

Genetics of Borderline Personality Disorder



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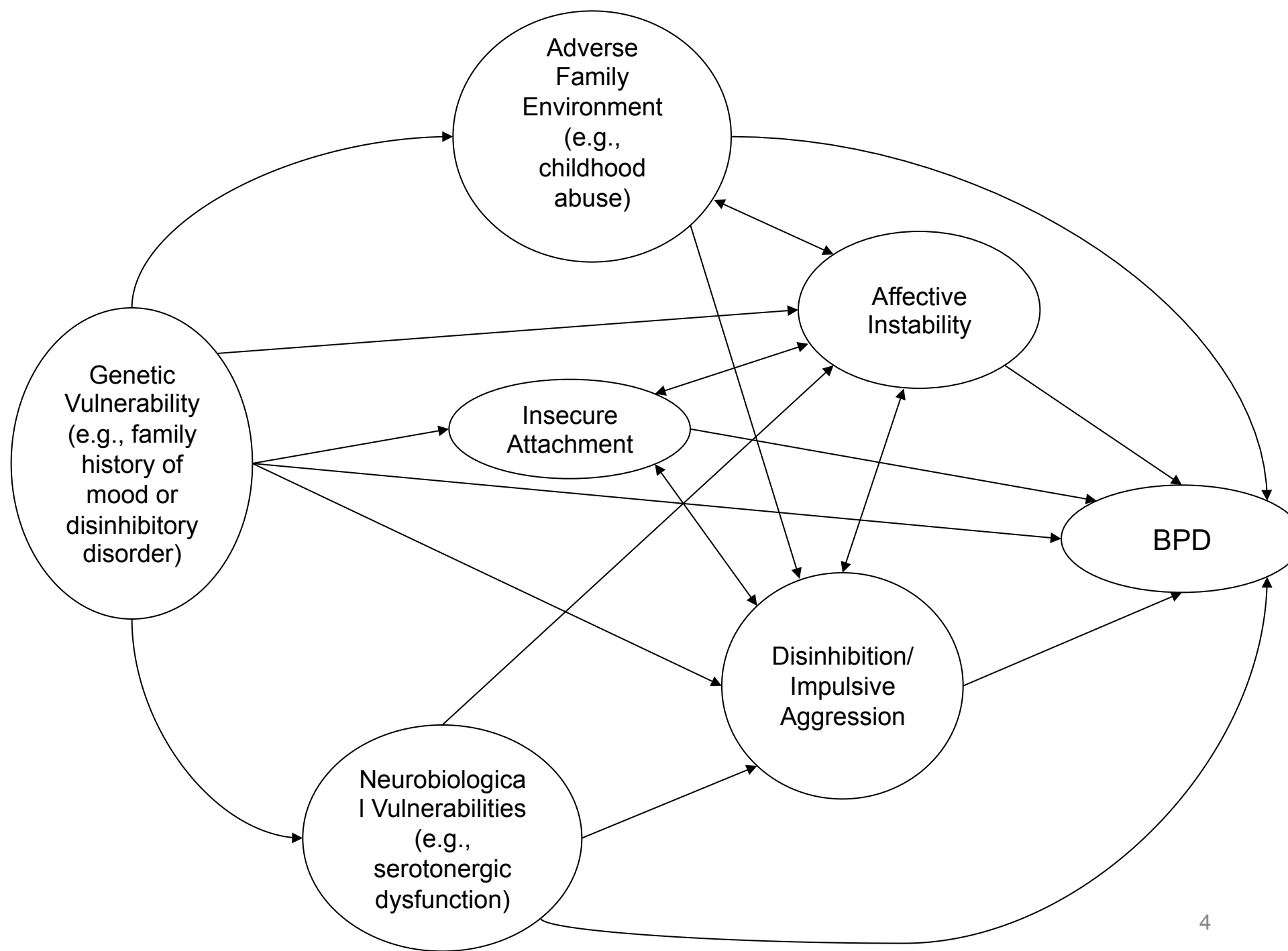
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DSM-IV Diagnostic Criteria for Borderline Personality Disorder (APA, 2000)

