

# The 5-HTTLPR Polymorphism in the Serotonin Transporter Gene Moderates the Association Between Emotional Behavior and Changes in Marital Satisfaction Over Time

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Why do some individuals become dissatisfied with their marriages when levels of negative emotion are high and levels of positive emotions are low, whereas others remain unaffected? Using data from a 13-year longitudinal study of middle-aged and older adults in long-term marriages, we examined whether the 5-HTTLPR polymorphism in the serotonin transporter gene moderates the association between negative and positive emotional behavior (objectively measured during marital conflict) and changes in marital satisfaction over time. For individuals with two short alleles of 5-HTTLPR, higher negative and lower positive emotional behavior at Time 1 predicted declines in marital satisfaction over time (even after controlling for depression and other covariates). For individuals with one or two long alleles, emotional behavior did not predict changes in marital satisfaction. We also found evidence for a crossover interaction (individuals with two short alleles of 5-HTTLPR and low levels of negative or high levels of positive emotion had the highest levels of marital satisfaction). These findings provide the first evidence of a specific genetic polymorphism that moderates the association between emotional behavior and changes in marital satisfaction over time and are consistent with increasing evidence that the short allele of this polymorphism serves as a susceptibility factor that amplifies sensitivity to both negative and positive emotional influences.

*Keywords:* emotional behavior, genetic polymorphisms, 5-HTTLPR, relationships

Marriage plays an important role in the social fabric of our lives. According to the 2009 census, 96% of Americans over the age of 65 had been married at least once in their life. The quality of

marriage is an important contributor to health and well-being (Proulx, Helms, & Buehler, 2007). Spouses who are unhappy in their marriages are at a heightened risk for mental health problems (Whisman, 2007), physical health problems (Kiecolt-Glaser & Newton, 2001), as well as divorce (Gottman & Levenson, 2000). Thus, marital dissatisfaction has negative consequences for individuals, family members, and society alike.

In considering the foundations and functioning of marriage, emotions play an enormously important role (e.g., Gottman, 1994; Greenberg & Johnson, 1988). Early research on marriage was largely the province of sociologists, who developed excellent questionnaires to quantify spouses' level of satisfaction with their marriages (e.g., Locke & Wallace, 1959). These questionnaires were used in survey studies that explored the association between marital satisfaction and select demographic, personality, and other variables (e.g., Bentler & Newcomb, 1978). As marriage became of increasing interest to psychologists, researchers began directly observing marital interaction and discovered the powerful role that emotion plays in marital satisfaction and marital stability (Birchler, Weiss, & Vincent, 1975; Levenson & Gottman, 1983). A large body of research, conducted over the ensuing decades, docu-

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mented the general finding that high levels of negative emotion and low levels of positive emotion during marital interactions are associated with low levels of concurrent and future marital satisfaction (e.g., Carstensen, Gottman, & Levenson, 1995; Gottman & Levenson, 1992; Karney & Bradbury, 1995; Levenson & Gottman, 1983; Levenson & Gottman, 1985). Given this, it is not surprising that emotions have become a primary target for contemporary therapies that attempt to improve couples' relationships (e.g., Gottman & Gottman, 2008; Greenberg & Johnson, 1988).

To paraphrase Longfellow, into each marriage some negative emotion must fall. Nonetheless, the effects of these negative emotions may not be as bad for some spouses as for others. Most of us know couples who seem to thrive in a negative emotional climate that is characterized by arguments and bickering. For other couples, even a droplet of negative emotion is toxic. Similarly, some couples may become much more satisfied with their marriages in the face of positive emotions, while others may remain relatively unaffected. Thus, the link between emotional behavior and marital satisfaction may differ greatly among individual couples (Bradbury, Fincham, & Beach, 2000; Karney & Bradbury, 1995; see also Sullivan, Pasch, Johnson, & Bradbury, 2010).

Previous research has pointed toward important moderating factors that could influence the association between emotion and marital satisfaction. These include attachment styles (Mikulincer & Shaver, 2003), coping processes (Karney & Bradbury, 1995; Randall & Bodenmann, 2009), and capacity for forgiveness (Fincham, Stanley, & Beach, 2007). Thus, for example, a person with an insecure attachment style, low coping skills, and an inability to forgive would be expected to experience particularly sharp declines in marital satisfaction in the face of high levels of negative and low levels of positive emotions.

Surprisingly, there has been almost no research on genetic moderators of the association between emotion and marital satisfaction. This is even more puzzling because marital satisfaction is known to be partly heritable (with heritabilities for aspects of marital quality ranging from 13% to 28%; Kendler & Baker, 2007; e.g., Spotts et al., 2004). Moreover, there is good evidence for genetic influences on attachment (Caspers et al., 2009), pair bonding (Walum et al., 2008), and sensitivity to spousal affect (Schoebi, Way, Karney, & Bradbury, 2012). Yet, even though these kinds of studies confirm heritability, they do not establish the role that specific genes play in the association between emotion and marital satisfaction.

In the present study, we examined the 5-HTTLPR polymorphism in the promoter region of the serotonin transporter gene (*SLC6A4*) (e.g., Caspi et al., 2003; Lesch et al., 1996) as a potential moderator of the association between negative and positive emotional behavior and marital satisfaction. The 5-HTTLPR polymorphism has two common alleles (short and long), with the short allele leading to lower transcriptional efficiency of the serotonin transporter protein (Lesch et al., 1996) and lower levels of serotonin uptake.

Early studies on 5-HTTLPR typically used gene  $\times$  environment designs (often focusing on gene  $\times$  stress interactions) to study its influence on distal outcomes such as the development of psychopathology. In this early work, the short allele was found to be associated with a heightened risk for depression and suicide in the face of stress and adversity (e.g., Caspi et al., 2003; Kendler, Kuhn, Vittum, Prescott, & Riley, 2005; Roy, Hu, Janal, & Gold-

man, 2007; Taylor et al., 2006). Initial enthusiasm for these findings was dampened by replication issues and the appearance of two meta-analyses that concluded that the interaction effect of 5-HTTLPR  $\times$  stress did not predict depression (Munafò, Durrant, Lewis, & Flint, 2009; Risch et al., 2009). However, a more recent meta-analysis (Karg, Burmeister, Shedden, & Sen, 2011) that included all available studies (the earlier meta-analyses were more selective) found strong support for the effect. Similar affirmative meta-analytic findings have been obtained for effects of 5-HTTLPR on anxiety (Sen, Burmeister, & Ghosh, 2004).

