Data Supplement for Mullins et al., GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. Am J Psychiatry (doi: 10.1176/appi.ajp.2019.18080957).

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Supplementary Note

Phenotype definition

Items from structured clinical interviews provided information on self-harm, suicidal ideation, plans and suicide attempt for Psychiatric Genomics Consortium (PGC) cohorts (Supplementary Table 4). Lifetime suicide attempt (SA) was defined across cohorts as a deliberate act of self-harm with at least some intent to result in death. Individuals who did not endorse suicide attempt were included in the non-attempter group and individuals missing information on suicide attempt were excluded. Phenotypic analyses were performed to assess the consistency of the suicide attempt phenotype across the 46 discovery cohorts. First, the association between the psychiatric interview used and the prevalence of suicide attempt in the cohorts was tested, using a linear regression model including psychiatric disorder as a covariate. The only psychiatric interview which showed an association with the prevalence of suicide attempt was the Schedule for Affective Disorders and Schizophrenia (SADS), which was used by eight cohorts and correlated with a higher prevalence of suicide attempt (P = 0.038). We note that the SADS interview has a specific item on suicide attempt and does not include self-harm or suicidal ideation (which are assessed using other items in the SADS) (Supplementary Table 4). The interview also specifies that evidence of intent to die is essential for suicide attempt and in the absence of intent "non-suicidal self-injurious behavior" is instead indicated. Therefore, we do not anticipate that the suicide attempter groups from cohorts assessed using the SADS may incorrectly include cases of suicidal ideation only. Furthermore, of the eight cohorts which used the SADS interview, five recruited psychiatric cases from inpatient hospital admissions and hence these cohorts may be more severe and have a higher prevalence of suicide attempt than other cohorts (Supplementary Tables 1-3).

Second, we assessed whether there was any association between prevalence of suicide attempt and interview items which are specific to suicide attempt only or those which assess a spectrum of suicidality (ranging from absence, to self-harm, ideation, plan and suicide attempt) in a single item

(Supplementary Table 4). In the latter, psychiatric cases were split into suicide attempters only versus other psychiatric cases according to the coding of the item. Individuals with missing data were excluded. Using a linear regression model, controlling for psychiatric disorder, there was no association between the prevalence of suicide attempt and the specificity of the item used. Since interviews assessing the spectrum of suicidality in a single item did not produce higher estimates of suicide attempt, this suggests that these do not report cases of suicidal ideation only as suicide attempters. In summary, these analyses suggest that the psychiatric interview used across cohorts does not result in heterogeneity in the definition of suicide attempt.

Polygenic risk scoring

Polygenic risk scoring was used to investigate the genetic relationship between suicide attempt and the psychiatric disorders and to test for overlap in the genetic etiology of suicide attempt between major depressive disorder (MDD), bipolar disorder (BIP) and schizophrenia (SCZ). Table S5 summarises the polygenic scoring analyses conducted, showing the discovery and test datasets used to investigate these hypotheses. PRSice software was used to generate polygenic risk scores (PRS), according to standard protocol (1). Discovery GWAS results were pruned for linkage disequilibrium (LD) using the P value informed clumping method in PLINK (--clump-p1 1 --clump-p2 1 --clump-r2 0.1 --clump-kb 250). This preferentially retains SNPs with the strongest evidence of association and removes SNPs in LD ($r^2 > 0.1$) that show weaker evidence of association within 250Kb windows, based on the LD structure in the test dataset. Subsets of SNPs were selected from the results at nine increasingly liberal P value thresholds (P < 0.0001, P < 0.001, P < 0.01, P < 0.05, P < 0.1, P < 0.2, P < 0.3, P < 0.4, P < 0.5). In the test datasets, the SNP probabilities were converted to best-guess data with a genotype call probability cut-off of 0.8. Sets of alleles, weighted by their log odds ratios (OR) from the discovery GWAS, were summed into PRS for each individual in the test datasets using PLINK. PRS were tested for association with suicide attempter status in the test datasets using a logistic regression model, including five genetic principal components (PCs) and a covariate for each cohort in the test dataset.

The amount of variance explained by the PRS (R²) is presented on the liability scale, which accounts for the proportion of cases in the test dataset (2).

First, PRS for BIP, major depression and SCZ were used to investigate whether suicide attempters and non-attempters differ in genetic liability for the psychiatric disorder they are affected by. To ensure no overlap between the discovery and test datasets, PRS for psychiatric disorders were generated using PGC cohorts not included in the suicide attempt analyses. All cohorts have been described in previous publications on GWAS of psychiatric disorders conducted by the PGC (3-5). The discovery GWAS for BIP consisted of 11 PGC cohorts totaling 8,711 BIP cases and 15,283 controls, and for SCZ included 25,756 SCZ cases and 35,686 controls from 40 PGC cohorts. The discovery GWAS for major depression is a recent meta-analysis of PGC MDD cohorts and samples from deCODE, GERA, iPSYCH, Generation Scotland and UK Biobank (3). The phenotype analysed in this study included clinically defined MDD cases as well as self-reported MDD symptoms or treatment and thus is referred to as 'major depression' (3). Results of this meta-analysis were available in turn excluding each of the 16 PGC MDD cohorts in the suicide attempt study, which allowed us to generate independent PRS for each of the 16 cohorts while maximising the discovery GWAS sample size. These discovery GWAS had approximately 59,000 cases and 112,000 controls. The PRS for BIP, major depression and SCZ were tested for association with suicide attempter versus non-attempter status in the same disorder using logistic regression as described previously. Second, based on the results of these analyses, PRS for major depression were also tested for association with suicide attempt in BIP and SCZ.

Third, in order to investigate genetic overlap in suicide attempt across psychiatric disorders, the results of the three GWAS on suicide attempt (SA in MDD, SA in BIP and SA in SCZ) were used in turn as discovery studies and PRS for suicide attempt were tested for association with SA in the other disorders. The Bonferroni corrected significance threshold for the polygenic scoring analyses is 0.006, adjusting for eight independent tests (Supplementary Table S5).

Replication studies

UK Biobank

Genetic associations with suicide attempt were tested for replication in two independent samples of patients with mood disorders drawn from the UK Biobank and iPSYCH. The UK Biobank is a prospective cohort study of 501,726 individuals, recruited at 23 centres across the United Kingdom (6). Genotypic data were available for 488,380 individuals and were imputed to the HRC, UK10K and 1,000 Genomes Phase 3 reference panels using IMPUTE4 to identify \approx 93M variants for 487,409 individuals (7). Variants for analysis were limited to those with minor allele frequency >= 0.01, imputation INFO-score >= 0.4, and which were either genotyped or imputed to the HRC reference panel, leaving a total of 7794483 SNPs for analysis. Using the genotyped SNPs, individuals were removed if: recommended by the UK Biobank core analysis team for unusual levels of missingness or heterozygosity; SNP genotype call rate < 98%; related to another individual in the dataset (KING r < 0.044, equivalent to removing up to third-degree relatives inclusive); phenotypic and genotypic gender information was discordant (X-chromosome homozygosity (FX) < 0.9 for phenotypic males, FX > 0.5 for phenotypic females). Removal of relatives was performed using a greedy algorithm, which minimises exclusions (for example, by excluding the child in a mother-father-child trio). All analyses were limited to individuals of White Western European ancestry, as defined by 4-means clustering on the first two genetic principal components provided by the UK Biobank (7). Principal component analysis was also performed on the European-only subset of the data using the software flashpca2 (8).

Extensive phenotypic data are available for UK Biobank participants from health records and questionnaires, including an online follow-up questionnaire focussing on mental health (Mental Health Questionnaire, MHQ). Participants were classified as having a mood disorder if they either self-reported a professional diagnosis of depression or bipolar disorder as part of the MHQ [UK Biobank field 20544] or if they met criteria for depression on MHQ questions derived from the Composite International Diagnostic Interview (CIDI). To meet these latter criteria, participants must have

reported ever feeling depressed [UK Biobank field 20446] or anhedonic [UK Biobank field 20441] for two weeks in a row, for at least most of the day [UK Biobank field 20436] almost every day [UK Biobank field 20439] with more than a little interference with daily activities [UK Biobank field 20440]. In addition, they must have reported experiencing at least five of the following symptoms in this period of depression or anhedonia: depression [UK Biobank field 20446], anhedonia [UK Biobank field 20441], tiredness [UK Biobank field 20449], weight change [UK Biobank field 20536], sleep change [UK Biobank field 20532], loss of concentration [UK Biobank field 20435], worthlessness [UK Biobank field 20450] and thoughts of death [UK Biobank field 20437]. The MHQ additionally contained screening questions for bipolar disorder (9). However, for the purpose of defining potential bipolar disorder, all individuals scoring positively on these screening questions were also required to meet the CIDI depression criteria defined above, and as such participants with potential bipolar disorder were a subset of those meeting criteria for depression. Individuals who self-reported a professional diagnosis of psychosis on the MHQ [UK Biobank field 20544] were excluded. Suicide attempters with mood disorders (n=2149) were defined as those who answered yes to the question "Have you ever harmed yourself with the intention to end your life?" [UK Biobank field 20483]. Non-attempters with mood disorders were defined as those who reported no self-harm on the MHQ (n=35912). A genome-wide association study was performed comparing suicide attempters versus non-attempters with mood disorders using BGenie v.1.2 (7), covarying for 6 PCs, and factors capturing site of recruitment and genotyping batch.

*i*PSYCH

The *i*PSYCH study was approved by the regional Danish ethics committee and the Danish Data Protection Agency (10). DNA preparation, genotyping on the Illumina PsychChip array and quality control were performed as described previously (11, 12). Individuals with mood disorders were identified based on ICD-10 codes (F30-F39) from the Danish Psychiatric Central Research Register and the National Registry of Patients, both complete until December 31, 2016 (10). Suicide attempters with mood disorders (n=4943) were defined as those with diagnoses of suicide attempt (ICD-10: X60-

X84, equivalent to intentional self-harm), those with suicide attempt indicated as 'reason for contact', and with a main diagnosis of poisoning (ICD-10: T39, T42, T43, and T58) or those with a diagnosis in the ICD-10: F chapter as main diagnosis and report of poisoning by drugs or other substances (ICD-10: T36–T50, T52–T60) or injuries to hand, wrist, and forearm (ICD-10: S51, S55, S59, S61, S65, S69). Only contacts starting at age 10 years old or older were considered. Individuals who died by suicide according to the Cause of Death Register were also included in the suicide attempter group. Non-attempters were defined as mood disorder cases not fulfilling any of these criteria (n=15849).

Analysis of depressive symptoms and suicide attempt in schizophrenia

Data on symptoms of illness in schizophrenia were available for eight of the PGC SCZ cohorts included in the GWAS on suicide attempt. Clinical symptoms were assessed using the OPCRIT (Operational Criteria for Psychotic Illness), PANSS (Positive and Negative Syndrome Scale), Lifetime Dimensions of Psychosis Scale (LDPS) or the Comprehensive Assessment of Symptoms and History (CASH) (13-16). As previously described, factor analyses were performed on these data to identify a quantitative depressive symptom dimension, harmonized across instruments and cohorts (17). Data were available for 1426 suicide attempters and 2428 non-attempters with schizophrenia. The association between standardized depressive symptom-based factor scores and suicide attempt was investigated using a logistic regression model, covarying for sex and cohort. Higher depressive symptom factor scores were significantly associated with suicide attempt (OR = 1.67, C.I. 1.56 -1.79, P = 2.7 x 10⁻⁴⁷).

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Supplementary Tables and Figures

Table S1: Description of 16 major depressive disorder cohorts

Cohort (References)	PGC label	Country	Ascertainment and Evaluation of Cases	Inclusion criteria (lifetime) of Cases	Exclusion criteria (lifetime) of Cases
BOMA 1-3	boma	Germany	Consecutive inpatients; SCID or SADS interview; medical records	DSM-IV MDD; German ancestry; age ≥ 18	BIP, hypomania, NAP, MDD related to SUD
CoFaMS 4	cof3	Australia	Opportunistic; inpatient & outpatient; SCID or MINI	DSM-IV MDD	BIP, NAP, MDD related to SUD
PsyCoLaus 5	col3	Switzerland	Random population sample; DIGS	DSM-IV MDD, age 35-66	BIP, hypomania, NAP, MDD related to SUD
GenRED1 6,7	gens	USA	Opportunistic; DIGS3; medical records or informant (subset)	DSM-IV MDD (recurrent or episode >3 yrs) & onset <31 yrs; FHx MDD in sibling or parent	BIP, NAP, mod-severe ID; FHx BIP; if SUD, MDD onsets without <2y of sobriety
GenRED2 6	grnd	USA	Opportunistic; DIGS; medical records or informant (subset)	DSM-IV MDD (recurrent or episode >3 yrs) & onset <31 yrs; FHx MDD in sibling or parent	BIP, NAP, mod-severe ID; FHx BIP; if SUD, MDD onsets without <2y of sobriety
GSK/MPIP 8	gsk2	Germany	Inpatients; SCAN	DSM-IV MDD (recurrent, mod-severe)	BIP, NAP, SUD, mood-incongruent psychosis, OCD, PTSD, secondary MD
MARS 9-11	mmi2 mmo4	Germany	Inpatients; CIDI	DSM-IV MDD	BIP, SUD, secondary MD, severe medical conditions
NESDA/NTR: NESDA 12,13	nes1	Netherlands	Psychiatric outpatients, primary care, & population; CIDI	DSM-IV MDD	BIP, NAP, SUD
NESDA/NTR: NTR 12,13	nes1	Netherlands	Twin registry; longitudinal MDD sx; CIDI (subset)	DSM-IV MDD	Mania (if interviewed)
QIMR 14,15	qi3c qi6c qio2	Australia	Australian Twin Registry (proband most severe, sx, or earlier onset); SSAGA	DSM-IV MDD	MDD related to SUD
RADIANT-UK 16	rad3	UK	UK outpatients from DeNT, DeCC, GENDEP studies; SCAN	DSM-IV MDD (recurrent in DeCC & DeNT; MDD FHx in DeNT)	BIP, NAP, MDD related to SUD; BIP FHx
RADIANT-GER 16	rage	Germany	German outpatients from DeNT, DeCC, GENDEP studies; SCAN	DSM-IV MDD (recurrent in DeCC & DeNT; MDD FHx in DeNT)	BIP, NAP, MDD related to SUD; BIP FHx
SHIP 0 17	shp0	Germany	Study of Health in Pomerania; CIDI	DSM-IV MDD	BIP, MDD related to SUD
STAR*D 18	stm2	USA	Outpatients in clinical trial; clinical interviews	DSM-IV MDD	BIP, NAP

Abbreviations: SCID=Structured Clinical Interview for DSM-IV, SADS=Schedule for Affective Disorders and Schizophrenia, MDD = major depressive disorder, BIP = bipolar disorder, NAP=non-affective psychosis, SUD = substance use disorder, MINI = MINI International Neuropsychiatric Interview,
DIGS=Diagnostic Interview for Genetic Studies, Fhx = family history, ID = intellectual disability, SCAN = Schedules for Clinical Assessment in Neuropsychiatry, OCD = obsessive compulsive disorder, PTSD = post-traumatic stress disorder, CIDI = Composite International Diagnostic Interview, sx = symptoms

References	Number	Citation
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	16	Lewis CM et al. Genome-wide association study of major recurrent depression in the U.K. population. Am J Psychiatry 167, 949-57 (2010).
	17	Volzke H et al. Cohort profile: the study of health in Pomerania. Int J Epidemiol 40, 294-307 (2011).
	18	Shyn SI et al. Novel loci for major depression identified by genome-wide association study of Sequenced Treatment Alternatives to Relieve Depression and meta-analysis of three studies. Mol Psychiatry 16, 202-15 (2011).