- Frantic efforts to avoid real or imagined abandonment
- Unstable and intense interpersonal relationships
- Persistently unstable self-image
- Impulsivity in at least two areas that are potentially self-damaging (e.g. sex and substance abuse)
- Suicidal behavior, gestures, or threats; Self-mutilating behavior
- Affective instability due to a marked reactivity of mood
- Feelings of emptiness
- Inappropriate and intense anger
- Stress-related dissociative symptoms

Borderline Personality Disorder Etiology

- A comprehensive model of the etiology of BPD includes the influences of genetics and family history, neurobiological factors, adverse environmental conditions, and personality factors.
- Figure 1 depicts a conceptualization of how these influences may interact and lead to the development of BPD symptoms
- Unidirectional and bidirectional effects.



Evidence for genetic effects

- Family studies
- Adoption studies
- Twin studies
- Candidate gene studies
- Genome wide association studies
- Gene-Environment studies
- What is inherited?

Borderline Personality Disorder Family Studies

- Family studies report increased rates of BPD in the relatives of individuals with BPD compared to relatives of control probands (e.g., Baron et al. 1985; Johnson et al. 1995; Zanarini et al. 2004; Bandelow et al. 2005; Zanarini et al. 1988; Loranger et al. 1982).
- Prevalences/morbidity risks for BPD in relatives of BPD probands ranged from 9.1% (Bandelow et al. 2005) to 24.9% (Zanarini et al. 1988).

Borderline Personality Disorder Family Studies (cont.)

- Prevalence of individual borderline symptoms or features in relatives of BPD probands.
- Silverman et al. (1991)
 - prevalence rates for **affective and impulsive personality disorder traits** were significantly higher in the relatives of BPD probands than in the relatives of probands with other personality disorders or in the relatives of schizophrenic probands.
- Zanarini et al. (2004)
 - in first degree relatives of BPD patients, the prevalence rates of five **(inappropriate anger, affective instability, paranoia/dissociation, general impulsivity, and intense, unstable relationships)** were significantly higher in first degree relatives of BPD patients than in first degree relatives of axis-II comparison subjects.

Borderline Personality Disorder Family Studies (cont.)

- Limitations
 - Sample sizes small, varying from 17 (Baron et al. 1985) to 83 BPD probands (Loranger et al. 1982)
 - Often not representative of the population (e.g. Loranger et al. 1982 assessed only female BPD probands).
 - Information on psychopathology of relatives was derived from the BPD probands themselves (e.g., Zanarini et al., 2004).
 - Family studies CANNOT RULE OUT ENVIRONMENTAL INFLUENCES

