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Gene-Environment Interaction and the Intergenerational Transmission of Parenting: Testing the Differential-Susceptibility Hypothesis

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Abstract The current study evaluated the differential-susceptibility hypothesis in explaining the intergenerational transmission of parenting, using data from the National Longitudinal Study of Adolescent Health (Add Health). Exposure to maternal parenting was measured prospectively when respondents were adolescents and parental stress was measured when they were parents themselves, some 14 years later, on average. Cumulative-genetic plasticity was measured by dominantly coding the presence of putative plasticity alleles from four genes: the 10R allele of DAT1, the A1 allele of DRD2, the 7R allele of DRD4, and the short allele of 5HTTLPR. Results showed that the more plasticity alleles individuals carried (range 0–4), the more that parenting experienced in adolescence predicted future parenting experience. Those respondents with the most plasticity alleles not only experienced the highest levels of parental stress when exposed to negative maternal parenting in adolescence but the lowest levels when exposed to positive maternal parenting in adolescence. These results indicate that differential susceptibility is operative in the case of the intergenerational transmission of parenting, which could explain why estimates of such transmission have proven so modest in studies which fail to consider GXE interactions.

Keywords Differential susceptibility · Genetics · Intergenerational · Parenting

Introduction

The question of why parents parent the way they do is one that has interested developmental scholars and lay persons for quite some time [7]. Although it is widely appreciated that parenting is multiply determined by individual, child and contextual factors [11],

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perhaps the pre-eminent hypothesis is that how parents were themselves treated as children shapes the ways in which they care for their progeny [60]. Indeed, a variety of theoretical perspectives embrace the view that parenting is intergenerationally transmitted, including life-course [35], attachment [19] and social-learning [3, 53] theories, even if they differ in terms of mechanisms presumed to account for the intergenerational transmission process.

There exists extensive evidence consistent with the claim that that parenting is transmitted across generations. Perhaps the earliest and best known is that pertaining to child maltreatment [5, 6, 26, 62], though much of the relevant data base has been called into question due to reliance on retrospective reports by adults of their rearing experiences; and this is because adults' recollections of how they were treated in their families of origin are prone to memory errors of omission (i.e., not remembering) and commission (i.e., falsely remembering), as well as to distortion based on current life circumstances (e.g., [6, 8, 40, 65]).

Fortunately, longitudinal studies not reliant on adult recall provide evidence that harsh parenting or high levels of family discord are intergenerationally transmitted (e.g., [23, 44, 50, 55]). Oregon Youth Study investigators studying boys growing up in the highest crime-rate areas of a medium-sized city from age nine followed them up approximately 12 years later, observing that the more these fathers, as boys, experienced poor parental supervision and harsh discipline, the more they provided their toddlers with harsh, inconsistent discipline [22]. Conger and associates [29] obtained similar results for angry and aggressive parenting behavior in a subsample of rural Iowan adolescents followed up 5–7 years later when some of them had become parents. And drawing on a sample from high-risk Seattle neighborhoods, Bailey and colleagues [1] found that both monitoring and harsh discipline experienced around 13.5 years of age predicted the same kind of parenting some 14 years later when children were adults rearing nine-year olds.

It is not just angry-aggressive-hostile parenting, however, that seems to be intergenerationally transmitted. Chen and Kaplan [25] observed that in a large, random sample of 13-year-old Houston school children who were re-contacted in their 30s, the experience of good parenting in early adolescence, defined in terms of consistent discipline and parental acceptance, predicted the provision of constructive parenting in adulthood (i.e., monitoring, communication, involvement, positive affection, inductive discipline). Similar results emerged in a study of adolescent girls (age 13–18) growing up in Rochester, NY whose parenting was studied when they were 20–22 years old [64]. More recently, Belsky and associates [14] observed the parenting of New Zealand women followed since age three as they interacted with their own 3-year olds. These mothers were more likely to behave in a warm, sensitive, stimulating manner if, during early childhood, their own mothers did not hold authoritarian childrearing attitudes; if, during the middle-childhood years, the emotional climate of the family was marked by cohesion, positive expressiveness, and low levels of conflict; and/or if, during the early adolescent years, they experienced a trusting, openly communicative and non-alienated relationship with their parents.