The studies reviewed thus far focused on distal effects that may take decades to develop (e.g., depression). Another group of studies focused on more proximal effects of 5-HTTLPR, examining responses to well-controlled laboratory stimuli. These studies have consistently shown that individuals with the short allele exhibit heightened reactivity to negative stimuli including: (a) greater amygdala reactivity to negative faces (Hariri et al., 2002), (b) greater cortisol reactivity to social stress (Way & Taylor, 2010), (c) greater startle responses to loud noise (Brocke et al., 2006), and, of direct relevance to the present study, (d) greater reactivity to a partner's negative affect (Schoebi et al., 2012). In two large laboratory studies using independent subject samples, we (Gyurak et al., 2013) recently found that the short allele of 5-HTTLPR was associated with two different kinds of heightened emotional reactivity. In the first study, individuals with two short alleles showed greater personal distress and greater physiological responding when watching films of others in distress. In the second study, individuals with two short alleles reported more anger and amusement and showed greater emotional behavior in response to an embarrassing situation (viewing a video of oneself singing in a Karaoke-like task).

As research on the distal and proximal effects of 5-HTTLPR has matured, a model has emerged that moves away from early views of the short allele of 5-HTTLPR as a risk factor and instead views it as a plasticity or susceptibility factor (for a review see Belsky & Pluess, 2009). In this view, the short allele is not directly linked to good or bad outcomes, but rather is associated with amplified reactions. Thus, the short allele is not only associated with greater reactivity to the kinds of negative stimuli reviewed above, but also with greater reactivity to more positive stimuli such as positive images (Beevers et al., 2011) and positive partner affect (Schoebi et al., 2012). Developing this model a bit further, individuals with the short allele would be likely to have the worst outcomes under unfavorable conditions (e.g., those high in negative and low in positive features) and the best outcomes under favorable conditions (e.g., those low in negative and high in positive features). Statistically, this differential susceptibility takes the form of "crossover interactions",<sup>1</sup> which have now been demonstrated between 5-HTTLPR and various factors including early family environment and current adversity (Taylor et al., 2006), parenting (Hankin et al., 2011), socioeconomic status (Mitchell et al., 2011), and current life events (Pluess, Belsky, Way, & Taylor, 2010) when predicting outcomes such as well-being, neuroticism, and depression. In contrast to earlier studies that almost always focused on gene  $\times$  stress interaction effects, these studies explicitly con-

<sup>1</sup> However, not all crossover interactions indicate differential susceptibility (for details see Belsky & Pluess, 2009).

sider gene  $\times$  positive factor (or lack of stress) interaction effects (Belsky & Pluess, 2009).

An important developmental theme has also emerged in studies of genetic polymorphisms (e.g., Lenroot & Giedd, 2011). Lifespan developmental theories propose that human aging magnifies the importance of biological factors (Baltes, 1997), and empirical studies suggest that genes in particular become increasingly influential later in life (Lindenberger et al., 2008). For example, human aging appears to magnify the effects of genetic polymorphisms associated with the dopamine system as the efficiency of the dopamine system declines with age (e.g., Nagel et al., 2008; Stürmer, Passow, Biesenack, & Li, 2012). The efficiency of the serotonin system has likewise been shown to decline with age in terms of both central serotonin transporter availability (van Dyck et al., 2000) and postsynaptic serotonin receptors (Meltzer et al., 1998). Thus, it is possible that the effects of serotonin-related genes such as 5-HTTLPR are also magnified with age. The longitudinal design utilized in the present study, which focuses on middle age and late life, may be particularly useful for examining the influence of aging on the effects of 5-HTTLPR (see also McArdle & Prescott, 2010).

### The Present Study

We examined how 5-HTTLPR moderates the association between positive and negative emotional behavior (rated by trained coders) during marital conflict and changes in marital satisfaction in a 13-year longitudinal study of long-term marriages in middle age and late life. This study extends previous research on 5-HTTLPR by focusing on marital satisfaction, objectively coding emotional behavior that occurred during naturalistic marital interactions, including middle-aged and older participants, and using a longitudinal design. Building on prior work (Caspi, Hariri, Holmes, Uher, & Moffitt, 2010), our primary hypothesis was that for individuals with the short allele of 5-HTTLPR, higher negative and lower positive emotional behavior at Time 1 (T1) would predict greater declines in marital satisfaction over time. We designed our analyses to take into account: (a) associations between positive and negative emotional behavior; (b) different possibilities for the dominance structure of 5-HTTLPR (Caspi et al., 2010); (c) statistical nonindependence between husbands' and wives' data; (d) the alternative explanation that the effect of 5-HTTLPR  $\times$  emotional behavior on marital satisfaction would be driven by depression (cf. Caspi et al., 2003) or other covariates; (e) the possibility of a crossover interaction of 5-HTTLPR  $\times$  emotional behavior (Belsky & Pluess, 2009); and (f) whether findings generalized across gender (Caspi et al., 2010; Williams et al., 2003), cohort (Nagel et al., 2008), and ethnicity (Way & Lieberman, 2010; Williams et al., 2003).

### Method

#### Participants

We analyzed data from a longitudinal study of long-term married couples consisting of a middle-aged (T1: age 40–50) and an older (T1: age 60–70) cohort. This sample of couples was recruited originally in 1989–1990 by a survey research company to be representative of a random sample of marriages in the San

Francisco Bay Area in terms of socioeconomic status, religion, ethnicity, and marital satisfaction (for full details see Levenson, Carstensen, & Gottman, 1993). We examined a subsample ( $N = 125$ ) who participated in genetic testing in 2009 (in 51 couples both spouses participated; in 23 couples one spouse participated). Spouses who participated in genetic testing did not differ from the other participants on the variables examined here except for showing higher positive emotional behavior at T1,  $p = .025$ . There were 74 middle-aged adults and 51 older adults (50% females; years of education:  $M = 15.97$ ,  $SD = 2.74$ ; 84.8% Caucasian, 4.8% African American, 4.0% Hispanic, 4.8% Latino/a, 1.6% Other). The genetic data from this sample have not been used in any prior publications.

Analyses draw from three waves of assessment, Time 1 (T1): 1989/90,  $n = 123$  with complete data on all variables analyzed in the present study; Time 2 (T2): 1995/96,  $n = 117$ ; Time 3 (T3): 2001/02,  $n = 107$ . None of the variables examined here predicted drop-out over time ( $ps > .05$ ).

