

PERSPECTIVE

Vulnerability genes or plasticity genes?

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The classic diathesis–stress framework, which views some individuals as particularly vulnerable to adversity, informs virtually all psychiatric research on behavior–gene–environment ($G \times E$) interaction. An alternative framework of ‘differential susceptibility’ is proposed, one which regards those most susceptible to adversity because of their genetic make up as *simultaneously* most likely to benefit from supportive or enriching experiences—or even just the absence of adversity. Recent $G \times E$ findings consistent with this perspective and involving monoamine oxidase-A, 5-HTTLPR (5-hydroxytryptamine-linked polymorphic region polymorphism) and dopamine receptor D4 (DRD4) are reviewed for illustrative purposes. Results considered suggest that putative ‘vulnerability genes’ or ‘risk alleles’ might, at times, be more appropriately conceptualized as ‘plasticity genes’, because they seem to make individuals more susceptible to environmental influences—for better *and* for worse.

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Introduction

Central to the field of psychiatric genetics is the search for ‘vulnerability genes’.¹ Particularly important in this search is uncovering the mechanisms whereby such genes influence disease risk, and determining whether they are directly associated with more general psychological and behavioral disturbance (that is, act through ‘main effects’) or whether they are principally related to dysfunction only under specific environmental conditions or in response to particular developmental experiences (that is, act through ‘gene- \times -environment ($G \times E$) interaction’). Findings of studies linking candidate vulnerability genes (or ‘risk alleles’) directly to specific psychopathological conditions have proven notoriously difficult to replicate.¹

Failure to replicate direct effects of candidate vulnerability genes on specific psychopathological conditions suggests that genes may not influence behavior directly, leading many investigators to examine how genes may moderate effects of the environment on human $G \times E$ interaction. One well-studied $G \times E$ interaction involves monoamine oxidase-A (MAOA), contextual adversity and antisocial behavior. The interaction of a functional MAOA gene polymorphism (MAOA-uVNTR) and childhood adversity, first detected by Caspi *et al.*² in their research

on child maltreatment, has now been replicated enough times that meta-analysis reveals it to be reliable³—despite claims by some to the contrary.⁴ Thus, there is growing evidence that individuals possessing the low-MAOA-activity allele are predisposed to become antisocial when they experience a variety of adverse experiences, most notably maltreatment in childhood.

It is not just in the case of the MAOA gene that the notion of genetic vulnerability takes center stage in research on $G \times E$ interactions. Consider the evidence showing that the ‘s’ allele of the 5-hydroxytryptamine-linked polymorphic region polymorphism (5-HTTLPR) is associated with increased depression in a high-stress context, with the ‘l’ allele functioning protectively.⁵ Several studies have replicated this finding, providing further support for the conclusion that 5-HTTLPR increases vulnerability to depression in the context of environmental stress.⁶

The fundamental premise of this essay, however, is that viewing relations among genes, behavior and the environment from the perspective of the classic diathesis–stress model of psychopathology, as so much psychiatric genetic research on $G \times E$ interaction does, may distort these relations and thereby undermine, rather than advance, the understanding of how genes and environment collectively operate to shape behavior and development, including risk of mental illness. Central to the diathesis–stress model is the postulate that some individuals are at heightened risk—because of their genetic make up—of succumbing to psychological disturbance when they encounter adversity, whereas others, lacking the genetic vulnerability, are not so affected even when

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exposed to the very same adversity. Thus, whereas an individual with a particular genetic vulnerability will be prone to develop a genetically specific disorder if he or she experiences what could be some particular or any of a variety of environmental stressors (for example, child abuse, negative life events, death of parent), the same environmental exposure will not engender psychopathology in an individual possessing a different version of the candidate gene in question.⁷