A very recent set of reports published in a special section of the journal *Developmental Psychology* sought to illuminate the mediators of the intergenerational transmission process (e.g., [49]). Studying a normative sample beginning when children were 10 years old and then as parents some two decades later, Schaffer et al. [59] observed not only that parenting quality was intergenerationally transmitted, but that this process could be at least partially accounted for by the social competence which the children manifested in young adulthood before becoming parents. Relatedly, Neppl and associates [51] reported, after finding that the harsh and positive parenting that rural Iowan adolescents experienced predicted, respectively, the harsh and positive parenting they provided to their own

preschool children, that whereas externalizing problems mediated the intergenerational transmission of harsh parenting, it was academic attainment which mediated the cross-generational linkage of positive parenting.

Despite long-standing evidence that parenting is intergenerationally transmitted, including recent work addressing mechanisms of influence, it remains indisputable that the parenting experienced in one generation is by no means inevitably repeated in the next [16, 48]. In fact, several recent studies indicate that only about 15% of the variance in parenting can be accounted for by one's childrearing history [14, 22, 29], complementing earlier work chronicling a great deal of discontinuity in child maltreatment across generations (e.g., [34, 45]). These observations raise the question of why some parents do and some do not repeat the parenting they experienced while growing up. Although the editors of the aforementioned special section of *Developmental Psychology* bemoaned the lack of submissions addressing this fundamental concern [16, 30], there exists research investigating this topic (e.g., [13, 34]) discovered, for example, that mistreated children who did not mistreat their own offspring experienced supportive close relationships somewhere along their life-course journey, some with a therapist, others with a romantic partner, including a spouse. Quinton and Rutter [55, 56] called attention to relationship factors as well when it came to accounting for why some English girls whose experience in dysfunctional families which resulted in them growing up in residential institutions parented so much better than their own parents, whereas others did not.

Not considered to date, but central to the research reported herein, is the prospect that genetic factors might play a role in determining which individuals do and do not sustain parenting across generations, a possibility raised by Conger et al. [30] when highlighting the need for additional research on moderators of the intergenerational transmission of parenting. Here we test the gene-X-environment (GXE) interaction hypothesis that individuals carrying more of what Belsky and associates [12, 16] have hypothesized to be "plasticity alleles" prove to be those most likely to have their parenting experience influenced by the parenting they experienced while growing up. Important to appreciate is that the present work examining GXE interaction involving effects of mothering during adolescence on parenting stress in adulthood diverges from most prior GXE work in two fundamental respects. Whereas most GXE research to date is based on the diathesis-stress view that some individuals are more vulnerable than others to the negative effects of contextual adversity [12, 67], the work reported herein is based on the differential-susceptibility hypothesis [9, 10]. This stipulates that not only are certain individuals, often for genetic reasons [12], more prone to function poorly (e.g., become depressed) when confronted with stressful conditions (e.g., negative life events), but that the very same putatively "vulnerable" individuals are also those most likely to benefit from supportive experiences [20].

A recent review of GXE findings chronicling differential susceptibility [17] underscored the need for research that would make it easier to determine whether GXE findings are consistent with diathesis-stress or differential-susceptibility models of environmental action. Considered essential, among other things, was not treating the absence of contextual stress (e.g., not abused) as the positive pole of the environmental continuum being measured. Thus, rather than just determining, for example, whether a child is harshly treated *or not*, GXE studies should seek to measure indisputably supportive environmental conditions (e.g., positive parenting). In the present work, a composite measure of parenting quality in adolescence ranging from very positive and supportive to very negative/disengaged and unsupportive is used to predict a measure of parenting stress in adulthood, one importantly known to relate systematically to actual parenting behavior [31, 32, 37, 46].

