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## High-resolution chromosome ideogram representation of recognized genes for bipolar disorder

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### Abstract

Bipolar disorder (BPD) is genetically heterogeneous with a growing list of BPD associated genes reported in recent years resulting from increased genetic testing using advanced genetic technology, expanded genomic databases, and better awareness of the disorder. We compiled a master list of recognized susceptibility and genes associated with BPD identified from peer-reviewed medical literature sources using PubMed and by searching online databases, such as OMIM. Searched keywords were related to bipolar disorder and genetics. Our compiled list consisted of 290 genes with gene names arranged in alphabetical order in tabular form with source documents and their chromosome location and gene symbols plotted on high-resolution human chromosome ideograms. The identified genes impacted a broad range of biological pathways and processes including cellular signaling pathways particularly cAMP and calcium (e.g., *CACNA1C*, *CAMK2A*, *CAMK2D*, *ADCY1*, *ADCY2*); glutamatergic (e.g., *GRIK1*, *GRM3*, *GRM7*), dopaminergic (e.g., *DRD2*, *DRD4*, *COMT*, *MAOA*) and serotonergic (e.g., *HTR1A*, *HTR2A*, *HTR3B*) neurotransmission; molecular transporters (e.g., *SLC39A3*, *SLC6A3*, *SLC8A1*); and neuronal growth (e.g., *BDNF*, *IGFBP1*, *NRG1*, *NRG3*). The increasing prevalence of BPD calls for better understanding of the genetic etiology of this disorder and associations between the observed BPD phenotype and genes. Visual representation of genes for bipolar disorder becomes a tool enabling clinical and laboratory geneticists, genetic counselors, and other health care providers and researchers easy access to the location and distribution of currently recognized BPD associated genes. Our study may also help inform diagnosis and advance treatment developments for those affected with this disorder and improve genetic counseling for families.

### Keywords

Bipolar disorder; Chromosome ideograms; Major affective disorder; Manic-depressive illness; Susceptibility genes

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Conflicts of interest

The authors declare no conflict of interest.

## 1. Introduction

Bipolar disorder (BPD) or major affective disorder is largely undiagnosed but known to cause unusual shifts in behavior, including mood, activity levels, and ability to perform everyday tasks. According to the Diagnostic and Statistical Manual for Mental Disorders, fifth edition (DSM-5), BPD can be separated into four basic types: Bipolar I Disorder (BP-I), Bipolar II Disorder (BP-II), Bipolar Disorder Not Otherwise Specified (BP-NOS), and Cyclothymic Disorder. The diagnosis is dependent on severity and length of the manic and depressive phases [Hirschfeld et al., 2000, 2003a, 2003b; Juvenile Bipolar Research Foundation (<http://www.jbrf.org/diagnosis-by-the-dsm/>); National Institute of Mental Health ([http://www.nimh.nih.gov/health/topics/bipolar-disorder/index.shtml#part\\_145403](http://www.nimh.nih.gov/health/topics/bipolar-disorder/index.shtml#part_145403))]. The twelve month prevalence of BPD is 1.5% for the adult population in the U.S., 0.4% for BP-I, 0.3% for BP-II, and 0.8% for BP-NOS. The lifetime prevalence of BPD in the U.S. population is 2.4% for the total population; 0.6% for BP-I, 0.4% for BP-II, and 1.4% for BP-NOS (Merikangas et al., 2011). Men and women are equally affected by BPD, but women are three times more likely than men to experience the rapid cycling process and tend to experience the depressive and mixed episodes more often than men (Merikangas et al., 2011). Most individuals are diagnosed with bipolar disorder around the age of 25 years, but some individuals are diagnosed as young as six years of age or as late as 40 and 50 years [Birmaher et al., 2006; Geller et al., 2004; Depression and Bipolar Support Alliance Depression and Bipolar Support Alliance ([http://www.dbsalliance.org/site/PageServer?pagename=education\\_statistics\\_bipolar\\_disorder](http://www.dbsalliance.org/site/PageServer?pagename=education_statistics_bipolar_disorder))]. Given the nature of the disorder, it is common for individuals with BPD to be misdiagnosed as having another mental illness, which can hinder recognition and treatment [Depression and Bipolar Support Alliance ([http://www.dbsalliance.org/site/PageServer?pagename=education\\_statistics\\_bipolar\\_disorder](http://www.dbsalliance.org/site/PageServer?pagename=education_statistics_bipolar_disorder))]. BPD often coexists with other Axis I and Axis II disorders [e.g. substance abuse, anxiety disorders (such as agoraphobia, post-traumatic stress disorder, and social phobia), and eating disorders] with reported rates of lifetime psychiatric comorbidity in bipolar I samples ranging from 50% to 70% (McElroy et al., 2001). Approximately 20% of the children and adolescents experiencing major depression will develop bipolar disorder within five years of being diagnosed with depression (Birmaher et al., 2006; McClellan et al., 2007). Almost one-third of adolescents with depression are actually experiencing early onset of bipolar disorder.

Numerous twin and family studies have supported a genetic contribution to the risk of developing BPD. Monozygotic twins show greater concordance with psychopathology than dizygotic twins and individuals with affected relatives show an increased chance of developing bipolar and/or unipolar depression (Smoller and Finn, 2003). One affected parent is associated with a 15 to 25% elevation in risk of BPD while two affected parents shows a 50 to 75% increase [Smoller and Finn, 2003; Depression and Bipolar Support Alliance ([http://www.dbsalliance.org/site/PageServer?pagename=education\\_statistics\\_bipolar\\_disorder](http://www.dbsalliance.org/site/PageServer?pagename=education_statistics_bipolar_disorder))]. Never-the-less, bipolar disorder is a complex genetic disorder that does not follow Mendelian inheritance patterns with no identifiable “gene of major effect” found for the majority of BPD cases. There are also several chromosomal regions, susceptibility loci and implicated genes for BPD that have

been repeatedly reported as associated with the disorder through linkage, candidate and genome-wide association studies (Craddock and Jones, 1999; Craddock et al., 2005). Interactions between numerous genes in multiple overlapping pathways and functions reduce the likelihood of one-to-one correspondence for any single “causal” bipolar disorder gene mutation (Potash and DePaulo, 2000).

Over the last few decades, researchers have identified numerous candidate genes associated with BPD using various methodologies, including genotyping single-nucleotide polymorphisms (SNPs), identifying cytogenetic abnormalities (i.e., chromosomal breakpoints which lead to the loss or alteration of specific genes), or convergent functional genomics (Cichon et al., 2009). We used existing literature and genomic databases to obtain evidence to compile a master list of currently recognized genes with their locations plotted on high-resolution chromosome ideograms (850 band level) associated with BPD. In tabular form, we listed the individual gene symbol, expanded name, and chromosome location alongside the reference source providing the information.

## 2. Material and methods

We used computer-based internet websites and PubMed (<https://www.ncbi.nlm.nih.gov/pubmed>) to search for key words based on the genetics of bipolar disorder. Articles were obtained by searching PubMed and Online Mendelian Inheritance in Man (OMIM: <http://www.ncbi.nlm.nih.gov/omim>) databases with the following search words: *bipolar disorder (BPD)*, *bipolar*, *major affective disorder*, *bipolar affective disorder (BPAD)*, *bipolar syndrome*, and *manic depressive psychosis*. We examined literature found in medical journals after our search for genetic involvement for BPD. Some articles we found had their own compilation of susceptibility genes for BPD with references to their own sources which we then further studied (Thomson et al., 2005; Le-Niculescu et al., 2009; Cichon et al., 2009; Shinozaki and Potash, 2014). The articles were then prioritized based on the following considerations: *sample size*, *use of standardized diagnostic criteria*, *types of genetic testing such as genome wide association studies and validated methods (e.g., convergent functional genomics, genotyping SNPs or identifying cytogenetic abnormalities)*, and *quality with reliability of the genetic data and presentation*. GeneCards (<http://www.genecards.org/>) was the primary source for determining the location of the gene or gene locus. The cytogenetic location of the gene was provided by Ensembl, Entrez Gene, or the Human Genome Organization Gene Nomenclature Committee (HGNC).