#### Procedure

At each assessment, couples completed a set of questionnaires including measures of marital satisfaction (see below). They then came to our laboratory at Berkeley and participated in a well-established procedure for studying marital interaction (Levenson & Gottman, 1983). This procedure consisted of having couples engage in three 15-min unrehearsed discussions of topics related to their marriage: (a) events of the day (at T1) or events since the last assessment (at subsequent waves); (b) a topic of continuing disagreement in their marriage (conflict topic); and (c) something they enjoyed doing together (pleasant topic). Conversations were recorded on videotape for subsequent analysis of emotional behavior and a number of physiological measures were recorded continuously from each spouse. For the present study, the emotional behavioral data obtained during the conflict conversation were used.

#### Measures

**5-HTTLPR.** DNA was collected and extracted from saliva using Oragene kits (DNA Genotek, Kanata, Ontario, Canada) according to manufacturer's protocol. Anonymized DNA samples were extracted and purified by the Abbott-UCSF Viral Diagnostics and Discovery Center, San Francisco, CA. The extracted DNA was genotyped using the procedure described in Assal et al. (2004) with slight modifications. A PCR product was amplified with primers (5'-GGCGTTGCCGCTCTGAATGC-3' and 5'-GAGGGACTGAGCTGGACAACCA-3') flanking the region containing the gene variation. The PCR conditions consisted of a 2-min denaturation step at 94 °C, 35 cycles of 30-s denaturation at 95 °C, 30-s annealing at 60 °C, and 30-s extension at 72 °C, and a final 7-min extension step at 72 °C. Genotype distribution (two short alleles:  $n = 21$ , 16.8%; one long allele:  $n = 70$ , 56%; two long alleles:  $n = 34$ , 27.2%) was consistent with previous studies (e.g., Caspi et al., 2003). Table 1 presents the distribution of the 5-HTTLPR genotypes in the total sample broken down by ethnicity as well as results for the Hardy-Weinberg equilibrium test. No deviations from Hardy-Weinberg equilibrium were detected,  $ps > .05$ , similar to other samples (e.g., Schoebi et al., 2012).

Table 1  
5-HTTLPR Genotype Frequencies by Ethnicity

	5-HTTLPR Genotype			HWE $\chi^2$ (df = 1)
	s/s	s/l	l/l	
Caucasians ( $n = 106$ )	16 (15.1%)	59 (55.7%)	31 (29.2%)	1.96, $p = .162$
Other ( $n = 19$ )	5 (26.3%)	11 (57.9%)	3 (15.8%)	.55, $p = .457$
Total sample ( $N = 125$ )	21 (16.8%)	70 (56.0%)	34 (27.2%)	2.19, $p = .139$

Note. Absolute (relative) frequencies (s/s = short/short, s/l = short/long, l/l = long/long). HWE = Hardy-Weinberg equilibrium.

5-HTTLPR genotyping using the methods we used has generally shown high reliability. Specifically, when we conducted a blind reanalysis of 8 DNA samples from a larger genotyping project that included samples from all of the participants analyzed here, genotype reproducibility was 100%.

**Emotional behavior.** Emotional behavioral during the T1 conflict conversation was rated by trained coders (blind to participants' genetic polymorphisms, marital satisfaction, and study hypotheses) using the Specific Affect Coding System (SPAFF; Gottman, 1996). SPAFF classifies emotional behaviors on a second-by-second basis based on verbal content, vocal tone, context, facial expression, gesture, and body movement. Interrater reliability for SPAFF was satisfactory in the present study (for detailed information, see Carstensen et al., 1995). There were nine negative speaker codes, five positive speaker codes, and one neutral speaker code. For the present study, we focused on the negative and positive speaker codes and counted the relative frequency of each negative and positive speaker emotion coded during each second of the 15-min conflict conversation (i.e., percentage of seconds in which each emotion code was present during the conversation). We obtained a composite score for *negative emotional behavior* by averaging across all negative speaker codes (i.e., anger, belligerence, contempt, defensiveness, disgust, domineering, fear/tension/worry, sadness, whining) and a composite score for *positive emotional behavior* by averaging across all positive emotion speaker codes (i.e., affection, humor, interest, joy, validation). As would be expected given the nature of the conflict conversation, the levels of negative emotional behavior ( $M = 4.86$ ,  $SD = 3.34$ ) were greater than the levels of positive emotional behavior ( $M = 2.23$ ,  $SD = 1.86$ ),  $t(122) = 6.38$ ,  $p < .001$ .

**Marital satisfaction.** Marital satisfaction was measured at T1, T2, and T3 by averaging two well-established self-report measures: (a) the Marital Adjustment Test (15 items; T1:  $\alpha = .77$ ; T2:  $\alpha = .77$ ; T3:  $\alpha = .76$ ; Locke & Wallace, 1959) and (b) the Marital Relationship Inventory (22 items; T1:  $\alpha = .87$ ; T2:  $\alpha = .85$ ; T3:  $\alpha = .87$ ; Burgess, Locke, & Thomes, 1971). Across waves of data collection, these two self-report measures were highly correlated, T1:  $r = .89$ ; T2:  $r = .85$ ; T3:  $r = .83$ . Across waves of measurement, there were no changes in mean levels of marital satisfaction, as indicated by nonsignificant slope means ( $ps > .05$ ). However, there were sizable interindividual differences in changes in marital satisfaction, as indicated by significant slope variances ( $ps < .05$ ).

**Covariates.** Covariates were depression (using the respective subscale from the SCL-90; 13 items; T1:  $\alpha = .88$ ; T2:  $\alpha = .90$ ; T3:  $\alpha = .92$ ; Derogatis & Cleary, 1977), ethnicity (0 = *else*; 1 = *Caucasian*), gender (0 = *male*; 1 = *female*), cohort (0 = *middle-*

*aged*; 1 = *older*), and education (in years). Analyses were also controlled for positive emotional behavior when analyzing negative emotional behavior (and vice versa).