Indeed, in studies of parents of children from toddlerhood to adolescence [2, 28, 47, 66], parental stress has been linked to more punitive and less emotionally supporting parenting and to children's greater internalizing and externalizing problems. Unfortunately, parenting behavior itself was not assessed in the Add Health project from which this study obtained its data.

Virtually all GXE work to date has examined, for theoretically sensible reasons, the moderating effect of single genes [21]. This is typically due to the fact that investigators have a theory of gene action stipulating how particular genes influence particular neuro-transmitters and thus particular phenotypes [24]. In this work we adopt a different strategy given evidence that multiple genes operate as "plasticity genes" in some GXE research, not just "vulnerability genes" [17], specifically the 10R allele of DAT1 [61], the A1 allele of DRD2 [36], the 7R allele of DRD4 [54] and the short allele of 5HTTLPR [63]. In at least some GXE studies, though certainly not all of them, individuals carrying these particular alleles seem to be affected by a variety of environmental factors in a "for-better-*and*-forworse" manner [15], having their functioning both disproportionately undermined and enhanced by, respectively, unsupportive and supportive environmental conditions relative to those not carrying those alleles [12]. Indeed, the latter individuals appear much less and sometimes not at all affected by the very same environmental factors under investigation.

Rather than considering each gene one at a time and testing a series of single gene GXE interactions which might illuminate the genetic conditions under which parenting experienced in childhood proves related to parenting stress experienced in adulthood, we build on the recent work of several investigatory teams that have created *composite* measures of diverse alleles to predict phenotypic outcomes, based on the view that individual genes have very small effects [33, 41]. But rather than generating a summary index of genetic *risk* as Beaver and associates [10] did when predicting antisocial phenotypes, we create an index of *cumulative-genetic plasticity* which reflects the number of putative plasticity alleles in total that an adolescent carries of the set of four listed in the preceding paragraph. This affords testing the hypothesis that not only will cumulative-genetic plasticity genes an individual carries, the stronger the parenting effect being evaluated will prove to be.

Methods

Data

The data analyzed in this study were drawn from the National Longitudinal Study of Adolescent Health (Add Health). Previously published reports provide detailed discussions of the data and the sampling design [42, 43, 57]. Briefly, Add Health involves a longitudinal and nationally representative sample of American youths who were enrolled in middle or high school during the 1994–1995 academic year. Four waves of data have been collected thus far; two in adolescence, one in early adulthood, and one when the respondents were in their late 20s or early 30s. Individuals were assessed on a variety of different phenotypes, including their behaviors, personalities, and family environments. Overall, more than 15,000 respondents were interviewed across these four waves [42].

Add Health also contains a subsample of respondents who were genotyped for genetic polymorphisms related to neurotransmission. During wave 3 interviews, respondents who had a sibling or a co-twin, who was also participating in the Add Health Study were asked to submit samples of their buccal cells. In total, 2,612 respondents were included in the

DNA subsample of Add Health. Because more than one sibling per household was sampled, the DNA subsample lacked independence in observations, which can lead to downwardly biased standard errors. The lack of independence was corrected in two ways. First, one twin from each monozygotic twin pair was randomly selected and removed from the final analytical sample [38]. Second, all tests of statistical significance for the regression coefficients were estimated using Huber/White standard errors.

Measures

Predictor 1: Parenting Quality

The Add Health data contain a range of items designed to tap the quality of parenting that the adolescent received from their mother. Specifically, prior Add Health researchers have developed three indexes/scales that tap different dimensions of maternal parenting: one that taps maternal involvement, one that taps maternal disengagement, and one that taps maternal attachment [4]. The maternal involvement index measures the extent to which the mother is involved in her child's life. During wave 1 interviews, youths were asked whether they and their mother had participated in ten different activities during the past month, including playing a sport, going shopping, or watching a movie. Items were coded dichotomously, where a value of "0" indicated that the adolescent and their mother did not engage in the activity. Responses to these ten items were then added together to create the maternal involvement index (Cronbach's $\alpha = .66$).

