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Female-specific intergenerational transmission patterns of the human corticolimbic circuitry

(Abbreviated title: Intergenerational effects in the human brain)

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Abstract

Parents have large genetic and environmental influences on offspring's cognition, behavior, and brain. These intergenerational effects are observed in mood disorders, with particularly robust association in depression between mothers and daughters. No studies have thus far examined the neural bases of these intergenerational effects in humans. Corticolimbic circuitry is known to be highly relevant in a wide range of processes including mood regulation and depression. These findings suggest that corticolimbic circuitry may also show matrilineal transmission patterns. We therefore examined human parent-offspring association in this neurocircuitry, and investigated the degree of association in gray matter volume between parent and offspring. We used voxel-wise correlation analysis in a total of 35 healthy families, consisting of parents and their biological offspring. We found positive associations of regional grey matter volume in the corticolimbic circuit including the amygdala, hippocampus, anterior cingulate cortex, and ventromedial prefrontal cortex between biological mothers and daughters. This association was significantly greater than mother-son, father-daughter, and father-son associations. The current study suggests that the corticolimbic circuitry, which has been implicated in mood regulation, shows a matrilineal specific transmission patterns. Our preliminary findings are consistent with what has been found behaviorally in depression, and may have clinical implications for disorders known to have dysfunction in mood regulation such as depression. Studies such as ours will likely bridge animal work examining gene expression in the brains and clinical symptom-based observations, and provide promising ways to investigate intergenerational transmission patterns in the human brain.

Significance Statement

Parents have large genetic and environmental influences on the offspring, known as intergenerational effects. Specifically, depression has been shown to exhibit strong matrilineal transmission patterns. While intergenerational transmission patterns in the human brain are virtually unknown, this would suggest that the corticolimbic circuitry relevant to a wide range of processes including mood regulation may also show matrilineal transmission patterns. We therefore examined the degree of association in corticolimbic grey matter volume (GMV) between parent and offspring in 35 healthy families. We found that positive correlations in maternal corticolimbic GMV with daughters were significantly greater than other parent-offspring dyads. Our findings provide new insight into the potential neuroanatomical basis of circuit-based female-specific intergenerational transmission patterns in depression.

Introduction

Parents have large genetic and environmental influences on offspring's cognition, behavior, and brain (Curley and Mashoodh, 2010; Cabrera et al., 2011; Curley, 2011). Furthermore, intergenerational trait transmission is often observed in the psychopathology of major psychiatric conditions, such as mood disorders (Curley and Mashoodh, 2010; Guilmatre and Sharp, 2012). For example, offspring of depressed parents are two to three times more likely than the offspring of non-depressed controls to exhibit elevated levels of depression (Weissman et al., 2006). If so, the neurocircuitry involved in depression may also show intergenerational transmission effects. A large body of human neuroimaging studies indicate that corticolimbic circuitry, which includes the amygdala, hippocampus, anterior cingulate cortex (ACC) and ventromedial prefrontal cortex (vmPFC - orbitofrontal cortex [OFC] and rectus gyri), is central in mood disorders (Price and Drevets, 2010). Growing animal and human neuroimaging literature has similarly implicated this corticolimbic circuitry as the biological substrate of emotion regulation (Banks et al., 2007). This points to corticolimbic circuitry as an important target for investigating depression and its intergenerational effects.

In depression, a maternal transmission pattern is seen, with maternal depression associated with numerous adverse outcomes among offspring (Downey and Coyne, 1990; Beardslee et al., 1998), with particularly robust association between maternal and adolescent daughter depression (Davies and Windle, 1997; Goodman and Gotlib, 1999; Goodman, 2007; Thompson et al., 2014). More specifically, maternal depressive symptoms are correlated with that of daughters ($r = 0.44$) but not sons ($r = -0.01$) (Fergusson et al., 1995), as well as maternal depressive/anxiety disorder correlating with daughters' but not sons' internalizing symptoms (Piche et al., 2011). Accordingly, we would expect that there is a female-specific neurobiological transmission pattern in the