BPD is a heterogeneous disorder involving many genes acting individually or in combination and responsive to environmental stimuli. We compiled a list of genes from the major sources and their references for a total of 290 genes. Our paper focused on genes associated with BPD by at least one mechanism of proven association with or susceptibility to BPD. Research articles were not limited to causal relationships to BPD. For example, Le-Niculescu and others used convergence of microsatellite markers for which at least one published study showed evidence for linkage for BPD, or a positive association study for the gene itself reported in previous literature (Le-Niculescu et al., 2009). Many of the genes on our list were found in multiple research studies and were reported more than once for being associated with BPD. Some studies considered target neurochemical pathways rather than

susceptibility loci. These genes were included on our list if they were also recognized by a peer-reviewed publication (e.g., PubMed) with supporting evidence (e.g., GWAS, informative SNPs, genetic linkage, or identified gene mutations) as a BPD associated gene (Le-Niculescu et al., 2009; Cichon et al., 2011). Other supporting genetic evidence can be found in the National Center for Biotechnology Resources at <https://www.ncbi.nlm.nih.gov/gene>.

### 3. Results

By using online databases and existing literature in peer-reviewed journals, we were able to compile a master list of 290 recognized genes associated with BPD. These genes were clinically relevant, or related to susceptibility of BPD. The position for each of the recognized or susceptibility genes for BPD were plotted on the high-resolution chromosome ideograms (850 band level), as shown in Fig. 1. We have also included gene symbols, expanded names, chromosome band location and reference sources in Table 1 for the 290 genes recognized as playing a role in BPD. The distribution of BPD genes is shown in Table 2 among individual chromosomes and chromosome arms arranged by the size for each chromosome (largest chromosome represented by the smallest number) in relationship to the proportion of total genes for BPD.

The majority of the genes listed in Table 1 are located on chromosomes 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 13, 16, 18, and 22 in accordance with previous reports examining polymorphisms and chromosomal breakpoints of known and candidate genes for BPD (Potash and DePaulo, 2000; Etain et al., 2012). One of the smallest chromosomes (i.e., 22) contained more BPD genes in relationship to its size (i.e., 4.1%) than other comparable chromosomes. The X chromosome which accounts for about 5.6% of the genome contained only 1.0% of the BPD genes. The genes identified in our investigation impacted a broad range of biological pathways and processes including cellular signaling pathways for cAMP and calcium (e.g., *CACNA1C*, *CAMK2A*, *CAMK2D*, *ADCY1*, *ADCY2*); glutamatergic (e.g., *GRIK1*, *GRM3*, *GRM7*), dopaminergic (e.g., *DRD2*, *DRD4*, *COMT*, *MAOA*) and serotonergic (e.g., *HTR1A*, *HTR2A*, *HTR3B*) neurotransmission; molecular transporters (e.g., *SLC39A3*, *SLC6A3*, *SLC8A1*); and neuronal growth (e.g., *BDNF*, *IGFBP1*, *NRG1*, *NRG3*). These gene classes influence neurotransmission and psychological functioning through direct and indirect effects on neuronal activity, growth, development, maintenance and remodeling. These genes overlap with biological markers and genetic factors associated with other psychiatric disorders such as schizophrenia, (e.g., *BDNF*, *CACNA1*, *DISC1*), major depressive and anxiety disorders (e.g., *TPH2*, *HTR2A*) involving developmental processes (e.g., *NRG1*, *NRG3*).

### 4. Discussion

The advent of genetic testing to identify the predisposition for BPD has increased understanding of the genetic etiology of BPD and the application of genetic testing should facilitate research and treatment development. We illustrated a master list of susceptibility genes and genes that are associated with BPD in our results by plotting the individual genes on high-resolution chromosome ideograms and generated a tabular form with references in

order to inform more individuals about the necessity for genetic testing and treatment for BPD.

The transmission of BPD does not appear to manifest through a simple Mendelian inheritance with a single allele (dominant) or two alleles (recessive) pattern instead showing incomplete penetrance, etiological heterogeneity and high prevalence (Kerner, 2015). Similarly, multivariate threshold models conceptualized as an accumulation of traits normally distributed throughout the genome are not strongly supported to achieve some threshold for expression. Rather, BPD appears to be a product of a combination of genetic, neurochemical, and environmental influences. The expression or development of bipolar disorder may be due to epistasis (interaction of multiple genes) or other complex mechanisms such as genomic imprinting or dynamic mutations with environmental contributions (Craddock and Jones, 1999). An intricate oligogenic quasi-Mendelian pattern has been proposed whereby a small number of mutations accumulate in a select biological pathway that is only tied to expression of the phenotype when released by environmental influences (Kerner, 2015). Symptoms of bipolar disorder parallel other genetically influenced psychiatric disorders including schizophrenia, depression and anxiety (Krishnan, 2005) and is, not surprisingly, impacted by overlapping genetic constructs (e.g., *BDNF*, *DISC1*, *HTR2A*, *TPH2*).

Our list of genes for BPD reflects the current status of recognized genes with clinical relevance but susceptibility and new genes are continually being identified. Not all genes in the list are equally significant or certainly causative for all individuals with BPD. Similarly, our results reflect gene-level associations and do not provide evidence of individual SNP- or CNV-level contributions to pathology. The list is particularly suited to the evaluation of structural genomic data (e.g., DNA microarrays) for copy number variations impacting genomic regions and genes of interest which may involve large regions and multiple candidate genes or encompass known genetic syndromes. Phenotype and severity can be predicted to be proportional to the number of candidate genes impacted by the CNV – highlighting the clinical value of spatial representation of the gene list and whether the region is duplicated or deleted. The effect of any individual SNP depends upon both the physiological relevance of the gene to neurodevelopmental processes and the impact of the specific sequence variation or mutation on the expression and function of the gene product (i.e., synonymous vs. non-synonymous). Non-synonymous variations leading to a change of the codon or reading frame (e.g., frameshift mutation, stop codon generation) have greater predicted impact on gene expression and thus stronger ties to pathology. Advances in genomic technology will facilitate the identification and characterization of novel SNPs among the gene candidates and improve understanding of the relative contributions of selected SNP or CNVs to the general disease prevalence.

Combinations of single nucleotide polymorphisms (SNPs) which alter the genetic code, creating a higher likelihood of developing bipolar disorder, are reportedly more common than chromosomal breakpoints at the susceptibility regions for BPD. For example, SNPs within the brain derived neurotrophic factor (*BDNF*) gene, located in the 11p14, region are reported to result in a modification of the processing and trafficking of the *BDNF* gene and increased susceptibility to BPD (Craddock et al., 2005). The *ABCA13* gene was also

identified as a candidate gene for both bipolar and schizophrenia disorders once it was initially discovered by a chromosome abnormality in a schizophrenic patient (Knight et al., 2009). The *ABCA13* gene was then resequenced and multiple rare coding variants identified. These variants were genotyped in bipolar cohorts, which led to the conclusion that 4.0% of the population contained a variant which could contribute to bipolar disorder (Knight et al., 2009). Multiple genome-wide association studies (GWAS) testing for SNPs in independent bipolar cases and controls often generate strong signals for gene associations with bipolar disorder (Sklar et al., 2011). For example, results from the first GWAS for BPD reported in 2007 determined that the strongest signals for an association were found in five genes: *BDNF* at 11p14, *DAOA* at 13q33, *DTNBP1* at 6p22, *DISC1* at 1q42, and *NRG1* at 8p12, and the *SNPs*rs420259 at 16p12 (Burton et al., 2007). Since this initial study, additional GWAS reports (e.g., Sklar et al., 2008; Ferreira et al., 2008) and a collection of genetic studies examining bipolar disorder did implicate genes that were consolidated in 2014 by Shinozaki and Potash (Shinozaki and Potash, 2014) with a list of candidate genes summarized from 15 studies (e.g., Baum et al., 2008a, 2008b; Sklar et al., 2011; Ou et al., 2015). These genes overlapped with biological markers and genetic factors associated with other psychiatric disorders such as schizophrenia, (e.g., *BDNF*, *CACNA1*, *DISC1*), major depressive and anxiety disorders (e.g., *TPH2*, *HTR2A*) and involved in developmental processes.