An alternative to the diathesis–stress framework at the heart of genetic vulnerability thinking is the one which presupposes that it is not so much that individuals vary only in their susceptibility to adversity vis-à-vis psychopathology, but rather that these putatively ‘vulnerable’ individuals are actually more susceptible and, thus, responsive to both positive and negative environmental conditions, that is, in a ‘for better and for worse’ manner. This differential-susceptibility perspective does not just contend, as many have, that genes are neither inherently good or bad, or even that their developmental and behavioral effects depend on person–environment fit,^{8,9} but rather—and distinctively—that individuals vary in their plasticity or susceptibility to environmental influences. Thus, the very genes that seem—in so much psychiatric genetic research—to make individuals disproportionately vulnerable to adversity vis-à-vis psychopathology may, *simultaneously*, confer on them an advantage when it comes to benefiting from exposure to environmental support or enrichment (for example, nurturance), including just the absence of adversity. Were this the case, it would seem more appropriate to speak of ‘plasticity genes’ rather than ‘vulnerability genes’ and of highly plastic or malleable individuals rather than the so-called vulnerable ones. Boyce and Ellis,¹⁰ although not directly concerned with $G \times E$ effects, which is the focus of this paper, also have argued that individuals vary in their susceptibility to environmental influences, what they refer to as ‘biological sensitivity to context.’ However, there is no presumption in their work that such malleability is a function of genotype, as they intriguingly entertain the prospect that experience can shape plasticity.

Ultimately, the purpose of this paper is not so much to challenge the view that diathesis–stress phenomena exist or that processes related to them operate. That seems indisputable. However, it is to contend—and illustrate empirically—that in many cases, wherein this may seem to be so, something different may be occurring, yet go virtually unnoticed as a result of expectations derived from the prevailing conceptual perspective, which guides both inquiry and interpretation of findings. Indeed, a central claim of this paper is that the disproportionate attention paid to the negative effects of contextual adversity, broadly defined and varied in its operationalization, on the problematic functioning and on disturbances in development and mental health, may actually lead scholars to mischaracterize environmental influences, as well as human development processes and phe-

nomena. And this is because, as stipulated by the differential-susceptibility hypothesis,^{11–14} ‘the very same individuals who may be most adversely affected by many kinds of stressors, may simultaneously reap the most benefit from environmental support and enrichment (including the absence of adversity)’.

In the primary body of this paper, we provide extensive but still illustrative $G \times E$ evidence to this effect, most of it very recent and much of which has gone unnoticed, even at times by the investigators generating it. What follows should not be regarded as an exhaustive review of the literature; however, nor should it be seen to imply, much less demonstrate, that evidence of differential susceptibility outweighs evidence of diathesis–stress, either in the literature as a whole or even in each and every study cited for illustrative purposes. To make the case, as we exclusively seek to, that differential susceptibility seems operative in human development and functioning, but that individual differences in plasticity have been largely overlooked—in favor of prevailing views that some individuals are simply more vulnerable to adversity than others—it is our contention that an admittedly selective compilation of illustrative $G \times E$ findings is exactly what is appropriate at the present time. This would seem especially so in light of the fact that almost all the available human $G \times E$ research focuses on both a restricted range of environments, typically emphasizing the negative end of the spectrum and failing to measure at all the positive (except for the absence of adversity), and a restricted range of psychological and behavioral outcomes, also typically emphasizing the negative, thereby failing to assess competent functioning (except for the absence of dysfunction). As a result of these design characteristics of so many $G \times E$ investigations, it remains unknown whether extensive evidence consistent with a diathesis–stress model and seemingly inconsistent with a differential-susceptibility framework is an accurate reflection of $G \times E$ processes or an artefact of study designs. Quite conceivably, simply treating the absence of adversity as the ‘good’ end of the environmental-exposure continuum and/or absence of a disorder as the ‘good’ end of the psychological functioning continuum may lead to the under-detection of differential-susceptibility findings and an over representation of vulnerability ones. It is for these reasons that it is considered appropriate at the present time to provide illustrative evidence of apparent differential-susceptibility effects rather than undertake a formal meta-analysis of $G \times E$ findings in hopes of determining which model fits the data better.