corticolimbic circuitry. Additionally, in the animal literature, maternal gestational stress significantly decreased the number of hippocampal neurons due to increases in calcium content and oxidant generation in the hippocampal CA3 region in female offspring but not in male offspring. Females may have a higher generation of oxidants due to higher corticosterone caused by prenatal stress (Zhu Z et al., 2004). Moreover, the hypothalamus-pituitary-adrenal (HPA) axis responses to gestational stress are different between male and female offspring. Basal and stress-induced increases in corticosterone levels are found only in prenatally-stressed female offspring (Weinstock M et al., 1992; Szuran et al., 2000). Such female-dominant HPA axis activity may be caused by a greater transport of corticosterone from maternal blood across the placenta of female fetuses than of male fetuses (Montano et al., 1993). These animal studies have revealed the underlying neurobiological underpinnings of female-specific intergenerational effects. However, to date, there have been no neuroimaging studies that examined the neural evidence of intergenerational transmission patterns in humans. Studies such as these can contribute unique information that can bridge animal research and human clinical studies that is different from the more common approach, i.e., to examine heritability in twins, where sex-specific intergenerational transmission patterns cannot be assessed (Alemany et al., 2013).

The purpose of the current study was to examine female-specific intergenerational transmission patterns on brain structure in parent-offspring pairs in healthy individuals. We hypothesized that mother-daughter pairs would reveal significantly positive grey matter volume (GMV) associations in the corticolimbic brain regions (implicated in mood disorders) compared to other parent-offspring pairs. Our hypothesis was based on findings from previous studies demonstrating that depression exhibits strong female-specific intergenerational transmission patterns indicated above (Fergusson et al., 1995; Goodman and Gotlib, 1999; Goodman, 2007; Piche et al.,

2011). We therefore focused on corticolimbic circuitry including the amygdala, hippocampus, ACC, and vmPFC GMV and analyzed the degree of association between mothers and daughters compared to mothers and sons, fathers and sons, and fathers and daughters, in these brain regions.

Materials and Methods

Participants

A total of 35 healthy families, consisting of parents ages 33 to 48 years and their biological offspring ages 5 to 13 years, were included for the present study from a larger study that was not focused on corticolimbic system and related behavior. Participants were recruited from local newspapers, school mailings, flyers, and mother's clubs. Exclusion criteria for this study were the presence of neurological or psychiatric disorders including substance abuse, the use of medication, and/or contraindication to MRI (e.g., metal in the subject's body), in either parent or offspring reported by the parents. It is still possible however that mothers omitted reporting past or current use of steroids including contraceptives which may effect brain morphometry (Pletzer et al., 2010), because these are commonly taken among healthy people. The Stanford University and University of California San Francisco (UCSF) Panel on Human Subjects in Medical Research approved the study. After complete description of the study to the participants, written informed consent and assent were obtained from parents and children, respectively. Thirty-one families had one child and 4 families had two children. We created 4 parent-offspring subgroups including 16 mother-daughter, 18 mother-son, 15 father-daughter, and 18 father-son pairs.

Demographic, cognitive, and behavioral assessments

Behavioral data (such as mental health status) of offspring were obtained by the Behavioral Assessment System for Children-2nd Edition (BASC-2) (Reynolds and Kamphaus, 2004), which is a multimethod, multidimensional system used to evaluate the adaptive and problematic behavior of children and young adults aged 2 through 25 years. We collected parent-rating scales (PRS) of nine clinical scales including Aggression, Anxiety, Attention Problems, Atypicality, Conduct Problems, Depression, Hyperactivity, Somatization, and Withdrawal and five adaptive scales including Activities of Daily Living, Adaptability, Functional Communication, Leadership, and Social Skills. The BASC-2 was normed in the United States using a large representative sample of youth (including over 13,000 cases) from across the nation. Aged-normed T-scores are continuously distributed within the population, with a mean of 50 and a standard deviation of 10. A high score implies more emotional and behavioral problems, except for the adaptive skills scale for which a high score implies more attributes that contribute to healthy functioning. More specifically, on the clinical scales, T-scores from 60–69 indicate “at-risk” behaviors, and 70 and above indicate “clinically significant” concerns. On the other hand, analogous cutoffs are established for the Adaptive Skills scale whereby T-scores from 31–40 are considered to be “at-risk”, and scores 30 or less are considered to be “clinically significant”. To evaluate potential sex differences in age in our samples, we utilized independent Student’s t-tests for age in parents and offspring, respectively. Since the BASC-2 PRS consists of fourteen subscales, we performed a multivariate analysis of covariance (MANCOVA) to compare group differences between male and female offspring using these BASC-2 PRS subscales as dependent variables. Then, if a main effect of group on the MANCOVA was confirmed, a post-hoc ANCOVA was performed. While there were no significant differences in age, age was included as covariate in the analyses because of the relatively wide range in age of the children (age range: 5 - 13). Since all data were originally collected for the purpose of a