Our list of candidate genes and visual representation of their locations on high-resolution chromosome ideograms for bipolar disorder provide an informative perspective of the pathophysiology of BPD to facilitate research, accurate diagnosis and genetic counseling, treatment development and classification of BPD (Craddock and Jones, 1999). The authors encourage the use of this collection of currently associated recognized susceptibility genes for BPD in the evaluation of patients presenting for genetic services and for a more accurate understanding of the role of genetics in BPD, a genetically heterogeneous disorder.

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## Abbreviations:

<b>BPD</b>	bipolar disorder
<b>BP-I</b>	Bipolar I Disorder
<b>BP-II</b>	Bipolar II Disorder
<b>BP-NOS</b>	Bipolar Disorder Not Otherwise Specified
<b>cAMP</b>	cyclic adenosine monophosphate



<b>DNA</b>	deoxyribonucleic acid
<b>DSM-5</b>	Diagnostic and Statistical Manual for Mental Disorders, fifth edition
<b>FDA</b>	Food and Drug Administration
<b>GWAS</b>	genome-wide association studies
<b>OMIM</b>	online Mendelian inheritance in man
<b>SNPs</b>	single nucleotide polymorphisms

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## Web Resources

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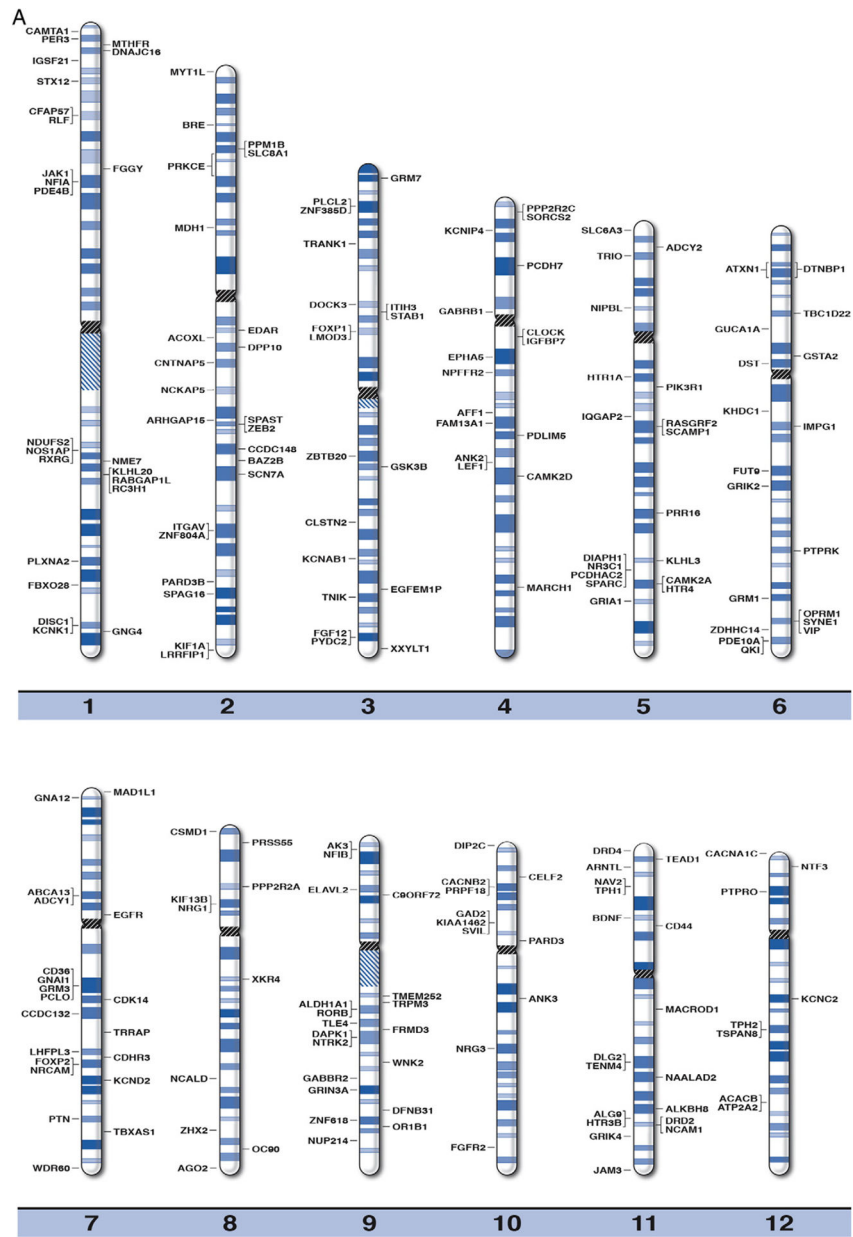
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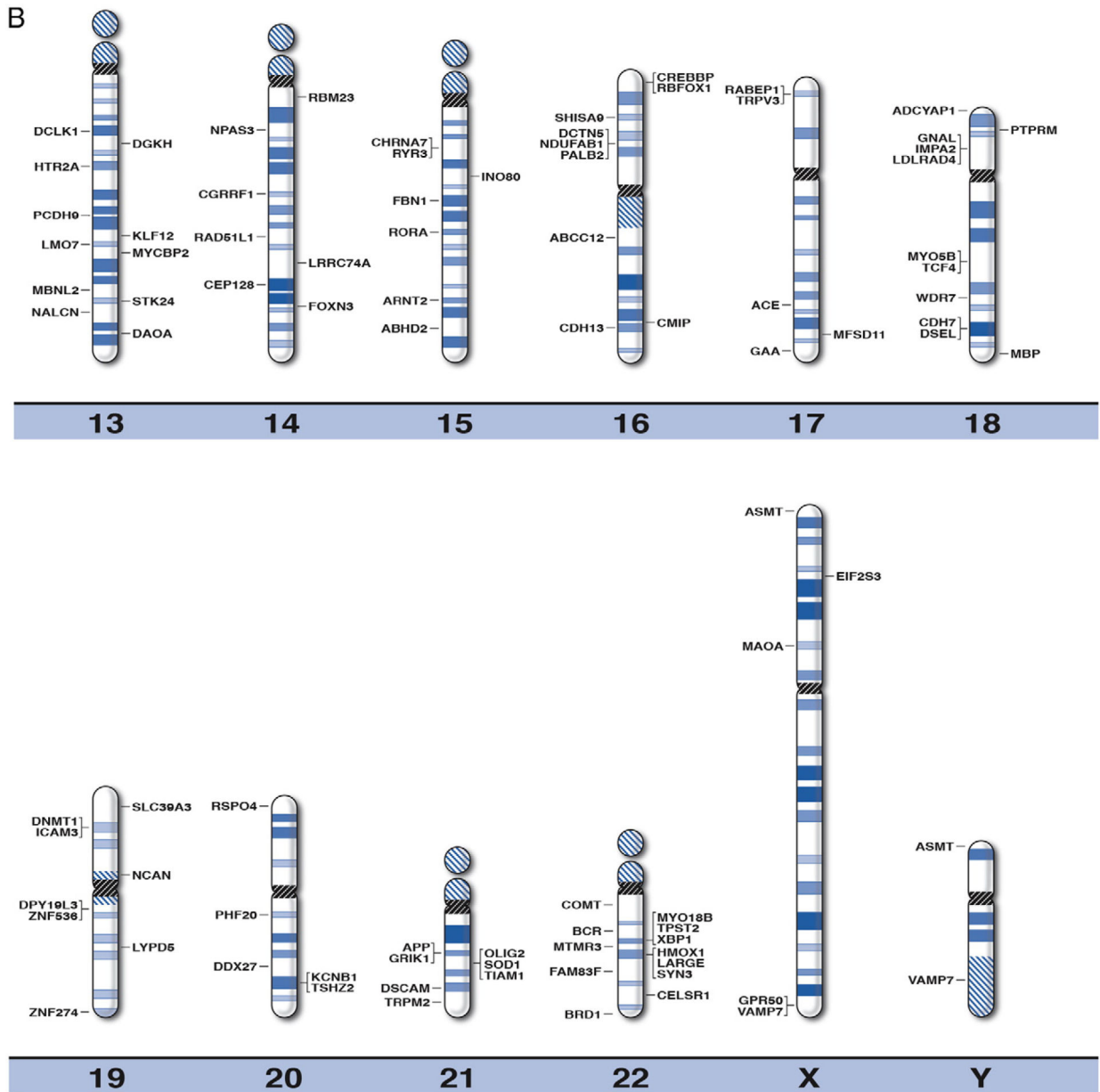
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**Fig. 1.** High-resolution chromosome ideograms (850 band level) with the BPD gene symbol placed at the chromosomal band location. The centromere area, highlighted in black, separates the upper short “p” arm and the lower long “q” arm for each chromosome. The gene symbols are arranged in alphabetical order with the expanded name and chromosome band position listed in Table 1.