### Statistical Analyses

Statistical analyses involving the three 5-HTTLPR genotypes (that is, short/short, short/long, long/long alleles) have been handled in different ways, reflecting different views as to the dominance or codominance of the alleles. A recent review (Caspi et al., 2010) found dominance of the short allele in 9 studies, dominance of the long allele in 10 studies, and codominance in 14 studies. Because this evidence is so mixed, we chose a statistical approach that enabled us to consider all possible dominance structures of 5-HTTLPR. Building on established "thermometer-coding" procedures (see Kendler et al., 2005) that we have used in previous studies (Gyurak et al., 2013), we coded 5-HTTLPR using two dummy variables, 5-HTTLPR (H1; 0 = *long/long*; 1 = *short/long*, *short/short*) and 5-HTTLPR (H2; 0 = *short/long*, *long/long*; 1 = *short/short*). This coding is well-suited for studying the 5-HTTLPR genotype because it anchors the analyses at the most sensitive group, individuals with two short alleles (coded as 1, 1), extends all the way to individuals with two long alleles (coded as 0, 0), and assigns intermediate levels to individuals with one short and one long allele (coded as 1, 0). A significant effect for 5-HTTLPR (H1) indicates that individuals with two long alleles (l/l) differ from individuals with one or two short alleles (s/l or s/s), thus indicating dominance of the short allele. A significant effect for 5-HTTLPR (H2) indicates that individuals with two short alleles (s/s) differ from individuals with one or two long allele (s/l or l/l), thus indicating dominance of the long allele.

We conducted separate multiple linear regressions predicting marital satisfaction at T2 and T3 from (a) negative or positive emotional behavior at T1, (b) 5-HTTLPR thermometer-coded variables (H1 and H2), and (c) interaction terms between 5-HTTLPR (H1 and H2)  $\times$  negative or positive emotional behavior. Covariates were depression, ethnicity, gender, cohort, education, the other kind of emotional behavior (positive when analyzing negative emotional behavior and vice versa), and marital satisfaction at T1. In follow-up analyses, to investigate whether findings generalized across gender and cohort, we examined 3-way interactions between gender or cohort, 5-HTTLPR (H2), and negative or positive emotional behavior. To evaluate effects of ethnicity, we also examined whether findings held when repeating analyses using only the Caucasian participants ( $n = 106$ ). Finally, in a follow-up analysis, we investigated whether findings remained stable when controlling for changes in depression across the respective time

interval instead of depression at T1. Unless noted otherwise, all findings remained stable. All continuous variables were mean-centered for the regression analyses.

Analyses were conducted using Stata 10 (StataCorp., 2007) and focused on within-spouse associations. Because of the dyadic nature of our data, we used Stata's cluster tool, which corrects all results for nonindependence between husbands and wives. Because we had complete data for only 51 couples (for the other 23 couples only one spouse per couple participated in genetic testing) we were underpowered to examine cross-spouse associations in greater depth. However, to obtain some idea of whether our primary hypothesis would also hold for cross-spouse associations, we employed a multigroup actor-partner modeling approach within a structural equation modeling framework (Olsen & Kenny, 2006) using AMOS (Arbuckle, 2008) to study how wives' and husbands' emotional behavior predicted changes in wives' and husbands' marital satisfaction depending on wives' 5-HTTLPR genotype (the models studying the effects of husbands' 5-HTTLPR genotype did not converge because of small cell sizes and thus were not tested.). For the actor-partner model, we used residualized change scores to model changes in marital satisfaction (by predicting marital satisfaction at T3 from marital satisfaction at T1 and saving the residuals for further analysis) to reduce the number of parameters to be estimated.

## Results

### Preliminary Analyses

Examination of intercorrelations between key study variables at T1 (Table 2) showed that negative and positive emotional behavior were negatively correlated. As expected (Carstensen et al., 1995), low levels of marital satisfaction were associated with high levels of negative emotional behavior and low levels of positive emotional behavior at T1. The 5-HTTLPR variables (H1 and H2) were not correlated with negative or positive emotional behavior or with marital satisfaction.

### 5-HTTLPR Moderation of the Association Between Emotional Behavior and Changes in Marital Satisfaction

Results from the regression analyses examining emotional behavior and 5-HTTLPR predicting changes in marital satisfaction

are presented in Table 3 and 4. Consistent with our primary hypothesis, an interaction effect was found between negative emotional behavior at T1 and 5-HTTLPR (H2) when predicting changes in marital satisfaction over the longest (13 years) time interval, from T1 to T3,  $B = -2.65$ ,  $SE(B) = 1.05$ ,  $p = .015$ , Partial  $R^2 = .08$ .<sup>2</sup> Similarly, an interaction effect was found between positive emotional behavior at T1 and 5-HTTLPR (H2) when predicting changes in marital satisfaction from T1 to T3,  $B = 3.48$ ,  $SE(B) = 1.27$ ,  $p = .008$ , Partial  $R^2 = .06$ . Statistical power for detecting these interaction effects (determined using GPower; Faul, Erdfelder, Lang, & Buchner, 2007) was .87 and .75, respectively. Interaction effects between negative or positive emotional behavior and 5-HTTLPR (H2) when predicting changes in marital satisfaction from T1 to T2 were not significant,  $ps > .05$ . No interaction effects emerged between negative or positive emotional behavior and the other dummy variable, 5-HTTLPR (H1), at any of the waves of data collection,  $ps > .05$ . Thus, in accordance with prior research (Kendler et al., 2005), we collapsed individuals with one or two long alleles into one group.

To control for possible spurious effects due to multiple testing (8 tests total, consisting of 4 interaction tests predicting changes in marital satisfaction at T2 and T3) we used procedures developed by Storey and colleagues (e.g., Storey & Tibshirani, 2003). Results showed that 5-HTTLPR still interacted with both negative and positive emotional behavior to predict changes in marital satisfaction (all  $qs < .05$  at a false discovery rate of 5%).

Follow-up analyses were then conducted to decompose the significant interaction effects, contrasting individuals with two short alleles with the other 5-HTTLPR variants. These analyses provided strong support for our primary hypothesis. For individuals with two short alleles of 5-HTTLPR, higher negative emotional behavior at T1 predicted declines in marital satisfaction from T1 to T3,  $B = -2.41$ ,  $SE(B) = 1.00$ ,  $p = .019$ . For individuals with one or two long alleles of 5-HTTLPR, negative emotional behavior at T1 did not predict changes in marital satisfaction from T1 to T3,  $B = .02$ ,  $SE(B) = .29$ ,  $p = .947$ . In a similar vein, for individuals with two short alleles of 5-HTTLPR, lower positive emotional behavior at T1 predicted declines in marital satisfaction from T1 to T3,  $B = 2.10$ ,  $SE(B) = 1.04$ ,  $p = .048$ . Somewhat unexpectedly, for individuals with one or two long alleles, higher positive emotional behavior at T1 predicted declines in marital satisfaction,  $B = -1.23$ ,  $SE(B) = .58$ ,  $p = .037$ . However, this association was no longer statistically significant at the .05 level when controlling for changes in depression from T1 to T3 (instead of depression at T1),  $B = -1.08$ ,  $SE(B) = .58$ ,  $p = .067$ .