Illustrative evidence: MAOA, 5-HTTLPR, dopamine receptor D4 (DRD4)

Belsky *et al.*¹³ recently delineated a series of empirical requirements, or steps, for convincingly establishing evidence of differential susceptibility to environmental influence, that is, individual differences in

plasticity. The first concerns the application of conventional statistical criteria for evaluating genuine moderation of a putative environmental influence by an organismic plasticity or susceptibility factor,¹⁵ including genotype, with some emphasis on excluding interactions with regression lines that do not cross (sometimes referred to as removable interactions). The next steps distinguish differential susceptibility from person–environment correlations, including Gene–Environment ones, which may reflect evocative effects of person characteristics on environmental experiences and from diathesis–stress models. If the (genetic) susceptibility factor and the (problematic) outcome are related, diathesis–stress is suggested. The specificity of the differential-susceptibility effect is demonstrated if the model is not replicated when other (genetic) susceptibility factors (that is, moderators) and outcomes are used.^{16,17} Differential susceptibility is thus demonstrated when the moderation reflects a crossover interaction that covers both the positive and the negative aspects of the environment, with the positive typically (and unfortunately) represented by the mere absence of adversity. The slope for the susceptible subgroup should be significantly different from zero and at the same time significantly steeper than the slope for the non-(or less-) susceptible subgroup.

In the remainder of this section, we present illustrative $G \times E$ evidence of differential susceptibility to environmental influence that are consistent with the view that individuals differ in their plasticity, with some being more affected than others by experiential influences in a for-better-and-for-worse manner. Perhaps, because so much of the work to be cited is new—and often conducted with a diathesis–stress frame of reference in mind—it is actually rare for investigations to address all or even most of the statistical criteria highlighted by Belsky *et al.*¹³ for providing convincing evidence of differential susceptibility to environmental influence. Indeed, even when investigators detect statistical interactions of a crossover nature, as is the case in all the researches to be cited, different strategies of following up such interactions are adopted to illuminate their nature. Whereas some investigations adopt a grouping approach for dealing with the interacting predictor variables, plotting or tabling sub-group means, others calculate and contrast slopes reflecting the differential predictive relation between the continuously measured environmental predictor and outcome for groups that differ on the moderating susceptibility factor. Only rarely is it reported whether such slopes differ significantly from each other, as would be preferable when the moderator does not have a natural break point, but is a continuous dimension (but as is not required when the moderator is naturally categorical with only two categories). Perhaps analogously, it is not always reported when the subclass means are plotted and exactly which mean differs significantly from which others.

In the service of illustrating what seems to be individual differences in plasticity, and thus, differ-

ential susceptibility in existing $G \times E$ research, we adopt a liberal standard of evidence once a significant crossover interaction has been detected when it comes to regarding the results as evidence of differential susceptibility to environmental influences. Specifically, and with regard to subgroup means, if one subgroup shows both the highest and the lowest mean of all susceptibility-factor-defined subgroups (for example, short vs long 5-HTTLPR allele) on an outcome with regard to the environmental effect in question, this is interpreted as in line with the for-better-and-for-worse differential-susceptibility patterning of results. Similarly, but with regard to slopes, whenever they indicate that one subgroup defined on the basis of the susceptibility factor in question would score highest and lowest given the environmental influence under investigation (that is, steepest slope), this too is interpreted as evidence of differential susceptibility. All the findings to be presented meet these criteria, with some pertaining to MAOA, some to 5-HTTLPR and some to DRD4.

Monoamine oxidase-A

Often unnoticed in Caspi *et al.*'s² groundbreaking $G \times E$ research showing that males with the less active version of the MAOA gene proved most antisocial in young adulthood when they experienced maltreatment in childhood, is that individuals with the same MAOA allele scored *lowest* in anti-social behavior when *not* exposed to child maltreatment, even if not by much. A re-interpretation of this study's results in terms of plasticity and differential susceptibility rather than vulnerability and diathesis stress would seem viable given results of a significant number of efforts to replicate the findings. For example, Kim-Cohen *et al.*³ studied 975 boys to determine whether the MAOA polymorphism moderated effects of mother-reported physical abuse in early childhood on later mental health problems. At age 7 years, boys with the low-MAOA-activity variant were rated by mothers and teachers as having more mental health problems—and specifically attention-deficit hyperactivity disorder (ADHD) symptoms—if they had been victims of abuse, but fewer problems if they had not, compared with boys with the high-MAOA-activity genotype. In another longitudinal study, this one of 514 adolescent twin boys aged 8–17 years, Foley *et al.*¹⁸ found that childhood adversity—based on parent and child report—predicted a 3-month history of conduct disorder (DSM-III, Diagnostic and Statistical Manual (of mental disorders)) differently for children with and without the low-activity-MAOA allele. Once again, boys with the low-MAOA-activity allele were more likely to be diagnosed with conduct disorder if exposed to higher levels of childhood adversity and less likely if exposed to lower levels of adversity, compared with boys with the high-MAOA-activity allele. Similar results emerged in Nilsson *et al.*'s¹⁹ cross-sectional investigation of 81 adolescent boys when the predictor was psychosocial risk, operationalized in terms of maltreatment experience

and living arrangement. Only boys with the low-MAOA-activity allele were affected by such risk, such that those with a history of adversity engaged in more criminal behavior (composite of vandalism, violence, stealing) and those lacking this history engaged in less.