**Table 1**  
Known and candidate genes for bipolar disorder (BPD) and overlap with schizophrenia and autism.

Gene symbol	Gene name	Location	Source
<i>ABCA13</i> *	ATP-binding cassette, subfamily A, member 13	7p12.3F	Knighet et al. (2009)
<i>ABCC12</i>	ATP-binding cassette, subfamily C, member 12	16q12.1	Xu et al. (2014)
<i>ABHD2</i>	Abhydrolase domain containing 2	15q26.1	Xu et al. (2014)
<i>ACACB</i>	Acetyl-coenzyme A carboxylase beta	12q24.11	Le-Niculescu et al. (2009)
<i>ACE</i> **	Angiotensin I converting enzyme	17q23.3	Zou et al. (2011)
<i>ACOXL</i>	Acyl-CoA oxidase-like	2q13	Xu et al. (2014)
<i>ADCY1</i> *	Adenylate cyclase 1 (brain)	7p12.3	Le-Niculescu et al. (2009)
<i>ADCY2</i>	Adenylate cyclase 2	5p15.31	Mühlaisen et al. (2014)
<i>ADCYAP1</i> *	Adenylate cyclase activating polypeptide 1 (pituitary)	18p11.32	Le-Niculescu et al. (2009)
<i>AFF1</i>	AF4/FMR2 family, member 1	4q21.3	Xu et al. (2014)
<i>AGO2</i>	Argonaute RISC catalytic component 2	8q24.3	Le-Niculescu et al. (2009)
<i>AK3</i>	Adenylate kinase 3	9p24.1	Le-Niculescu et al. (2009)
<i>ALDH1A1</i>	Aldehyde dehydrogenase 1 family, member A1	9q21.13	Le-Niculescu et al. (2009)
<i>ALG9</i>	ALG9, alpha-1,2-mannosyltransferase	11q23.1	Baysal et al. (2002)
<i>ALKBH8</i>	ALKB, alkylation repair homolog 8 ( <i>Escherichia coli</i> )	11q22.3	Xu et al. (2014)
<i>ANK2</i> #	Ankyrin 2, neuronal	4q25	Le-Niculescu et al. (2009)
<i>ANK3</i> **	Ankyrin 3	10q21.2	Takata et al. (2011)
<i>APP</i> #	Amyloid beta (A4) precursor protein	21q21.3	Le-Niculescu et al. (2009)
<i>ARHGAP15</i> #	Rho GTPase activating protein 15	2q22.2	Xu et al. (2014)
<i>ARNT2</i> #	Aryl-hydrocarbon receptor nuclear translocator 2	15q25.1	Xu et al. (2014)
<i>ARNTL</i> *	Aryl hydrocarbon receptor nuclear translocator-like	11p15.3	Le-Niculescu et al. (2009)
<i>ASMT</i> #	Acetylserotonin O-methyltransferase	Xp22.33 or Yp11.32	Etain et al. (2012)
<i>ATP2A2</i>	ATPase, Ca++ transporting, cardiac muscle, slow twitch 2	12q24.11	Xu et al. (2014)
<i>ATXN1</i> *	Ataxin 1	6p22.3	Le-Niculescu et al. (2009)
<i>BAZ2B</i>	Bromodomain adjacent to zinc finger domain, 2B	2q24.2	Xu et al. (2014)
<i>BCR</i>	Breakpoint cluster region	22q11.23	Hashimoto et al. (2005)

Gene symbol	Gene name	Location	Source
<i>BDNF</i> <sup>**</sup>	Brain-derived neurotrophic factor	11p14.1	Fan and Sklar (2008)
<i>BRDJ</i> <sup>*</sup>	Bromodomain containing 1/zinc finger, BED-type containing 4	22q13.33	Nyegaard et al. (2010)
<i>BRE</i>	Brain and reproductive organ-expressed (TNFRSF1Amodulator)	2p23.2	Cichon et al. (2011)
<i>C9ORF72</i>	Chromosome 9 open reading frame 72	9p21.2	Meisler et al. (2013)
<i>CACNA1C</i> <sup>**</sup>	Calcium channel, voltage-dependent, L type, alpha 1C subunit	12p13.33	Ou et al. (2015)
<i>CACNB2</i> <sup>**</sup>	Calcium channel, voltage dependent, beta 2 subunit	10p12.33	Le-Niculescu et al. (2009)
<i>CAMK2A</i>	Calcium/calmodulin-dependent protein kinase II, alpha	5q32	Le-Niculescu et al. (2009)
<i>CAMK2D</i>	Calcium/calmodulin-dependent protein kinase II, delta	4q26	Le-Niculescu et al. (2009)
<i>CAMTA1</i> <sup>#</sup>	Calmodulin binding transcription activator 1	1p36.31	Cichon et al. (2011)
<i>CCDC132</i>	Coiled-coil domain containing 132	7q21.3	Cichon et al. (2011)
<i>CCDC148</i>	Coiled-coil domain containing 148	2q24.1	Xu et al. (2014)
<i>CD36</i>	CD36 molecule (Thrombospondin receptor)	7q21.11	Blair et al. (2006)
<i>CD44</i> <sup>#</sup>	CD44 molecule (Indian blood group)	11p13	Le-Niculescu et al. (2009)
<i>CDH7</i>	Cadherin 7, type 2	18q22.1	Soronen et al. (2010)
<i>CDH13</i>	Cadherin 13	16q23.3	Le-Niculescu et al. (2009)
<i>CDHR3</i>	Cadherin-related family member 3	7q22.3	Xu et al. (2014)
<i>CDK14</i>	Cyclin-dependent kinase 14	7q21.13	Le-Niculescu et al. (2009)
<i>CELF2</i>	CUGBP, elav-like family member 2	10p13	Le-Niculescu et al. (2009)
<i>CELSR1</i>	Cadherin, EGF LAG seven-pass G-type receptor 1	22q13.31	Le-Niculescu et al. (2009)
<i>CEP128</i>	Centrosomal protein 128 kDa	14q31.1	Le-Niculescu et al. (2009)
<i>CFAP57</i>	Cilia and flagella associated protein 57	1p342	Xu et al. (2014)
<i>CGRRF1</i>	Cell growth regulator with ring finger domain 1	14q22.2	Xu et al. (2014)
<i>CHRNA7</i> <sup>**</sup>	Cholinergic receptor, nicotinic, alpha 7 (neuronal)	15q13.3	Le-Niculescu et al. (2009)
<i>CLOCK</i> <sup>*</sup>	Clock circadian regulator	4q12	Benedetti et al. (2003)
<i>CLSTN2</i>	Calsyntenin 2	3q23	Le-Niculescu et al. (2009)
<i>CMIP</i> <sup>#</sup>	C-Maf inducing protein	16q23.2	Xu et al. (2014)
<i>CNTNAPS</i> <sup>**</sup>	Contactin associated protein-like 5	2q14.3	Baum et al. (2008a, 2008b)
<i>COMT</i> <sup>*</sup>	Catechol-O-methyltransferase	22q11.21	Zhang et al. (2009)
<i>CREBBP</i> <sup>#</sup>	CREB binding protein	16p13.3	Le-Niculescu et al. (2009)