study examining the neural bases of academic achievement in the offspring, there were no relevant behavioral data in parents (except for measures such as IQ).

Image acquisition and preprocessing

MR images were acquired using a 3T whole-body GE-Signa HDxt scanner (GE Medical Systems, Milwaukee, WI) with a quadrature head coil at the Lucas Center for Imaging at Stanford University. For voxel-based morphometry (VBM), high-resolution T1 anatomical scans were obtained using a spoiled gradient recalled (FSPGR) echo pulse sequence (160 coronal slices; TR/TE, 8.5/1.7 ms; inversion time, 300 ms; flip angle, 15°; NEX, 1; FOV, 22 cm; matrix, 256 × 256; thickness, 1.2 mm; acquisition time, 4 min 34 sec). Images were first visually checked for scanner artifacts and anatomical anomalies. Pre-processing was done using DARTEL (Ashburner, 2007) for VBM in SPM8 (<http://www.fil.ion.ucl.ac.uk/spm>). We further selected the modulated normalized-non linear only option, which results in an analysis of relative differences in regional GMV, corrected for individual brain size. This allows comparing the absolute amount of tissue corrected for individual brain sizes.

Parent-offspring correlation in corticolimbic circuitry

First, we performed voxel-wise correlation analysis of volumes between mother and daughter, controlling for daughter's age and limiting the analysis to the region of interest (ROI) using the Biological Parametric Mapping (BPM) toolbox (Casanova et al., 2007). We created one corticolimbic ROI including bilateral amygdala, ACC, vmPFC including the OFC and rectus gyrus, hippocampus, and parahippocampus gyrus defined by Automated Talairach Atlas Label (Lancaster et al., 2000) in the WFU PickAtlas toolbox (<http://fmri.wfubmc.edu/cms/software#PickAtlas>). ROI

analyses were performed using a statistical threshold of $P = 0.05$ for voxel-height, which then was thresholded at a cluster threshold of $P = 0.05$ for family-wise error (FWE). This resulted in brain map showing significant positive GMV correlations between mother and daughter.

Second, because our primary goal was to examine the specificity of the parent-offspring GMV relationship, the next steps were to create brain maps that compared correlation coefficients (r) between different parent-offspring groups on a voxel-wise basis in the corticolimbic ROI. Based on our a priori hypothesis, we generated three brain maps: 1. correlation coefficients (association) that were significantly stronger between mother-daughter compared to mother-son pairs to examine whether mothers show stronger effects on daughters than sons, 2. mother-daughter greater than father-daughter association to examine whether mothers compared to fathers show stronger effects on daughters, and 3. mother-daughter greater than father-son association to show, to the extent possible, sex-specific effects. One challenge in testing for statistical differences between correlations in our data is the partial dependency of the sample. For example, when we compare mother-daughter correlation with mother-son correlation, some observations are dependent (e.g., a mother who has a daughter and a son) but some others are not (e.g., a mother who has a daughter only). To address this statistical issue, we used a bootstrapping method (Efron and Tibshirani, 1993) to assess the significance of the difference between the two correlation coefficients. Specifically, we derived the standard errors of the differences in correlations (e.g., standard error of the differences in mother-daughter correlation and mother-son correlation) by repeatedly resampling the observed data. Importantly, the resampling procedure takes into account the (partial) dependency of the data, making it possible to precisely compute the standard errors. Before implementing bootstrapping, each individual's processed voxel-wise GMV values within the corticolimbic ROI were extracted and converted to matrices using an in-house Matlab-based MVPA toolbox (Hoefl et al., 2008; 2011).

