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Salivary Alpha Amylase-Cortisol Asymmetry in Maltreated Youth

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Abstract

Background—Maltreatment represents a major stressor in the lives of many youth. Given the known effects of stress exposure on subsequent biological stress response systems, researchers have been interested in the effects of maltreatment on the functioning of these systems. Experimental studies reveal that previous exposure to stress affects the symmetry between components of the physiological stress response to subsequent stress. The present study examined asymmetry between salivary alpha amylase (sAA), a sympathetic indicator, and cortisol reactivity to a social stressor among maltreated and comparison youth age 9 to 14 years. Consistent with earlier studies suggesting that stress leads to asymmetry between hypothalamic-pituitary-adrenal axis and sympathetic nervous system activity, we expected that maltreated youth would exhibit greater sAA-cortisol asymmetry than would comparison youth.

Methods—Forty-seven maltreated and 37 comparison youth visited the lab and engaged in a social stress protocol. We collected 2 saliva samples before the stressor and 4 after, at 0 minutes post stress and every 10 minutes for 30 minutes.

Results—Maltreatment status moderated the relation between sAA and cortisol activity in response to the stressor. Comparison youth showed significant links between the sAA and cortisol responses; maltreated youth had no significant associations between responses in the two biomarkers.

Conclusion—The data were consistent with sAA-cortisol asymmetry among maltreated youth. Further research should seek to replicate this finding and investigate its implication for developmental trajectories.

The present study examined the relation between child maltreatment and asymmetry between salivary alpha-amylase (sAA) and cortisol responses to stress. Given connections between the HPA axis and the SNS, one would anticipate symmetry between these systems, meaning that relatively strong responses in one system should correspond with relatively strong responses in the other. Recently researchers have suggested that the degree of asymmetry between

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biological stress systems may have implications for developmental outcomes (Bauer, Quas, and Boyce, 2002). Factors affecting degree of asymmetry between these biological stress systems remain poorly understood.

The biological stress response consists primarily of two systems: The locus cereleus-norepinephrine/sympathetic nervous system (LC-NE/SNS) and the hypothalamic-pituitary-adrenal (HPA) axis (Chrousos and Gold, 1992). The LC-NE/SNS is responsible for effects often referred to as the “fight or flight” response, which include increased cardiovascular tone, faster breathing rate, and increased blood flow to muscles (Cannon, 1914). The HPA axis involves multiple steps, including the release of corticotropin releasing hormone (CRH) from the hypothalamus, triggering the release of adrenocorticotrophic hormone (ACTH) from the anterior pituitary, causing the release of the steroid hormone cortisol from the adrenal glands.

The negative impact of maltreatment on psychological/behavioral development may be at least partially due to an altered stress response. Childhood trauma in general and maltreatment in particular are linked with alterations in both the HPA axis and in the LC-NE/SNS. Maltreated youth have demonstrated reduced stress reactivity in the HPA axis (Hart, Gunnar, and Cicchetti, 1995). One theory is that chronic stress leads to attenuation of the stress response (Goldstein & McEwen, 2002; Susman, 2006). Though few studies have examined the effect of maltreatment on the functioning of the LC-NE/SNS among children, some researchers report elevated urinary catecholamine levels among maltreated and traumatized children (De Bellis, Baum, Birmaher, et al. 1999; De Bellis, Lefter, Trickett, and Putnam, 1994). Thus, some research links maltreatment to alterations of both the HPA axis and the LC-NE/SNS, though the specific nature of the impact of maltreatment on stress reactivity remains unclear.

The HPA axis and the SNS are connected at multiple neural levels, and thus activity in these two systems should demonstrate some degree of symmetry (Chrousos and Gold, 1992; Lovallo and Thomas, 2000; Young, Abelson, and Cameron, 2005). Factors leading to individual differences in the degree of symmetry between these systems remain unclear. One possible factor may be different habituation rates of these systems given previous exposure to stress. Some evidence suggests faster habituation rates in the HPA axis (Gerra, Zaimovic, and Mascetti, 2001; Schommer, Hellhammer, and Kirschbaum, 2003). Schommer et al. (2003) and Gerra et al. (2001) both found decreased ACTH and cortisol responses over successive repetitions of the Trier Social Stress Test (TSST; Kirschbaum, Pirke, and Hellhammer, 1993), but unchanged responses in level of epinephrine and NE. Animal research also suggests that prolonged stress may lead to asymmetry between these systems (Britton, Segal, Kuczynski, and Hauger, 1992).