Borderline Personality Disorder Adoption Studies

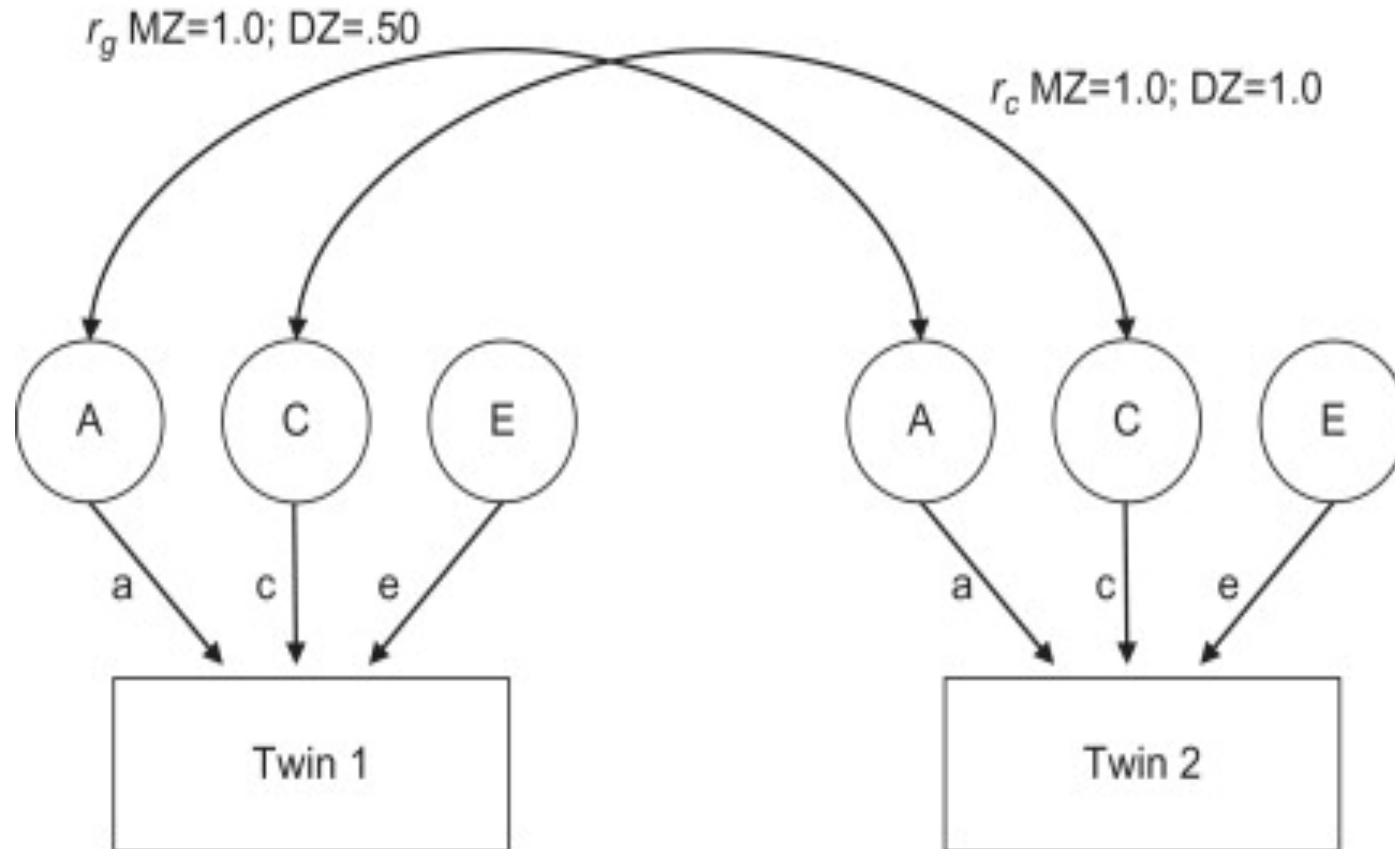
- Very compelling design
- Genetic influences from biological parents; no environmental influence from biological parents (not raised in this home)
- Environmental influences from foster home (and unique environment).
- Foster siblings outcome can be used in comparison; same shared environment.
- Unfortunately, no adoption studies of BPD

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Twin Studies

- Family studies cannot disentangle the effects of genes from the effects of environment shared by family members, social interaction and cultural inheritance.
- Twin studies can disentangle the effects of common environment and genes by making use of the different genetic relatedness of monozygotic (MZ) and dizygotic (DZ) twins.
- MZ twins are genetically (nearly) identical while DZ twins and siblings share on average 50% of their segregating genes.
- If genetic factors are important for a trait, MZ twins must be more similar than DZ twins or other first degree relatives.
- If MZ twins are as similar as DZ twins, familiarity is mainly due to common environmental factors.

Univariate genetic model



Borderline Personality Disorder Twin Studies (cont.)

- Few twin studies on BPD available using structured interviews.
- Torgersen et al. (2000) found a concordance rate for definite BPD at 35% for MZ pairs and 7% for DZ pairs.
 - $h^2=.69$
 - A number of methodological problems with this study (including the sampling of those who were being treated for a mental disorder, the small number of twin pairs, and the interviewers' awareness of both zygosity and diagnostic status of the co-twin).
- Torgersen et al, (2008): approx 1400 twin pairs
 - h^2 estimate for BPD=.35

Borderline Personality Disorder Twin Studies (cont.)

