reflects unique temporal features. The ICA-derived individual participant SN maps can be regarded as FC maps, in which the highest values are recorded in voxels with time courses that are highly correlated with the mean temporal trace for the SN for that participant, and thus areas of high FC to key regions within that network.

We used the ICA-based analytic procedure in this study because, unlike univariate methods (e.g., regression analysis), ICA does not rely on drawing inferences based on a priori regions of interest (ROIs). Selecting ROI seed points for children and adolescents based on previous work with adult samples is problematic. Although recent studies have examined properties of the SN in young populations (Fair et al., 2007; Stevens et al., 2009; Thomason et al., 2011), none of these studies provides an *independent* published set of SN peak coordinates that we could use to run a seed-based analysis in our youth sample. We have, however, recently published SN peak coordinates in a young sample (Thomason et al., 2011), and these coordinates may be useful in future work in which a seed-region-based analysis is more appropriate.

Random effects analysis

SPM8 (<http://www.fil.ion.ucl.ac.uk/spm/software/spm8/>) was used to create a group-wise statistical parametric map of the SN network. Back-reconstructed single subject spatial maps that corresponded to the SN were

Table 1 Summary of regions that comprise the salience network of the brain in youth

		X	Y	Z	No. of voxels	T	Z	p
Limbic/frontal								
Cingulate/MFG ^a	L32/9	-2	26	26	1,906	28.25	>7.5	<0.001
Subcortex/frontal								
Insula/IFG	R13/47	39	18	5	250	19.5	>7.5	<0.001
Insula/IFG	L13/47	-40	11	-6	192	18.94	>7.5	<0.001
Parietal								
Inferior	R40	62	-34	29	178	14.61	>7.5	<0.001
Inferior	L40	-64	-37	33	145	13.72	>7.5	0.001
Precuneus	R7	15	-70	35	15	9.71	7.19	<0.001
Cerebellum								
Culmen/anterior lobe		-30	-60	-26	15	8.8	6.76	<0.001

Coordinates are given in Talairach and Tournoux convention. BA, Brodmann's area; MFG, middle frontal gyrus; IFG, inferior frontal gyrus.

^aSecondary peaks with Euclidean distance >30 mm from the peak of this cluster were observed at -34, 43, 34 and 36, 43, 34, bilateral BA9 MFG.

Table 2 Regions for which FC was significantly positively correlated with cortisol response

		X	Y	Z	No. of voxels	T	Z	p
Limbic								
	Cingulate	L24	-13	3	16	3.37	3.17	0.001
	Cingulate	R25	8	18	2	2.87	2.78	0.003
Frontal								
	Medial	R9	15	36	15	3.03	2.88	0.002
	Medial	L25	-2	27	4	3.16	3.2	0.001
	Medial	R11	8	28	3	2.69	2.58	0.005
	Superior	R9/10	II	60	3	2.93	2.74	0.003
	Middle	L11	-43	44	2	2.92	2.79	0.003
Subcortex								
	Thalamus – ventral anterior nucleus		-13	-5	6	2.85	2.72	0.003

Coordinates are given in Talairach and Tournoux convention. BA, Brodmann’s area.

subjected to a one-sample *t*-test; a method that has been validated in previous studies (Stevens et al., 2009). Results were used to derive maximally significant peaks comprising nodes of the SN; these results are summarized in Table 2. Results reported are significant at a family wise error (FWE)-corrected level of $p < .001$.

Regression analysis

We conducted a voxel-wise regression analysis with the aim of identifying regions in which SN connectivity correlated significantly with stress-induced cortisol response. Cortisol response was computed using the incremental area under the curve (AUCi), an index of time-dependent change measuring the increase in cortisol from baseline across a series of measurements (as described by Pruessner, Kirschbaum, Meinlschmid, & Hellhammer, 2003). Before calculating AUCi, we excluded outlying cortisol responses ($>2 SD$) from the analysis, as has been done in previous cortisol studies (Gotlib, Joormann, Minor, & Hallmayer, 2008). Following this, SPM8 was used to conduct a whole-brain regression analysis examining the relation between HPA axis cortisol reactivity and FC of the SN of the brain. We controlled for age by including this variable as a regressor of noninterest in the regression model. Results for the whole brain are given for $p < .005$, uncorrected.