Whereas experimental manipulations suggest that repeated stressors may lead to asymmetry between the HPA axis and the SNS, we know little about the effects of real life stressors on the symmetry between activity in these two systems in response to subsequent stress. If previous exposure to stress leads to asymmetry between the systems, we might expect individuals with a history of significant stress to show more asymmetry. This asymmetry may be important for understanding the impact of stress on atypical development. Recently, researchers have argued that examination of each stress system alone limits what we know about the role of the stress response in atypical development (Bauer et al., 2002). Each system independently has been linked to developmental trajectories in psychological and behavioral characteristics. Activity in the HPA axis covaries with aggression, disruptive behavior, depression, and inhibited behavior (e.g., De Souza, 1995; Duval et al., 2006; Shirtcliff et al., 2005; Shoal et al., 2003). Autonomic arousal has been associated with aggression, delinquency, and antisocial behavior (Lorber, 2004; Ortiz and Raine, 2004). Bauer et al. suggest that beyond relations with each system independently, we will learn more about the role of the biological

stress response in mediating developmental outcomes if we consider the functioning of these systems together and examine the relations between them.

Examination of the HPA axis is long-established through salivary cortisol (Kirschbaum, Bartussek, and Strasburger, 1992). Recently, a marker of autonomic activity has been possible via salivary alpha-amylase (sAA), an enzyme released by the parotid gland (Granger, Kivlighan, El-Sheikh, Gordis, and Stroud, 2007). Protein release into the saliva is primarily controlled by sympathetic arousal, and in particular by beta-adrenergic activity (Bosch, de Geus, Veerman, Hoogstraten, and Amerongen, 2003). In addition, parasympathetic activity appears to augment release of salivary proteins, including sAA, due to the effect of parasympathetic activity on salivary flow rate (Bosch et al., 2003; Gjørstrup, 1979). Animal studies have demonstrated that stimulation of the sympathetic system leads to increased secretion of sAA in the saliva in the presence of parasympathetic activity (Gjørstrup, 1979). In humans, sAA increases in response to stress (Bosch et al., 2003; Nater, Rohleder, Gaab, et al., 2005; Rohleder, Nater, Wolf, Ehlert, and Kirschbaum, 2004). Some research shows that sAA correlates with plasma NE (Chatterton, Vogrldon, Ellman, and Hudgens, 1996), though this link has not been completely consistent (Nater, La Marca, Florin, et al., 2006). Stress-dependent release of sAA correlates with cardiovascular autonomic measures (Bosch et al., 2003). Beta adrenergic antagonist propranolol leads to lower levels of sAA at rest and in response to a stressor (van Stegeren, Rohleder, and Everaerd, 2006). Thus, saliva can provide indices of both SNS and HPA axis activity.

The present study examined the symmetry between HPA and SNS activity in maltreated and comparison adolescents. We expected that indices of HPA and SNS activity in response to stress would correlate positively among youth with no history of maltreatment. Consistent with the idea that stress may cause asymmetry between the systems, we expected that maltreated youth would demonstrate lower correlations between cortisol and sAA indices, and that maltreatment status would moderate the link between cortisol and sAA response to stress.

Materials and Methods

Participants

Eighty-four adolescents were selected from a larger, longitudinal study of the effects of maltreatment on adolescent development. Youth visited the lab with a parent or guardian for the first or second of three yearly interviews. Maltreated youth were recruited from select zip codes in a large urban area on the basis of recent (within the previous 2 months) referrals to the Department of Children and Family Services (DCFS) for child maltreatment (physical or sexual abuse or neglect). Comparison youth were recruited from the same or comparable census blocks. We excluded youth who used medications that would interfere with cortisol or sAA measures (e.g., synthetic corticosteroids or beta adrenergic agonists) or who reported smoking. In addition, based on parents'/guardians' reports, no child had a history of cancer, diabetes, or thyroid problems. In the final sample of 84 adolescents, the mean age was 12.1 years ($SD = 1.2$, range = 9.1-14.5). The maltreated group included 47 youth (26 male), and the comparison group included 37 youth (18 male). Demographic information regarding the sample appears in Table 1. Ethnic composition did not differ by maltreatment status. Among the maltreated youth, 50% had been referred to DCFS for physical abuse, 34.2 for sexual abuse, 31.5 for emotional abuse, and 63.2 for neglect. Consistent with these percentages, most youth (78.9%) had more than one referral to DCFS (mean number of referrals = 3.8, $SD = 2.7$, range = 1-12, median = 3).