Three additional studies extend the Caspi *et al.*² findings: one was a prospective investigation of 631 male and female, and white and black victims of (court-substantiated) child abuse and neglect, along with a comparison group matched on age, sex, race/ethnicity and social class background;²⁰ the second one a retrospective study of 235 adult psychiatric outpatients and healthy controls who reported on trauma experienced in childhood and physical aggression in adulthood;²¹ and the last one a cross-sectional retrospective study with an American Indian sample of 291 adult women, 50% of whom had a history of childhood sexual abuse.²² White (but not black) males and females with the low-MAOA-activity allele in the longitudinal study manifested the most lifetime violent and antisocial behavior during adolescence, as well as around age 40, if they had been maltreated, but the least (at both times of measurement) if they had not been victims of abuse. In the second study, men (only) with the low-MAOA-activity variant reported more physical aggression if they experienced one or more (retrospectively reported) objective traumatic events while growing up (for example, death of mother, severe physical handicap of sibling) and less physical aggression if there was no history of trauma, compared with high-MAOA-activity men (for whom trauma proved unrelated to aggression). In the third study, women homozygous for the low-MAOA-activity variant had the highest count of antisocial personality disorder symptoms when reporting childhood sexual abuse and the lowest count when having no history of sexual abuse, compared with women homozygous for the high-activity allele. In all inquiries except for the one by Kim Cohen *et al.*,³ the MAOA polymorphism proved unrelated to the environmental predictor and to the outcome investigated, consistent with a differential-susceptibility interpretation.

The 5-hydroxytryptamine-linked polymorphic region polymorphism

Again breaking the empirical ground in G × E research, Caspi *et al.*⁵ were the first to show that the 5-HTTLPR moderates effects of stressful life events during early adulthood on depressive symptoms as well as on probability of suicide ideation/attempts, and of major depression episode at age 26 years. Individuals with two 's' alleles proved most adversely affected, whereas effects on l/l genotypes were weaker or entirely absent. Of special significance, given our focus on differential susceptibility, is that carriers of the s/s genotype scored best on the outcomes just mentioned when stressful life events were absent, though this was just as true among low-MAOA activity individuals in Caspi *et al.*,² although not by very much.

Several research groups have attempted to replicate Caspi *et al.*'s⁵ findings of increased vulnerability to depression in response to stressful life events for individuals with one or more copies of the 's' allele, with most succeeding (see below), even if not all (for example, Surtees *et al.*²³). Going unnoticed in most, even if not all, of this work to be summarized below, however, is the fact that those carrying short alleles (s/s, s/l) did not just function most poorly when exposed to many stressors, but best—showing least problems—when encountering few or none of the stressors. Consider, for example, Taylor *et al.*'s²⁴ findings (appreciated by the investigators) showing that young adults homozygous for short alleles (s/s) manifested greater depressive symptomatology than individuals with other allelic variants when exposed to early adversity (that is, problematic childrearing history) as well as many recent negative life events, consistent with a diathesis–stress framework, yet the fewest symptoms when they experienced a supportive early environment or recent positive experiences, that is—and importantly—not just the absence of adversity. A similar for-better-and-for-worse pattern of environmental effects emerged in still other investigations of stressful life events and depression, including one targeting depressed patients, healthy controls and experiences during the 6 months before study enrollment,²⁵ and another of a sizeable community sample (*n* = 567) and life events up to 2 years before the assessment of depression.²⁶