Table S2: Description of 21 bipolar disorder cohorts

Cohort (References)	PGC label	Country	Ascertainment and Evaluation of Cases	Inclusion criteria (lifetime) of Cases	Exclusion criteria (lifetime) of Cases
BOMA-Germany I1-4	bonn	Germany	Consecutive admissions to in-patent units, SCID-I, SADS-L, medical records, FHx, OPCRIT	DSM-IV BIP I or BIP II	I.
Trinity College Dublin 5	dub1	Ireland	Hospitals and Community psychiatric facilities, SCID, case note review	DSM-IV BIP I	
FaST, TGEN1, TGEN2 6	fat2	USA	Hospitals, ADE, MINI	DSM-IV BIP I or BIP II	
French PGC2 6	fran	France	DIGS, FIGS, medical case notes, mood scales, self-rating questionnaires assessing dimensions	DSM-IV BIP I or BIP II	
			Advertisements in hospitals, clinics, primary care physician offices, patient support groups,		Dx of intravenous drug dependency/use, mood incongruent psychotic sx, manic
BACCs 7	gsk1	UK, Canada	SCAN CATEGO algorithm	DSM-IV or ICD-10 BIP I or BIP II	episo des only with alcohol/substance abuse/dependence/medical illnesses/medications
			Mayo Clinic Bipolar Biobank, patients ascertained through routine clinical appointments, in-patients in		
Mayo Clinic 8	may1	USA	mood disorder units and recruitment advertising, SCID	DSM-IV-TR BIP I/BIP II/ schizoaffective	•
Pritzker Neuropsychiatric Disorde	ers				
Research Consortium 7, 9	mich	USA	NIMH Genetics Initiative Repository, DIGS, FIGS, medical record review	DSM-III or IV BIP I	Suspected major depression
STEP1 5,7	stp1	USA	ADE, MINI	DSM-IV BIP I	
STEP2	st2c	USA	Hospitals, ADE, MINI	DSM-IV BIP I or BIP II	
TOP 7 10	top7	Norway	Out-patient and in-patient psychiatric units, SCID-I, case note review, follow up interview	DSM-IV BIP I, BIP II, SAB, BIP-NOS	IQ score < 70
TOP8 10	top8	Norway	Out-patient and in-patient psychiatric units, SCID-I, case note review, follow up interview	DSM-IV BIP I, BIP II, SAB, BIP-NOS	IQ score < 70
UCL5, 11	uclo	UK	Clinical diagnosis according to UK National Health Service (NHS) psychiatrists at interview, SADS-L, OPCRI	T DSM-IV BIP I	
UMEA	ume4	Sweden	MINI, DIGS, FIGS, SCAN	DSM-IV-TR BIP	
WTCCC 5, 7, 12	wtcc	UK	Individuals in contact with mental health services, SCAN	RDC BIP I, BIP II, SAB, BIP-NOS	
GAIN 7, 13	gain	USA	Multiplex families, sibling pair families or individuals, DIGS, FIGS, medical records	DSM IIR & IV BIP I or SAB	
BOMA-Germany II 14	bmg2	Germany	Consecutive admissions to in-patient units, AMDP, medical records, family history, OPCRIT	DSM-IV lifetime BIP	
BOMA-Germany III 14	bmg3	Germany	Recruited from psychiatric hospitals, AMDP, CID-S, SADS-L, SCID, medical records, family history, OPCRIT	DSM-IV lifetime BIP	
BOMA-Poland 14	bmpo	Poland	Recruited from Department of Psychiatry, SCID	DSM-IV lifetime BIP	
BOMA-Spain14	bmsp	Spain	Recruited from hospital mental health departments, SADS-L, OPCRIT, medical records, FISC	DSM-IV and RDC BIP	
NovaScotia	hal 2	Canada	Recruited from specialty mood disorder clinics, SADS-L	DSM-IV and RDC BIP	
BOMA-Romania15	rom3	Romania	Consecutive admissions to psychiatric hospital, DIGS, FIGS, medical records, family reports	DSM-IV BIP I	

Abbreviations: SCID = Structured Clinical Interview for DSM-IV, SADS-L = Schedule for Affective Disorders and Schizophrenia Lifetime Version, FHx = family history, OPCRIT = Operational Criteria Checklist, BIP = bipolar disorder, ADE = Affective Disorders Evaluatio
MINI = MINI International Neuropsychiatric Interview, DIGS = Diagnostic Interview for Genetic Studies, FIGS = Family Interview for Genetic Studies, SCAN = Schedules for Clinical Assessment in Neuropsychiatry, Dx = diagnosis, sx = symptoms, SAB = Seasonal aff
BIP-NOS = bipolar disorder not otherwise specified, AMDP = Association Methodology and Documentation in Psychiatry, CID-S = Composite International Diagnostic Screener, FISC = Family Informant Schedule and Criteria, RDC = Research Diagnostic Criteria

References	Number	Citation
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	11	Sklar, P. et al. Whole-genome association study of bipolar disorder. Mol Psychiatry 13, 558-69 (2008).
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Table S3: Description of 9 schizophrenia cohorts

Cohort (References)	PGC label	Country	Ascertainment and Evaluation of Cases	Inclusion criteria (lifetime) of Cases	Exclusion criteria (lifetime) of Cases
Bonn/ Mannheim 1	boco	Germany	Consecutive hospital admissions, SCID, SADS-L, OPCRIT, medical records, FHx	DSM-IV SCZ	<u>.</u>
Bulgaria 2	butr	Bulgaria	Family trios where proband had SCZ/ SA, SCAN	DSM-IV SCZ / SA	Mental retardation
Denmark 1	denm	Denmark	Psychiatric departments and twin pair studies, OPCRIT	ICD-10 SCZ	Mania/bipolar illness
Molecular Genetics of Schizophrenia 3	mgs2	USA, Australia	Clinical settings and community residences, DIGS 2.0, FIGS 2.0, Medical records	DSM-IV SCZ / SA	
Munich 1	munc	Germany	Cases diagnosed with SCZ from the Munich area, SCID interview	DSM-IV SCZ	Head injury/ neurological diseases
Portugal 4	port	Portugal	Probands from families segregating SCZ, DIGS, SIS, SANS, SAPS, OPCRIT	DSM-IV SCZ	Bipolar disorder
Thematic Organized Psychosis Research	top8	Norway	Out-patient and in-patient psychiatric units, SCID-I interview	DSM-IV SCZ/ SA/ schizophreniform disorder	IQ score < 70
					Short-term drug-induced psychoses, psychoses with learning
UCLA 1	ucla	Netherlands	Inpatients and outpatients recruited through psychiatric hospitals and institutions, CASH	DSM-IV SCZ	disability/ head injury, other symptomatic psychoses
University College London 4	uclo	UK	SCZ diagnosis recorded in medical case-history, SADS-L, RDC	ICD-10 SCZ	SA, bipolar disorder, schizomania

Abbreviations: SCID = Structured Clinical Interview for DSM-IV, SADS = Schedule for Affective Disorders and Schizophrenia, OPCRIT = Operational Criteria Checklist, FHx = family history, SCZ = schizophrenia, SA = schizoaffective disorder,
SCAN = Schedules for Clinical Assessment in Neuropsychiatry, DIGS = Diagnostic Interview for Genetic Studies, FIGS = Family Interview for Genetic Studies, SIS = Kendler's Structured Interview for Schizotypy, SANS = Schedule for the Assessment of Negative Symptoms,
SAPS = Schedule for the Assessment of Positive Symptoms, CASH = Comprehensive Assessment of Symptoms and History, RDC = Research Diagnostic Criteria

References	Number	Citation
	1	Stefansson, H. et al. Common variants conferring risk of schizophrenia. Nature 460, 744-7 (2009).
	2	Kirov G. et al. De novo CNV analysis implicates specific abnormalities of postsynaptic signalling complexes in the pathogenesis of schizophrenia. Molecular Psychiatry 17, 142–153 (2012).
	3	Shi, J. et al. Common variants on chromosome 6p22.1 are associated with schizophrenia. Nature 460, 753-7 (2009).
	4	International Schizophrenia Consortium. Rare chromosomal deletions and duplications increase risk of schizophrenia. Nature 455, 237-41 (2008).
	5	Athanasiu, L. et al. Gene variants associated with schizophrenia in a Norwegian genome-wide study are replicated in a large European cohort. Journal of Psychiatric Research 44, 748-53 (2010).

Table 64. Reins on saledae nom psychiatric interviews	1	Ti and the second secon
Psychiatric Interview	Section/ Question	Information Collected
SCAN (Schedules for Clinical Assessment in Neuropsychiatry)	6.011 Suicide attempt and self-harm during episode of depression	0=absent, 1=deliberately considered suicide or self-injury but made no attempt, 2= injured self or made an attempt but no serious harm results, 3 = as 2 but with serious self-harm, 4 = made an attempt at suicide designed to result in death
SCID (Structured Clinical Interview for DSM-IV)	Sections on Depression, Mania, Mixed states	Recurrent thoughts of death (not just fear of dying), recurrent suicidal ideation without a specific plan, or a suicide attempt or a specific plan for committing suicide
SADS (Schedule for Affective Disorders and Schizophrenia)	Section O Suicidal Behavior	Ever made a suicide attempt, describe the most serious attempt, most serious attempt is rated by the interviewer in terms of intent and lethality
DIGS (Diagnostic Interview for Genetic Studies)	Section O Suicidal Behavior	Ever made a suicide attempt, describe the most serious attempt, medical treatment or hospitalisation required, whether the patient wanted to die or thought they would die, most serious attempt is rated by the interviewer in terms of intent and lethality
OPCRIT (Operational Criteria Checklist)	Past psychiatric history	Ever made suicide attempt
MINI (MINI International Neuropsychiatric Interview)	Section C Suicidality	Lifetime suicide attempt, in the past month thoughts about suicide, suicide plan, suicide attempt, hoped to survive or expected to die
CIDI (Composite International Diagnostic Interview)	Section on Major Depression	During worst two weeks in the last 12 months - thought about committing suicide, suicide plan, suicide attempt
SSAGA (Semi-Structured Assessment for the Genetics of Alcoholism)	Section I Depression, Section N Suicidal Behavior	Thoughts of death or suicide, sucide plan, suicide attempt, describe the most serious attempt, method, medical treatment, hospitalisation, whether the patient wanted to die or thought they would die, interviewer rates both the lethality and intent from unclear to extreme
FIGS (Family Interview for Genetic Studies)	During depression	Did the family member talk about death or suicide, try suicide
CASH (Comprehensive Assessment of Symptoms and History)	Major Depressive Syndrome	Thoughts about death and suicide, plus possible wishes to be dead, suicide plans, suicide attempts, rated from mild to severe

Table S5: Summary of polygenic risk scoring analyses with arrows showing direction from discovery to test dataset

Test for genetic overlap between psychiatric disorders and suicide attempt in the same disorder				
PGC BIP 1	\rightarrow	Suicide attempt vs non-attempt in BIP		
PGC Major Depression 2	\rightarrow	Suicide attempt vs non-attempt in MDD		
PGC SCZ 3	\rightarrow	Suicide attempt vs non-attempt in SCZ		
Test for genetic overlap between major depression and suicide attempt in other disorders				
PGC Major Depression 2	\rightarrow	Suicide attempt vs non-attempt in BIP		
PGC Major Depression 2	\rightarrow	Suicide attempt vs non-attempt in SCZ		
Test for genetic overlap in suicide attem	pt across	psychiatric disorders		
Suicide attempt vs non-attempt in BIP	\leftrightarrow	Suicide attempt vs non-attempt in MDD		
Suicide attempt vs non-attempt in BIP	\leftrightarrow	Suicide attempt vs non-attempt in SCZ		
Suicide attempt vs non-attempt in SCZ	\leftrightarrow	Suicide attempt vs non-attempt in MDD		
DCC Developing Committee Committee DID binder disorder MDD project demonstrate disorder CC7				

PGC, Psychiatric Genomics Consortium; BIP, bipolar disorder; MDD, major depressive disorder, SCZ, schizophrenia. 1. Stahl et al. Genomewide association study identifies 30 loci associated with bipolar disorder. bioRxiv, 2017. 2. Wray et al. Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. Nat Genet. 2018;50(5):668-81. 3. Schizophrenia Working Group of the Psychiatric Genomics Consortium. Biological insights from 108 schizophrenia-associated genetic loci. Nature. 2014;511(7510):421-7.

Table S6: Summary of suicide attempt in major depressive disorder cohorts st

Cohort	N Suicide attempters	N Non-attempters
CoFaMS	27	74
PsyCoLaus	65	442
GenRED2	168	653
GSK MPIP	115	763
MARS 650	137	407
MARS OMNIex	38	187
NTR/NESDA	229	1146
QIMR I317	61	521
QIMR 1610	32	263
QIMR COEX	48	299
RADIANT-UK	150	1424
RADIANT-Ger	35	276
STAR*D	126	807
BOMA	170	361
SHIP 0	18	348
GenRED1	203	815
Total	1622	8786

^{*}Individuals missing information on suicide attempt were excluded.

Table S7: Summary of suicide attempt in bipolar disorder cohorts*

Cohort	N Suicide attempters	N Non-attempters
BOMA-Germany I	241	365
Trinity College Dublin	26	26
FaST, TGEN1, TGEN2	120	124
French PGC2	185	254
BACCs	77	505
Mayo Clinic	307	610
Pritzker Neuropsychiatric Disorders Research		
Consortium	161	310
STEP2	170	363
STEP1	392	453
TOP7	116	207
TOP8	48	94
UCL	182	74
UMEA	74	124
WTCCC	423	673
GAIN	233	277
BOMA-Germany II	62	119
BOMA-Germany III	132	234
Boma Poland	150	251
Boma Spain	22	66
Nova Scotia Canada	78	223
BOMA Romania	65	148
Total	3264	5500

 $^{{\}bf *Individuals\ missing\ information\ on\ suicide\ attempt\ were\ excluded}.$

 ${\it Table S8: Summary of suicide attempt in schizophrenia cohorts*}$

Cohorts	N Suicide attempters	N Non-attempters
Bonn/ Mannheim	287	310
Bulgaria	103	208
Denmark	28	98
Molecular Genetics of Schizophrenia	754	1202
Munich	156	263
Portugal	80	243
Thematic Organized Psychosis Research Study	100	214
UCLA	86	304
University College London	89	104
Total	1683	2946

 $[\]hbox{*Individuals missing information on suicide attempt were excluded.}$

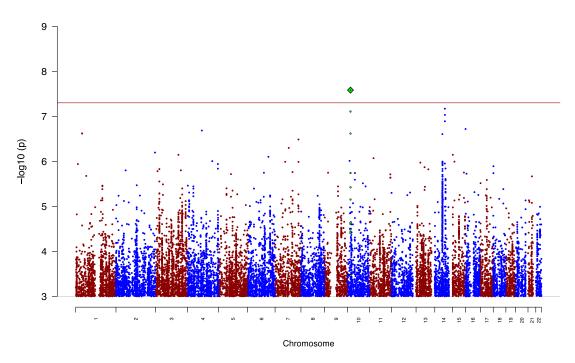


Figure S1: Manhattan plot from GWAS of suicide attempt in major depressive disorder

Table S9: Top 20 results from GWAS of suicide attempt in major depressive disorder showing the most significant SNP from each genomic region

				A1 freq	A1 freg non-			
Variant	CHR	ВР	A1/A2	attempters	attempters	P value	OR (C.I.)	Direction in each cohort
rs45593736	10	18954937	A/G	0.02	0.01	2.61E-08	2.38(1.75-3.23)	+++?++++-++?+-?+
rs111625585	14	82804332	T/C	0.08	0.06	6.75E-08	1.57(1.33-1.84)	+++++++-+
rs116428372	16	589359	A/G	0.07	0.06	1.91E-07	1.74(1.41-2.14)	++++++++++++++++++
rs77033326	4	89777618	A/G	0.03	0.02	2.07E-07	2.24(1.65-3.03)	+-++-+-+++++
rs183414028	1	40442026	T/C	0.98	0.99	2.39E-07	0.39(0.27-0.56)	
rs113330417	14	67249421	A/G	0.96	0.98	2.48E-07	0.57(0.46-0.71)	
rs111367251	7	144968289	C/G	0.98	0.99	3.26E-07	0.45(0.33-0.61)	-?-++
rs62460873	7	84977966	T/C	0.98	0.99	5.01E-07	0.37(0.25-0.54)	????++
rs186736781	2	240473090	T/C	0.03	0.02	6.37E-07	2.27(1.64-3.13)	+?+?-+-++
chr15_24344805_D	15	24344805	D/I3	0.37	0.41	7.14E-07	0.79(0.73-0.87)	
rs111326206	3	142438431	T/C	0.95	0.97	7.16E-07	0.62(0.51-0.75)	-++?+
chr6_128178230_I	6	128178230	12/D	0.08	0.06	7.92E-07	1.58(1.32-1.90)	
rs184924771	11	25885205	A/C	0.98	0.99	8.51E-07	0.39(0.27-0.57)	-?-?+?+-??-?
rs113386487	10	13358583	A/T	0.97	0.98	9.73E-07	0.47(0.35-0.64)	-?-??+
rs13137453	4	153907879	A/G	0.97	0.98	9.88E-07	0.46(0.34-0.63)	-?-?+?+
rs9972552	15	34396913	A/C	0.03	0.02	1.01E-06	2.18(1.59-2.98)	+-++-+++++++++
rs191852465	7	63164142	T/C	0.96	0.97	1.02E-06	0.46(0.34-0.63)	-++
chr13_46256859_D	13	46256859	D/I6	0.02	0.01	1.07E-06	2.27(1.63-3.16)	++?+?-++++++
rs145440507	4	188400537	A/T	0.98	0.99	1.15E-06	0.37(0.25-0.55)	-?-????-?-
rs76347430	1	14395819	A/G	0.97	0.98	1.15E-06	0.50(0.38-0.66)	?+

 $CHR, chromosome; BP, base pair position; freq, frequency; OR, odds \ ratio; CI, confidence interval$

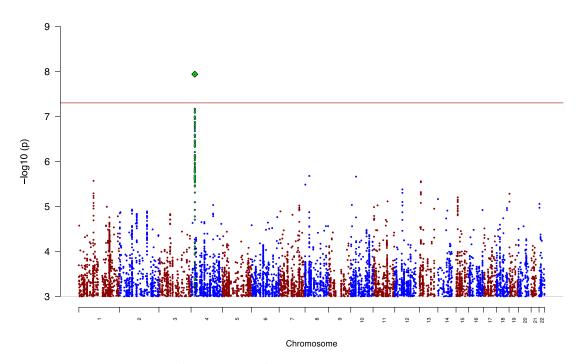


Figure S2: Manhattan plot from GWAS of suicide attempt in bipolar disorder

Table S10: Top 20 results from GWAS of suicide attempt in bipolar disorder showing the most significant SNP from each genomic region

Table S10: Top 20 rest	uits from Gi	was of suicide a	ttempt in bip	olar alsoraer sn	owing the most :	signijicant SNP	from each genomic region
				A1 freq	A1 freq non-		
Variant	CHR	BP	A1/A2	attempters	attempters	P value	OR (C.I.) Direction in each cohort
chr4_23273116_D	4	23273116	D/I10	0.20	0.17	1.15E-08	1.29(1.18-1.40) -+++++++++++++
rs1052873	8	27667793	T/C	0.19	0.22	2.10E-06	0.83(0.76-0.89)
rs118167891	10	32946009	T/C	0.02	0.02	2.17E-06	1.93(1.47-2.53) -+++++-?++-+
rs6428588	1	90825206	T/C	0.30	0.34	2.69E-06	0.85(0.79-0.91)+
rs7982251	13	28909835	T/C	0.85	0.87	2.76E-06	0.79(0.72-0.87)++-+-
rs67658161	8	3286733	A/C	0.48	0.44	3.27E-06	1.16(1.09-1.24) -+-++-++++++-+-
rs7979008	12	47479528	A/C	0.31	0.34	4.18E-06	0.85(0.79-0.91)+
rs55893662	19	2955759	T/C	0.11	0.10	5.22E-06	1.46(1.24-1.73)+++++++++
rs5016373	15	34314041	T/C	0.61	0.65	6.28E-06	0.86(0.80-0.92)++++
rs190572487	14	19675351	T/C	0.75	0.73	6.86E-06	1.35(1.18-1.54) -++++++++++++++
rs3847511	11	91454034	T/G	0.08	0.06	7.70E-06	1.36(1.19-1.55) -++++++++
rs165774	22	19952561	A/G	0.33	0.30	8.78E-06	1.18(1.10-1.27) -+++-+-+-+-+-
chr10_7228436_I	10	7228436	12/D	0.14	0.12	9.25E-06	1.31(1.16-1.47)+++
rs12639760	4	135958271	A/T	0.22	0.19	9.31E-06	1.21(1.11-1.31) -++-+++++++++++
rs141199126	11	29761274	T/C	0.02	0.01	9.43E-06	2.05(1.49-2.83) -++++++?+?+-?++
chr7_123745464_I	7	123745464	12/D	0.08	0.07	9.54E-06	1.32(1.17-1.49)
chr1_172543621_I	1	172543621	12/D	0.03	0.04	1.02E-05	0.60(0.47-0.75) -+++++++++++++
rs12799429	11	8097070	T/C	0.54	0.56	1.04E-05	0.86(0.80-0.92)+++-
rs17077064	18	64976306	T/C	0.83	0.80	1.08E-05	1.21(1.11-1.31) -++++++++++++++
rs117018753	13	110581806	T/C	0.07	0.06	1.12E-05	1.37(1.19-1.58) -+++++++++++++++