Procedures

All procedures were approved by the university institutional review board and were carried out with the written understanding and assent of the adolescent participants and consent of their parents or guardians. Each adolescent and a parent or guardian attended a lab session lasting approximately 4 hours. As part of the procedures, adolescents engaged in a modified version of the TSST (Kirschbaum et al., 1993). During this procedure, the interviewer and a panel of 2 judges entered the room. The interviewer told the participant that he or she would read the beginning of a story, and the participant would have 5 minutes to develop the next part of the story, after which the participant would present the next part of the story to the panel of judges for 4 minutes. The interviewer and judges then left the room for 5 minutes. When they re-entered, the interviewer asked the adolescent to present the story. If the youth did not fill the 4 minutes, the interviewer used a standard set of prompts to encourage him or her to continue. After the 4 minutes, the interviewer asked the youth to perform a serial subtraction task before the judges. This task was designed to be challenging for the particular age of the adolescent (e.g., subtracting 7 serially beginning with 758).

We collected saliva samples 6 times, twice before and 4 times after the TSST. The first sample occurred 45 minutes before the stressor. The second sample was 10 minutes before the stressor, immediately after a 5 minute relaxation protocol involving listening to soft music while viewing a still slide of a beach scene. The third sample occurred immediately after the stressor, and the fourth, fifth, and sixth samples occurred 10, 20, and 30 minutes after the end of the stressor, respectively. We collected saliva samples via passive drool through a short straw into a vial. Data collection occurred primarily in the afternoon, with 96% of participants providing the baseline sample between 1200h and 1700h. The earliest baseline sample was collected at 1223h, and the latest was collected at 1735h. We control for time of day in regression analyses. Saliva samples were immediately frozen.

Samples were transported on ice to Penn State University and stored frozen at -80°C until assayed for cortisol and sAA. On the day of testing, all samples were centrifuged at 3000 rpm for 15 minutes to remove mucins.

α -Amylase—Samples were assayed for sAA using a commercially available kinetic reaction assay (Salimetrics, State College PA). The assay employs a chromagenic substrate, 2-chloro-p-nitrophenol, linked to maltotriose. The enzymatic action of sAA on this substrate yields 2-chloro-p-nitrophenol, which can be spectrophotometrically measured at 405 nm using a standard laboratory plate reader. The amount of α -amylase activity present in the sample is directly proportional to the increase (over a 2 minute period) in absorbance at 405 nm. Results are computed in U/mL of α -amylase using the formula: [Absorbance difference per minute \times total assay volume (328 ml) \times dilution factor (200)] / [millimolar absorptivity of 2-chloro-p-nitrophenol (12.9) \times sample volume (.008 ml) \times light path (.97)]. Intrassay variation (CV) computed for the mean of 30 replicate tests was less than 7.5%. Inter-assay variation computed for the mean of average duplicates for 16 separate runs was less than 6%.

Cortisol—Samples were assayed for salivary cortisol using a highly-sensitive enzyme immunoassay US FDA (510k) cleared for use as an in vitro diagnostic measure of adrenal function (Salimetrics, State College, PA). The test used 25 μ l of saliva (for singlet determinations), had a lower limit of sensitivity of .007 μ g/dl (.19 nmol/L), range of sensitivity from .007 to 1.8 μ g/dl (.19 to 49.7 nmol/L), and average intra- and inter-assay coefficients of variation of less than 5% and 10%, respectively.

Statistical Analyses

We conducted data analyses using SPSS statistical analysis software (version 15.0 for Windows). We first examined the effect of the TSST on cortisol and sAA levels. We conducted repeated measures ANCOVAs to examine whether level of cortisol or sAA varied by saliva sample. We examined the within subjects effect of saliva sample using Wilks' lambda, as suggested by Stevens (1979). Because the data violated sphericity assumptions, we used multivariate tests, which do not require sphericity, unlike the univariate tests. Covariates included maltreatment status, time of day, sex, and pubertal stage, assessed via self- and parent/guardian-reported Tanner staging (Marshall and Tanner, 1969; 1970), which has been shown to demonstrate reasonable agreement with nurse practitioner ratings (Dorn, Susman, Nottelmann, Inoff-Germain, and Chrousos, 1990). In addition, we found higher sAA levels among African-American youth vs. Latino/Latina and Caucasian youth. Thus, we also controlled for ethnicity (African-American vs. Latino/Latina and White) in the regression analyses, though this did not affect the substantive findings.