Additionally, youths were also asked two questions that measured the child-mother relationship. In particular, during wave 1 interviews, youths reported on how close they felt to their mother and how much they thought their mother cared about them. Responses to these two questions were coded such that higher values indicated greater attachment. Responses to these questions were summed to create the maternal attachment scale (Cronbach's $\alpha = .49$). The third parenting scale available in Add Health measures maternal disengagement. During wave 1 interviews, five questions were asked to adolescents that tapped the extent to which their mothers lacked interest in and/or involvement with them. For instance, youths were asked to report how warm and loving their mother was, how often they talked with their mother, and the overall quality of their relationship with their mother. These questions were coded so that higher values represented more maternal disengagement. Responses to the items were summed together to create the maternal disengagement scale (Cronbach's $\alpha = .83$).

A principal components factor analysis with varimax rotation was then conducted on these three parenting composite measures. The results indicated that the variance– covariance matrix of these the three composites could be accounted for by a single-factor solution. The maternal disengagement scale was reverse-coded such that higher values reflected lower disengagement. Following prior research, [4], the three maternal parenting scales composites were transformed into a weighted factor score to create a measure of parenting quality. The parenting quality factor scale was then divided into quartiles to allow for more stable parameter estimates for the multiplicative interaction terms.

Predictor 2: Cumulative Genetic Plasticity

The genotyping of Add Health subjects was carried out at the Institute for Behavioral Genetics at the University of Colorado [43]. Prior research has identified four of the genetic

polymorphisms available in the Add Health¹ as being related to plasticity: DAT1, DRD2, DRD4, and 5HTTLPR [16]. In particular, the plasticity alleles for each of the genes were as follows: the 10R allele of DAT1, the A1 allele of DRD2, the 7R allele of DRD4, and the short allele of 5HTTLPR. Each polymorphism was assigned a value of "1" if at least one putative plasticity allele was present. The values for each of the genetic polymorphisms were added to create an index of "cumulative-genetic plasticity." The distribution of the cumulative-plasticity alleles was: 0 plasticity alleles = 0.2% (n = 2), 1 plasticity allele = 12.2% (n = 118), 2 plasticity alleles = 38.8% (n = 376), 3 plasticity alleles = 38.0% (n = 368), and 4 plasticity alleles = 10.8% (n = 105). Because only two cases had 0 plasticity alleles, all of the models were recalculated removing these two cases from the sample and also pooling them with the 1 plasticity allele group. The results were virtually identical and thus the two cases were included in the analyses with their original coding scheme.

Dependent Construct: Parental Stress

During wave 4 interviews, respondents were asked whether they had a biological child. If they responded affirmatively, they then responded to four questions drawn from the Parental Stress Scale [18] gauging, on a 5-point basis (1 = strongly disagree, 5 = strongly agree), the amount of stress that they experienced in being a parent. Specifically, they rated how (a) happy they were in the role of parent and (b) close they felt to their child(ren) (both reverse-coded), and the extent to which their children (c) were a major source of stress in their life and (d) they felt overwhelmed by the responsibility of being a parent. Responses to the four items were summed together to create the parental stress scale (Cronbach's $\alpha = .54$). Importantly, prior psychometric research of the Parental Stress Scale has revealed that it is both a reliable and valid way to assess parental stress [18].

Control Variables

Two control variables, gender (0 = female; 1 = male) and race, were included in the analyses. To avoid population stratification effects, race was coded by means of a series of dichotomous dummy variables. Caucasian was coded 0 = non-Caucasian and 1 = Caucasian; African-American was coded 0 = non-African-American and 1 = African-American; and Other was coded 0 = Caucasian or African-American and 1 = Other race. Other race was omitted from the equations and served as the comparison group for Caucasian and African-American.

Results

Statistical analysis involved a series of linked steps. To test for gene-environment correlation between the cumulative-genetic-plasticity index and parenting quality (measured in quartiles), bivariate correlations were estimated first (r = -.05, p > .05). The lack of

¹ The Add Health respondents were also genotyped for a polymorphism in the promoter region of the MAOA gene. However, since MAOA is X-linked, including this polymorphism in the plasticity index would necessitate separate models for males and females. Given that there is not a theoretical reason to believe that plasticity would differentially affect males and females in terms of parenting, we opted to exclude MAOA from the analyses and analyze males and females simultaneously.