CHR, chromosome; BP, basepair position; freq, frequency; OR, odds ratio; CI, confidence interval

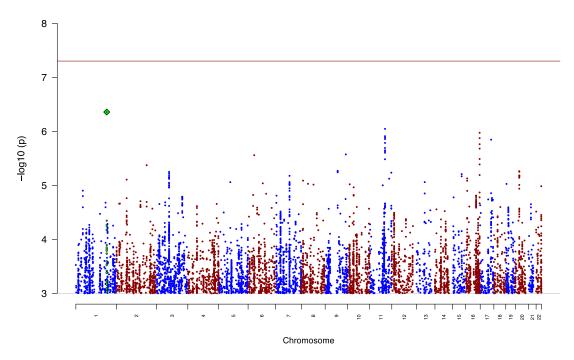


Figure S3: Manhattan plot from GWAS of suicide attempt in schizophrenia

 $Table\,S11: Top\,20\,results\,from\,GWAS\,of\,suicide\,attempt\,in\,schizophrenia\,showing\,the\,most\,significant\,SNP\,from\,each\,genomic\,region$

				A1 freq	A1 freq non-		
Variant	CHR	ВР	A1/A2	attempters	attempters	P value	OR (C.I.) Direction in each cohort
rs482039	1	190777567	T/C	0.03	0.02	4.39E-07	2.37(1.70-3.31) +++-++++
rs3858375	11	95077167	T/C	0.07	0.05	8.99E-07	1.60(1.33-1.94) ++++-+++
rs4843180	16	86753070	T/C	0.44	0.39	1.06E-06	1.25(1.14-1.36) ++++++++
rs180697792	17	67294430	A/G	0.23	0.20	1.43E-06	1.34(1.19-1.50) ++++++++
rs72756712	9	128902906	A/G	0.93	0.95	2.67E-06	0.60(0.49-0.74)
rs191312301	6	38975727	A/C	0.86	0.89	2.76E-06	0.71(0.62-0.82)+
chr2_188481671_D	2	188481671	D/I3	0.34	0.38	4.23E-06	0.79(0.71-0.87)
rs73650494	9	78831443	T/C	0.96	0.98	5.39E-06	0.56(0.43-0.72)
rs6114731	20	24427180	A/G	0.04	0.03	5.46E-06	1.78(1.39-2.27) ++++++++
rs75305337	3	84333133	A/C	0.12	0.09	5.63E-06	1.39(1.20-1.60) ++++++++
chr11_133764404_I	11	133764404	15/D	0.79	0.75	5.82E-06	1.29(1.15-1.43)+
rs57729539	15	78524199	A/G	0.80	0.83	6.18E-06	0.76(0.68-0.86)+
rs875777	7	86745380	T/C	0.16	0.20	6.66E-06	0.76(0.67-0.86)+
rs6497871	16	10364163	A/G	0.64	0.60	7.44E-06	1.24(1.13-1.37) +++++++
rs151336980	11	120470737	T/C	0.04	0.03	7.52E-06	1.78(1.38-2.29) ++++-++
rs4494728	2	65589513	T/C	0.50	0.55	7.82E-06	0.82(0.75-0.89)
rs2739958	8	12232534	T/C	0.58	0.60	8.18E-06	0.68(0.57-0.80)++-
rs73215273	13	72403250	A/C	0.88	0.90	8.71E-06	0.66(0.54-0.79) -+
rs11739808	5	72745041	A/G	0.03	0.02	8.72E-06	2.30(1.59-3.31) ++-+++++
rs13198361	6	91673413	T/C	0.11	0.14	9.17E-06	0.65(0.54-0.79)+

 $CHR, chromosome; BP, base pair position; freq, frequency; OR, odds \, ratio; CI, confidence \, interval$

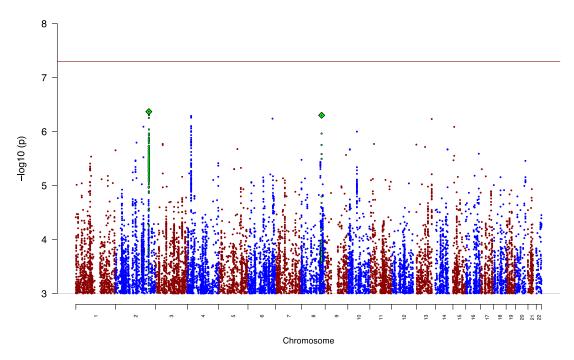


Figure S4: Manhattan plot from meta-analysis of suicide attempt in major depressive disorder, bipolar disorder and schizophrenia

Table S12: Top 20 results from meta-analysis of suicide attempt in MDD, BIP and SCZ showing the most significant SNP from each genomic region

				A1 freq	A1 freq non-		
Variant	CHR	ВР	A1/A2	attempters	attempters	P value	OR (C.I.) Direction in each cohort
rs149268645	2	203833018	A/G	0.14	0.15	4.28E-07	0.85(0.80-0.90)
rs4870888	8	125108977	T/C	0.52	0.54	5.00E-07	0.89(0.86-0.93)
chr4_23273116_D	4	23273116	D/I	0.19	0.18	5.13E-07	1.16(1.09-1.23) +++
rs141252918	6	151828058	A/G	0.02	0.01	5.76E-07	1.73(1.39-2.14) ++?
rs9577511	13	113991823	A/G	0.86	0.87	5.86E-07	0.83(0.77-0.89)
rs62173322	2	170611029	A/G	0.86	0.87	8.13E-07	0.84(0.78-0.90)
rs76371172	15	31814455	T/G	0.98	0.98	8.20E-07	0.61(0.50-0.74)
rs11004733	10	56849344	T/C	0.04	0.04	1.00E-06	1.34(1.19-1.50) +++
rs138689899	2	128288162	T/C	0.02	0.02	1.61E-06	1.53(1.29-1.82) ++-
rs142055939	3	45995554	T/C	0.02	0.02	1.69E-06	1.48(1.26-1.75) +++
rs35107435	11	27249330	A/T	0.16	0.15	1.70E-06	1.17(1.10-1.24) +++
rs113988902	13	19525105	T/C	0.05	0.04	1.74E-06	1.52(1.28-1.80) ?++
chr13_73243177_I	13	73243177	I/D	0.04	0.03	1.93E-06	1.59(1.32-1.93) ?
rs186672572	5	116878032	T/C	0.02	0.01	2.11E-06	1.64(1.33-2.00) +++
rs113386487	10	13358583	A/T	0.98	0.98	2.12E-06	0.67(0.56-0.79)
rs73348245	14	96158072	A/G	0.06	0.06	2.13E-06	1.27(1.15-1.40) +++
rs6426297	1	246538381	T/C	0.02	0.01	2.22E-06	1.65(1.34-2.04) +++
rs73577700	16	80280761	A/T	0.83	0.85	2.57E-06	0.87(0.82-0.92)
rs72756712	9	128902906	A/G	0.94	0.95	2.71E-06	0.78(0.70-0.86)
rs75633108	1	96735185	T/C	0.02	0.02	2.91E-06	1.59(1.31-1.93) +++

MDD, major depressive disorder; BIP, bipolar disorder; SCZ, schizophrenia; CHR, chromosome; BP, basepair position; OR, odds ratio; because of the control of the control

CI, confidence interval

 $Table \, S13: Top \, 20 \, results \, from \, meta-analysis \, of \, suicide \, at tempt \, in \, mood \, disorders \, showing \, the \, most \, significant \, SNP \, from \, each \, genomic \, region \, and \, from \, each \, genomic \, region \, from \,$

				A1 freq	A1 freq			
Variant	CHR	ВР	A1/A2		non-attempters		OR (C.I.)	Direction in each cohort
rs138689899	2	128288162	T/C	0.02				
rs28591567	4	23253912	A/G	0.78	0.80	3.11E-08	0.84(0.79-0.89)	
chr6_151835609_D	6	151835609	I/D	0.94	0.95	3.66E-07	0.74(0.66-0.83)	++
chr13_61834504_D	13	61834504	I/D	0.91	0.92	5.77E-07	0.78(0.71-0.86)	++
rs186672572	5	116878032	T/C	0.02	0.01	7.19E-07	1.78(1.42-2.24)	++
rs112944737	8	134677667	T/C	0.91	0.92	1.11E-06	0.80(0.73-0.87)	
rs9577511	13	113991823	A/G	0.86	0.87	1.35E-06	0.81(0.75-0.88)	
rs150795632	6	37439376	A/G	0.98	0.98	1.69E-06	0.59(0.48-0.73)	
rs113051785	11	20167807	C/G	0.02	0.01	1.70E-06	1.75(1.39-2.20)	++
rs1355048	1	90830490	T/C	0.37	0.40	1.91E-06	0.88(0.84-0.93)	
rs115833694	4	99918226	C/G	0.98	0.98	1.99E-06	0.63(0.52-0.76)	
rs17764923	6	159820779	A/G	0.16	0.14	2.04E-06	1.19(1.11-1.27)	++
rs117020391	12	107099751	T/C	0.98	0.98	2.09E-06	0.63(0.52-0.76)	
rs72832403	2	115551269	A/G	0.92	0.93	2.16E-06	0.79(0.72-0.87)	
rs150320200	4	23450330	A/G	0.97	0.98	2.19E-06	0.66(0.55-0.78)	
rs114598476	1	224438315	A/G	0.96	0.97	2.30E-06	0.67(0.57-0.79)	
rs76400344	22	26697120	A/C	0.03	0.02	3.01E-06	1.52(1.28-1.82)	++
chr2_124851208_D	2	124851208	I/D	0.96	0.96	3.48E-06	0.72(0.62-0.82)	++
rs143457262	7	82072544	A/G	0.03	0.03	4.16E-06	1.51(1.26-1.79)	++
rs3829881	1	117531813	T/C	0.49	0.47	4.45E-06	1.12(1.07-1.18)	++

CHR, chromosome; BP, basepair position; freq, frequency; OR, odds ratio; CI, confidence interval

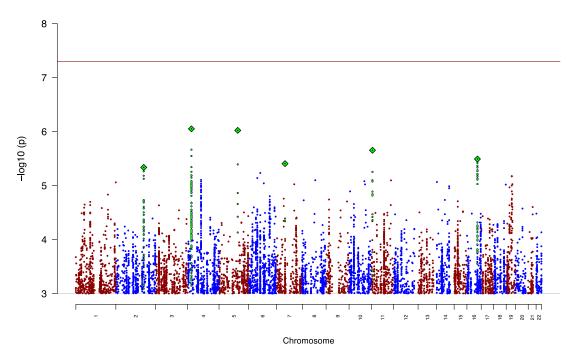


Figure S5: Manhattan plot from meta-analysis of suicide attempt in bipolar disorder and schizophrenia

 $Table \, S14: Top \, 20 \, results \, from \, meta-analysis \, of \, suicide \, attempt \, in \, BIP \, and \, SCZ \, showing \, the \, most \, significant \, SNP \, from \, each \, genomic \, region \, and \, SCZ \, showing \, the \, most \, significant \, SNP \, from \, each \, genomic \, region \, and \, SCZ \, showing \, the \, most \, significant \, SNP \, from \, each \, genomic \, region \, and \, SCZ \, showing \, the \, most \, significant \, SNP \, from \, each \, genomic \, region \, and \, SCZ \, showing \, the \, most \, significant \, SNP \, from \, each \, genomic \, region \, and \, SCZ \, showing \, the \, most \, significant \, SNP \, from \, each \, genomic \, region \, and \, SCZ \, showing \, the \, most \, significant \, SNP \, from \, each \, genomic \, region \, and \, SCZ \, showing \, the \, subject \, and \, SCZ \, showing \, the \, subject \, and \, SCZ \, showing \, subject \, and \, SCZ \,$

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				A1 freq	A1 freq		
Variant	CHR	ВР	A1/A2	attempters	non-attempters	P value	OR (C.I.) Direction in each cohort
chr4_23273116_D	4	23273116	D/I	0.19	0.17	8.90E-07	1.19(1.11-1.27) ++
rs26318	5	115687905	T/C	0.98	0.99	9.50E-07	0.52(0.40-0.67)
rs118102650	11	9099847	A/G	0.02	0.01	2.22E-06	1.79(1.41-2.28) ++
rs12925656	16	65107920	T/C	0.88	0.90	3.22E-06	0.82(0.76-0.89)
rs73122740	7	53695512	T/G	0.92	0.91	3.93E-06	1.26(1.14-1.39) ++
rs2353181	2	170305256	T/C	0.95	0.94	4.61E-06	1.33(1.17-1.49) ++
rs75237141	6	72966596	A/G	0.90	0.92	5.84E-06	0.80(0.73-0.88)
rs56342621	19	35071781	C/G	0.98	0.99	6.67E-06	0.61(0.49-0.75)
rs9475195	6	55061018	T/C	0.58	0.61	7.25E-06	0.89(0.84-0.93)
chr4_82106147_I	4	82106147	I/D	0.07	0.06	7.85E-06	1.29(1.15-1.44)
chr8_80591790_D	8	80591790	I/D	0.61	0.63	7.95E-06	0.88(0.83-0.93) ++
rs10892827	11	122210990	T/G	0.90	0.92	8.01E-06	0.80(0.73-0.88)
rs189924441	10	94182243	A/G	0.97	0.97	8.30E-06	0.67(0.56-0.80)
rs8022689	14	21608986	A/G	0.38	0.35	8.58E-06	1.14(1.07-1.20) ++
rs3007305	1	246862572	C/G	0.66	0.68	8.72E-06	0.88(0.83-0.93)
rs72924216	6	94570858	A/G	0.95	0.96	9.08E-06	0.72(0.63-0.83)
rs66666015	19	38717143	T/C	0.76	0.74	9.42E-06	1.16(1.09-1.24) ++
rs38758	7	109943767	A/C	0.47	0.44	9.46E-06	1.12(1.07-1.18) ++
rs117559494	10	98008526	A/G	0.03	0.03	9.50E-06	1.53(1.27-1.85) ++
rs117637007	18	75827605	T/C	0.11	0.13	9.58E-06	0.80(0.72-0.88)

CHR, chromosome; BP, basepair position; freq, frequency; OR, odds ratio; CI, confidence interval

Table S15: Genome-wide significant loci for suicide attempt tested in independent replication cohorts

Cohort	Variant	CHR	A1/A2	P value	OR (C.I.)
UK Biobank	rs45593736	10	A/G	0.985	1.00 (0.83-1.21)
UK Biobank	rs138689899	2	T/C	0.677	0.96 (0.79-1.16)
UK Biobank	rs4626184^	4	C/A	0.663	1.04 (0.86-1.26)
iPSYCH	rs117607218*	10	C/T	0.776	0.97 (0.79-1.18)
iPSYCH	rs138689899	2	T/C	0.228	0.87 (0.69-1.09)
iPSYCH	rs28591567	4	G/A	0.521	1.02 (0.96-1.07)

[^]rs28591567 was not present in the UK Biobank dataset, so the SNP in highest LD ($R^2 = 0.88$) was chosen.

CHR, chromosome; OR, odds ratio; CI, confidence interval

^{*}rs45593736 was not present in the iPSYCH dataset, so the SNP in highest LD ($R^2 = 0.60$) was chosen.

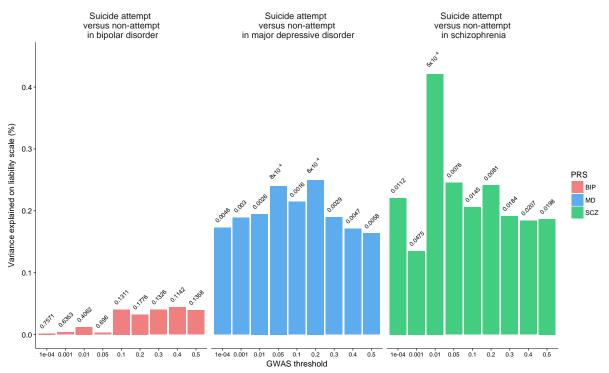


Figure S6: Polygenic risk scores for bipolar disorder, major depression and schizophrenia tested for association with suicide attempt in the same disorder.

PRS-polygenic risk score, BIP-bipolar disorder, MD-major depression, SCZ-schizophrenia. P values of association between polygenic scores and suicide attempt are shown above each bar.

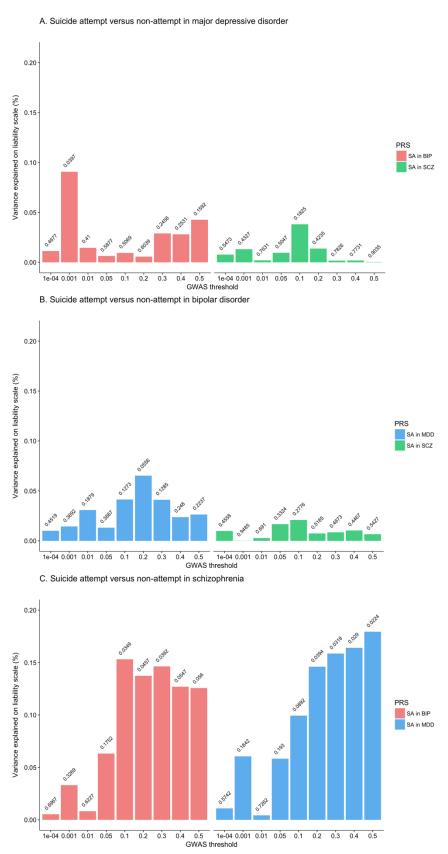


Figure S7: Polygenic risk scores for suicide attempt tested for association with suicide attempt in A - major depressive disorder, B - bipolar disorder and C - schizophrenia. PRS-polygenic risk score, SA-suicide attempt, MDD-major depressive disorder, BIP-bipolar disorder, SCZ-schizophrenia. P values of association between polygenic scores and suicide attempt are shown above each bar.