We then ran bootstrapping with an in-house *R* code (R Core Team, 2015). As a result, we obtained three brain maps representing significantly greater positive GMV correlation in mother-daughter pairs ($P < 0.05$) compared to mother-son, father-daughter and father-son pairs, respectively. Finally, we overlapped these three brain maps and the FWE-corrected mother-daughter correlation map to identify the voxels that showed significantly greater positive correlations between mother-daughter pairs than for the other three pairings.

Correlation between regional brain volume and behavioral data

To examine whether the degree of GMV in brain regions that show significant positive association between mothers and daughters is associated with emotional and behavioral functioning in daughters, Pearson's correlation analyses were performed between GMV data and standardized scores from all BASC-2 scales obtained from the daughters. Based on previous behavioral reports showing female-specific transmission patterns in depression (Davies and Windle, 1997; Goodman and Gotlib, 1999; Goodman, 2007; Thompson et al., 2014), we hypothesized that there may be associations between daughters' GMV and their BASC-2 depression subscales. GMV data were extracted from clusters in the amygdala, ACC, OFC, rectus gyms, and hippocampus that showed significant positive associations between mothers and female offspring compared to other parent-offspring dyads within the corticolimbic ROI. All behavioral data were originally collected for the purpose of a different study, and hence we only had data from 10 out of 16 daughters, making these results tentative and exploratory at best.

Results

Corticolimbic ROI analysis

In offspring, a MANCOVA showed no significant sex differences for T-scores in the BASC-2 PRS scales (Wilks' Lambda 0.079, $F_{(14, 4)} = 3.33$, $P = 0.127$) (Table 1), hence, no post-hoc analysis was performed. We performed voxel-wise statistical comparisons of correlation coefficients between different parent-offspring pairs to find regions within the corticolimbic ROI in parent-offspring correlations were strongest for mother-daughter pairs. Analyses comparing mother-daughter vs. mother-son, mother-daughter vs. father-daughter, and mother-daughter vs. father-son showed quite similar morphometric association patterns. Specifically, positive correlations between mothers and daughters were significantly greater than for other parent-offspring pairs in many of the corticolimbic structures, including the right amygdala, the bilateral ACC (more prominent on the left), vmPFC including the bilateral rectus gyrus (more prominent on the left) and bilateral OFC (more prominent on the right), and the right hippocampus and the bilateral parahippocampus gyrus (Table 2, Fig. 1). We also overlapped these brain maps to identify the brain areas that are most strongly correlated in mother-daughter relationships. Although the extent of brain regions became smaller (perhaps due to the conservative nature of conjoint analysis), we still observed significant voxels in the same brain areas (Fig. 2).

Correlation between regional brain volume and behavioral data in daughters

There was a positive correlation between GMV of the rectus gyrus and adaptability T-scores ($r = 0.710$, $P = 0.021$ uncorrected); however, these results did not reach statistical significance when applying Bonferroni correction. There was no significant correlation between GMV of the other brain regions and any behavioral measures despite other subscales showing similar or even greater variance (all P 's > 0.1 corrected).

Discussion

The present study found female-specific intergenerational effects on brain structure between biological parent-offspring dyads. Specifically, consistent with our hypothesis, positive associations of regional grey matter volume between mothers and daughters were significantly greater than mother-son, father-daughter, and father-son associations in the amygdala, hippocampus, ACC, and vmPFC, which included the rectus gyrus and OFC. These findings may indicate that the daughter's corticolimbic circuitry is associated specifically with that of the mother's. A growing number of animal and human neuroimaging studies have implicated corticolimbic circuitry, specifically the amygdala, ACC, and vmPFC, as the biological substrates of emotion regulation. The amygdala regulates a variety of emotions, including depression, anxiety, and fear, and plays a major role in emotional memory processing and reactivity of the hypothalamus pituitary adrenal (HPA) axis to stress (Banks et al., 2007). The ventral ACC, in particular the pregenual and subgenual ACC, has functional connections with OFC, amygdala, and hippocampus and participates generally in emotion regulation (Drevets et al., 2008; Pizzagalli, 2011). The vmPFC, via its direct structural and functional connections with the amygdala (Ghashghaei et al., 2007) modulates emotional responses through GABAergic inhibitory projections (Akirav et al., 2006). Dysfunction in this circuitry is strongly associated with conditions such as mood and anxiety disorders (Price and Drevets, 2010). In addition, the finding in the hippocampus further supports our hypothesis. The hippocampus is one of the key components of emotional regulatory networks in the brain. Previous studies have reported that depression is associated with smaller hippocampal volume due to glucocorticoid neurotoxicity and stress induced reduction in neurotrophic factors and neurogenesis (Sheline, 2011).