	Model 1			Model 2		
	b	Beta	SE	b	Beta	SE
Genetic plasticity × parenting quality				19*	27	.08
Genetic plasticity	06	02	.09	.23	.08	.13
Parenting quality	24*	11	.07	.25	.12	.20
Gender	20	04	.16	20	04	.16
Caucasian	05	01	.23	.04	.01	.23
African-American	19	03	.28	18	03	.28
Caucasian African-American	05 19	01 03	.23 .28	.04 18	.01 03	

Table 1 OLS regression models predicting parental stress (N = 969)

Note: All models estimated using Huber/White standard errors

* p < .05, two-tailed tests

association meant that any discerned GXE interaction did not simply reflect G:E correlation and thus a possible evocative effect of cumulative genetic plasticity on mothering.

The next set of analyses estimated ordinary least squares (OLS) regression models to examine the interrelationships among the cumulative-genetic-plasticity index, parenting quality, and parental stress. Inspection of the results of the additive model presented in Model 1 of Table 1 reveals no association between cumulative genetic plasticity and parental stress ($\beta = .02$, p > .05), but a statistically significant and positive association between parenting quality and parental stress ($\beta = .11$, p < .05). Model 2 displays the results of the OLS equation, where a two-way multiplicative interaction term between cumulative genetic plasticity and parenting quality proved to be statistically significant ($\beta = .27$, p < .05).

Supplementary analyses evaluated the two-way interaction between cumulative genetic plasticity and parenting quality in closer detail. First, simple slopes were calculated for each of the plasticity groups and difference-in-coefficients *z*-tests were estimated to determine which of the slopes differed significantly from each other [52]; see Table 2. Of particular importance, the simple slope for the 4-plasticity-allele group was statistically significant and differed significantly from the simple slopes for the 1-, the 2- and the 3-plasticity-allele groups (z = 3.17, 2.29, 2.73, respectively, all p < .05). (Because only two cases carried 0 plasticity alleles, they were excluded from these analyses.) The simple slopes among the 1-, 2-, and 3-plasticity-allele groups were not significantly different from each other.

 Table 2
 Post-hoc analysis of the gene-environment interaction between cumulative genetic plasticity and parenting quality

	Simple slope	Comparison group for Z-test for equality in regression slopes				
	Parenting quality ^a	1 plasticity allele	2 plasticity alleles	3 plasticity alleles		
1 plasticity allele	0.10	_	_	_		
2 plasticity alleles	-0.24*	1.62	-	-		
3 plasticity alleles	-0.13	1.09	0.81	-		
4 plasticity alleles	-0.80*	3.17*	2.29*	2.73*		

* p < .05, two-tailed tests

^a corrected for race and gender



Fig. 1 The association between parenting quality and parental stress as a function of plasticity alleles

Last, the slopes for each of the plasticity allele subgroups were plotted across the four quartiles of the parenting quality scale. Figure 1 portrays these plots and shows a clear pattern of results consistent with the differential susceptibility hypothesis. In particular, individuals in the 4-plasticity-allele group scored lowest on parental stress when reared by the most supportive mothers *and* highest on the parental stress scale when reared by the most negative mothers. Inspection of the Figure and the estimates presented in Table 2 indicate that individuals with 2 and 3 plasticity alleles appeared to be more affected by their rearing history, also in a "for-better-and-for-worse" manner, depending on the quality of parenting experienced in adolescence, than those with only 1 plasticity allele.

Discussion

The fact that parenting has been found to be transmitted across generations, but only to a modest extent, inspired the work presented here by raising the possibility that this could be due to the fact that individuals differ in their susceptibility to rearing [9, 10, 20]. After all, if individuals more and less susceptible are not distinguished, then average estimates of intergenerational transmission could both under- and over-estimate effects of parenting experienced on the experience of parenting. The current study investigated whether gene-environment interaction might contribute to understanding the intergenerational transmission of parenting.