- Twin studies using self-report questionnaire measures of BPD
- Distel et al. (2008) were able to assess BPD features in 5,496 twins (1,852 complete pairs) between the ages of 18 and 86 years from the Netherlands, Belgium and Australia.
 - Results showed that genetic influences explained 42% of the variation in BPD features in both men and women.
 - The heritability was equal between the three countries suggesting no interaction between genotype and country.

Borderline Personality Disorder Twin studies

Distel et al. (2008), Psychological Medicine

	A	E
Netherlands	42.3%	57.7%
Belgium	42.5%	57.5%
Australia	41.6%	57.8%

Estimates constrained to be equal

A=42.2%; E=57.8%

Borderline Personality Disorder Twin Studies (cont.)

- Twin studies using self-report questionnaire measures of BPD
- Torgersen et al. (in press)
- ~2,800 Norwegian Twins; used items from the Dysfunction Personality Questionnaire that tapped BPD criteria.
- Heritability of questionnaire scores estimated to be .46

Borderline Personality Disorder Twin Studies (cont.)

- Limitations:
- Aggregate effect of genes only
- Cannot identify specific genetic influences
- Equal shared environments for MZ versus DZ twins assumed

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Candidate gene studies

- Most candidate genes are functional genes that have biological consequences related to the trait, disorder or disease.
- Reduced serotonergic function in anger (Giegling et al. 2006), aggression (Siever 2008), suicidal behaviour (Bah et al. 2008; Zaboli et al. 2006) and impulsivity (Passamonti et al. 2008; New et al. 1998), and increased serotonergic function in emotional lability (Hoefgen et al. 2005) have led to several **serotonergic candidate genes for BPD**.
- Tryptophan hydroxylase (TPH) and the serotonin transporter gene (5-HTT), are the most studied candidate genes.

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Candidate gene studies

- There is some evidence that **dopamine (DA) dysfunction** may be associated with BPD. DA dysfunction is associated with emotional dysregulation, impulsivity and cognitive-perceptual impairment (for a review see Friedel 2004), three important dimensions of BPD.
- Joyce et al. (2006) found a significant replicated association between the 9-repeat allele of dopamine transporter 1 (dopamine active transporter, DAT1) and BPD in depressed patients.

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Candidate gene studies

- Genes involved in the production of **monoamine oxidase-A (MAOA)**, which degrades amongst others 5-HTT and DA, are suggested to be involved in BPD because it is shown to be associated with aggression (Buckholtz and Meyer-Lindenberg 2008), impulsivity (Manuck et al. 2000) and mood lability (Furlong et al. 1999).
- To test whether MAOA is also associated with the BPD diagnosis Ni et al. (2007) genotyped two MAOA polymorphisms (promoter VNTR and rs6323) in 111 BPD patients and 289 control subjects.
- A high frequency of the high activity VNTR alleles and a low frequency of the low activity haplotype was found in BPD patients suggesting that the high activity allelic variant may play a role in the etiological development of BPD.

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Candidate gene studies

- **Some Limitations inherent in the candidate gene approach:**
- Must have a good understanding of the function of a candidate gene
- Most traits/features are influenced by many genes, each contributing only a very small amount to overall genetic risk
- Therefore, need extremely large samples to detect these very small effects, and have to target the “correct” genes
- Gene variants that are relevant will be missed if they are not within the specific gene region of investigation

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Genome wide association studies

- Genome wide association (GWA) studies. GWA searches the whole genome for small variations (single-nucleotide polymorphisms; SNPs) that occur more frequently in people with a particular disorder than in people without the disorder.
- Each study can look at thousands of SNPs at the same time.
- However, association analysis measures statistical associations, cannot be used to test for causality, and is prone to population stratification. Cases and controls should ideally come from the same population.

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Genome wide association studies

- Large samples are required, and most findings for other disorders have not been replicated.
- Only a small percentage of the variance in the phenotype explained (<5%); only **common** genetic variants detected

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Gene-Environment Studies

- Three major ways that genes and environment can jointly influence the overall vulnerability to psychopathology.
 - First, an individual's liability for a disorder may be the sum of the contributions of genes and environment (**the additive model**).
 - Second, genes and environment may interact, such that genes control the sensitivity to the environment or, alternatively, one can say the environment controls gene expression (**the gene-environment interaction model; GxE**).
 - Third, genes and environment can be correlated such that genes influence environmental exposure (**the gene-environment correlation model; rGE**).