Because sAA and cortisol show a time differential in response to stressors, we examined several indices of salivary cortisol and sAA. Thus, to examine whether overall HPA and autonomic activity and relative response to stress are correlated between the two systems, we calculated area under the curve scores with respect to ground (AUC_G) and with respect to increase (AUC_I) according to formulas provided by Pruessner, Kirschbaum, Meinlschmid, and Hellhammer (2003). The AUC_G and AUC_I allow examination of summary scores over time, reflecting total output and total response to the stressor, respectively. These scores are based on trapezoid formulas. Specifically, AUC_G is calculated by summing the areas of the trapezoids under the plot of the biomarker level at each time point for each case. AUC_I is calculated by subtracting the rectangular area representing baseline from the AUC_G score. We also examined baseline levels, peak values, and raw change from baseline to peak in both systems.

To examine whether the data are consistent with maltreatment status as a moderator of the symmetry between responses of cortisol and sAA, we conducted multiple regression analyses. Specifically, for each index of cortisol and sAA, we examined whether the interaction between maltreatment status and the sAA index accounted for variance in the cortisol index after including covariates and main effects of sAA and maltreatment status. Covariates included time of day, wave of visit, sex, pubertal stage, and ethnicity. We regressed each cortisol index onto the centered sAA index, maltreatment (-1 = comparison, +1 = maltreated), and the interaction between maltreatment and sAA, after controlling for covariates. To address positive skew in cortisol indexes (skew = 3.5, 2.0, and 1.6, respectively for baseline, peak and AUC_G ; $SD = .27$), we conducted log transformations of cortisol baseline, peak and AUC_G . We conducted square root transformations of sAA baseline, peak, and AUC_G as these indexes were moderately skewed (skew = 1.1, .8, and .8, respectively; $SD = .27$). In addition, to eliminate multivariate outliers, we eliminated cases with centered leverage values greater than $3k/n$ (where k = the number of predictors and n = the sample size) for each equation, as suggested by Cohen, Cohen, West, and Aiken (2003) for regression analyses with small samples. Elimination of multivariate outliers resulted in dropping one case in the equation predicting change in cortisol and two cases in predicting cortisol AUC_I . To understand significant interaction effects, we compared correlation coefficients in the separate groups using Fisher's R transformation to Z .

Results

Response curves among maltreated and comparison youth appear in Figure 1. Cortisol and sAA have different response times, with cortisol responding to and recovering from stress more gradually, and sAA responding more immediately and recovering more quickly. For sAA, repeated measures analysis of variance revealed a significant within subjects main effect for

saliva sample (Wilks' $\lambda = .83$, Exact $F(5,74) = 2.99$, $p < .05$). Levels were lowest at pre-relaxation, increased after relaxation, peaked immediately post stressor, and returned to baseline levels by 10 minutes post stressor. A majority of the sample, 63.1%, had greater than 10% increases in sAA from baseline to post-stress peak.

Repeated measures analysis of variance revealed an effect for time of sample on cortisol as well. Examination of the within subjects effect of saliva sample across all 6 samples revealed a marginally significant effect for saliva sample (Wilks' $\lambda = .88$, Exact $F(5,74) = 1.97$, $p < .10$). A follow-up repeated measures ANCOVA examining change over time only during the linear rise during from pre-relaxation to 10 minutes post stressor, when responses peaked on average, revealed a significant effect for time (Wilks' $\lambda = .89$, Exact $F(3,76) = 3.17$, $p < .05$). Cortisol peaked at 10 minutes post-stressor and returned to baseline levels at 20 minutes post-stressor. Similar to sAA, 64.5% of the sample had cortisol increases of great than 10% from baseline to post-stress peak.

Regression analyses reveal that in equations predicting cortisol AUC_I, baseline, peak, and raw change scores, the maltreatment status by corresponding sAA index interaction was significant, consistent with the hypothesis that maltreatment status moderates the relation between cortisol and sAA activity. In the equation predicting cortisol AUC_G, the interaction term was marginally significant. Correlations among biomarker indices across the whole sample of youth appear in Table 2. Several corresponding cortisol and sAA indices are correlated across the sample, including AUC_I, baseline, and change. For each biomarker, the different indices are correlated with each other. Regression analyses appear in Table 3. Table 4 presents correlations between corresponding cortisol and sAA indexes among maltreated and comparison youth. Consistent with the hypothesis, among the comparison youth, the indexes are significantly correlated, whereas among the maltreated youth the correlations are not significant. Comparison of these correlation coefficients reveals that they are significantly different between the two groups. To illustrate the interaction, scatterplots of the relation between peak cortisol and peak sAA among maltreated and comparison youth appear in Figure 2. In these plots, peak cortisol is log transformed and peak sAA is square root transformed.