In this study, we show the intriguing finding that mother and daughter's corticolimbic morphology was more similar than other parent-offspring pairs. Our study provides novel

neurobiological support for female-specific transmission patterns in depression. Several previous studies have revealed that healthy controls with a family history of depression (high risk group) compared to those with no family history of any psychiatric disease (low risk group) had structural abnormalities in corticolimbic circuitry including smaller hippocampus, smaller ACC, and larger amygdala volumes. They suggested these structural anomalies may be potential neurobiological markers of vulnerability for depression (Boes et al., 2008; Chen et al., 2010; Romanczuk-Seiferth et al., 2014). Moreover, other data suggest that the combination of early-life adversity and smaller hippocampal size significantly increases the vulnerability for depression (Rao et al., 2010). Together, our findings indicate that if mothers have brain structural anomalies in the corticolimbic circuitry, their female but not male offspring are more likely to have similar abnormal structural patterns in the same brain regions. While further research is warranted, these matrilineal associations may be tightly linked to greater vulnerability for daughters but not sons in developing depression when their mothers have depression.

It is currently difficult to point to the precise underlying mechanisms of our findings due to lack of prior studies. Nonetheless, one may speculate that genetic, prenatal and postnatal environmental factors and their interactions may underly female-specific intergenerational patterns in depression symptoms, brain morphometry and gene expression patterns in the corticolimbic circuitry. In addition to animal studies mentioned in the introduction, gestational stress leads to increased corticotropin releasing hormone (CRH) messenger RNA (mRNA) in the amygdala (Cratty et al., 1995), and the amount of maternal care influences epigenetic modification (methylation) to hypothalamic ER α in female rats (Champagne et al., 2006). These studies provide the underlying neurobiological evidence of female-specific intergenerational effects by examining gene expression in brain tissue. Furthermore, female but not male offspring of mice show a reduction in hippocampal

glial count in the pyramidal layer following prenatal stress (Behan et al., 2011). In humans, one multi-site survey in both genetically related and unrelated parent-offspring pairs reported that girls may be more prone to the negative influences of maternal depression symptoms than boys through environmental processes (Lewis et al., 2011). A previous study revealed that higher maternal prenatal cortisol level and postnatal depression problems were associated with larger right amygdala volumes in girls but not boys (Buss et al., 2012). These findings indicate the presence of female-specific placental adaptation to stress exposure (Clifton, 2010), which may influence more similar neurodevelopmental trajectory among females compared to males in the corticolimbic circuitry. Longitudinal studies in families with different genetic, prenatal and postnatal environmental influences are needed to examine the relationship between maternal stress, endocrine effects and brain development in mothers and their daughters.

Furthermore, in the rectus gyrus, which is the medial part of the OFC, a positive correlation between GMV of daughters and their adaptability in the BASC-2 was found, such that larger volumes of the OFC may be related to higher levels of adaptability. This finding may support previous animal studies indicating the OFC contributes to emotional and social adaptive behavior (Roberts, 2006). Behavioral measures in the current study were collected for the broader purpose of another study and were not optimized to examine intergenerational effects, hence together with the small sample and lack of measures in parents, the results of this exploratory analysis should be interpreted with caution.