Table 2  
Intercorrelations of Key Study Variables at T1

	1.	2.	3.	4.	5.
1. Negative emotional behavior T1	—				
2. Positive emotional behavior T1	-.51***	—			
3. 5-HTTLPR (H1) <sup>a</sup>	-.01	.01	—		
4. 5-HTTLPR (H2) <sup>a</sup>	.07	-.06	.27**	—	
5. Marital satisfaction T1	-.27**	.21*	.05	.00	—

<sup>a</sup> 5-HTTLPR genotype was thermometer-coded (see Kendler et al., 2005; for a detailed description see text) using two dummy variables: 5-HTTLPR (H1) (0 = long/long; 1 = short/long, short/short) and 5-HTTLPR (H2) (0 = short/long, long/long; 1 = short/short).

\*  $p < .05$ . \*\*  $p < .01$ . \*\*\*  $p < .001$ .

<sup>2</sup> Our hypotheses focused on predicting changes in marital satisfaction over time. Note that an interaction effect between negative emotional behavior at T1 and 5-HTTLPR (H2) was already found when predicting marital satisfaction at T1,  $B = -2.44$ ,  $SE(B) = 1.18$ ,  $p = .042$ . When decomposing the interaction, a similar pattern of results was obtained. For individuals with two short alleles of 5-HTTLPR, higher negative emotional behavior predicted lower marital satisfaction,  $B = -3.37$ ,  $SE(B) = 1.12$ ,  $p = .004$ . For individuals with one or two long alleles of 5-HTTLPR, negative emotional behavior did not predict marital satisfaction,  $B = -.59$ ,  $SE(B) = .48$ ,  $p = .220$ .

Table 3  
*Negative Emotional Behavior, 5-HTTLPR, and Their Interaction as Predictors of Changes in Marital Satisfaction Over Time*

	T2 Marital satisfaction				T3 Marital satisfaction			
	<i>B</i>	<i>SE (B)</i>	95% Confidence interval		<i>B</i>	<i>SE (B)</i>	95% Confidence interval	
Negative emotional behavior T1	-.30	.78	-1.86	1.25	-.29	.62	-1.53	.94
5-HTTLPR (H1) <sup>a</sup>	3.93	2.15	-.37	8.23	-.14	1.95	-4.04	3.77
5-HTTLPR (H2) <sup>b</sup>	-3.96	2.35	-8.66	.73	-3.97	2.39	-8.75	.81
Negative emotional behavior T1 × 5-HTTLPR (H1) <sup>a</sup>	.65	.81	-.97	2.27	.47	.71	-.96	1.90
Negative emotional behavior T1 × 5-HTTLPR (H2) <sup>b</sup>	-.66	1.21	-3.08	1.77	-2.65*	1.05	-4.76	-.54
	<i>F</i> (12, 66) = 18.51; <i>R</i> <sup>2</sup> = .77				<i>F</i> (12, 58) = 14.27; <i>R</i> <sup>2</sup> = .78			

Note. Results from regression analyses controlling for depression, ethnicity, gender, cohort, education, positive emotional behavior, and marital satisfaction at T1. Standard errors corrected for clustering within couples.

<sup>a</sup> 5-HTTLPR (H1) (0 = long/long; 1 = short/long, short/short). <sup>b</sup> 5-HTTLPR (H2) (0 = short/long, long/long; 1 = short/short).

\*  $p < .05$ .

### Crossover Interactions

As shown in Figure 1 and 2, interaction effects between 5-HTTLPR and emotional behavior were in fact crossover interactions. That is, individuals with two short alleles of 5-HTTLPR had: (a) the lowest marital satisfaction at high levels of negative and low levels of positive emotional behavior and (b) the highest marital satisfaction at low levels of negative and high levels of positive emotional behavior. To illustrate the nature of the crossover interaction further, Figure 3 plots the changes in marital satisfaction from T1 to T3 at low versus high levels of negative emotional behavior at T1 depending on the 5-HTTLPR genotype.

### Effects of Gender, Cohort, and Ethnicity

We explored the generalizability of these findings across gender, cohort, and ethnicity. Findings for positive emotional behavior generalized across gender and cohort, as indicated by nonsignificant 3-way interactions involving gender or cohort, 5-HTTLPR (H2), and positive emotional behavior when predicting changes in marital satisfaction from T1 to T3,  $ps > .05$ .

In contrast, findings for negative emotional behavior were qualified by gender and cohort, as indicated by significant 3-way interactions,  $ps < .05$ . In terms of gender, the prediction of declines in marital satisfaction by having two short alleles appeared stronger for wives,  $B = -3.47$ ,  $SE(B) = 1.21$ ,  $p = .006$ , than for husbands,  $B = -2.13$ ,  $SE(B) = 1.03$ ,  $p = .043$ . Among individuals with one or two long alleles, negative emotional behavior did not predict declines in marital satisfaction for wives,  $B = -.24$ ,  $SE(B) = .31$ ,  $p = .439$ , or for husbands,  $B = -.37$ ,  $SE(B) = .41$ ,  $p = .379$ . In terms of cohort, higher negative emotional behavior predicted a decline in marital satisfaction for older adults with two short alleles,  $B = -3.07$ ,  $SE(B) = .69$ ,  $p < .001$ , but not for middle-aged adults with two short alleles,  $B = 1.22$ ,  $SE(B) = 1.39$ ,  $p = .385$ . Among individuals with one or two long alleles, negative emotional behavior did not predict declines in marital satisfaction for older adults,  $B = -.11$ ,  $SE(B) = .47$ ,  $p = .821$ , or for middle-aged adults,  $B = .20$ ,  $SE(B) = .32$ ,  $p = .527$ . To summarize, the moderating effect of 5-HTTLPR on the association between negative emotional behavior and changes in marital satisfaction appeared stronger for wives than for husbands and was present for older spouses but not for middle-aged spouses.