Discussion

The results of this study suggest that among comparison youth, sAA and cortisol responses to stress demonstrate symmetry, whereas among maltreated youth, they demonstrate asymmetry. Among comparison youth, sAA baseline and peak levels, AUC scores with respect to both ground and increase, and change from baseline to peak are correlated with corresponding indexes in cortisol. The correlations suggest that among comparison youth, the magnitude of the response to stress in the ANS covaries with the magnitude of response in the HPA axis. In contrast, among maltreated youth, responses across the two systems are uncorrelated, suggesting asymmetry. Regression analyses reveal significant or marginally significant sAA index by maltreatment status interactions in predicting the corresponding cortisol index, consistent with the hypothesis that asymmetry between cortisol and sAA response is moderated by maltreatment status. Whereas the overall activity of the two systems in response to the TSST appears to demonstrate some symmetry, as indicated by the correlations between these systems in the whole sample, the maltreatment by sAA interactions and follow-up correlations reveal that the cortisol and sAA indexes were correlated only in the comparison group. Among the maltreated adolescents, the systems demonstrate asymmetry. These data are consistent with the notion that maltreatment may cause asymmetry between these systems, possibly by interrupting the connections between them or by causing attenuation in one system but not the other (Ressler and Nemeroff, 2000).

Given the connections between the two systems, symmetry between their activity makes sense. For example, the lateral paraventricular nucleus (PVN) connects to areas in the hindbrain responsible for sympathetic activity via neurons that secrete CRH (Chrousos and Gold, 1992). Catecholaminergic pathways project from the LC to the PVN (Brown et al., 1982; Chrousos and Gold, 1992; Dunn and Berridge, 1990). Experimental manipulations support the interconnectedness of these systems. For example, administration of CRH increases catecholaminergic activity and norepinephrine (NE) levels (Brown, Fisher, Speiss, Rivier, and Vale, 1982; Dunn and Berridge, 1990). CRH antagonists cause reduced responsiveness in the LC-NE system (Dunn and Berridge, 1990). NE stimulates the release of CRH in the PVN (Brown et al., 1982; Dunn and Berridge, 1990), and β -adrenergic blockers reduce the behavioral consequences of CRH (Chrousos and Gold, 1992; Dunn and Berridge, 1990).

Despite these connections, the two systems may be activated by different kinds of stressors and may respond to stress differently over time. Some research suggests that CRH pathways function differently in responding to metabolic versus psychological stressors, and thus that CRH may affect the autonomic nervous system (ANS) under conditions of metabolic but not psychological stress (Malarkey, Lipkus, and Cacioppo, 1995). Moreover, the HPA axis may be particularly sensitive to fear and frustration, whereas the ANS may be more generally responsive (Frankenhaeuser, 1982; Lovallo and Thomas, 2000).

Chronic stress may have different implications for the HPA axis and ANS and may thus affect the degree of symmetry between them. Chronically elevated levels of cortisol and CRH may promote long term changes in the functioning of CRH in response to stress (Tarullo and Gunnar, 2006). Given connections between CRH and the ANS, these changes may have implications for the relation between the HPA axis and the ANS. Experimental manipulations have revealed that repeated stress leads to asymmetry between the two systems. Some researchers report attenuated responses in the HPA axis to experimental manipulations but not in the LC-NE/SNS (Gerra et al., 2001; Schommer et al., 2003). In our sample, peak cortisol levels in the maltreated group were marginally lower than peak cortisol levels in the comparison group ($F(1,78) = 3.40, p = .07$), whereas no such difference emerged in sAA. In contrast, some researchers have found a decreased response over time in the SNS, with sustained HPA axis responding (Britton et al., 1992). Beyond effects of lab-based stressed exposure on these stress response systems, different types of psychopathology that are linked with stress and maltreatment may relate to different kinds of alterations in the stress response systems (Gunnar and Quevedo, 2007). Understanding the role that alterations in both stress response systems and their degree of symmetry play in the link between maltreatment and different forms of psychopathology is an important future direction.

Though our data do not clarify the cause of asymmetry between the systems or the role of CRH in each of these systems, the asymmetry of these systems among the maltreated youth may be due to a reduced response in the HPA axis but not in the LC-NE/SNS. A reduced responsiveness to stress among maltreated youth would be consistent with the hypothesis that stress systems might attenuate in responsivity due to allostatic load (Goldstein and McEwen, 2002; Susman, 2006). As Susman notes, a downregulation of the stress response would protect the individual from chronic energy expenditure and arousal in the face of repeated stressors. Given the potential harmful effects of chronically high levels of cortisol, an attenuation in the responsiveness of the HPA axis in the face of chronic stress may be protective (Fries, Hesse, Hellhammer and Hellhammer, 2005; Goldstein & McEwen, 2002; Sapolsky, 2004). Consistent with lab based studies (Gerra et al. 2001; Schommer et al., 2003), data consistent with such attenuation occurred in the HPA axis but not the LC-NE/SNS, which may be more sensitive to certain types of arousal and more resistant to attenuation.