To do so, the DNA subsample of Add Health was analyzed to examine the interrelationships among maternal parenting experienced by the respondent during adolescence, the respondent's own experience of parental stress—obtained on average 14 years after the parenting-in-adolescence measurement—and a measure of the respondent's cumulative genetic plasticity. Although it would have been preferable to have measurements of actual parenting rather than of parenting stress, no such data were available for analysis in Add Health. The fact that parental stress is a known predictor of parenting mitigates this study limitation somewhat [31, 32, 46]. Nevertheless, it will be important to determine whether the results reported here for what might be regarded as a proxy measure of parenting—parenting stress—extend to actual parenting.

Two broad findings emerged from this inquiry. First, there was a statistically significant interaction between parenting quality and cumulative genetic plasticity in the prediction of parental stress during adulthood. This finding suggests that parenting quality differentially affects future parental stress depending on genotype-and thus could at least partially explain why estimates of transmission across generations are not as strong as might perhaps be expected on the basis of theory. Second, and perhaps even more importantly, analyses conducted to examine whether the GXE interaction was consistent with the diathesis-stress or differential-susceptibility model of environmental action proved indisputably consistent with differential-susceptibility. Recall that respondents in the 4-plasticity-allele group who were exposed to the most positive maternal parenting as adolescents scored the lowest on parental stress during adulthood and, at the same time, respondents in the 4-plasticity-allele group who were exposed to the most negative maternal parenting as adolescents scored the highest on parental stress during adulthood (see Fig. 1). Although not nearly as pronounced, a similar pattern of findings surfaced for the 2- and 3- plasticity-allele groups. This "for-better-and-for-worse" finding is precisely what the differential-susceptibility hypothesis predicts [12, 15]; see also [20].

Even though the results reported extend the range of phenotypes that the differentialsusceptibility hypothesis applies to, the findings need to be interpreted with caution for a number of reasons. To begin with, the cumulative genetic plasticity index employed in this study was created from only four genes. Belsky and associates [16, 12] have called attention to others, however, that appear to operate in a differential-susceptibility, not just diathesis-stress manner, including COMT and DRD3. Unfortunately, Add Health respondents were only genotyped for a handful of genes and this thus constrained the cumulative plasticity index that could be used in this inquiry. Given that single genes tend to have such small effects [58], one cannot but wonder, had additional putative plasticity genes been available for analysis, whether findings consistent with differential susceptibility would have been even more pronounced.

Limiting this study, too, was the fact that the outcome to be explained, parental stress (in the absence of a direct measure of parenting), was based on only four items and, as a result, had only moderate internal consistency. Such fallibility of measurement of the dependent variable could also have attenuated the discerned effects of the predictor variables used in this inquiry [39]. This seems quite conceivable because the power needed to detect statistical interactions is significantly greater than needed for evaluation of main effects [27].

An additional limitation of the measurement was that the dependent construct, parental stress, only assessed the negative side of the parenting experience. A more appropriate measurement strategy when testing for differential-susceptibility is to include measures that range from positive to negative [12]. In most GXE studies, especially those conducted in psychiatric genetics, however, the absence of a negative environment (e.g., not maltreated) or outcome (e.g., not depressed) is treated as positive. Central to the differential-susceptibility hypothesis is the proposition that plasticity alleles contribute to positive outcomes in the face of positive environments, not just negative outcomes in the face of measurements like parental stress that do not capture the positive pole of the parenting experience (other than absence of stress), the ability to detect the "for better" part of the "for-better-and-for-worse" differential-susceptibility hypothesis becomes extremely difficult. That

such a pattern of findings emerged in this inquiry despite this measurement handicap also provides a basis for wondering whether findings would have been stronger had the parenting-experience outcome been evaluated across the full negative-to-positive range, just as the parenting-in-adolescence predictor was.

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Conflict of interest The authors have no competing financial interests in relation to the work described.

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