There has been increasing consensus on sex differences in grey matter developmental trajectories (Giedd et al., 1999; De Bellis et al., 2001). Cerebral and grey matter volume in the frontal and parietal lobes peak earlier in girls than in boys (Lenroot and Giedd, 2006). Moreover, sex differences have been demonstrated in the hippocampus (larger in females), amygdala (larger in

males), and thalamus (larger in males) (Koolschijn and Crone, 2013); however, our study found that mother-daughter pairs consistently demonstrated significantly greater positive GMV correlations compared to the other parent-offspring pairs not only in the amygdala and medial PFC where volume is larger in males than in females, but also in the OFC and hippocampus where volume is larger in females than in males (Cosgrove et al., 2007; Koolschijn and Crone, 2013). These results indicate that sex difference in these brain volumes cannot account for the pattern of associations we observed. Further research is needed to examine the mechanisms underlying the consistent female-specific association patterns in each brain region showing a pronounced sexual dimorphism in development within the corticolimbic circuitry.

Several limitations of this study should be noted. First, since we only utilized genetically-related families, where genetic, prenatal and postnatal environments are shared, and did not include in-vitro fertilization (IVF) and adoptive families, we were unable to dissociate the influence of these three factors on our neurobiological findings of intergenerational effects. Second, we did not directly examine how functional and anatomical connectivities within corticolimbic circuitry are mediated by the intergenerational effects in this sample. Therefore, further studies using multimodal imaging in three different groups (genetic, IVF, and adoptive groups) are needed. Third, we only focused on matrilineal transmission patterns in the human brain based on our hypothesis; however, Rodgers et al. found that stressed male mice show an increase in microRNAs in their sperm, leading to reduced HPA stress axis responsivity in adult offspring (Rodgers et al., 2013; 2015). The result indicates novel epigenetic mechanisms of paternal transmission patterns in neuropsychiatric disorders including depression. Thus, it is also important to examine paternal intergenerational transmission effects in the corticolimbic circuitry in future studies. Fourth, the human amygdala appears to undergo rapid development early in life (Tottenham et al., 2009). Amygdala structural

growth is complete by four years old (Giedd et al., 1996) and functional development reaches its peak during adolescence (Hare et al., 2008). On the other hand, in rodent models, the vmPFC develops late in life (Van Eden and Uylings, 1985) and connectivity between the amygdala and the vmPFC does not emerge until adolescence (Cunningham et al., 2002). Therefore, longitudinal studies are required to explore how our findings of female-specific intergenerational patterns are affected by the developmental trajectory in the corticolimbic circuitry. In the current study, while cross-sectional, we attempted to capture children with minimal influence of puberty and extra-family environment on the brain (Dahl, 2004). Finally, because of the relatively small sample size and exploratory nature of our analysis, the current findings are preliminary and require further research with a larger sample. The current study however, represents the first study to examine parental transmission patterns of corticolimbic morphometry in humans *in vivo*.

In sum, to our knowledge, this is the first study to investigate intergenerational transmission patterns in the human brain using neuroimaging. We demonstrate a significant matrilineal association between parents and their offspring in a brain circuit associated with emotion regulation as measured by regional GMV. Our results provide new insights into the potential neuroanatomical basis of female-specific intergenerational effects of psychiatric conditions, especially in depression. Findings from our study have the potential to bridge animal work and clinical observations. Ultimately, findings such as ours may have important implications for furthering the understanding and the creation of developmental-stage specific and circuit-based treatment intervention of mental illnesses, such as depression, that show intergenerational effects.

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Figure 1. Grey matter volume associations between parents and offspring. **Top**, Brain regions showing significantly greater positive correlations in mother-daughter pairs compared to mother-son pairs. **Second row**, Brain regions showing significantly greater positive correlations in mother-daughter pairs compared to father-daughter pairs. **Third row**, Brain regions showing significantly greater positive correlations in mother-daughter pairs compared to father-son pairs. All comparisons showed similar morphometric association patterns. OFC, orbitofrontal cortex; ACC, anterior cingulate cortex. Axial view of Talairach coordinate (TC), z: -27, -22, -17, -12, -7, -2; sagittal view TC, x: 29, 13, -3. Lt, left; Rt, right. For all analyses, the statistical threshold is set at $P = 0.05$ corrected.