Replication of these findings will be important before we can understand the implications of the asymmetry for maltreated youth. Different profiles of symmetry or asymmetry in HPA axis/SNS activity may have different implications for behavioral outcomes (Bauer et al., 2002). For example low or high activity in both systems may put youth at risk for aggression or anxiety, respectively. Further studies should examine whether asymmetry is a risk factor for negative outcomes. Conversely, asymmetry may protect youth who have extremely high or low reactivity in one system. A moderate response in one system may serve a protective function among youth with extremely high or low reactivity in the other stress response system.

The study has several limitations. First, the small sample size and cross-sectional design of the study limit power to test interactions and differences in effect sizes between groups. In addition, cortisol and sAA response curves to stress are timed differently. Thus, examination of their asymmetry must consider response over time. The AUC ground and change scores reflect ways to examine the relation between the sum totals of each system in response to stress. The small sample size limits our ability to examine asymmetry regarding other characteristics of the response curves over time in cortisol and sAA. The sample size also precludes our examining whether psychopathology plays a role in HPA and SNS patterns of responding to stressors. The non-experimental, cross-sectional design precludes clarification of the causal relationships between maltreatment and asymmetry between these two stress responses systems. We also did not conduct physical examinations of the youth to rule out potentially confounding medical conditions, although we did obtain some information regarding medical history and medication use.

Nonetheless, this study is new in examining the effect of maltreatment on asymmetry between the SNS and HPA axis in adolescents. Maltreatment is linked with alterations in both of these biological response systems and with a variety of negative outcomes. Researchers recently have suggested that the pattern of asymmetry between the HPA axis and SNS may be related to atypical development (Bauer et al., 2002). Thus, the role of maltreatment in asymmetry may have relevance for the links between maltreatment as a major life stressor and youth outcomes.

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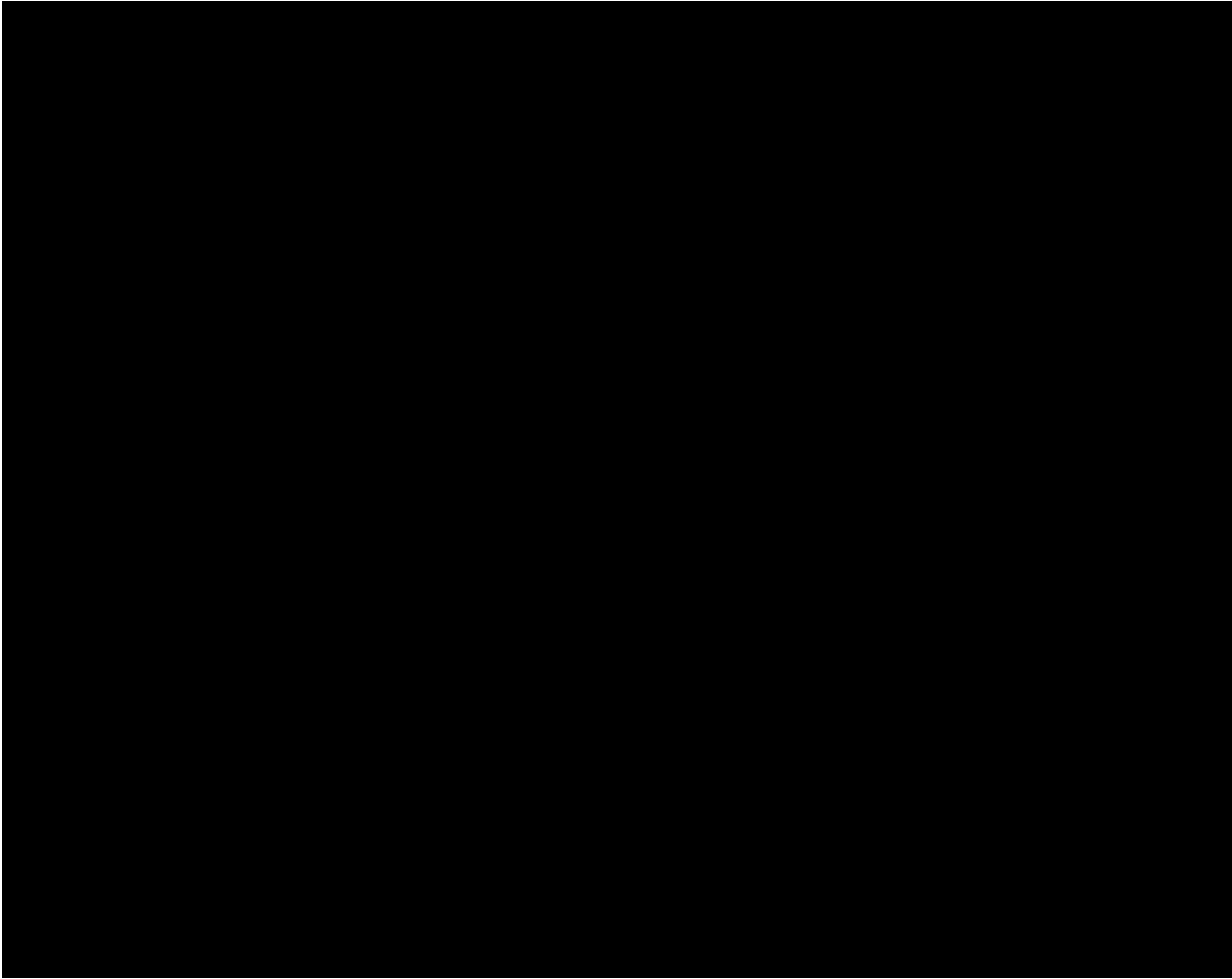