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Some data used in this study were obtained from dbGaP. Funding support for the Genome-Wide Association of Schizophrenia Study was provided by the National Institute of Mental Health (R01 MH67257, R01 MH59588, R01 MH59571, R01 MH59565, R01 MH59587, R01 MH60870, R01 MH59566, R01 MH59586, R01 MH61675, R01 MH60879, R01 MH81800, U01 MH46276, U01 MH46289 U01 MH46318, U01 MH79469, and U01 MH79470) and the genotyping of samples was provided through the Genetic Association Information Network (GAIN). The datasets used for the analyses described in this manuscript were obtained from the database of Genotypes and Phenotypes (dbGaP) found at http://www.ncbi.nlm.nih.gov/gap through dbGaP accession number phs000021.v3.p2. Samples and associated phenotype data for the Genome-Wide Association of Schizophrenia Study were provided by the Molecular Genetics of Schizophrenia Collaboration (PI: Pablo V. Gejman, Evanston Northwestern Healthcare (ENH) and Northwestern University, Evanston, IL, USA)." Funding support for the Whole Genome Association Study of Bipolar Disorder was provided by the National Institute of Mental Health (NIMH) and the genotyping of samples was provided through the Genetic Association Information Network (GAIN). The datasets used for the analyses described in this manuscript were obtained from the database of Genotypes and Phenotypes (dbGaP) found at http://www.ncbi.nlm.nih.gov/gap through dbGaP accession number phs000017.v3.p1. Samples and associated phenotype data for the Collaborative Genomic Study of Bipolar Disorder were provided by the The NIMH Genetics Initiative for Bipolar Disorder. Data and biomaterials were collected in four projects that participated in NIMH Bipolar Disorder Genetics Initiative. From 1991-98, the Principal Investigators and Co-Investigators were: Indiana University, Indianapolis, IN, U01 MH46282, John Nurnberger, M.D., Ph.D., Marvin Miller, M.D., and Elizabeth Bowman, M.D.; Washington University, St. Louis, MO, U01 MH46280, Theodore Reich, M.D., Allison Goate, Ph.D., and John Rice, Ph.D.; Johns Hopkins University, Baltimore, MD U01 MH46274, J. Raymond DePaulo, Jr., M.D., Sylvia Simpson, M.D., MPH, and Colin Stine, Ph.D.; NIMH Intramural Research Program, Clinical Neurogenetics Branch, Bethesda, MD, Elliot Gershon, M.D., Diane Kazuba, B.A., and Elizabeth Maxwell, M.S.W. Data and biomaterials were collected as part of ten projects that participated in the NIMH Bipolar Disorder Genetics Initiative. From 1999-03, the Principal Investigators and Co-Investigators were: Indiana University, Indianapolis, IN, R01 MH59545, John Nurnberger, M.D., Ph.D., Marvin J. Miller, M.D., Elizabeth S. Bowman, M.D., N. Leela Rau, M.D., P. Ryan Moe, M.D., Nalini Samavedy, M.D., Rif El-Mallakh, M.D. (at University of Louisville), Husseini Manji, M.D. (at Wayne State University), Debra A. Glitz, M.D. (at Wayne State University), Eric T. Meyer, M.S., Carrie Smiley, R.N., Tatiana Foroud, Ph.D., Leah Flury, M.S., Danielle M. Dick, Ph.D., Howard Edenberg, Ph.D.; Washington University, St. Louis, MO, R01 MH059534, John Rice, Ph.D, Theodore Reich, M.D., Allison Goate, Ph.D., Laura Bierut, M.D.; Johns Hopkins University, Baltimore, MD, R01 MH59533, Melvin McInnis M.D., J. Raymond DePaulo, Jr., M.D., Dean F. MacKinnon, M.D., Francis M. Mondimore, M.D., James B. Potash, M.D., Peter P. Zandi, Ph.D, Dimitrios Avramopoulos, and Jennifer Payne; University of Pennsylvania, PA, R01 MH59553, Wade Berrettini M.D., Ph.D.; University of California at Irvine, CA, R01 MH60068, William Byerley M.D., and Mark Vawter M.D.; University of Iowa, IA, R01 MH059548, William Coryell M.D., and Raymond Crowe M.D.; University of Chicago, IL, R01 MH59535, Elliot Gershon, M.D., Judith Badner Ph.D., Francis McMahon M.D., Chunyu Liu Ph.D., Alan Sanders M.D., Maria Caserta, Steven Dinwiddie M.D., Tu Nguyen, Donna Harakal; University of California at San Diego, CA, R01 MH59567, John Kelsoe, M.D., Rebecca McKinney, B.A.; Rush University, IL, R01 MH059556, William Scheftner M.D., Howard M. Kravitz, D.O., M.P.H., Diana Marta, B.S., Annette Vaughn-Brown, MSN, RN, and Laurie Bederow, MA; NIMH Intramural Research Program, Bethesda, MD, 1Z01MH002810-01, Francis J. McMahon, M.D., Layla Kassem, PsyD, Sevilla Detera-Wadleigh, Ph.D, Lisa Austin, Ph.D, Dennis L. Murphy, M.D.

Psychiatric Genomics Consortium Major Depressive Disorder Cohorts

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BoMa Rietschel, S Cichon 01ZX1314G/01ZX1614G, BMBF Integrament Germa BoMa MM Nöthen, M Rietschel, S Cichon 01GS08144, 01GS08147 BMBF NGFNplus MooDS Germa	F	ImmunoSensation	Forschungsgemeinschaft	Germany	
BoMa MN Nöthen, M Rietschel, S Cichon 012X1314G/012X1614G, 01GS08144, 01GS08147 BMBF NGFNplus MooDS Germa	l _B	' '	RMRF Integrament	Germany	
Rietschel, S Cichon 01GS08144, 01GS08147 BMBF NGFNplus MooDS Germa	Ľ	01ZX1314G/01ZX1614G,	Dividi integrament	Cermany	
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Psychiatric Genomics Consortium Bipolar Disorder Cohorts

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Psychiatric Genomics Consortium Schizophrenia cohorts

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Replication cohorts

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Consortium Authorship

Major Depressive Disorder Working Group of the Psychiatric Genomics Consortium

Naomi R Wray* 1, 2 Stephan Ripke* 3, 4, 5 Manuel Mattheisen* 6, 7, 8, 9 Maciej Trzaskowski* 1 Enda M Byrne 1 Abdel Abdellaoui 10 Mark J Adams 11 Esben Agerbo 9, 12, 13 Tracy M Air 14

Till F M Andlauer 15, 16
Silviu-Alin Bacanu 17
Marie Bækvad-Hansen 9, 18
Aartjan T F Beekman 19
Tim B Bigdeli 17, 20
Elisabeth B Binder 15, 21
Douglas H R Blackwood 11

Julien Bryois 22

Henriette N Buttenschøn 8, 9, 23 Jonas Bybjerg-Grauholm 9, 18

Na Cai 24, 25 Enrique Castelao 26

Jane Hvarregaard Christensen 7,

8,9

Toni-Kim Clarke 11 Jonathan R I Coleman 27 Lucía Colodro-Conde 28 Baptiste Couvy-Duchesne 2, 29

Nick Craddock 30

Gregory E Crawford 31, 32

Gail Davies 33 Ian J Deary 33

Franziska Degenhardt 34, 35

Eske M Derks 28 Nese Direk 36, 37 Conor V Dolan 10 Erin C Dunn 38, 39, 40 Thalia C Eley 27

Valentina Escott-Price 41 Farnush Farhadi Hassan Kiadeh

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Hilary K Finucane 43, 44

Jerome C Foo 45

Andreas J Forstner 34, 35, 46, 47

Josef Frank 45 Héléna A Gaspar 27 Michael Gill 48 Fernando S Goes 49 Scott D Gordon 28 Jakob Grove 7, 8, 9, 50 Lynsey S Hall 11, 51

Christine Søholm Hansen 9, 18 Thomas F Hansen 52, 53, 54 Stefan Herms 34, 35, 47

Ian B Hickie 55 Per Hoffmann 34, 35, 47 Georg Homuth 56 Carsten Horn 57

Jouke-Jan Hottenga 10
David M Hougaard 9, 18
David M Howard 11
Marcus Ising 58
Rick Jansen 19
Ian Jones 59
Lisa A Jones 60
Eric Jorgenson 61
James A Knowles 62
Isaac S Kohane 63, 64, 65

Julia Kraft 4

Warren W. Kretzschmar 66

Jesper Krogh 67 Zoltán Kutalik 68, 69

Yihan Li 66

Penelope A Lind 28
Donald J MacIntyre 70, 71
Dean F MacKinnon 49
Robert M Maier 2
Wolfgang Maier 72
Jonathan Marchini 73
Hamdi Mbarek 10
Patrick McGrath 74
Peter McGuffin 27
Sarah E Medland 28

Christel M Middeldorp 10, 76, 77

Evelin Mihailov 78 Yuri Milaneschi 19 Lili Milani 78

Divya Mehta 2, 75

Francis M Mondimore 49
Grant W Montgomery 1
Sara Mostafavi 79, 80
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Matthias Nauck 81, 82
Bernard Ng 80
Michel G Nivard 10
Dale R Nyholt 83
Paul F O'Reilly 27
Hogni Oskarsson 84
Michael J Owen 59

Carsten Bøcker Pedersen 9, 12, 13 Marianne Giørtz Pedersen 9, 12,

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Roseann E. Peterson 17, 85

Erik Pettersson 22

Jodie N Painter 28

Wouter J Peyrot 19 Giorgio Pistis 26

Danielle Posthuma 86, 87

Jorge A Quiroz 88
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Brien P. Riley 17
Margarita Rivera 27, 90
Saira Saeed Mirza 36
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Stacy Steinberg 99
Fabian Streit 45
Jana Strohmaier 45
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Jordan W Smoller 38, 39, 40

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Henry Völzke 102

Myrna M Weissman 74, 141

Thomas Werge 9, 53, 142

Cathryn M Lewis 27, 143

Douglas F Levinson 144

Gerome Breen 27, 145

Anders D Børglum 7, 8, 9

Patrick F Sullivan 22, 146, 147

- 1, Institute for Molecular Bioscience, The University of Queensland, Brisbane, QLD, AU
- 2, Queensland Brain Institute, The University of Queensland, Brisbane, QLD, AU
- 3, Analytic and Translational Genetics Unit, Massachusetts General Hospital, Boston, MA, US
- 4, Department of Psychiatry and Psychotherapy, Universitätsmedizin Berlin Campus Charité Mitte, Berlin, DE
- 5, Medical and Population Genetics, Broad Institute, Cambridge, MA, US
- 6, Centre for Psychiatry Research, Department of Clinical Neuroscience, Karolinska Institutet, Stockholm, SE
- 7, Department of Biomedicine, Aarhus University, Aarhus, DK
- 8, iSEQ, Centre for Integrative Sequencing, Aarhus University, Aarhus, DK
- 9, iPSYCH, The Lundbeck Foundation Initiative for Integrative Psychiatric Research, DK
- 10, Dept of Biological Psychology & EMGO+ Institute for Health and Care Research, Vrije Universiteit Amsterdam, Amsterdam, NL
- 11, Division of Psychiatry, University of Edinburgh, Edinburgh, GB
- 12, Centre for Integrated Register-based Research, Aarhus University, Aarhus, DK
- 13, National Centre for Register-Based Research, Aarhus University, Aarhus, DK
- 14, Discipline of Psychiatry, University of Adelaide, Adelaide, SA, AU
- 15, Department of Translational Research in Psychiatry, Max Planck Institute of Psychiatry, Munich, DE
- 16, Munich Cluster for Systems Neurology (SyNergy), Munich, DE
- 17, Department of Psychiatry, Virginia Commonwealth University, Richmond, VA, US
- 18, Center for Neonatal Screening, Department for Congenital Disorders, Statens Serum Institut, Copenhagen, DK
- 19, Department of Psychiatry, Vrije Universiteit Medical Center and GGZ inGeest, Amsterdam, NL
- 20, Virginia Institute for Psychiatric and Behavior Genetics, Richmond, VA, US
- 21, Department of Psychiatry and Behavioral Sciences, Emory University School of Medicine, Atlanta, GA, US
- 22, Department of Medical Epidemiology and Biostatistics, Karolinska Institutet, Stockholm, SE
- 23, Department of Clinical Medicine, Translational Neuropsychiatry Unit, Aarhus University, Aarhus, DK
- 24, Human Genetics, Wellcome Trust Sanger Institute, Cambridge, GB
- 25, Statistical genomics and systems genetics, European Bioinformatics Institute (EMBL-EBI), Cambridge, GB
- 26, Department of Psychiatry, University Hospital of Lausanne, Prilly, Vaud, CH
- 27, Social Genetic and Developmental Psychiatry Centre, King's College London, London, GB
- 28, Genetics and Computational Biology, QIMR Berghofer Medical Research Institute, Brisbane, QLD, AU
- 29, Centre for Advanced Imaging, The University of Queensland, Brisbane, QLD, AU
- 30, Psychological Medicine, Cardiff University, Cardiff, GB
- 31, Center for Genomic and Computational Biology, Duke University, Durham, NC, US
- 32, Department of Pediatrics, Division of Medical Genetics, Duke University, Durham, NC, US
- 33, Centre for Cognitive Ageing and Cognitive Epidemiology, University of Edinburgh, Edinburgh, GB
- 34, Institute of Human Genetics, University of Bonn, Bonn, DE
- 35, Life&Brain Center, Department of Genomics, University of Bonn, Bonn, DE
- 36, Epidemiology, Erasmus MC, Rotterdam, Zuid-Holland, NL
- 37, Psychiatry, Dokuz Eylul University School Of Medicine, Izmir, TR
- 38, Department of Psychiatry, Massachusetts General Hospital, Boston, MA, US
- 39, Psychiatric and Neurodevelopmental Genetics Unit (PNGU), Massachusetts General Hospital, Boston, MA, US
- 40, Stanley Center for Psychiatric Research, Broad Institute, Cambridge, MA, US
- 41, Neuroscience and Mental Health, Cardiff University, Cardiff, GB
- 42, Bioinformatics, University of British Columbia, Vancouver, BC, CA
- 43, Department of Epidemiology, Harvard T.H. Chan School of Public Health, Boston, MA, US
- 44, Department of Mathematics, Massachusetts Institute of Technology, Cambridge, MA, US
- 45, Department of Genetic Epidemiology in Psychiatry, Central Institute of Mental Health, Medical Faculty
- Mannheim, Heidelberg University, Mannheim, Baden-Württemberg, DE
- 46, Department of Psychiatry (UPK), University of Basel, Basel, CH
- 47, Human Genomics Research Group, Department of Biomedicine, University of Basel, Basel, CH
- 48, Department of Psychiatry, Trinity College Dublin, Dublin, IE
- 49, Psychiatry & Behavioral Sciences, Johns Hopkins University, Baltimore, MD, US
- 50, Bioinformatics Research Centre, Aarhus University, Aarhus, DK
- 51, Institute of Genetic Medicine, Newcastle University, Newcastle upon Tyne, GB
- 52, Danish Headache Centre, Department of Neurology, Rigshospitalet, Glostrup, DK

- 53, Institute of Biological Psychiatry, Mental Health Center Sct. Hans, Mental Health Services Capital Region of Denmark, Copenhagen, DK
- 54, iPSYCH, The Lundbeck Foundation Initiative for Psychiatric Research, Copenhagen, DK
- 55, Brain and Mind Centre, University of Sydney, Sydney, NSW, AU
- 56, Interfaculty Institute for Genetics and Functional Genomics, Department of Functional Genomics,
- University Medicine and Ernst Moritz Arndt University Greifswald, Greifswald, Mecklenburg-Vorpommern, DE
- 57, Roche Pharmaceutical Research and Early Development, Pharmaceutical Sciences, Roche Innovation Center Basel, F. Hoffmann-La Roche Ltd, Basel, CH
- 58, Max Planck Institute of Psychiatry, Munich, DE
- 59, MRC Centre for Neuropsychiatric Genetics and Genomics, Cardiff University, Cardiff, GB
- 60, Department of Psychological Medicine, University of Worcester, Worcester, GB
- 61, Division of Research, Kaiser Permanente Northern California, Oakland, CA, US
- 62, Psychiatry & The Behavioral Sciences, University of Southern California, Los Angeles, CA, US
- 63, Department of Biomedical Informatics, Harvard Medical School, Boston, MA, US
- 64, Department of Medicine, Brigham and Women's Hospital, Boston, MA, US
- 65, Informatics Program, Boston Children's Hospital, Boston, MA, US
- 66, Wellcome Trust Centre for Human Genetics, University of Oxford, Oxford, GB
- 67, Department of Endocrinology at Herlev University Hospital, University of Copenhagen, Copenhagen, DK
- 68, Institute of Social and Preventive Medicine (IUMSP), University Hospital of Lausanne, Lausanne, VD, CH
- 69, Swiss Institute of Bioinformatics, Lausanne, VD, CH
- 70, Division of Psychiatry, Centre for Clinical Brain Sciences, University of Edinburgh, Edinburgh, GB
- 71, Mental Health, NHS 24, Glasgow, GB
- 72, Department of Psychiatry and Psychotherapy, University of Bonn, Bonn, DE
- 73, Statistics, University of Oxford, Oxford, GB
- 74, Psychiatry, Columbia University College of Physicians and Surgeons, New York, NY, US
- 75, School of Psychology and Counseling, Queensland University of Technology, Brisbane, QLD, AU
- 76, Child and Youth Mental Health Service, Children's Health Queensland Hospital and Health Service, South Brisbane, QLD, AU
- 77, Child Health Research Centre, University of Queensland, Brisbane, QLD, AU
- 78, Estonian Genome Center, University of Tartu, Tartu, EE
- 79, Medical Genetics, University of British Columbia, Vancouver, BC, CA
- 80, Statistics, University of British Columbia, Vancouver, BC, CA
- 81, DZHK (German Centre for Cardiovascular Research), Partner Site Greifswald, University Medicine, University Medicine Greifswald, Greifswald, Mecklenburg-Vorpommern, DE
- 82, Institute of Clinical Chemistry and Laboratory Medicine, University Medicine Greifswald, Greifswald, Mecklenburg-Vorpommern, DE
- 83, Institute of Health and Biomedical Innovation, Queensland University of Technology, Brisbane, QLD, AU
- 84, Humus, Reykjavik, IS
- 85, Virginia Institute for Psychiatric & Behavioral Genetics, Virginia Commonwealth University, Richmond, VA, US
- 86, Clinical Genetics, Vrije Universiteit Medical Center, Amsterdam, NL
- 87, Complex Trait Genetics, Vrije Universiteit Amsterdam, Amsterdam, NL
- 88, Solid Biosciences, Boston, MA, US
- 89, Department of Psychiatry, Washington University in Saint Louis School of Medicine, Saint Louis, MO, US
- 90, Department of Biochemistry and Molecular Biology II, Institute of Neurosciences, Center for Biomedical Research, University of Granada, Granada, ES
- 91, Department of Psychiatry, University of Groningen, University Medical Center Groningen, Groningen, NL
- 92, Department of Psychiatry and Psychotherapy, Medical Center of the University of Munich, Campus Innenstadt, Munich, DE
- 93, Institute of Psychiatric Phenomics and Genomics (IPPG), Medical Center of the University of Munich, Campus Innenstadt, Munich, DE
- 94, Division of Cancer Epidemiology and Genetics, National Cancer Institute, Bethesda, MD, US
- 95, Behavioral Health Services, Kaiser Permanente Washington, Seattle, WA, US
- 96, Faculty of Medicine, Department of Psychiatry, University of Iceland, Reykjavik, IS
- 97, School of Medicine and Dentistry, James Cook University, Townsville, QLD, AU
- 98, Institute of Health and Wellbeing, University of Glasgow, Glasgow, GB
- 99, deCODE Genetics / Amgen, Reykjavik, IS