Results

Participant characteristics

Participants were 34 Caucasians (69%), 3 Hispanic Americans (6%), and 12 participants of multiracial or biracial descent (25%). Participants were 9–15 years of age, covering a range over which the processes of brain maturation are still occurring (Sowell, Thompson, Holmes, Jernigan, & Toga, 1999).

Stress reactivity

Participants showed a characteristic gradual and incremental rise in cortisol at 15 and 30 min following baseline and onset of the behavioral stressor, followed by a decrease at the fourth measurement

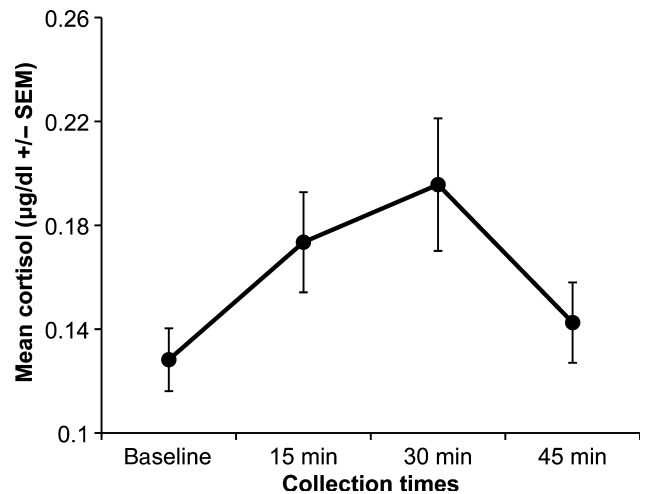


Figure 1 Stress reactivity measured across all participants, $N = 49$, ages 9–15

time point, 45 min after baseline (see Figure 1). The elevated cortisol production in response to stress observed in this sample is similar to what we have measured in our previous work with other samples of children (Gotlib et al., 2008). Interassay coefficient of variation (CV%)s were 11.46, 7.76, and 6.19 for low, medium, and high cortisol concentrations, respectively; and intra-assay CV%*s* were 7.73, 6.08, and 4.48 for low, medium, and high cortisol concentrations, respectively.

Cortisol AUCi was not significantly related to participants’ age, $r(49) = .21$, $p = .15$, nor did male and female participants differ with respect to cortisol AUCi, $t(47) = .74$, $p = .46$. Furthermore, the three ethnic groups did not differ in AUCi, $F(2,46) = .69$, $p = .5$. In addition, AUCi was not significantly correlated with physiological parameters (HR and respiration rate) measured during scanning, all $p > .5$.

Movement and physiological responses during rs-fMRI scans

Participant age was not significantly related to translational or rotational mean movement (all