Figure 1. Cortisol and sAA at Pre Relaxation (T = -45), Pre Stress (T = -10), and at 0 (T = +15), 10 (T = +25), 20 (T = +35), and 30 (T = +45) Min Post Stress in Maltreated and Comparison Youth
Note: For sAA, repeated measures ANCOVA (within subjects factor saliva sample controlling for group, time of day, pubertal status, and ethnicity) Wilks' $\lambda = .83$, Exact $F(5,74) = 2.99$, $p < .05$). Levels were lowest at pre-relaxation, increased after relaxation, peaked immediately post stressor, and returned to baseline levels by 10 minutes post stressor. For cortisol, repeated measures ANCOVA (within subjects factor saliva sample controlling for group, time of day, pubertal status, and ethnicity) Wilks' $\lambda = .88$, Exact $F(5,74) = 1.97$, $p < .10$). Cortisol peaked at 10 minutes post-stressor and returned to baseline levels at 20 minutes post-stressor.

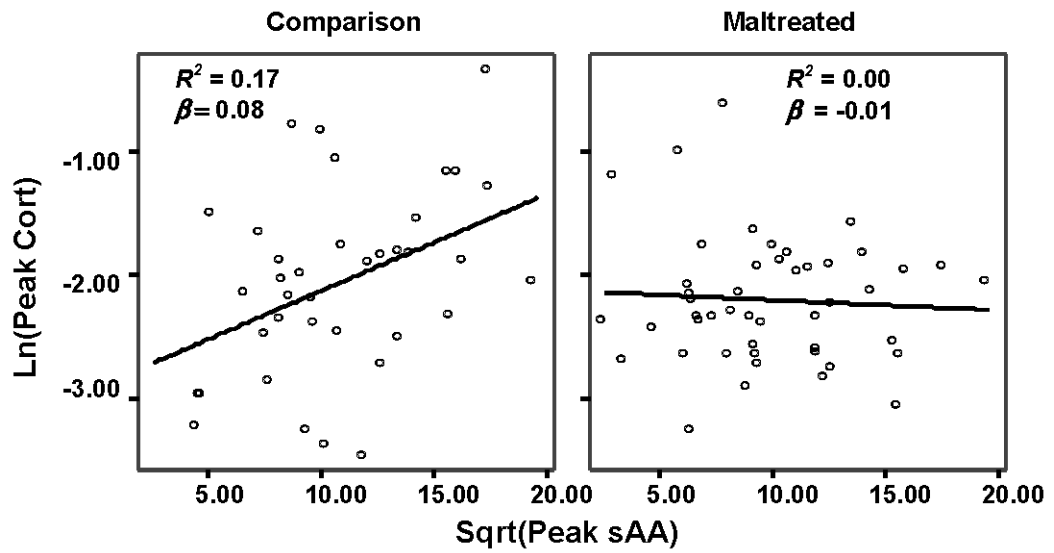


Figure 2. Scatterplots of the Relation between Peak Cortisol and Peak sAA among Maltreated and Comparison Youth

Note: Fitted lines reflect linear regression lines. Peak cortisol is log transformed. Peak sAA is square root transformed.