- 100, College of Biomedical and Life Sciences, Cardiff University, Cardiff, GB
- 101, Institute of Epidemiology and Social Medicine, University of Münster, Münster, Nordrhein-Westfalen, DE
- 102, Institute for Community Medicine, University Medicine Greifswald, Greifswald, Mecklenburg-Vorpommern, DE
- 103, Department of Psychiatry, University of California, San Diego, San Diego, CA, US
- 104, KG Jebsen Centre for Psychosis Research, Norway Division of Mental Health and Addiction, Oslo University Hospital, Oslo, NO
- 105, Medical Genetics Section, CGEM, IGMM, University of Edinburgh, Edinburgh, GB
- 106, Clinical Neurosciences, University of Cambridge, Cambridge, GB
- 107, Internal Medicine, Erasmus MC, Rotterdam, Zuid-Holland, NL
- 108, Roche Pharmaceutical Research and Early Development, Neuroscience, Ophthalmology and Rare Diseases Discovery & Translational Medicine Area, Roche Innovation Center Basel, F. Hoffmann-La Roche Ltd, Basel, CH
- 109, Department of Psychiatry and Psychotherapy, University Medicine Greifswald, Greifswald, Mecklenburg-Vorpommern, DE
- 110, Department of Psychiatry, Leiden University Medical Center, Leiden, NL
- 111, Virginia Institute for Psychiatric & Behavioral Genetics, Virginia Commonwealth University, Richmond, VA, US
- 112, Computational Sciences Center of Emphasis, Pfizer Global Research and Development, Cambridge, MA, US
- 113, Institute for Molecular Bioscience; Queensland Brain Institute, The University of Queensland, Brisbane, QLD, AU
- 114, Department of Psychiatry, University of Münster, Münster, Nordrhein-Westfalen, DE
- 115, Department of Psychiatry, Melbourne Medical School, University of Melbourne, Melbourne, AU
- 116, Institute of Medical Genetics and Pathology, University Hospital Basel, University of Basel, Basel, CH
- 117, Institute of Neuroscience and Medicine (INM-1), Research Center Juelich, Juelich, DE
- 118, Amsterdam Public Health Institute, Vrije Universiteit Medical Center, Amsterdam, NL
- 119, Centre for Integrative Biology, Università degli Studi di Trento, Trento, Trentino-Alto Adige, IT
- 120, Department of Psychiatry and Psychotherapy, Medical Center, University of Freiburg, Faculty of Medicine, University of Freiburg, Freiburg, DE
- 121, Psychiatry, Kaiser Permanente Northern California, San Francisco, CA, US
- 122, Medical Research Council Human Genetics Unit, Institute of Genetics and Molecular Medicine, University of Edinburgh, Edinburgh, GB
- 123, Department of Psychiatry, University of Toronto, Toronto, ON, CA
- 124, Centre for Addiction and Mental Health, Toronto, ON, CA
- 125, Division of Psychiatry, University College London, London, GB
- 126, Neuroscience Therapeutic Area, Janssen Research and Development, LLC, Titusville, NJ, US
- 127, Institute of Molecular and Cell Biology, University of Tartu, Tartu, EE
- 128, Psychosis Research Unit, Aarhus University Hospital, Risskov, Aarhus, DK
- 129, University of Liverpool, Liverpool, GB
- 130, Mental Health Center Copenhagen, Copenhagen University Hospital, Copenhagen, DK
- 131, Human Genetics and Computational Biomedicine, Pfizer Global Research and Development, Groton, CT, US
- 132, Psychiatry, Harvard Medical School, Boston, MA, US
- 133, Psychiatry, University of Iowa, Iowa City, IA, US
- 134, Department of Psychiatry and Behavioral Sciences, Johns Hopkins University, Baltimore, MD, US
- 135, Department of Psychiatry and Psychotherapy, University Medical Center Göttingen, Goettingen, Niedersachsen, DE
- 136, Human Genetics Branch, NIMH Division of Intramural Research Programs, Bethesda, MD, US
- 137, Faculty of Medicine, University of Iceland, Reykjavik, IS
- 138, Child and Adolescent Psychiatry, Erasmus MC, Rotterdam, Zuid-Holland, NL
- 139, Psychiatry, Erasmus MC, Rotterdam, Zuid-Holland, NL
- 140, Psychiatry, Dalhousie University, Halifax, NS, CA
- 141, Division of Epidemiology, New York State Psychiatric Institute, New York, NY, US
- 142, Department of Clinical Medicine, University of Copenhagen, Copenhagen, DK
- 143, Department of Medical & Molecular Genetics, King's College London, London, GB
- 144, Psychiatry & Behavioral Sciences, Stanford University, Stanford, CA, US
- 145, NIHR Maudsley Biomedical Research Centre, King's College London, London, GB

Bipolar Disorder Working Group of the Psychiatric Genomics Consortium

Eli A Stahl 1, 2, 3, Gerome Breen 4, 5, Andreas J Forstner 6, 7, 8, 9, Andrew McQuillin 10, Stephan Ripke 11, 12, 13, Vassily Trubetskoy 12, Manuel Mattheisen 14, 15, 16, 17, 18, Yunpeng Wang 19, 20, Jonathan R I Coleman 4, 5, Héléna A Gaspar 4, Christiaan A de Leeuw 21, Stacy Steinberg 22, Jennifer M Whitehead Pavlides 23, Maciej Trzaskowski 24, Tune H Pers 3, 25, Peter A Holmans 26, Liam Abbott 27, Esben Agerbo 18, 28, 29, Huda Akil 30, Diego Albani 31, Ney Alliey-Rodriguez 32, Thomas D Als 14, 15, 18, Adebayo Anjorin 33, Verneri Antilla 27, Swapnil Awasthi 12, Judith A Badner 34, Marie Bækvad-Hansen 18, 35, Jack D Barchas 36, Nicholas Bass 10, Michael Bauer 37, Sarah E Bergen 38, Carsten Bøcker Pedersen 18, 28, 29, Erlend Bøen 39, Marco Boks 40, James Boocock 41, Monika Budde 42, William Bunney 43, Margit Burmeister 44, Jonas Bybjerg-Grauholm 18, 35, William Byerley 45, Miquel Casas 46, 47, 48, 49, Felecia Cerrito 27, Pablo Cervantes 50, Alexander W Charney 2, Danfeng Chen 27, Claire Churchhouse 13, 27, Toni-Kim Clarke 51, William Coryell 52, David W Craig 53, Cristiana Cruceanu 54, David Curtis 55, 56, Piotr M Czerski 57, Anders M Dale 58, 59, 60, 61, Simone de Jong 4, 5, Franziska Degenhardt 8, 9, Jurgen Del-Favero 62, J Raymond DePaulo 63, Srdjan Djurovic 64, 65, Amanda L Dobbyn 1, 2, Ashley Dumont 27, Torbjørn Elvsåshagen 66, 67, Valentina Escott-Price 26, Chun Chieh Fan 61, Sascha B Fischer 6, Matthew Flickinger 68, Tatiana M Foroud 69, Liz Forty 26, Josef Frank 70, Christine Fraser 26, Nelson B Freimer 71, Louise Frisén 72, 73, 74, Katrin Gade 42, 75, Julie Garnham 76, Claudia Giambartolomei 41, Marianne Giørtz Pedersen 18, 28, 29, Scott D Gordon 77, Katherine Gordon-Smith 78, Elaine K Green 79, Melissa J Green 80, Tiffany A Greenwood 60, Jakob Grove 14, 15, 18, 81, Weihua Guan 82, José Guzman Parra 83, Marian L Hamshere 26, Martin Hautzinger 84, Urs Heilbronner 42, Stefan Herms 6, 8, 9, Maria Hipolito 85, Per Hoffmann 6, 8, 9, 86, Dominic Holland 58, 87, Laura Huckins 1, 2, Stéphane Jamain 88, 89, Anders Juréus 38, Radhika Kandaswamy 4, Robert Karlsson 38, James L Kennedy 90, 91, 92, 93, Sarah Kittel-Schneider 94, Sarah V Knott 78, James A Knowles 95, 96, Manolis Kogevinas 97, Anna C Koller 8, 9, Ralph Kupka 98, 99, 100, Catharina Lavebratt 72, Jacob Lawrence 101, William B Lawson 85, Markus Leber 102, Phil H Lee 11, 13, 103, Shawn E Levy 104, Jun Z Li 105, Chunyu Liu 106, Susanne Lucae 107, Anna Maaser 8, 9, Donald J MacIntyre 108, 109, Pamela B Mahon 63, 110, Wolfgang Maier 111, Lina Martinsson 73, Steve McCarroll 11, 112, Peter McGuffin 4, Melvin G McInnis 113, James D McKay 114, Helena Medeiros 96, Sarah E Medland 77, Fan Meng 30, 113, Lili Milani 115, Grant W Montgomery 24, Derek W Morris 116, 117, Thomas W Mühleisen 6, 118, Niamh Mullins 1, 4, Hoang Nguyen 1, 2, Caroline M Nievergelt 60, 119, Annelie Nordin Adolfsson 120, Evaristus A Nwulia 85, Claire O'Donovan 76, Loes M Olde Loohuis 71, Anil P S Ori 71, Lilijana Oruc 121, Urban Ösby 122, Roy H Perlis 123, 124, Amy Perry 78, Andrea Pfennig 37, James B Potash 63, Shaun M Purcell 2, 110, Eline J Regeer 125, Andreas Reif 94, Céline S Reinbold 6, John P Rice 126, Fabio Rivas 83, Margarita Rivera 4, 127, Panos Roussos 1, 2, 128, Douglas M Ruderfer 129, Cristina Sánchez-Mora 46, 47, 49, Alan F Schatzberg 130, William A Scheftner 131, Nicholas J Schork 132, Cynthia Shannon Weickert 80, 133, Tatyana Shehktman 60, Paul D Shilling 60, Engilbert Sigurdsson 134, Claire Slaney 76, Olav B Smeland 58, 135, 136, Janet L Sobell 137, Christine Søholm Hansen 18, 35, Anne T Spijker 138, David St Clair 139, Michael Steffens 140, John S Strauss 92, 141, Fabian Streit 70, Jana Strohmaier 70, Szabolcs Szelinger 142, Robert C Thompson 113, Thorgeir E Thorgeirsson 22, Jens Treutlein 70, Helmut Vedder 143, Weiqing Wang 1, 2, Stanley J Watson 113, Thomas W Weickert 80, 133, Stephanie H Witt 70, Simon Xi 144, Wei Xu 145, 146, Allan H Young 147, Peter Zandi 148, Peng Zhang 149, Sebastian Zollner 113, Rolf Adolfsson 120, Ingrid Agartz 16, 39, 150, Martin Alda 76, 151, Lena Backlund 73, Bernhard T Baune 152, Frank Bellivier 153, 154, 155, 156, Wade H Berrettini 157, Joanna M Biernacka 158, Douglas H R Blackwood 51, Michael Boehnke 68, Anders D Børglum 14, 15, 18, Aiden Corvin 117, Nicholas Craddock 26, Mark J Daly 13, 27, Udo Dannlowski 159, Tõnu Esko 3, 112, 115, 160, Bruno Etain 153, 155, 156, 161, Mark Frye 162, Janice M Fullerton 133, 163, Elliot S Gershon 32, 164, Michael Gill 117, Fernando Goes 165, Maria Grigoroiu-Serbanescu 166, Joanna Hauser 57, David M Hougaard 18, 35, Christina M Hultman 38, Ian Jones 26, Lisa A Jones 78, René S Kahn 2, 40, George Kirov 26, Mikael Landén 38, 167, Marion Leboyer 89, 153, 168, Cathryn M Lewis 4, 169, Qingqin S Li 170, Jolanta Lissowska 171, Nicholas G Martin 77, 172, Fermin Mayoral 83, Susan L McElroy 173, Andrew M McIntosh 51, 174, Francis J McMahon 175, Ingrid Melle 176, 177, Andres Metspalu 115, 178, Philip B Mitchell 80, Gunnar Morken 179, 180, Ole Mors 18, 181, Preben Bo Mortensen 14, 18, 28, 29, Bertram Müller-Myhsok 54, 182, 183, Richard M Myers 104, Benjamin M Neale 3, 11, 13, 13, 27, Vishwajit Nimgaonkar 184, Merete Nordentoft 18, 185, Markus M Nöthen

- 8, 9, Michael C O'Donovan 26, Ketil J Oedegaard 186, 187, Michael J Owen 26, Sara A Paciga 188, Carlos Pato 96, 189, Michael T Pato 96, Danielle Posthuma 190, 191, Josep Antoni Ramos-Quiroga 46, 47, 48, 49, Marta Ribasés 46, 47, 49, Marcella Rietschel 70, Guy A Rouleau 192, 193, Martin Schalling 72, Peter R Schofield 133, 163, Thomas G Schulze 42, 63, 70, 75, 175, Alessandro Serretti 194, Jordan W Smoller 11, 195, 196, Hreinn Stefansson 22, Kari Stefansson 22, 197, Eystein Stordal 198, 199, Patrick F Sullivan 38, 200, 201, Gustavo Turecki 202, Arne E Vaaler 203, Eduard Vieta 204, John B Vincent 141, Thomas Werge 18, 205, 206, John I Nurnberger 207, Naomi R Wray 23, 24, Arianna Di Florio 26, 201, Howard J Edenberg 208, Sven Cichon 6, 8, 86, 118, Roel A Ophoff 209, 210, Laura J Scott 68, Ole A Andreassen 135, 136, John Kelsoe 60, Pamela Sklar 1, 2, 128
- 1, Department of Genetics and Genomic Sciences, Icahn School of Medicine at Mount Sinai, New York, NY, US
- 2, Department of Psychiatry, Icahn School of Medicine at Mount Sinai, New York, NY, US
- 3, Medical and Population Genetics, Broad Institute, Cambridge, MA, US
- 4, MRC Social, Genetic and Developmental Psychiatry Centre, King's College London, London, GB
- 5, NIHR BRC for Mental Health, King's College London, London, GB
- 6, Department of Biomedicine, University of Basel, Basel, CH
- 7, Department of Psychiatry (UPK), University of Basel, Basel, CH
- 8, Institute of Human Genetics, University of Bonn, Bonn, DE
- 9, Life&Brain Center, Department of Genomics, University of Bonn, Bonn, DE
- 10, Division of Psychiatry, University College London, London, GB
- 11, Stanley Center for Psychiatric Research, Broad Institute, Cambridge, MA, US
- 12, Department of Psychiatry and Psychotherapy, Charité Universitätsmedizin, Berlin, DE
- 13, Analytic and Translational Genetics Unit, Massachusetts General Hospital, Boston, MA, US
- 14, iSEQ, Center for Integrative Sequencing, Aarhus University, Aarhus, DK
- 15, Department of Biomedicine Human Genetics, Aarhus University, Aarhus, DK
- 16, Department of Clinical Neuroscience, Centre for Psychiatry Research, Karolinska Institutet, Stockholm, SE
- 17, Department of Psychiatry, Psychosomatics and Psychotherapy, Center of Mental Health, University Hospital Würzburg, Würzburg, DE
- 18, iPSYCH, The Lundbeck Foundation Initiative for Integrative Psychiatric Research, DK
- 19, Institute of Biological Psychiatry, Mental Health Centre Sct. Hans, Copenhagen, DK
- 20, Institute of Clinical Medicine, University of Oslo, Oslo, NO
- 21, Department of Complex Trait Genetics, Center for Neurogenomics and Cognitive Research Neuroscience, Vrije Universiteit Amsterdam, Amsterdam, NL
- 22, deCODE Genetics / Amgen, Reykjavik, IS
- 23, Queensland Brain Institute, The University of Queensland, Brisbane, QLD, AU
- 24, Institute for Molecular Bioscience, The University of Queensland, Brisbane, QLD, AU
- 25, Division of Endocrinology and Center for Basic and Translational Obesity Research, Boston Children's Hospital, Boston, MA, US
- 26, Medical Research Council Centre for Neuropsychiatric Genetics and Genomics, Division of Psychological Medicine and Clinical Neurosciences, Cardiff University, Cardiff, GB
- 27, Stanley Center for Psychiatric Research, Broad Institute, Boston, MA, US
- 28, National Centre for Register-Based Research, Aarhus University, Aarhus, DK
- 29, Centre for Integrated Register-based Research, Aarhus University, Aarhus, DK
- 30, Molecular & Behavioral Neuroscience Institute, University of Michigan, Ann Arbor, MI, US
- 31, NEUROSCIENCE, Istituto Di Ricerche Farmacologiche Mario Negri, Milano, IT
- 32, Department of Psychiatry and Behavioral Neuroscience, University of Chicago, Chicago, IL, US
- 33, Psychiatry, Berkshire Healthcare NHS Foundation Trust, Bracknell, GB
- 34, Psychiatry, Rush University Medical Center, Chicago, IL, US
- 35, Center for Neonatal Screening, Department for Congenital Disorders, Statens Serum Institut, Copenhagen, DK
- 36, Department of Psychiatry, Weill Cornell Medical College, New York, NY, US
- 37, Department of Psychiatry and Psychotherapy, University Hospital Carl Gustav Carus, Technische Universität Dresden, DE
- 38, Department of Medical Epidemiology and Biostatistics, Karolinska Institutet, Stockholm, SE
- 39, Department of Psychiatric Research, Diakonhjemmet Hospital, Oslo, NO
- 40, Psychiatry, UMC Utrecht Hersencentrum Rudolf Magnus, Utrecht, NL
- 41, Human Genetics, University of California Los Angeles, Los Angeles, CA, US