Gene symbol	Gene name	Location	Source
<i>CSMD1</i> *#	CUB and sushi multiple domains 1	8p23.2	Sklar et al. (2008)
<i>DAOA</i> *	D-Amino acid oxidase activator	13q33.2	Detera-Wadleigh and McMahon (2006)
<i>DAPK1</i> #	Death-associated protein kinase 1	9q21.33	Le-Niculescu et al. (2009)
<i>DCLK1</i>	Doublecortin-like kinase 1	13q13.3	Le-Niculescu et al. (2009)
<i>DCTN5</i> #	Dynactin 5 (P25)	16p12.2	Burton et al. (2007)
<i>DDX27</i>	DEAD (Asp-Glu-Ala-Asp) box polypeptide 27	20q13.13	Xu et al. (2014)
<i>DFNB31</i>	Deafness, autosomal recessive 31	9q32	Ollila et al. (2009)
<i>DGKH</i>	Diacylglycerol kinase, eta	13q14.11	Baum et al. (2008a, 2008b)
<i>DIAPI1</i>	Diaphanous-related formin 1	5q31.3	Le-Niculescu et al. (2009)
<i>DIP2C</i>	Disco-interacting protein 2 homolog C ( <i>Drosophila</i> )	10p15.3	Djurovic et al. (2010)
<i>DISC1</i> *#	Disruption in schizophrenia 1	1q42.2	Hodgkinson et al. (2004)
<i>DLG2</i> *	Dises, large homolog 2 ( <i>Drosophila</i> )	11q14.1	Xu et al. (2014)
<i>DNAJC16</i>	DnaJ (Hsp40) homolog, subfamily C, member 16	1p36.21	Xu et al. (2014)
<i>DNM1</i> *	DNA methyltransferase 1	19p13.2	Veldic et al. (2005)
<i>DOCK3</i>	Dedicator of cytokinesis 3	3p212	Baum et al. (2008a, 2008b)
<i>DPP10</i> *#	Dipeptidyl-peptidase 10 (non-functional)	2q14.1	Le-Niculescu et al. (2009)
<i>DPY19L3</i>	Dpy-19-like 3 ( <i>Caenorhabditis elegans</i> )	19q13.11	Smith et al. (2009)
<i>DRD2</i> *#	Dopamine receptor D2	11q23.2	Le-Niculescu et al. (2009)
<i>DRD4</i> *	Dopamine receptor D4	11p15.5	López León et al. (2005)
<i>DSCAM</i> #	Down syndrome cell adhesion molecule	21q22.2	Xu et al. (2014)
<i>DSEL</i>	Dermatan sulfate epimerase-like	18q22.1	Goossens et al. (2003)
<i>DST</i> #	Dystonin	6p12.1	Le-Niculescu et al. (2009)
<i>DTNBP1</i> *	Dystrobrevin binding protein 1	6p22.3	Gaysina et al. (2009)
<i>EDAR</i>	Ectodysplasin A receptor	2q12.3	Xu et al. (2014)
<i>EGFEMP1</i>	EGF-Like and EMI domain containing 1, pseudogene	3q26.2	Xu et al. (2014)
<i>EGFR</i>	Epidermal growth factor receptor	7p11.2	Sklaret al. (2008)
<i>EIF2S3</i> #	Eukaryotic translation initiation factor 2, subunit 3 gamma, 52 kDa	Xp22.11	Cichon et al., (2011)
<i>ELAVL2</i>	ELAV like neuron-specific RNA binding protein 2	9p21.3	Le-Niculescu et al. (2009)
<i>EPHA5</i>	EPH receptor 5	4q13.1	Le-Niculescu et al. (2009)

Gene symbol	Gene name	Location	Source
<i>FAM13A1</i>	Family with sequence similarity 13, member A	4q22.1	Le-Niculescu et al. (2009)
<i>FAM83F</i>	Family with sequence similarity 83, member F	22q13.1	Xu et al. (2014)
<i>FBN1</i>	Fibrillin 1	15q21.1	Djurovic et al. (2010)
<i>FBXO28</i>	F-Box protein 28	1q42.11	Xu et al. (2014)
<i>FGF12</i>	Fibroblast growth factor 12	3q28	Cichon et al. (2011)
<i>FGFR2</i>	Fibroblast growth factor receptor 2	10q26.13	Xu et al. (2014)
<i>FGGY</i>	FGGY carbohydrate kinase domain containing	1p32.1	Kerner et al. (2013)
<i>FRMD3</i>	FERM domain containing 3	9q21.32	Djurovic et al. (2010)
<i>FOXN3</i>	Forkhead box N3	14q32.11	Baum et al. (2008a, 2008b)
<i>FOXP1#</i>	Forkhead box P1	3p14.1	Le-Niculescu et al. (2009)
<i>FOXP2*#</i>	Forkhead box P2	7q31.1	Xu et al. (2014)
<i>FUT9</i>	Fucosyltransferase 9 (alpha (1,3) fucosyltransferase)	6q16.1	Le-Niculescu et al. (2009)
<i>GAA</i>	Glucosidase, alpha; acid	17q25.3	Le-Niculescu et al. (2009)
<i>GABBR2</i>	Gamma-aminobutyric acid (GABA) B receptor 2	9q22.33	Djurovic et al. (2010)
<i>GABRB1#</i>	Gamma-aminobutyric acid (GABA) A receptor, beta 1	4p12	Burton et al. (2007)
<i>GAD2*</i>	Glutamate decarboxylase 2 (pancreatic islets and brain, 65 kDa)	10p11.23	Heckers et al. (2002)
<i>GNAI1</i>	Guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 1	7q21.11	Le-Niculescu et al. (2009)
<i>GNAI2</i>	Guanine nucleotide binding protein (G protein) alpha 12	7p22.2	Le-Niculescu et al. (2009)
<i>GNAL</i>	Guanine nucleotide binding protein (G protein), alpha activating activity polypeptide, olfactory type	18p11.21	Corradi et al. (2005)
<i>GNG4</i>	Guanine nucleotide binding protein (G protein), gamma 4	1q42.3	Cichon et al. (2011)
<i>GPR50*</i>	G protein-coupled receptor 50	Xq28	Thomson et al. (2005)
<i>GRIA1*</i>	Glutamate receptor, ionotropic, AMPA1 (alpha 1)	5q33.2	Le-Niculescu et al. (2009)
<i>GRIK1</i>	Glutamate receptor, ionotropic, kainate 1	21q21.3	Le-Niculescu et al. (2009)
<i>GRIK2#</i>	Glutamate receptor, ionotropic, kainate 2	6q16.3	Shaltiel et al. (2008)
<i>GRIK4*</i>	Glutamate receptor, ionotropic, kainate 4	11q23.3	Blackwood et al. (2007)
<i>GRIN3A*</i>	Glutamate receptor, ionotropic, N-methyl-D-aspartate 3A	9q31.1	Cichon et al. (2011)
<i>GRM1#</i>	Glutamate receptor, metabotropic 1	6q24.3	Le-Niculescu et al. (2009)
<i>GRM3*</i>	Glutamate receptor, metabotropic 3	7q21.11	Le-Niculescu et al. (2009)