The same for-better-and-for-worse pattern of results are evident—and noted—in Brummett *et al.*'s²⁷ investigation of more than 200 adults (mean age 58 years) who differed in whether or not they served as caregiver of a relative with Alzheimer's disease (see Figure 1) and in Eley *et al.*'s²⁸ research on adolescent girls who were and were not exposed to risky family environments. Indeed, careful consideration of Figure 1 in Eley *et al.*'s²⁸ report reveals a beneficial

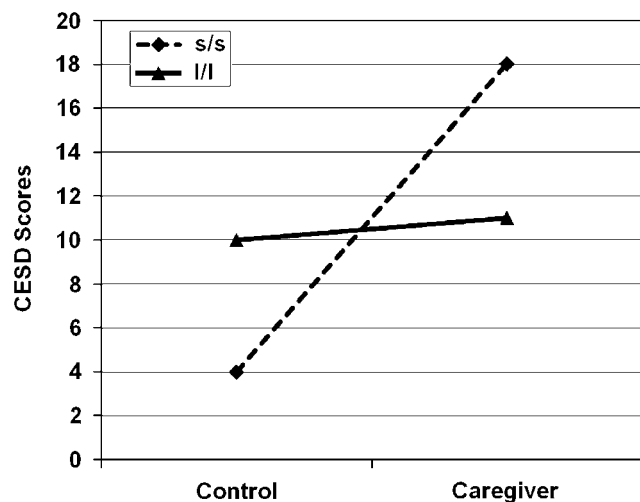


Figure 1 Center for Epidemiological Studies-Depression (CESD) scores for female caregivers and non-caregiver controls by 5-HTTLPR genotype (Brummett *et al.*²⁷).

effect of the *s/s* genotype in low-stress environments, with females homozygous for short alleles being 25% less likely to be in the high depression group than *l/l* participants. This positive effect is actually greater than the negative one described by the authors in the case of the *s/s* genotype in a high stress environment (that is, 20% more likely to be in high depression group than *l/l*). Re-graphing Eley *et al.*'s²⁸ Figure 1 using the format adopted originally by Caspi *et al.*⁵ brings this fact to the fore (see Figure 2 below). Comparison of the two figures highlights graphically the point being made repeatedly: the *s/s* genotype is associated with elevated depressive symptoms or risk among women in high-stress environments, yet among those in low-stress environments the *s/s* genotype is associated with reduced depressive symptom levels or risk relative to women with the *l/l* genotype. Wilhelm *et al.*'s²⁹ longitudinal data document the same pattern in an investigation of probability of life-time major depression and exposure to adverse events across a 5-year study period.

The effect of 5-HTTLPR in moderating environmental influences in a manner consistent with differential susceptibility is not restricted to depression and its symptoms, but also—and perhaps unsurprisingly—to anxiety and ADHD. Gunthert *et al.*³⁰ documented the former result in a longitudinal study of 350 college students. At study entry and a year later, participants reported anxiety and negative events daily for 30 days. Genotyping distinguished three alleles, but the L_G allele was grouped with 's' alleles owing to its functional equivalence vis-à-vis promoter activity. Individuals judged homozygous for short alleles (including s/L_G and L_G/L_G) reported more anxiety in the evening when daily-event stress was high compared with individuals with different genotypes, but also less anxiety than other genotypes when experiencing little daily-event stress, a pattern consistent across measurement occasions. Once again the fact that the susceptibility factor did not predict the environmental measure or the outcome is considered important.

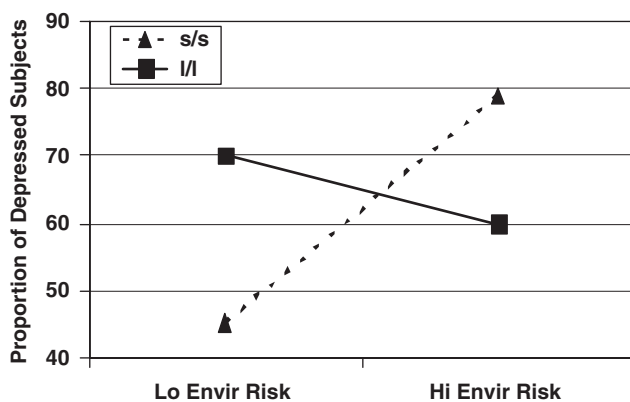


Figure 2 Proportion of female participants with a high level of depression by environmental risk group and 5-HTTLPR genotype (Eley *et al.*²⁸).