- 42, Institute of Psychiatric Phenomics and Genomics (IPPG), Medical Center of the University of Munich, Munich, DE
- 43, Department of Psychiatry and Human Behavior, University of California, Irvine, Irvine, CA, US
- 44, Molecular & Behavioral Neuroscience Institute and Department of Computational Medicine & Bioinformatics, University of Michigan, Ann Arbor, MI, US
- 45, Psychiatry, University of California San Francisco, San Francisco, CA, US
- 46, Instituto de Salud Carlos III, Biomedical Network Research Centre on Mental Health (CIBERSAM), Madrid, ES
- 47, Department of Psychiatry, Hospital Universitari Vall d'Hebron, Barcelona, ES
- 48, Department of Psychiatry and Forensic Medicine, Universitat Autònoma de Barcelona, Barcelona, ES
- 49, Psychiatric Genetics Unit, Group of Psychiatry Mental Health and Addictions, Vall d'Hebron Research Institut (VHIR), Universitat Autònoma de Barcelona, Barcelona, ES
- 50, Department of Psychiatry, Mood Disorders Program, McGill University Health Center, Montreal, QC, CA
- 51, Division of Psychiatry, University of Edinburgh, Edinburgh, GB
- 52, University of Iowa Hospitals and Clinics, Iowa City, IA, US
- 53, Translational Genomics, USC, Phoenix, AZ, US
- 54, Department of Translational Research in Psychiatry, Max Planck Institute of Psychiatry, Munich, DE
- 55, Centre for Psychiatry, Queen Mary University of London, London, GB
- 56, UCL Genetics Institute, University College London, London, GB
- 57, Department of Psychiatry, Laboratory of Psychiatric Genetics, Poznan University of Medical Sciences, Poznan, PL
- 58, Department of Neurosciences, University of California San Diego, La Jolla, CA, US
- 59, Department of Radiology, University of California San Diego, La Jolla, CA, US
- 60, Department of Psychiatry, University of California San Diego, La Jolla, CA, US
- 61, Department of Cognitive Science, University of California San Diego, La Jolla, CA, US
- 62, Applied Molecular Genomics Unit, VIB Department of Molecular Genetics, University of Antwerp, Antwerp, Belgium
- 63, Department of Psychiatry and Behavioral Sciences, Johns Hopkins University School of Medicine, Baltimore, MD, US
- 64, Department of Medical Genetics, Oslo University Hospital Ullevål, Oslo, NO
- 65, NORMENT, KG Jebsen Centre for Psychosis Research, Department of Clinical Science, University of Bergen, Bergen, NO
- 66, Department of Neurology, Oslo University Hospital, Oslo, NO
- 67, NORMENT, KG Jebsen Centre for Psychosis Research, Oslo University Hospital, Oslo, NO
- 68, Center for Statistical Genetics and Department of Biostatistics, University of Michigan, Ann Arbor, MI, US
- 69, Department of Medical & Molecular Genetics, Indiana University, Indianapolis, IN, US
- 70, Department of Genetic Epidemiology in Psychiatry, Central Institute of Mental Health, Medical Faculty Mannheim, Heidelberg University, Mannheim, DE
- 71, Center for Neurobehavioral Genetics, University of California Los Angeles, Los Angeles, CA, US
- 72, Department of Molecular Medicine and Surgery, Karolinska Institutet and Center for Molecular Medicine, Karolinska University Hospital, Stockholm, SE
- 73, Department of Clinical Neuroscience, Karolinska Institutet and Center for Molecular Medicine, Karolinska University Hospital, Stockholm, SE
- 74, Child and Adolescent Psychiatry Research Center, Stockholm, SE
- 75, Department of Psychiatry and Psychotherapy, University Medical Center Göttingen, Göttingen, DE
- 76, Department of Psychiatry, Dalhousie University, Halifax, NS, CA
- 77, Genetics and Computational Biology, QIMR Berghofer Medical Research Institute, Brisbane, QLD, AU
- 78, Department of Psychological Medicine, University of Worcester, Worcester, GB
- 79, School of Biomedical and Healthcare Sciences, Plymouth University Peninsula Schools of Medicine and Dentistry, Plymouth, GB
- 80, School of Psychiatry, University of New South Wales, Sydney, NSW, AU
- 81, Bioinformatics Research Centre, Aarhus University, Aarhus, DK
- 82, Biostatistics, University of Minnesota System, Minneapolis, MN, US
- 83, Mental Health Department, University Regional Hospital. Biomedicine Institute (IBIMA), Málaga, ES
- 84, Department of Psychology, Eberhard Karls Universität Tübingen, Tubingen, DE
- 85, Department of Psychiatry and Behavioral Sciences, Howard University Hospital, Washington, DC, US
- 86, Institute of Medical Genetics and Pathology, University Hospital Basel, Basel, CH

- 87, Center for Multimodal Imaging and Genetics, University of California San Diego, La Jolla, CA, US
- 88, Psychiatrie Translationnelle, Inserm U955, Créteil, FR
- 89, Faculté de Médecine, Université Paris Est, Créteil, FR
- 90, Campbell Family Mental Health Research Institute, Centre for Addiction and Mental Health, Toronto, ON, CA
- 91, Neurogenetics Section, Centre for Addiction and Mental Health, Toronto, ON, CA
- 92, Department of Psychiatry, University of Toronto, Toronto, ON, CA
- 93, Institute of Medical Sciences, University of Toronto, Toronto, ON, CA
- 94, Department of Psychiatry, Psychosomatic Medicine and Psychotherapy, University Hospital Frankfurt, Frankfurt am Main, DE
- 95, Cell Biology, SUNY Downstate Medical Center College of Medicine, Brooklyn, NY, US
- 96, Institute for Genomic Health, SUNY Downstate Medical Center College of Medicine, Brooklyn, NY, US
- 97, Center for Research in Environmental Epidemiology (CREAL), Barcelona, ES
- 98, Psychiatry, Altrecht, Utrecht, NL
- 99, Psychiatry, GGZ inGeest, Amsterdam, NL
- 100, Psychiatry, VU medisch centrum, Amsterdam, NL
- 101, Psychiatry, North East London NHS Foundation Trust, Ilford, GB
- 102, Clinic for Psychiatry and Psychotherapy, University Hospital Cologne, Cologne, DE
- 103, Psychiatric and Neurodevelopmental Genetics Unit, Massachusetts General Hospital, Boston, MA, US
- 104, HudsonAlpha Institute for Biotechnology, Huntsville, AL, US
- 105, Department of Human Genetics, University of Michigan, Ann Arbor, MI, US
- 106, Psychiatry, University of Illinois at Chicago College of Medicine, Chicago, IL, US
- 107, Max Planck Institute of Psychiatry, Munich, DE
- 108, Mental Health, NHS 24, Glasgow, GB
- 109, Division of Psychiatry, Centre for Clinical Brain Sciences, University of Edinburgh, Edinburgh, GB
- 110, Psychiatry, Brigham and Women's Hospital, Boston, MA, US
- 111, Department of Psychiatry and Psychotherapy, University of Bonn, Bonn, DE
- 112, Department of Genetics, Harvard Medical School, Boston, MA, US
- 113, Department of Psychiatry, University of Michigan, Ann Arbor, MI, US
- 114, Genetic Cancer Susceptibility Group, International Agency for Research on Cancer, Lyon, FR
- 115, Estonian Genome Center, University of Tartu, Tartu, EE
- 116, Discipline of Biochemistry, Neuroimaging and Cognitive Genomics (NICOG) Centre, National University of Ireland, Galway, Galway, IE
- 117, Neuropsychiatric Genetics Research Group, Dept of Psychiatry and Trinity Translational Medicine Institute, Trinity College Dublin, Dublin, IE
- 118, Institute of Neuroscience and Medicine (INM-1), Research Centre Jülich, Jülich, DE
- 119, Research/Psychiatry, Veterans Affairs San Diego Healthcare System, San Diego, CA, US
- 120, Department of Clinical Sciences, Psychiatry, Umeå University Medical Faculty, Umeå, SE
- 121, Department of Clinical Psychiatry, Psychiatry Clinic, Clinical Center University of Sarajevo, Sarajevo, BA
- 122, Department of Neurobiology, Care sciences, and Society, Karolinska Institutet and Center for Molecular Medicine, Karolinska University Hospital, Stockholm, SE
- 123, Psychiatry, Harvard Medical School, Boston, MA, US
- 124, Division of Clinical Research, Massachusetts General Hospital, Boston, MA, US
- 125, Outpatient Clinic for Bipolar Disorder, Altrecht, Utrecht, NL
- 126, Department of Psychiatry, Washington University in Saint Louis, Saint Louis, MO, US
- 127, Department of Biochemistry and Molecular Biology II, Institute of Neurosciences, Center for Biomedical Research, University of Granada, Granada, ES
- 128, Department of Neuroscience, Icahn School of Medicine at Mount Sinai, New York, NY, US
- 129, Medicine, Psychiatry, Biomedical Informatics, Vanderbilt University Medical Center, Nashville, TN, US
- 130, Psychiatry and Behavioral Sciences, Stanford University School of Medicine, Stanford, CA, US
- 131, Rush University Medical Center, Chicago, IL, US
- 132, Scripps Translational Science Institute, La Jolla, CA, US
- 133, Neuroscience Research Australia, Sydney, NSW, AU
- 134, Faculty of Medicine, Department of Psychiatry, School of Health Sciences, University of Iceland, Reykjavik, IS
- 135, Div Mental Health and Addiction, Oslo University Hospital, Oslo, NO
- 136, NORMENT, University of Oslo, Oslo, NO

- 137, Psychiatry and the Behavioral Sciences, University of Southern California, Los Angeles, CA, US
- 138, Mood Disorders, PsyQ, Rotterdam, NL
- 139, Institute for Medical Sciences, University of Aberdeen, Aberdeen, UK
- 140, Research Division, Federal Institute for Drugs and Medical Devices (BfArM), Bonn, DE
- 141, Centre for Addiction and Mental Health, Toronto, ON, CA
- 142, Neurogenomics, TGen, Los Angeles, AZ, US
- 143, Psychiatry, Psychiatrisches Zentrum Nordbaden, Wiesloch, DE
- 144, Computational Sciences Center of Emphasis, Pfizer Global Research and Development, Cambridge, MA, US
- 145, Department of Biostatistics, Princess Margaret Cancer Centre, Toronto, ON, CA
- 146, Dalla Lana School of Public Health, University of Toronto, Toronto, ON, CA
- 147, Psychological Medicine, Institute of Psychiatry, Psychology & Neuroscience, King's College London, London, GB
- 148, Department of Mental Health, Johns Hopkins University Bloomberg School of Public Health, Baltimore, MD, US
- 149, Institute of Genetic Medicine, Johns Hopkins University School of Medicine, Baltimore, MD, US
- 150, NORMENT, KG Jebsen Centre for Psychosis Research, Division of Mental Health and Addiction, Institute of Clinical Medicine and Diakonhjemmet Hospital, University of Oslo, Oslo, NO
- 151, National Institute of Mental Health, Klecany, CZ
- 152, Discipline of Psychiatry, University of Adelaide, Adelaide, SA, AU
- 153, Department of Psychiatry and Addiction Medicine, Assistance Publique Hôpitaux de Paris, Paris, FR
- 154, Paris Bipolar and TRD Expert Centres, FondaMental Foundation, Paris, FR
- 155, UMR-S1144 Team 1 : Biomarkers of relapse and therapeutic response in addiction and mood disorders, INSERM, Paris, FR
- 156, Psychiatry, Université Paris Diderot, Paris, FR
- 157, Psychiatry, University of Pennsylvania, Philadelphia, PA, US
- 158, Health Sciences Research, Mayo Clinic, Rochester, MN, US
- 159, Department of Psychiatry, University of Münster, Münster, DE
- 160, Division of Endocrinology, Children's Hospital Boston, Boston, MA, US
- 161, Centre for Affective Disorders, Institute of Psychiatry, Psychology and Neuroscience, London, GB
- 162, Department of Psychiatry & Psychology, Mayo Clinic, Rochester, MN, US
- 163, School of Medical Sciences, University of New South Wales, Sydney, NSW, AU
- 164, Department of Human Genetics, University of Chicago, Chicago, IL, US
- 165, Department of Psychiatry and Behavioral Sciences, Johns Hopkins University School of Medicine, Baltimore, MD, US
- 166, Biometric Psychiatric Genetics Research Unit, Alexandru Obregia Clinical Psychiatric Hospital, Bucharest, RO
- 167, Institute of Neuroscience and Physiology, University of Gothenburg, Gothenburg, SE
- 168, INSERM, Paris, FR
- 169, Department of Medical & Molecular Genetics, King's College London, London, GB
- 170, Neuroscience Therapeutic Area, Janssen Research and Development, LLC, Titusville, NJ, US
- 171, Cancer Epidemiology and Prevention, M. Sklodowska-Curie Cancer Center and Institute of Oncology, Warsaw, PL
- 172, School of Psychology, The University of Queensland, Brisbane, QLD, AU
- 173, Research Institute, Lindner Center of HOPE, Mason, OH, US
- 174, Centre for Cognitive Ageing and Cognitive Epidemiology, University of Edinburgh, Edinburgh, GB
- 175, Human Genetics Branch, Intramural Research Program, National Institute of Mental Health, Bethesda, MD, US
- 176, Division of Mental Health and Addiction, Oslo University Hospital, Oslo, NO
- 177, Division of Mental Health and Addiction, University of Oslo, Institute of Clinical Medicine, Oslo, NO
- 178, Institute of Molecular and Cell Biology, University of Tartu, Tartu, EE
- 179, Mental Health, Faculty of Medicine and Health Sciences, Norwegian University of Science and Technology NTNU, Trondheim, NO
- 180, Psychiatry, St Olavs University Hospital, Trondheim, NO
- 181, Psychosis Research Unit, Aarhus University Hospital, Risskov, DK
- 182, Munich Cluster for Systems Neurology (SyNergy), Munich, DE
- 183, University of Liverpool, Liverpool, GB

- 184, Psychiatry and Human Genetics, University of Pittsburgh, Pittsburgh, PA, US
- 185, Mental Health Services in the Capital Region of Denmark, Mental Health Center Copenhagen, University of Copenhagen, Copenhagen, DK
- 186, Division of Psychiatry, Haukeland Universitetssjukehus, Bergen, NO
- 187, Faculty of Medicine and Dentistry, University of Bergen, Bergen, NO
- 188, Human Genetics and Computational Biomedicine, Pfizer Global Research and Development, Groton, CT, US
- 189, College of Medicine Institute for Genomic Health, SUNY Downstate Medical Center College of Medicine, Brooklyn, NY, US
- 190, Department of Complex Trait Genetics, Center for Neurogenomics and Cognitive Research, Amsterdam Neuroscience, Vrije Universiteit Amsterdam, Amsterdam, NL
- 191, Department of Clinical Genetics, Amsterdam Neuroscience, Vrije Universiteit Medical Center, Amsterdam, NI
- 192, Department of Neurology and Neurosurgery, McGill University, Faculty of Medicine, Montreal, QC, CA
- 193, Montreal Neurological Institute and Hospital, Montreal, QC, CA
- 194, Department of Biomedical and NeuroMotor Sciences, University of Bologna, Bologna, IT
- 195, Department of Psychiatry, Massachusetts General Hospital, Boston, MA, US
- 196, Psychiatric and Neurodevelopmental Genetics Unit (PNGU), Massachusetts General Hospital, Boston, MA, US
- 197, Faculty of Medicine, University of Iceland, Reykjavik, IS
- 198, Department of Psychiatry, Hospital Namsos, Namsos, NO
- 199, Department of Neuroscience, Norges Teknisk Naturvitenskapelige Universitet Fakultet for naturvitenskap og teknologi, Trondheim, NO
- 200, Department of Genetics, University of North Carolina at Chapel Hill, Chapel Hill, NC, US
- 201, Department of Psychiatry, University of North Carolina at Chapel Hill, Chapel Hill, NC, US
- 202, Department of Psychiatry, McGill University, Montreal, QC, CA
- 203, Dept of Psychiatry, Sankt Olavs Hospital Universitetssykehuset i Trondheim, Trondheim, NO
- 204, Clinical Institute of Neuroscience, Hospital Clinic, University of Barcelona, IDIBAPS, CIBERSAM, Barcelona, SP
- 205, Institute of Biological Psychiatry, MHC Sct. Hans, Mental Health Services Copenhagen, Roskilde, DK
- 206, Department of Clinical Medicine, University of Copenhagen, Copenhagen, DK
- 207, Psychiatry, Indiana University School of Medicine, Indianapolis, IN, US
- 208, Biochemistry and Molecular Biology, Indiana University School of Medicine, Indianapolis, IN, US
- 209, Jane and Terry Semel Institute for Neuroscience and Human Behavior, Los Angeles, CA, US
- 210, UMC Utrecht Hersencentrum Rudolf Magnus, Utrecht, NL