Finally, when repeating all analyses including only Caucasian participants ( $n = 106$ ), the results remained essentially unchanged. One association (negative emotional behavior predicting decline in marital satisfaction from T1 to T3 among husbands with two short alleles) was reduced to trend level,  $p = .070$ .

### Cross-Spouse Associations

Exploratory actor-partner analyses provided some evidence for cross-spouse associations in the expected direction.<sup>3</sup> For wives with two short alleles, both wives' higher negative emotional behavior,  $B = -.26$ ,  $SE(B) = .08$ ,  $p = .001$  as well as husbands' higher negative emotional behavior,  $B = -.19$ ,  $SE(B) = .08$ ,  $p = .017$ , predicted declines in wives' marital satisfaction from T1 to T3 (controlling for husbands' and wives' positive emotional behavior). For wives with one or two long alleles, neither wives' own negative emotional behavior,  $B = -.03$ ,  $SE(B) = .06$ ,  $p = .641$ , nor husbands' negative emotional behavior,  $B = .03$ ,  $SE(B) = .09$ ,  $p = .715$ , predicted changes in wives' marital satisfaction from T1 to T3. These results remained stable when controlling for covariates. The multigroup model exploring the effects of husbands' 5-HTTLPR genotype were not tested because of small cell sizes.

### Discussion

To our knowledge, the present findings are the first to implicate a specific genetic polymorphism as playing an important role in moderating the association between emotional behavior and changes in marital satisfaction. Consistent with our primary hypothesis, for individuals who had two short alleles of 5-HTTLPR, higher negative and lower positive emotional behavior at T1 predicted declines in marital satisfaction over time. In contrast, for individuals with one or two long alleles, levels of negative and positive emotional behavior did not predict

<sup>3</sup> In addition, this actor-partner model approach revealed that changes in wives' and husbands' marital satisfaction from T1 to T3 were not correlated ( $r = .03$ ,  $p = .854$ ), consistent with the notion that there may be two marriages, his and hers (Barnard, 1982). Moreover, both change residuals for wives' and husbands' marital satisfaction showed significant variability,  $\sigma^2 = 1.02$ ,  $p < .001$  and  $\sigma^2 = .85$ ,  $p < .001$ , indicating that different individuals changed in different directions over time.

Table 4  
Positive Emotional Behavior, 5-HTTLPR, and Their Interaction as Predictors of Changes in Marital Satisfaction Over Time

	T2 Marital satisfaction			T3 Marital satisfaction		
	B	SE (B)	95% Confidence interval	B	SE (B)	95% Confidence interval
Positive emotional behavior T1	2.51*	1.05	.42 4.61	-.86	1.37	-3.61 1.89
5-HTTLPR (H1) <sup>a</sup>	3.94	2.06	-.17 8.06	-.18	1.95	-4.07 3.72
5-HTTLPR (H2) <sup>b</sup>	-3.85	2.19	-8.21 .52	-3.11	2.33	-7.77 1.55
Positive emotional behavior T1 × 5-HTTLPR (H1) <sup>a</sup>	-2.36	1.22	-4.79 .07	-.50	1.43	-3.37 2.37
Positive emotional behavior T1 × 5-HTTLPR (H2) <sup>b</sup>	2.23	1.41	-.60 5.04	3.48**	1.27	.95 6.02
	$F(12, 66) = 16.75; R^2 = .78$			$F(12, 58) = 14.27; R^2 = .78$		

Note. Results from regression analyses controlling for depression, ethnicity, gender, cohort, education, negative emotional behavior, and marital satisfaction at T1. Standard errors corrected for clustering within couples.

<sup>a</sup> 5-HTTLPR (H1) (0 = long/long; 1 = short/long, short/short). <sup>b</sup> 5-HTTLPR (H2) (0 = short/long, long/long; 1 = short/short).

\*  $p < .05$ . \*\*  $p < .01$ .

changes in marital satisfaction. Although one effect could already be seen at T1, the findings were most pronounced when predicting changes in marital satisfaction across the full 13-year range of data collection. This is consistent with a view that genetic influences may increase over time and/or age (Lindenberg et al., 2008; McArdle & Prescott, 2010). Given earlier concerns about the replicability of these kinds of single-gene effects (e.g., Risch et al., 2009), we designed our analyses to be maximally conservative by: (a) controlling for variance in marital satisfaction shared by positive and negative emotional behavior; (b) considering different possibilities for the dominance structure of 5-HTTLPR; (c) correcting for nonindependence between husbands and wives; and (d) controlling for the pos-

sible effects of depression as well as ethnicity, gender, cohort, and education.

Although our primary hypothesis involved the association between an unfavorable emotional climate (i.e., high levels of negative emotion and low levels of positive emotion) and declines in marital satisfaction, our findings also provided evidence for the kind of crossover interaction involving 5-HTTLPR that has been reported by others (e.g., Belsky & Pluess, 2009; Mitchell et al., 2011; Taylor et al., 2006). In our study, individuals with two short alleles of 5-HTTLPR also had the highest marital satisfaction in a favorable emotional climate (i.e., low levels of negative emotion and high levels of positive emotion). This finding is consistent with a growing body of empirical work showing that the short

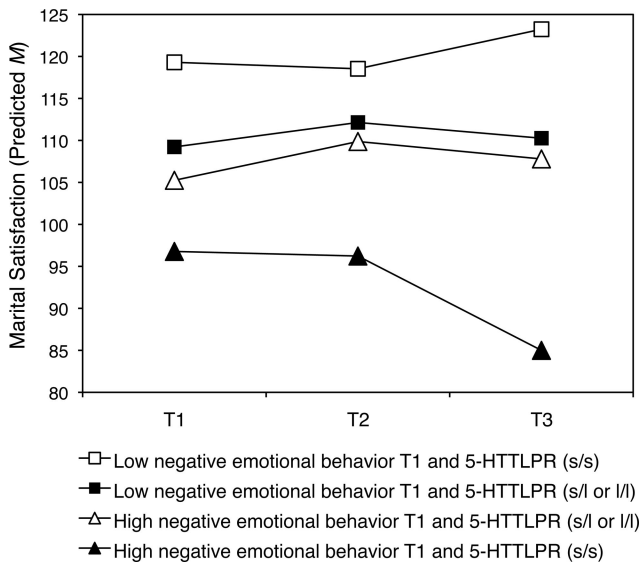


Figure 1. Marital satisfaction over 13 years by negative emotional behavior and 5-HTTLPR. Note. Predicted uncentered means of marital satisfaction at T1, T2, and T3 for individuals with high ( $M + 1 SD$ ) or low ( $M - 1 SD$ ) levels of negative emotional behavior at T1 and two short alleles (s/s) or one or two long alleles (s/l or l/l) of 5-HTTLPR. Analyses controlled for depression, ethnicity, gender, cohort, education, and positive emotional behavior at T1. T1: 1989/90. T2: 1995/96. T3: 2001/02.