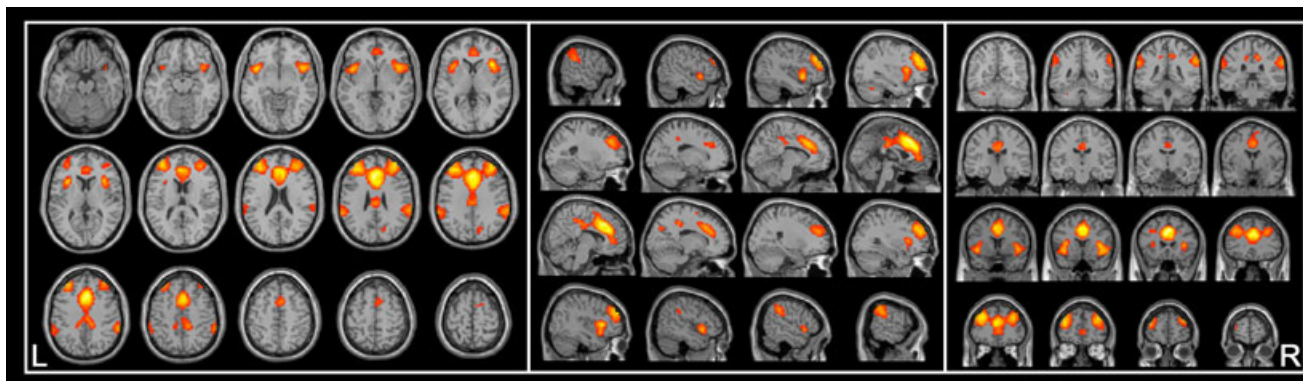


Figure 2 Saliency network (SN) functional connectivity across all study participants. Results of one-sample t -test are presented for $p < .001$, family wise error-corrected. Transverse, sagittal, and coronal slices of the SN are projected over a template brain to provide anatomical reference. L, left; R, right. Corresponding network peak coordinates are provided in Table 1

$p > .4$), max excursion (all $p > .2$), and mean jitter (all $p > .3$). In addition, men and women did not differ in movement for any of the six motion parameters, all $p > .05$. Across the participant sample, none of the six motion parameters were correlated with AUCi measurements, all $p > .15$. Finally, the number of frames retained for each participant after removing frames corresponding to movement spikes (described before) was not correlated with AUCi, $r(49) = .1$, $p = .58$.

As predicted, participant age was significantly negatively correlated both with HR ($M = 74$ beats/min), $r(48) = -.39$, $p = .007$, and with respiration ($M = 21$ breaths/min), $r(49) = .39$, $p = .006$. The relation between movement and physiological data was not significant for 11 of 12 correlations tested (six motion parameters, two physiological measures), all $p > .05$; there was, however, a significant

positive correlation between average HR and mean translational movement, $r(48) = .3$, $p = .04$. There was not a significant relation between the number of frames retained for the rs-fMRI analysis (after removing movement spikes) and physiological measurements, all $p > .05$. Finally, male and female participants did not differ in HR or respiration measures, or in number of frames removed for motion correction, all $p > .4$.

Saliency network

Regions we defined as comprising the SN in youth (see Figure 2, Table 1) are similar to the SN regions that have been reported in adults (White, Joseph, Francis, & Liddle, 2010) Brain regions in which HPA axis reactivity was significantly correlated with FC to the SN are listed in Table 2. Consistent with our

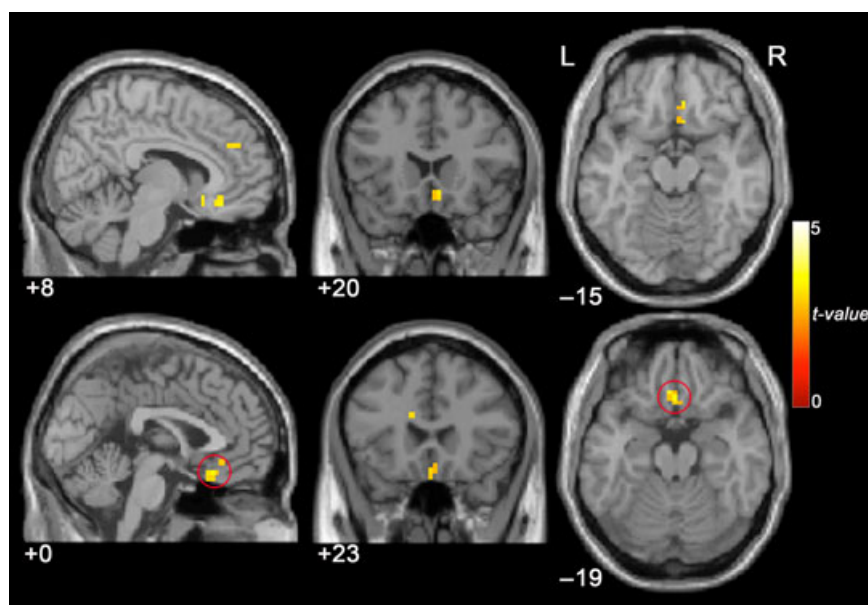


Figure 3 Hypothalamic–pituitary–adrenal (HPA) axis reactivity was significantly correlated with saliency network activity during resting-state functional magnetic resonance imaging. Participants with higher HPA axis response (measured using cortisol incremental area under the curve) showed significantly higher ($p < .01$) temporal coupling in regions of the medial frontal cortices and subgenual cingulate (Cg25; encircled by red rings)

hypothesis, greater stress reactivity was associated with greater temporal coupling between time-series data from the subgenual anterior cingulate cortex (Cg25) and from areas of the mPFC and the resting time course of the SN. Regions showing significant HPA response-by-SN FC correlation are shown in Figure 3.