Schizophrenia Working Group of the Psychiatric Genomics Consortium

Stephan Ripke^{1,2}, Benjamin M. Neale^{1,2,3,4}, Aiden Corvin⁵, James T. R. Walters⁶, Kai-How Farh¹, Peter A. Holmans^{6,7}, Phil Lee^{1,2,4}, Brendan Bulik-Sullivan^{1,2}, David A. Collier^{8,9}, Hailiang Huang^{1,3}, Tune H. Pers^{3,10,11}, Ingrid Agartz^{12,13,14}, Esben Agerbo^{15,16,17}, Margot Albus¹⁸, Madeline Alexander¹⁹, Farooq Amin^{20,21}, Silviu A. Bacanu²², Martin Begemann²³, Richard A. Belliveau Jr², Judit Bene^{24,25}, Sarah E. Bergen^{2,26}, Elizabeth Bevilacqua², Tim B. Bigdeli²², Donald W. Black²⁷, Richard Bruggeman²⁸, Nancy G. Buccola²⁹, Randy L. Buckner^{30,31,32}, William Byerley³³, Wiepke Cahn³⁴, Guiqing Cai^{35,36}, Dominique Campion³⁷, Rita M. Cantor³⁸, Vaughan J. Carr^{39,40}, Noa Carrera⁶, Stanley V. Catts^{39,41}, Kimberly D. Chambert², Raymond C. K. Chan⁴², Ronald Y. L. Chen⁴³, Eric Y. H. Chen^{43,44}, Wei Cheng⁴⁵, Eric F. C. Cheung⁴⁶, Siow Ann Chong⁴⁷, C. Robert Cloninger⁴⁸, David Cohen⁴⁹, Nadine Cohen⁵⁰, Paul Cormican⁵, Nick Craddock^{6,7}, James J. Crowley⁵¹, David Curtis^{52,53}, Michael Davidson⁵⁴, Kenneth L. Davis³⁶, Franziska Degenhardt^{55,56}, Jurgen Del Favero⁵⁷, Ditte Demontis^{17,58,59}, Dimitris Dikeos⁶⁰, Timothy Dinan⁶¹, Srdjan

Djurovic^{14,62}, Gary Donohoe^{5,63}, Elodie Drapeau³⁶, Jubao Duan^{64,65}, Frank Dudbridge⁶⁶, Naser ${\sf Durmishi}^{67}, {\sf Peter Eichhammer}^{68}, {\sf Johan Eriksson}^{69,70,71}, {\sf Valentina Escott-Price}^6, {\sf Laurent Essioux}^{72}, {\sf Ayman}$ H. Fanous^{73,74,75,76}, Martilias S. Farrell⁵¹, Josef Frank⁷⁷, Lude Franke⁷⁸, Robert Freedman⁷⁹, Nelson B. Freimer⁸⁰, Marion Friedl⁸¹, Joseph I. Friedman³⁶, Menachem Fromer^{1,2,4,82}, Giulio Genovese², Lyudmila Georgieva⁶, Ina Giegling^{81,83}, Paola Giusti-Rodr iguez⁵¹, Stephanie Godard⁸⁴, Jacqueline I. Goldstein^{1,3}, Vera Golimbet⁸⁵, Srihari Gopal⁸⁶, Jacob Gratten⁸⁷, Lieuwe de Haan⁸⁸, Christian Hammer²³, Marian L. Hamshere⁶, Mark Hansen⁸⁹, Thomas Hansen^{17,90}, Vahram Haroutunian^{36,91,92}, Annette M. Hartmann⁸¹, Frans A. Henskens^{39,93,94}, Stefan Herms^{55,56,95}, Joel N. Hirschhorn^{3,11,96}, Per Hoffmann^{55,56,95}, Andrea Hofman^{55,56}, Mads V. Hollegaard⁹⁷, David M. Hougaard⁹⁷, Masashi Ikeda⁹⁸, Inge Joa⁹⁹, Antonio Julia¹⁰⁰, Rene' S. Kahn³⁴, Luba Kalaydjieva^{101,102}, Sena Karachanak-Yankova¹⁰³, Juha Karjalainen⁷⁸, David Kavanagh⁶, Matthew C. Keller¹⁰⁴, James L. Kennedy^{105,106,107}, Andrey Khrunin¹⁰⁸, Yunjung Kim⁵¹, Janis Klovins¹⁰⁹, James A. Knowles¹¹⁰, Bettina Konte⁸¹, Vaidutis Kucinskas¹¹¹, Zita Ausrele Kucinskiene¹¹¹, Hana Kuzelova-Ptackova¹¹², Anna K. Ka hler²⁶, Claudine Laurent^{19,113}, Jimmy Lee Chee Keong^{47,114}, S. Hong Lee⁸⁷, Sophie E. Legge⁶, Bernard Lerer¹¹⁵, Miaoxin Li^{43,44,116}, Tao Li¹¹⁷, Kung-Yee Liang¹¹⁸, Jeffrey Lieberman 119, Svetlana Limborska 108, Carmel M. Loughland 39,120, Jan Lubinski 121, Jouko Lonnqvist 122, Milan Macek Jr¹¹², Patrik K. E. Magnusson²⁶, Brion S. Maher¹²³, Wolfgang Maier¹²⁴, Jacques Mallet¹²⁵, Sara Marsal 100, Manuel Mattheisen 17,58,59,126, Morten Mattingsdal 14,127, Robert W. McCarley 128,129, Colm McDonald 130, Andrew M. McIntosh 131,132, Sandra Meier 77, Carin J. Meijer 88, Bela Melegh 24,25, Ingrid Melle 14,133, Raquelle I. Mesholam-Gately 128,134, Andres Metspalu 135, Patricia T. Michie 39,136, Lili Milani¹³⁵, Vihra Milanova¹³⁷, Younes Mokrab⁸, Derek W. Morris^{5,63}, Ole Mors^{17,58,138}, Kieran C. Murphy¹³⁹, Robin M. Murray¹⁴⁰, Inez Myin-Germeys¹⁴¹, Bertram Muller-Myhsok^{142,143,144}, Mari Nelis¹³⁵, Igor Nenadic¹⁴⁵, Deborah A. Nertney¹⁴⁶, Gerald Nestadt¹⁴⁷, Kristin K. Nicodemus¹⁴⁸, Liene Nikitina-Zake¹⁰⁹, Laura Nisenbaum¹⁴⁹, Annelie Nordin¹⁵⁰, Eadbhard O'Callaghan¹⁵¹, Colm O'Dushlaine², F. Anthony O'Neill¹⁵², Sang-Yun Oh¹⁵³, Ann Olincy⁷⁹, Line Olsen^{17,90}, Jim Van Os^{141,154}, Psychosis Endophenotypes International Consortium¹⁵⁵, Christos Pantelis^{39,156}, George N. Papadimitriou⁶⁰, Sergi Papiol²³, Elena Parkhomenko³⁶, Michele T. Pato¹¹⁰, Tiina Paunio^{157,158}, Milica Pejovic-Milovancevic¹⁵⁹, Diana O. Perkins¹⁶⁰, Olli Pietila inen^{158,161}, Jonathan Pimm⁵³, Andrew J. Pocklington⁶, John Powell¹⁴⁰, Alkes Price^{3,162}, Ann E. Pulver¹⁴⁷, Shaun M. Purcell⁸², Digby Quested¹⁶³, Henrik B. Rasmussen^{17,90}, Abraham Reichenberg³⁶, Mark A. Reimers¹⁶⁴, Alexander L. Richards⁶, Joshua L. Roffman^{30,32}, Panos Roussos^{82,165}, Douglas M. Ruderfer^{6,82}, Veikko Salomaa⁷¹, Alan R. Sanders^{64,65}, Ulrich Schall^{39,120}, Christian R. Schubert ¹⁶⁶, Thomas G. Schulze ^{77,167}, Sibylle G. Schwab ¹⁶⁸, Edward M. Scolnick ², Rodney J. Scott^{39,169,170}, Larry J. Seidman^{128,134}, Jianxin Shi¹⁷¹, Engilbert Sigurdsson¹⁷², Teimuraz Silagadze¹⁷³, Jeremy M. Silverman^{36,174}, Kang Sim⁴⁷, Petr Slominsky¹⁰⁸, Jordan W. Smoller^{2,4}, Hon-Cheong So⁴³, Chris C. A. Spencer¹⁷⁵, Eli A. Stahl^{3,82}, Hreinn Stefansson¹⁷⁶, Stacy Steinberg¹⁷⁶, Elisabeth Stogmann¹⁷⁷, Richard E. Straub¹⁷⁸, Eric Strengman^{179,34}, Jana Strohmaier⁷⁷, T. Scott Stroup¹¹⁹, Mythily Subramaniam⁴⁷, Jaana Suvisaari¹²², Dragan M. Svrakic⁴⁸, Jin P. Szatkiewicz⁵¹, Erik So derman¹², Srinivas Thirumalai¹⁸⁰, Draga Toncheva¹⁰³, Sarah Tosato¹⁸¹, Juha Veijola^{182,183}, John Waddington¹⁸⁴, Dermot Walsh¹⁸⁵, Dai Wang⁸⁶, Qiang Wang 117, Bradley T. Webb 22, Mark Weiser 54, Dieter B. Wildenauer 186, Nigel M. Williams 6, Stephanie Williams⁵¹, Stephanie H. Witt⁷⁷, Aaron R. Wolen¹⁶⁴, Emily H. M. Wong⁴³, Brandon K. Wormley²², Hualin Simon Xi¹⁸⁷, Clement C. Zai^{105,106}, Xuebin Zheng¹⁸⁸, Fritz Zimprich¹⁷⁷, Naomi R. Wray⁸⁷, Kari Stefansson¹⁷⁶, Peter M. Visscher⁸⁷, Wellcome Trust Case-Control Consortium 2, Rolf Adolfsson¹⁵⁰, Ole A. Andreassen 14,133, Douglas H. R. Blackwood 132, Elvira Bramon 190, Joseph D. Buxbaum 35,36,91,191, Anders D. Børglum^{17,58,59,138}, Sven Cichon^{55,56,95,192}, Ariel Darvasi¹⁹³, Enrico Domenici¹⁹⁴, Hannelore Ehrenreich²³, To nu Esko^{3,11,96,135}, Pablo V. Gejman^{64,65}, Michael Gill⁵, Hugh Gurling⁵³, Christina M. Hultman²⁶, Nakao Iwata⁹⁸, Assen V. Jablensky^{39,102,186,195}, Erik G. Jo nsson^{12,14}, Kenneth S. Kendler¹⁹⁶,

George Kirov⁶, Jo Knight^{105,106,107}, Todd Lencz^{197,198,199}, Douglas F. Levinson¹⁹, Qingqin S. Li⁸⁶, Jianjun Liu^{188,200}, Anil K. Malhotra^{197,198,199}, Steven A. McCarroll^{2,96}, Andrew McQuillin⁵³, Jennifer L. Moran², Preben B. Mortensen^{15,16,17}, Bryan J. Mowry^{87,201}, Markus M. Noʻʻthen^{55,56}, Roel A. Ophoff^{38,80,34}, Michael J. Owen^{6,7}, Aarno Palotie^{2,4,161}, Carlos N. Pato¹¹⁰, Tracey L. Petryshen^{2,128,202}, Danielle Posthuma^{203,204,205}, Marcella Rietschel⁷⁷, Brien P. Riley¹⁹⁶, Dan Rujescu^{81,83}, Pak C. Sham^{43,44,116}, Pamela Sklar^{82,91,165}, David St Clair²⁰⁶, Daniel R. Weinberger^{178,207}, Jens R. Wendland¹⁶⁶, Thomas Werge^{17,90,208}, Mark J. Daly^{1,2,3}, Patrick F. Sullivan^{26,51,160} & Michael C. OʻDonovan^{6,7}

1 Analytic and Translational Genetics Unit, Massachusetts General Hospital, Boston, Massachusetts 02114, USA. ²Stanley Center for Psychiatric Research, Broad Institute of MIT and Harvard, Cambridge, Massachusetts 02142, USA. ³Medical and Population Genetics Program, Broad Institute of MIT and Harvard, Cambridge, Massachusetts 02142, USA. ⁴Psychiatric and Neurodevelopmental Genetics Unit, Massachusetts General Hospital, Boston, Massachusetts 02114, USA. ⁵Neuropsychiatric Genetics Research Group, Department of Psychiatry, Trinity College Dublin, Dublin 8, Ireland. ⁶MRC Centre for Neuropsychiatric Genetics and Genomics, Institute of Psychological Medicine and Clinical Neurosciences, School of Medicine, Cardiff University, Cardiff CF24 4HQ, UK. ⁷National Centre for Mental Health, Cardiff University, Cardiff CF24 4HQ, UK. ⁸Eli Lilly and Company Limited, Erl Wood Manor, Sunninghill Road, Windlesham, Surrey GU20 6PH, UK. ⁹Social, Genetic and Developmental Psychiatry Centre, Institute of Psychiatry, King's College London, London SE5 8AF, UK. ¹⁰Center for Biological Sequence Analysis, Department of Systems Biology, Technical University of Denmark, DK-2800, Denmark. 11 Division of Endocrinology and Center for Basic and Translational Obesity Research, Boston Children's Hospital, Boston, Massachusetts 02115, USA. ¹²Department of Clinical Neuroscience, Psychiatry Section, Karolinska Institutet, SE-17176 Stockholm, Sweden, ¹³Department of Psychiatry, Diakonhiemmet Hospital, 0319 Oslo, Norway. ¹⁴NORMENT, KG Jebsen Centre for Psychosis Research, Institute of Clinical Medicine, University of Oslo, 0424 Oslo, Norway. ¹⁵Centre for Integrative Register-based Research, CIRRAU, Aarhus University, DK-8210 Aarhus, Denmark. ¹⁶National Centre for Register-based Research, Aarhus University, DK-8210 Aarhus, Denmark. ¹⁷The Lundbeck Foundation Initiative for Integrative Psychiatric Research, iPSYCH, Denmark. ¹⁸State Mental Hospital, 85540 Haar, Germany. ¹⁹Department of Psychiatry and Behavioral Sciences, Stanford University, Stanford, California 94305, USA. ²⁰Department of Psychiatry and Behavioral Sciences, Atlanta Veterans Affairs Medical Center, Atlanta, Georgia 30033, USA. ²¹Department of Psychiatry and Behavioral Sciences, Emory University, Atlanta, Georgia 30322, USA. ²²Virginia Institute for Psychiatric and Behavioral Genetics, Department of Psychiatry, Virginia Commonwealth University, Richmond, Virginia 23298, USA. ²³Clinical Neuroscience, Max Planck Institute of Experimental Medicine, Go¨ttingen 37075, Germany. ²⁴Department of Medical Genetics, University of Pe´cs, Pe´cs H-7624, Hungary. ²⁵Szentagothai Research Center, University of Pé cs, Pé cs H-7624, Hungary. ²⁶Department of Medical Epidemiology and Biostatistics, Karolinska Institutet, Stockholm SE-17177, Sweden. ²⁷Department of Psychiatry, University of Iowa Carver College of Medicine, Iowa City, Iowa 52242, USA. ²⁸University Medical Center Groningen, Department of Psychiatry, University of Groningen NL-9700 RB, The Netherlands. ²⁹School of Nursing, Louisiana State University Health Sciences Center, New Orleans, Louisiana 70112, USA. ³⁰Athinoula A. Martinos Center, Massachusetts General Hospital, Boston, Massachusetts 02129, USA. ³¹Center for Brain Science, Harvard University, Cambridge, Massachusetts 02138, USA. ³²Department of Psychiatry, Massachusetts General Hospital, Boston, Massachusetts 02114, USA. ³³Department of Psychiatry, University of California at San Francisco, San Francisco, California 94143, USA. ³⁴University Medical Center Utrecht, Department of Psychiatry, Rudolf Magnus Institute of Neuroscience, 3584 Utrecht, The Netherlands. ³⁵Department of Human Genetics, Icahn School of Medicine at Mount Sinai, New York, New York 10029, USA. ³⁶Department of Psychiatry, Icahn School of Medicine at Mount Sinai, New York, New York 10029, USA. ³⁷Centre Hospitalier du Rouvray and INSERM U1079 Faculty of Medicine, 76301 Rouen, France. ³⁸Department of Human Genetics, David Geffen School of Medicine, University of California, Los Angeles, California 90095,