Table 1

Demographic Information

	Comparison	Maltreated
Yearly Family Income		
Less than \$30,000	17	34
\$30,000-60,000	13	8
More than \$60,000	7	5
Parent's Education		
Less than high school	4	22
High School Diploma or GED	11	9
Some College	9	12
College degree or advanced degree	13	4
Ethnicity		
Black	45.9%	38.3%
Caucasian	8.1	12.8
Latino/Latina	37.8	40.4
Other/Multiple ethnicities	8.1	8.5
Parent/Guardian		
Biological or Adoptive Parent	34	26
Step-parent	1	3
Other family member	2	9
Foster parent	0	9

Table 2

Partial Correlations Among sAA and Cortisol Indices

	1	2	3	4	5	6	7	8	9
1. sAA-AUC _G	--								
2. sAA-AUC ₁	.31**	--							
3. Baseline sAA	.74**	-.38**							
4. Peak sAA	.96**	.38**	--						
5. sAA Change	.43	.92**	.64**	--					
6. Cort-AUC _G	.15	-.07	.20 ⁺	.58**	--				
7. Cort-AUC ₁	.07	.30*	-.10	.15	-.00	--			
8. Baseline Cort	.05	-.26*	.23*	.01	.36**	.10	--		
9. Peak Cort	.11	.01	.13	.14	-.23*	.77**	-.46**	--	
10. Cort Change	.05	.19 ⁺	-.06	.13	.08	.86**	.34**	.46**	--
					.27*	.28*	.89**	-.29**	.61**

Note: *df* = 76 for all coefficients. Coefficients control for maltreatment status, time of day, ethnicity, pubertal status, sex, and wave of visit. sAA = Salivary alpha-amylase. Cort = Cortisol. AUCG = Area under the curve with respect to ground. AUCI = Area under the curve with respect to increase.

* $p < .05$.

** $p < .01$.

+ $p < .10$.

Table 3
Regression Equations Predicting Cortisol from sAA, Maltreatment, and Maltreatment by sAA Interactions

	β	T	F	$Adj R^2$
Cortisol AUC _I				
sAA AUC _I	.42	3.91**		
Maltreatment status	-.14	-1.25		
sAA AUC _I by Maltreatment Status	-.42	-3.88**		
Equation			8.49 ^{a**}	.25
Cortisol AUC _G				
sAA AUC _G	.18	1.59		
Maltreatment status	-.19	-1.60		
Salivary A-A AUC _G by Maltreatment Status	-.21	-1.96 ⁺		
Equation			3.04 ^{b*}	.10
Baseline Cortisol				
Baseline sAA	.33	2.79**		
Maltreatment Status	.04	.35		
Baseline sAA by Maltreatment Status	-.32	-2.80**		
Equation			4.12 ^{b**}	.14
Peak Cortisol				
Peak sAA	.20	1.69		
Maltreatment Status	-.17	-1.49		
Peak sAA by Maltreatment Status	-.24	-2.26*		
Equation			3.41 ^{b*}	.10
Cortisol Change				
sAA Change	.29	2.70*		
Maltreatment status	-.09	-.78**		
sAA Change by Maltreatment Status	-.33	-3.14 ^{c**}		
Equation			6.43 ^{c**}	.19

Note: Equations control for pubertal status, sex, time of day, ethnicity (African-American vs. Latino/Latina or white) and wave of visit. To reduce positive skew, cortisol AUC_G was subjected to log transformation, and sAA AUC_G was subjected to square root transformation. Predictors were centered.

Maltreatment status coded as -1 = comparison and +1 = maltreated. sAA and cortisol change are raw change from baseline to post-stress peak levels.

* $p < .05$.

** $p < .01$.

⁺ $p < .10$.

^a $df = 3,73$

^b $df = 3,75$

^c $df = 3,74$

Table 4
Correlations between corresponding cortisol and sAA indices

	Comparison (n = 37)	Maltreated (n = 47)
AUC _I	.48 ^{a**}	.11 ^a
AUC _G	.37 ^{a*}	-.04 ^a
Baseline	.44 ^{a**}	-.01 ^a
Peak	.41 ^{a*}	-.06 ^a
Change (Baseline to Peak)	.51 ^{b**}	.01 ^b

AUC_I = Area under the curve with respect to increase.

AUC_G = Area under the curve with respect to ground.

*
 $p < .05$.

**
 $p < .01$.

^a coefficients sharing this subscript differ at $p < .05$ (one-tailed).

^b coefficients sharing this subscript differ at $p < .01$ (one-tailed).