In a second study focused on undergraduate students ($n=247$) and anxiety,³¹ but this time concerned with (retrospectively reported) emotional abuse in childhood, a $G \times E$ interaction once more emerged, with genotype importantly proving unrelated to the environmental predictor and the outcome, anxiety sensitivity. The significantly steeper abuse-anxiety slope in the case of students homozygous for short alleles relative to those with one or more long alleles indicated that *s/s* individuals scored highest in anxiety sensitivity when exposed to abuse and lowest when not exposed.

Moving on to consider ADHD (in childhood and adulthood), Retz *et al.*³² focused on the moderated effects of an adverse childhood environment in their study of 184 male delinquents who averaged 34 years of age. Using a retrospective assessment of childhood ADHD as well as of early adversity, but a clinical interview to assess functioning in adulthood, these investigators detected a crossover interaction with respect to the persistence of ADHD over time. Compared with *l/l* genotypes, individuals with 's' alleles had more and less persistent ADHD, depending on whether or not, respectively, they experienced an adverse early environment.

One observation that makes the findings under consideration particularly interesting is that short alleles on the serotonin transporter gene have also been found, in at least some molecular-genetic research,³³ to be associated with negative emotionality in young infants. What makes this gene-behavior linkage particularly important is that, even though negative emotionality or difficult temperament in infants has long been conceptualized as a risk or vulnerability factor for the development of behavior problems in childhood,³⁴ a growing literature on differential susceptibility reveals negatively emotional or difficult infants to be more plastic or malleable than other infants—in a for-better-and-for-worse manner.³⁵ That is, they do worse than others under poor rearing conditions, but better than others under good ones. Consider in this regard the evidence that infants rated by mothers as highly negatively emotional at 6 months of age not only manifest, relative to other children, more behavior problems in early childhood when experiencing low-quality parenting³⁶ or low-quality child care,^{37,38} but also fewer problems and more social skills than other children when exposed to high-quality parenting or child care. Relatedly, Kochanska *et al.*³⁹ observed that highly fearful 15-month-olds experiencing high levels of power-assertive paternal discipline were most likely to cheat in a game at 38 months, yet when cared for in a supportive manner, such negatively emotional, fearful and putatively vulnerable toddlers manifested the most rule-compatible conduct. Even more noteworthy than such longitudinal/correlational evidence is the clinical-trial finding of a parenting intervention designed to promote secure attachment of infant to mother showing that the principle beneficiaries of the documented experimental effect were highly negative

and fearful infants.⁴⁰ These findings pertaining to negatively emotional infants demonstrate not only that such putatively vulnerable children are more beneficially affected than other infants by some enriching rearing conditions but, in so doing, raise the prospect that this could be because they are carriers of short alleles on the serotonin transporter gene. Only when investigators evaluate interactions involving both the environment and genetics, and the environment and temperament will it be possible to determine whether the same highly plastic individuals are being identified in different studies using different (behavioral and genetic) markers.

DRD4

Recent research on $G \times E$ interaction involving the 7-repeat allele of the dopamine receptor gene, DRD4, which meta-analysis reveals to be reliably associated with ADHD,¹ also provides support for the claim that the so-called vulnerability genes may be better conceptualized as plasticity genes. What makes the first two studies to be considered especially important is that the predictor variable, parenting, ranges from quite limited to very competent, thus meaning that a supportive environment is not just the one in which adverse experiences are absent. In a longitudinal investigation of 47 infants, greater maternal insensitivity observed when children were 10 months predicted greater externalizing problems reported by the mother more than 2 years later, but only for children carrying the 7-repeat DRD4 allele.⁴¹ Moreover, although children with the 7-repeat DRD4 allele displayed, consistent with a diathesis–stress model, the most externalizing behavior of all children when mothers were judged insensitive, they also manifested the least externalizing behavior when mothers were highly sensitive (but see, for contradictory results, Propper *et al.*⁴²). Similar results emerged in a cross-sectional investigation of sensation seeking involving 45 children who were 18–21-month-olds, with toddlers carrying the 7-repeat allele rated by parents as showing, compared with children without the 7-repeat allele, less sensation seeking behavior when parenting quality was high and more when parenting quality was low.⁴³ Although parenting proved significantly associated with sensation seeking in the 7-repeat individuals, it did not in other children. Of importance is the fact that genotype did not predict parenting or sensation seeking, fulfilling important differential-susceptibility criteria.