Gene symbol	Gene name	Location	Source
<i>GRM7</i> *	Glutamate receptor, metabotropic 7	3p26.1	Burton et al. (2007)
<i>GSK3B</i> **#	Glycogen synthase kinase 3 beta	3q13.33	Le-Niculescu et al. (2009)
<i>GSTA2</i>	Glutathione S-transferase alpha 2	6p12.2	Le-Niculescu et al. (2009)
<i>GUCY1A</i>	Guanylate cyclase activator 1 A	6p21.1	Xu et al. (2014)
<i>HMOX1</i>	Heme oxygenase 1	22q12.3	Le-Niculescu et al. (2009)
<i>HTR1A</i> *	5-Hydroxytryptamine (serotonin) receptor 1A, G protein-coupled	5q12.3	Kishi et al. (2011)
<i>HTR2A</i> **#	5-Hydroxytryptamine (serotonin) receptor 2A, G-protein-coupled	13q14.2	Le-Niculescu et al. (2009)
<i>HTR3B</i> *	5-Hydroxytryptamine (serotonin) receptor 3B, ionotropic	11q23.1	Hammer et al. (2012)
<i>HTR4</i> *	5-Hydroxytryptamine (serotonin) receptor 4, G-protein coupled	5q32	Ohtsuki et al. (2002)
<i>ICAM3</i>	Intercellular adhesion molecule 3	19p13.2	Cichon et al. (2011)
<i>IGFBP7</i>	Insulin-like growth factor binding protein 7	4q12	Xu et al. (2014)
<i>IGSF21</i>	Immunoglobulin superfamily, member 21	1p36.13	Cichon et al. (2011)
<i>IMP2</i> *	Myo-inositol monophosphatase 2	18p11.21	Sjøholt et al. (2004)
<i>IMPG1</i>	Interphotoreceptor matrix proteoglycan 1	6q14.1	Xu et al. (2014)
<i>INO80</i>	INO80 complex subunit	15q15.1	Xu et al. (2014)
<i>IQGAP2</i>	IQmotif containing GTPase activating protein 2	5q13.3	Le-Niculescu et al. (2009)
<i>ITGAV</i>	Integrin, alpha V	2q32.1	Le-Niculescu et al. (2009)
<i>ITIH3</i> *	Inter-alpha-trypsin inhibitor, heavy chain 3	3p21.1	Hamshire et al. (2013)
<i>JAK1</i>	Janus kinase 1	1p31.3	Xu et al. (2014)
<i>JAM3</i>	Junctional adhesion molecule 3	11q25	Baum et al. (2008a, 2008b)
<i>KCNAB1</i>	Potassium channel, voltage gated subfamily A regulatory beta subunit 1	3q25.31	Le-Niculescu et al. (2009)
<i>KCNB1</i>	Potassium channel, voltage gated Shab-related subfamily B, member 1	20q13.2	Le-Niculescu et al. (2009)
<i>KCNC2</i>	Potassium channel, voltage gated Shaw related subfamily C, member 2	12q14.1	Burton et al. (2007)
<i>KCND2</i> **#	Potassium voltage-gated channel, Shal-related family, member 2	7q31.31	Le-Niculescu et al. (2009)
<i>KCNIP4</i>	Kv channel interacting protein 4	4p15.32	Xu et al. (2014)
<i>KCNK1</i>	Potassium channel, subfamily K, member 1	1q42	Le-Niculescu et al. (2009)
<i>KHDC1</i>	KH homology domain containing 1	6q13	Xu et al. (2014)
<i>KIAA1462</i>	KIAA1462 gene	10p11.23	Xu et al. (2014)

Gene symbol	Gene name	Location	Source
<i>KIF13B</i>	Kinesin family member 13B	8p12	Xu et al. (2014)
<i>KLF12</i>	Kruppel-like factor 12	13q22.1	Le-Niculescu et al. (2009)
<i>KIF1A</i>	Kinesin family member 1 A	2q37.3	Le-Niculescu et al. (2009)
<i>KLHL3</i>	Kelch-like family member 3	5q31.2	Cichon et al. (2011)
<i>KLHL20</i>	Kelch-like family member 20	1q25.1	Xu et al. (2014)
<i>LARGE</i>	Like-glycosyltransferase	22q12.3	Le-Niculescu et al. (2009)
<i>LDLRAD4</i>	Low density lipoprotein receptor class A domain containing 4	18p11.21	Le-Niculescu et al. (2009)
<i>LEF1</i>	Lymphoid enhancer-binding factor 1	4q25	Le-Niculescu et al. (2009)
<i>LHFPL3</i>	Lipoma HMGIC fusion partner-like 3	7q22.2	Cichon et al. (2011)
<i>LMO7</i>	LIM domain 7	13q22.2	Le-Niculescu et al. (2009)
<i>LMOD3</i>	Letomodlin 3	3p14.1	Xu et al. (2014)
<i>LRRC74A</i>	Leucine rich repeat containing 74A	14q24.3	Xu et al. (2014)
<i>LRRFIP1</i>	Leucine rich repeat (In FLII) interacting protein 1	2q37.3	Xu et al. (2014)
<i>LYPD5</i>	LY6/PLAUR domain containing 5	19q13.31	Cichon et al. (2011)
<i>MACRODMACRO</i>	domain containing 1	1q13.1	Cichon et al. (2011)
<i>MAD1L1</i>	*MAD1 mitotic arrest deficient-like 1 (yeast)	7p22.3	Cichon et al. (2011)
<i>MAOA</i>	*# Monoamine oxidase A	Xp11.3	Fan et al. (2010)
<i>MARCH1</i>	Membrane-associated ring finger (C3HC4) 1, E3 ubiquitin protein ligase	4q32.2	Xu et al. (2014)
<i>MBNL2</i>	Musclebind-like splicing regulator 2	13q32.1	Le-Niculescu et al. (2009)
<i>MBP</i>	Myelin basic protein	18q23	Le-Niculescu et al. (2009)
<i>MDHI</i>	Malate dehydrogenase 1, NAD (soluble)	2p13.3	Le-Niculescu et al. (2009)
<i>MFSD11</i>	Major facilitator superfamily domain containing 11	17q25.1	Xu et al. (2014)
<i>MTHFR</i>	*#Methylenetetrahydrofolate reductase (NAD(P)H)	1p36.22	El-Hadidy et al. (2014)
<i>MTMR3</i>	Myotubularin related protein 3	22q12.2	Xu et al. (2014)
<i>MYCBP2</i>	MYC binding protein 2, E3 ubiquitin protein ligase	13q22.3	Le-Niculescu et al. (2009)
<i>MYO5B</i>	Myosin VB	18q21.1	Sklar et al. (2008)
<i>MYO18B</i>	Myosin XVIIIIB	22q12.1	Xu et al. (2014)
<i>MYT1L</i>	*# Myelin transcription factor 1-like	2p25.3	Le-Niculescu et al. (2009)
<i>NAALAD2N</i>	acetylated alpha-linked acidic dipeptidase 2	11q14.3	Xu et al. (2014)
<i>NALCN</i>	*# Sodium leak channel, non-selective	13q32.3	Sklar et al. (2008)