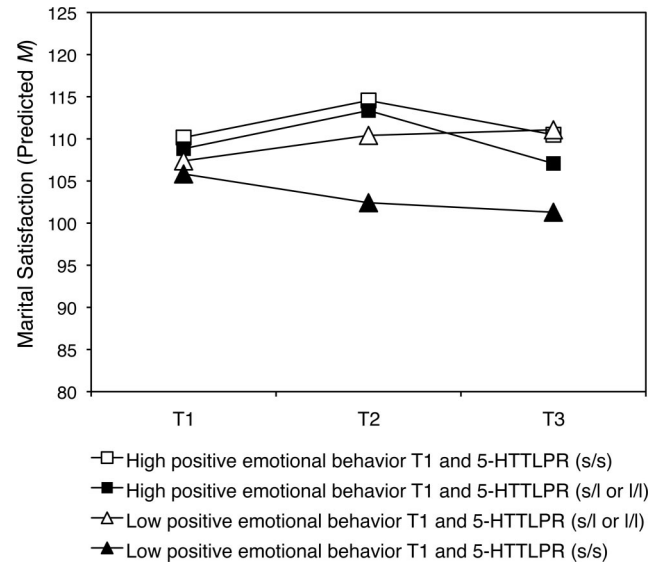
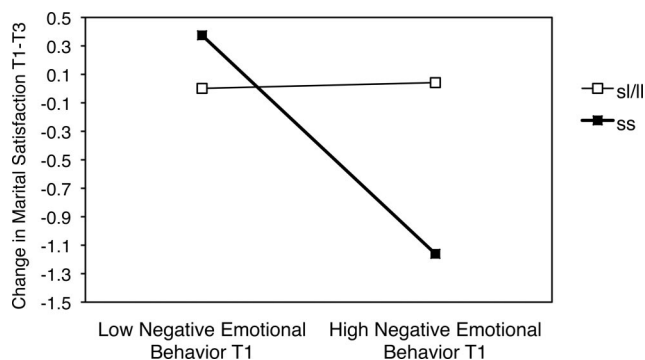


Figure 2. Marital satisfaction over 13 years by positive emotional behavior and 5-HTTLPR. Note. Predicted uncentered means of marital satisfaction at T1, T2, and T3 for individuals with high ( $M + 1 SD$ ) or low ( $M - 1 SD$ ) levels of positive emotional behavior at T1 and two short alleles (s/s) or one or two long alleles (s/l or l/l) of 5-HTTLPR. Analyses controlled for depression, ethnicity, gender, cohort, education, and negative emotional behavior at T1. T1: 1989/90. T2: 1995/96. T3: 2001/02.



**Figure 3.** Crossover interaction between negative emotional behavior and 5-HTTLPR predicting changes in marital satisfaction over 13 years. *Note.* Residualized change in marital satisfaction from T1 to T3 for individuals with low ( $M - 1 SD$ ) or high ( $M + 1 SD$ ) levels of negative emotional behavior at T1 and two short alleles (s/s) or one or two long alleles (s/l or l/l) of 5-HTTLPR. Analyses controlled for depression, ethnicity, gender, cohort, education, and positive emotional behavior at T1. T1: 1989/90. T3: 2001/02.

allele amplifies reactivity to positive as well as negative emotional stimuli (e.g., Beevers et al., 2011; Gyurak et al., 2013; Hariri et al., 2002; Homberg & Lesch, 2011; Schoebi et al., 2012; Way & Taylor, 2010). These findings suggest that viewing a particular 5-HTTLPR variant as inherently “bad” or “good” is unwise. Rather, whether the short allele results in bad or good outcomes in the long term will depend on the presence or absence of positive and negative factors.

For negative emotional behavior, our analyses also revealed several interesting differences related to gender and age. In terms of gender, the 5-HTTLPR effects on the relationship between negative emotional behavior and marital satisfaction were stronger for females than for males. This is consistent with prior findings that 5-HTTLPR effects can be more pronounced for females; however, this pattern has not always been found (for an overview see, e.g., Caspi et al., 2010). In terms of age, the 5-HTTLPR effects on the association between negative emotional behavior and marital satisfaction were present for older spouses but not for middle-aged spouses. These findings are consistent with both theory (Lindenberger et al., 2008) and empirical findings (e.g., Nagel et al., 2008; Stürmer et al., 2012) that suggest that genes become increasingly influential in late life. Interestingly, these gender and age effects were only found for negative emotions and not for positive emotions. This is consistent with findings that negative behaviors have a greater overall effect on marital satisfaction than positive behaviors (see Karney & Bradbury, 1995) and with the more generalized notion that “bad is stronger than good” (Baumeister, Bratslavsky, Finkenauer, & Vohs, 2001). However, it is important to realize that the present study was based on emotional behaviors that occurred when couples discussed a relationship problem. Thus, it is possible that moderation effects for positive emotions might have emerged in a context more conducive to the production of positive emotional behaviors.

Understanding the role that specific genes play in moderating the association between emotional behaviors and marital satisfaction may help explain the variability in this association that is found in popular lore (e.g., some loving couples love to bicker) and

in the marital research literature (Bradbury et al., 2000; Karney & Bradbury, 1995). Our study examined broad classes of emotional behaviors (negative and positive) during a particular kind of marital interaction (discussing a marital problem). We expect that this association could be profitably explored in terms of more specific emotions and other contexts. For example, some negative emotions, such as sadness, may promote marital satisfaction when they involve revealing vulnerabilities and attempting to elicit support (Clark, Ouellette, Powell, & Milberg, 1987; Graham, Huang, Clark, & Helgeson, 2008). Some positive emotions, for example amusement at the expense of the partner, could undermine marital satisfaction (Levenson, Haase, Bloch, Holley, & Seider, *in press*). Cultural variations likely also come into play, with different traditions of emotional expression (Soto, Levenson, & Ebling, 2005) and different views as to which emotions and levels of emotional intensity are most desirable (Tsai, Knutson, & Fung, 2006). One implication of the present study is that short-allele carriers of 5-HTTLPR would be expected to show stronger associations between emotional behavior and marital satisfaction regardless of the direction of that association, as determined by specific emotions, contexts, and cultures.