Figure 2. Brain areas specific to mother-daughter relationships. Brain map representing voxels that overlap between three maps presented in Figure 1 to illustrate a specific positive association in mother-daughter pairs. Red circle represents amygdala. Blue circle represents rectus gyrus. Green circle represents orbitofrontal cortex. Pink circle represents anterior cingulate cortex. Yellow circle represents parahippocampus gyrus. Although the extent of brain regions are smaller compared to Figure 1 (due to the conservative nature of conjoint analysis), we still observed significant voxels in the same brain areas. OFC, orbitofrontal cortex; ACC, anterior cingulate cortex. Lt, left; Rt, right. For all analyses, the statistical threshold is set at $P = 0.05$ corrected.

Table 1. Characteristics of parents and offspring

Characteristic: mean (SD)	Parent				Offspring			
	Mother	Father	T value	P Value	Daughter	Son	T value	P Value
n	30	29			19	20		
Age	41.3 (3.4)	42.4 (4.6)	-0.76	0.45	8.9 (1.3)	8.0 (2.3)	1.36	0.19
BASC-2 PRS subscales: mean T-score (SD)					(n = 10)	(n = 10)		0.127
Hyperactivity					51.4 (8.3)	49.8 (8.5)		
Aggression					51.5 (8.8)	56.9 (10.7)		
Conduct Problems					50.0 (6.9)	53.8 (4.9)		
Anxiety					51.5 (9.3)	53.3 (11.1)		
Depression					50.9 (8.4)	46.9 (19.1)		
Somatization					56.8 (14.2)	49.0 (11.7)		
Atypicality					51.2 (9.7)	50.9 (11.4)		
Withdrawal					52.1 (9.9)	61.3 (13.2)		
Attention Problems					48.8 (7.9)	51.9 (9.1)		
Adaptability					51.5 (8.5)	43.4 (7.3)		
Social Skills					54.5 (7.9)	42.5 (7.3)		
Leadership					54.5 (8.3)	49.6 (5.1)		
Activities of Daily Living					51.0 (7.7)	44.3 (5.3)		
Functional Communication					51.5 (8.7)	49.3 (7.5)		

Note: The table presents means with standard deviations (SD) in age and standardized T-scores for all BASC-2 PRS subscales. In the BASC-2 PRS subscales, a MANCOVA controlling for age demonstrated no significant sex difference for T-scores (Wilks' Lambda 0.079, $F_{(14, 4)} = 3.33$, $P = 0.127$). BASC-2, Behavior Assessment System for Children Second Edition; PRS, Parent Rating Scale.

Table 2. Brain regions showing significantly greater positive correlations in mother-daughter pairs compared to other parent-offspring pairs

Brain Region	MNI Coordinates			mm ³	Z score	P value
	X	Y	Z			
Mother-Daughter > Mother-Son						
Rt-Amygdala	32	-2	-18	320	2.13	0.016
Rt-OFC	30	27	-18	1320	2.10	0.017
Bi-Rectus gyrus	-4	30	-21	920	2.29	0.011
Rt-Rectus gyrus	7	52	-23	944	3.19	0.001
Rt-ACC	3	18	-10	72	2.22	0.013
Lt-ACC	-2	52	1	232	2.46	0.006
Rt-Hippocampus	37	-10	-16	368	2.35	0.009
Mother-Daughter > Father-Daughter						
Rt-Amygdala	24	2	-20	360	1.88	0.030
Rt-OFC	33	27	-17	2032	3.37	0.001
Lt-OFC	-15	12	-22	112	1.81	0.035
Bi-Rectus gyrus	-5	29	-22	1048	2.93	0.001
Rt-Rectus gyrus	13	19	-16	624	2.14	0.016
Lt-Rectus gyrus	-8	14	-19	616	1.98	0.023
Lt-ACC	-6	20	-8	72	2.11	0.017
Rt-Hippocampus	22	-6	-25	1280	2.13	0.016
Mom-Daughter > Father-Son						
Rt-Amygdala	26	0	-19	832	2.28	0.011
Rt-OFC	35	25	-20	1456	3.23	0.001
Bi-Rectus gyrus	0	30	-21	1784	1.94	0.026
Rt-Rectus gyrus	7	51	-23	736	2.93	0.001
Bi-ACC	-1	27	-7	1536	2.55	0.005
Rt-Hippocampus	24	-6	-18	944	1.71	0.043
Rt-Parahippocampus	20	-12	-28	256	2.39	0.008

Lt, Left; Rt, Right; Bi, Bilateral; OFC, orbitofrontal cortex; ACC, Anterior cingulate cortex.