Gene symbol	Gene name	Location	Source
<i>NAV2</i>	Neuron navigator 2	<i>11p15.1</i>	Le-Niculescu et al. (2009)
<i>NCALD</i>	Neurocalcin delta	<i>8q22.3</i>	Xu et al. (2014)
<i>NCAM1*</i>	Neural cell adhesion molecule 1	<i>11q23.2</i>	Le-Niculescu et al. (2009)
<i>NCAN</i>	Neurocan	<i>19p12</i>	Cichon et al. (2011)
<i>NCKAP5#</i>	NCK-associated protein 5	<i>2q21.2</i>	Smith et al. (2009)
<i>NDUFAB1</i>	NADH dehydrogenase (ubiquinone) 1, alpha/beta subcomplex, 1, 8 kDa	<i>16p12.2</i>	Burton et al. (2007)
<i>NDUFS2</i>	NADH dehydrogenase (ubiquinone) Fe-S protein 2, 49 kDa (NADH-coenzyme Q reductase)	<i>1q23.3</i>	Le-Niculescu et al. (2009)
<i>NFLA#</i>	Nuclear factor I/A	<i>1p31.3</i>	Le-Niculescu et al. (2009)
<i>NFIB</i>	Nuclear factor I/B	<i>9p24.1</i>	Le-Niculescu et al. (2009)
<i>NIPBL</i>	Nipped-B homolog ( <i>Drosophila</i> )	<i>5p132</i>	Xu et al. (2014)
<i>NME7</i>	NME/NM23 family member 7	<i>1q24.2</i>	Xu et al. (2014)
<i>NOS1AP*</i>	Nitric oxide synthase 1 (neuronal) adaptor protein	<i>1q23.3</i>	Xu et al. (2014)
<i>NPAS3*</i>	Neuronal PAS domain protein 3	<i>14q13.1</i>	Pickard et al. (2009)
<i>NPFRR2</i>	Neuropeptide FF receptor 2 isoform 1	<i>4q13.3</i>	Xu et al. (2014)
<i>NR3C1</i>	Nuclear receptor subfamily 3, group C, member 1 (glucocorticoid receptor)	<i>5q31.3</i>	Le-Niculescu et al. (2009)
<i>NRCAM#</i>	Neuronal cell adhesion molecule	<i>7q31.1</i>	(Le-Niculescu et al., (2009)
<i>NRG1*</i>	Neuregulin 1	<i>8p12</i>	Prata et al. (2009)
<i>NRG3*</i>	Neuregulin 3	<i>10q23.1</i>	Xu et al. (2014)
<i>NTF3*</i>	Neurotrophin 3	<i>12p13.31</i>	Cichon et al. (2011)
<i>NTRK2*</i>	Neurotrophic tyrosine kinase, receptor, type 2	<i>9q21.33</i>	Smith et al. (2009)
<i>NUP214</i>	Nucleoporin 214 kDa	<i>9q34.13</i>	Xu et al. (2014)
<i>OC90</i>	Otoconin 90	<i>8q24.22</i>	Xu et al. (2014)
<i>OLIG2*</i>	Oligodendrocyte lineage transcription factor 2	<i>21q22.11</i>	Le-Niculescu et al. (2009)
<i>OPRM1*</i>	Opioid receptor, mu 1	<i>6q25.2</i>	Le-Niculescu et al. (2009)
<i>OR1B1</i>	Olfactory receptor, family 1, subfamily B, member 1	<i>9q33.2</i>	Xu et al. (2014)
<i>PALB2</i>	Partner and localizer of BRCA2	<i>16p12.2</i>	Tesli et al. (2010)
<i>PARD3</i>	Par-3 family cell polarity regulator	<i>10p11.21</i>	Le-Niculescu et al. (2009)
<i>PARD3B#</i>	Par-3 family cell polarity regulator beta	<i>2q33.3</i>	Cichon et al. (2011)

Gene symbol	Gene name	Location	Source
<i>PCDHAC2</i> <sup>#</sup>	Protocadherin-alpha, subfamily C, 2	5q31.3	Pedrosa et al. (2008)
<i>PCDH7</i>	Protocadherin 7	4p15.1	Le-Niculescu et al. (2009)
<i>PCDH9</i> <sup>#</sup>	Protocadherin 9	13q21.32	Le-Niculescu et al. (2009)
<i>PCLO</i>	Piccolo presynaptic cytomatrix protein	7q21.11	Choi et al. (2011)
<i>PDE10A</i>	Phosphodiesterase 10 A	6q26	Le-Niculescu et al. (2009)
<i>PDE4B</i> <sup>**#</sup>	Phosphodiesterase 4B, CAMP-specific	1p31.3	Xu et al. (2014)
<i>PER3</i> <sup>*</sup>	Period circadian clock 3	1p3623	Benedetti et al. (2008)
<i>PDLIM5</i> <sup>*</sup>	PDZ and LIM domain 5	4q22.3	Shi et al. (2008)
<i>PHF20</i>	PHD finger protein 20	20q11.22	Xu et al. (2014)
<i>PIK3R1</i>	Phosphoinositide-3-kinase, regulatory subunit 1 (alpha)	5q13.1	Le-Niculescu et al. (2009)
<i>PLCL2</i>	Phospholipase C-like 2	3p24.3	Xu et al. (2014)
<i>PLXNA2</i> <sup>*</sup>	Plexin A2	1q32.2	Le-Niculescu et al. (2009)
<i>PPM1B</i>	Protein phosphatase, Mg <sup>2+</sup> /Mn <sup>2+</sup> dependent, 1B	2p22.1	Le-Niculescu et al. (2009)
<i>PPP2R2A</i>	Protein phosphatase 2, regulatory subunit B, alpha	8p212	Xu et al. (2014)
<i>PPP2R2C</i>	Protein phosphatase 2, regulatory subunit B, gamma	4p16.1	Baum et al. (2008a, 2008b)
<i>PRKCE</i>	Protein kinase C, epsilon	2p21	Le-Niculescu et al. (2009)
<i>PRPF18</i>	Pre-mRNA processing factor 18	10p12.33	Xu et al. (2014)
<i>PRR16</i>	Proline rich 16	5q23.1	Xu et al. (2014)
<i>PRSS55</i>	Protease, serine, 55	8p23.1	Xu et al. (2014)
<i>PTN</i>	Pleiotrophin	7q33	Le-Niculescu et al. (2009)
<i>PTPRK</i>	Protein tyrosine phosphatase, receptor type, K	6q22.33	Le-Niculescu et al. (2009)
<i>PTPRM</i> <sup>#</sup>	Protein tyrosine phosphatase, receptor type, M	18p11.23	Le-Niculescu et al. (2009)
<i>PTPRO</i>	Protein tyrosine phosphatase, receptor type, O	12p12.3	Xu et al. (2014)
<i>PYDC2</i>	Pyridin domain containing 2	3q28	Cichon et al. (2011)
<i>QKI</i> <sup>*</sup>	Quaking homolog, KH domain RNA binding	6q26	Le-Niculescu et al. (2009)
<i>RABEP1</i>	Rabaptin, RAB GTPase binding effector protein 1	17p13.2	Djurovic et al. (2010)
<i>RABGAP1</i> <sup>RAB</sup>	GTPase activating protein 1-like	1q25.1	Xu et al. (2014)
<i>RADRMAD5</i> 1, paralog B		14q24.1	Xu et al. (2014)
<i>RASGEF1</i> <sup>RAB</sup>	protein-specific guanine nucleotide-releasing factor 2	5q14.1	Le-Niculescu et al. (2009)
<i>RBF1</i> <sup>RAB</sup>	binding protein, fox-1 homolog ( <i>C. elegans</i> ) 1	16p13.3	Le-Niculescu et al. (2009)