Discussion

The present study was designed to examine the relation between activation of the HPA axis in response to stress and neural network brain connectivity in young adolescents. We found that those adolescents who exhibited higher HPA axis reactivity to a structured social stress interview administered in the laboratory had higher functional coupling between the subgenual anterior cingulate (Cg25) and the salience network (SN). Cg25 has been identified as a key component in studies of self-referential processing and emotional monitoring (Zald, Mattson, & Pardo, 2002). There is considerable evidence to suggest that aberrant Cg25 neural activation is associated with disturbed emotional (Gotlib et al., 2005) and physiological function (Wager et al., 2009). Aberrant metabolism in Cg25 has been reported in sad mood induction paradigms in both normal and depressed subjects (Damasio et al., 2000; Liotti, Mayberg, Mcginnis, Brannan, & Jerabek, 2002; Mayberg et al., 1999). Furthermore, rs-fMRI has revealed increased default-mode network (DMN) FC in Cg25 in adults with major depression (Greicius et al., 2007). In fact, the Cg25 region has been the successful target of deep brain stimulation for treatment-resistant depression (Mayberg et al., 2005), further corroborating the unique centrality of this region in the neuropathophysiology of mood disorders. The subgenual cortex has recently been associated with HPA system activity in rhesus monkeys (Jahn et al., 2010), but to our knowledge the present study is the first demonstration of an association between HPA response magnitude and Cg25 neural network function in humans.

To identify patterns of brain network connectivity that may correspond to heightened stress reactivity, we took an approach that was both theory- and data-driven. While a more extensive analysis may have included several brain rs-fMRI networks accumulated from group ICA analysis, we chose to limit our analysis to the SN, with special emphasis on the theoretical construct of physical response to psychological stress (i.e., HPA activation). SN circuitry has been implicated in self-generated emotional processing (Damasio et al., 2000), homeostatic processing (Seeley et al., 2007; Taylor et al., 2008), resolving uncertainty (Grinband et al., 2006), anticipatory anxiety (Chua, Krams, Toni, Passingham, & Dolan, 1999), and error detection (Menon, Adleman, White, Glover, & Reiss, 2001). In

particular, the ACC component of the SN may be specialized in generating autonomic changes, whereas the anterior insular component may be specialized for monitoring visceral sensory inputs (reviewed by Critchley, 2005). Research using rs-fMRI has begun to establish a critical role for insular components of the SN in switching between activity in the central executive network [also known as the 'task-positive network' (TPN)] and the DMN [also called the 'task-negative network'] (Sridharan, Levitin, & Menon, 2008). Work by our group has demonstrated that individuals with major depression who adopt positive, adaptive rumination strategies engaged the TPN more often, and that engagement of TPN was associated with increased activation of the anterior insula (Hamilton et al., 2011). Our previous results, therefore, implicated an adaptive relation between the TPN and DMN in depression that may be instigated by insula-mediated awareness of negative emotional states. In the present study, we have extended this formulation by demonstrating that the SN, a system that is critical to awareness and that may drive action in other brain networks, is characterized by increased functional coupling with medial prefrontal circuitry that is implicated in key aspects of emotional (Mayberg, 2003) and physiological function (Wager et al., 2009) in children and adolescents with greater HPA axis response to psychological stress. It is noteworthy that the results obtained for this unmasked, whole-brain analysis demonstrate specificity, in that significant results reside almost exclusively in medial frontal brain areas. Importantly, there were no regions of the brain in which there was a negative correlation between FC and cortisol reactivity.