Experimental intervention research involving the enhancement of parenting also documents a moderating effect of the 7-repeat allele; once again, then, this $G \times E$ work does not simply define the ‘good’ environment in terms of the absence of adversity. Thus, when Bakermans-Kranenburg *et al.*⁴⁴ looked at the change over time in parenting—from before to well after a video-feedback parenting intervention was provided on a random basis to 157 mothers of 1–3-year-olds who scored high on externalizing problems—they not only found that the intervention

succeeded in promoting more sensitive parenting and positive discipline, but that experimental effects extended to improvements in child behavior. This proved to be the case, however, only for those children carrying the DRD4 7-repeat allele, with most of the experimental effect being carried by these genotypically susceptible children whose mothers showed the most improvement in their parenting. Much the same was the case when, at post-treatment follow up, stress reactivity was measured by means of change in salivary cortisol before and after administration of an experimental stressor (that is, area under the curve⁴⁴). Indeed, DRD4 7-repeat children in the experimental group not only showed the least physiological stress reactivity of all children, but the most if their mothers had been assigned to the control group.

The same team of Dutch investigators whose $G \times E$ research has focused upon indisputably positive environmental effects also found evidence that the DRD4 7-repeat allele moderated the effect of a maternal psychological condition, unresolved loss or trauma (as measured by means of the Adult Attachment Interview) on early infant development. More specifically, unresolved loss predicted infant attachment disorganization, an early developmental marker of psychological disturbance later in life,⁴⁵ but only in the case of infants carrying the 7-repeat allele.⁴⁶ Indeed, these infants manifest both the most and least disorganized attachment behavior when stressed depending on whether their mothers had or had not experienced unresolved loss or trauma in their own lives. Importantly, genotype predicted neither unresolved loss nor disorganization, with the data thus meeting criteria for differential susceptibility (rather than just genetic vulnerability).

Finally, in a study with a focus rather different than the ones just considered, Seeger *et al.*⁴⁷ evaluated whether the season of the year in which a child was born interacted with the dopamine DRD4 polymorphism in predicting the hyperkinetic conduct disorder (ADHD). Employing a cross-sectional design involving 64 children with the disorder and 163 healthy controls (mean age 11–12 ± 3 years), they found that it did—and in ways consistent with what is known about photoperiod exposure during pregnancy. When comparing patients with controls, children with one copy of the DRD4 7-repeat allele born in autumn and winter (that is, long photoperiod during pregnancy) had a 5.4-fold decreased relative risk for hyperkinetic conduct disorder, whereas children with the same genotype born in spring and summer (that is, short photoperiod) had a 2.8-fold increased relative risk for hyperkinetic conduct disorder. Neither season of birth nor the presence of DRD4 7-repeat allele represented a risk factor for hyperkinetic conduct disorder *per se*.

Conclusion

In some respects, it should not be surprising that putative vulnerability genes may actually function

more like plasticity genes, resulting in certain individuals being more responsive than others to both positive and negative environmental experiences, including the simple absence of contextual adversity. Not only has plasticity been found to be heritable in many species,^{48,49} functioning perhaps as a selectable character in and of itself,⁵⁰ but recent computer simulations show that individual differences in responsiveness to the environment could most certainly evolve.⁵¹ In fact, one wild bird population shows evidence that selection favoring individuals who are highly plastic with regard to the timing of reproduction has intensified over the past three decades, perhaps in response to climate change causing a mismatch between the breeding times of the birds and their caterpillar prey.⁵² Of note too is Suomi's⁵³ observation that only two species of primates fill diverse ecological niches around the world, humans and rhesus macaques, and that what distinguishes both of these 'weed species', as he calls them, from all other primates is the presence of 5-HTTLPR short allele in some individuals. It seems unlikely that that which might afford these two species such an adaptive advantage would only be 'vulnerability genes' that predispose carriers to depression in the face of contextual stress.