USA. ³⁹Schizophrenia Research Institute, Sydney NSW 2010, Australia. ⁴⁰School of Psychiatry, University of New South Wales, Sydney NSW 2031, Australia. ⁴¹Royal Brisbane and Women's Hospital, University of Queensland, Brisbane, St Lucia QLD 4072, Australia. ⁴²Institute of Psychology, Chinese Academy of Science, Beijing 100101, China. ⁴³Department of Psychiatry, Li Ka Shing Faculty of Medicine, The University of Hong Kong, Hong Kong, China. 44State Key Laboratory for Brain and Cognitive Sciences, Li Ka Shing Faculty of Medicine, The University of Hong Kong, Hong Kong, China. ⁴⁵Department of Computer Science, University of North Carolina, Chapel Hill, North Carolina 27514, USA. ⁴⁶Castle Peak Hospital, Hong Kong, China. ⁴⁷Institute of Mental Health, Singapore 539747, Singapore. ⁴⁸Department of Psychiatry, Washington University, St. Louis, Missouri 63110, USA. ⁴⁹Department of Child and Adolescent Psychiatry, Assistance Publique Hopitaux de Paris, Pierre and Marie Curie Faculty of Medicine and Institute for Intelligent Systems and Robotics, Paris 75013, France. ⁵⁰Blue Note Biosciences, Princeton, New Jersey 08540, USA ⁵¹Department of Genetics, University of North Carolina, Chapel Hill, North Carolina 27599-7264, USA. ⁵²Department of Psychological Medicine, Queen Mary University of London, London E1 1BB, UK. ⁵³Molecular Psychiatry Laboratory, Division of Psychiatry, University College London, London WC1E 6JJ, UK. ⁵⁴Sheba Medical Center, Tel Hashomer 52621, Israel. ⁵⁵Department of Genomics, Life and Brain Center, D-53127 Bonn, Germany. ⁵⁶Institute of Human Genetics, University of Bonn, D-53127 Bonn, Germany. ⁵⁷Applied Molecular Genomics Unit, VIB Department of Molecular Genetics, University of Antwerp, B-2610 Antwerp, Belgium. ⁵⁸Centre for Integrative Sequencing, iSEQ, Aarhus University, DK-8000 Aarhus C, Denmark. ⁵⁹Department of Biomedicine, Aarhus University, DK-8000 Aarhus C, Denmark. ⁶⁰First Department of Psychiatry, University of Athens Medical School, Athens 11528, Greece. ⁶¹Department of Psychiatry, University College Cork, Co. Cork, Ireland. ⁶²Department of Medical Genetics, Oslo University Hospital, 0424 Oslo, Norway. ⁶³Cognitive Genetics and Therapy Group. School of Psychology and Discipline of Biochemistry, National University of Ireland Galway, Co. Galway, Ireland. ⁶⁴Department of Psychiatry and Behavioral Neuroscience, University of Chicago, Chicago, Illinois 60637, USA. 65 Department of Psychiatry and Behavioral Sciences, NorthShore University HealthSystem, Evanston, Illinois 60201, USA. ⁶⁶Department of Non-Communicable Disease Epidemiology, London School of Hygiene and Tropical Medicine, London WC1E 7HT, UK. ⁶⁷Department of Child and Adolescent Psychiatry, University Clinic of Psychiatry, Skopje 1000, Republic of Macedonia. ⁶⁸Department of Psychiatry, University of Regensburg, 93053 Regensburg, Germany. ⁶⁹Department of General Practice, Helsinki University Central Hospital, University of Helsinki P.O. Box 20, Tukholmankatu 8 B, Fl-00014, Helsinki, Finland ⁷⁰Folkha Isan Research Center, Helsinki, Finland, Biomedicum Helsinki 1, Haartmaninkatu 8, FI-00290, Helsinki, Finland. ⁷¹National Institute for Health and Welfare, P.O. Box 30, FI-00271 Helsinki, Finland. ⁷²Translational Technologies and Bioinformatics, Pharma Research and Early Development, F. Hoffman-La Roche, CH-4070 Basel, Switzerland. ⁷³Department of Psychiatry, Georgetown University School of Medicine, Washington DC 20057, USA. ⁷⁴Department of Psychiatry, Keck School of Medicine of the University of Southern California, Los Angeles, California 90033, USA. ⁷⁵Department of Psychiatry, Virginia Commonwealth University School of Medicine, Richmond, Virginia 23298, USA. ⁷⁶Mental Health Service Line, Washington VA Medical Center. Washington DC 20422, USA. ⁷⁷Department of Genetic Epidemiology in Psychiatry, Central Institute of Mental Health, Medical Faculty Mannheim, University of Heidelberg, Heidelberg, D-68159 Mannheim, Germany. ⁷⁸Department of Genetics, University of Groningen, University Medical Centre Groningen, 9700 RB Groningen, The Netherlands. ⁷⁹Department of Psychiatry, University of Colorado Denver, Aurora, Colorado 80045, USA. ⁸⁰Center for Neurobehavioral Genetics, Semel Institute for Neuroscience and Human Behavior, University of California, Los Angeles, California 90095, USA. 81 Department of Psychiatry, University of Halle, 06112 Halle, Germany. ⁸²Division of Psychiatric Genomics, Department of Psychiatry, Icahn School of Medicine at Mount Sinai, New York, New York, New York 10029, USA. ⁸³Department of Psychiatry, University of Munich, 80336, Munich, Germany. ⁸⁴Departments of Psychiatry and Human and Molecular Genetics, INSERM, Institut de Myologie, Ho[^] pital de la Pitie[^] -Salpe[^] trie[^] re, Paris 75013, France. ⁸⁵Mental Health Research Centre, Russian

Academy of Medical Sciences, 115522 Moscow, Russia. ⁸⁶Neuroscience Therapeutic Area, Janssen Research and Development, Raritan, New Jersey 08869, USA. ⁸⁷Queensland Brain Institute, The University of Queensland, Brisbane, Queensland, QLD 4072, Australia. ⁸⁸Academic Medical Centre University of Amsterdam, Department of Psychiatry, 1105 AZ Amsterdam, The Netherlands. ⁸⁹Illumina, La Jolla, California, California 92122, USA. ⁹⁰Institute of Biological Psychiatry, Mental Health Centre Sct. Hans, Mental Health Services Copenhagen, DK-4000, Denmark. ⁹¹Friedman Brain Institute, Icahn School of Medicine at Mount Sinai, New York, New York 10029, USA. 92J. J. Peters VA Medical Center, Bronx, New York, New York 10468, USA. ⁹³Priority Research Centre for Health Behaviour, University of Newcastle, Newcastle NSW 2308, Australia. ⁹⁴School of Electrical Engineering and Computer Science, University of Newcastle, Newcastle NSW 2308, Australia. ⁹⁵Division of Medical Genetics, Department of Biomedicine, University of Basel, Basel CH-4058, Switzerland. ⁹⁶Department of Genetics, Harvard Medical School, Boston, Massachusetts, Massachusetts 02115, USA. 97 Section of Neonatal Screening and Hormones, Department of Clinical Biochemistry, Immunology and Genetics, Statens Serum Institut, Copenhagen DK-2300, Denmark. ⁹⁸Department of Psychiatry, Fujita Health University School of Medicine, Toyoake, Aichi, 470-1192, Japan. ⁹⁹Regional Centre for Clinical Research in Psychosis, Department of Psychiatry, Stavanger University Hospital, 4011 Stavanger, Norway. ¹⁰⁰Rheumatology Research Group, Vall d'Hebron Research Institute, Barcelona 08035, Spain. ¹⁰¹Centre for Medical Research, The University of Western Australia, Perth WA6009, Australia. ¹⁰²The Perkins Institute for Medical Research, The University of Western Australia, Perth WA6009, Australia. ¹⁰³Department of Medical Genetics, Medical University, Sofia 1431, Bulgaria. ¹⁰⁴Department of Psychology, University of Colorado Boulder, Boulder, Colorado 80309, USA. ¹⁰⁵Campbell Family Mental Health Research Institute, Centre for Addiction and Mental Health, Toronto, Ontario M5T 1R8, Canada. ¹⁰⁶Department of Psychiatry, University of Toronto, Toronto, Ontario M5T 1R8, Canada. ¹⁰⁷Institute of Medical Science, University of Toronto, Toronto, Ontario M5S 1A8, Canada. ¹⁰⁸Institute of Molecular Genetics, Russian Academy of Sciences, Moscow 123182, Russia. 109 Latvian Biomedical Research and Study Centre, Riga, LV-1067, Latvia. ¹¹⁰Department of Psychiatry and Zilkha Neurogenetics Institute, Keck School of Medicine at University of Southern California, Los Angeles, California 90089, USA. ¹¹¹Faculty of Medicine, Vilnius University, LT-01513 Vilnius, Lithuania. ¹¹²Department of Biology and Medical Genetics, 2nd Faculty of Medicine and University Hospital Motol, 150 06 Prague, Czech Republic. ¹¹³Department of Child and Adolescent Psychiatry, Pierre and Marie Curie Faculty of Medicine, Paris 75013, France. 114 Duke-NUS Graduate Medical School, Singapore 169857. ¹¹⁵Department of Psychiatry, Hadassah-Hebrew University Medical Center, Jerusalem 91120, Israel. ¹¹⁶Centre for Genomic Sciences, The University of Hong Kong, Hong Kong, China. ¹¹⁷Mental Health Centre and Psychiatric Laboratory, West China Hospital, Sichuan University, Chengdu, 610041 Sichuan, China. ¹¹⁸Department of Biostatistics, Johns Hopkins University Bloomberg School of Public Health, Baltimore, Maryland 21205, USA. ¹¹⁹Department of Psychiatry, Columbia University, New York, New York 10032, USA. ¹²⁰Priority Centre for Translational Neuroscience and Mental Health, University of Newcastle, Newcastle NSW 2300, Australia. ¹²¹Department of Genetics and Pathology, International Hereditary Cancer Center, Pomeranian Medical University in Szczecin, 70-453 Szczecin, Poland. ¹²²Department of Mental Health and Substance Abuse Services; National Institute for Health and Welfare, P.O. BOX 30, FI-00271 Helsinki, Finland. ¹²³Department of Mental Health, Bloomberg School of Public Health, Johns Hopkins University, Baltimore, Maryland 21205, USA. ¹²⁴Department of Psychiatry, University of Bonn, D-53127 Bonn, Germany. ¹²⁵Centre National de la Recherche Scientifique, Laboratoire de Gé né tique Molé culaire de la Neurotransmission et des Processus Neurodé gé né ratifs, Ho^pital de la Pitié Salpe^trie`re, 75013 Paris, France. ¹²⁶Department of Genomics Mathematics, University of Bonn, D-53127 Bonn, Germany. ¹²⁷Research Unit, Sørlandet Hospital, 4604 Kristiansand, Norway. ¹²⁸Department of Psychiatry, Harvard Medical School, Boston, Massachusetts 02115, USA. 129VA Boston Health Care System, Brockton, Massachusetts 02301, USA. ¹³⁰Department of Psychiatry, National University of Ireland Galway, Co. Galway,

Ireland. ¹³¹Centre for Cognitive Ageing and Cognitive Epidemiology, University of Edinburgh, Edinburgh EH16 4SB. UK. ¹³²Division of Psychiatry, University of Edinburgh, Edinburgh, EH16 4SB, UK. ¹³³Division of Mental Health and Addiction, Oslo University Hospital, 0424 Oslo, Norway. ¹³⁴Massachusetts Mental Health Center Public Psychiatry Division of the Beth Israel Deaconess Medical Center, Boston, Massachusetts 02114, USA. ¹³⁵Estonian Genome Center, University of Tartu, Tartu 50090, Estonia. ¹³⁶School of Psychology, University of Newcastle, Newcastle NSW 2308, Australia. ¹³⁷First Psychiatric Clinic, Medical University, Sofia 1431, Bulgaria. ¹³⁸Department P, Aarhus University Hospital, DK-8240 Risskov, Denmark. ¹³⁹Department of Psychiatry, Royal College of Surgeons in Ireland, Dublin 2, Ireland. 140King's College London, London SE5 8AF, UK. ¹⁴¹Maastricht University Medical Centre, South Limburg Mental Health Research and Teaching Network, EURON, 6229 HX Maastricht, The Netherlands. ¹⁴²Institute of Translational Medicine, University of Liverpool, Liverpool L69 3BX, UK. ¹⁴³Max Planck Institute of Psychiatry, 80336 Munich, Germany. ¹⁴⁴Munich Cluster for Systems Neurology (SyNergy), 80336 Munich, Germany. ¹⁴⁵Department of Psychiatry and Psychotherapy. Jena University Hospital, 07743 Jena, Germany, ¹⁴⁶Department of Psychiatry, Queensland Brain Institute and Queensland Centre for Mental Health Research, University of Queensland, Brisbane, Queensland, St Lucia QLD 4072, Australia. ¹⁴⁷Department of Psychiatry and Behavioral Sciences, Johns Hopkins University School of Medicine, Baltimore, Maryland 21205, USA. ¹⁴⁸Department of Psychiatry, Trinity College Dublin, Dublin 2, Ireland. ¹⁴⁹Eli Lilly and Company, Lilly Corporate Center, Indianapolis, 46285 Indiana, USA. ¹⁵⁰Department of Clinical Sciences, Psychiatry, Umea[®] University, SE-901 87 Umea[®], Sweden. ¹⁵¹DETECT Early Intervention Service for Psychosis, Blackrock, Co. Dublin, Ireland. ¹⁵²Centre for Public Health, Institute of Clinical Sciences, Queen's University Belfast, Belfast BT12 6AB, UK. ¹⁵³Lawrence Berkeley National Laboratory, University of California at Berkeley, Berkeley, California 94720, USA. ¹⁵⁴Institute of Psychiatry, King's College London, London SE5 8AF, UK. ¹⁵⁵A list of authors and affiliations appear in the Supplementary Information. ¹⁵⁶Melbourne Neuropsychiatry Centre, University of Melbourne & Melbourne Health, Melbourne, Vic 3053, Australia. ¹⁵⁷Department of Psychiatry, University of Helsinki, P.O. Box 590, FI-00029 HUS, Helsinki, Finland. ¹⁵⁸Public Health Genomics Unit, National Institute for Health and Welfare, P.O. BOX 30, FI-00271 Helsinki, Finland ¹⁵⁹Medical Faculty, University of Belgrade, 11000 Belgrade, Serbia, ¹⁶⁰Department of Psychiatry, University of North Carolina, Chapel Hill, North Carolina 27599-7160, USA. ¹⁶¹Institute for Molecular Medicine Finland, FIMM, University of Helsinki, P.O. Box 20FI-00014, Helsinki, Finland ¹⁶²Department of Epidemiology, Harvard School of Public Health, Boston, Massachusetts 02115, USA. ¹⁶³Department of Psychiatry, University of Oxford, Oxford, OX3 7JX, UK. ¹⁶⁴Virginia Institute for Psychiatric and Behavioral Genetics, Virginia Commonwealth University, Richmond, Virginia 23298, USA. ¹⁶⁵Institute for Multiscale Biology, Icahn School of Medicine at Mount Sinai, New York, New York 10029, USA. 166 PharmaTherapeutics Clinical Research, Pfizer Worldwide Research and Development, Cambridge, Massachusetts 02139, USA. ¹⁶⁷Department of Psychiatry and Psychotherapy, University of Gottingen, 37073 Gottingen, Germany. ¹⁶⁸Psychiatry and Psychotherapy Clinic, University of Erlangen, 91054 Erlangen, Germany. ¹⁶⁹Hunter New England Health Service, Newcastle NSW 2308, Australia. ¹⁷⁰School of Biomedical Sciences, University of Newcastle, Newcastle NSW 2308, Australia. ¹⁷¹Division of Cancer Epidemiology and Genetics, National Cancer Institute, Bethesda, Maryland 20892, USA. ¹⁷²University of Iceland, Landspitali, National University Hospital, 101 Reykjavik, Iceland. ¹⁷³Department of Psychiatry and Drug Addiction, Tbilisi State Medical University (TSMU), N33, 0177 Tbilisi, Georgia. ¹⁷⁴Research and Development, Bronx Veterans Affairs Medical Center, New York, New York 10468, USA. ¹⁷⁵Wellcome Trust Centre for Human Genetics, Oxford OX3 7BN, UK. ¹⁷⁶deCODE Genetics, 101 Reykjavik, Iceland. ¹⁷⁷Department of Clinical Neurology, Medical University of Vienna, 1090 Wien, Austria. ¹⁷⁸Lieber Institute for Brain Development, Baltimore, Maryland 21205, USA. ¹⁷⁹Department of Medical Genetics, University Medical Centre Utrecht, Universiteitsweg 100, 3584 CG, Utrecht, The Netherlands. ¹⁸⁰Berkshire Healthcare NHS Foundation Trust, Bracknell RG12 1BQ, UK. ¹⁸¹Section of Psychiatry, University of Verona, 37134 Verona, Italy. ¹⁸²Department of Psychiatry, University of Oulu, P.O. Box 5000, 90014,

Finland. ¹⁸³University Hospital of Oulu, P.O. Box 20, 90029 OYS, Finland. ¹⁸⁴Molecular and Cellular Therapeutics, Royal College of Surgeons in Ireland, Dublin 2, Ireland, ¹⁸⁵Health Research Board, Dublin 2, Ireland. ¹⁸⁶ School of Psychiatry and Clinical Neurosciences, The University of Western Australia, Perth WA6009, Australia. ¹⁸⁷Computational Sciences CoE, Pfizer Worldwide Research and Development, Cambridge, Massachusetts, 02139, USA. ¹⁸⁸Human Genetics, Genome Institute of Singapore, A*STAR, Singapore 138672. ¹⁹⁰University College London, London WC1E 6BT, UK. ¹⁹¹Department of Neuroscience, Icahn School of Medicine at Mount Sinai, New York, New York 10029, USA. ¹⁹²Institute of Neuroscience and Medicine (INM-1), Research Center Juelich, 52428 Juelich, Germany. ¹⁹³Department of Genetics, The Hebrew University of Jerusalem, 91905 Jerusalem, Israel. ¹⁹⁴Neuroscience Discovery and Translational Area, Pharma Research and Early Development, F. Hoffman-La Roche, CH-4070 Basel, Switzerland, ¹⁹⁵Centre for Clinical Research in Neuropsychiatry, School of Psychiatry and Clinical Neurosciences, The University of Western Australia, Medical Research Foundation Building, Perth WA6000, Australia. ¹⁹⁶Virginia Institute for Psychiatric and Behavioral Genetics, Departments of Psychiatry and Human and Molecular Genetics, Virginia Commonwealth University, Richmond, Virginia 23298, USA. ¹⁹⁷The Feinstein Institute for Medical Research, Manhasset, New York 11030, USA. ¹⁹⁸The Hofstra NS-LIJ School of Medicine, Hempstead, New York 11549, USA. ¹⁹⁹The Zucker Hillside Hospital, Glen Oaks, New York 11004, USA, 200 Saw Swee Hock School of Public Health, National University of Singapore, Singapore 117597, Singapore. ²⁰¹Queensland Centre for Mental Health Research, University of Queensland, Brisbane 4076, Queensland, Australia. ²⁰²Center for Human Genetic Research and Department of Psychiatry, Massachusetts General Hospital, Boston, Massachusetts 02114, USA, ²⁰³Department of Child and Adolescent Psychiatry, Erasmus University Medical Centre, Rotterdam 3000, The Netherlands. ²⁰⁴Department of Complex Trait Genetics, Neuroscience Campus Amsterdam, VU University Medical Center Amsterdam, Amsterdam 1081, The Netherlands. ²⁰⁵Department of Functional Genomics. Center for Neurogenomics and Cognitive Research, Neuroscience Campus Amsterdam, VU University, Amsterdam 1081, The Netherlands. ²⁰⁶University of Aberdeen, Institute of Medical Sciences, Aberdeen AB25 2ZD, UK. ²⁰⁷Departments of Psychiatry, Neurology, Neuroscience and Institute of Genetic Medicine, Johns Hopkins School of Medicine, Baltimore, Maryland 21205, USA. ²⁰⁸Department of Clinical Medicine, University of Copenhagen, Copenhagen 2200, Denmark.