By identifying neural processes that are associated with maladaptive stress responses in late childhood and early adolescence, our results elucidate differences in brain connectivity that may influence the development of adaptive emotional responses. Investigators have demonstrated that maladaptive behaviors that have been found to be associated with emotional disorders are not simply correlates of the disorder, but instead may play a critical role in the etiology and maintenance of psychopathology (e.g., Joormann, Talbot, & Gotlib, 2007). We complement these findings by focusing on neural developmental processes and, in this context, document the significance of limbic and prefrontal circuitry that is continuing to develop across the age range studied here (Cunningham et al., 2002; Sowell, Thompson, & Toga, 2004). Brain white-matter tracts continue to mature through adolescence (reviewed by Thomason & Thompson, 2011), and with this maturation, there is a strengthening of long-range, and a weakening of short-range functional connections (Fair et al., 2007). We suggest that altered FC influences individuals to experience and express maladaptive responses to stress, and that repeated experiences of a heightened stress response will, in turn, strength-

en the activation of brain regions that will further exacerbate these effects.

While the present results highlight neural circuitry that may underlie the association between psychological experience and visceral reactions of the body, we should note a limitation of this research and an important difference between this study and previous work. First, our measure of HPA axis reactivity was obtained within 1 week of the scan session, but on a different day. Although we found an association between HPA axis reactivity and neural FC, it will be important in future research to assess neural connectivity and cortisol reactivity concurrently. Recent empirical studies have documented that participants in MRI studies experience a range of stress in response to the scanner environment (Muehlhan, Lueken, Wittchen, & Kirschbaum, 2010), and it would be of interest to distinguish transient vs. trait-like HPA axis activation effects on FC.

Conclusion

In the present study we used an FC neuroimaging strategy to identify neural system-level mechanisms corresponding to heightened HPA axis activity in response to stress. Whole-brain analysis implicated significant FC differences that occurred almost exclusively in midline frontal cortical brain regions (e.g., Cg25 and BA11; see Table 2) that have been found in previous studies to be associated both with mood and anxiety disorders and with physiological responses to stress. While investigations of functional abnormalities during task-provocation-based BOLD fMRI studies have provided insights about heightened HPA axis reactivity, our data underscore the importance of using rs-fMRI to examine dynamic relations in neural systems that are not constrained by a particular type of stress-induced thought or activity. We think that this analysis cap-

tures unique aspects of the altered neurodevelopmental processes that can lead to maladaptive behaviors or represent a vulnerability to developing a psychiatric disorder. This result provides evidence that in early life, greater stress reactivity is associated with network-level neural patterns that may hinder the development of healthy emotional reactivity.

Supporting information

Additional Supporting Information may be found in the online version of this article:

Appendix S1

Please note: Wiley-Blackwell are not responsible for the content or functionality of any supporting materials supplied by the authors. Any queries (other than missing material) should be directed to the corresponding author for the article.

Acknowledgements

This project was supported by awards from the National Institute of Mental Health (MH081583 to M.E.T., MH079651 to J.P.H., and MH074849 to I.H.G.), and by NARSAD Young Investigator Awards to M.E.T. and to J.P.H. The authors thank Melissa Henry, Emily Dennis, and Rebecca Johnson, for their assistance in acquiring the scan data, and Yamanda Wright, and Hannah Burley for their assistance in participant recruitment, screening, and conducting structured behavioral interviews. The authors also thank Jutta Joormann, Ben Varasteh, and Michael Chen for consultation on cortisol analyses.

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Key points

- Intrinsic brain connectivity is related to HPA axis stress response in adolescents.
- Subgenual cingulate connectivity is higher in adolescents with a greater stress response.
- Anomalous patterns of neural network connectivity may dispose children and adolescents to develop maladaptive responses to stress.
- The present findings are among the first to examine how frontolimbic circuitry is related to HPA axis reactivity in humans.
- The results support the use of task-free fMRI for interrogating tonic aspects of neural connectivity in children and adolescents.

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Accepted for publication: 15 March 2011

Published online: 3 June 2011