Figure 1.

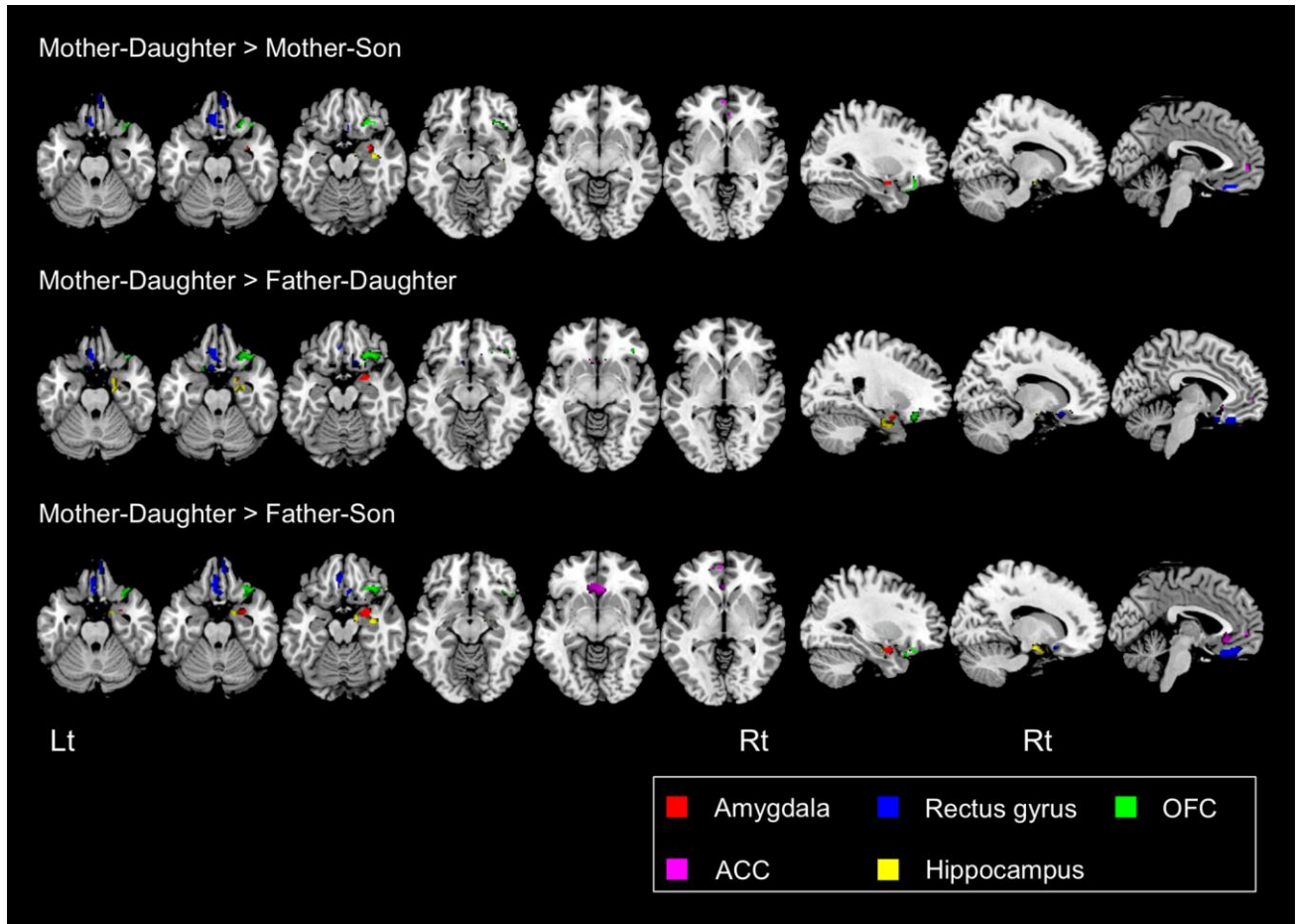


Figure 2.