Given the focus in psychiatric genetics on adversity in the form of environmental risk factors and vulnerability in the form of genes associated with pathological conditions, it is not surprising that the possibility that the so-called vulnerability genes actually function more like plasticity genes could go unnoticed. It is almost as if, metaphorically speaking, sailors are so busy—and wisely—looking under the water line for extensions of icebergs that could sink their ship that they fail to appreciate that by climbing on top of the iceberg it might prove possible to chart a clear passage through the ice-laden sea. To the extent that it is appropriate to think in terms of plasticity rather than vulnerability, research will be required that extends the purview of the molecular-genetic study of behavior well beyond the investigation of dysfunction and environmental adversity. Once again Caspi *et al.*⁵⁴ have served as groundbreakers, discovering that, with respect to intelligence, children carrying one variant of FADS2, a gene involved in the genetic control of fatty acid pathways, benefit from breastfeeding, whereas those carrying a different allele are not so affected. What has not been determined as of yet, because all mothers in this study were apparently in relatively good nutritional state, is whether those individuals found to benefit from breastfeeding for genetic reasons actually are adversely affected by it when mothers are in poor nutritional condition. Not only is this just what a differential-susceptibility perspective would predict, but is also what might explain why the gene associated with increased intelligence in the context of breastfeeding, most certainly the universal condition in ancestral times, has not gone to fixation.

Although the primary purpose of this essay has been to argue that the virtually exclusive focus on adverse environmental conditions and psychopathology in psychiatric genetic research risks mischaracterizing individuals who are more susceptible than others to the negative consequences of adversity and to the benefits of environmental support and enrichment as being exclusively the former—that is, genetically vulnerable—it would be a mistake to view these orientations as mutually exclusive. Indeed, thinking about plasticity and vulnerability together raises the following three interrelated questions: Are there some polymorphisms that make some individuals more responsive than others to supportive and adverse environments, just as a differential-susceptibility framework presupposes? Are there some polymorphisms that only cause some individuals to be more susceptible than others to adversity, just as a diathesis–stress/genetic-vulnerability framework presupposes? And are there still other polymorphisms that only make individuals more susceptible than others to enriching environmental conditions? Of note, with regard to the last possibility is that, although the English language has terms to characterize those highly susceptible to both positive and negative conditions (that is, plastic/malleable) and highly susceptible to adversity (that is, vulnerable), it is difficult to find a term which would characterize those disproportionately responsive to supportive conditions only—besides lucky!

Owing to the inherent limits of so many of the studies that we have considered, both in terms of what has been measured and how the data have been analyzed and presented in primary publications, it remains impossible to be certain whether the extensive findings considered throughout this paper as evidence of differential susceptibility should be regarded as such. Recall in this regard that rather liberal standards of interpretation have, by necessity, been applied to G × E findings, most of which emerged from investigations designed to evaluate diathesis–stress hypotheses. To enable both primary researchers and reviewers of the literature in the future, including meta-analysts, to address this fundamental issue about how human development operates, investigatory and reporting practices will need to change; and hopefully, this selective review, by calling attention to the possibility of differential susceptibility, will stimulate such change.

Thus, in addition to meeting the Belsky *et al.*¹³ criteria for establishing differential susceptibility summarized earlier, which informed the interpretation of study findings considered herein, several other research desiderata are called for. First, studies should measure not just the presence of adversity and its absence, but environmental support, like Taylor *et al.*²⁴ did in assessing positive life events and Bakermans-Kranenburg and van IJzendoorn⁴¹ did in measuring sensitive parenting. Second and relatedly, human functioning should be measured along a continuum ranging from dysfunction to competence,

not just from dysfunction to its absence or just from competence to its absence, to avoid the masking of differential susceptibility by ceiling or floor effects; should this not prove possible for some reason, separate measurements of negative and positive functioning should be obtained and examined vis-à-vis $G \times E$. In addition, once a $G \times E$ interaction has been discerned, follow-up analysis should determine whether significant differences in the functioning of individuals hypothesized to be more and less susceptible to environmental influences are obtained when the environmental circumstances are adverse as well as when they are supportive (that is, at both ends of the environmental continuum). It is when significant differences are obtained for both comparisons that differential susceptibility rather than diathesis–stress would be the correct inference.

Conflict of interest

The authors report no biomedical financial interests or potential conflicts of interest.

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