Gene symbol	Gene name	Location	Source
<i>RBM3</i>	mRNA binding motif protein 23	14q11.2	Xu et al. (2014)
<i>RC3H1</i>	ring finger and CCCH-type domains 1	1q25.1	Xu et al. (2014)
<i>RLF</i>	Rearranged L-Myc fusion	1p34.2	Xu et al. (2014)
<i>ROR1</i>	RAR-related orphan receptor alpha	15q22.2	Le-Niculescu et al. (2009)
<i>ROR2</i>	RAR-related orphan receptor beta	9q21.13	Le-Niculescu et al. (2009)
<i>RSPOR1</i>	spondin 4	20p13	Xu et al. (2014)
<i>RXR1</i>	retinoid X receptor, gamma	1q23.3	Le-Niculescu et al. (2009)
<i>RYR1</i>	Ryanodine receptor 3	15q13.3	Le-Niculescu et al. (2009)
<i>SCA1</i>	secretory carrier membrane protein 1	5q14.1	Le-Niculescu et al. (2009)
<i>SCN1A</i>	sodium channel, voltage gated, type VII alpha subunit	2q24.3	Xu et al. (2014)
<i>SHISA1</i>	sa family member 9	16p13.12	Xu et al. (2014)
<i>SLC39A1</i>	zinc transporter, member 3	19p13.3	Baum et al. (2008a, 2008b)
<i>SLC6A1</i>	neurotransmitter transporter, dopamine, member 3	5p15.33	Greenwood et al. (2006)
<i>SLC8A1</i>	sodium/calcium exchanger, member 1	2p22.1	Le-Niculescu et al. (2009)
<i>SOD1</i>	superoxide dismutase 1, soluble	21q22.11	Le-Niculescu et al. (2009)
<i>SORCS1</i>	sorting nexin-related VPS10 domain containing receptor 2	4p16.1	Soronen et al. (2010)
<i>SPARC</i>	secretory associated antigen 16	2q34	Xu et al. (2014)
<i>SPARC</i>	secreted protein, acidic, cysteine-rich (osteonectin)	5q31.3	Le-Niculescu et al. (2009)
<i>SPAS1</i>	spastin	2q22.3	Le-Niculescu et al. (2009)
<i>STAB1</i>	stabilin 1	3p21.1	Baum et al. (2008a, 2008b)
<i>STK2</i>	serine/threonine kinase 24	13q32.2	Le-Niculescu et al. (2009)
<i>STX12</i>	syntaxin 12	1p35.3	Xu et al. (2014)
<i>SVIL</i>	Supervillin	10p11.23	Purcell et al. (2009)
<i>SYN1</i>	synapsin III	22q12.3	Le-Niculescu et al. (2009)
<i>SYN1</i>	synapsin repeat containing, nuclear envelope 1	6q25.2	Ferreira et al. (2008)
<i>TBC1D22B</i>	TBC domain family, member 22B	6p21.2	Xu et al. (2014)
<i>TBX1</i>	transcription factor 1	7q34	Xu et al. (2014)
<i>TCF11</i>	transcription factor 4	18q21.1	Del-Favero et al. (2002)
<i>TEAD1</i>	TEA domain family member 1	11p15.4	Xu et al. (2014)
<i>TEN1</i>	neurin transmembrane protein 4	11q14.1	Sklar et al., 2011

Gene symbol	Gene name	Location	Source
<i>TLAM</i>	cell lymphoma invasion and metastasis 1	21q22.11	Le-Niculescu et al. (2009)
<i>TLE4</i>	Transducin-like enhancer of split 4	9q21.31	Cichon et al. (2011)
<i>TMEM45</i>	transmembrane protein 252	9q21.11	Xu et al. (2014)
<i>TNIKTRAF2</i>	and NCK interacting kinase	3q26.31	Le-Niculescu et al. (2009)
<i>TPH1</i>	Tryptophan hydroxylase 1	11p15.1	Chen et al. (2012)
<i>TPH2</i>	Tryptophan hydroxylase 2	12q21.1	Cichon et al. (2008)
<i>TPST1</i>	Tyrosylprotein sulfotransferase 2	22q12.1	Le-Niculescu et al. (2009)
<i>TRAF1</i>	NF- $\kappa$ B-inducing kinase 1	3p22.2	Mühlhausen et al. (2014)
<i>TRIO</i>	the guanine nucleotide exchange factor	5p15.2	Xu et al. (2014)
<i>TRPM2</i>	transient receptor potential cation channel, subfamily M, member 2	21q22.3	McQuillin et al. (2006)
<i>TRPM3</i>	transient receptor potential cation channel, subfamily M, member 3	9q21.12	Le-Niculescu et al. (2009)
<i>TRPV1</i>	transient receptor potential cation channel, subfamily V member 1	17p13.2	Cichon et al. (2011)
<i>TRRAP</i>	transformation/transcription domain-associated protein	7q22.1	Xu et al. (2014)
<i>TSHZ1</i>	zinc finger homeobox 2	20q13.2	Le-Niculescu et al. (2009)
<i>TSPAT1</i>	transcriptin 8	12q21.1	Scholz et al. (2010)
<i>VAMP5</i>	vesicle-associated membrane protein 5	Xq28 or Yq12	Saito et al. (2000)
<i>VIP1</i>	Vasoactive Intestinal Peptide	6q25.2	Soria et al. (2010)
<i>WDR60</i>	WD repeat domain 60	7q36.3	Xu et al. (2014)
<i>WDR7</i>	WD repeat domain 7	18q21.31	Xu et al. (2014)
<i>WNLK</i>	lysine deficient protein kinase 2	9q22.31	Cichon et al. (2011)
<i>XBPA</i>	box-binding protein 1	22q12.1	Kakinuchi et al. (2003)
<i>XKR2</i>	Kell blood group complex subunit-related family, member 2	8q12.1	Djurovic et al. (2010)
<i>XYL1</i>	xylosyltransferase 1	3q29	Xu et al. (2014)
<i>ZEB2</i>	zinc finger E-box binding homeobox 2	2q22.3	Cichon et al. (2011)
<i>ZBTB20</i>	zinc finger and BTB domain containing 20	3q13.31	Xu et al. (2014)
<i>ZDHHC1</i>	zinc finger, DHHC-type containing 1	6q25.3	Le-Niculescu et al. (2009)
<i>ZHX2</i>	zinc fingers and homeoboxes 2	8q24.13	Le-Niculescu et al. (2009)
<i>ZNF274</i>	zinc finger protein 274	19q13.43	Xu et al. (2014)
<i>ZNF385D</i>	zinc finger protein 385D	3p24.3	Xu et al. (2014)
<i>ZNF536</i>	zinc finger protein 536	19q13.11	Djurovic et al. (2010)



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Gene symbol	Gene name	Location	Source
<i>ZNF628</i>	zinc finger protein 618	9q33.1	Cichon et al. (2011)
<i>ZNF804</i>	zinc finger protein 804 A	2q32.1	Williams et al.(2011)

\* denotes BPD genes that overlap with clinically relevant genes for schizophrenia.

# denotes BPD genes that overlap with clinically relevant genes for autism spectrum disorder.

**Table 2**

Distribution of bipolar disorder (BPD) genes among chromosomes.

Chromosome	Total	Proportion of total BPD genes	P arm	Q arm
1	24	8.3%	12	12
2	23	7.9%	6	17
3	18	6.2%	9	9
4	16	5.5%	5	11
5	18	6.2%	4	14
6	18	6.2%	6	12
7	20	6.9%	5	15
8	10	3.4%	5	5
9	19	6.6%	4	15
10	11	3.8%	8	3
11	18	6.2%	7	11
12	8	2.8%	3	5
13	11	3.8%	NA	11
14	7	2.4%	NA	7
15	7	2.4%	NA	7
16	9	3.1%	6	3
17	5	1.7%	2	3
18	11	3.8%	5	6
19	8	2.8%	4	4
20	5	1.7%	1	4
21	7	2.4%	NA	7
22	12	4.1%	NA	12
X	3	1.0%	2	1
Y	0	0.0%	0	0
Both X and Y	2	0.7%	NA	NA
Total	290	100%	94	194

Total number of BPD genes were counted for each chromosome and chromosome arm.

p = short arm; q = long arm.

NA = not applicable due to chromosome structure or location.