Borderline Personality Disorder Gene-Environment Studies

- Gene-environment interaction (GxE) implies that genes determine the degree to which a subject is sensitive to an environment. In the presence of interaction, individuals with a 'sensitive' genotype will be of greater risk to develop BPD if the predisposing environment is present, than individuals with an 'insensitive' genotype.
- GxE can be detected by determining if the heritability of BPD varies in groups with different environmental conditions (for example, experiencing sexual abuse).

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Gene-Environment Interaction

- Many studies purporting to find GxE effects on a phenotype are unable to do so definitely for one or more of the following reasons:
 - (1) the number of participants is much too small to reliably detect GxE effects (effects which are notoriously small);
 - (2) factors that are considered “environmental” may actually be under some degree of genetic influence (i.e., a gene-environment correlation);
 - (3) “environments” are measured retrospectively and imprecisely; and
 - (4) the inadequate scaling of environments (i.e., how these are quantified) can lead one to conclude that a GxE effect exists when in fact it does not.

Borderline Personality Disorder Gene-Environment Interaction

- There is no strong, replicated empirical evidence yet supporting a gene-environment interaction (GxE) model of liability to borderline pathology. Regardless of whether....
 - Studies have examined genetic underpinnings of features of BPD (impulsivity, emotional sensitivity, anger and aggression, suicidal behavior)
 - Studies have examined the BPD phenotype itself (via questionnaire or interview)

Borderline Personality Disorder Gene-Environment Correlation

- Distel et al. (2011) tested the GxE and rGE models in a large twin/sibling sample.
- rGE, the correlation between the genetic influence on environmental risk factors and the genetic influence on BPD traits
- Twin and sibling pairs from the Netherlands and Belgium (n=6368).
 - Participants self-reported their symptoms of BPD, using the Personality Assessment Inventory-Borderline Scale (PAI-BOR), and
 - the experience of SLEs (i.e., divorce, car accident, assault, robbery, and job loss).

Borderline Personality Disorder Gene-Environment Correlation (cont.)

- Distel et al. (2011)
- Evidence suggested significant **gene-environment correlation** for divorce/break-up, violent assault, sexual assault, and job loss and borderline personality.
- That is, *the genes influencing borderline features increased the likelihood of being exposed to these adverse life events.*
- However, it is not possible to determine the direction of causality.

What is inherited or under genetic control?

- Neurobiological Vulnerabilities (serotonin, dopamine, endogenous opioid system)
- Personality traits: impulsivity/aggression and emotional dysregulation
- Adverse life events
- Attachment styles?

Borderline Personality Disorder Personality

Distel et al. (2009)

- All genetic variance in borderline personality is shared with FFM personality traits (esp. with N and A)
- 33% of environmental effect is NOT shared with FFM traits
- Unique environmental influence may influence normal personality → borderline personality

The future of genetics research for BPD

- Well-designed, large twin and family studies that assess the aggregate effects of genes.
- In order to reliably characterize a G-E effects, it is necessary to use very large samples (thousands).
- It is necessary to also evaluate gene-environment correlation effects, and a failure to do so could lead to erroneous conclusions about a GxE effect

The future of genetics research for BPD (cont)

- Great need for improved measures of environmental influences (prospective, cumulative, and proximal)
- The genes that have been considered include a relatively small range of candidate genes, whose status as risk factors for BPD is uncertain.
- There are no large scale genome wide association (GWA) studies yet for BPD.

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Final Comments

- The expression of BPD features is genetically mediated, although genetic factors cannot explain the entire picture.
- These issues can only be addressed with extremely large population-based samples and clinical case-controls.
- Research on the genetics of BPD is a very important next step
- Identifying genes that influence the development of BPD will help to develop better strategies to diagnose, treat and prevent the disorder.
- Our measures of environmental influences are woefully inadequate.

Thank you!



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Further reading

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