Viewed in a broader framework, the present study extends previous research that has demonstrated genetic influences on relationship outcomes in adulthood (e.g., Schoebi et al., 2012; Walum et al., 2008) into the realm of marital satisfaction. Similarly, it extends research that has demonstrated that 5-HTTLPR moderates the association between risk factors and outcomes in the domain of individual psychopathology (e.g., Caspi et al., 2003; Karg et al., 2011) to show similar effects in the social, interpersonal domain of marriage. Our study can be seen as falling within the gene  $\times$  stress framework that characterized early studies of 5-HTTLPR (Caspi et al., 2003), although the early studies focused on more “external” factors (e.g., stressful life events) and ours focused more on “internal” factors (e.g., emotional behaviors that emerged during spousal interactions). The present findings thus contribute to a biopsychosocial view of marriage and family (e.g., Horwitz & Neiderhiser, 2011; Kiecolt-Glaser & Newton, 2001).

### Strengths and Limitations

To our knowledge, this is the first study that shows that a specific genetic polymorphism (5-HTTLPR) moderates the association between objectively coded emotional behaviors and marital satisfaction in a longitudinal design. The study had several additional strengths: (a) the duration of the longitudinal follow-up was substantial (13 years), (b) participants were in well-established long-term marriages in middle age and late life, and (c) emotional behaviors were sampled during a naturalistic discussion of a marital problem. Our finding that having two short alleles of 5-HTTLPR was associated with a stronger association between the emotional quality of marital interaction and spousal marital satisfaction is consistent with newer views of the role this allele variant plays as a plasticity factor (Belsky & Pluess, 2009) that increases susceptibility to negative as well as positive conditions. The findings also extend our prior work investigating the influence of this polymorphism in the realm of emotional reactivity (Gyurak et al., 2013). In one of these studies, we found that having two short alleles was associated with greater emotional reactivity (self-report and physiology) to viewing the distress of others. In the other



study, we found that having two short alleles was associated with greater emotional reactivity (self-report and objectively coded behavior) in an embarrassing situation. The present study extended this work into the realm of gene  $\times$  stress associations, finding that having two short alleles strengthened the association between emotional behavior and a more distal outcome (marital satisfaction). Given that these three studies were performed on three independent samples of participants and utilized three different experimental paradigms underscores the robustness of the effects of 5-HTTLPR in the realm of emotion (or at least when using these kinds of well-controlled laboratory procedures and multimethod assessments of emotion).

This study also had several limitations worthy of note. Our sample was recruited to be representative of the San Francisco Bay Area and thus was primarily Caucasian and well-educated. Thus, the generalizability of these findings to other populations needs to be established. Our sample size ( $N = 125$  individuals) was small by the standards of population-based genetics studies (although it is quite respectable by the standards of observational studies of marital interaction) and thus we might have been relatively underpowered to explore the full range of influences of factors such as gender and age. A larger sample size would have enabled us to explore cross-spouse associations in greater depth (our exploratory analyses suggested that these might exist and that they parallel those found in within-spouse analyses). A larger sample size would have also given us greater power to detect possible codominant effects of the 5-HTTLPR alleles (i.e., additional differences between individuals with one and two long alleles) and possible 3-way interaction effects involving positive emotional behavior (we found that some associations only held for negative emotional behavior). Finally, there are additional subdivisions of 5-HTTLPR (e.g., SNP rs25531; Wendland, Martin, Kruse, Lesch, & Murphy, 2006) that merit additional study using these kinds of longitudinal experimental designs and behavioral data.

### Conclusions

The search for the recipe that produces a successful marriage has long occupied scientists, therapists, and laypeople alike. Among the key ingredients, an important role for emotion has been endorsed in many empirical studies (e.g., Carstensen et al., 1995; Gottman & Levenson, 1992; Karney & Bradbury, 1995; Levenson & Gottman, 1983), in a number of marital therapies (Gottman & Gottman, 2008; Lebow, Chambers, Christensen, & Johnson, 2012), and in folk wisdom. Here we present the first evidence that a common genetic polymorphism in the serotonin transporter gene plays an important role in determining how powerfully negative and positive emotions will influence marital satisfaction over time. For those with two short alleles of 5-HTTLPR, we would expect that marital satisfaction will be most likely: (a) to decline in a climate characterized by high levels of negative emotional behavior and low levels of positive emotional behavior; and (b) to increase in a climate characterized by low levels of negative emotional behavior and high levels of positive emotional behavior. For those with one or more long alleles, this association between emotional behavior and marital satisfaction should be much weaker.

What are we to do with information about how sensitive a marriage might be to the emotions expressed by spouses? We

are probably a long way from being able to use it to make important life decisions such as whom to marry and how much attention to pay to our partner's emotion (e.g., "marry someone with long alleles of 5-HTTLPR and you won't have to worry about how they feel"). Clearly, these kinds of genetic effects do not fully determine the fate of marriage, but rather represent small biases in a particular direction. Our findings (and those of others) that suggest that genetic influences on behavior become stronger with age suggest that such biases do aggregate and become more substantial over the decades of adult development. Nonetheless, many, many other factors affect the fate of an individual marriage and can override or counteract the biases associated with particular genetic influences. Still, these findings do raise the intriguing question of whether knowing you and your partner's 5-HTTLPR genotype would be useful information for relationship partners, much in the same way as knowing their temperament, attachment style, or level of neuroticism might be.

As conceptual models and experimental paradigms for studying common functional genetic polymorphisms continue to improve, we expect that the importance of this approach in emotion research will increase. Clearly, this research has great potential to help us understand individual differences in emotional functioning and the implications these differences have for the lives people live. The present study shows the important role that one particular genetic polymorphism plays in influencing the association between emotion and the quality of marriage, arguably one of the most important intimate relationships of adulthood. We expect that these kinds of influences will be found to extend to other kinds of intimate adult relationships as well. Given the critical role that social relationships play in physical health (e.g., Cacioppo & Patrick, 2008; House, Landis, & Umberson, 1988; Robles & Kiecolt-Glaser, 2003), mental health (e.g., Whisman, 2007), and general well-being (e.g., Coan, 2008; Proulx et al., 2007), there will be many opportunities to explore the way that our genes influence the complex relationships between emotions, intimate relationships, and health.

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